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# Selección de cepas de *Rhizobium leguminosarum* por variables simbióticas y productivas en *Pisum sativum* L.

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## ABSTRACT

Field pea (*Pisum sativum* L.) is a winter symbiotic legume that associates with *Rhizobium leguminosarum* sv viciae. This work aimed to evaluate strains of *R. leguminosarum* for their infective ability and early-plant growth, BNF contribution, biomass and grain yield. Seventy-eight specific strains and four pea cultivars were evaluated in a growth chamber, five strains and three cultivars were evaluated in a greenhouse, and three strains and two cultivars were evaluated in a field experiment. Only 44-55% of all evaluated strains were infective in the four cultivars. In the greenhouse, D70 and D156 strains showed the best nodulation variables as well as higher N content and yield. The field experiment showed D156 and D70 yielded a similar behavior for N content in canopy biomass and individual nodule biomass, whereas D191 had a higher nodule number per plant, aerial biomass and grain yields. D70 provided good nodulation, N content in biomass, and yield in the growth chamber, greenhouse, and field experiments. Therefore, D156 could constitute a good candidate for bacterial single-strain inoculants, as well as for formulating microbial consortia.

## **Keywords**

Field pea • Rhizobium leguminosarum • strain selection • symbiotic efficiency

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## RESUMEN

La arveja (*Pisum sativum L.*) es una leguminosa simbiótica del invierno que establece asociación con *Rhizobium leguminosarum* sv viciae. Los objetivos de trabajo fueron evaluar cepas de *R. leguminosarum* por infectividad y crecimiento temprano, aporte de BNF, producción de biomasa y rendimiento. Setenta y ocho cepas y cuatro cultivares se evaluaron en cámara de crecimiento, cinco cepas y tres cultivares en invernadero, mientras que tres cepas y dos cultivares en experimento de campo. Solo el 44-55% de las cepas fueron infectivas en los cultivares. En invernadero, las cepas D70 y D156 mostraron los mejores valores de nodulación y los mayores contenidos de N en biomasa y rendimiento. En campo, D156 y D70 mostraron comportamientos similares que en invernadero, mientras que D191 tuvo más nódulos por planta, mayor biomasa aérea y mayor rendimiento. La cepa D70 mantuvo su comportamiento de nodulación, contenido de N en biomasa y rendimiento en cámara de crecimiento, invernadero y campo; mientras que D156 tuvo igual o mejor comportamiento que D70 en invernadero y campo. Por lo tanto, D156 podría ser una buena candidata para formular inoculantes bacterianos con esta cepa o para formular consorcios microbianos.

### **Palabras clave**

Arveja • Rhizobium leguminosarum • selección de cepas • eficiencia simbiótica

## INTRODUCTION

Field pea (*Pisum sativum* L.) is a winter symbiotic legume able to establish a specific mutualistic association with *Rhizobium leguminosarum* symbiotic variant (sv) *viciae*. This association contributes to satisfying part of the crop nitrogen (N) demand via biological nitrogen fixation (BNF). From an agronomic perspective, field peas may be included as an alternative winter crop in agricultural rotations, with a shorter cycle duration and lower water consumption than other winter crops like wheat or barley, and the N-fixing ability (19). The N-fixing efficiency of the legume-rhizobium relationship depends mainly on crop genotype (1, 2), rhizobium strain (23), and soil-water interactions associated with management practices (10, 13, 14). Contributions of BNF in field pea range between 45 and 286 kg N ha<sup>-1</sup> (1, 13, 16, 22, 31, 36), representing 33-91% of total N uptake (1, 13, 16, 21, 22, 26, 31, 36).

*Rhizobium leguminosarum* strains have three distinct symbionts: (i) *R. leguminosarum* sv *phaseoli*, which nodulates beans (*Phaseolus vulgaris*); (ii) *R. leguminosarum sv trifolii*, which nodulates clovers (*Trifolium* sp); and (iii) *R. leguminosarum sv viciae*, which nodulates field pea, vetch (*Vicia* sp), lentil (*Lens* sp), faba beans (*Vicia faba*) and Lathyrus (*Lathyrus* sp) (7, 8, 27). In field peas, statistical interactions between strains and crop genotypes have been observed for traits associated with BNF, plant growth and photosynthesis (11). Rhizobia strains have greater relevance on BNF and N content, biomass and grains, than plant genotype (23). This strain-dependent behavior is associated with the expression of genes related to *R. leguminosarum* sv viciae nodulation, ensuring nodule establishment (7).

In Argentina, although field pea is not a native plant, native soil N-fixing rhizobia can nodulate this crop. Currently, the *R. leguminosarum* native strain D70 is the elite strain inoculating field peas, vetches, and lentils. This strain can nodulate all commercial cultivars and has shown greater BNF contribution in all agro-ecological environments evaluated in Argentina (13). However, even showing good behavior regarding grain yield and biomass production, it does not necessarily respond equally in all agricultural environments. Newly incorporated commercial cultivars still require evaluation regarding symbiotic behavior. In this regard, strains of the collection of the Instituto de Microbiología y Zoología Agrícola (IMyZA) of the Instituto Nacional de Tecnología Agropecuaria (INTA) could potentially perform better than D70 and be used in new inoculant formulations. Thus, this work aimed to i) evaluate *Rhizobium* sp. soil isolates and strains from the IMyZA-INTA collection for infective ability and early growth promotion under controlled conditions (growth chamber), ii) evaluate symbiotic efficiency of strains selected from the growth chamber, under greenhouse conditions, and iii) evaluate inoculation effects of the greenhouse selected strains on BNF contribution, nodulation, and production variables under field conditions.

## **MATERIALS AND METHODS**

## Growth chamber experiments

Seventy-one to seventy-eight specific strains and/or soil isolates from the IMyZA-INTA collection (Additonal material 1) were combined with four field pea cultivars (Facón, Bicentenario, Manantiales, and Pampa). Rhizobium sp. inocula were grown in glass tubes (20 mm x 195 mm) with 10 ml of yeast extract mannitol (YEM) broth (35) on an orbital shaker (180 rpm, 48 h at 28°C). Each inoculum contained at least 3 x  $10^8$ colony-forming units per milliliter (CFU ml<sup>-1</sup>). Field pea seeds were disinfected superficially with 80% ethanol for 1 min, 4% sodium hypochlorite for 2 min, at 80% ethanol for 0.5 min, and rinsed 5 times with sterile distilled water. Then, these seeds were placed in plastic trays with moistened absorbent paper for 72 h in an incubation stove at 28°C for germination. Two germinated seeds were inoculated with 1 ml of inoculant and planted in a 220-ml pot with sterile vermiculite substrate and daily irrigated with N-free nutrient solution (17). Each pot was kept at a constant temperature of 25°C and 16 h of light. Each strain and cultivar combination was considered a treatment, and a non-inoculated control was added for each field pea cultivar. The experimental design was set as completely randomized with three replications. Finally, the number of nodules per plant ( $N^{\circ}_{Nod}$ ) and biomass per plant (Biom<sub>Total</sub>) were evaluated 25 days after planting (DAP).

## **Greenhouse experiment**

Five strains selected from the growth chamber experiment (D70, D156, D191, D192, and D193) were evaluated for nodulation and early growth promotion abilities in the greenhouse. The experiment was planted in 2 L pots on August 21st, 2013, using four seeds per pot and sterile vermiculite as substrate. Seeds were disinfected as described above. Treatments consisted of a factorial combination of the five selected strains plus a non-inoculated control, with three commercial field pea cultivars (Viper, Facón, and Bicentenario) arranged in a complete randomized block design totaling 17 replicates. The inoculant was peat-based with a count of at least  $1 \ge 10^{8}$  CFU g<sup>-1</sup> of product. Sugared water (20% w/v) was used as adherent at a rate of 10 ml kg seed<sup>-1</sup>. The inoculant was applied to seeds, mixed to homogenize, and allowed to stand for 30 min before planting. Pots were irrigated three days a week with an N-free nutrient solution (17). Variables per plant were evaluated at 56 DAP and 78, 92, and 103 DAP per pot. Root biomass (Biom<sub>Root</sub>), aerial biomass ( $Biom_{Aerial}$ ), plant biomass ( $Biom_{Plant}$ ), number of nodules ( $N^{\circ}_{Nod}$ ), dry nodule biomass (Biom<sub>Nod</sub>) and dry biomass per nodule (Biom<sub>Nod-I</sub>) were evaluated at two sampling moments (at 56 and 78 DAP), with additional measurements such as root and aerial length (Leng<sub>Root</sub> and Leng<sub>Stem</sub>, respectively) at 56 DAP. Then, the following variables were also evaluated: vegetative, grain, and aerial biomass (Biom<sub>veg</sub>, Biom<sub>Grain</sub>, and Biom<sub>Aerial</sub>, respectively), and plant yield components as pod numbers, number of grains, number of grains per pod, and individual grain biomass  $(N^{\circ}_{_{Pod}}, N^{\circ}_{_{Grain}}, N^{\circ}_{_{Grain-pod}}, and Biom_{_{Grain-1'}}$  respectively) at harvest (ca. 103 DAP). Additionally, N content  $(\%N_{_{upt}})$  and isotopic composition ( $\delta^{15}N_{unt}$ ) on whole plant biomass were determined at 92 DAP to estimate the  $\beta$ factor for each strain and field pea cultivar combination according to Boddey *et al.* (2000).

## **Field experiment**

Strains D70, D156, and D191, first evaluated in the greenhouse, were then compared in a field experiment. All field treatments (*i.e.* the combinations between these three strains and two pea cultivars, Facón and Viper) were carried out in triplicate. The field experiment was performed in the locality of Rueda (33°21'48.48" S and 60°27'39.12" W), in Santa Fe province (Argentina), on a Typic Argiudoll soil. The experimental unit (plot) had six furrows (at 0.175 m interrow) and 5 m in length. The experiment was sown on July 29<sup>th</sup>, 2014, with 119 plants m<sup>-2</sup> (110-127 plants m<sup>-2</sup>) and harvested on November 14<sup>th</sup> (108 DAP). Plants were fertilized with 100 kg ha<sup>-1</sup> of monoammonium phosphate (*i.e.* 20 kg P ha<sup>-1</sup>). Seeds were inoculated as described in the greenhouse experiment. Weeds and pests were controlled. During the cycle, total rainfall was 284 mm and mean temperature was 17.3°C.

At 60 DAP (September 14<sup>th</sup>), 0.5 m of two rows (0.18 m<sup>-2</sup>) was sampled and the number of nodules in each plant ( $N^{\circ}_{Nod}$ ) was counted. Then, plants were oven-dried at 65°C for 72 h

to determine nodule biomass per plant. At 95 DAP, 1 m<sup>2</sup> of the total aboveground biomass of field pea of each plot (N-fixing crop) plus one linear meter of wheat (non-fixing crop) were sampled. These samples were dried in an air circulation stove at 65°C for 72 h. A subsample of each mentioned sample was ground with a Wiley mill to determine N content in biomass ( $(N_{upt})$ ) according to the micro-Kjeldahl method. The N uptake ( $N_{upt}$ ) was obtained by multiplying  $(N_{upt})$  by Biom<sub>Aerial</sub>. BNF was determined by the <sup>15</sup>N natural abundance method, using an elemental analyzer Carlo Erba EA 1108 coupled to a ThermoScientific Delta V Advantage isotopic mass spectrometer of continuous flow through a ConFLo IV interface. Then,  $(N_{FBN} = (N_{BNF} * N_{upt}) = (N_{BNF} * N_{upt})$ 

At harvest (108 DAP), two samples of aboveground biomass (Biom<sub>Aerial</sub>) were taken from each plot of 1 m<sup>2</sup>. The, grained (Biom<sub>Grain</sub>) and vegetative structures (Biom<sub>Veg</sub>) were separated, dried in an air circulation stove at 65°C for 72 h and weighed. The harvest index was calculated. Grain yield (Yield<sub>Grain</sub>) was adjusted to grain moisture of 0.135 kg water kg grain<sup>-1</sup>.

## Molecular characterization

Three of the strains evaluated (D70, D156 and D191) were characterized by partial sequencing of the 16S rRNA gene by amplification reactions. Pure colonies were grown on plates in YEM culture medium, then suspended in 50 µl of ultrapure water and boiled in a water bath for 10 min to obtain DNA extracts. The universal primers fD1 and rP2 proposed by Weisburg *et al.* (1991) were used and amplification was carried out in a volume of 25 µl of the reaction containing 2.5 ml 1X Buffer, 0.75 µl of 50 mM MgCl<sub>2</sub>, 0.5 µl of 10 mM dNTP, 0.25 µl Taq DNA Polymerase, 0.5 µl of each primer (fD1 and rP2), 1 µl of tempered genomic DNA and 19 µl of ultrapure water. Amplification conditions consisted of an initial denaturation of 2 min at 94°C, followed by 35 cycles of denaturation (94°C, 40 seconds), annealing (52°C, 40 seconds) and extension (72°C, 1.5 min) and a final extension at 72°C for 10 min. A negative control without template DNA was included. Electrophoresis was performed at 90 V. The amplified products were analyzed by 0.8% agarose electrophoresis stained with SYBR Safe DNA gel stain (Invitrogen<sup>TM</sup>) for 30 min and then purified with the commercial Gel Extraction kit QIAEX II (Qiagen). Partial sequences of 16S rRNA were compared with those deposited in the NCBI BLAST database (http://blast.ncbi.nlm.nih.gov/).

## Data analysis

Statistical analysis of the growth chamber, greenhouse, and field experiments was carried out using a two-way ANOVA. Means were compared with the DGC test ( $p \le 0.05$ ). All analysis was performed using Infostat software version 2018p (Di Rienzo *et al.*, 2018).

## RESULTS

## Growth chamber experiment

Only 44-55% of all the strains tested (71-78) presented nodules in the four cultivars (*i.e.* they were infective), varying according to the cultivar. Facón and Pampa cultivars had the highest number of infectious strains (41), whereas Bicentenario and Manantiales had the lowest (32 and 31, respectively).

Twenty-three strains were able to infect the four cultivars evaluated. The isolates obtained from nodules of the genus *Lathyrus* had a higher proportion of infective and effective nodules in the four cultivars evaluated (60%), followed by those from *Lens*, with 50%, and those from *Vicia*, with 29%, whereas the isolates from *Pisum*, with 8%. Soil isolates had 38% nodulation ability in the four cultivars.

Results from the growth chamber showed highly significant (p<0.001) variability in the N°<sub>Nod</sub> per plant (*i.e.* strain x cultivar), whereas the effects of strains on Biom<sub>Total</sub> were significant at p=0.025 (Additional material 2). N°<sub>Nod</sub> ranged from 76 to 1. D70 presented the highest N°<sub>Nod</sub> values, with an average of 43 nod pl<sup>-1</sup>, followed by D191, with 35 nod pl<sup>-1</sup>, and higher N°<sub>Nod</sub> than the rest. These responses were 97% and 67% higher, on average, at D70 and D191, respectively (Additional material 1). Biom<sub>Total</sub> ranged from 287 to 149 mg pl<sup>-1</sup>, with D156, D155, and D191 showing the highest values (233-221 mg pl<sup>-1</sup>), *i.e.* 6% higher than the Biom<sub>Total</sub> of D70 with the four cultivars (Additional material 1).

Based on these results, greenhouse experiments included strain D156 selected for biomass production ability, D191 for nodulation behavior, and D192 and D193 for fast initial growth (not shown) as compared to the reference strain D70.

## **Greenhouse experiment**

Strains D70, D156, D191, D192, and D193 were evaluated in the greenhouse experiment. At 56 DAP, a highly significant interaction between strain and cultivar was detected in all variables, except root biomass. Strain D192 had the best behavior in Leng<sub>Root</sub> (358 mm pl<sup>-1</sup>) followed by D156 and D193 (322 mm pl<sup>-1</sup>, on average) with cultivar Facón, with 28 and 15% differences in comparison to D70 and D191. D192 and D193 showed better Leng<sub>Aerial</sub> values, with 235 mm pl<sup>-1</sup>, followed by D156 and D70 (206 mm pl<sup>-1</sup>), all with cultivar Bicentenario, being 66% for 192 and D193 and 46% for D156 and D70. Instead, strain D193 had higher Leng<sub>Plant</sub> (515 mm pl<sup>-1</sup>), followed by D192 (508 mm pl<sup>-1</sup>) with Bicentenario and Facón. Likewise, strains D192 and D193 had higher Biom<sub>Plant</sub> (473 mg pl<sup>-1</sup>) with Bicentenario, with significant differences with the rest of 55%. These strains also had the highest Biom<sub>Aerial</sub> values (271 mg pl<sup>-1</sup>), followed by D156 and D70 (214 mg pl<sup>-1</sup>), all with Bicentenario. In contrast, regarding nodule variables, D70 and D156 showed higher N°<sub>Nod</sub> (32 nod pl<sup>-1</sup> on average) with Viper, 100% higher than the rest, as well as higher  $Biom_{Nod}$  (9 mg pl<sup>-1</sup> on average), 141% higher than the rest, and highest Biom<sub>Nod-I</sub> (0.44 mg nod<sup>-1</sup> in average), 144% higher than the rest, all with Bicentenario.

At 78 DAP, a significant interaction between strain and cultivar was observed in the nodulation variables, except in individual nodule biomass. Strain D70 had the highest N°<sub>Nod</sub>, together with strains D192 and D193, which were better associated with Bicentenario (109, 129, and 110 nod pot<sup>-1</sup>, respectively). Also, strain D70 showed better Biom<sub>Nod</sub> with Viper (40 mg pot<sup>-1</sup>) and Bicentenario (38 mg pot<sup>-1</sup>), and the best Biom<sub>Nod-I</sub> (0.53 mg nod<sup>-1</sup>) concerning other strains (75%). Differently, plant growth variables showed significant differences among strains. Strain D191 showed the lowest values of Biom<sub>Aerial</sub> (2.0 g pot<sup>-1</sup>) and Biom<sub>Root</sub> (0.7 g pot<sup>-1</sup>) with respect to the rest (40 and 39%, respectively).

At harvest (103 DAP), the five strains were evaluated for Biom<sub>Veg</sub>, Biom<sub>Grain</sub>, and Biom<sub>Aerial</sub> Only Biom<sub>vee</sub>, showed cultivar and strain interaction. All plant yield components (*i.e.* N° N° Grain-pod and Biom Grain showed significant interaction between both factors, although only measured in Facón and Viper with strains D70, D156, and D191. Strain D70 produced  $3.9 \text{ g pot}^{-1}$  of Biom<sub>veg</sub> with Viper cultivar, significantly higher than the rest (45% on average), followed by D156 (3.5 g pot<sup>-1</sup>) also with Viper cultivar. In contrast, strain D191 showed the worst performance (1.6 g pot<sup>1</sup>) with Facón (figure 1 A, page 6). Regarding Biom<sub>Grain</sub>, D156, D70 and D193 had higher values (1.7, 1.6, and 1.5 g pot<sup>-1</sup>, respectively) than D191 and D192 strains (figure 1B, page 6). Regarding Biom<sub>Aerial</sub>, D191 (3.6 g pot<sup>-1</sup>) showed significantly lower values (22%) than other strains (figure 1C, page 6). Considering yield components, strain D156, followed by D70, threw the best results compared to D191, except in Biom<sub>Grain-I</sub> where D191 had the second-best weight after D156. In addition, D156 stran had 6.5 pods pot<sup>-1</sup>, followed by D70 strain (5.8 pods pot<sup>-1</sup>), both with Facón cultivar having more pod number per pot ( $N^{\circ}_{_{Pod}}$ ) than the rest (figure 1D, page 6). Regarding grain number per pod (N°<sub>Grain-pod</sub>), D156 also performed better with Facón (4.0 grains pod<sup>-1</sup>) and D70 with Viper (3.8 grains pod<sup>-1</sup>) (figure 1E, page 6). Likewise, D156 followed by D191, had the highest individual grain biomasses (Biom<sub>Grain-I</sub>) with 136 and 132 mg grain<sup>-1</sup> respectively, with Facón (figure 1F, page 6). D156 with Facón had higher N°<sub>Grain</sub> (18.8 grains pot<sup>-1</sup>) than the rest (44%) (figure 1G, page 6).

Total N ( $N_{upt}$ ) and  $\delta_{15}N$  ( $M_{00}$ ) contents of whole plants were determined at 92 DAP for Facón and Viper inoculated with D70, D156, and D191 (Additional material 3). D156 (2.2%) and D70 (2.1%) strains had higher biomass  $_{unt}$ %Nupt than D191 (p<0.01). The mean  $\beta$  value was -0.66‰, where D70 had lower depletion of  $\delta_{15}N$  (-0.45‰), without differences with D156 (-0.55‰), being both over D191 (-0.88‰) with

a significance level of p=0.06 (Additional material 3).



A) vegetative weight (g), B) grain weight (g), C) aerial biomass (g), D) pod number, E) grain per pod, F) individual grain biomass (mg) andG) grain number per pod.

Different letters indicate significant differences between treatments at p≤0.05. Results shown in panel D correspond to cultivars Facón and Viper average values.

A) biomasa de grano (g), B) biomasa vegetativa (g) y C) biomasa aérea total (g), D) número de vainas por maceta, E) número de granos por vaina, F) biomasa individual del grano (mg) y G) número de granos por maceta.

Letras diferentes indican diferencias significativas entre tratamientos con valor p≤0,05. Los resultados mostrados en el panel D corresponden al promedio de los cultivares Facón y Viper.

Figure 1. Plant variables and yield components per pot at harvest (103 days after planting) as dry matter.
 Figura 1. Variables de la planta y componentes del rendimiento por maceta a la cosecha (103 días después de la siembra).

## **Field experiment**

In the field experiment, only nodule number had significant interaction between factors. Regarding strain behavior significant differences were found for  $\text{Biom}_{\text{Veg'}}$ ,  $\text{Biom}_{\text{Aerial'}}$  and  $\text{Biom}_{\text{Nod-I'}}$ .

Strain D191 behaved differently than in the greenhouse regarding  $\operatorname{Biom}_{\operatorname{Veg}}$ , Yield<sub>Grain</sub>,  $\operatorname{Biom}_{\operatorname{Aerial}}$  and  $\operatorname{N}_{upt}$ . Likewise, strain D191 strain had better behavior in  $\operatorname{Biom}_{\operatorname{Veg}}$  (3.3 Mg ha<sup>-1</sup>) and  $\operatorname{Biom}_{\operatorname{Aerial}}$  (6.7 Mg ha<sup>-1</sup>) than D156 (15% in both variables) and D70 (33 and 25%, respectively, table 1, page 7). In addition, D191 strain had higher N content in aerial biomass (212 kg N ha<sup>-1</sup>) than D156 (8%) and D70 (16%). Average Yield<sub>Grain</sub> was 3.5 Mg ha<sup>-1</sup>, %N<sub>FBN</sub> was 70.5% and N°<sub>FBN</sub> was 139 kg N ha<sup>-1</sup> (table 1, page 7). Instead, regarding %N<sub>upt</sub>, D156 (3.43%) and D70 (3.41%) showed higher values than D191 (8 and 7%, respectively), maintaining the same behavior as in the greenhouse (table 1, page 7). Strain D191 had the highest N°<sub>Nod</sub> per plant (10.5 nod pl<sup>-1</sup>) differing 105% with the rest. Instead, D156 and D70 had highly significant differences (p≤0.01) in  $\operatorname{Biom}_{\operatorname{Nod-I}}$  (2.5 and 1.7 mg nod<sup>-1</sup>, respectively) with respect to D191 (162 and 85%, respectively), but with no differences in  $\operatorname{Biom}_{\operatorname{Nod}}$  (table 1, page 7).

**Table 1.** Nodulation at 60 DAP, nitrogen plant variables at 95 DAP and plant harvest at108 DAP.

	Plant	Nodulation	n (60 DAP)		
Strain / Variable	Biom <sub>veg</sub> Mg ha <sup>-1</sup>	Yield <sub>Grain</sub> * Mg ha <sup>.1</sup>	Biom <sub>Aerial</sub> Mg ha <sup>-1</sup>	Biom <sub>Nod</sub> mg	Biom <sub>Nod-I</sub> mg
D70	2.5 b	3.3 a	5.3 b	97.2 a	1.74 a
D156	2.9 b	3.3 a	5.8 b	104.3 a	2.46 a
D191	3.3 a	3.8 a	6.7 a	67.3 a	0.94 b
p-value	0.006	0.09	0.004	0.56	0.008
	Nitrogen pla	nt variables at	harvest (108 DA	AP)	
Strain / Variable	N <sub>upt</sub> kg ha <sup>.1</sup>	%N <sub>fbn</sub> %	N <sub>FBN</sub> kg ha <sup>-1</sup>	%N <sub>above</sub> %	
D70	182 b	72 a	132 a	3.43 a	
D156	196 b	73 a	143 a	3.41 a	
D191	212 a	67 a	142 a	3.18 b	
p-value	0.03	0.25	0.51	0.02	

**Tabla 1.** Nodulación a los 60 DDP, de las variables de nitrógeno en planta (95 DDS) yrendimiento de granos a cosecha (108 DDS).

\*Adjusted to 0.135 kg  $H_2O$  kg grain<sup>-1</sup> grain moisture. Values followed by different letters in columns indicate significant differences at  $p \le 0.05$ . \*Ajustado a una humedad de grano de 0,135 kg  $H_2O$  kg grano<sup>-1</sup>. Los valores seguidos de letras diferentes en la columna indican diferencias significativas en  $p \le 0,05$ .

## Molecular characterization

Molecular identification of D70, D156, and D191 strains via partial sequencing of the 16S rRNA gene showed 100% identity with *Rhizobium leguminosarum*, and deposited in the same database under accession numbers KU933357, KX 066064, and KX346599, respectively. Sequences were then processed to obtain multiple alignments then concatenated for the construction of a phylogenetic tree. In turn, strains Az39 (*Azospirillum argentinense*) and E109 (*Bradyrhizobium japonicum*) were incorporated as members of other groups. Results of this analysis showed that D70 and D156 are more closely related than D191.

## DISCUSSION

Our results showed that nodulation ability of both native and naturalized isolates of rhizobia in field pea was similar to that observed by other authors (3, 8, 23, 38). For example, Ballard et al. (2004) found that 67% (22 of 33) of naturalized populations of rhizobia (applied to pea plants as soil suspensions) caused early and abundant nodulation in one cultivar of field pea, while our work found that 44-55% of native isolates of Argentina or introduced strains kept at the IMYZA-INTA collection were infective in four field pea cultivars. Likewise, Boivin et al. (2020b) observed that Rhizobium leguminosarum sy viciae are all potential symbionts of Fabaceae hosts (field pea, vetch, and lens) but display variable competitiveness to form root nodules. However, when Fabaceae legumes are exposed to natural soil bacterial populations, symbiotic efficiency is often suboptimal. In a recent study, Boivin et al. (2020a) determined that small genetic differences in nodD genes allowed observing a higher specificity degree among R. leguminosarun sv viciae in pea, faba bean, and lentil. In the present study, we observed that rhizobia nodule isolates from Lathyrus sp and *Lens* sp were the most infective, followed by *Vicia* sp. Instead, isolates from pea plants did not ensure infection in all cultivars of the same species (*Pisum sativum*), since only 1 of 12 isolates/strains caused nodules in the four cultivars evaluated. Therefore, field pea

cultivars could have genetic differences (2, 36), which, associated with the rhizobial genome, may influence competitiveness for nodulation as well as rhizosphere colonization, decisive steps in the competition for nodule occupancy by *R. leguminosarum* in soil populations (23).

The elite strain D70, isolated from *Vicia* sp, is most used in inoculant formulations. This strain has high nodulation ability and BNF contribution. Enrico *et al.* (2020) determined an average of 134 kg ha<sup>-1</sup> of N<sub>FBN</sub> in field pea (8 experiments, average). In this work, D70 had a contribution of 131 kg N ha<sup>-1</sup>, while D156, also isolated from *Vicia* sp, showed a similar and better BNF behavior in both greenhouse and field experiments (*i.e.* 143 kg N ha<sup>-1</sup>). Instead, the high nodulation ability observed in D191 (obtained from *Lens* sp) in the growth chamber, then resulted the least effective regarding yield variables in the greenhouse experiment. However, this strain performed better in the field experiment regarding biomass production (6.7 Mg ha<sup>-1</sup>), grain yield (3.8 Mg ha<sup>-1</sup>) and N absorbed (212 kg ha<sup>-1</sup>), considering the other strains. Table 2 shows BNF contributions from different studies, indicating the selected strains had high N contribution. Likewise, D156 and D70 had better N content in biomass ( $^{\circ}N_{upt}$ ), which, according to Laguerre *et al.* (2007), depends on BNF efficiency of rhizobia derived from a robust nodular system.

BNF contribution (%)	N derived from BNF kg N ha <sup>-1</sup>	References
72	131	D70 strain (this work)
73	143	D156 strain (this work)
67	142	D191 strain (this work)
69-71	166-286	Kumar and Goh 2000
84	238	Voisin <i>et al.,</i> 2002
47-81	64-81	Hauggaard-Nielsen et al., 2009
87-91	64-130	Abi-Ghanem <i>et al.</i> , 2011
37-85	45-106	Ruisi <i>et al.,</i> 2012
41-45	-	Keneni <i>et al.,</i> 2013
43-62	58-69	Hossain et al., 2016
35-65	-	Yang <i>et al.</i> , 2017
55	63	Liu <i>et al.,</i> 2019
33-83	11-71	Enrico <i>et al.,</i> 2020

Table 2. Contribution of biological nitrogen fixation (BNF) on field pea.Tabla 2. Contribución de la fijación biológica de nitrógeno (FBN) en arveja sobre<br/>experimentos de campo.

In the present study, strains D70 and D191 had high nodulation ability in controlled conditions (growth chamber and greenhouse). In contrast, both D156 and D70 had higher individual nodule biomass than D191, similar to controlled conditions. This behavior may be due to differences observed by the 16S genes concerning D70 or the larger number of specific genes that control nodulation competitiveness (8). This behavior can also be due to plant energetic regulation in association since, according to Rainbird *et al.* (1984), sustaining nodular system involves 22% of total energy, and nitrogenase activity (BNF) consumes an additional 52% (in soybean). Likewise, Ryle *et al.* (1986) determined that the energy consumption of white clover nodules was 22-27%. Therefore, nodule numbers adapt to plant possibility to maintain a symbiotic system benefiting yield.

Early growth effects in the growth chamber (D156) and greenhouse experiment (D156, D192, and D193 strains) could be associated with strain-dependant phytohormone effects, a fact not addressed in this study. Therefore, strains tending to increase auxins would be associated with stem and rootelongation during post-emerging growth (6, 27, 33). In addition, cytokine balance in the early nodulation stages may be associated with a larger number of nodules. Also, final number of nodules may depend on ethylene production (4, 15). Strains D192, D193 and D156 stood out in stem and root elongation and nodule number and

could have an adequate profile of phytohormones helping plant implantation. Then, at an advanced stage (78 DAP), the most relevant aspect would be symbiotic functionality, where the contribution of biological N translates into greater biomass (11, 34). Both strains D156 and D70 showed differences with D191, possibly due to symbiotic effectiveness. Differences were also observed in grain yield when the only N source was biological. Likewise, the only source of <sup>15</sup>N abundance in biomass resulted to be the biological source ( $\beta$  value), adjusting the contribution of BNF in field conditions. These results are consistent with those obtained by Unkovich *et al.* (1994) in clovers under different environments and with those by López-Bellido *et al.* (2010) in chickpea and faba beans between selected and natural soil strains. For non-different nodule structures (*i.e.* the same cultivar), differences in <sup>15</sup>N abundance in the plant would be associated with strain genotype, generating a distinctive behavior (23). This study confirms the importance of *Rhizobium* strains in <sup>15</sup>N isotopic discrimination in field pea, due to isolate origin, where adaptive or microevolutionary changes would be expected after genetic divergences (20, 28).

In our field experiment, aerial biomass (Biom<sub>Aerial</sub>) and yield (Yield<sub>Grain</sub>) had no interaction (same as in the greenhouse). Yields obtained from the field experiment (3.2 Mg ha<sup>-1</sup> for D70, 3.3 for D156 Mg ha<sup>-1</sup>, and 3.8 Mg ha<sup>-1</sup> for D191; 3.5 Mg ha<sup>-1</sup> on average) were higher than those obtained in the 2014 crop cycle by Prieto *et al.* (2015) (2.7 Mg ha<sup>-1</sup>) and Enrico *et al.* (2020) (1.3 Mg ha<sup>-1</sup>) in eight experiments performed in different environments of Argentina between 2012 and 2017. Unlike that observed in the greenhouse, inoculation with D191 showed the best behavior in aerial and vegetative biomass and nodule number per plant. In the field experiment, D156 and D70 had the same behavior on the specific variables derived from the *Rhizobium* strain, such as  $%N_{BNF}$  and/or the contribution of  $N_{BNF}$ , and maintained the efficiency demonstrated in the greenhouse (23). In this study, in the field experiment, we observed 70.5% (range 67 to 73%) of  $%N_{BNF}$  on average with 67% for D91, 72% for D70, and 73% for D156, all higher values than those established by Enrico *et al.* (2020), who found 59% inoculated with D70, but within the interquartile range obtained for field pea (33 to 83%).

The incorporation of a winter legume such as field pea (inoculated with efficient N-fixing strains) preceding a summer crop (*e.g.* maize) may constitute a valid agronomic management. Enrico *et al.* (2020) observed a positive effect on maize yield when the previous crop was inoculated (field pea with strain D70). Having rich in N and low C:N ratio stubble allowed rapid soil N availability. This higher availability of N considerably reduced the need for chemical nitrogen fertilization.

## CONCLUSIONS

Most of the isolates that were infective and effective for *Pisum sativum* cultivars here evaluated came from species other than field pea.

Strain D70 maintained good behavior in the growth chamber, greenhouse, and field experiments concerning nodule, nitrogen, biomass, and yield variables, whereas, D156 had similar or better behavior in the greenhouse and field experiments. In contrast, D191 had only better behavior in the field experiment.

The  $\beta$  value calculation of <sup>15</sup>N natural abundance model to determine  $N_{FBN}$  was key to adjusting  $N_{FBN}$  contribution. Determining differences in strain behavior results an essential requirement and would allow a more precise adjustment of N input.

Finally, strains selected for BNF efficiency should implicitly carry infectivity ability in all commercial cultivars. This will allow certainty in their recommendation for the formulation of inoculants. If new evaluations are required in different agroecological environments, strain D156 could be a good candidate for bacterial single-strain inoculants, as well as for a microbial consortium for *Pisum sativum*.

#### SUPPLEMMENTARY MATERIAL

Squematic figure of the dual-choice olfatometer utilized in choice test with stored grain pests: https://drive.google.com/drive/folders/17vBzpdAKT\_WP2l\_SdnicLF\_EH-puPzCy?usp=sharing

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## Effects of geomorphology and distribution of water sources for livestock on the floristic composition and livestock receptivity of the Arid Chaco

## Efecto de la geomorfología y la distribución de las fuentes de agua para el ganado en la composición florística y receptividad ganadera del Chaco Árido

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## ABSTRACT

Livestock production in semi-arid areas is possible due to the presence of permanent water sources, which create a radial pattern of grazing intensity known as the piosphere. For this reason, we predicted that permanent water sources would negatively impact the ecological conditions of plant communities, leading to variations in livestock receptivity. To test this prediction, we compared grazing gradients in two geomorphological units, using distance to water sources as an indicator of accumulated livestock pressure. We assessed variations in the botanical composition of both areas by analysis of variance and principal components analysis. Additionally, we modeled the relationship between distance to water source and livestock receptivity. Our results revealed significant differences in the contribution of different species based on their distance to water sources. Notably, a non-linear regression model provided the best fit for the relationship between water source and livestock receptivity in both geomorphological units. These findings demonstrate that the distance to permanent water sources serves as a reliable indicator of accumulated livestock pressure in semi-arid regions like the study area.

## Keywords

arid Chaco • natural grasslands • piosphere • geomorphology

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## RESUMEN

La producción de ganado en zonas semiáridas es posible debido a la existencia de fuentes de agua permanentes, lo que genera un patrón radial de intensidad de pastoreo llamada piosfera. Por tal motivo, nuestra predicción se basa en que las fuentes de agua permanente influyen negativamente en las condiciones ecológicas de las comunidades vegetales provocando diferentes receptividades ganaderas. Comparamos gradientes de pastoreo tomando la distancia a las fuentes de agua como indicador de presión ganadera acumulada en dos unidades geomorfológicas. La variación en la composición botánica de las diferentes áreas se realizó utilizando análisis de la varianza y un análisis de componentes principales. Se efectuó un modelo de la relación entre la distancia a la fuente de agua y la receptividad ganadera. Nuestros resultados mostraron diferencias significativas en la contribución de las diferentes especies en relación con la distancia a las fuentes de agua. El modelo de regresión no lineal fue el que mejor se ajustó entre la fuente de agua y la receptividad ganadera para ambas unidades geomorfológicas. La distancia a las fuentes de agua permanente es un buen indicador de la presión ganadera acumulada en regiones semiáridas como el área de estudio.

## **Palabras clave**

Chaco árido • pastizales naturales • piosfera • geomorfología

## INTRODUCTION

Like other arid and semi-arid regions, the Arid Chaco exhibits heterogeneous spatial patterns of degradation driven by the uneven distribution of drinking throughs (4, 19). The availability of permanent water sources is critical for livestock production in these areas. Consequently, a radial pattern of grazing intensity, known as 'piosphere,' develops around water sources. Analysis of piospheres allows to quantify the effects of radial attenuation of a disturbance on the system's condition (8).

Piospheres are areas around water sources that suffer from heavy grazing and trampling, making it difficult for vegetation to establish (degraded zones) (22). Management practices near these water bodies often involve continuous grazing without rest periods for the land, leading to changes in plant composition (10). As a result, plant communities near water sources are dominated by annual grasses and/or species with little grazing value (23).

Grassland dynamics can be conceptualized using a state-and-transition model (STM), which describes vegetation as existing in discrete states with transitions triggered by natural events or management practices (5, 38). Within an STM framework, each state represents a distinct plant community reflecting the current ecological conditions relative to a climax community. Transitions within a state are considered reversible, while transitions between states can be irreversible depending on disturbance severity (26, 36).

In interaction with livestock, geomorphology shapes the spatial patterns and dynamics of plant communities (27). In arid and semi-arid environments, water availability is the primary control of vegetation structure and function, surpassing even the influence of the physical and chemical characteristics of each geomorphological unit (24). Recent sedimentary environments, are particularly sensitive to the colonization strategies employed by vegetation given the specific arrangement and structures of their deposits (15).

Evaluating the impact of grazing on ecosystem integrity becomes difficult, especially in systems where the original condition is unknown given widespread and irreversible transformations of plant communities (10). Furthermore, when a system has surpassed a critical threshold and transitioned to an alternative stable state, experiments utilizing grazing exclusion methods may yield misleading interpretations (9).

An alternative approach to evaluate grazing is to interpret current vegetation assemblages in the context of accumulated livestock pressure and associated management practices. Proxy indicators such as proximity to water sources (1), livestock posts (32), and even specific plant growth forms or species abundance (13, 18) can be employed for this purpose. These approaches often provide solid information to inform management decisions, particularly when time constraints prevent reliance on long-term grazing trials (10). This study investigates the impact of extensive livestock farming on botanical composition and livestock receptivity within distinct geomorphological units of the Arid Chaco region in San Juan, Argentina. We aim to contribute to the understanding of the ecological state of plant communities in semi-arid areas assessing livestock receptivity potential and informing sustainable management strategies. The study area exhibits diverse plant communities with varying physiognomies, shaped by the combined effects of livestock farming, forestry exploitation, and interactions with geomorphological processes.

Within this framework, our study hypothesizes that botanical composition within each geomorphological unit varies in response to distance from water sources. This prediction is based on permanent water sources negatively influencing the ecological states of plant communities, leading to differential livestock receptivity.

#### MATERIALS AND METHODS

## Study area

The study area covers approximately 1000 km<sup>2</sup> within the Valle Fértil department, located between parallels 30°50' and 30°29' S and meridians 67°27' and 67°12' W. It is located east of the Sierras de Valle Fértil-La Huerta within the Bajo Oriental depression, bordering La Rioja to the east. This region represents a unique and characteristic expression of the Arid Chaco in San Juan, exhibiting distinctive physiographic, climatic, social, and productive features. Its climate is classified as arid (BWk) according to the Köppen-Geiger system (20), with an average annual temperature of 17.9°C. Precipitation falls within the 200-300 mm isohyets. The vegetation comprises an open forest dominated by *Aspidosperma quebracho blanco, Neltuma flexuosa*, and *Bulnesia retama*. A shrub layer rich in *Larrea divaricata* is present, while the herbaceous layer is well-represented by genera such as *Leptochloa, Setaria, Aristida*, and *Pappophorum*, among others.

## **Basin delimitation**

A high-resolution (12.5-meter pixel size) Alos Palsar Digital Elevation Model (DEM) was used to delineate the basins. Flow direction derived from the DEM was employed to identify the basins associated with the main channels within the study area. GRASS GIS software, a free and open-source module linked to QGIS 3.14.15 (30), was used for the digital processing of the DEM.

Given the study area corresponds to a plain, vectors representing the mountain range were excluded from the analysis. Consequently, sampling was focused on the geomorphological units of the foothills (Piedemonte) and the river floodplain (Floodplain) (figure 1, page 15).

## Location of livestock posts

The study area encompasses a diversity of fenced enclosures with varying surface areas. Each enclosure has a water source (dam, well, or drinker) for livestock located in the vicinity of the post. All sampled fields employed a continuous grazing regime, where cattle roam freely with unrestricted access to a central water source. During periods of extreme drought, ranchers are forced to either sell animals at a reduced price or pay to graze them in fields with superior forage availability.

This study adopts the methodology proposed by Cingolani *et al.* (2008), who emphasizes the value of distance to permanent water sources as an objective, measurable, and precise indicator for assessing the long-term effects of extensive grazing.

## **Establishment of distance ranges**

Given the extensive grazing behavior of livestock in the Arid Chaco region (16), we categorized sampling locations according to their distance to water sources: close range (0-1000 m), intermediate range (1001-2000 m), and far range (>2001 m). This approach was applied within each geomorphological unit (Piedemonte and Floodplain). Thirty 50-meter transects were established within each unit, with readings taken at 50 cm intervals along the transects (figure 1, page 15). In the Piedemonte, ten transects were located close to water sources (within 1000 m, designated PMC), ten at intermediate distances (1001-2000 m, PMI),

and ten at a far range (>2001 m, PML). Similarly, in the Floodplain, ten transects were placed near water sources (<1000 m, PLC), ten at intermediate distances (1001-2000 m, PLI), and ten at a far range (>2001 m, PLL) (figure 1).



Fuente/Source: IGN (Instituto Geográfico Nacional, República Argentina).

**Figure 1.** Location of study sites on the Arid Chaco phytogeographic province of San Juan and La Rioja, Argentina. Projection: Posgar 2007; Argentina/2; EPSG: 5344.

**Figura 1.** Ubicación de los sitios de estudio en la Provincia Fitogeográfica del Chaco Árido de la provincia de San Juan y La Rioja, Argentina. Proyección: Posgar 2007; Argentina/2; EPSG: 5344.

## **Determination of botanical composition**

Botanical composition was quantified at peak growing season using the modified Point Quadrat method (28) in April and May, following the rainy season. The resulting data were then used to calculate percent coverage and specific contribution per contact (CSC) for each species.

$$CSC = \frac{C}{\sum C} * 100$$

where:

C = Contacts of a species

 $\Sigma C$ = Sum of the contacts of all species

## Determination of field forage receptivity for bovine cattle

Livestock receptivity refers to the amount of forage required to sustain an Equivalent Cow Unit (EV). An EV is defined as a 400 kg cow that gestates and raises a calf to 6 months old (weighing 160 kg), including the forage consumed by the calf (Passera, C.P.Q.). One EV can be supported by 100 Pastoral Value (VP) units.

Plant species for calculating Pastoral Value (VP) were selected based on established criteria for determining specific quality indices (19, 29). Table 1 (page 16-19), presents their classification according to livestock preference: Preferential (P) - readily consumed species without selection; Good (G) - species initially rejected compared to Preferential ones; Regular (R) - species presenting some difficulty in consumption.

**Table 1**. Contribution of each species, total coverage, and diversity indices in each sector considered for sampling within each Geomorphological Unit separately.

**Tabla 1**. Aporte de cada especie, cobertura total e índices de diversidad en cada uno de los sectores considerados para el muestreo en cada Unidad Geomorfológica por separado.

	Geomorphological Units								
		Piede	monte		Floodplain				
	PML	PMI	РМС	С	PLL	PLI	PLC	С	S
Species				Perei	nnial Gras	ses			·
Leptochloa pluriflora	11.03c	-	2.48b	0.54	15.25d	10.36d	-	0.63	Р
Gouinia paraguayensis	2.49b	0.63a	0.52a	0.54	1.34e	2.45d	2.78d	-0.48	Р
Setaria lachnea	5.23b	5.99b	0.16c	0.5	1.21d	0.54d	-	0.62	Р
Setaria leucopila	5.93b	6.73b	3.84a	0.32	6.96d	6.00d	-	0.49	Р
Setaria cordobensis	0.64	-	-	0.44	-	-	-	-	Р
Digitaria californica	1.71b	2.16b	0.47a	0.41	0.60d		0.52d	0.19	Р
Cenchrus ciliaris	-	0.16	-	0.04	-	-	-	-	Р
Cottea pappophoroides	1.63b	0.68a	1.60b	-0.09	-	-	-	-	G
Setaria hunzikeri	1.9	-	-	0.55	-	-	-	-	Р
Pappophorum philippianum	1.32a	4.85b	11.12b	-0.48	-	-	-	-	Р
Neobouteloua lophostachia	0.87a	2.41b	4.06b	-0.43	0.92d	2.34d	10.75e	-0.59	R
Leptochloa crinita	16.34a	1.73b	0.53b	0.56	1.13d	1.84d	-	0.33	Р
Leptochloa dubia	0.34	-	-	0.56	-	-	-	-	Р
Chloris parvispicula	0.79	-	-	0.34	-	-	-	-	Р
Aristida medocina	-	-	0.18	-0.58	11.14d	1.14e	2.78e	0.71	Р
Setaria sp	-	-	-		0.81	-	-	0.65	Р
Sporobolus pyramidatus	0.17a	0.31a	-	0.29	0.40d	-	1.08d	-0.3	G

Different letters indicate significant differences (P < 0.05). The column labeled C represents the correlation of each species, total coverage, and diversity indices with respect to the distance from water sources. Significant correlations (P < 0.05) are indicated by grayed cells. Letras diferentes indican diferencias significativas (P<0,05). La columna indicada con C representa la correlación de cada especie, cobertura total e índices de diversidad, respecto de la distancia con las fuentes de agua. Las correlaciones significativas (P<0,05) se indican con los casilleros pintados de gris.

Geomorphological Units										
Piedemonte					Floodplain					
	PML	РМІ	РМС	С	PLL	PLI	PLC	С	S	
Annual Grasses										
Bouteloua aristoides	7.21a	19.85b	31.13b	-0.55	-	-	-	-	R	
Eragrostis mexicana var. virescens	0.84b	-	0.38b	0.34	-	-	-	_	G	
Aristida adscensionis	0.13a	0.81a	1.28a	-0.49	2.36d	-	0.36d	0.38	R	
Tripogon spicatus	2.03	-	-	0.55	-	-	-	-	R	
			SI	nrubbery						
Larrea divaricata	9.05a	17.86b	20.26b	-0.75	30.82d	33.97d	55.71e	-0.74	-	
Lippia turbinata	0.48b	-	0.38b	0.16	-	-	-	-	-	
Mimozyganthus carinatus	1.30a	5.92a	1.26a	0.05	4.54	-	-	-	R	
Cordobia argentea	0.60ab	0.68b	0.16a	0.43	5.60d	16.51e	8.65d	-0.24	G	
Capparis atamisquea	0.48	-	-	0.55	-	-	-	-	R	
Senna aphyla	1.95a	7.49b	0.49c	0.17	0.46d	5.65e	1.20d	-0.1	R	
Pithecoctenium cynanchoides	0.67	-	-	0.56	-	-	-	-	-	
Lantana fucata	1.74a	0.54b	-	0.67	-	-	-	-	-	
Senegalia gilliesii	0.12a	-	0.57a	-0.35	-	-	-	-	R	
Lippia integrifolia	1.00a	-	1.88a	-0.24	1.38	-	-	0.56	-	
Ximenia americana	-	0.51a	0.65a	-0.58	-	3.57	-	-0.02	-	
Aloysia gratissima	1.05a	0.16b	-	0.72	-	-	-	-	-	
Condalia microphylla	0.51	-	-	0.56	-	0.54	-	-0.06	R	
Tillandsia sp	0.13a	-	0.16a	-0.18	-	-	-	-	R	
Lantana grisebachii	-	2.93a	5.00a	-0.67	-	-	-	-	-	
Strombocarpa torquata		2.38a	0.47b	-0.1	0.44d	5.66e	8.34e	-0.74	R	
Salvia gilliesii	-	-	-	-	0.23	-	-	0.56	-	
Cereus aethiops	-	-	-	-	0.23	-	-	0.56	-	
Ephedra triandra	-	-	-	-	0.21	-	-	0.45	G	
Lycium chilensis var. filifolium	1.02c	0.33a	0.34a	0.53	0.23	-	-	0.56	R	
Justicia gilliesii	2.89a	-	0.38b	0.55	-	-	-	-	G	

significant differences (P<0.05). The column labeled C represents the correlation of each species, total coverage, and diversity indices with respect to the distance from water sources. Significant correlations (P<0.05) are indicated by grayed cells. Letras diferentes indican diferencias significativas (P<0,05). La columna indicada con C representa la correlación de cada especie, cobertura total e índices de diversidad, respecto de la distancia con las fuentes de agua. Las correlaciones significativas (P<0,05) se indican con los casilleros pintados de gris.

Different letters indicate

	Geomorphological Units								
	Piedemonte Floodplain								
	PML	РМІ	РМС	С	PLL	PLI	PLC	С	S
Mimozyganthus carinatus	1.30a	5.92a	1.26a	0.05	4.54	-	-	0.66	R
Lantana xenica	-	-	0.18	-0.58	-	0.31d	3.31e	-0.68	-
Tricomaria usillo	-	-	0.18	-0.58	1.38d	3.92e	0.96d	0.07	G
			l	Forestry					
Parkinsonia praecox var praecox	3.53a	2.55a	3.65a	3.00E- 03	0.42e	1.92d	-	0.06	R
Celtis tala	0.66a	0.16a	-	0.36	-	-	-	-	R
Neltuma flexuosa	8.27a	0.33b	1.30b	0.53	2.11d	2.01d	3.13d	-0.26	G
Geoffroea decorticans	-	-	-		6.75d	1.18d	0.52e	0.81	G
A quebracho blanco	-	2.44b	0.19b	0.01	-	-	-	-	-
Bulnesia retama	0.51a	4.39b	0.16a	0.06	-	-	-	-	-
		Dice	otyledono	us herbac	eous plan	ts			
Bidens subalternans	0.29b	-	0.38b	-0.06	-	-	-	-	-
Pseudabutilon virgatum	0.25a	0.18a	0.95a	-0.37	-	-	-	-	-
Zinnia pruviana	0.25a	-	0.19a	0.1	-	-	-	-	-
Talinum polygaloides	-	-	0.18	-0.58	-	-	-	-	-
Gaya parvifolia	0.46a	0.16a	-	0.72	0.69d	-	0.52d	0.14	-
Amaranthus standleyanus	0.26	-	-	0.34	-	-	-	-	-
Gomphrena pulchella	0.34a	-	0.19a	0.27	-	-	-	-	-
Tribulus terrestris	0.12	-	-	0.56	-	-	-	-	-
Parthenium hysterophorus	0.13a	0.18a	-	0.25	-	-	-	-	-
Euphorbia catamarcensis	0.67	-	-	0.56	-	-	-	-	-
Sphaeralcea miniata	0.56a	1.88a	1.90a	-0.29	-	-	-	-	-
Allionia incarnata	0.13a	1.93b	1.14ab	-0.27	-	-	-	-	-
Heliotropium mendocinum	-	0.18	-	0.03	-	-	-	-	-
Porophyllum lanceolatum	-	0.16	-	0.04	-	-	-	-	-
Portulaca grandiflora	-	-	0.18	-0.58	-	-	-	-	-
Solanum elaeagnifolium	-	-	0.18	-0.58	-	-	0.52	-0.5	-

Different letters indicate significant differences (P<0.05). The column labeled C represents the correlation of each species, total coverage, and diversity indices with respect to the distance from water Significant correlations (P<0.05) are indicated by grayed cells. Letras diferentes indican diferencias significativas (P<0,05). La columna indicada con C representa la correlación de cada especie, cobertura total e índices de diversidad, respecto de la distancia con las fuentes de agua. Las correlaciones significativas (P<0,05) se indican con los casilleros pintados

de gris.

Different letters indicate significant differences (P<0.05). The column labeled C represents the correlation of each species, total coverage, and diversity indices with respect to the distance from water sources. Significant correlations (P<0.05) are indicated by grayed cells. Letras diferentes indican diferencias significativas (P<0,05). La columna indicada con C representa la correlación de cada especie, cobertura total e índices de diversidad, respecto de la distancia con las fuentes de agua. Las correlaciones significativas (P<0,05) se indican con los casilleros pintados de gris.

	Geomorphological Units								
	Piedemonte				Floodplain				
	PML	РМІ	РМС	С	PLL	PLI	PLC	С	S
		Di	versity Co	verage an	d Indexes				
Coverage	94.89a	88.96b	70.37c	-0.21	73.14d	67.78d	53.45e	0.63	
Shannon - Weaver	2.92a	2.54b	2.23c	0.68	2.21d	2.07d	1.58e	0.78	
Simpson	0.08a	0.12a	0.2b	-0.57	0.16c	0.19c	0.34d	-0.73	

The Pastoral Value (VP) is calculated by the following formula:

$$VP = 0.1 * (CSC * Is) Cf$$

where:

0.1 = constant coefficient

*CSC* = Specific Contribution per Contact

*Cf* = Forage Coverage

*Is (Specific Quality Index)* = ranging from 1 to 10, representing the classification of species based on their suitability and potential as forage.

## Statistical analysis

Variation in botanical composition across different areas was explored using principal component analysis (PCA) based on a correlation matrix among all present species and their respective contact-specific contributions (CSC) within the community.

The influence of distance to water sources on botanical composition, diversity (Shannon-Weaver and Simpson), and total cover was assessed using analysis of variance (ANOVA) with a completely randomized design. Separate ANOVAs were conducted for each geomorphological unit (Piedemonte and Floodplain), treating distance categories (PMC, PMI, PML and PLC, PLI, PLL) as fixed effects. Additionally, a Pearson correlation analysis was performed to explore relationships between plant species, diversity indices, and total coverage. Data were square root transformed to address non-homoscedasticity and normality violations. Means were compared using Tukey's test. Data results are presented using untransformed means for ease of interpretation.

Visual exploration of the data involved plotting livestock receptivity against distance to water sources for each geomorphological unit. This analysis revealed a negative correlation up to a certain distance, followed by a plateau effect. Consequently, non-linear models were fitted for each unit to account for this observed pattern (12).

## RESULTS

## **Botanical composition**

A total of 26 families were recorded, with *Poaceae* being the best-represented family, comprising 22 species. *Fabaceae, Verbenaceae*, and *Asteraceae* followed in abundance, with 7, 6, and 4 species, respectively.

The Piedemonte plant community is dominated by *Leptochloa crinita*. Additionally, *Leptochloa pluriflora, Bouteloua aristoides, Setaria leucopila, Setaria lachnea,* and *Gouinia paraguayensis* are present in PML sectors. In PMI and PMC sectors, the *Bouteloua aristoides* contribution increases, while the contributions of other grass species (except *Setaria leucopila* and *Setaria lachnea*, which maintain their presence in the PMI sector) decrease.

The Piedemonte shrub layer is dominated by *Larrea divaricata*, accompanied by *Mimosiganthus carinatus*, and *Senna aphylla*. The forest layer exhibits a distinct zonation: *Prosopis flexuosa* dominates in PML sectors, *Bulnesia retama* in PMI sectors, and *Aspidosperma quebracho blanco* occurs at low densities across all sectors.

Within the Floodplain, shrub communities dominate the vegetation structure. *Larrea divaricata* is the most abundant species, with accompanying species *Cordobia argentea* and *Mimozyganthus carinatus* in PLL sectors. In PLI sectors, shrub species with increased contributions include *Cordobia argentea*, *Senna aphyla*, and *Prosopis torquata*. Similarly, *Larrea divaricata* shows a significant increase in contribution within PLC sectors, accompanied by *Prosopis torquata*, *Lantana xenica*, and *Cordobia argentea*.

Following the shrub layer, grasses contribute significantly to the Floodplain community. *Aristida mendocina* and *Leptochloa pluriflora* are the most prominent grasses, with *Setaria leucopila* present to a lesser extent in PLL sectors. However, in PLC sectors, *Leptochloa pluriflora* disappears, and *Aristida mendocina* abundance declines significantly. Conversely, *Neobouteloua lophostachia* becomes the dominant grass species.

The forest layer plays a minor role in the Floodplain, with *Geoffroea decorticans* being the most prevalent species in PLL sectors, while *Prosopis flexuosa* dominates in PLI and PLC sectors (table 1, page 16-19).

Across both geomorphological units, total coverage and the Shannon-Weaver diversity index exhibit a significant rise with increasing distance from water sources. Conversely, Simpson's dominance index shows a significant positive correlation with proximity to water sources (table 1, page 16-19).

Principal Component Analysis (PCA) revealed a variation pattern in botanical composition across the sampled geomorphological units. The first two principal components explained 61% of the total species variance, effectively separating sectors within the Floodplain from those in the Piedmont. Within the Piedmont, sectors located close and halfway to water sources (PMC and PMI) were distinct from those situated further away (PML) (figure 2).



Figure 2. Principal Component Analysis (PCA) ordination diagram of sampling sectors (PMC, PMI, PML, PLC, PLI, PLL) and associated plant species.
Figura 2. Diagrama de la ordenación del Análisis de Componentes Principales (ACP) de los sectores seleccionados para los muestreos (PMC, PMI, PML, PLC, PLI, PLL) y las especies asociadas.

Within the alluvial plain, CP1 and CP2 displayed the most negative values for *L. divaricata* (Lar div), *C. argentea* (Cor arg), *T. usillo* (Tri usi), and *G. paraguayensis* (Ga par). In contrast, within the floodplain and specifically PML sectors, CP1 exhibited the most positive values for *J. gilliesii* (Jus gui), *L. chilensis var. filifolium* (Lyc chi), *S. cordobensis* (Set cor), *L. crinita* (Lep cri), *C. tala* (Cel tal), *N. flexuosa* (Ne fle), and *L. pluriflora* (Lep plu). Similarly, positive values in CP2 were associated with PMI and PMC sectors for *B. aristoides* (Bou aris), *A. quebracho blanco* (Asp que), *S. miniata* (Spha mini), *A. incarnata* (All inc), *L. grisebachii* (Lan gri), *P. philippianum* (Pap phi), *H. mendocinum* (Hel men), and *P. lanceolatum* (Po lan) (figure 2, page 20).

## Livestock receptivity and distance to water sources

Non-linear regression analysis proved the best fit between livestock receptivity (Ha.EV<sup>-1</sup>) and distance to water sources (DP), for both the Piedemonte (PM) and Floodplain (PL) units (figure 3).



**Figure 3.** Non-linear regression model between livestock receptivity (Ha.EV<sup>-1</sup>) and distance to water source (DP) in the Piedemonte (A) and Floodplain (B).

**Figura 3.** Modelo de regresión no lineal entre la receptividad ganadera (Ha.EV<sup>-1</sup>) y la distancia a la fuente de agua (DP) en el Piedemonte (A) y en la Planicie de inundación (B).

## DISCUSSION

Our findings reveal changes in floristic composition across both geomorphological units, with distance to water sources playing a significant role (figure 2, page 20 and table 1, page 16-19). The study area is characterized by fenced fields of varying sizes and shapes, along with an irregular distribution of watering holes, dams, and/or livestock shelters. This heterogeneity contributes to a highly diverse botanical composition within plant communities. Szymañski *et al.* (2022) attribute similar variations primarily to livestock management practices. Specifically, our results indicate the presence of plant communities dominated by perennial forage grasses and others dominated by non-forage annual grasses. In such cases, the potential for transition towards improved ecological conditions exists due to the presence of seed sources from desirable species, facilitating the natural rehabilitation of nearby degraded areas (10).

The results align with established patterns of overgrazing documented by Briske *et al.* (2006), where plant communities near water sources (PMC and PLC) are dominated by annual grasses and/or non-forage species (table 1, page 16-19). While grazing responses vary among species, when grazing pressure alters biotic structures and interactions changes in species coverage can be attributed to underlying biotic mechanisms of grazing impact. Several studies support this notion, demonstrating that grazing pressure favors plant traits associated with rapid growth, regeneration, annual life cycles, and a ruderal strategy (11, 14, 32).

Across the geomorphological units, perennial grasses exhibited a significant trend of increased colonization away from water sources (table 1, page 16-19). Conversely, annual grasses and species with low forage value showed the opposite pattern. This aligns with established knowledge on grazing pressure inducing species turnover, favoring some species while hindering others (10). Briske *et al.* (2006), further suggests that heterogeneous livestock use is common in homogeneous landscapes (*i.e.*, within a single geomorphological unit). This heterogeneity contributes to an increase in landscape physiognomic diversity. However, increased livestock density can lead to a decline in unused areas (37). While maintaining some level of unused sites is important for overall community coexistence (10), excessively large paddocks with few water sources or shelters may allow even high livestock densities to coexist with unused areas (21, 26).

Total coverage and Shannon-Weaver diversity in the geomorphological units exhibited a significant decrease near water sources (table 1, page 16-19). Conversely, Simpson's dominance index shows a positive trend in these areas. This pattern aligns with observations in other semi-arid African grasslands, where grazing-sensitive forage species are replaced by those more resistant to grazing. However, this replacement may not fully compensate for diversity losses, potentially leading to environments dominated by a few resilient species. In this context, grazing regimes exceeding the historical range experienced by the ecosystem are likely to induce a significant decline in overall diversity (16).

The observed variations in floristic composition may be explained by mechanisms of resilience to harsh conditions proposed by Cingolani *et al.* (2008) for arid and semi-arid grasslands. These mechanisms, such as reduced sprouting and production of long-lived seeds, can confer some resistance to short periods of intense grazing. However, this resilience is insufficient to withstand continuous grazing for extended periods, particularly under moderate to high grazing pressure. A study evaluating intensity versus grazing strategies in perennial forage grasses of the Arid Chaco region found that while increased defoliation intensity may maximize short-term biomass yield, it negatively impacts long-term sustainability and produces detrimental residual effects on grasslands (31). Importantly, the study identified continuous grazing combined with high defoliation intensity as the least sustainable management practice for this ecosystem (31).

In these systems, continuous grazing by introduced herbivores without proper stocking management is highly likely to cause widespread extinctions and/or significant declines of specific plant species (10, 21, 25). However, a contrasting recovery trajectory may emerge when degradation and/or extinction of desirable species occurs homogenously across large areas (several square kilometers) and grazing pressure is subsequently relieved (10, 21). An alternative stability perspective for degraded ecosystems emphasizes persistence relative to the livelihood needs of ecosystem users. If the timeframe for ecosystem recovery surpasses

user tolerance, typically a few years to a decade, the degraded state can be considered persistent from a practical viewpoint. Therefore, a 'persistent decline' of an ecosystem service may not always be associated with a critical threshold and a shift to a new ecological state. Slow recovery within a single stability domain can also lead to persistent declines (23).

Livestock receptivity values in this study range from 47.02 to 2.86 Ha.EV<sup>-1</sup> (figure 3, page 21). These values are consistent with observations from other arid and semi-arid regions in Patagonia (23), Puna, Monte, and the driest parts of the Arid Chaco, where reported livestock receptivities typically exceed 6 Ha.EV<sup>-1</sup> (3). Nevertheless, the value of 8.92 Ha.EV<sup>-1</sup> measured in the PMC sector deviates from historical data for degraded foothills, which reported values around 24 Ha.EV<sup>-1</sup> (33). This discrepancy may be attributed to interannual precipitation variations, known to cause substantial differences (>300%) in grassland productivity within degraded areas of the Arid Chaco (11).

Despite using the same model for both geomorphological units, the estimated values and rates of change differed (figure 3, page 21). These non-linear relationships align with the findings of Sasaki *et al.* (2008), who highlight the non-linear nature of ecological patterns and processes in response to grazing pressure. Furthermore, the estimated values from our models are consistent with other studies conducted in the region (2, 7, 11). Additionally, the observed spatial patterns of vegetation change correspond to documented livestock grazing behavior in the Arid Chaco (16). This distribution pattern of livestock receptivity informs spatial dynamics within grazing plots, allowing for the establishment of criteria for plot size and water source distribution.

## **CONCLUSIONS**

Across both geomorphological units, distance to water sources significantly influences floristic composition, which in turn affects livestock receptivity. Areas closer to water sources are dominated by annual grasses and/or non-forage species, resulting in lower livestock receptivity. Conversely, areas farther from water sources are dominated by perennial grasses with higher forage value, leading to increased livestock receptivity.

A long history of domestic grazing has shaped the heterogeneity of plant communities, influencing species diversity and dominance. This translates to a landscape with varying ecological conditions (good, fair, and bad) distributed in different proportions. Quantifying the extent of each condition is crucial for optimizing water source distribution and achieving biologically and economically sustainable grazing management strategies. Notably, non-linear models provided the best fit between distance to water sources and livestock receptivity for both geomorphological units under study (figure 3, page 21).

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## Landform heterogeneity drives multi-stemmed Neltuma flexuosa growth dynamics. Implication for the Central Monte Desert forest management

## La heterogeneidad de paisaje modula la dinámica en el crecimiento de *Neltuma flexuosa*. Implicancias para el manejo forestal de los bosques del Desierto del Monte Central

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## ABSTRACT

Drylands represent the main earth biome, providing ecosytemic services to a large number of people. Along these environments, woodlands are often dominated by multi-stemmed trees, which are exploited by local inhabitants to obtain forest products for their livelihood. In central-west Argentina, *Neltuma flexuosa* (algarrobo) woodlands are distributed across different landform units, varying in topographical and soil characteristics. This research aimed to reconstruct stem-growth time until harvestable diameter was achieved, and biological rotation age according to topo-edaphic variability in three algarrobo forests using dendrochronological methods. Results indicated that landform heterogeneity modulated species radial growth, influencing stem increments and cutting cycle period. In this sense, a decreasing trend in tree productivity emerged along a loamy-to-sandy textured soil gradient. These findings provide useful novel information for *N. flexuosa* forest management, suggesting the need to account for spatial landform/soil heterogeneity when examining desert forest dynamics.

### **Keywords**

arid woodlands • forest management • growth form • tree-rings

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## RESUMEN

Las tierras secas representan el principal bioma terrestre, proveyendo numerosos servicios ecosistémicos a un ingente número de personas. En estos ambientes, los bosques son a menudo dominados por árboles de porte multi-fustal, explotados por los habitantes locales para obtener productos forestales útiles para su sustento. En los territorios del centro-oeste de Argentina, los bosques de Neltuma flexuosa (algarrobo) están distribuidos en diferentes unidades de paisaje, los que presentan variaciones en las características topográficas y edáficas. En este estudio reconstruimos, a través de métodos dendrocronológicos, el crecimiento, el intervalo para obtener productos maderables, y el turno biológico de corta en función de la variabilidad topo-edáfica en tres bosques diferentes de algarrobo. Los resultados indican que la heterogeneidad del paisaje modula el crecimiento radial de la especie, influenciando el incremento diamétrico y el turno de corta. En este sentido, se evidencia una tendencia decreciente en valores de productividad según un gradiente edáfico desde textura limosa a arenosa. Estos hallazgos proveen información novedosa y de utilidad para los planes de manejo forestal de los bosques de N. flexuosa, indicando la necesidad por considerar la heterogeneidad espacial establecidas por las unidades de paisaje/suelos en estudios relativos a la dinámica forestal de los bosques de tierras secas.

## **Palabras claves**

bosques de tierras secas • manejo forestal • forma de crecimiento • ancho de anillo

## INTRODUCTION

Worldwide, drylands represent the largest biome on Earth, covering more than 40% of land surface (7, 29). These environments, characterized by precipitation/evapotranspiration ratio below 0.2, include 20% of plant diversity hotspots, play a fundamental role in various biogeochemical cycles, and contribute to approximately 40% of global net primary productivity (11, 16, 29).

Drylands host more than two billion people (29). Along these biomes, forests cover approximately 1.400 Mha, providing several ecosystemic services like water and soil regulation, food, biochemical, and raw material provision, as well as cultural services (7, 24). Physiognomically, many desert forests are dominated by tree species capable of regeneration through development of new tissue following disturbance, a physiological trait known as resprouting (8, 30). This results in vegetation with coexisting one- and multi-stemmed trees, which differ in growth, productivity, reproduction, and survival ability (30).

Dryland forests have suffered various types of natural and anthropogenic disturbances, such as, fire, insect outbreak, and drought, in addition to high deforestation rates due to agricultural and livestock expansion (6, 20). In this sense, it is estimated that, globally, about 220,000 km<sup>2</sup> of tree-covered drylands were converted into other land cover types between 1992 and 2015 (29). This suggests the need to sustainably exploit and manage these natural resources, particularly in the actual climate change scenario (16).

The Central Monte Desert (hereafter, CMD) is a dryland biome located in central-west Argentina (2). Along these districts, the algarrobo dulce, *Neltuma flexuosa* (DC.) C.E. Hughes & G.P. Lewis (formerly designated as *Prosopis flexuosa* DC), is a dominant tree species growing on a variety of landform units (20). These woodlands were severely exploited during the first half of the 20<sup>th</sup> century, mainly for domestic wood demand, railway construction and, afterwards, vineyard expansion, with a resulted extraction of approximately 1 million tons of wood (1, 28).

Nowadays, the *N. flexuosa* CMD forests are mostly managed by the local Huarpe native community, through forestry practices that consider the typical low tree growth rates of these woodlands (28). As a result, logging and thinning are extremely rare activities in this area, whereas other silvicultural approaches, such as pruning and deadwood removal, represent valuable forest management alternatives (4, 27). Pruning plays a prominent role due to the high presence of multi-stemmed trees in CMD forests, as well as their higher productivity with respect to one-stemmed individuals (3, 4).

Previous analyses examined productivity in one- and multi-stemmed *N. flexuosa* trees, providing valuable information regarding the CMD algarrobo forest management, but solely at local scale (4). As previously mentioned, the CMD N. flexuosa woodlands grow on a mosaic of landform units, expression of soil variability and topographic characteristics (21). This environmental heterogeneity is reflected in differences of tree-ring development, as well as its relation with precipitation regime and disturbance (21, 22). It could be hypothesized, therefore, that the CMD landform variability would represent a factor modulating multi-stemmed *N. flexuosa* tree productivity, as well as stem-cutting cycle periods. Studying how topo-edaphic characteristics influence multi-stemmed algarrobo growth dynamics can enhance CMD silvicultural management by identifying landforms that can sustain intense forest exploitation without ecological imbalance. For this reason, this research analyzed stem growth rates of several multi-stemmed *N. flexuosa* trees distributed along three different landform units, aiming to reconstruct diametric cumulative increment, time (expressed in years) until forest products can be obtained, and biological rotation age according to variations in topography and soil characteristics of the CMD algarrobo forest stands.

## MATERIALS AND METHODS

In this study, we examined three sites within the CMD located in the Province of Mendoza, Argentina, representing distinct landform units (figure 1). Along these districts, the algarrobo woodlands grow under semi-arid climatic conditions, with mean annual precipitation of 155 mm, and large variability in temperature at both daily and seasonal timescales (2). Vegetation consists of the typical CMD plant association, with *N. flexousa* dominating the tree layer, occasionally accompanied by *Geoffroea decorticans*, along with shrub presence of *Larrea divaricata* Cav., (Hook. And Arn.), *Atamisquea emarginata* Miers ex Hook. & Arn and *Bulnesia retama* (Hook.) Griseb (28).



Monthly total precipitation (continuous line) and monthly air temperature (dashed line) data belong to the Encón climatic station (32°15' S, 67°47' W), covering the 1971-1987 period. RI, PR, and ID represent River, Paleo-river, and Inter-dune valley units, respectively. Los datos de precipitación media mensual (línea continua) y temperatura promedio mensual (línea puntuada) pertenecen a la estación meteorológica de Encón (32°15' S, 67°47' W), para el período 1971-1987. RI, PR, e ID representan respectivamente las unidades Riparia, Paleo-riparia, y de Valle inter-medanoso.

Sites were selected based on prior geomorphological classification (23). In this study, River, Paleo-river, and Inter-dune valley landform units were examined, differing in topographic characteristics and edaphic features (table 1). The selected algarrobo woodlands were in relative proximity (within 50 km), allowing us to assume a common climatic influence upon the species stem growth. Soil types varied among the examined stands, with sandy soils in the River and Interdune valley units and loamy soils in the Paleo-river site (table 1). On the other hand, soil permeability differed among the selected woodlands, showing high to moderate and moderate-hydrophobic characteristics (table 1).

**Table 1.** Geographical and environmental characteristics of the sampled sites.**Tabla 1.** Características geográficas y ambientales de los sitios de muestreo.

Landform Unit	Altitude (m a. s. l.)	Latitude (°S)	Longitude (°W)	Soil texture	Soil permeability
River	534	32°06'26,6''	68°07'26,4''	Sandy	Moderate- hydrophobic
Paleo-river	497	32°18'51,9''	67°38'18,1''	Loamy	Moderate
Inter-dune valley	511	32°19'17,1''	67°42'17,0''	Sandy	High

m a. s. l.: meters above sea level. m a. s. l.: metros sobre el nivel del mar.

> At each site, 1-2 rectangular plots of  $1.000m^2$  (50m x 20m) were established to examine the algarrobo forest dynamics in terms of tree radial growth relation with rainfall and its response to thinning, results already published (21, 22). The examined stands exhibited variations in structural characteristics, including differences in multi-stemmed tree density, mean basal stem diameter, and tree height, whereas the number of stems per tree was similar among the selected algarrobo forests (table 2).

> > **Table 2.** Stand structural characteristics of the sampled sites.**Tabla 2.** Características estructurales de los sitios muestreados.

Landform Unit	Density (ms/ha)	Dbase (cm)	H (m)	N° stems/tree
River	60	14,9	4,8	3,8
Paleo-river	110	13,8	5.3	3,2
Inter-dune valley	70	17,2	7,1	3,7

For each multi-stemmed tree, two samples were extracted with a gas-powered drill (TED\_262R, Tanaka Kogyo Co. Ltd, Chiba, Japan), at approximately 50 cm above the ground from the stem with the highest basal diameter. Samples were air-drained, mounted on a wooden support, and polished with progressively finer sandpaper to highlight wood anatomy. Rings were dated following the Schulman criteria for the Southern Hemisphere tree species (25). Tree-ring widths were measured from pith to bark with a 0.01 mm resolution Velmex table connected to a computer. The correct dating of these measurements was statistically validated with COFECHA software (13). Then, we estimated individual tree diametric increments by averaging ring measurements from samples belonging to each algarrobo tree and multiplying these values by two. Finally, we calculated the cumulative

ms: multi-stemmed tree. Dbase: mean multi-stemmed tree basal diameter; H: mean multi-stemmed tree height.

ms: árbol multi-fustal. Dbase: diámetro basal promedio del árbol multi-fustal. H: altura promedio del árbol multi-fustal. diametric increment (CDI) as the sum of the current annual increments in diameter for each biological year. This information allowed us to estimate the time, in absolute tree age, required for the stem to reach the minimum diameter corresponding to forest products in CMD algarrobo woodlands, based on available information for the studied area (poles: stem  $\emptyset$ = 10 cm) (19). When the core center was not reached, tree age was estimated through a geometric method (10). We used ANOVA with post-hoc Fisher's least significant difference test through the Infostat software (9) to compare tree age data across landform units. Five trees were excluded from this analysis since they did not reach minimum CDI values of 10 cm (figure 2).



Dashed horizontal lines show the minimum stem ø = 10 cm. / Las líneas puntuadas indican el ø mínimo del fuste = 10 cm.

Figure 2. Cumulative diameter increment for the River, Paleo-river, and Inter-dune valley landform units.Figura 2. Crecimiento diamétrico cumulativo para las unidades de paisaje de Ripario, Paleo-ripario, y valle inter-medanoso.

Biological rotation age of the sampled algarrobo trees was estimated considering raw ring widths converted to basal area increment (BAI). First, tree-ring measurements from the two radii, corresponding to the two samples extracted for each tree, were averaged at tree level to obtain individual ring-width chronologies. Then, the AGE routine included in the DPL software (Dendrochronological Program Library) (14) was used to estimate current (CBAI) and mean (MBAI) basal area, according to the following equation:

**CBAI**<sub>t</sub> = 
$$\pi * (\mathbf{r}_{t}^2 - \mathbf{r}_{t-1}^2)$$
 and **MBAI**<sub>t</sub> =  $\pi * (\mathbf{r}_{t}^2)/t$ 

where:

 $r_t$  represents ring width in year *t*. Biological rotation age (cutting cycle period) was determined at the age when MBAI and CBAI intersected (5).

## RESULTS

Tree-ring width chronologies were constructed using 36 samples corresponding to 24 *N. flexuosa* multi-stemmed trees. The relatively low number of selected trees at River and Inter-dune valley depended on landowner sampling permission. Multi-stemmed algarrobo site chronologies spanned from 44 (Paleo-river unit) to 96 years (Inter-dune valley unit). Mean annual radial growth rates oscillated between 1.27 (Inter-dune valley unit) and 1.77 mm/yr (Paleo-river unit). Mean correlation values (MC) between individual tree-ring chronologies ranked from 0.446 (Inter-dune valley unit) to 0.490 (Paleo-river unit), with all values being significant at p < 0.05 (table 3, page 31).

CDI analysis revealed that multi-stemmed trees achieved, on average, a minimum diameter of 10 cm at different ages: 37 years for River unit, 24 years for Paleo-river environment, and 44 years for Inter-dune valley landform (figure 2).

<b>Table 3.</b> Characteristics of the tree-ring chronologies.
Fabla 3. Características de las cronologías de ancho de anillo

Landform Unit	N	Period	RW (mm)	МС
River	6 (9)	1954-2011	1.38	0.463
Paleo-river	11 (15)	1967-2010	1.77	0.490
Inter-dune valley	7 (12)	1916-2011	1.27	0.446

N = Number of sampled trees per site (in parenthesis: total number of dendrochronological samples); Period = time range of the sampled cores; RW = mean ring-width value; MC = mean correlation between individual tree-ring series at each stand.

N = número de los árboles muestreados (entre paréntesis: número total de las muestras dendrocronológicas); Period = intervalo temporal de las muestras; RW = valor de ancho de anillo promedio; MC = correlación promedio entre las series dendrocronológicas individuales para cada sitio.

Statistically significant differences emerged regarding tree age corresponding to CDI  $\phi$ = 10 cm among Paleo-river units on one side, and River and Inter-dune valley landforms on the other (one-way ANOVA, F = 5.40, *p* = 0.0161, *df* = 2, n = 19; figure 3).



**Figure 3.** Box and whiskers plot for tree age corresponding to the minimum harvestable diameter of 10 cm across landform units.

**Figura 3.** Gráficos de caja-bigote para los valores del diámetro mínimo maderable de 10 cm según la unidad de paisaje.

Landform heterogeneity was reflected in biological rotation age among the selected units. In this sense, the cutting cycle period occurred at ages 53 and 41 years at River and Paleo-river units, corresponding to stem diameters of 19 and 18 cm, respectively. In contrast, at the Inter-dune valley landform, the culmination age exceeded 96 years (stem  $\phi$  = 24.5 cm), corresponding to the age of the two oldest sampled trees (figure 4, page 32).

interquartile distance (ID 25% above and below the median). The median is shown as a black horizontal bar in the boxes. Whiskers represent values reaching 1.5 times the IDs. Different letters indicate significant differences at p < 0.05. Cada caja muestra los valores comprendidos en la distancia de un intercuartil (ID 25% por encima y por debajo del promedio). El promedio se muestra como línea negra. Los bigotes corresponden a un valor de 1,5 veces la distancia de un ID y se muestran como líneas negras. Letras diferentes indican diferencias significativas por *p* < 0,05.

Each box shows the

values within one



Coloured areas show the number of trees used for estimating MBAI and CBAI. Las áreas coloreadas indican el número de árboles utilizados en el cálculo de MBAI y CBAI.

Figure 4. Mean (MBAI) and current (CBAI) annual basal area increments in relation to tree ages across landform units.Figura 4. Incrementos anuales en área basal promedio (MBAI) y corriente (CBAI) en relación a la edad del árbol según la unidad de paisaje.

## DISCUSSION

In this study, we employed dendrochronological methods to reconstruct growth dynamics of multi-stemmed *N. flexuosa* across three distinct landform units within the CMD. This research provides pioneering evidence of how topographical and edaphic factors influence growth rates of multi-stemmed algarrobo trees in the studied area. Our investigation delivered novel insights concerning the most important forest resource that sustains local communities residing in arid central-west Argentina. The low number of examined multi-stemmed *N. flexuosa* individuals at two sites is not considered an analytic limitation, aimed to explore for the first time tree growth form-dependent dynamics at regional scale. Future studies increasing site repetitions as well total sampled tree number could validate our findings.

Our analyses demonstrated that landform variations were mirrored by growth rates of multi-stemmed algarrobo trees, evidenced by differences in the time required to attain harvestable stem diameters, and the length of the cutting cycle period. Previous dendrochronological assessments suggested that soil attributes and environmental heterogeneity play pivotal roles in shaping moisture availability in the CMD districts, thereby influencing growth dynamics of algarrobo woodlands (21). In this sense, it is worth mentioning that *N. flexuosa* exhibits characteristics of a facultative phreatophyte species, with distinctive root architecture that enables algarrobo trees to access deep phreatic water while also exploring shallow soil horizons (12). This trait allows N. flexuosa trees to use both precipitation and groundwater for tree growth (15). Across the woodlands examined in our study, the phreatic water table consistently lies at similar depths, approximately between 4.5 and 6 meters (26), suggesting that factors beyond water table accessibility, likely soil characteristics and impact on water retention of superficial horizons, probably contributed to the observed variations in tree growth. In arid ecosystems like the CMD, sandy soils tend to facilitate deep water infiltration, whereas loamy edaphic layers promote water retention in superficial horizons, resulting in enhanced stem growth (17, 18, 21). Moreover, soil permeability directly influences water storage capacity, with moderately permeable soils exhibiting better precipitation retention compared to those with higher permeability (18, 21). Both of these edaphic attributes may collectively account for our research outcomes.

Previous analyses have indicated that the biological rotation age of *N. flexuosa* is growth-form dependent, with multi-stemmed trees reaching the cutting cycle period at younger age than their one-stemmed counterparts (4). According to Alvarez *et al.* (2011), multi-stemmed algarrobo trees could be harvested at approximately 80 years of age.
However, our study underscores that landform clearly modulated the cutting cycle period. Our findings revealed different values for biological rotation age across the examined stand. Regarding the discrepancy observed for similar environments, that is, between the Inter-dune valley stand analyzed in this research and the woodland examined by Alvarez *et al.* (2011), belonging to the same landform unit, it could be hypothesized that variations in topographical features, such as the presence (4) or absence (our study) of a complete dune system surrounding the Inter-dune valley site, may affect runoff processes and hence water availability. This landform variability could potentially account for this particular divergent result, although further investigations are warranted to explore the specific topographic influence at a smaller spatial scale within the Inter-dune valley unit, to address this difference more comprehensively.

At local scale, our results indicated that wood extraction from multi-stemmed *N. flexuosa* in CMD districts should consider landform heterogeneity. Specifically, while multi-stemmed algarrobo trees in loamy Paleo-river soil and, to a lesser extent, River units could be managed for pole production, pruning should be avoided in Inter-dune valley *N. flexuosa* forest stands. This recommendation is grounded in the exceptionally extended time required to reach biological rotation age in this landform, which is approximately 100 years. Additionally, the slow radial growth rates in the Inter dune valley environment translated to an extended time window for obtaining forest products used by local inhabitants in the studied area.

## CONCLUSIONS

This research sheds crucial light on a strategically vital forest resource for the communities residing in the CMD area. These findings should be integrated with the development of future forest management strategies for the semi-arid *N. flexuosa* woodlands located in the central-western region of Argentina. On a broader scale, our study underscores the paramount importance of considering landform heterogeneity in the management of desert resprouting woody vegetation.

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# Rural abandonment and its drivers in an irrigated area of Mendoza (Argentina)

# Abandono rural y sus causas en un área irrigada de Mendoza (Argentina)

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#### ABSTRACT

Rural abandonment is a global phenomenon promoted by biophysical, socio-economic, and socio-productive causes, leading to the disappearance of traditional agricultural practices and serious impacts on food security and local livelihoods. This phenomenon is more complex in drylands since the lost of productive land is unlikely to be recovered due to the limited availability of water resources. This study aimed to identify abandoned agricultural lands in a sector located east of the northern oasis of Mendoza (Argentina) and determine the main driving forces leading this process. The interdisciplinary perspective employed included the Normalized Difference Vegetation Index (NDVI) difference technique implemented on Landsat images, the boosted regression trees analysis of spatially explicit drivers, and a digital survey providing perception assessments from local producers and their technical advisors. Abandoned agricultural land has increased by 92% between 2002 and 2020, being accessibility, crop type, vulnerable living conditions of the local population, availability of irrigation water and labor, and the lack of profitability, the main drivers identified by both sources of information (spatial model and social perception). The proposed approach contributes to monitore productive resources and land-use planning with a holistic and long-term vision.

# Keywords

agricultural land abandonment • NDVI difference technique • spatial modeling • social actors' perception

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#### RESUMEN

El abandono rural es un fenómeno global promovido por causas biofísicas, socioeconómicas y socioproductivas, que conduce a la desaparición de prácticas agrícolas tradicionales y tiene graves impactos en la seguridad alimentaria y los medios de vida locales. Este fenómeno es más complejo en las tierras secas, ya que es poco probable que se recuperen las tierras productivas perdidas debido a la disponibilidad limitada de recursos hídricos. Los objetivos de este estudio son identificar tierras agrícolas abandonadas en un sector del este del oasis norte de Mendoza (Argentina) y determinar las principales causas que conducen este proceso. Se empleó una perspectiva interdisciplinaria, que incluye la implementación de la técnica de diferencia del Índice de Vegetación de Diferencia Normalizada (NDVI) a partir de imágenes Landsat, el análisis de boosted regression trees para evaluar causas espacialmente explícitas y la evaluación de la percepción de los productores locales y sus asesores técnicos mediante una encuesta digital. Las tierras agrícolas abandonadas aumentaron un 92% entre 2002 y 2020 y los principales factores identificados por ambas fuentes de información (modelo espacial y percepción social) fueron la accesibilidad, el tipo de cultivo, las condiciones de vida vulnerables de la población local, la disponibilidad de agua de riego, la disponibilidad de mano de obra y la falta de rentabilidad. El enfoque propuesto busca contribuir al monitoreo de los recursos productivos y a un ordenamiento territorial con una visión holística y de largo plazo.

#### **Palabras clave**

abandono de tierra agrícola • técnica de diferencia del NDVI • modelado espacial • percepción de los actores sociales

# INTRODUCTION

Rural abandonment is a global phenomenon detected in diverse regions, such as Europe, the United States, Asia, and Latin America (40). In general terms, rural abandonment is mainly promoted by three types of causes: biophysical (including factors such as elevation and slope, soil, or climate), socio-economic (including market incentives, migration processes, inefficient technology, land tenure, accessibility or characteristics of the producers) and, thirdly, socio-productive (associated with inadequate agricultural management leading to land degradation, overexploitation or loss of productivity) (35). Abandonment of productive land generally occurs in marginal agricultural areas and is associated with low levels of profitability (3, 40). This affects crop productivity even more and undermines investment, technological advancement, and the achievement of acceptable economic returns deepening the abandonment processes (37). Some authors see this change in agricultural land use as an opportunity for the restoration of ecosystem services and the recovery of biodiversity (19, 40), depending on the level of degradation. However, the social consequences of rural abandonment are extremely negative and include the disappearance of traditional practices, seriously impacting food security and local livelihoods (9, 12).

According to the UNCCD (2014), 52% of the world's agricultural land is moderately or severely affected by land degradation. Based on current consumption trends, it is estimated that agricultural production will need to increase by 70% globally, and 100% in developing countries, to meet population demands in 2050. Paradoxically, in Latin America, 50% of agricultural land could be affected by desertification by the same year (42). In this context and especially in drylands, inadequate management of productive resources puts the producers' subsistence and livelihoods at risk. Drylands account for 47% of the earth's land surface, covering hyper-arid, arid, semi-arid, and dry sub-humid climate regions where 2.6 billion people live (23). In Argentina, 63% of the national territory constitutes drylands (43). In these regions, where different irrigation systems transform arid ecosystems into cultivable irrigated areas known as "oases", water limits agricultural production. This limitation further complicates the phenomenon of rural abandonment. In arid to semi-arid Mendoza (Argentina) with 250 mm average rainfall, irrigation depends on water from snowmelt in the Andes Mountains, which feeds surface and underground water sources. Thus, after years of gaining desert land through costly installations of irrigation systems, productive land is lost with low replacement prospects.

In this context, information systems have become valuable tools for managing datasets and identifying critical areas affected by land degradation processes (37). In particular, rural abandonment involves complex dynamics that, as such, must be addressed from an interdisciplinary perspective, including biophysical and socio-economic aspects. Therefore, the construction of spatial models in geographic information systems (GIS) and their statistical analysis makes it possible to associate local data on the occurrence of rural abandonment with variables that describe the biophysical environment and anthropogenic uses. Moreover, each variable has a relative assigned weight reflecting its influence on the process. This approach has allowed several case studies (12, 30, 33, 40). Given this multicausal problem, producer perspectives are valuable assets for understanding perceived constraints, threats, and opportunities (10, 13). Some studies have explored their perceptions of abandonment and plausible causes (4, 24, 31), while others have integrated quantitative data derived from spatial analysis with qualitative data derived from social studies to analyze landscape changes (14, 16, 32). Sustainability demands combining knowledge from different sources to achieve deeper understanding of any given problem.

Rural abandonment has been identified as an environmental problem in the oases of Mendoza but has not yet been thoroughly quantified (1). A previous study identified land-use changes between 2003 and 2019 in a district of the Guaymallén department located in the northern oases (15). Although the authors did identify the category of abandoned land, the employed methodology had limited scalability since it consisted of a visual interpretation of high-definition satellite images available on Google Earth. Both the observer training as well as image availability may hamper analysis extension to the entire oases. Another recent study identified land use and land cover changes between 1986 and 2018 at a larger scale, the Mendoza and Tunuyán River Basins (36). In this case, the methodology enabled an extensive analysis but did not explicitly identify the category of abandoned land. Regarding causes of productive land abandonment, Guida-Johnson et al. (2020) investigated the perception of local actors but did not intend to perform spatial modeling. Our study complements previous work and deepens the level of analysis. Rural abandonment is a complex process that can be addressed, quantified, and expressed spatially through an appropriate methodology. In this sense, detecting crop abandonment and investigating possible causes will significantly contribute to understanding rural abandonment dynamics at a local scale. Our objectives were: (1) to identify abandoned land in an irrigated area in the north of Mendoza province between 2002 and 2020 and (2) to determine the main biophysical, socio-economic, and socio-productive drivers considering spatial modeling and social actors perception.

## **MATERIALS AND METHODS**

## Study area

The productive activities in Mendoza province (Argentina) are concentrated in three oases: north (irrigated by the Mendoza and lower Tunuyán rivers), center or Valle de Uco (irrigated by the upper Tunuyán river), and south (irrigated by the Diamante and Atuel rivers). In particular, the northern oasis can be divided into two zones with distinctive characteristics: east and center. The study area was the department of General San Martín (figure 1, page 38), good representative of the process of rural abandonment in eastern Mendoza. Located in the Mendoza plain, the department has an area of 1,507 km<sup>2</sup>, with its capital city located 43 km from the provincial capital. San Martín is characterized by a great diversity of soils and subsoil water availability, which makes it suitable for crop development. The predominant natural vegetation is xeric, psammophilous, or halophilous shrub steppe, scrubland, and some small relicts of forest (5, 44). Land degradation processes affect the entire area and accelerate the natural phenomena of water and wind erosion (2). According to the latest census (18), the department has almost 140,000 inhabitants and a population density of 92.8 inhabitants/km<sup>2</sup>, most of them concentrated in the capital city, and the rest distributed in small or medium-sized towns surrounding the main city.

Approximately 60% of the population over 5 years of age attended some educational establishment (18). Considering population density, health and service infrastructure, and logistics, San Martín is the main department of eastern Mendoza. However, inequalities between urban and rural populations regarding access to social infrastructure (such as drinking water, public transportation, sewage, natural gas, electricity, or waste collection) are profound. Considered one of the main grape-growing areas in the country, the main economic activity is viticulture, followed by olive and fruit production, as well as horticulture. The main economic units are associated industries and family businesses. Historically, urban expansion over rural areas has been carried out in a chaotic or disorderly manner, transforming lands with high agricultural potential and accentuating territorial imbalances.



Figure 1. Location of the study area. Figura 1. Localización del área de estudio.

#### **Detecting rural abandonment**

A land-use land-cover (LULC) map was used as a reference and satellite images were analyzed by the Normalized Difference Vegetation Index (NDVI) difference technique to detect changes (22, 27, 41). The production map of Mendoza province constituted our reference. The map, prepared and provided by the Institute for Rural Development [Instituto de Desarrollo Rural, IDR], was created by visual interpretation of high-resolution images available on Google Earth. LULC categories were: abandoned land, non-productive tree cover, fruit trees, vegetables, natural vegetation, olives, pastures, other uses (including swimming pools, and campsites), urban, vines, and vines with olives.

The NDVI difference technique is one algebraic change detection technique, simple and easy to implement and interpret (25). Considering the reference map, the study period was defined between 2002 and 2020. This study used two atmospherically corrected satellite images transformed to surface reflectance: one from the Landsat 5 TM sensor obtained in 2002 and one from Landsat 8 OLI obtained in 2020. A preliminary assessment of monthly NDVI variation associated with different LULC categories allowed image date selection. In

December 2019, 200 field points were surveyed. The largest differences between cultivated and abandoned land were recorded in February when both selected satellite images were obtained. The NDVI was calculated for each scene from the reflectance values in the near-infrared (NIR) and red (R) regions, following Eq. (1).

$$NDVI = \frac{NIR - R}{NIR + R} \tag{1}$$

The NDVI difference was then calculated ( $\Delta$ NDVI) (Eq. 2). In the resulting image, pixels were divided into no-change pixels with values around zero and change pixels with values sufficiently far from zero (positive and negative) (29).

$$\Delta NDVI = NDVIt2 - NDVIt1 \tag{2}$$

Where NDVIt2 is calculated for the Landsat scene obtained in 2020 and NDVIt1 is calculated for the Landsat scene obtained in 2002. One key issue is threshold definition, *i.e.*, which value change is to be considered significant. In general, thresholds associated with different numbers of standard deviations for NDVI values ( $\sigma$ ) are tested, visually choosing the one that best detects changes (27, 29, 41). In this study, four thresholds were tested:  $1\sigma$ ,  $1.5\sigma$ ,  $2\sigma$ , and the one determined by the Jenks natural breaks classification method.

The changed pixels interpretation was based on comparisons with the reference map considering whether  $\Delta$ NDVI was positive or negative. Detection of abandoned rural land was performed as detailed in Table 1: no-change pixels preserve their category, whereas change pixels are converted into abandoned land (AL), cultivated land (CL) or not cultivated land (NCL). A rural abandonment map was produced for each  $\Delta$ NDVI threshold. Original LULC categories from the reference map were regrouped into abandoned land, cultivated land (including fruit trees, vegetables, olives, pastures, vines, and vines with olives) and not cultivated land (non-productive tree cover, natural vegetation, other uses, and urban).

**Table 1.** Criteria to detect abandoned agricultural land based on NDVI differencetechnique and a reference map.

**Tabla 1.** Criterios para detectar tierra agrícola abandonada basándose en la técnica dediferencia del NDVI y un mapa de referencia.

			Rural abandonment m	ap	
Reference map		No-change pixels	Change pixels		
		ΔNDVI near 0	Significantly (-) $\Delta$ NDVI	Significantly (+) $\Delta$ NDVI	
Abandoned land	AL	AL	AL	CL	
Fruit crops					
Vegetable crops					
Olive crops	CI		AL	CL	
Pasture	CL	CL			
Vine crops					
Vine and olive crops					
Natural vegetation		NCL	NCL	CL	
Unproductive tree cover	NCI				
Other land uses	INCL	NCL	NCL	NCL	
Urban					

The  $\Delta$ NDVI thresholds were assessed by validating the rural abandonment maps from ground truth data. Three campaigns surveying GPS points in the study area were conducted in December 2019, March 2021, and June 2021, collecting 537 sample points. An error matrix was constructed for each map with these data, while descriptive statistics were calculated. Total accuracy represents the proportion of reference points correctly classified. Producer's accuracy indicates the probability of a reference point correctly classified, *i.e.* how properly a particular area can be classified. Finaly, user's accuracy estimates the probability of a point on the map representing the class on the ground (6). A rural abandonment map was selected based on statistics, and the area associated with each category was calculated.

#### Identifying rural abandonment drivers

Rural abandonment drivers were identified and assessed using two sources of information: a spatial model, and social actors perceptions. A GIS model weighed the relative contribution of spatially explicit driving forces (12, 33). To that end, 2000 points were randomly plotted on the selected rural abandonment map to gather data and run a machine learning algorithm. For each point, the response variable was presence or absence of rural abandonment. Predictor variables were compiled from available geo-referenced secondary data of three types, namely attributes of the biophysical environment, and socio-economic and socio-productive aspects. The former included: (1) elevation obtained from Shuttle Radar Topography Mission (SRTM) data (20); (2) aridity index values, freely available from the Territorial Environmental Information System of Mendoza [Sistema de Información Ambiental Territorial de Mendoza] (38); and (3) frequency of storm damage events recorded during 2002-2017, available at the Directorate of Agriculture and Climatic Contingencies of Mendoza website [Dirección de Agricultura y Contingencias Climáticas de Mendoza] (7). The socio-economic and socio-productive aspects included: (4) the original type of crop obtained from the productive map of Mendoza provided by the Institute for Rural Development; (5) population density, that is the number of inhabitants registered by census unit, (6) the percentage of households with unsatisfied basic needs (UBN), including overcrowded homes, precarious housing, poor sanitary conditions, non schooled children or household head without complete schooling, and (7) percentage of the working-age population who only reached primary education level (*i.e.* those not meeting certain minimum conditions - secondary education - in order to access formal employment), with data obtained from the National Census of Population, Households and Dwellings 2010 [Censo Nacional de Población, Hogares y Viviendas 2010] (17); (8) the Euclidean distance to national and provincial roads available in the Territorial Environmental Information System of Mendoza (39); (9) the Euclidean distance to irrigation network and (10) to groundwater extraction wells and (11) the density of canals and (12) wells, with data provided by the General Department of Irrigation of Mendoza [Departamento General de Irrigación (DGI)]. The values of all the predictor variables were recorded at each random point.

The data were analyzed by boosted regression trees (BRT) (30, 40) through R software (34) using the dismo package version 1.3-5. This approach uses boosting to combine large numbers of relatively simple tree models to optimize predictive performance and identify relevant variables, explaining an observed pattern (11). Model parametrization was performed with tree complexity, which controls the number of fitted interactions; the learning rate, which determines tree contribution to the model; and the bag fraction, which defines the proportion of data selected at each step (11). For this, different combinations of tree complexity (1 to 9), learning rate (0.01, 0.005, 0.001, and 0.0005), and bag fraction (0.5 and 0.75) were tested (21, 28). Model selecting criteria intended to maximize the explained deviance, *i.e.* variability explained by the model, and minimized the difference between the training data AUC score and the cross-validated AUC score, to reduce overfitting (8, 11). Variables with relative influence under the expected by chance (100 divided by the number of explanatory variables) were considered non-relevant for interpretation (30). Considering the 12 variables tested, in this case, the relative influence threshold was 8.33%.

Regarding social perception, questions addressed abandonment causes and the areas and years in which the phenomenon was evident in the department of San Martín. Focus was placed on producers and their technical advisors. For this purpose, a pilot study was conducted and a qualitative online survey was administered through a Google Form. A total of 50 producers and/or advisors from San Martín were contacted through different social networks. The survey consisted of five open and closed questions on (1) major drivers of rural abandonment in the department of San Martín (considering depth of the water table, crop type, population density, access to road network, and water and labor availability), (2) additional environmental or socio-economic and cultural factors impacting the abandonment process; (3) localities with abandoned agricultural plots, (4) localities in which this process occurred with greater intensity between 2002 and 2020, and (5) the time in which the process of abandonment was intensified (dividing the period under analysis into three equal parts: 2002-2008, 2009-2014 or 2015-2020). The surveys were analyzed using descriptive statistics and compared with the spatial analysis.

#### RESULTS

Rural abandonment maps derived from the four  $\Delta$ NDVI thresholds found values of overall accuracy between 0.70 and 0.77 (table 2). The differences between these values lay in the accuracy with which abandoned and cultivated land was identified. At one extreme, the map derived from the threshold given by the Jenks Natural Breaks function correctly classified 63% of sample points identified as abandoned land (producer's accuracy 0.63), while it misclassified 29% of sample points identified as cultivated land (producer's accuracy 0.71). On the opposite, the map derived from the threshold defined as  $2\sigma$ , correctly classified 36% of sample points identified as abandoned land (producer's accuracy 0.36), with 95% of the sample points identified as cultivated land correctly classified (producer's accuracy 0.95). Considering this paper aims to detect rural abandonment in the study area, the Jenks Natural Breaks function was selected as  $\Delta$ NDVI threshold.

**Table 2.** Total, producer's, and user's accuracy associated with the four rural abandonment maps derived from assessed  $\Delta$ NDVI thresholds: Jenks Natural Breaks (NB), 1 $\sigma$ , 1.5 $\sigma$  y 2 $\sigma$ .

**Tabla 2.** Exactitud total, del productor y del usuario asociadas a los cuatro mapas de abandono rural generados a partir de los umbrales de  $\Delta$ NDVI analizados: Quiebres Naturales de Jenks (NB), 1 $\sigma$ , 1.5 $\sigma$  y 2 $\sigma$ .

Producer's accuracy					User's accuracy				
ΔNDVI threshold	Total accuracy	Abandoned land	Cultivated land	Not cultivated land	Abandoned land	Cultivated land	Not cultivated land		
NB	0.70	0.63	0.71	0.84	0.50	0.82	0.79		
1σ	0.75	0.50	0.86	0.84	0.61	0.80	0.81		
1.5σ	0.77	0.42	0.93	0.84	0.71	0.78	0.79		
2σ	0.77	0.36	0.95	0.84	0.74	0.77	0.79		

According to the selected rural abandonment map, abandoned land increased by 92% during the study period. This was calculated considering the 13,492 ha in the reference map which was incremented to 25,869 ha in the 2020 rural abandonment map (table 3, page 42). Four areas concentrate abandoned land in north, central-west, central-east, and south San Martín (figure 2, page 42).

**Table 3.** Area (ha) for each land use category in the referencemap (Mendoza productive map elaborated by IDR) and in the rural abandonment map(detected with NDVI difference technique). Variations are indicated in %.

**Tabla 3.** Área (ha) para cada categoría de uso de suelo en el mapa de referencia (mapa productivo de Mendoza elaborado por el IDR) y en el mapa de abandono rural (detectado con la técnica de diferencia del NDVI). Se indican los cambios (%) entre ellos.

Land use category	Reference map	Rural abandonment map	Changes (%)
Abandoned land	13,492	25,869	+92
Cultivated land	42,397	30,169	-29
Non-cultivated land	11,941	11,791	-1



**Figure 2.** Rural abandonment map for the study area in 2020. **Figura 2.** Mapa de abandono rural para el área de estudio en 2020.

The spatial analysis indicated that the Euclidean distance to national and provincial roads, crop type, percentage of households with unsatisfied basic needs (UBN), distance to wells and irrigation canals, and density of canals and groundwater extraction wells mostly explained the spatial distribution of abandoned land in the study area (table 4, page 43). The model explained 24.62% of the observed variability, probably due to the complexity of the analyzed problem. Distance to roads indicates accessibility to agricultural plots and accounts for management and harvesting difficulties. The crop would be linked to production profitability and local, national, and international market incentives and disincentives. The percentage of UBN households provides idea of structural poverty conditions in which some producers or their employees live (*e.g.*, contractors who work and live on the farm and receive a share of production profit), as well as households of local people working in companies, especially during harvest time. Finally, the four variables associated with irrigation canals and groundwater extraction wells account for irrigation availability. It should be noted that, geographically, San Martín largely covers the eastern boundary of the northern oasis of Mendoza.

**Table 4.** Relative influence of explanatory variables for a BRT model developedwith a tree complexity of 3, a learning rate of 0.005, and a bag fraction of 0.5.Explained deviance was 24.62%

**Tabla 4.** Influencia relativa de las variables explicativas para el modelo de BRT desarrollado con complejidad de árbol de 3, tasa de aprendizaje de 0,005 y *bag fraction* de 0,5. La devianza explicada fue del 24,62%.

Explanatory variable	<b>Relative influence</b>
Distance to roads	14.03
Type of crop	13.41
% Households with unsatisfied basic needs	12.86
Distance to wells	11.34
Channel density	10.78
Well density	10.63
Distance to channels	8.80
Elevation	6.32
Storm damage	4.29
Population density	4.05
% Working-age population with only primary education	3.35
Aridity index	0.15

Twenty-seven out of 50 producers and/or advisors completed the survey. Concerning the most important drivers of land abandonment in San Martín, 85% of the respondents pointed to irrigation availability, 81% to labor availability, and 67% to crop type, in agreement with the spatial analysis. When asked what other biophysical or socio-economic and cultural factors impact the abandonment process, 63% of the respondents pointed to a lack of profitability. In addition, 15% indicated difficulties associated with generational changes (in the words of the interviewees "(...) aging of the vine-growers and lack of desire of the new generations to continue with the activity"), 15% related to the regional economy and another 15% associated with insecurity (robberies, vandalism, and theft). When respondents were asked to indicate which districts in San Martín currently had abandoned plots, 74% indicated Montecaseros, 56% El Divisadero and 48% El Ramblón. Particularly, regarding the process between 2002 and 2020, 77% of respondents indicated Montecaseros, 58% El Ramblón and 46% El Divisadero. Data from spatial analysis and surveys point to Montecaseros and El Ramblón as two districts with intermediate evidence of rural abandonment (figure 3, page 44). Nueva California district was identified by some respondents, but showed strong evidence of rural abandonment in the spatial analysis. Conversely, El Divisadero was noted by numerous respondents but showed less evidence of abandonment in the spatial analysis. It is worth noting that southern districts, where a large amount of abandoned land was detected according to the rural abandonment map, were slightly mentioned by respondents. Finally, 65% of respondents indicated that rural abandonment has intensified in the last period, 2015-2020.



Figure 3. Location of rural abandonment in the study area according to remote sensing and respondents perception.

Figura 3. Localización del abandono rural en el área de estudio según la teledetección y la percepción de los encuestados.

# DISCUSSION

Rural abandonment is a global phenomenon associated with biophysical, socio-economic, and socio-productive causes (35, 40). In this respect, our results are in line with other studies associating this phenomenon with soil characteristics, topography and accessibility (12), the presence of soils with agricultural potential and agricultural subsidies (9), topography, accessibility, tractor and cropland density (30), crop yields and accessibility (33), physical environmental conditions, accessibility and global market pressures (3), equipment and materials costs and property taxes (24) or lack of state support, lack of occupation, demotivation of young people and lack of educational centers (31). In the study area, coincidences were found for both sources of information examined and also with these previous studies. According to the spatial analysis, rural abandonment would be associated with accessibility, type of crop, vulnerable living conditions of the local population, and availability of irrigation water. Whereas, the study of perception pointed to irrigation water and labor availability, type of crop, and the lack of profitability as major causes. Crop type is related to both profitability and agricultural management.

Regarding the territorial dimension, the NDVI difference technique showed that abandonment had increased in the analysed period at an alarming rate. All respondents acknowledged rural abandonment in the study area, recently intensified and associated with different districts in the department. The spatial pattern of rural abandonment showed some divergences between satellite image analysis and respondent perceptions. In some cases, both sources of information associated the process of agricultural land abandonment with different districts. It is worth noting that the sector where the greatest divergences were found (the south) overlaps with a zone defined by intense urban and industrial land uses in a matrix of cultivated areas. Two main, fast-growing cities in the department are located there: Ciudad de San Martín and Palmira. Moreover, the sector is crossed by National Route 7, one important commercial and tourist highway in the country, connecting the Atlantic and the Pacific oceans and integrating the Central Bioceanic Corridor. We could hypothesize that the development of this area interferes with the perception of rural abandonment. Although remote sensing analysis showed moderate to high levels of abandonment in south San Martín, respondents associated the phenomenon with locations farther away from these urban centers.

Remotely sensed information provides up-to-date databases supporting land management. Regarding rural abandonment, early detection of this type of change is particularly relevant to prevent and mitigate land degradation, which becomes the most efficient option. Thus, starting from an updated LULC map, detecting and monitoring changes may simplify and speed up the analysis (26). The methodology of LULC change detection used in this study resulted in a rapid application and easy implementation and interpretation. Compared with a previous study detecting abandoned land in a sector of the north oases (15), the implemented methodology could enable a larger-scale monitoring of the process. Therefore, the provided information could be useful for local authorities (among other social actors) for detecting the most compromised districts, speeding up the diagnosis, an invaluable phase of land use policy-making.

Rural abandonment becomes more complex in drylands, naturally vulnerable to degradation processes. In these regions, abandonment implies a loss of productive land and the entire associated technological system that is unlikely to be replaced due to the limited availability of water resources. Food security and local livelihoods are seriously at risk (9, 12). In this context, the complementation of quantitative data on landscape change with the perception of the local community is highly relevant and leads to understanding views, beliefs, and decisions of different social actors. This, in turn, is key to formulating public policies addressing processes of landscape change (14). On the other hand, studying the perception of social actors promotes a better understanding and interpretation of changes, enriches the explanation of driving factors, and leads to a comprehension of the complexity of the socio-economic and cultural context in which they occur (14, 16, 32). The incorporation of local community knowledge in decision-making contributes to strengthening alliances between the social actors involved, optimizing land-use planning processes with a holistic and long-term vision in pursuit of sustainable use of productive resources.

# **CONCLUSIONS**

In the department of San Martín, located east of the northern oasis of Mendoza (Argentina), cropland abandonment increased by 92% between 2002 and 2020. As in other places around the world, this phenomenon is multi-causal. Accessibility, crop type, vulnerable living conditions of the local population, availability of irrigation water and labor, and lack of profitability constitute main drivers for land abandonment. Addressing this complex problem needs to integrate different sources of information and skills to develop public land-use planning policies that promote sustainable local development. The loss of productive land in drylands threatens food security, determining the urgency of preserving existing cultural and productive resources. The multidisciplinary approach implemented in this study used different methodologies and sources of information enabling a better understanding of the environmental problem. Moreover, it allowed defining differences between the perception of local social actors and the results derived from spatial analysis tools. Understanding and managing territorial complexity from an interdisciplinary perspective is essential to contributing knowledge to the local social reality.

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# Agrochemical characterization of *Vitis labrusca* grape pomace and its effect on a soil-plant system

# Caracterización agroquímica del orujo de uva de *Vitis labrusca* y su efecto sobre el sistema suelo-planta

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# ABSTRACT

This study characterized the agrochemical properties of *V. labrusca* grape pomace (GP) and evaluated the effect on the rhizobacteria *Azospirillum brasiliense* and horticultural crops, determining safety as fertilizer and/or mulching. Two first bioassays were performed with the GP at different concentrations evaluating toxicity on *A. brasiliense*, and on tomato and lettuce seeds. A third bioassay evaluated GP mulching effects on tomato and lettuce plants growing with amounts varying between 20 and 80 t ha<sup>-1</sup>. Agrochemical characterization showed that GP is rich in potassium and phosphorus, with a low content of Na<sup>+</sup> salts (SAR < 15). The GP at 2.5% (w v<sup>-1</sup>) significantly increased survival rates of N<sub>2</sub>-fixing rhizobacteria. Results on seed germination revealed lettuce was more susceptible to increasing GP concentrations. The application of 20 t ha<sup>-1</sup> of GP in greenhouse experiments increased lettuce and tomato root biomass. Furthermore, the aerial part of tomato showed no toxicity symptoms. These results open the possibility of considering *V. labrusca* GP as mulching without prior treatment in tomato crops.

#### **Keywords**

agrowaste • revalorization • macronutrients • Azospirillum brasiliense • tomato • lettuce

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#### RESUMEN

Este estudio tuvo como objetivo caracterizar las propiedades agroquímicas del orujo de uva (GP) de *V. labrusca* y evaluar su efecto sobre *Azospirillum brasiliense* y cultivos hortícolas para determinar su uso seguro como fertilizante y/o mulching. Los bioensayos se realizaron con el GP a diferentes concentraciones para evaluar su toxicidad sobre *A. brasiliense* y semillas de tomate y lechuga. Para evaluar el efecto de GP sobre el crecimiento de plantas de tomate y lechuga se aplicaron diferentes dosis como mulching en el rango de 20-80 t ha<sup>-1</sup>. La caracterización agroquímica mostró que GP es rico en potasio y fósforo; y tiene bajo contenido de sales de Na<sup>+</sup> (SAR < 15). El GP al 2,5% (p v<sup>-1</sup>) fue responsable de un aumento significativo en la tasa de supervivencia de las rizobacterias fijadoras de N<sub>2</sub>. Los resultados de la germinación de las semillas revelaron que la lechuga fue más susceptible al aumento de las concentraciones de GP. La aplicación de 20 t ha<sup>-1</sup> de GP en los experimentos de invernadero incrementó la biomasa radicular de lechuga y tomate. Además, la parte aérea del tomate no presentó síntomas de toxicidad. Estos resultados abren la posibilidad de considerar el aprovechamiento de *V. labrusca* GP como mulching, sin utilizar ninguna tecnología de tratamiento antes de su aplicación al suelo en cultivos de tomate.

#### **Palabras claves**

residuo agronómico • revalorización • macronutrientes • *Azospirillum brasiliense* • tomate • lechuga

#### INTRODUCTION

The wine industry is one of the oldest and most important in the world. In 2019, wine production was estimated at around 258 Mhl (27). Grapes for winemaking belong to the Genus *Vitis* sp, being *V. vinifera* and *V. labrusca* the two main producing species (20). *V. vinifera* is cultivated in arid soils, poor in organic matter, while the hybrid *V. labrusca* originated in eastern USA, grows in different edaphoclimatic regions and is highly resistant to diseases (18, 24). In South America, mainly in Argentina and Brazil, *V. labrusca* cultivated areas have increased in the last decade (24, 40).

Annually, the wine industry produces 13 x 10<sup>6</sup> t of solid waste or by-products worldwide, with grape pomace (GP) being the most abundant (10-13 Mt per year). In terms of circular economy, GP could generate new products of different values (7, 21), closing the cycle as mulching or organic fertilizer due to its lignocellulosic nature (8). Mulching improves soil structure, keeps moisture, moderates temperature, and increases organic matter (OM), returning soil nutrient cycling environments to pristine soil patterns (25). On the other hand, when the agronomic purpose is to increase crop yield, organic fertilizers of plant origin constitute an interesting agroecological alternative for intensive crops like vegetables, flowers, and orchards (22). Nevertheless, a proper agronomic management program incorporating lignocellulosic by-products, such as GP, needs to characterize the product and establish an appropriate use. Unlike the well-known physicochemical characteristics of V. vinifera GP (6, 16, 34), the relevant agrochemical properties of V. labrusca GP have not been studied yet. Direct incorporation of V. vinifera GP into the soil could negatively affect plant growth due to inhibitory compounds or given competition between soil microorganisms for essential nutrients such as nitrogen (6). However, this study only considers GP physicochemical properties, leaving aside other sensitive parameters in the soil-plant systems.

Biological indicators are living organisms, such as plants, animals, and/or microorganisms exploited to detect toxic substances in terrestrial and aquatic ecosystems (45). The effect of agrowastes on the soil-plant system can be evaluated by lettuce (*Lactuca sativa*) and tomato (*Solanum lycopersicum*) as biological indicators, as well as by the free-soil nitrogen-fixing bacterium *Azospirillum brasiliense* (14, 36, 37, 41). The combination of agrochemical characterization together with biological indicators enables the development of appropriate agronomic management programs (22).

This study characterized the agrochemical properties of *V. labrusca* var. *Isabella* GP and evaluated toxicity effects on *A. brasiliense*, tomato and lettuce determining GP safety as fertilizer or/and mulching.

#### MATERIAL AND METHODS

## Study area and GP sampling

The study considered twenty-four ha of *V. labrusca* var *Isabella* at "Cooperativa de la Costa" wine-cellar located in Berisso, Buenos Aires, Argentina (34°53'22.79" S; 57°49'21.11" W (28). The soil is Rendzic Leptosol (11) with an A-horizon of 20-40 cm and a high content of well-humified organic matter, under which there is an AC or C-horizon constituted by layers of small shells (1). In 2013, the winery produced 350 hL of wine and a total of 12 t of GP, representing 25% of grape total weight (29).

Samplings were carried out in 2015, 2016 and 2017 seasons. Each year, 10 random GP samples were collected immediately after grape pressing and blended to obtain a compound sample. One part of the sample was separated to determine moisture content. The rest was oven-dried at 60°C to constant weight and then divided into sub-samples. These were stored in hermetically sealed in containers until GP physicochemical characterization. Part of fresh material from the 2017 campaign was separated and stored at -20°C for subsequent bioassays.

#### Agrochemical GP characterization

Ashes and OM were determined by calcination (6). Percentage contents of organic carbon (C) and total nitrogen (N) were determined after Walkley and Black (1934) and micro-Kjeldahl methods (5) respectively, and the C/N ratio was calculated. Total phosphorus (P) was determined by the yellow vanadate-molybdate method (19). Extractable and exchange cations were obtained by the saturated paste extraction method and the Ammonium acetate method 1N pH7, respectively (35). Calcium (Ca<sup>+2</sup>) and Magnesium (Mg<sup>+2</sup>) were estimated by complexometry with 0.02 N EDTA; Sodium (Na<sup>+</sup>) and Potassium (K<sup>+</sup>) by flame photometry. Sodium adsorption ratio (SAR) was calculated using the following equation (31) (eq. 1).

The  $Mg^{+2}/K^+$ ,  $Ca^{+2}/Mg^{+2}$ , and  $Ca^{+2}/K^+$  ratios were calculated using the obtained values for individual exchangeable cations (17). Cation exchange capacity (CEC) was determined by steam distillation and percentage base saturation (S) was calculated by eq. 2 (33).

# S (%) = ( $\Sigma$ exchangeable cations/CEC) \*100 (2)

Electric conductivity (EC) and pH values were measured in the aqueous soluble fraction (1:5 ratio) by conductometry and potentiometry, respectively (19). Moisture content (%) was determined by oven-drying samples at 60°C until constant weight and calculating the difference between wet and dry weight. Water-soluble phenols (WSP) were determined as described by Osono and Takeda (2001). Three technical replicates were used to determine each physicochemical parameter. Values were averaged to obtain a unique value for each year. Mean, range and t variance were estimated with three annual values. The classification criterion proposed by Fernández Linares *et al.* (2006) and Havlin *et al.* (1999) allowed interpretation.

# **Bioassays with GP application**

Three bioassays determined GP safety as fertilizer or/and mulching: a) GP toxicity evaluation on a N fixing microorganism (*A. brasiliense*), b) GP toxicity evaluation on germination of tomato and lettuce seeds; and c) GP effect on growth parameters of tomato and lettuce plants.

a)- GP toxicity evaluation on a N fixing microorganisms (A. brasiliense)

GP effects on survival of A. brasiliense CECT 590 T were evaluated according to

Saparrat *et al.* (2010). A bacterial suspension (100  $\mu$ l) was inoculated in 900  $\mu$ l of the GP sterilized by filtration (0.22  $\mu$ m millipore membrane) at different concentrations: 2.5, 3.5, 5, and 10% (w v<sup>-1</sup>). Controls were made using sterile distilled water and 4 replicates per treatment were utilized. Cultures were grown at 28°C and 150 rpm for 24 h. Colony-forming units (CFU mL<sup>-1</sup>) were estimated by the dilution and plating method using the selective medium Congo red. The data were transformed to logarithms and analyzed by ANOVA and a Tukey test (*p* < 0.05).

b)- GP toxicity evaluation on germination of tomato and lettuce seeds

GP phytotoxicity was evaluated after the effects observed on germination and root growth of lettuce and tomato according to Tiquia *et al.* (1996) with modifications. Seeds of each species were placed on filter paper inside Petri dishes (9 mm diameter) in contact with 3 ml of the GP at different concentrations: 2.5, 5, 10, and 20% w v<sup>-1</sup>. Negative and positive controls consisted of paper soaked with sterile distilled water and 1 M of a CuSO<sub>4</sub> solution, respectively. Four replicates per treatment were incubated at 25°C for 7 days in the dark.

Number of germinated seeds and root length (mm) were determined. Relative germination percentage (G) (eq. 3), relative root length (RL) (eq. 4), and germination index (GI) (eq. 5) were calculated using the following equations:

# G (%) = (number of germinated seeds at X GP concentration/ number of seeds germinated in negative control) \*100 (3)

# RL (%) = (mean root length at X GP concentration/ mean root length in negative control) \*100 (4)

$$IG = (G * RL) / 100$$
 (5)

A 2 mm primary root defined seed germination. The data obtained were analyzed by ANOVA and Tukey test (p < 0.05). Percentual data were transformed using arc sen  $\sqrt{p}$ , before statistical analysis.

c) GP effect on growth of tomato and lettuce plants

The effect of GP on lettuce and tomato plants was determined using ten plants per treatment and according to Sampedro *et al.* (2004) with modifications. Seedlings with three expanded leaves were placed in 5 L pots filled with a mixture of soil and sand in a 1:1 ratio (v v<sup>-1</sup>). GP mulching was added to the pots at different doses: 20, 40, and 80 t h<sup>-1</sup> (wet weight equivalent to 60, 120 and 240 g of GP per pot). The plants were grown for 1 month in a greenhouse and periodically watered. Pots without GP were set up as control. The number of expanded leaves per plant was estimated by direct counting. Leaf greenness (GrI) was measured using a portable chlorophyll meter (SPAD-502, Minolta Corp. Japan), randomly selecting three expanded leaves per plant. Total leaf area per plant was captured by photography and analyzed using J image software (38). Plants were harvested and soil adhered to roots was washed with running water. The aerial part of the plants and roots were oven-dried at 60°C to constant weight determining aerial and root dry biomass. The data were analyzed for each plant species comparing dose effects. A Kruskal-Wallis test and a non-parametric multiple contrast as *post hoc* test (*p* < 0.05) analyzed the number of expanded leaves. The remaining data were analyzed by ANOVA and Tukey test (*p* < 0.05).

#### RESULTS

#### Agrochemical GP characterization

The C/N ratio < 25 suggested an equilibrate decomposition of OM. Among extractable cations, GP showed high levels of K<sup>+</sup> and low Na<sup>+</sup>, high CEC (> 40), and a varied macronutrient supply (exchangeable cations, high content of K<sup>+</sup>, followed by Ca<sup>+2</sup>, and low Mg<sup>+2</sup>). The GP exhibited a high base saturation index (S > 40), high EC (> 4 dS m<sup>-1</sup>), extremely acidic pH, and high hygroscopicityy (> 70% humidity, table 1, page 52).

Parameter	Mean	Range	VC <sup>1</sup>
OM (g kg <sup>-1</sup> )	9.3	9.1-9.4	0.2
Ash (%)	7.5	6.4-9.1	18.9
TOC (g kg <sup>-1</sup> )	4.71	3.42-5.51	2.39
TN (g kg <sup>-1</sup> )	0.21	0.2-0.22	0.58
C/N	22.4	16.6-26.3	26.5
TP (g kg <sup>-1</sup> )	3.1	2.9-3.4	8.6
Ext K <sup>+</sup> (g kg <sup>-1</sup> )	37.6	27.7-51.7	33.3
Ext Na <sup>+</sup> (mg kg <sup>-1</sup> )	64	57-75	15.1
Ext Ca <sup>+2</sup> (g kg <sup>-1</sup> )	4.4	3.7-5.0	15.0
Ext Mg <sup>+2</sup> (g kg <sup>-1</sup> )	0.7	0.5-0.8	23.2
SAR <sup>2</sup>	1.3-3	1.1-3- 1.51-3	14.7
Exch K <sup>+</sup> (cmol <sub>c</sub> Kg <sup>-1</sup> )	49.3	43.9-54.1	10.4
Exch Ca <sup>+2</sup> (cmol <sub>c</sub> Kg <sup>-1</sup> )	7.2	6.6-7.6	7.2
Exch Mg <sup>+2</sup> (cmol <sub>c</sub> Kg <sup>-1</sup> )	0.9	0.4-1.7	72.9
Exch Na <sup>+</sup> (cmol <sub>c</sub> Kg <sup>-1</sup> )	0.4	0.4-0.5	21.0
Exch Ca <sup>+2</sup> / Exch Mg <sup>+2</sup>	10.47	4.5-16.5	57.8
Exch K <sup>+</sup> / Exch Ca <sup>+2</sup>	6.9	5.8-7.6	14.2
Exch K <sup>+</sup> / Exch Mg <sup>+2</sup>	75.9	25.8-124.5	65.0
CEC (cmol <sub>c</sub> Kg <sup>-1</sup> )	43.7	41.5-46.7	6.2
S <sup>3</sup> (%)	132.4	125.7-137.7	4.64
EC (dS m <sup>-1</sup> )	5.8	5.4-6.1	6.1
рН	3.7	3.6-3.7	0.7
Moisture (%)	79.2	78.1-80.1	1.3
WSP <sup>4</sup> (mg g <sup>-1</sup> )	74.0	73.4-75.4	1.5

**Table 1.** Physicochemical characteristics of *V. labrusca* var. *isabella* GP.**Tabla 1.** Características fisicoquímicas del OU de *V. labrusca* var. *isabella*.

# GP toxicity evaluation on a N fixing microorganism (A. brasiliense)

GP concentrations showed significant differences in *A. brasiliense* growth (F= 684.50; p < 0.001). No CFU of the diazotrophic bacteria was recovered at 10% GP concentration (figure 1).



Means followed by the same letter are not significantly different (Tukey test, p < 0.05). Los datos son medias de cuatro réplicas ± D.E. Medias seguidas por la misma letra no presentan diferencias significativas (Test de Tukey, p < 0.05).

1, Variance coefficient. 2, Sodium adsorption ratio. 3, Base saturation index. 4, Water-soluble phenols. Extractable (Ext) and interchange (Exch) cations. n= 3 annual samples. 1, Coeficiente de variación. 2, Radio de adsorción de sodio. 3, Índice de saturación de bases. 4, Fenoles hidrosolubles. Cationes extractables (Ext) e intercambiables (Exch). n= 3 muestras anuales.



**Figura 1**. Crecimiento de *A. brasiliense* después de 24 h de incubación en distintas concentraciones de OU (%, p v<sup>-1</sup>).

# GP toxicity evaluation on germination of tomato and lettuce seeds

Significant differences were found in relative germination (G) (F= 86.04; p < 0.001), relative root length (RL) (F= 583.56; p < 0.001) and germination index (GI) (F= 348.69; p < 0.001) when lettuce seeds were incubated at different GP concentrations. The G values at 2.5, 5 and 10% GP concentration equalled or exceeded 90%. The RL and GI, registered values below 50% indicating a toxic effect of GP at all concentrations evaluated (figure 2a, c, e).

For tomato, significant differences were found in G (F= 368.32; p < 0.001), RL (F= 145.96; p < 0.001) and GI (F= 278.02; p < 0.001). The toxic effect in G, RL and GI increased with GP concentration (figure 2b, d, f).



Mean  $\pm$  S.D. of four replicates followed by the same letter are not significantly different (Tukey test, p < 0.05). Los datos son medias de cuatro réplicas  $\pm$  D.E. Medias seguidas por la misma letra no presentan diferencias significativas (Test de Tukey, p < 0.05).

Figure 2. Effect of GP addition on germination, root length and germination index of lettuce and tomato plants.Figura 2. Efecto de la adición del OU sobre la germinación, longitud radicular e índice de germinación de plantas de lechuga y tomate.

# GP effect on growth of tomato and lettuce plants.

Significant differences were found in expanded leaf number (EL) (H= 8.54; p= 0.0259), greenness index (GrI) (F= 14.21; p < 0.0001), total leaf area (TLA) (F= 95.71; p < 0.0001), aerial dry weight (ADW) (F= 77.05; p < 0.0001) and root dry weight (RDW) (F= 40.54; p < 0.0001) of lettuce. The TLA, ADW and RDW significantly decreased with GP doses over 40%. The addition of 20 t ha<sup>-1</sup> of GP as mulch showed an increase in RDW.

For tomato plants, significant differences were found in GrI (F= 6.72; p= 0.0038), TLA (F= 46.47; p < 0.0001), ADW (F= 21.00; p < 0.0001) and RDW (F= 30.54; p < 0.0001). The GP presented the highest toxic effect on ADW and TLA at doses of 40-80 t ha<sup>-1</sup>. In contrast, GrI significantly increased at the maximum dose. Also, a non-toxic effect of GP was observed on RDW at 20 t ha<sup>-1</sup> (table 2).

**Table 2.** Expanded leaf number (EL), greenness index (GrI), total leaf area (TLA), aerialdry weight (ADW) and root dry weight (RDW) of lettuce and tomato plants grown withincreasing GP doses.

**Tabla 2**. Número de hojas expandidas (EL), Índice de verdor (GrI), Área foliar total (TLA), Biomasa seca área (ADW) y biomasa seca radicular (RDW) de plantas de lechuga y tomate en respuesta a diferentes dosis crecientes de OU.

Doses (t ha <sup>.1</sup> )	EL (n° plan	t1)	GrI (SPAD units)		TLA (cm² plant⁻¹)		ADW (g plant <sup>-1</sup> )		RDW (g plant <sup>1</sup> )	
					Lettuce					
0	11	а	28.2 ± 1.8	а	656.9 ± 85.3	а	3.6 ± 0.4	a	0.6 ± 0.1	b
20	9	b	19.8 ± 2.9	b	295.7 ± 18.9	b	2.3 ± 0.3	b	0.8 ± 0.1	а
40	9	b	23.3 ± 1.9	b	180.0 ± 30.7	с	$1.0 \pm 0.2$	с	$0.3 \pm 0.0$	с
80	9	b	22.1 ± 1.7	b	195.9 ± 41.7	с	$1.0 \pm 0.3$	с	$0.3 \pm 0.0$	с
	Tomato									
0	11	Α	$31.2 \pm 4.0$	В	552.1 ± 57.2	A	9.1 ± 2.2	A	0.6 ± 0.1	В
20	11	А	31.2 ± 1.9	В	548.7 ± 43.9	A	9.5 ± 0.6	A	0.8 ± 0.1	A
40	10	A	36.3 ± 4.8	AB	275.7 ± 56.5	В	3.7 ± 1.1	В	$0.4 \pm 0.0$	С
80	11	Α	38.1 ± 3.5	Α	295.6 ± 41.2	В	4.0 ± 1.8	В	$0.4 \pm 0.1$	С

Data means  $\pm$  S.D. of 10 replicates. For each row and vegetal species, different letters indicate significant differences (p < 0.05; Tukey test or Kruskal-Wallis test).

Los datos son medias de 10 réplicas  $\pm$ D.E. Para cada fila y especie vegetal, diferentes letras indican diferencias significativas (p < 0,05; Test de Tukey o Kruskal-Wallis).

#### DISCUSSION

This study found high OM contents in *V. labrusca* GP, as reported for *V. vinifera* and other agroindustrial by-products like solid olive mill "alperujo" (*Olea europaea* L.) and coffee pulp (2, 3, 4, 6). Unlike oil and coffee by-products, grape pomaces have high NT content and a C/N ratio nearly under the recommended limits for organic fertilizing (< 25) (6, 22).

Depending on the origin, commercial organic fertilizers have diverse amounts of essential nutrients (23). Bustamante *et al.* (2008) found 1.15 g kg<sup>-1</sup> total P for *V. vinifera*, while in this study *V. labrusca* showed a higher total P content, resulting in a more attractive by-product for the agronomic industry.

While most organic fertilizers usually require EC values under 4 dS m<sup>-1</sup>, the value obtained in this study was higher and considered detrimental to plant growth (22). However, the main salts found in *V. labrusca* GP are Ca<sup>+2</sup> and/or Mg<sup>+2</sup> chlorides and sulfates (SAR < 15). In this sense, the pomace could be considered a good alternative to animal manure causing soil disintegration due to concentrated Na<sup>+</sup> salts (9, 10). Furthermore, *V. labrusca* pomace exhibited a CEC value similar to that of highly productive soils (>45), probably given by the available functional groups negatively charged (phenols, carboxylic acids, etc.) found in OM (15). Hence, soil addition of GP might reduce leaching, thus increasing essential cations availability (39). Soils with a saturation base close to 100% exhibit alkaline pH (14). In this study, the pomace showed an elevated base saturation while being an acidic by-product, probably given to the presence of organic acids like malic and tartaric, common in grapes, that together with K+ and its effect over the diminished free tartaric acid, define by-product acidity (3). On the other hand, the high moisture content in this by-product (79.2%) is within the range reported for *V. vinifera* pomace and other agroindustrial wastes such as alperujo and coffee pulp ranging between 64-80% (3, 4).

Introducing OM into the system could differentially affect soil microorganisms and plants, depending on application dose and tolerance ranges (26, 41). *A. brasiliense* is a soil-free nitrogen-fixing bacteria producing several plant signalling molecules like phytohormones (auxins and gibberellins) (13, 32). The positive effect of GP on this diazotrophic bacteria has not been previously reported using other by-products (37).

When seeds were exposed to 10% GP concentration, primary roots showed a brownish colour in both plant species, probably given by cell necrosis caused by toxic compounds in GP or polymerization of compounds (chromophores) in root exudates as a defence response. A similar symptom was reported in tomato roots as an allelochemical effect of *Sicyos deppei* (Cucurbitaceae) (33). Some phenols act as allelochemical compounds related to polar narcosis (structural and functional alteration of cell membrane), oxidation uncoupling, alterations in electrophilicity, hydrophobicity, and dissociation and union of H in biomolecules (30, 44). Since *V. labrusca* pomace is rich in soluble phenolic compounds, these substances could be related to the mentioned root symptoms.

In the plant experiments, the stimulant effect of GP on root systems at the lowest dose could be due to greater nutrient availability (9). In intensive crops, such as lettuce and tomato, the recommended dose of an organic fertilizer aimed at maintaining soil productive capacity is 40 t ha<sup>-1</sup> (42). The results obtained in this study would limit GP utilization as an organic fertilizer, but applications at the lowest dose could be used as mulching in tomatoes and possibly in other crops with similar tolerance ranges.

#### **CONCLUSIONS**

*V. labrusca* GP presents physicochemical characteristics associated with soil health. Its high content of phosphorus and potassium, as well as low sodium and low SAR values differentiate this winery by-product from the one derived from *V. vinifera*. Low concentrations of *V. labrusca* GP promotes *A. brasiliensis* and tomato root system without altering aerial biomass, an effect known as hormesis. Toxicity symptoms of *V. labrusca* GP on plant growth at highest doses restricts its usage as an organic fertilizer. However, further ongoing field experiments will evaluate the effect of GP as mulching on tomato growth and other microorganisms indicating soil biological quality.

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# A polyphasic study of non-aflatoxigenic *Aspergillus flavus* Link, isolates from maize in the Chaco semi-arid region of Argentina

# Estudio polifásico de aislados no aflatoxigénicos de *Aspergillus flavus* Link, en maices de la región del Chaco semiárido argentino

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# ABSTRACT

Maize (Zea mays L.) is one of the most widely planted crops globally with Argentina leading world production and exportation. Santiago del Estero province, east of Tucumán and north of Córdoba encompasses eight agro-climatic zones in the Chaco Semi-arid region, agro-ecologically characterized by a wide temperature range and frequent drought periods that expose the crop to pathogens, particularly Aspergillus flavus. This pathogen is responsible for ear rot and grain contamination with mycotoxins such as aflatoxin B, and cyclopiazonic acid. This study obtained fungal isolates from ears of maize and characterized them according to toxigenic capability and morphotype of sclerotia (S < 400  $\mu$ m, associated with high levels of aflatoxins and L > 400  $\mu$ m, related to variable levels of aflatoxins). In addition, those not producing aflatoxins were studied to determine phylogenetic relationships based on sequences of a segment of the *CaM* gene. Fifty-eight isolates were obtained in eight localities representing each agro-climatic zone, 30 of which were non-aflatoxigenic, 28 aflatoxigenic, and all producers of ciclopiazonic acid. Six isolates did not produce sclerotia, 51 were L and only one was S, the latter being a non-producer of aflatoxins. The number of sclerotia was positively correlated with the production of aflatoxin B<sub>1</sub>, while size was negatively correlated. The CaM gene sequences corroborated that the isolates belonged to the A. flavus clade and the high nucleotide similarity among them (99.4% to 100%) revealed almost zero genetic diversity in this geographic region. No significant differences were observed in the proportion of isolates between growing seasons or among agroclimatic districts. This research revealed characteristics of fungus populations in this agricultural region of north Argentina.

# **Keywords**

non-aflatoxigenic isolate • cyclopiazonic acid • sclerotia production • phylogenetic relationships

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#### RESUMEN

El maíz (Zea mays L.) es uno de los cultivos más sembrados en el mundo, siendo Argentina líder mundial en producción y exportación. La Provincia de Santiago del Estero, el este de Tucumán y el norte de Córdoba abarcan ocho zonas agroclimáticas en la región Chaco semiárida, la cual se caracteriza por un amplio rango de temperatura y frecuentes sequías que exponen al cultivo a patógenos, particularmente Aspergillus flavus, responsable de la podredumbre de la espiga y la contaminación de granos con micotoxinas como aflatoxina B, y ácido ciclopiazónico. Se obtuvieron aislados del hongo desde espigas de maíz y se caracterizaron según capacidad toxigénica y morfotipos de esclerocios (S < 400  $\mu$ m, asociados con elevados niveles de aflatoxinas y L > 400 µm, relacionados con niveles variables de aflatoxinas). Además, aquellos que no producían aflatoxinas se estudiaron para determinar sus relaciones filogenéticas sobre la base de un segmento del gen CaM. Cincuenta y ocho aislados fueron obtenidos en ocho localidades representativa de cada zona agroclimática, 30 de los cuales resultaron no aflatoxigénicos, 28 aflatoxigénicos y todos productores de ácido ciclopiazónico. Seis aislados no produjeron esclerocios, 51 fueron L y solo uno fue S, siendo este último no aflatoxigénico. El número de esclerocios correlacionó positivamente con la producción de aflatoxina B<sub>1</sub>, mientras que el tamaño lo hizo de manera negativa. Las secuencias del gen CaM corroboraron que los aislados pertenecen al clado A. flavus y la alta similitud nucleotídica (99,4% a 100%) reveló casi nula diversidad genética en esta región. No se observaron diferencias en la proporción de aislados entre campañas agrícolas estudiadas ni distritos agroclimáticos. Esta investigación reveló características de las poblaciones de hongos en esta región agrícola del norte argentino.

## **Palabras clave**

aislado no aflatoxigénico • ácido ciclopiazónico • producción de esclerocios • relaciones filogenéticas

#### INTRODUCTION

Argentina ranks sixth in maize production, with approximately 60 million tons a year. The region encompassing Santiago del Estero province, east of Tucumán and north of Córdoba, in the Chaco Semi-arid region, produces about 6 million tons (58). This geographical area is a critical climatic zone, where climatic factors are highly variable (57). Thus, maize crop development in these agroecosystems is constrained by stressful environmental factors that expose the crop to different pathogens, such as those responsible for ear rot (27, 67). The most dangerous of these pathogens is Aspergillus flavus Link, a globally distributed filamentous, cosmopolitan pathogen that causes opportunistic infections in animals and plants (12, 42). This pathogen is commonly present in air and soil mycobiota. Under certain water and thermal stress conditions, some isolates produce mycotoxins like aflatoxins and cyclopiazonic acid (CPA). Aflatoxins are secondary metabolites with strong hepatotoxic, teratogenic, immunosuppressive and mutagenic activity when inhaled, ingested or absorbed through animal and human skin (51); A. flavus is the main driver of aflatoxin contamination in maize in the world (45). While CPA produces liver necrosis, convulsions myocardium lesions in animals (62, 63), the specific exposure response to this toxin differs among species, without sufficient evidence about natural contamination of foods (81).

The maximum safety limit of aflatoxins for maize commercialization varies between countries, being 20 ng. g<sup>-1</sup> in Argentina, Brazil and USA, 4 ng. g<sup>-1</sup> in the European Union, and 15 ng. g<sup>-1</sup> in African countries like Ghana (4, 9, 16, 34, 77).

Species of the genus *Aspergillus* produce sclerotia, resistance structures consisting of darkly pigmented and hardened hyphal mats (50) conferring different phenotypic characteristics (1). Thus, *Aspergillus* species can be classified into two morphotypes: S isolates, which produce numerous sclerotia under 400  $\mu$ m in diameter, associated with high levels of aflatoxins, and L isolates, with fewer sclerotia, diameter over 400  $\mu$ m, and related to variable levels of aflatoxins (75). A third type of isolate not producing sclerotia under laboratory conditions, has been identified in Argentina and Italy (13, 38).

The variability among *A. flavus* populations can be estimated by establishing phylogenetic relationships based on morphological and molecular markers (51), as in the case of the calmodulin gene (*CaM*) (36). Calmodulin (*CaM*) is a highly conserved polypeptide present in eukaryotic cells. In the genus *Aspergillus, CaM* is important in phosphorylation and dephosphorylation of proteins involved in the biosynthesis of aflatoxins (47).

Aflatoxin contamination usually starts in the field, during harvest of infected grains, becoming more severe during storage (35). In the field, spores of *A. flavus* are transported by biotic agents (insects, birds) and/or abiotic agents (wind, rain splash) to the maize ear. Spores enter through the silk during flowering and through the wounds of the ear cover or of developing kernels (66). Water stress, high temperatures, insect attack, certain sowing dates and high crop density favor contamination with the pathogen and the possible production of aflatoxins (48).

Some members of *A. flavus* were found unable to biosynthesize aflatoxins being, therefore, plausible biocontrol tools (23). Biological control applied in maize and peanut (6, 73), among other crops, is based on the competition for infection sites and essential nutrients between non-aflatoxigenic and aflatoxigenic isolates. Thus, non-aflatoxigenic isolates displace aflatoxigenic ones and reduce the latter's capacity to produce aflatoxins (70, 79).

Non-aflatoxin producing strains selected as biocontrol agents must meet some requirements, such as being native to the area of application, being able to displace the toxicogenic isolates in the ecosystem and thereby reduce infections (23) and be genetically incompatible with the remaining fungal population. Therefore, studies of the native fungal population are crucial (42).

This work aimed to study the diversity of *A. flavus* populations in the Chaco Semi-arid region of Argentina through the morphological, toxicological and molecular characterization of isolates obtained from maize ears.

#### MATERIALS AND METHODS

#### Sampling

Sampling was conducted during the 2015/16 and 2016/17 growing seasons in representative localities of eight agro-climatic zones: Sachayoj (Zone 4: Río Muerto), Quimilí (Zone 5: Hickmann), Bandera (Zone 17: Bandera), Santiago del Estero (Zone 18: Monteagudo), Sumampa (Zone 37: Soto), Santa Rosa de Leales (Zone 18A: Monteagudo), Rayo Cortado (Zone 37C: Soto-north of Córdoba) and Villa de Tulumba (Zone 61B: Deán Funes), located between -30.38 and -26.55 S and -61.82 and -65.27 W (25) (figure 1, page 61). Ten ears were randomly collected along a diagonal line in the plot, at plant physiological maturity before harvest. They were threshed to form a composite sample and oven-dried at 38°C for 72 hours until final humidity was under 12 %, therefore, minimizing fungus development (26).

# Isolation and morphological and molecular identification

From each sample, 100 kernels were taken, surface-desinfested with a 1% sodium hypochlorite wash followed by three washes with distilled water and plated in DG-18 medium. Kernels were incubated for 7 days at 25°C and cultures with characteristics similar to those of *A. flavus* were transferred to MEA medium. They were incubated for 7 days at 25°C (65) and then identified through taxonomic keys (48). Spores of each isolate were serially diluted, and the most diluted concentration was cultured in 2 % Water - Agar (WA) medium and incubated for 18 hours at 27°C (65). The germinated conidia were identified under Nikon SMZ-10 stereoscopic microscope (15X). Two conidia were extracted from each cultured Petri dish, along with a portion of culture medium, and transferred to Petri dishes containing MEA medium. They were incubated for 7 days at 25°C. Then, one of them was stored to obtain the single spore cultures corresponding to each isolate.



Localidades evaluadas: Sachayoj. Quimilí. Santiago del Estero. Bandera. Sumampa (Santiago del Estero province). Santa Rosa de Leales (Tucumán province). Rayo Cortado. Villa de Tulumba (Córdoba province).

Sampling localities /

Figure 1. Sampling localities in the Chaco Semi-arid region of Argentina. Figura 1. Localidades evaluadas en la región del Chaco semiárido argentino.

The identity of *A. flavus* was confirmed by PCR using the primers FLA1, 5'-GTAGG-GTTCCTAGCGAGCC-3'; FLA2 5'-GGAAAAAGATTGATTTGCGTTC-3' (41), with the ITS region as target, using the isolate *A. flavus* CCC 101-92 < NRRL 3251 as positive control, and the isolate *A. parasiticus* NRRL 2999 as negative control. Each single-spore isolate was cultured in MEA and incubated for 7 days at 25°C. Then, an aliquot of mycelium was taken, and DNA was extracted (55). For each reaction, 1 µL (final concentration of 20-100 ng of DNA.µL<sup>-1</sup>, quantified in Nano Drop, Thermo Fisher Scientific, USA) of DNA solution was mixed with 24 µL of a solution composed of 5 µL 5x Green GoTaq® reaction buffer (Promega®, Madison, WI, USA), 0.5 µL of a mixture of dNTP (10 mM of dATP, dCTP, dGTP and dTTP), 1 µL of the primers (20 µM), 1 µL of polymerase enzyme GoTaq® (5U.µL<sup>-1</sup>, Promega®, Madison, WI, USA) and 16.5 µL DEPC-treated water. The PCR reaction was performed following González Salgado *et al.* (2011). PCR products were revealed by agarose gel electrophoresis, visualized in UV transilluminator, with previous staining in GelRed<sup>TM</sup> Biotium solution (2.5 ng. µL<sup>-1</sup>) and confirmed by the expected band size of 500 pb. Molecular size of the DNA fragments was estimated using the "qLadder 100 pb precision" marker (PB-L®, Bs As, Argentina).

#### Sclerotia production

Each isolate was cultured in 6-cm Petri dishes containing Czapeck dox (Cz) medium. They were inoculated in the center of the dish with 10  $\mu$ L of a spore suspension and incubated for 14 days at 30°C (64). Sclerotia were recovered, their diameters measured under a microscope (Nikon eclipse Cs1 spectral) and characterized following Cotty (1989) into L or S morphotypes or isolates not producing (NP) sclerotia (60). Isolates not producing sclerotia after an incubation period were cultured in 5/2 medium for 5 to 7 days at 31°C to induce their production (38).

# **Toxigenic capability**

Considering Aflatoxin B<sub>1</sub> (AFB<sub>1</sub>) was one of the greatest frequency and toxicity of the four most important aflatoxins identified, its production was quantified (23, 82). *A. flavus* isolates were inoculated in duplicate on MEA slants for 7 days at 28°C. These cultures were used to prepare spore suspensions, following the method described by Alaniz Zanon *et al.* (2013). Spore concentration was measured in a Neubauer chamber and adjusted to  $10^5$  spores. mL<sup>-1</sup>. Four-milliliter vials containing 1 mL broth from a medium containing 150 g sucrose, 20 g yeast extract, 10 g soytone, and 1 L distilled water, were inoculated with 100 µL of each spore suspension. Medium pH was adjusted to 5.9 with HCl. Cultures were incubated for 7 days at 30°C. Vial cultures were extracted according to Horn *et al.* (1996), by adding 1 mL of chloroform to each vial and vortexing for 30 s.

The first group of dried extracts was evaluated in their capacity to produce aflatoxins by HPLC according to Horn *et al.* (1996). Aflatoxins were analyzed by injecting 50  $\mu$ L of extract from each vial into an HPLC system consisting of a Hewlett Packard model 1100 pump (Palo Alto, CA) connected to a Hewlett Packard model 1046A programmable fluorescence detector and a data module Hewlett Packard Kayak XA (HP ChemStation Rev.A.06.01). Chromatographic separations were performed on a stainless steel, C18 reversed-phase column (150 mm × 4.6 mm i.d., 5  $\mu$ m particle size; Luna Phenomenex, Torrance, CA, USA) connected to a precolumn Security Guard (20 mm × 4.6 mm i.d., 5  $\mu$ m particle sizes, Phenomenex). The mobile phase was water:methanol:acetonitrile (4:1:1, v/v/v) at a flow rate of 1.5 mL. min<sup>-1</sup>. Pure aflatoxin solutions were used as external standards (Sigma-Aldrich, St. Louis, MO, USA).

The second group of dried extracts was resuspended in 500  $\mu$ L of mobile phase of acetonitrile:  $ZnSO_4$  4 mM buffer solution (65:35, V/V) and CPA was determined in the HPLC system. Chromatographic separations were performed in Agilent ZORBAX RX-SIL column (250 mm x 4.6 mm i.d., 5  $\mu$ m particle size) connected to a pre-column Security Guard (20 mm x 4.6 mm i.d., 5  $\mu$ m particle sizes, Phenomenex) and 50  $\mu$ l of each sample were analyzed at a flow velocity of 0.8 mL. min<sup>-1</sup>.

The quantitative analysis was performed by normalization of peak areas. A calibration curve was elaborated from the areas obtained for the different concentrations of standards of aflatoxins and CPA, as applicable (Sigma Aldrich, St. Louis, MO, USA). The AFB<sub>1</sub> and CPA detection limit was 1 ng.  $g^{-1}$  (24, 32).

#### Sequencing and Phylogenetic relationships

Molecular variability of the non-aflatoxigenic isolates was explored using a 688-pb fragment of the *CaM* gene. We studied only this type of isolate given their potential for biocontrol strategies. Spores of the isolates cultivated in MEA medium were suspended for 7 days at 25°C. Conidia ( $1x10^6$  conidia. mL<sup>-1</sup>) were inoculated in 100 mL of lixiviated medium of potato glucose agar (10% of potato infusion from 200 g potato, 2% glucose, 4.5 pH). They were incubated in orbital shaker at 150 rpm for 48 hours at 25°C. Mycelium was collected by filtration and powdered with liquid N<sub>2</sub> (28). This culture medium allows one to obtain enough amount of mycelium. DNA was extracted using the cetyl-trimethylammonium bromide (CTAB) method (54), quantified by spectrophotometry and a segment of the *CaM* was amplified with the primers CL1, 5'- GA(GA)T(AT)CAAGGAGGCCTTCTC -3'; CL2A, 5'- TTTTGCATCATGAGTTGGAC -3' (61). The obtained fragments of the expected size (688 pb) were purified using DNA columns of Wizard® SV Gel columns and PCR clean - Up system (Promega, Madison, WI, USA), following manufacturer instructions. Then, they were quantified by spectrophotometry. Both DNA chains were sent for sequencing by Sanger

method (Macrogen, Seoul, South Korea). The obtained sequences were aligned using the software Clustal X2 (54). The software MEGA version 7 (53) was used to select the best nucleotide substitution model and to obtain the phylogenetic trees using Neighbor-Joining (NJ) (74) and Maximum-Likelihood (ML) methods (39, 76), both with 10,000 replicates bootstrap. The sequences were compared with the reference NW\_002477238.1 and *A. niger* MH645004.1 was used as outgroup, both from the GenBank.

#### Statistical analysis

Data on sclerotia morphotypes, toxigenic capacity and variability of the sequences of the *CaM* genes were analyzed using InfoStat statistical software as well as correlation analyses (29). Mean values were obtained using analysis of variance (ANOVA) and differences between means were compared using Fisher's LSD test (P< 0.05).

#### **RESULTS AND DISCUSSION**

# Morphological and molecular identification

A total of 58 isolates were obtained in the sampling localities from the eight agro-climatic zones (table 1, page 63-64) with *A. flavus* morphological characteristics, according to Klich (2002) (figure 2, page 64). *A. flavus* identity was confirmed using the specific primers FLA1 and FLA2 through amplification of a 500-pb band (41) (figure 3, page 65). The presence of isolates in the eight zones, each one being considered a different agroecological environment given different meteorological scenarios at each growing season shows the broad distribution of the fungus in the study region, as reported for other regions of the country (8, 17) and the world (11, 30, 32).

Table 1. Isolates of Aspergillus flavus from maize ears collected from representative localities of eight agro-climatic zones of the Chaco Semi-arid region of Argentina.
Production of aflatoxin B<sub>1</sub>, cyclopiazonic acid and sclerotia morphotype during the 2015/16 and 2016/17 growing seasons.

**Tabla 1.** Aislados de *Aspergillus flavus* de espigas de maíz colectadas en localidades representativas de ocho zonas agroclimáticas del Chaco semiárido argentino. Producción de aflatoxina B<sub>1</sub>, ácido ciclopiazónico y morfotipo de esclerocios durante las campañas agrícolas 2015/16 y 2016/17.

Isolate	Agro-climatic Zone	Locality	Sampling year	Sclerotia Morphotype	AFB <sub>1</sub> (ng. g <sup>-1</sup> )	CPA (ng. g <sup>-1</sup> )
ASRC1	37C	Rayo Cortado	2017	L	6.4	3.3
ASRC2	37C	Rayo Cortado	2017	NP	0.0	6.2
ASRC3	37C	Rayo Cortado	2017	NP	0.0	0.6
ASVT1	61B	Tulumba	2017	L	278.4	0.3
ASVT2	61B	Tulumba	2017	L	0.0	2.8
ASVT3	61B	Tulumba	2017	L	405.1	8.6
ASVT4	61B	Tulumba	2017	L	60.6	2.3
ASBA12	17	Bandera	2016	L	16.3	9.9
ASBA3	17	Bandera	2016	L	0.0	2.8
ASBA4	17	Bandera	2016	L	7.2	9.2
ASBA8	17	Bandera	2016	L	0.0	4.7
ASBA1	17	Bandera	2017	L	0.0	2.3
ASBA10	17	Bandera	2017	L	5.9	9.3
ASBA11	17	Bandera	2017	L	0.0	2.3
ASBA2	17	Bandera	2017	L	0.0	2.3
ASBA5	17	Bandera	2017	L	0.0	6.7
ASBA6	17	Bandera	2017	L	90.7	2.9
ASBA7	17	Bandera	2017	L	6.7	5.4
ASBA9	17	Bandera	2017	NP	0.0	6.9

Agro-climatic zones: 4: Río Muerto, 5: Hickmann, 17: Bandera, 18 and 18A: Monteagudo, 37: Soto, 37C: Soto - North of Córdoba. 61B: Dean Funes. L: sclerotia of > 400 µm; S: sclerotia of < 400 µm; NP: non-producer of sclerotia. AFB1: Aflatoxin B; CPA: cyclopiazonic acid. Zonas agroclimáticas: 4: Río Muerto, 5: Hickmann, 17: Bandera, 18 y 18A: Monteagudo, 37: Soto, 37C: Soto - Norte de Córdoba, 61B: Deán Funes. L: esclerocios de tamaño > 400 µm; S: esclerocios de tamaño < 400 µm; NP: no productor de esclerocios.

AFB<sub>1</sub>: Aflatoxina B; CPA: ácido ciclopiazónico.

	Isolate	Agro- climatic Zone	Locality	Sampling year	Sclerotia Morphotype	AFB <sub>1</sub> (ng. g <sup>-1</sup> )	CPA (ng. g <sup>-1</sup> )
	ASQU1	5	Quimilí	2016	L	7.2	5.6
	ASQU10	5	Quimilí	2016	L	0.0	54
	ASQU11	5	Quimilí	2016	L	2521.5	7.5
	ASQU12	5	Quimilí	2016	L	17.6	5.8
	ASQU13	5	Quimilí	2016	L	5.6	1.2
	ASQU14	5	Quimilí	2016	L	5.6	8.6
	ASQU15	5	Quimilí	2016	L	0.0	3.2
	ASQU16	5	Quimilí	2016	L	0.0	4.2
	ASQU17	5	Quimilí	2016	L	0.0	8.7
	ASQU18	5	Quimilí	2016	L	0.0	8.7
	ASQU19	5	Quimilí	2016	L	0.0	0.6
	ASQU2	5	Quimilí	2016	L	0.0	0.6
	ASQU20	5	Quimilí	2016	L	0.0	7.1
	ASQU21	5	Quimilí	2016	L	499.1	5.9
	ASQU3	5	Quimilí	2016	L	34	0.4
Agro-climatic zones:	ASQU4	5	Quimilí	2016	L	9.3	8.1
4: Río Muerto,	ASQU5	5	Quimilí	2016	L	6.3	16.4
5: Hickmann, 17: Bandera, 18 and	ASQU6	5	Quimilí	2016	L	6	6.2
18A: Monteagudo,	ASOU8	5	Ouimilí	2016	L	5.7	6.9
37: Soto, 37C: Soto	ASQU9	5	Quimilí	2016	L	0.0	8.6
- North of Córdoba,	ASQU7	5	Quimilí	2017	L	0.0	1.6
61B: Dean Funes.	ASSA1	4	Sachavoj	2016	L	9.8	8.5
L: sclerotia of > 400 $\mu$ m; S: sclerotia of < 400 $\mu$ m;	ASSA3	4	Sachavoi	2016	L	13.2	9.9
NP: non-producer of	ASSA4	4	Sachavoi	2016	L	0.0	7.6
sclerotia.	ASSA6	4	Sachavoi	2016	L	9.1	9.1
AFB1: Aflatoxin B; CPA:	ASSA2	4	Sachavoi	2017	L	70.1	7.9
cyclopiazonic acid.	ASSA5	4	Sachavoi	2017	L	0.0	3.8
Zonas agroclimáticas:	ASSE5	18	Sgo del Estero	2016	L	5.8	8.9
4: Río Muerto,	ASSE1	18	Sgo del Estero	2017	S	0.0	1.8
Bandera, 18 v 18A:	ASSE2	18	Sgo del Estero	2017	L	0.0	0.6
Monteagudo, 37: Soto,	ASSE3	18	Sgo del Estero	2017	L	48958.1	2.2
37C: Soto - Norte de	ASSE4	18	Sgo del Estero	2017	L	290.9	4.9
Córdoba, 61B: Deán	ASSE6	18	Sgo del Estero	2017	NP	0.0	4.9
Funes.	ASSE7	18	Sgo del Estero	2017	NP	0.0	0.6
L: escierocios de tamano	ASSE8	18	Sgo del Estero	2017	L	0.0	2.0
de tamaño < 400 µm;	ASSE9	18	Sgo del Estero	2017	L.	0.0	0.8
NP: no productor de	ASSUI	37	Sumamna	2017	NP	0.0	2.9
esclerocios.	ASSU2	37	Sumampa	2017	L	12.3	51
AFB <sub>1</sub> : Aflatoxina B; CPA: ácido ciclopiazónico.	ASLE1	18A	Leales	2017	L	0.0	1.3



Figure 2. Aspergillus flavus Link colony on malt extract agar (MEA) culture medium. Figura 2. Colonia de Aspergillus flavus Link en medio de cultivo agar extracto de malta (MEA).



**Figure 3.** Electrophoresis gel showing PRC products obtained using the specific primers FLA1 and FLA2 (González Salgado *et al.*, 2011).

**Figura 3.** Gel de electroforesis mostrando productos de PCR obtenidos utilizando los iniciadores específicos FLA1 y FLA2 (González Salgado *et al.*, 2011).

## Sclerotia production

Of the 58 *A. flavus* isolates identified, 6 (10%) produced no sclerotia, whereas 52 (90%) produced sclerotia. Of the latter, 51 (98%) were identified as L morphotype (figure 4), with > 400  $\mu$ m in diameter, and only isolate, ASSE1, (2%) corresponded to S morphotype, with < 400  $\mu$ m in diameter (table 1, page 63-64). Similar proportions between L, S and NP morphotypes in maize kernels were reported by Moreno (2004) in Mexico and by Pildain *et al.* (2005) in Argentina in peanut isolates. The low or null presence of S morphotype in maize was also reported by Giorni *et al.* (2007) in Italy. These authors identified one S isolate out of 70 isolates, with the remaining ones being NP. Similarly, Alaniz Zanon *et al.* (2018) reported a high proportion of L morphotype in Argentina with respect to NP, with absent S morphotype.

Sclerotia production was reported in isolates from all the agro-climatic zones and in both evaluated growing seasons. Dominance of L morphotype among isolates obtained in the Chaco Semi-arid region agrees with previous records in Argentina (7, 18) and in other countries, like Brazil, Portugal, Nigeria and Sub-Saharan Africa regions (5, 20, 33, 68, 72).



**Figure 4.** Sclerotia L morphotype in aflatoxigenic isolate of *Aspergillus flavus*. Observation under stereo microscope Nikon SMZ-10 (15X).

Figura 4. Esclerocios morfotipo L en aislado aflatoxigénico de *Aspergillus flavus.* Observación bajo lupa estereoscópica Nikon SMZ-10 (15X).

M: molecular marker (100-pb DNA Ladder); +: Aspergillus flavus NRRL 3251 (positive control); -: A. parasiticus NRRL2999 (negative control); 1: ASSA2; 2: ASQU7; 3: ASQU21; 4: ASQU10; 5: ASQU15; 6: ASRC2; 7: ASQU12; 8: ASQU6; 9: ASBA3; 10: ASVT1.

M: marcador molecular (100-pb DNA Ladder); +: Aspergillus flavus NRRL 3251 (control positivo); -: A. parasiticus NRRL2999 (control negativo); 1: ASSA2; 2: ASQU7; 3: ASQU21; 4: ASQU10; 5: ASQU15; 6: ASRC2; 7: ASQU15; 6: ASRC2; 9: ASBA3; 10: ASVT1. The high incidence of S morphotype is associated with regions of low precipitation and high temperatures, where high number of small sclerotia can be a survival trait facing rapid temperature and humidity fluctuations (19). However, and even though the study region is dry and hot, this was not observed.

## **Toxigenic capability**

Based on the production of  $AFB_1$  and CPA of the 58 isolates obtained, we identified 30 (52%) non-aflatoxigenic and 28 (48%) aflatoxigenic. This similarity of proportions was indicated by Martins *et al.* (2017) in peanut crop in Brazil and Camiletti *et al.* (2018) in maize ears in Argentina.

Aflatoxin-producer and non-producer isolates were obtained in all the agro-climatic zones and growing seasons, except Santa Rosa de Leales, where the only isolate found was non-producer.

Of the 28 isolates producing AFB<sub>1</sub>, 26 (93%) had concentrations below 500 ng.g<sup>-1</sup>, whereas ASQU11 and ASSE3 were higher aflatoxin producers of 2,521.5 ng. g<sup>-1</sup> and 48,958.1 ng. g<sup>-1</sup> respectively (table 1, page 63-64). Although the capacity to produce mycotoxins may depend on geographical origin and environmental conditions (62), our results do not show any pattern associated with zones or growing seasons. This agrees with Bayman and Cotty (1993), who did not detect differential patterns between nearby zones.

All isolates were CPA producers, indicating toxicity importance (78) and plausible use as aflatoxin biocontrol strategies. This is the case of the AF36 agent used in competitive exclusion strategies and found responsible for CPA increase in maize kernels inoculated in the field, and peanuts (2, 31). By contrast, Camiletti *et al.* (2018) detected 19% of CPA non-producer isolates in maize kernels in Argentina. On the other hand, Vaamonde *et al.* (2003) studied wheat, soybean and peanut in Argentina and found between 6% and 27% of isolates not producing this mycotoxin. In Italy, Giorni *et al.* (2007) reported 39% CPA non-producer isolates in maize kernels, whereas, in organic nut plantations in Brazil, Reis *et al.* (2014) found that 34% of the *A. flavus* isolates were non-producers of that mycotoxin. In addition, Jamali *et al.* (2012) identified 19% of the isolates as CPA non-producers in soil samples of pistachio plantations in Czech Republic.

The presence of isolates producing both CPA and aflatoxins was reported in Argentina and other parts of the world (8, 18, 37, 71). The isolates with the highest AFB<sub>1</sub> production obtained in the Chaco Semi-arid region do not correspond to those with the highest CPA production.

Correlations between  $AFB_1$  production and the number (r= 062) and size of sclerotia (r= -0.38) indicate that the produced amount of  $AFB_1$  increases as sclerotia size decreases. These results are similar to those reported by Arrúa Alvarenga *et al.* (2012) and Pildain *et al.* (2005), but differ from Bouti *et al.* (2020), who did not find any correlation between sclerotia and aflatoxin production. No correlation was observed between CPA production and sclerotia morphotype.

The only S morphotype identified in this study (ASSE1), collected in 2016/17 growing season, did not produce  $AFB_1$ . While the presence of non-aflatoxin-producing isolates was also reported for USA, Ghana and Brazil (3, 22, 40, 44), in general, S morphotype isolates produce higher aflatoxin concentrations than L or NP morphotypes and are identified as major causal agents of severe contamination in maize and of most human deaths due to aflatoxicosis (13, 23). The S isolates are worldwide distributed, associated with aflatoxins B production in the USA and Africa (68), and of both B and G aflatoxins (3, 38).

#### Sequencing and phylogenetic analysis

All the *CaM* gene sequences studied in this work, using both NJ and ML, belong to the *A. flavus* clade (figure 5, page 67 and figure 6, page 68). The high similarity among isolates for that character showed limited genetic diversity in this geographic region. In addition, the only S morphotype is located together with L morphotype isolates, all of which are aflatoxin non-producers. This finding may be attributed to the fact that these are young populations and, therefore, may have not undergone sufficient mutations or recombination events leading to variability among isolates from the region (42, 62). However, only the northern zones of the region (4 and 5) correspond to land recently converted to agriculture, whereas in the remaining districts, both commercial hybrids and maize for self-consumption have been cultivated for several decades.

	D5QuA16 (3)
	D5QuA16 (2)
	D17BaA17 (6)
	D37CRCA17 (3)
	D17BaA17 (5)
	D4SaC16 (2)
	D18SEB17 (2)
h	D5QuA16 (8)
	D37CRCA17 (2)
	D375uA17 (1)
	D18SEB17(1)
	D18ALeC17
	D17BaB16 (1)
	D5QuA16 (12)
l.	D18SEA17 (4)
	D4SaA17 (2)
ľ	D61BTuA17 (3)
l i	D5QuA16(17)
4	lspergillus flavus NW 002477238.1
1	18SEA17(3)
I I I I I I I I I I I I I I I I I I I	017BaA17 (8)
1	)17BaA17 (9)
I I I I I I I I I I I I I I I I I I I	017BaA17 (2)
1	95QuC17
1	)5QuA16 (5)
I I I I I I I I I I I I I I I I I I I	017BaC16
I	018SEA17(6)
I	05QuA16 (1)
I	05QuA16 (6)
I I	05QuA16(15)
l <sub>D</sub>	18SEA17 (2)
	Aspergill us niger MH645004.1

corresponds to bootstrap support values (10000 replicates). A. flavus NW 002477238.1 was selected as reference sequence and A. niger MH645004.1 as outgroup sequence. El número de nodos corresponde a un valor de bootstrap de 10000 réplicas. A. flavus NW 002477238.1 se utilizó como secuencia de referencia y A. niger MH645004.1 como secuencia externa al grupo.

Node number

**Figure 5.** Phylogenetic tree built using the Neighbor-Joining (NJ) statistic method based on the relationship among segments of the sequence of the *CaM* gene from *A. flavus* isolates from agro-climatic zones of the Chaco Semi-arid region in Argentina.

0.05

**Figura 5.** Árbol filogenético realizado con el método estadístico Neighbor-Joining (NJ) basado en las relaciones entre las secuencias de segmentos del gen *CaM* de aislados de *A. flavus* provenientes de zonas agroclimáticas del Chaco semiárido argentino.

	D5QuA16 (3)
	D5QuA16 (2)
	D17BaA17 (6)
	D18SEA17(4)
	D5QuA16 (12)
	D37CRCA17 (3)
	D17BaA17 (5)
	D4SaC16 (2)
ή	D18SEB17(2)
	D5QuA16 (8)
	D37CRCA17 (2)
	D37SuA17(1)
	D18SEB17(1)
	D18ALeC17
	D17BaB16 (1)
	D4SaA17 (2)
	D61BTuA17 (3)
1	D5QuA16 (17)
	Aspergillus flavus NW 002477238.1
	D18SEA17(3)
	D17BaA17 (8)
	D17BaA17 (9)
	D17BaA17 (2)
	D5QuC17
L	D5QuA16 (5)
	D17BaC16
	D18SEA17(6)
	D5QuA16 (1)
	D5QuA16 (6)
	D5QuA16 (15)
	D18SEA17(2)

selected as reference sequence and A. niger MH645004.1 as outgroup sequence. El número de nodos corresponde a un valor de bootstrap de 10000 réplicas. A. flavus NW 002477238.1 se utilizó como secuencia de referencia y A. niger MH645004.1 como secuencia externa al grupo.

Node number corresponds to bootstrap support values (10000 replicates). *A. flavus* NW 002477238.1 was

0.05

**Figure 6.** Phylogenetic tree built using the Maximum-Likelihood (ML) statistical method based on the relationship among segments of sequences of the *CaM* gene from *A. flavus* isolates from agro-climatic zones in the Chaco Semi-arid region in Argentina.

**Figura 6.** Árbol filogenético realizado con el método estadístico Maximum-Likelihood (ML) basado en las relaciones entre las secuencias de segmentos del gen *CaM* de aislados de *A. flavus* provenientes de zonas agroclimáticas del Chaco semiárido argentino.
#### CONCLUSIONS

*A. flavus* is present in all localities of the sampled agro-climatic zones and both growing seasons, evidencing the wide distribution of the pathogen in the Chaco Semi-arid region.

The population of *A. flavus* exhibits diversity in terms of sclerotia morphotypes and production of AFB<sub>1</sub> and CPA.

Low genetic variability among the isolates was observed, all belonging to the *A. flavus* clade.

In the future, genes from more variable regions will be used to perform phylogenetic analyses among *A. flavus* isolates.

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## Weed control in different germination fluxes with preemergent herbicides on sugarcane straw under dry periods

### Control de malezas en diferentes flujos de germinación a través de herbicidas preemergentes en aplicaciones sobre paja de caña de azúcar y períodos de seco

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#### ABSTRACT

Preemergent herbicides are a frequent weed control strategy. Considering different crop germinative fluxes, these products must present long-lasting weed control. This study evaluated preemergent herbicides in different germination fluxes of Merremia aegyptia, Mucuna aterrima and Ricinus communis when applied to different quantities of straw and different simulated dry periods. The experiment was conducted in a  $4 \times 2 \times 2$  factorial design with four replications. The treatments included four dry periods (0, 30, 60, and 90 days), two straw quantities (0 and 10 t ha<sup>-1</sup>), and two germination fluxes. The herbicides amicarbazone (1225 g  $ha^{-1}$ ), imazapic (147 g  $ha^{-1}$ ), sulfentrazone (800 g ha<sup>-1</sup>), and tebuthiuron (900 g ha<sup>-1</sup>) were applied for preemergence weed control, and germination flush fluxes were evaluated at 7, 14, 21, 28, and 35 days after emergence (DAE) while verifying plant dry mass. Amicarbazone controlled less than 80% of the studied species at the 90-day dry period in the presence of straw. Imazapic did not present control residue for any of the species analyzed. Sulfentrazone showed the same control pattern at all germination fluxes, regardless of the amount of straw. Tebuthiuron successfully controlled all species in the first germination flush, exceeding 80% regardless of the amount of straw. Herbicides associated with straw quantities and dry periods have a significant impact on *M. aegyptiaca*, *M. aterrima* and *R. communis*.

#### Keywords

amicarbazone • flush • germination • imazapic • precipitation • residue • straw • sulfentrazone • tebuthiuron

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#### RESUMEN

La aplicación de herbicidas preemergentes es una estrategia de control de malezas, sin embargo, estos productos deben presentar residualidad para el control de diferentes flujos germinativos. Este estudio tuvo como objetivo evaluar la eficacia de herbicidas preemergentes en diferentes flujos de germinación de Merremia aegyptia, Mucuna aterrima y Ricinus communis, cuando se aplican sobre diferentes cantidades de paja y diferentes períodos secos simulados. El experimento se realizó en un diseño factorial  $4 \times 2 \times 2$  con cuatro repeticiones, el tratamiento incluye cuatro períodos secos (0, 30, 60 y 90), dos cantidades de paja (0 y 10 t ha<sup>-1</sup>) y dos flujos de germinación. Los herbicidas amicarbazona (1225 g ha<sup>-1</sup>); imazapic (147 g ha<sup>-1</sup>), sulfentrazona (800 g ha<sup>-1</sup>) y tebuthiuron (900 g ha<sup>-1</sup>), se aplicaron para el control de malezas antes de la emergencia, y los flujos de flujo de germinación se evaluaron a los 7, 14, 21, 28 y 35 días después de la emergencia de especies (DAE), mientras se verifica la masa seca. La amicarbazona presentó una reducción del control para todas las especies en los periodos secos más prolongados y presencia de paja. La amicarbazona mostró menos del 80% de control para todas las especies a los 90 días del período seco en presencia de paja. Imazapic no presentó residuo control para ninguna de las especies analizadas. Para sulfentrazona, la cantidad de paja no afectó el control de las malezas en diferentes flujos de germinación, mostrando el mismo patrón de control independientemente de la cantidad de paja, porcentajes de control superiores al 80% independientemente de la cantidad de paja. Los herbicidas asociados a cantidades de paja y periodos secos tienen impacto sobre las especies de malezas *M.aegyptia*, *M.aterrima* y R. communis.

#### **Palabras claves:**

amicarbazone • flush • germinación • imazapic • precipitación • residuo • paja • sulfentrazone • tebuthiuron

#### INTRODUCTION

Raw sugarcane straw (without preburning) on the soil surface promotes a favorable environment for seed germination and weed development, such as *Merremia aegyptia* (L.) Urb., *Mucuna aterrima* Piper & Tracy, and *Ricinus communis* L. (12, 13). In these productive systems in Brazil, these three weed species are popularly known as "the three M's" (MMM-castor bean, *morning glory*, and mucuna). These species are adapted to sugarcane production systems with straw deposition on the soil surface. These systems hinder weed control with herbicides, causing serious damage to sugarcane production (2, 5, 18, 28, 33).

In addition to favoring the establishment of these species, the straw that remains on the soil surface represents a physical barrier to the action of preemergent herbicides (29), that once intercepted by the straw, becomes vulnerable to volatilization and/or photodegradation (7, 17) before reaching the soil (9). Another important aspect is the permanence period of a product on the straw. In Brazilian sugarcane plantations, products are applied during the winter season, characterized by low rainfall, especially in the southeast region (23). In addition, the longer the herbicide stays in the straw, the more susceptible it will be to degradation, consequently decreasing its transport and bioavailability for weed control (9, 27, 29, 30).

Some specific physical-chemical characteristics of the herbicides may facilitate an efficient straw-soil transport of these products. This, in addition to high solubility in water, absence of photodegradation (being preferentially degraded by microorganisms), and low  $K_{ow}$  (octanol/water partition coefficient, *i.e.* not having lipophilic character), (10) constitute key features for a successful product. Some herbicides have these physical-chemical characteristics. Among these herbicides, amicarbazone, presents high water solubility of 4.6 g L<sup>-1</sup> at pH 4-9 and a low  $K_{ow}$  (1.23); sulfentrazone, has medium water solubility of 780 mg L<sup>-1</sup> at pH 7 and medium  $K_{ow}$  of 0.16; and tebuthiuron, high solubility in water of 2.500 ml L<sup>-1</sup> at 25°C and a high  $K_{ow}$  of 67.1 (26).

Several studies have reported effective control results in dry periods with the use of amicarbazone, imazapic, sulfentrazone and tebuthiuron herbicides (6, 11, 20, 22, 23, 30). Based on the above, this study evaluated the efficacy of preemergent herbicides in different germination flushes of *M. aegyptia*, *M. aterrima*, and *R. communis* when applied on different quantities of straw and with different simulated dry periods.

#### MATERIALS AND METHODS

The study was conducted under a greenhouse in the Department of Natural Resources of the Federal University of Sao Carlos at the agricultural science campus. The experiment was replicated twice, June/July 2016 and June/July 2017.

The herbicides were applied in preemergence in a completely randomized design following a  $4 \times 2 \times 2$  factorial scheme with four replications. The variables were four dry periods, two quantities of straw, and two germination fluxes. These factors were adopted for each of the three weed species (*M. aegyptia*, *M. aterrima*, and *R. communis* L.) and the four herbicide treatments (amicarbazone, imazapic, sulfentrazone, and tebuthiuron), individually. The experimental units were composed of 25 L polyethylene pots filled with soil from the arable layer of an Eutrophic Red Latosol (table 1).

**Table 1.** Soil chemical analysis (0 to 20 cm).**Tabla 1.** Análisis químico del suelo (0 a 20 cm).

Unit: Al, H+Al, K, Ca, Mg, SB and CTC (mmolc dm <sup>-3</sup>); P (resina) (mg dm <sup>-3</sup>); V, clay, silt, sand (%). Unidad: Al, H + Al, K, Ca, Mg, SB y CTC (mmolc dm <sup>-3</sup>); P (resina) (mg dm <sup>-3</sup>); V, arcilla, limo, arena (%).

рН	M.O.	Р	K	Ca	Mg	Al+H	SB	СТС	V	Argil	Silt	Sand
(CaCl <sub>2</sub> )	(g dm <sup>-3</sup> )	(mg dm <sup>-3</sup> )			(mm	olc dm	(%)		(g kg <sup>-1</sup> )			
5.2	15	12	1.9	15	4	20	20.9	40.9	51	175	55	770

After filling the pots, 0 and 10 t ha<sup>-1</sup> of sugar cane straw ('RB966928' variety) were allocated on the pot surface. Then, the herbicides amicarbazone (1225 g ha<sup>-1</sup>), imazapic (147 g ha<sup>-1</sup>), sulfentrazone (800 g ha<sup>-1</sup>), or tebuthiuron (900 g ha<sup>-1</sup>) were applied using a  $CO_2$  pressurized, constant-pressure spray with fan-type tips (XR 110.02) at a pressure of 2.0 x 10<sup>5</sup> Pa with a syrup volume of 200 L ha<sup>-1</sup>. During applications, the temperature was 17.1°C, the relative air humidity was 85%, and the wind velocity was 0.2 m s<sup>-1</sup>.

After treatment application, the pots were submitted to four different periods without rain (0, 30, 60, and 90 days after herbicide treatment). After these periods, the pots received a rainfall simulation of 30 mm (flow rate of 1 L min<sup>-1</sup>). Finally, the pots stood for 72 hours, enough time for the straw to dry and be carefully removed.

After removing the straw, the weed species *M. aegyptia*, *M. aterrima*, and *R. communis* were individually and carefully planted in the pots at 5 cm depth, aiming for minimum soil turnover and five plants per pot. Concerning *M. aterrima*, mechanical scarification broke dormancy.

The germination flux factor consisted of two different weed sowing times in the same experimental unit (pot). The first flux occurred immediately after rainfall simulation for each of the four dry periods (0, 30, 60, and 90 DAT with no water). At 35 days after emergence, for each dry period and first germination flow, the weeds were cut and removed for dry mass analysis. At this moment, a new germination flow began. For this purpose, in the same experimental units, the weed species *M. aegyptia*, *M. aterrima*, and *R. communis* were re-sown. Thus, for each dry period and experimental unit, two germination fluxes were simulated. The first one was related to weed sowing immediately after a 30 mm rain simulation (for each dry period: 0, 30, 60, and 90 days), and the second germination flux was sown after the first flush of germination.

Weed control percentage at each germination flux and within each dry period was evaluated at 7, 14, 21, 28, and 35 days after emergence (DAE) where 0 (zero) corresponded to no injury and 100 corresponded to plant death (1).

At 35 DAE for each germination flux and within each dry period, weeds were cut, packed in cardboard bags, taken to a greenhouse, and stored at 60°C for 72 hours. After those periods, the samples were weighed. For data analysis, dry mass values were expressed as reduction percentages in relation to the control without herbicide.

Statistics consisted of the reparametrized version of the logistic model with three parameters (3, 26) (Eq. 1):

$$Y = D\{1 + \exp[B(\log X - \log E)]\}$$
(1)

where:

Y = Control and Biomass Reduction percentages

X = dry period

*D* = maximum estimate of the response variable

Parameter E = dry days estimates at 50% response

*B* = slope of the curve fitting at the inflexion point.

All statistical analyses were performed in R software (2022). The *ggplot2* (33) and *drc* (25) packages were used for graphical presentation and for fitting the Equation 1 model, respectively.

#### RESULTS

According to ALAM (1974) and Vanhala *et al.* (2004), weed control percentages of 81-90% are classified as very good and 91-100% as excellent. For flow 1, control and biomass reduction of *M. aegypta* with amicarbazone were below 80% at 90 days of drought and on straw (figure 1, page 78). The other treatments controlled more than 90%, regardless of the amount of straw. For amicarbazone, in flow 2 with straw, control of *M. aegypta* was superior to 80% at 0 and 30 dry periods, but at 60 and 90, it was under 80% (ineffective). In the application without straw, control and reduction of *M. aegypta* biomass for the same germination flow was lower than 40% only at 90 dry periods.

In flow 1, control and biomass reduction of *M. aterrima* with amicarbazone was less than 80%, only at 90 days of dry periods, regardless of the amount of straw. At 60 days of dry period biomass reduction was lower than 80%, while for flow 2, control and biomass reduction were greater than 80% at 0 dry periods with and without straw (figure 1, page 78).

At flow 1, control and biomass reduction of *R. communis* with amicarbazone was below 80% at 90 DAT of drought, with or without straw (figure 1, page 78). Control of *R. communis* at flow 2 with amicarbazone was greater than 80% in the 0 dry periods without straw. In flow 2, biomass reduction of *R. communis* with amicarbazone was inadequate, with percentages below 80% in all dry periods and amounts of straw. Control of *M. aegyptia* with imazapic at flow 1 was over 80% in all dry periods with straw in the biomass reduction was inadequate with percentages below 80% at 60 and 90 days. Flow 2 without straw showed control over 80% at the 0 dry period, while not exceeding 60% with straw. For flow 2, biomass reduction was less than 60% in all dry periods and amounts of straw (figure 1, page 78).

In flow 1, control and biomass reduction of *M. aterrima* with imazapic was greater than 80% at 0 dry period, while control of *R. communis* was greater than 80% in the dry period with and without straw, and biomass reduction was over 80% at 0, 30 and 60 days regardless of straw. Flow 2 showed biomass reduction under 80% in all dry periods and amounts of straw (figure 1, page 78).

Control of M. aegyptia for flow 1 was greater than 80% in all dry periods, regardless of the amount of straw. The reduction of biomass in flow 1 was greater than 80% at 0 and 30 dry periods with straw and 0, 30 and 60 dry periods without straw (figure 1, page 78). In flow 2, control was over 80% at 0 dry periods without straw, and biomass reduction was greater than 80% at 0 and 30 dry periods with and without straw.





Flow 1 with and without straw, and flow 2 without straw, showed control and biomass reduction of *M. aterrima* with sufentrazone over 80% at 0, 30 and 60 dry periods, regardless of the amount of straw. Flow 2 with straw, resulted in control and biomass reduction of *M. aterrima* over 80% at 0 and 30 dry periods, regardless of the amount of straw (figure 1). Flow 1, had control and biomass reduction of *R. communis* under 80% at the 90 dry period with and without straw. Control of *R. communis* in flow 2, achieved over 80% at 0 dry periods. In flow 2, biomass reduction was greater than 80% at 0 and 30 dry periods, with and without straw (figure 1).

Control and biomass reduction of tebuthiuron at flow 1 was greater than 80% in all dry periods regardless of straw, while at flow 2, control and biomass reduction of thebuthiuron were greater than 80% at 0, 30 and 60 dry periods with or without straw (figure 1, page 78).

In flow 1, control and biomass reduction of *M. aterrima* was greater than 80% at 0 and 30 dry periods with straw, and 0, 30 and 60 dry periods without straw. For flow 1, biomass reduction of *M. aterrima* without straw exceeded 80% at all dry periods while at flow 2, this species control exceeded 80% at 0 and 30 dry periods, regardless of straw. Biomass reduction exceeded 80% at dry periods 0 and 30 without straw and dry period 0 with straw (figure 1, page 78). For flow 1, control of *R. communis* was over 80% in all dry periods with and without straw, while biomass reduction was less than 80% at 90 dry periods. In flow 2, control exceeded 80% at 0 and 30 dry periods, while biomass reduction was greater than 80% at the 0 dry period (figure 1, page 78).

#### DISCUSSION

Weed control efficiency of amicarbazone was gradually reduced with longer dry periods and increasing amounts of straw. This reduction was higher at the 90 dry period and 10 t ha<sup>-1</sup> sugarcane straw. Thus, it can be noted that longer dry periods and the presence of straw on the soil surface at the time of application, reduced the efficacy of amicarbazone. Contrasting results showed how 90 days after application resulted in over 90% control of *M. aterrima* (14).

Efficacy of pre-emergent amicarbazone over *I. grandifolia, B. plantaginea, B. decumbens,* and *C. rotundus* was reduced when applied on sugarcane straw, compared to bare soil applications (19). However, this herbide showed higher efficiencies when leached from the straw by simulated rain after application (19). These results are in agreement with our study, where longer dry periods associated with amicarbazone on straw resulted in reduced weed control efficiency, probably explained by amicarbazone having higher water solubility (4.6 g L<sup>-1</sup>, pH 4-9) and low Kow (Log Kow of 1.23) (26). This contributes to low absorption and/or retention on straw and easier recovery of herbicide action by rain simulation. Thus, the higher control percentages during the first germination flux when the herbicide was directly applied to soil can be given by lower retention by straw and the consequent higher soil solution availability.

Studies on amicarbazone dynamics in sugarcane straw through HPLC/MS/MS showed that straw quantities equal to or greater than 5 t ha<sup>-1</sup> retained almost all of the herbicide at the time of application, while increasing straw quantity (mainly at 15 and 20 t ha<sup>-1</sup> sugarcane straw) reduced herbicide transport from straw to soil (9). The longer the period between herbicide application and the first rain, the lower the transport from straw to soil. However, 20 mm of rainfall at 7 and 14 days after application allowed enough recovery of the intercepted product.

Due to its high solubility, amicarbazone is easily washed from straw to soil. However, longer periods between product application on straw and the first rain may reduce product mobility, reducing weed control effectiveness. Amicarbazone's solubility can also explain the lower control percentages obtained in the second germination flux, where greater leaching in the soil solution reduced herbicide quantity in the root zone. A second eventual factor related to the lower efficiency in the second germination flux is microbial degradation of amicarbazone influenced by soil humidity and higher temperatures.

The absent residual activity of imazapyr over a second weed emergence flux, regardless of species, dry periods, and/or straw quantities, constitutes a disadvantage considering the critical period of weed infestation in sugarcane exceeding 150 days after planting (21). Therefore, herbicides with prolonged residual activity within this period are more appropriate.

Long dry periods after application of preemergent herbicides resulted in the control efficient control of different species of morning glory (*Ipomoea purpurea*) (23). Control effectiveness of imazapic diminished by 40% between 30 and 60 days of dry periods after application on *M. aegyptia*, presumably due to the high solubility (2.200 mg L<sup>-1</sup> at 25°C). In addition, this herbicide presents weak acid behavior, and low dissociation in the soil pH range between 5.0 and 7.0 (4, 15, 16). Since soil pH in this experiment was 5.2, dissociation

and bioavailability of imazapic would be practically null (16). Additionally, the experimental units received simulations of daily and constant rainfall in the greenhouse solubilizing Imazapic. Other studies showed Imazapic applied to columns with clay soil and pH of 4.7 resulting in an average of approximately 46 and 23% phytotoxicity in cucumber plants at depths of 30 and 40 cm, respectively, through an 80 mm rainfall simulation, showing the high mobility of this herbicide in acidic soils (11). Therefore, interactions between dissociation and solubility of imazapic may have resulted in greater leaching and/or degradation of this herbicide, decreasing weed absorption in the sowing period. Finally, we must also consider the quantity of herbicide absorbed in the first germination flux. This reinforces the possible high mobility of the herbicide beyond the weed seeding zone, a possible reason for the absence of residual herbicide in a second weed germination flush.

Sugarcane straw did not influence sulfentrazone effects in the first and second weed germination fluxes (for all the plants), meaning control was similar in both quantities of straw (0 and 10 t ha<sup>-1</sup>), regardless of the simulated dry period. This product efficiently controls levels in both the first and second germination fluxes. However, drought influenced weed control efficiency, since in general, control percentages decreased as dry periods increased. Difference abscence between applications may be related to the high solubility (490 mg L<sup>-1</sup>) and low Kow (1.48) of sulfentrazone, inducing low interception and/or absorption of this herbicide in sugarcane straw, in addition to favoring a good recovery of sulfentrazone initially retained by straw. This behavior results in higher soil solution availability. These results are in agreement with those of Carbonari *et al.* (2016), who found that 20 mm of water released the maximum percentage of sulfentrazone, regardless of straw quantities. For the simulated dry period after sulfentrazone application, the authors obtained recoveries of 76.5, 61.7, and 42.3% for periods of 30 and 60 days after sulfentrazone application and rain simulation.

Tebuthiuron showed excellent control of the three evaluated weeds, *M. aegyptia*, *R. communis* and *M. aterrima*, in the first germination flux. However, a noticeable reduction in control efficiency was observed in the second weed germination flux when the product was positioned on sugarcane straw. The good tebuthiuron performance in the first and second weed germination fluxes may be related to its long half-life (up to 480 days), providing soil herbicide availability for proper control of a first germination flux and residual control of a second weed emergence flux. However, when tebuthiuron is applied on sugarcane straw during dry periods, higher amounts of rain are required for an adequate release from straw to soil.

Tebuthiuron applied at 5 or more t ha<sup>-1</sup> straw resulted in almost 100% interception (29). The authors also found that lower quantities of straw resulted in higher output of the initially intercepted product. They also observed that for rainfall exceeding 20 mm, there is a tendency for the data to be similar, regardless of the quantity of straw. That is, maximum recovery capacity of the herbicide occurs with 20 mm of rain. Longer dry periods between tebuthiuron application on sugarcane straw and rainfall simulation result in less transport from straw to soil solution. The larger quantities of straw present on the soil surface at the moment of application resulted in greater interception of tebuthiuron. Additionally, longer dry periods between applications and rainfall simulation resulted in less herbicide recovery.

#### CONCLUSION

Amicarbazone herbicide presented effective control over the first weed germination flush. Straw quantity had an influence when associated with longer dry periods, while in the second germination flush, residual effects were affected by longer dry periods and the presence of straw. For Imazapic, the species presented variable control over the first germination flush, with residual effect. For Sulfentrazone, straw quantity did not have a significant influence on weed control. For Tebuthiuron, straw associated with longer dry periods reduced control percentages. However, in general, this herbicide presented enough weed control efficacy.

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## Bio-efficacy of entomopathogenic fungi and vegetable oils against the pink pineapple mealybug: *Dysmicoccus brevipes* (Cockerell)

# Bioeficacia de hongos entomopatógenos y aceites vegetales contra el piojo harinoso rosado de la piña: *Dysmicoccus brevipes* (Cockerell)

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#### ABSTRACT

Dysmicoccus brevipes (Cockerell) (Hemiptera: Pseudococcidae) is an important insect pest of pineapple worldwide due to the direct damage it causes and because it is a vector of mealybug pineapple wilt. Entomopathogenic fungi are an alternative management tool for this pest. A preliminary experiment evaluated the lethal effects of two isolates of Beauveria bassiana (BbCT, BbCa) and one isolate of Metarhizium anisopliae (Ma) against adult female D. brevipes. Subsequently, the efficacy of the most virulent isolates and a commercial strain of Paecilomyces fumosoroceus (PAE-SIN) were evaluated under laboratory and greenhouse conditions, either alone or in combination with soybean oil or neem oil. Results showed variation amongst isolates and that *B. bassiana* was the most effective. Isolate BbCa at  $1 \times 10^7$  mL<sup>-1</sup> conidia, was the most effective against *D. brevipes* nymphs and adults at 26 ± 1°C, causing 66% ± 6% mortality 8 days after inoculation. BbCa was the most virulent with an LC<sub>50</sub> of 3.45x10<sup>7</sup> mL<sup>-1</sup> conidia and a LC<sub>95</sub> of 2.29x10<sup>8</sup> mL<sup>-1</sup> conidia, under controlled conditions. Efficacy of BbCa increased when combined with neem oil, causing 100% mortality 6 days after inoculation. In conclusion, a combination of B. bassiana isolate BbCa and neem oil achieved 100% mortality in D. brevipes under the experimental conditions reported in this study.

#### **Keywords**

entomopathogenic fungi • biological control • pineapple • Dysmicoccus • vegetable oils

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#### RESUMEN

Dismiccoccus brevipes (Cockerell) (Hemiptera: Pseudococcidae) es una de las plagas de la piña de mayor importancia a nivel mundial, no solo por los daños directos que ocasiona, sino por ser trasmisor del virus marchitez roja de la piña. Los hongos entomopatógenos son una alternativa para el manejo de este insecto. En un experimento preliminar se evaluó la patogenicidad y la virulencia de dos aislamientos de Beauveria bassiana (BbCT, BbCa) y uno de Metarhizium anisopliae (Ma), sobre hembras adultas de D. brevipes. Posteriormente, los aislamientos más virulentos y una cepa comercial de Paecelomyces fumosoroceus (PAE-SIN), fueron evaluados en otro experimento en condiciones de invernadero, solos o en combinación con aceite de soya y aceite de neem. Los aislamientos evaluados presentaron diferente grado de virulencia; sin embargo, B. bassiana resultó ser el más virulento. El aislamiento BbCa a una concentración inicial de 1×10<sup>7</sup> conidios mL<sup>-1</sup> fue más efectivo contra adultos de D. brevipes comparado con el control, causando mortalidad del  $66\% \pm 6\%$  a los 8 días pos inoculación a 26  $\pm$  1°C. BbCa presentó la mayor virulencia con una  $CL_{50}$ , de 3.45x10<sup>7</sup> conidios mL<sup>-1</sup> y una CL<sub>95</sub> de 2.29x10<sup>8</sup> conidios mL<sup>-1</sup>, bajo condiciones controladas. Sin embargo, la eficacia se incrementó para BbCa, cuando se combinó con aceite de neem, al causar el 100 % de mortalidad a los 6 días pos inoculación. En conclusión, la combinación *B. bassiana* (BbCa) y aceite de neem fue el mejor tratamiento, con una mortalidad de 100% de D. brevipes bajo las condiciones experimentales reportadas en este estudio.

#### **Palabras clave**

entomopatógenos • control biológico • piña • Dysmicoccus • aceite vegetal

#### INTRODUCTION

Pineapple production generates significant economic resources worldwide. Mexico's main pineapple exports are destined for the United States market, with an annual value in 2020 of \$30,602,000 USD (24). Unfortunately, the pineapple industry is affected by various phytosanitary problems. Since pineapple is grown intensively and in monoculture, pesticides are commonly applied for pest management, causing problems for human health, the environment, and agroecosystems. The mealybugs *Dysmicoccus brevipes* (Cockerell) (Hemiptera; Pseudococcidae) and *D. neobrevipes* Beardsley (Hemiptera: Pseudococcidae) are major pests of commercial pineapple cultivation (29) causing significant damage throughout the crop growth cycle; they are also vectors of Pineapple Mealybug Wilt associated Virus (PMWaV) which can cause up to 100% of export crop losses due to rejection of fruit (19). Recent management strategies for *D. brevipes* in pineapple are largely based on synthetic organophosphate insecticides. However, efficacy of chemical control is limited by the cryptic location and behavior of these insects on plants, and their waxy surface layer which is a barrier to the action of contact insecticides, even protecting eggs from residual effects. There is also increasing concern in general about the toxic risks of excessive pesticide use in agriculture. Therefore, exploration of economically viable and environmentally safe strategies is necessary. We hypothesize that commercial pineapple production could benefit from the use of botanical extracts and biological pest control agents, such as entomopathogenic fungi, within integrated pest management (33).

Entomopathogenic fungi can infect directly without the need for ingestion and so are effective against sucking pests such as aphids, mealybugs, whiteflies, and mosquitoes (4). Some entomopathogenic fungi have a combination of modes of action against arthropods: toxins; nutrient depletion; physiological disruption; and mechanical damage to internal tissues due to mycelium development (12). Efficacy of entomopathogenic fungi has been widely documented, particularly against mealybugs. For example, *Beauveria bassiana* (Bals.) Vuill. (Ascomycota: Hypocreales), *Lecanicillium lecanii* (Zimm.) and *Metarhizium anisopliae* (Metschnikoff) Sorokin (Ascomycota: Hypocreales) infect and kill *Paracoccus marginatus* Williams & Granara de Willinks (2). Despite this, there are few laboratory and field studies on management of *D. brevipes* using entomopathogenic fungi in pineapple. One study by Miranda Vindas and Blanco Getzler (2013) has evaluated a range of options in the laboratory

that included both entomopathogenic fungi and botanical oil extracts that are known for their repellent, anti-feeding, and growth inhibition properties; high degradability; and environmental safety (18). Specific evaluations included: *B. bassiana* (4.0 x10<sup>10</sup> conidia/g); *M. anisopliae* (1.0 x10<sup>10</sup> conidia/g); a mixture of both fungi (0.5 g + 0.5 g/L distilled water,  $4.0 \times 10^{10}$  conidia/g +  $1.0 \times 10^{10}$  conidia/g); potassium salts; fatty acids (7 mL/L); and botanical extracts (a mixture of hot chili, garlic, onion, mustard and jackass bitters) (7 mL/L). Results showed high efficiency of entomopathogenic fungi and that the botanical extract achieved the fastest mortality. In the same publication, the botanical extract was also evaluated in a commercial pineapple plantation in comparison with the typical chemical control options Diazinon® 60 EC (diazinon) (0.5 ml/L) and Sevin® 80 WP (carbaryl) (1 kg/ha); the lowest incidence of mealybugs was achieved in the botanical extract treatment (16).

These results suggest that combinations of entomopathogenic fungi and vegetable oil extracts have potential as control agents that may increase mortality of adult *D. brevipes* females. For this reason, the potential of two native fungal isolates was evaluated in comparison with a commercial product based on *Paecelomyces fumosoroseus*. Our specific objectives were to determine the pathogenicity and virulence of the entomopathogenic fungi, alone and in combination with neem oil or soybean oil, against the pineapple mealybug under laboratory and greenhouse conditions.

#### MATERIAL AND METHODS

Experiments were done at the Biological Control Laboratory, Postgraduate College, Tabasco Campus, Cárdenas, Tabasco, Mexico, between January and December 2021.

#### Collection and mass rearing of D. brevipes

*Dysmicoccus brevipes* adults were collected from two varieties of pineapple (MD2 and bighead) on commercial plantations in Huimanguillo, Tabasco. Insects were taken to the Biological Control laboratory of the Postgraduate College, Tabasco Campus, for laboratory breeding, following the methods of Pandey & Johnson (2006). For the breeding stock, 50 eight-month-old pineapple cloves (25 bighead and 25 MD2 variety) were transplanted from a commercial pineapple plantation in Huimanguillo, Tabasco, into plastic pots and kept in a greenhouse at 30-35°C. Twenty days after potting, they were infested with adult female *D. brevipes* (20 per plant) in the greenhouse.

#### **Entomopathogenic fungal isolates**

Pathogenicity and virulence evaluations were made on two *Beauveria bassiana* (BbCa, BbCT) isolates and one isolate of *Metarhizium anisopliae* (Ma); all were native entomopathogenic isolates from Tabasco, Mexico held in the collection of the Biological Control Laboratory of the Colegio de Postgraduados, Tabasco Campus (table 1). Subsequent bio-efficacy experiments included a commercial product based on *Paecilomyces fumosoroceus* (PAE-SIN®).

Mycelia from each isolate was grown on sterile Sabouraud Dextrose Agar (ADS, Bioxon, Mexico) in Petri dishes, 90 x 15 mm for 3 weeks at 26±1°C in darkness. Conidia were then scraped from the surface and suspended in 0.03% Tween 80. Conidial concentration was determined using a Neubauer chamber, following the method of Inglis *et al.* (2012).

**Table 1.** Reference of the fungi used in the evaluation of pathogenicity against *D. brevipes.*

 Tabla 1. Referencia de los hongos usados en la evaluación de patogenicidad contra

 D. brevipes.

Species Key		Host	Location
Beauveria bassiana	BbCT	Hypsiphilla grandella	Huimanguillo, Tab. Mexico
Beauveria bassiana	BbCa	Hypothenemus sp.	Huimanguillo, Tab. Mexico
Metarhizium anisopliae	Ма	Aeneolamia sp.	Cárdenas, Tab. Mexico

#### Pathogenicity and virulence of D. brevipes

Pathogenicity and virulence of *B. bassiana* (BbCa, BbCT) and *M. anisopliae* (Ma) against *D. brevipes* were determined experimentally using a completely randomized design with four replicates of each treatment and control; the entire experiment was repeated on three occasions. Groups of ten adult females were each placed on two basal pieces of MD2 pineapple leaf ( $8 \times 8 \text{ cm}$ ) inside a plastic box ( $20 \text{ cm} \times 10 \text{ cm} \times 10 \text{ cm}$ ) with openings covered with organdy mesh for ventilation. Wet filter paper was placed in each box to provide moisture. Each group of adults was sprayed (from a spray bottle) with 1.5 ml of conidia (either  $10^6$ ,  $10^7$  or  $10^8 \text{ mL}^{-1}$ ) suspended in 0.05% aqueous Tween 80; the control was sprayed with 0.05% aqueous Tween 80 only. Applications were made following the methodology of Ramírez-Sánchez *et al.* (2019). Boxes containing treated insects were incubated at  $26 \pm 1^\circ$ C, 65 - 70% RH and a 14:10 h light: dark regime). Mortality was assessed daily for 8 days. Dead insects were incubated to determine cause of death (mycosis), following the methodology of Butt and Goettel (2000). Abbott's formula was used to correct data for control mortality (1).

## Bio-efficacy of B. bassiana and *P. fumosoroseus*, either alone or in combination with vegetable oils, against *D. brevipes* under greenhouse conditions

An experiment was set up under greenhouse conditions based on the results of the aforementioned bioassays. Pineapple suckers (variety MD2 [40 cm in size]) were planted individually in replicate pots, each containing 2 kg of sandy soil collected from a commercial pineapple plantation in Huimanguillo, Tabasco, Mexico. To each pot twenty *D. brevipes* adults were inoculated at the base of the pineapple sucker and incubated for one month before experimental treatments were added.

A total of eight treatments were compared including the highly virulent *B. bassiana* isolate, BbCa, and a formulated strain of *P. fumosoroseus* (PAE-SIN®), either alone or in combination with soybean oil (CARRIER)® or neem extract oil (Nimicide 80®) (table 2). All treatments were applied as 20 ml solutions/ suspensions; all fungal treatments contained  $1 \times 10^7$  conidia mL<sup>-1</sup>. There were four replicates of each treatment arranged in a completely randomized design. After inoculation, insect mortality was assessed daily for 8 days.

Treatments	Key	Conidia mL <sup>.1</sup>
B. bassiana	BbCa	1 x 10 <sup>7</sup>
P. fumosoroceus	PAE-SIN	1 x 10 <sup>7</sup>
Soybean oil	SO (CARRIER®)	1.2 mL <sup>-1</sup> L water
Neem oil	NO (NIMICIDE 80®)	1 mL <sup>-1</sup> L water
BbCa + SO	BbCa + SO	1 x 10 <sup>7</sup>
BbCa + NO	BbCa + NO	1 x 10 <sup>7</sup>
PAE-SIN + SO	PAE-SIN + SO	1.0 x 10 <sup>7</sup>
PAE-SIN + NO	PAE-SIN + NO	1.0 x 10 <sup>7</sup>
Control	Tween 80	0.05 %

**Table 2.** Treatments evaluated in the greenhouse assay.**Tabla 2.** Tratamientos evaluados en el experimento de invernadero.

#### Statistical analysis

Probit analysis was used to estimate the  $LC_{50}$  and  $LC_{95}$  of each isolate with a 95% confidence limit. ANOVA and multiple comparisons of means for both isolates and their concentrations were also done with the Tukey test (p≤ 0.05) in SAS software (25). The probit regression model  $\Phi^{-1} [\Pi(x)] = \alpha + \beta x$  and the formula LC (P) = (qnorm (P) -  $\alpha$ ) /  $\beta$  were used to estimate lethal concentrations. Virulence graphs were constructed using R software v. 1.0.143 (22).

#### RESULTS

#### Pathogenicity and virulence

Both isolates of *B. bassiana* (BbCa and BbCT) caused higher levels of *D. brevipes* mortality at all conidia concentrations evaluated compared with *M. anisopliae* at the same concentrations. There were highly significant differences amongst treatments (p < 0.0001) in the mean daily mortality after 8 days at the  $1x10^6$  conidia concentration. The highest daily mean % mortality was achieved by isolate BbCa (32.2%), followed by isolate BbCT (16.3%) and then Ma (15.8%); in the control mortality was 4.7% (figure 1 A1). However, cumulative mortality at day 8 after inoculation was 45, 40, 32.5, and 4.5% for isolates BbCa, BbCT, Ma and the control treatment, respectively (figure 1 B1).



Bars with different letters are significantly different to each other (Tukey  $p \le 0.05$ ). Error bars represent  $\pm 1$ standard error of the mean (SEM), n=4. Barras con diferentes letras son significativamente diferentes unas de otras (Tukey  $p \le 0.05$ ). Barras de error representan  $\pm 1$ el estándar error de la media (SEM), n=4.



Figura 1. Dinámica de A) mortalidad media general del periodo de 1 - 8 días post inoculación y B) mortalidad acumulada diaria de hembras adultas de *Dysmicoccus brevipes* después de los tratamientos con 1 x 10<sup>6</sup> (A), 1 x 10<sup>7</sup> (B) o 1 x 10<sup>8</sup> (C) conidios mL<sup>-1</sup>, de *B. bassiana* (BbCa, BbCT) y *M. anisopliae* (Ma). Mortality data for the 1 x  $10^7$  conidia mL<sup>-1</sup> concentration showed an increase in % mortality compared with the 1 x  $10^6$  conidia mL<sup>-1</sup> concentration. There were highly significant differences amongst treatments (p < 0.0001); the highest mean daily % mortality over 8 days was achieved by isolates BbCa and BbCT, with 60% and 58.43% mortality, respectively, followed by isolate Ma with 49.06% and the control with 7.18% (figure 1 A2, page 87). Cumulative mortality on day 8 was 95, 85, 82, and 7.18% for isolates BbCa, BbCT, Ma and the control, respectively (figure 1 B2, page 87).

Mortality data for the  $1 \times 10^8$  conidia mL<sup>-1</sup> concentration showed an even greater increase in % mortality compared with the 1 x 10<sup>6</sup> and 1 x 10<sup>7</sup> conidia mL<sup>-1</sup>. Highly significant differences among treatments (p < 0.0001) were detected, with the highest mean daily % mortality achieved by isolates BbCa and BbCT being 69.4 and 62.5%, respectively, followed by Ma (51.9%), while in the control mortality was 5.9% (figure 1 A3, page 87). Cumulative % mortality on day 8 was 100, 95, 87.5 and 7.1% for isolates BbCa, BbCT, Ma and the control, respectively (figure 1 B3, page 87).

Concerning cumulative mortality, *B. bassiana* isolate BbCa was more effective from day 5 post-inoculation than isolate BbCT. However, both isolates of *B. bassiana*, at each concentration evaluated, showed high efficacy against pineapple mealybug with increasing cumulative mortality over time after inoculation (figure 1B, page 87).

Proliferation of mycelium and conidial structures was observed on cadavers of *D. brevipes* produced during the first 4 days after inoculation. Abundant sporulation was detected from day 8 after inoculation, particularly from cadavers killed by *B. bassiana* isolates (figure 2).



**Figure 2.** *Beauveria bassiana* isolates A) BbCa, B) BbCT and C) *Metarhizium anisopliae* (Ma) infecting adult female pineapple mealybug *Dysmicoccus brevipes* at 4 x magnification with an optical microscope.

Figura 2. Aislamientos de Beauveria bassiana A) BbCa, B) BbCT y C) Metarhizium anisopliae (Ma) infectando a hembras adultas del piojo harinoso de la piña Dysmicoccus brevipes a una magnificación de 4 x con un microscopio óptico.

#### **Determination of virulence**

 $LC_{50}$  values were  $3.4 \times 10^7$ ,  $5.2 \times 10^7$  and  $8.5 \times 10^7$  mL<sup>-1</sup> conidia for isolates BbCa, BbCT, and Ma, respectively.  $LC_{95}$  values were  $2.29 \times 10^8$ ,  $2.67 \times 10^8$  and  $3.77 \times 10^8$  conidia mL<sup>-1</sup> for isolates BbCa, BbCT and Ma, respectively (figure 3, page 89). Probit regression lines for *B. bassiana* were Y = - 0.2864+8.401<sup>-09</sup> (x) and Y= - 0.3915+7.602<sup>-09</sup>(x), for BbCa, BbCT, respectively, whereas for *M. anisopliae*, it was Y= - 0.4795+5.626<sup>-09</sup>(x), where 'Y' was the probit mortality and 'x' was the fungal concentration (figure 3, page 89). Data fitted well with the model and there was a positive correlation between conidial concentration evaluated and bioinsecticidal activity of both fungi.



**Figure 3**. Corrected probit plot mortality of adult female *D. brevipes* treated with different conidia concentrations (log dose) of *B. bassiana* (BbCa [A], BbCT [B]) and *M. anisopliae* (Ma[C]) visualized using R software v. 1.0.143 (22).

**Figura 3.** Mortalidad probit corregida de hembras adultas de *D. brevipes* tratadas con diferentes concentraciones de conidios (log dosis) de *B. bassiana* (BbCa [A], BbCT [B]) y *M. anisopliae* (Ma[C]) visualizadas con el uso del programa R v. 1.0.143 (22).

#### Bio-efficacy of entomopathogenic fungi and vegetable oils against D. brevipes

Mortality of adult *D. brevipes* females following treatment with isolate BbCa (at 1 x  $10^8$  conidia mL<sup>-1</sup>) or *P. fumosoroseus* alone or in combination with vegetable oils varied significantly amongst treatments (p < 0.0001). Mean mortality in the untreated control was 5.5%. The combination BbCa + neem oil achieved the highest mortality, 98.4%, (p < 0.0001) (figure 4). Neem oil treatment alone caused 81.4% mortality. The conidial concentrations of the commercial formulation of *P. fumosoroseus* used in this experiment, caused only 55.1% mortality. However, when *P. fumosoroseus* was combined with neem oil mortality was 66.1%.

Cumulative mortality data also showed that the combination of isolate BbCa and neem oil was sufficient to achieve high efficacy by day 6 after inoculation, which shortened the time to kill by 100% compared with neem oil alone.



Bars with different letters are significantly different to each other (Tukey p ≤ 0.05). Error bars represent ±1 standard error of the mean (SEM), n= 4. Barras con

diferentes letras son significativamente diferentes unas de otras (Tukey p ≤ 0,05). Barras de error representan ±1 el estándar error de la media (SEM), n= 4.

**Figure 4.** Overall mortality of *D. brevipes* after treatment with *B. bassiana* (isolate BbCa; 1 x 10<sup>7</sup> conidia mL<sup>-1</sup>) or *P. fumosoroseus* at 1 x 10<sup>7</sup> conidia mL<sup>-1</sup>(PAE-SIN<sup>®</sup>) either alone or in combination with soybean oil (SO) or neem oil (NO).

**Figura 4.** Mortalidad total de *D. brevipes* después de los tratamientos con *B. bassiana* (aislamiento BbCa; 1 x 10<sup>7</sup> conidios mL<sup>-1</sup>) y *P. fumosoroseus* a 1 x 10<sup>7</sup> conidios mL<sup>-1</sup> (PAE-SIN<sup>®</sup>) solos o en combinación con aceite de soya (SO) o aceite de neem (NO).

#### DISCUSSION

In the present study isolates of both local native entomopathogenic fungi, *B. bassiana* and *M. anisopliae*, were pathogenic. They caused mortality and variable mycosis in adult female *D. brevipes*. The BbCa isolate was significantly more virulent than the *M. anisopliae* isolate under laboratory conditions. *B. bassiana* isolates are widely used as biological control agents of a wide variety of insect pests, including sucking pests such as mealybugs (21, 26, 32). For example, previous research showed that two isolates of *B. bassiana* (FF and PPRC-56) caused 97% and 100% mortality in adults of the mealybug *Paraputo ensete* (Williams and Ferrero) (Hemiptera: Pseudococcidae) twenty days after inoculation (14). In another study, two isolates of *B. bassiana* (GAR 17 B3, GB AR 23133) caused 67.5% and 64% mortality in the citrus mealybug *Planococcus citri* (Risso) (Hemiptera: Pseudococcidae) (5). Another isolate of *B. bassiana* caused 93% mortality in *Pseudococcus jackbeardsleyi* Gimpel & Miller (Hemiptera: Pseudococcidae) nymphs 5 days after treatment (10). The results of our research are in agreement with Mohamed (2016), who evaluated the virulence of *B. bassiana, M. anisopliae* and *Lecanicillium lecanii* isolates against adults of the mealybug,

*Planococcus ficus* (Signoret) under laboratory conditions; they reported that virulence levels of *B. bassiana* were higher than those of *M. anisopliae* and *L. lecanii*, resulting in up to 98% mortality at a concentration of  $5 \times 10^8$  mL<sup>-1</sup> conidia. Moreover, Manjushree and Chellapan (2019) reported that isolates of *B. bassiana* caused higher mortality in *D. brevipes* (Cockerell) (Hemiptera: Pseudococcidae) at a concentration of  $10^9$  conidia mL<sup>-1</sup> than isolates of *M. anisopliae* and *L. lecanii*. Like Surulivelu *et al.* (2012), Manjushree and Chellapan (2019) also reported that the same fungi were also effective against papaya mealybug *Paracoccus marginatus* (Williams) (Hemiptera: Pseudococcidae) under field conditions.

Concentration of inoculum (conidia) is a very important aspect of fungal pathogenicity and virulence. Results of our research suggest that isolate BbCa was the most virulent of the evaluated isolates with an  $LC_{50}$  of  $3.4 \times 10^7$  conidia mL<sup>-1</sup>, and that the higher the concentration of conidia the greater the mortality of *D. brevipes.*; overall 98% mortality was achieved 6 days after inoculation at  $1 \times 10^7$  conidia mL<sup>-1</sup>. These results are consistent with other studies that report *B. bassiana* caused higher mortality of various mealybug species than *M. anisopliae* (11, 13). It has been reported that foliar applications of *V. lecanii* and *B. bassiana* ( $2 \times 10^8$  conidia mL<sup>-1</sup>) in approximately 5 g/mL per L water is sufficient to reduce mealybug populations during months when the relative humidity is high (28). High effectiveness of *B. bassiana* and *P. fumosoroseus* against adult *D. brevipes* females was recorded from the 6<sup>th</sup> day after inoculation. In a field-level study, Ugalde-Trejos (2010) found no differences in the efficacy of *B. bassiana*, *M. anisopliae*, *Trichoderma* spp., and *Bacillus thuringiensis* treatments against populations of *D. brevipes* infesting pineapple.

Results of the bio-efficacy assay showed that mortality of *D. brevipes* increased when B. bassiana (isolate BbCa) and neem oil were combined, making it possible to consider this treatment for future field trials. Fernández and Juncosa (2002) reported that the use of adjuvant oils improved effectiveness of entomopathogenic fungi. In the same way, Elósegui and Elizondo (2010) found that mixtures of entomopathogenic fungi and adjuvants increased efficacy and tolerance of the product to wider ranges of temperatures. Vásquez, (2000) evaluated in vitro effectiveness of B. bassiana, M. anisopliae, Entomophthora, soap, hydrated lime, garlic extract (Allium sativum) and neem extract for control of D. brevipes in an organic pineapple plantation, where the greatest efficacy was achieved with mixtures of entomopathogenic fungi and extracts of soap, garlic, neem, and hydrated lime. Results of the present study agree with Miranda Vindas & Blanco Getzler (2013), who found that botanical extracts were highly efficient in causing D. brevipes mortality under field conditions. Gopal et al. (2021), found that the maximum cumulative mortality of Maconellicoccus hirsutus (Green) (Hemiptera: Pseudococcidae) was achieved when entomopathogenic fungi B. bassiana + L. lecanii (6 g/L + 6 g/L) were applied together rather than individually, resulting in 57.6% mortality, while neem and pongamia vegetable oils at 15 mL/L caused cumulative mortality of 81.4%, compared with the standard dose of neem oil (10 g/L) which caused 78.1% mortality. Our results showed that neem oil in combination with entomopathogenic fungi such as *B. bassiana* was the most efficient in killing adult female *D. brevipes* in greenhouse tests with up to 100% mortality by day 8 post-inoculation.

#### CONCLUSION

Both local isolates of *B. bassiana* and *M. anisopliae* were pathogenic to adult female *D. brevipes*. When *B. bassiana* and *P. fumosoroseus* were combined with neem oil under greenhouse conditions bio-efficiency increased by 20 % and 11%, respectively. *B. bassiana* (BbCa) combined with neem oil resulted in the highest mortality of *D. brevipes* reaching up to 100% by 8 days after inoculation.

We suggest that more research is needed to evaluate effectiveness of entomopathogenic fungi in combination with vegetable oils under field conditions. Design of biocontrol programs against pineapple mealybug is recommended in pineapple-growing regions of Mexico, as a strategy within integrated pest management programs. This option could reduce the use of toxic insecticides, which are harmful to the environment and human health.

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# Application stages and doses of tembotrione herbicide in grain sorghum (*Sorghum bicolor*) crop

# Etapas de aplicación y dosis del herbicida tembotrione en el cultivo del sorgo granífero (*Sorghum bicolor*)

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#### ABSTRACT

The herbicide tembotrione is effective against grassy weeds constituting an important tool in sorghum crops. However, in Brazil, this herbicide is only registered for corn. This study aimed to evaluate the selectiveness of tembotrione combined with atrazine at different doses and developmental stages of grain sorghum. Two experiments were conducted in Rio Verde and Montividiu, state of Goiás, Brazil, in 2018. A randomized block design with four replications, in a 3x2+1+1 factorial arrangement, tested three developmental stages ( $V_3$ ,  $V_5$  and  $V_7$ ), two doses of tembotrione (37.8 and 75.6 g ha<sup>-1</sup>) combined with the herbicide atrazine (1,000 g ha<sup>-1</sup>), an additional treatment only with atrazine at  $V_3$  and a control, free of herbicide. Evident phytotoxicity was observed with the combination of tembotrione and atrazine at  $V_3$  and  $V_5$  stages. Symptoms included reductions in plant height, sorghum stem diameter, panicle length, and cumulative dry mass of sorghum plant shoots. However, there was no influence on thousand grains mass, regardless of application stages. Tembotrione at 37.8 g ha<sup>-1</sup> combined with atrazine at 1,000 g ha<sup>-1</sup> was selective for grain sorghum when applied at  $V_7$ , without affecting grain yield.

#### **Keywords**

atrazine • weeds • post-emergence • yield • Sorghum bicolor

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#### RESUMEN

El herbicida tembotrione tiene un buen control de malezas y puede ser una herramienta importante en el cultivo del sorgo. Sin embargo, este herbicida está registrado en Brasil solo para maíz. Este estudio tuvo como objetivo evaluar la selectividad de la tembotrione asociada a la atrazine en diferentes dosis y etapas de desarrollo del grano de sorgo. Se realizaron dos experimentos en Rio Verde-GO y Montividiu-GO, Brasil en 2018. Se adoptó un diseño de bloques al azar con cuatro repeticiones, en un factorial 3x2+1+1, con tres etapas de desarrollo (V<sub>3</sub>, V<sub>5</sub> y V<sub>7</sub>), dos dosis de tembotrione (37,8 y 75,6 g ha<sup>-1</sup>) asociada al herbicida atrazine (1,000 g ha<sup>-1</sup>), un tratamiento adicional con solo atrazine en V<sub>3</sub> más un testigo sin herbicida. Los resultados permitieron observar síntomas visibles de fitotoxicidad cuando se aplicó la asociación entre los herbicidas tembotrione y atrazine en la etapa  $V_3 y V_5$ . Los síntomas incluyeron reducciones en la altura de la planta y el diámetro del tallo del sorgo y reducciones en la longitud de la panícula con la aplicación en la etapa V<sub>s</sub>. La aplicación de tembotrione en asociación con atrazine en V<sub>3</sub> y V<sub>5</sub> resultó en una reducción de la acumulación de masa seca en la parte aérea de las plantas de sorgo, pero sin influencia en la masa de mil granos, independientemente de la etapa de aplicación. El herbicida tembotrione en dosis de 37,8 g ha<sup>-1</sup> en asociación con atrazine en dosis de 1,000 g ha<sup>-1</sup> fue selectivo para sorgo en grano cuando se aplicó en  $V_{\gamma}$ , sin causar reducción en el rendimiento de grano.

#### **Palabras clave**

atrazine • malezas • postemergencia • productividad • Sorghum bicolor

#### INTRODUCTION

Under water scarcity and considering food security, sorghum has become an increasingly strategic crop for both animal and human feed (9, 22, 24). In addition, sorghum has a significantly lower production cost than corn (10). Due to these factors, sorghum constitutes an option for second-crop cultivation, mainly in succession to soybean (15, 24).

Like sweet corn, sorghum greater sensitivity to herbicides constitutes one key factor limiting crop expansion (7, 20), especially considering herbicides effective against narrow-leaved weeds (grasses) (21). Evidence of this higher sensitivity refers to sorghum as an indicator plant for herbicide presence in soils (19) and substrates (6). This becomes more evident when analyzing the low number of herbicides registered for sorghum pre or postemergence in Brazil: 2,4-D, atrazine, S-metolachlor, imazethapyr, and imazapic. For imidazolinone herbicides, registers only include tolerant hybrids (17). In Brazil, no selective post-emergence product against weeds can ensure sorghum crops.

Tembotrione is registered in Brazil for post-emergence applications in corn (17). This herbicide is a triketone that inhibits the enzyme 4-hydroxyphenylpyruvate dioxygenase (HPPD), essential for carotenoid synthesis, causing leaf discoloration, necrosis, and plant death (12, 26). This highly selective herbicide for corn (11, 28) could be evaluated for sorghum, a taxonomically related species (Poaceae).

Studies on post-emergence herbicide selectivity should consider adequate dosages and application stages. According to Negrisoli *et al.* (2004), selectivity cannot solely be determined by isolated assessments of visual intoxication symptoms. These authors state that several herbicides reduce crop yield without causing visual injuries.

Complementing tembotrione with atrazine seeks sorghum selective improvement in weed control (8). In this sense, this research hypothesized that selectivity of the herbicide tembotrione in association with atrazine may vary depending on sorghum phenological stage and dose. The objective was to evaluate herbicide selectivity of tembotrione combined with atrazine, at post-emergence of sorghum in different doses and developmental stages.

#### MATERIAL AND METHODS

Two field experiments were conducted in the southwest region of the state of Goiás, an important sorghum-producing region in Brasil, in the municipalities of Rio Verde (17°52′55″ S; 50°55′43″ W, 740 m altitude) and Montividiu (17°22′58″ S; 51°22′40″ W, 905 m altitude). The experiments were installed in the second crop after soybean in a no-till system, from March to July 2018.

Soils were classified as *Latossolo Vermelho-Amarelo Distrófico* and *Latossolo Vermelho distrófico* in Montividiu and Rio Verde, respectively (23). Soil analysis was performed for chemical and physical characterization at the 0.0 - 0.20 m layer. Results for Rio Verde were: pH in CaCl<sub>2</sub>: 5.9; Ca, Mg, K, Al, H+Al: 4.2; 1.3; 0.2; 0.1; 4.3 in cmol<sub>c</sub> dm<sup>-3</sup>, respectively; P: 3.8 mg dm<sup>-3</sup>; Organic matter (OM): 277 g dm<sup>-3</sup>; clay, silt, and sand: 393; 125 and 482 in g kg<sup>-1</sup>, respectively. In Montividiu results were: pH in CaCl<sub>2</sub>: 5.6; Ca, Mg, K, Al, H+Al: 3.3; 0.9; 0.4, 0.05; 4.5 in cmol<sub>c</sub> dm<sup>-3</sup>, respectively; P: 37.6 mg dm<sup>-3</sup>; OM: 226 g dm<sup>-3</sup>; clay, silt, and sand: 249; 68 and 683 in g kg<sup>-1</sup>, respectively.

According to the Köppen classification, the climate is tropical (Aw), with dry winter and rainfall concentrated in summer (2). Annual rainfall and temperature averages in Rio Verde and Montividiu are, 1,493 mm and 23.4°C and 1,512 mm and 23.0°C, respectively (3). Figure 1 shows meteorological data recorded during the experiments.



Source/Fuente: INMET - Instituto Nacional de Meteorologia. Collection station: Rio Verde (Goiás State).

Figure 1. Rainfall (mm), relative humidity (%) and average temperature (°C) in Rio Verde and Montividiu (Brazil), 2018 second crop.

**Figura 1.** Datos de precipitación (mm), humedad relativa del aire (%) y temperatura media (°C) durante los experimentos realizados en Rio Verde y Montividiu (Brasil), 2018 fuera de temporada.

The experiments were conducted in a randomized block design with four replications, in a  $3x^2+1+1$  factorial arrangement, with three developmental stages ( $V_3$ ,  $V_5$  and  $V_7$ , with three, five and seven fully developed leaves, respectively), two doses of the herbicide tembotrione (37.8 and 75.6 g ha<sup>-1</sup>) combined with the herbicide atrazine (1,000 g ha<sup>-1</sup>), an additional treatment with only atrazine applied at  $V_3$  stage and a herbicide-free control. Experimental units consisted of four 6.0 m long rows spaced 0.5 m, constituting a total area of 12.0 m<sup>2</sup> and a useful area of 5.0 m<sup>2</sup>, with two central rows, ignoring the last 0.5 m.

In both locations, the grain sorghum hybrid BRS 330 (simple, early hybrid with red grains and no tannin) was used. This genotype is widely cultivated in the Southwest region of Goiás State. Mechanical sowing at 3 cm depth was carried out in March, with 200,000 plants per hectare. According to the regional cultivation system, no fertilization was carried out after the preceding soybean crop. Herbicides were applied using a  $CO_2$ -pressurized backpack sprayer equipped with a bar with four double-fan nozzles, 110.02, with air induction, spaced 0.5 m apart. The working pressure was 2 kgf cm<sup>-2</sup>, resulting in a spray volume of 150 L ha<sup>-1</sup>. Mixture preparation included herbicide and an adjuvant based on soybean oil methyl ester at 0.1% v v<sup>-1</sup>. Climate data during application (development stages V<sub>3</sub>, V<sub>5</sub> and V<sub>7</sub>, respectively) in Rio Verde were: temperature: 23.0; 23.8 and 28.3°C, relative humidity: 65; 56 and 53% and wind speed: 2.8; 2.4 and 2.5 m s<sup>-2</sup> at V<sub>3</sub>, V<sub>5</sub> and V<sub>7</sub>, respectively. In Montividiu, the recorded data were temperature: 27.0; 28.8 and 29.6°C, relative humidity: 55; 59 and 39% and wind speed: 3.2; 1.2 and 2.1 m s<sup>-2</sup>. Regardless of herbicide action, all plots were hand-weeded, leaving sorghum plants exposed to herbicide treatments. No pest or disease control was necessary.

Responses of sorghum plants to herbicide application, *i.e.* phytotoxicity, were visually evaluated at 2, 7, 14 and 28 days after application (DAA). According to the SBCPD scale (27), percentage scores were assigned where zero is symptoms absence and 100 is plant death. At harvest, five plants randomly chosen were evaluated for plant height (from neck to the upper end of the panicle); panicle length (between base and panicle tip); stem diameter (using a digital caliper, at the first node above the ground); and shoot dry mass (weighing oven-dried shoots, after forced air circulation at 65°C). In addition, evaluations were carried out for grain yield (harvesting and threshing panicles, with subsequent cleaning and weighing of grains) and thousand-grain mass, evaluated according to the Rules for Seed Analysis (1). For both variables, moisture was corrected to 13%.

Data were tested by ANOVA at 5% significance. Initially, phytotoxicity data were transformed by the expression ( $\sqrt{x+1}$ ), for homoscedasticity and normality followed by Tukey test (p<0.05) and Dunnet test (p<0.05) for means comparison. Statistical tests were run using the Assistant statistical software (25).

#### **RESULTS AND DISCUSSION**

Treatments containing tembotrione and atrazine caused phytotoxicity symptoms in grain sorghum plants in both locations (table 1, page 98 and table 2, page 99). Observed symptoms included leaf discoloration (whitening), with a slightly reddish tint, and in most severe cases, necrosis, in accordance with Karam *et al.* (2009). At 2, 7 and 14 DAA and in both locations, the use of 75.6 g ha<sup>-1</sup> tembotrione + atrazine, caused stronger effects compared to the lowest dose (37.8 g ha<sup>-1</sup>) of tembotrione, also combined with atrazine. Furthermore, later applications (V<sub>7</sub> stage) resulted in lower phytotoxicity comparing V<sub>3</sub> and V<sub>5</sub> applications (table 1, page 98 and table 2, page 99).

At 28 DAA, the highest dose of tembotrione (75.6 g ha<sup>-1</sup>) still showed the highest phytotoxicity (table 1, page 98 and table 2, page 99). However, a tendency towards recovery of sorghum plants was observed mainly in Rio Verde, compared to the previous evaluation at 14 DAA (table 1, page 98). In this evaluation, applications at V<sub>3</sub> caused greater phytotoxicity in sorghum than those at V<sub>5</sub> and V<sub>7</sub>. Therefore, early applications of the combination of tembotrione and atrazine cause greater toxicity to sorghum plants, in accordance with Dan *et al.* (2010b), who reported that sorghum plants receiving tembotrione at three-leaf stage showed higher phytotoxicity than at five-leaf stage. This leads to the conclusion that, at more advanced stages, sorghum better tolerates tembotrione.

Atrazine alone (1,000 g ha<sup>-1</sup> at V<sub>3</sub>) did not cause any phytotoxicity in sorghum plants (table 1, page 98 and table 2, page 99), demonstrating its high selectivity when used at the said dose. Takano *et al.* (2018) found high selectivity even at higher doses (2,000 g ha<sup>-1</sup>) when applied to grain sorghum plants with three to four leaves.

Increasing the dose of tembotrione from 37.8 to 75.6 g ha<sup>-1-</sup> in combination with atrazine did not reduce plant height, regardless of the experimental location (table 3, page 100). Applications of tembotrione and atrazine herbicides at  $V_3$  and  $V_5$ , in Rio Verde, induced lower growth rates, yielding smaller plants compared to the application at  $V_7$ . In Montividiu,  $V_3$  application resulted in the highest occurrence of smaller plants. In agreement with Dan *et al.* (2010b), who concluded that tembotrione reduced growth when applied at  $V_3$  stage.

**Table 1**. Visual phytotoxicity after application of tembotrione and atrazine herbicides atdifferent doses and developmental stages of grain sorghum. Rio Verde (Brazil), 2018.

**Tabla 1.** Fitotoxicidad visual después de la aplicación de los herbicidas tembotriona y atrazinaen diferentes dosis y estados de desarrollo de sorgo en grano. Rio Verde (Brasil), 2018.

Hardeladdau (a barl)	Developmental stages									
Herbicides (g ha <sup>+</sup> )	V <sub>3</sub> V <sub>5</sub>			r 5	v	7	Mean			
Ph	ytotoxio	city 2 D	AA (%)							
Tembotrione + atrazine (37.8 + 1,000)	5.0	*	7.5	*	4.2	*	5.5	а		
Tembotrione + atrazine (75.6 + 1,000)	8.5	*	11.3	*	4.0	*	7.9	b		
Means	6.7 B 9.3 C 4.1 A									
Atrazine (1,000) at V <sub>3</sub>	0.0									
Control				0	.0					
Coefficient of variation (%)				15	5.0					
Phytotoxicity 7 DAA (%)										
Tembotrione + atrazine (37.8 + 1,000)	14.7	*	10.5	*	5.2	*	10.1	а		
Tembotrione + atrazine (75.6 + 1,000)	17.5	*	15.0	*	9.0	*	13.8	b		
Means	16.1	С	12.7	В	7.1	А				
Atrazine (1,000) at V <sub>3</sub>	0.0									
Control	0.0									
Coefficient of variation (%)	5.2									
Phy	totoxic	ity 14 E	DAA (%)	)						
Tembotrione + atrazine (37.8 + 1,000)	11.2	*	16.2	*	6.5	*	11.3	а		
Tembotrione + atrazine (75.6 + 1,000)	15.0	*	18.7	*	6.7	*	13.5	b		
Means	13.1	В	17.5	С	6.6	А				
Atrazine (1,000) at V <sub>3</sub>				0	.0					
Control				0	.0					
Coefficient of variation (%)				10	).7					
Phy	totoxic	ity 28 E	DAA (%)	)						
Tembotrione + atrazine (37.8 + 1,000)	4.0*	Aa	4.5*	Aa	4.0*	Aa	4.1	а		
Tembotrione + atrazine (75.6 + 1,000)	10.0*	Bb	5.2*	Aa	4.2*	Aa	6.5	b		
Means	7.0	В	4.8	А	4.1	А				
Atrazine (1,000) at V <sub>3</sub>				0	.0					
Control				0	.0					
Coefficient of variation (%)				8	.4					

DAA: days after application. Means followed by different uppercases, in the same row, and lowercases, in the same column, are significantly different by Tukey's test (p≤0.05). \* Significant difference from the control by Dunnett test ( $p \le 0.05$ ). DAA: días después aplicación. Medias seguidas de diferentes letras mayúsculas, en la misma fila, y minúsculas, en la misma columna, son significativamente diferentes mediante prueba de Tukey (p≤0,05). \* Diferencia significativa respecto al control mediante prueta de Dunnett (p≤0,05).

Table 2. Visual phytotoxicity after application of tembotrione and atrazine herbicides at different doses and developmental stages of grain sorghum. Montividiu (Brazil), 2018.
Tabla 2. Fitotoxicidad visual después de la aplicación de los herbicidas tembotriona y atrazina en diferentes dosis y estados de desarrollo de sorgo en grano. Montividiu (Brasil), 2018.

	Developmental stages											
Herbicides (g ha <sup>-1</sup> )	V <sub>3</sub>		<b>V</b> <sub>5</sub>		V <sub>7</sub>		Means					
	Phytoto	oxicity 2	DAA (%	<b>b)</b>		·						
Tembotrione + atrazine (37.8 + 1,000)	8.7	*	6.2	*	3.2	*	6.0	а				
Tembotrione + atrazine (75.6 + 1,000)	11.2	*	9.5	*	4.0	*	8.2	b				
Means	10.0 B 7.8 B 3.6 A											
Atrazine (1,000) at $V_{3}$	0.0											
Control				0	.0							
Coefficient of variation (%)				14	.6							
	Phytotoxicity 7 DAA (%)											
Tembotrione + atrazine (37.8 + 1,000)	12.2	*	9.5	*	4.5	*	8.7	а				
Tembotrione + atrazine (75.6 + 1,000)	16.2	*	13.7	*	6.0	*	12.0	b				
Means	14.2	В	11.6	В	5.2	А						
Atrazine (1,000) at $V_3$	0.0											
Control	0.0											
Coefficient of variation (%)				11	.2							
I	Phytoto	xicity 14	DAA (%	6)								
Tembotrione + atrazine (37.8 + 1,000)	22.5*	Ва	9.0*	Aa	7.7*	Aa	13.0	а				
Tembotrione + atrazine (75.6 + 1,000)	31.2*	Cb	13.7*	Bb	7.7*	Aa	17.5	b				
Means	26.8	С	11.3	В	7.7	А						
Atrazine (1,000) at $V_3$				0	.0							
Control				0	.0							
Coefficient of variation (%)				8	.2							
Ι	Phytoto	xicity 28	DAA (%	6)								
Tembotrione + atrazine (37.8 + 1,000)	11.2*	Ва	3.7*	Aa	2.7*	Aa	5.9	а				
Tembotrione + atrazine (75.6 + 1,000)	20.0*	Cb	6.2*	Bb	3.0*	Aa	9.7	b				
Means	15.6	В	5.0	А	2.8	А						
Atrazine (1,000) at $V_3$				0	.0							
Control				0	.0							
Coefficient of variation (%)				9	.3							

DAA: days after application. Means followed by different uppercases, in the same row, and lowercases, in the same column, are significantly different by Tukey's test (p≤0.05). \* Significant difference from the control by Dunnett test (p≤0.05). DAA: días después aplicación. Medias seguidas de diferentes letras mayúsculas, en la misma fila, y minúsculas, en la misma columna, son significativamente diferentes mediante prueba de Tukey (p≤0,05). \* Diferencia significativa respecto al control mediante prueta de Dunnett (p≤0,05).

**Table 3.** Plant height and stem diameter after application of tembotrione and atrazineat different doses and developmental stages of grain sorghum. Rio Verde and Montividiu(Brazil), 2018.

**Tabla 3.** Altura de planta y diámetro de tallo después de la aplicación de tembotriona y atrazina en diferentes dosis y estados de desarrollo de sorgo en grano. Rio Verde y Montividiu (Brasil), 2018.

Herbisides (a beil)	Developmental stages												
Herbicides (g ha <sup>-</sup> )	v	3	v	5	V <sub>7</sub>		Means						
	Plant height (cm)												
Rio Verde													
Tembotrione + atrazine (37.8 + 1,000)	108.2	*	111.1	*	119.1		112.8	а					
Tembotrione + atrazine (75.6 + 1,000)	103.2	*	102.8	*	123.5		109.8	а					
Means	105.7	В	106.9	В	121.3	А							
Atrazine (1,000) at $V_{3}$				12	4.3								
Control				12	6.9								
Coefficient of variation (%)				6	.8								
	M	Iontivid	iu										
Tembotrione + atrazine (37.8 + 1,000)	113.4	*	120.5		119.9		117.9	а					
Tembotrione + atrazine (75.6 + 1,000)	111.5	*	120.8		120.6		117.6	а					
Means	112.4	В	120.6	А	120.2	А							
Atrazine (1,000) at $V_{3}$	123.2												
Control	122.3												
Coefficient of variation (%)				2	.5								
	Stem d	liamete	r (mm)										
	F	Rio Verd	le										
Tembotrione + atrazine (37.8 + 1,000)	13.0		13.3		13.5		13.2	а					
Tembotrione + atrazine (75.6 + 1,000)	11.9	*	13.3		13.3		12.8	а					
Means	12.4	А	13.3	А	13.4	А							
Atrazine (1,000) at $V_{3}$				14	ł.6								
Control				14	ł.9								
Coefficient of variation (%)				9	.5								
	M	lontivid	iu										
Tembotrione + atrazine (37.8 + 1,000)	9.9	*	12.2		12.8		11.6	а					
Tembotrione + atrazine (75.6 + 1,000)	9.7	*	13.0		12.6		11.8	а					
Means	9.8	В	12.6	А	12.7	А							
Atrazine (1,000) at $V_{3}$				12	2.3								
Control				13	3.0								
Coefficient of variation (%)				11	.3								

different uppercases, in the same row, and lowercases, in the same column, are significantly different by Tukey's test (p≤0.05). \* Significant difference from the control by Dunnett test (p≤0.05). Medias seguidas de diferentes letras mayúsculas, en la misma fila, y minúsculas, en la misma columna, son significativamente diferentes mediante prueba de Tukey (p≤0,05). \* Diferencia significativa respecto al control mediante prueta de Dunnett (p≤0,05).

Means followed by

In Rio Verde, only the treatment tembotrione + atrazine (75.6 + 1,000 g ha<sup>-1</sup>) applied at  $V_3$  reduced stem diameter of sorghum plants compared to the control without application and the treatment containing only atrazine (1,000 g ha<sup>-1</sup>) (table 3). This was also observed in Montividiu, independently of the dose of tembotrione and atrazine applied at  $V_3$ . The reduction in stem diameter may cause lodging and consequent losses during mechanized harvesting.

In contrast, applications of tembotrione combined with atrazine at  $V_5$  or  $V_7$  did not change stem diameter (table 3). In popcorn, regardless of stage ( $V_2$ ,  $V_4$  or  $V_6$ ), Maia *et al.* (2019) detected no differences in stem diameter after application of tembotrione + atrazine (76 + 2,000 g ha<sup>-1</sup>).

Tembotrione affected panicle development in grain sorghum plants. In Rio Verde, tembotrione + atrazine (37.8 + 1,000 g ha<sup>-1</sup>) applied at V<sub>5</sub>, and tembotrione + atrazine (75.6 + 1,000 g ha<sup>-1</sup>) at V<sub>3</sub> and V<sub>5</sub> reduced panicle length compared to atrazine alone (1,000 g ha<sup>-1</sup>) and the control without herbicide (table 4). In Montividiu, this was observed only when sorghum plants were subjected to the combination of tembotrione + atrazine (75.6 + 1,000 g ha<sup>-1</sup>) at the V<sub>5</sub>, demonstrating that applications with these two herbicides at V<sub>5</sub> are not recommended.

**Table 4.** Panicle length and shoot dry mass after application of tembotrione and atrazineat different doses and developmental stages of grain sorghum. Rio Verde and Montividiu(Brazil), 2018.

**Tabla 4.** Longitud de panícula y masa seca de brotes después de la aplicación de tembotriona y atrazina en diferentes dosis y estados de desarrollo de sorgo en grano. Rio Verde y Montividiu (Brasil), 2018.

	Developmental stages											
Herbicides (g ha <sup>+</sup> )	V	3	v	5	v	7	Means					
	Panicle length (cm)											
Rio Verde												
Tembotrione + atrazine (37.8 + 1,000)	22.3		18.7	*	22.6		21.2	а				
Tembotrione + atrazine (75.6 + 1,000)	20.9	*	17.7	*	23.0		20.5	а				
Means	21.6	А	18.2	В	22.8	А						
Atrazine (1,000) at $V_{3}$				24	1.0							
Control				24	4.3							
Coefficient of variation (%)				5	.8							
				Mont	ividiu							
Tembotrione + atrazine (37.8 + 1,000)	21.0		20.9		21.4		21.1	а				
Tembotrione + atrazine (75.6 + 1,000)	20.4		19.6	*	21.5		20.5	а				
Means	20.7	А	20.2	А	21.4	А						
Atrazine (1,000) at $V_{3}$	22.1											
Control	22.5											
Coefficient of variation (%)				5	.1							
S	hoot dr	y mass (	(g plant <sup>-</sup>	<sup>1</sup> )								
	I	Rio Verd	le									
Tembotrione + atrazine (37.8 + 1,000)	96.7	*	107.5	*	129.2		111.1	а				
Tembotrione + atrazine (75.6 + 1,000)	91.5	*	103.0	*	124.5		106.3	b				
Means	94.1	С	105.2	В	126.8	А						
Atrazine (1,000) at $V_3$				12	8.7							
Control				12	8.6							
Coefficient of variation (%)				2.	30							
	N	Iontivid	liu									
Tembotrione + atrazine (37.8 + 1,000)	85.7	*	93.0	*	107.0		95.2	а				
Tembotrione + atrazine (75.6 + 1,000)	82.7	*	91.2	*	106.7		93.5	а				
Means	84.2	С	92.12	В	106.8	А						
Atrazine (1,000) at $V_3$				10	7.2							
Control				10	7.3							
Coefficient of variation (%)				6.	20							

Means followed by different uppercases, in the same row, and lowercases, in the same column, are significantly different by Tukey's test  $(p \le 0.05)$ . \* Significant difference from the control by Dunnett test  $(p \le 0.05)$ .

Medias seguidas de diferentes letras mayúsculas, en la misma fila, y minúsculas, en la misma columna, son significativamente diferentes mediante prueba de Tukey (p≤0,05). \* Diferencia significativa respecto al control mediante prueta de Dunnett (p≤0,05). Negative effects of tembotrione and atrazine herbicides at  $V_3$  and  $V_5$  include significant reductions in shoot dry mass in both locations (table 4, page 101). However, effects observed at  $V_3$  were stronger than at  $V_5$ . On the other hand, applications of this mixture of herbicides at  $V_7$  did not influence shoot dry mass. Thus, applications at more advanced stages result safer for sorghum plants. Dan *et al.* (2010a) found that tembotrione doses equal to or less than 118 g ha<sup>-1</sup> applied to millet plants with seven leaves reduced shoot dry mass by less than 10%. Even focusing on grain yield, dry mass production is important for straw mulch, and the consequent no-till sustainability. Due to biomass production, the crop becomes an excellent option for crop rotation systems (16).

No dose of tembotrione (37.8 and 75.6 g ha<sup>-1</sup>) + atrazine (1,000 g ha<sup>-1</sup>), nor application stage ( $V_{3'}$ ,  $V_5$  and  $V_7$ ) affected the thousand-grain mass in either location (table 5, page 102-103). This yield component was hardly influenced by herbicides. Similarly, in corn, tembotrione at 100.8 g ha<sup>-1</sup> at  $V_2$ ,  $V_4$ ,  $V_7$  and  $V_{10'}$  did not affect thousand-grain mass (14).

In both locations, sorghum grain yield was significantly reduced with the application of tembotrione at  $V_3$  or  $V_5$ , either at 37.8 or 75.6 g ha<sup>-1</sup> combined with atrazine (table 5, page 102-103). On the other hand, applications of these herbicides at  $V_7$  did not reduce grain yield, except for the treatment tembotrione at 75.6 + atrazine 1,000 g ha<sup>-1</sup>, in Rio Verde. In this case, yield reduction was approximately 234 kg ha<sup>-1</sup>.

Tembotrione selectivity is related to the cytochrome P-450 complex, majorly responsible for the metabolism of HPPD-inhibiting herbicides in species tolerant to active ingredients. In this study, sorghum grain yields indicated that tembotrione at 37.8 g ha<sup>-1</sup>, in combination with atrazine at  $V_{\gamma}$ , was selective when applied in the second crop in Southwestern Goiás.

The application of atrazine alone at  $V_3$  did not reduce grain yield compared to the control without herbicide (table 5, page 102-103). Previous research states that post-emergence application of 2,000 g ha<sup>-1</sup> atrazine is selective for sorghum, without affecting grain yield (21). Given reductions in grain yield after tembotrione doses over 75.6 g ha<sup>-1</sup> combined with atrazine regardless of stage, using these herbicides becomes unfeasible. After lower reductions in yield when applying tembotrione to the most developed sorghum plants (five and seven developed leaves) (6), lower doses should be considered.

These evaluations should always be done in combination with atrazine, due to greater efficiency in weed control, even when used with higher doses (8). As grain sorghum is planted soon after soybean harvest, and grain losses are inevitable, atrazine becomes a fundamental herbicide for the control of volunteer plants of the legume when applied post-emergence (4).

**Table 5.** Thousand-grain mass and grain yield after application of tembotrione andatrazine at different doses and developmental stages of grain sorghum. Rio Verde andMontividiu (Brazil), 2018.

**Tabla 5.** Masa de mil granos y rendimiento de grano después de la aplicación de tembotriona y atrazina en diferentes dosis y estados de desarrollo de sorgo granífero. Rio Verde y Montividiu (Brasil), 2018.

Harbigidas (a bai)		Developmental stages									
Herbicides (g na <sup>-</sup> )	V	3	V <sub>5</sub>		<b>V</b> <sub>7</sub>		Means				
Thousand grain mass (g)											
Rio Verde											
Tembotrione + atrazine (37.8 + 1,000)	23.6		23.3		24.7		23.9	а			
Tembotrione + atrazine (75.6 + 1,000)	23.2		22.5		24.4		23.4	а			
Means	23.4	Α	22.9	А	24.5	A					
Atrazine (1,000) at $V_{3}$	24.2										
Control	24.4										
Coefficient of variation (%)				5.	69						

	Developmental stages									
Herbicides (g ha <sup>-1</sup> )	V	3	V <sub>5</sub>		<b>V</b> <sub>7</sub>		Mea	ans		
	Montividiu									
Tembotrione + atrazine (37.8 + 1,000)	25.1		25.0		25.1		25.1	а		
Tembotrione + atrazine (75.6 + 1,000)	24.9		24.1		25.0		24.7	а		
Means	24.0	А	24.6	А	25.1	А				
Atrazine (1,000) at $V_3$				2	5.5					
Control				2	5.7					
Coefficient of variation (%)	5.72									
Grain yield (kg ha <sup>-1</sup> )										
	Rio Verde									
Tembotrione + atrazine (37.8 + 1,000)	2,765	*	2,644	*	3,456		2,955	а		
Tembotrione + atrazine (75.6 + 1,000)	2,420	*	2,506	*	3,261	*	2,729	b		
Means	2,592	В	2,575	В	3,358	А				
Atrazine (1,000) at $V_3$				3,4	178					
Control				3,4	195					
Coefficient of variation (%)				3.	65					
	M	Iontivio	liu							
Tembotrione + atrazine (37.8 + 1,000)	2,232	*	2,573	*	3,192		2,666	а		
Tembotrione + atrazine (75.6 + 1,000)	2,132	*	2,284	*	3,120		2,512	а		
Means	2,182	С	2,428	В	3,156	А				
Atrazine (1,000) at $V_3$				3,3	350					
Control				3,3	381					
Coefficient of variation (%)		6.72								

Means followed by different uppercases, in the same row, and lowercases, in the same column, are significantly different by Tukey's test ( $p \le 0.05$ ). \* Significant difference from the control by Dunnett test ( $p \le 0.05$ ).

Medias seguidas de diferentes letras mayúsculas, en la misma fila, y minúsculas, en la misma columna, son significativamente diferentes mediante prueba de Tukey (p≤0,05). \* Diferencia significativa respecto al control mediante prueta de Dunnett (p≤0,05).

#### CONCLUSIONS

Phytotoxicity symptoms are more noticeable in grain sorghum plants when combinations of tembotrione and atrazine are applied at  $V_3$  stage.

Reductions in plant height and stem diameter of grain sorghum plants are observed when tembotrione (37.8 and 75.6 g ha<sup>-1</sup>) and atrazine (1,000 g ha<sup>-1</sup>) are applied in combination at  $V_3$  stage, while panicle reduction is evidenced when applications are at  $V_5$ .

Application of tembotrione (37.8 and 75.6 g ha<sup>-1</sup>) and atrazine (1,000 g ha<sup>-1</sup>) at  $V_3$  and  $V_5$  leads to a reduction in cumulative shoot dry mass of grain sorghum plants; with no effects on thousand-grain mass, regardless of application stage.

Tembotrione at 37.8 g ha<sup>-1</sup> with atrazine at 1,000 g ha<sup>-1</sup> is selective for grain sorghum (hybrid BRS 330) when applied at the  $V_7$  stage, not reducing grain yield.

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# Traditional cow-calf systems of the northern region of Santa Fe, Argentina: current situation and improvement opportunities

## Sistemas de cría tradicionales de la región norte de Santa Fe, Argentina: situación actual y oportunidades de mejora

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#### ABSTRACT

Cow-calf systems are at the core of Argentina's significant national beef industry. The objectives were: i) to characterize the productive state of traditional cow-calf systems, named BASE, from the northern region of Santa Fe province, ii) to identify technologies for the productive improvement of the BASE system, and iii) to quantify the productive and economic impact of the adoption of the identified technologies. To characterize the BASE system, the available published data were systematized and validated in a workshop with leading regional experts in the field. To identify the technologies for improvement, a survey was conducted among regional farm advisors. Finally, to quantify the impact of adopting improvements in the BASE system, a modelling study was conducted. The results showed that traditional cow-calf systems have low productive and reproductive efficiency (45 kg LW ha<sup>-1</sup> year<sup>-1</sup> and 48% weaning rate) and little adoption of herd management and forage production technologies. The technologies identified were grazing management, training of farmers and farm staff, and seasonal mating. The modelling study showed that improvements in the production and use of forage and herd management practices would increase beef production and the gross margin of the BASE system by 70% and 96%, respectively.

#### Keywords

beef production • survey • technologies • simulation • opportunities

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#### RESUMEN

Los sistemas de cría son el núcleo de la importante industria nacional de carne bovina de Argentina. Los objetivos fueron: i) caracterizar la situación productiva de sistemas de cría tradicionales, nombrado BASE, del norte de la provincia de Santa Fe ii) identificar tecnologías para su mejora productiva iii) cuantificar el impacto productivo y económico de la adopción de las tecnologías identificadas. Para caracterizar el sistema BASE se sistematizó la información disponible que fue validada en un taller con expertos referentes de la zona. Para identificar tecnologías de mejora, se implementó una encuesta a asesores referentes de la región. Finalmente, para cuantificar el impacto de la adopción de mejoras en el sistema BASE se realizó un estudio de simulación. Los resultados demostraron que los sistemas de cría tradicionales tienen baja eficiencia productiva y reproductiva (45 kg de PV ha<sup>-1</sup> año<sup>-1</sup> y 48% de destete, respectivamente) y baja adopción de tecnologías de manejo del rodeo y producción forrajera. Las tecnologías identificadas fueron manejo del pastoreo, capacitación del productor y el personal de campo y estacionamiento del servicio. La simulación demostró que mejoras en producción y uso de forrajes y manejo de rodeo podrían incrementar la producción de carne y el margen bruto del sistema BASE en un 70% y 96%, respectivamente.

#### **Palabras claves**

producción de carne • encuesta • tecnologías • simulación • oportunidades

#### INTRODUCTION

The livestock sector faces the challenge of producing food in the context of increasing global demand for meat, which is estimated to increase by 1.6% per year (30). Intensification has been a way to improve productivity and efficiency in the beef production sector and has contributed to an increase in food production since the mid-twentieth century (18, 38). Intensification of livestock systems is defined as an increase in meat and milk production per animal and per area of land (31). In beef production systems, there are mainly two ways of intensification: through the increase in pasture production and supplementation of animals in grazing systems, or through the confinement of animals in feedlots with high feed offers (20).

Argentina produces 3.1 thousand tonnes of beef and ranks fourth among the world's beef-producing and exporting countries (43). Beef production is a relevant activity for the country's economy because it contributes 28.7% of gross domestic income and 11% of private employment within the agricultural industry (14). However, the national average weaning rate (total of weaned calves/ total of cows × 100) is lower (63%) (26) than that in other beef-producing countries such as Australia (70%) (41) and New Zealand (80%) (42). In Argentina, more than 95% of the area used for cow-calf systems is based on natural grasslands (non-cultivated environments), with poor synchronization between forage supply and livestock nutrient demand, reduced control of animal diseases (15), mainly concerning venereal and reproductive diseases (2), and low stocking rates (less than 0.50 cow ha<sup>-1</sup>) (26), resulting in low productivity, in terms of beef production per hectare (less than 90 kg ha<sup>-1</sup>) (26).

Buenos Aires province is the main beef-producing region of Argentina, and different studies have evaluated the impact of technological improvements (3, 16), technical assistance on the productivity of systems (33), and pasture production (22), among others. However, for the second most important region in calves' provision, the northern region of Santa Fe province, which provides 10% of total Argentine calves (26), there is minimal information regarding the characterization of technified systems (21), but none for traditional systems. Therefore, the objectives of the present study were: i) to characterize the productive situation of the traditional cow-calf systems (hereafter BASE system) in the northern region of Santa Fe province, Argentina; ii) to identify technologies for improving productivity based on critical technologies; and iii) to quantify the productive and economic impacts of applying technologies.

#### MATERIALS AND METHODS

#### **Description of the region**

The cow-calf systems analysed in this study are located in the north-central region of Argentina, between 28° to 30° South and 59° to 60° West in the departments of General Obligado and Vera in the province of Santa Fe. This region has approximately 900,000 ha of agricultural use (10). The climate is subtropical with average, minimum, and maximum annual air temperature of 20.1°C, 10.1°C (July) and 28.2°C (January), respectively (23). The average annual rainfall (±SD) (over the last 50 years) is 1,294 ± 310 mm, concentrated in the warmest season (82% between October and April) (24). Predominant soils belong to the Natracualf and Alfacualf groups, with drainage deficiency and saline-sodium conditions (19).

#### Productive characterization of the traditional systems

Different sources of information (scientific literature, technical reports, national and regional statistics and a workshop with local experts) were used to characterize the traditional (BASE) system in terms of land use, herd management, forage production, and productive efficiency indicators such as stocking rate (cows ha<sup>-1</sup>), weaning rate (%), and beef production (kg of calves beef ha<sup>-1</sup> year<sup>-1</sup> and kg of LW ha<sup>-1</sup> year<sup>-1</sup>).

#### Survey design: Identification and ranking of technologies to improve productivity

A digital survey (Google Forms) was designed to identify and rank technologies that could promote the improvement of traditional systems. The project's interdisciplinary team identified most region-based farm advisors and extensionists with recognized expertise in the field (n = 22) and invited them to complete the survey. The survey was structured into 10 questions. Questions 1 to 6 refer to the degree of agreement that advisors had regarding the priority of improvements in forage resources, herd management, productive and economic records, and farm infrastructure. Questions 7 to 10 refer to the technologies that advisors prioritize to improve forage resources, herd management, and farm infrastructure.

To analyze the results, radar charts were created with the average priority for each option. The prioritization patterns for each question were analyzed using principal components analysis. In addition, the relationship between the prioritizations assigned according to expertise background (agriculture or veterinary science) and work environment (private or public) of the respondents was evaluated using ANOVA. Infostat software version 2018 (12) was used for statistical analyses.

#### Simulation of productive and economic impact of applying technologies

1- Simulation model: The productive and economic impact of the adoption of the identified technologies in the survey described above was quantified through a participatory modelling approach (17) using Baqueano Cría software (40). This deterministic simulation model represents stabilized cow-calf systems and allows for monthly estimations of herd dynamics, forage and energy balance between feed supply and animal requirements, and productive and economic results. The main inputs of this model include herd composition, prices and live weight of cattle categories, monthly availability of forage, and prices of the main inputs (food, health, and labour). The main outputs included beef production (kg LW ha<sup>-1</sup> year<sup>-1</sup>) and gross margin (U\$S ha<sup>-1</sup>).

2- Simulation of BASE and improved systems: The traditional cow-calf systems, characterized in the present study (objective i) and named as BASE system, were first simulated. It was used as the baseline to simulate three further scenarios, using technologies to improve productivity and economic results (improved systems) (table 1, page 109). Based on the technologies identified as critical by the experts (objective ii of this study), three improved systems were designed (table 1, page 109): +SR+S, which includes increased stocking rate and supplementation with hay (+39% SR and +173% of hay than the BASE), +EFFICIENCY, which includes higher pregnancy rates and lower mortality rates in cows and calves; and finally +SR+S+E system was simulated, which combined the alternatives +SR+S and +EFFICIENCY. It was assumed that the greater pregnancy efficiency was the result of strategic supplementation (2.5 kg of DM of cottonseed and 1 kg of DM sorghum seed cow<sup>-1</sup> d<sup>-1</sup> between May and September) due to its incidence on the body condition of

cows (35), and mortality rates were reduced due to better health management, with greater expenses on cow health (+77% compared to BASE).

 Table 1. Characteristics of the BASE system and improved systems include: increased stocking rate (+SR), increased reproductive efficiency (+EFFICIENCY) and the combination of both alternatives (+SR+E).

Tabla 1. Características del sistema BASE y sistemas mejorados incluyendo: aumentode carga animal (+SR), aumento de la eficiencia reproductiva (+EFFICIENCY) y lacombinación de ambas alternativas (+SR+E).

Variable	Base	+SR+S	+Efficiency	+SR+S+E
Stocking rate, cows/ha	0.30	0.42	0.30	0.42
Pregnancy rate, %	62	62	85	85
Culled cows, %	18	18	5	5
Weaning rate, %	50	50	81	81
Cow mortality rate %	3	3	1	1

3- Productive and economic assumptions: Forage production and utilization for the BASE system were obtained from the database reviewed for objective (i) of this study (table 2), and the same figures were assumed for the improved systems. The mating season was assumed to occur from November to February for all systems.

**Table 2.** Average forage production (Tn DM ha <sup>-1</sup> year <sup>-1</sup>) of the traditional cow-calf systemof the northern region of Santa Fe province.

**Tabla 2.** Valores de producción (Tn MS ha<sup>-1</sup> año<sup>-1</sup>) de los recursos forrajeros de los sistemas de cría tradicionales del norte de la provincia de Santa Fe.

Environment	Dominant species	Area (ha)	Production (t DM ha <sup>-1</sup> year <sup>-1</sup> )
Grasslands	Sorghastrum setosum	300	5.5
Forest	Stipa spp.	210	3.0
Low stratum vegetation	Leersia hexandra, Luziola peruviana	90	8.3

Economic values are expressed in U.S. dollars (U\$S dollars). A cost of US\$ 16 cow <sup>-1</sup> was assumed for animal health. Full-time employees were considered for all farm tasks (180 cows), with a monthly salary of US\$744. Herd live weights and farm prices are listed in table 3. The purchase and sale expenses of the different animal categories were 5% and 2% of the price, respectively. The annual gross margin, defined as the difference between net income and direct costs (1), was also simulated, considering the prices of the main products for the region (feed, health, and labour).

**Table 3.** Herd live weight (kg head <sup>-1</sup>) and farm price (U\$S kg <sup>-1</sup>) of different animal categories in a cow-calf system in the northern region of Santa Fe province.

**Tabla 3.** Peso (kg cabeza<sup>-1</sup>) y precio (U\$S kg<sup>-1</sup>) de las diferentes categorías en un sistema de cría bovina de la región norte de la provincia de Santa Fe.

Animal class	Live weight (Kg head <sup>.1</sup> )	Price (US\$ kg <sup>-1</sup> )
Culling cow	400	1.17
Weaned steer calf	200	2.30
Weaned heifer calf	180	2.09
Heifer	290	1.72
Purchased bulls	900	1.94
Sold bulls	800	0.89

#### **RESULTS AND DISCUSSION**

# Productive characterization of the traditional cow-calf system in northern region of Santa Fe

#### Use of area and forage resources

Three contrasting vegetation environments were differentiated in the region: grasslands, forests, and low-stratum vegetation (27). Such environments are usually found in each farm in proportions of 50%, 35%, and 15% of the total area, respectively (11). The aforementioned diversity of environments poses a challenge for livestock management as they have different herbage mass rates, which implies different grazing management in each environment.

1- Grasslands: It is defined as plant communities dominated by various species where it predominates *Sorghastrum setosum* (Grise.) Hitchc (5, 34). The forage contribution to livestock in these environments varies from 3,000 to 6,000 kg DM ha<sup>-1</sup>. Other species with high forage value, such as legumes (*i.e.*, genus *Desmodium*, *Desmanthus*, and *Vicia*) and grasses of the genus Paspalum (5), can be found in this environment.

2- Forest: The predominant species in this environment was *Schinopsis balansae* Engl. Plant communities in the forest are dominated by species of the genera *Stipa* and *Piptochaetium* (28). These environments provide forage for cattle in variable quantities and quality (1,000-5,000 kg MS ha<sup>-1</sup>) according to the state of forest conservation.

3- Low stratum vegetation: These environments are dominated by hygrophilous herbaceous communities dominated by grasses such as *Echinochloa helodes* (Hackel) Parodi, *Leersia hexandra* Sw., and *Luziola peruviana* Juss. Ex J.F. Gmel., with a dry matter production of 6,000 to 8,000 kg ha<sup>-1</sup> (34).

Improvement of forage production through fertilization or introduction of cultivated species such as perennial pastures or annual forage crops is almost null among the traditional farms in the northern region of Santa Fe province. Cultivated forage species are usually no more than 2% of total area in cow-calf systems some cultivated species are *Avena sativa* L., *Melilotus albus* Medik, *Medicago sativa* L., *Sorghum bicolor* L. Monech and *Chloris gayana* Kunth (6).

#### Productive and reproductive efficiency and herd management

Mating is continuous throughout the year, with little adoption of herd management and health technologies, such as venereal disease control (13). The age at the first mating is usually greater than 24 months. Supplementation of heifers is carried out occasionally with pasture hay (less than 1 kg DM animal<sup>-1</sup>) in winter and, to a lesser extent, energy concentrates, such as corn and sorghum grains (6). Calve weaning is performed at 8 months of age, the weaning rate is 48%, and beef production is approximately 45 kg LW ha<sup>-1</sup> year<sup>-1</sup> (6).

#### Survey results: opportunities for technological improvement

There was a high level of answers (86% of the invited regional consultants). Respondents were highly experienced experts in veterinary sciences (42%) and agriculture science (58%). The results are presented in table 4 (page 111) and figure 1 (page 112). Priority given to improve herd management was higher for professionals working in the private sector (p < 0.05), and in general, answers for each aspect (forage supply, herd management practices, and farm infrastructure) were independent of the career and the field of work of the respondents (p > 0.05).

Table 4. Questions 1 to 6 used in the survey to regional farm advisors and answers.
Tabla 4. Preguntas utilizadas en la encuesta a asesores referentes y respuestas.

Questions	Possible answers	Answers
	Disagree	0.0%
Do better and more precise information on the	Little agree	5.3%
production and quality of NATURAL forage resources be	Quite agree	21.1%
prioritized for the improvement of cow-calf systems?	Strongly agree	15.8%
	Totally agree	57.9%
	Disagree	0.0%
Do better and more information on production and	Little agree	21.1%
quality of cultivated forage species be a priority for the	Quite agree	26.3%
improvement of cow-calf systems in northern?	Strongly agree	47.4%
	Totally agree	5.3%
	Disagree	0.0%
Do herd management (health, seasonal mating,	Little agree	0.0%
pregnancy diagnosis, and weaning management) be a	Quite agree	5.3%
priority for the improvement of cow calf systems?	Strongly agree	36.8%
	Totally agree	57.9%
	Disagree	0.0%
	Little agree	0.0%
Are productive or economic records priorities for the improvement of cow-calf systems?	Quite agree	26.3%
	Strongly agree	36.8%
	Totally agree	36.8%
	Disagree	0.0%
	Little agree	0.0%
Do you think that farmers make little use of the available records?	Quite agree	15.8%
	Strongly agree	36.8%
	Totally agree	47.4%
	Disagree	0.0%
	Little agree	5.3%
Are infrastructure improvements a priority for the improvement of cow-calf systems?	Quite agree	47.4%
improvement of cow can systems.	Strongly agree	10.5%
	Totally agree	36.8%

#### Productive and economic impact of technological improvements

The results of the modelling studies are shown in figure 2 (page 112). All three improved systems resulted in higher beef production and a higher gross margin than those of the BASE system. The +SR+S+E alternative showed an increase of 70% and 96% in beef production and gross margin, respectively, compared with the BASE system, despite showing higher direct costs (figure 2, page 112). These results agree with previous simulation studies (16, 17) conducted in other regions of Argentina, which showed that the combination of increased SR increased supplementation, and better reproductive management (similar to +SR+S+E in this study) would increase productive and economic results to a greater extent than if they are implemented as sole alternatives.

A change in stocking rate directly influences income as it correlates with the growth of livestock capital. However, it's essential to note that the economic efficiency of agricultural systems can be significantly influenced by factors beyond the scope of this study, such as the land tenure regime (39).



Figure 1. Technologies prioritized by advisors, (10 maximum, 1 minimum).

(1.a.) Priority of potential technological improvements.

Figura 1. Tecnologías priorizadas por los asesores, (10 máximo, 1 mínimo). (1.a.) Prioridad de mejoras tecnológicas potenciales.



**Figure 2.** Beef production (kg LW ha<sup>-1</sup> year<sup>-1</sup>) and gross margin (MB, US\$ ha<sup>-1</sup> year<sup>-1</sup>) of BASE system and improved alternatives.

**Figura 2.** Producción de carne (kg PV ha<sup>-1</sup> año<sup>-1</sup>) y margen bruto (MB, U\$S ha<sup>-1</sup> año<sup>-1</sup>) del sistema BASE y las alternativas mejoradas.

supply, (1.c.) herd management, (1.d) infrastructure. Prioridad en: (1.b.) oferta forrajera, (1.c.) manejo del rodeo, (1.d.) infraestructura.

Priority in: (1.b.) forage



Table 5 shows previous studies and compares the contrasting productive parameters between the traditional system and existing top technological systems (high use of technologies) in the same region (21). The productive potential of current top cow-calf systems (those having greater technological adoption and management skills compared to traditional farmers in the region) in this region has been recently estimated (21) and the technological gap with the BASE system is 86% in beef productivity (kg/ha/year) and 44% in weaning rate (table 5). This difference is based on the application of technologies that increase forage supply (greater area of cultivated pastures and annual forage crops) and improve herd management techniques, such as greater supplementation of cows, higher stocking rate, seasonal mating, and shorter age for first mating and weaning applied in the top systems compared to the traditional systems.

**Table 5.** Productive differences between the traditional and top cow-calf systems of thenorthern region of Santa Fe province.

<b>Productive Variables</b>	Unit	traditional system	Top system (21)
Stocking rate	cows ha -1	0.30	0.46
Weaning rate	%	48	69
Calf beef production	Kg LW ha <sup>-1</sup> year <sup>-1</sup>	29	51
Beef production	Kg LW ha <sup>-1</sup> year <sup>-1</sup>	45	83
Grasslands area	% of total area	100	90
Area with cultivated species	% of total area	0	10
Age at first mating	month	27-36	27
Mating strategy		Continuous	Seasonal (4 months)
Weaning age	Months	8	6

**Tabla 5.** Diferencias productivas entre sistemas tradicionales y tecnificados de la regiónnorte de la provincia de Santa Fe.

Fernandez-Rosso *et al.* (2020) reported 63% more beef production and 340% higher gross margin in systems that combined herd management technologies such as early weaning (2 to 4 months) and implantation of cultivated forage species, in the southwest of Buenos Aires province, compared with traditional systems of that region.

Data available from net aerial primary productivity (NAPP) and the quality of forage available in the region under study are mainly reported for cultivated pastures (32, 36). The productive and economic simulations carried out in this study were based on NAPP data of natural forage resources using a combination of unpublished data of forage cuts validated by experts (table 2, page 109). However, alternative methodologies that allow for the estimation of NAPP have been applied with promising results in other regions of Argentina, such as the green index (22), simulation models (4), and regression equations for forage cuts (17), and could be used in future studies.

In the northern region of Santa Fe Province, there have been several public policies aimed at assisting farmers in improving the productive efficiency of cow-calf systems through subsidized loans and farm advisory support by applying and monitoring health, nutritional, and reproductive management technologies (7, 29). However, the low adoption of technologies and the current low productive and reproductive efficiency (table 5), which have remained stable for years (8, 9), reflect the low effectiveness of those policies. This situation encourages a deeper understanding of the causes of farmers' scarce technological adoption. In other important beef cattle breeding regions of the country, barriers to the adoption of technologies in farming systems are mentioned. In cow-calf system studies located in Buenos Aires province, it has stood out (17, 35) as adoption barriers of technologies, the absence of suitable public policies for the region, and the producers' partial dedication to the activity. Additionally, barriers related to the lack of agricultural vocation among heirs and the absence of technical assistance in low-tech systems have also been described (17).

The applied participatory modelling methodology (17) provided preliminary information and a "what if" analysis (25) of this important productive area. However, the productive characterization of cow-calf traditional systems carried out in this study will require additional research to refine farm information and to define barriers to technological adoption in breeding systems in northern Santa Fe. This understanding might aid in the better design of public policies, which should include the social and cultural conditions of farmers (37). This methodology was also key to the conservation and sustainable development of livestock systems in other countries (44).

#### CONCLUSIONS

We combined the available scarce data on traditional cow-calf systems in the northern region of Santa Fe Province with the qualified knowledge provided by highly experienced farm advisors, in order to establish a benchmark and to identify challenges for future studies. Experts prioritized the improvement of forage supply and herd management to increase the productivity of cow-calf systems. Implementation of a rational grazing system for grasslands, training the farmer and farm staff on herd management, and seasonal mating were the factors selected to be adopted in the first place. The modelling study showed that increased SR, higher supplementation and higher reproductive efficiency increased production and economic results by 70 and 96%, respectively. The participatory modelling methodology applied also allowed us to identify areas in which greater research efforts are needed, such as more precise research information on farm characterisation, forage production and quality, and farmers' constraints for technological adoption, which will be relevant inputs for designing and promoting effective policies for the livestock sector.

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# Nutritional quality of amaranth (*Amaranthus*) silage in response to forage airing and addition of lactic bacteria

## Calidad nutricional de amaranto (*Amaranthus*) ensilado en respuesta a la aireación del forraje y a la adición de bacterias lácticas

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#### ABSTRACT

Climate change is reducing forage availability for ruminants. Previous studies in Northern Patagonia, Argentina, have demonstrated the adaptation of the amaranth crop to these agroclimatic conditions under irrigation. Moreover, this crop is used as forage in marginal areas of the world, given its outstanding productive and nutritional qualities. The objective of this study was to evaluate the nutritional quality of amaranth silage in response to previous wilting and the addition of lactic acid bacteria. The crop was harvested at the milky grain stage and ensiled in experimental microsilos for 60 days. Before ensiling, different treatments (wilting and addition of lactic acid bacteria) were applied. Parameters related to nutritional quality were evaluated, including ash, crude protein (CP), neutral detergent fiber (NDF), acid detergent fiber (ADF), dry matter digestibility (DMD), and metabolizable energy (ME). Simultaneous treatment with air and the addition of lactic acid bacteria before ensiling resulted in the best nutritional quality characteristics of the silage. The most significant results were protein value of 12.7%, 41.1% NDF and 19.1% FDA. The DM and ME were 74% and 2.67 Mcal/kg, respectively. Thus, amaranth silage can be considered an alternative conserved forage for animal feed in this region.

#### Keywords

nutritional value • amaranth • silage • inoculation • quality

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#### RESUMEN

El cambio climático está reduciendo la disponibilidad de forraje para los rumiantes. Estudios previos en la Patagonia Norte Argentina demuestran la adaptación del cultivo de amaranto a estas condiciones agroclimáticas bajo riego. Sin embargo, este cultivo se utiliza como forraje en zonas marginales del mundo, dadas las destacadas cualidades productivas y nutricionales del cultivo de amaranto. El objetivo de este estudio fue evaluar la calidad nutricional del ensilado de amaranto en respuesta al oreado previo del forraje y a la adición de bacterias lácticas. El cultivo se cosechó en el estado grano lechoso y se ensiló en microsilos experimentales durante 60 días. Antes del ensilado, se llevaron a cabo diferentes tratamientos (oreado y adición de bacterias lácticas). Se evaluaron parámetros relacionados con la calidad nutricional: cenizas, proteína cruda (PC), Fibra Detergente Neutro (FDN), Fibra Detergente Ácido (FDA), digestibilidad de la materia seca (DMS) y energía metabolizable (EM). El tratamiento simultáneo de oreado y adición de bacterias lácticas antes del ensilado dio lugar a las mejores características de calidad nutricional del ensilado obtenido. Los resultados más importantes bajo estas condiciones fueron valores de proteína del 12,7%, FDN del 41,1% y FDA del 19,1%. La DMS y la EM fueron del 74% y 2,67 Mcal/kg, respectivamente. Así, el ensilaje de amaranto puede ser considerado como una alternativa forrajera conservada para la alimentación animal de la región.

#### **Palabras claves**

valor nutricional • amaranto • ensilado • inoculación • calidad

#### INTRODUCTION

The projected growth in global food demand has led to an increased focus on underutilized crops, with the potential to improve global food security and mitigate the adverse effects of climate change. Animal production in the Lower Rio Negro Valley is limited by a seasonal lack of forage. Feed and conserved fodder, such as silage, ensures forage quality and quantity. The main silage resource in the area is corn, with a biomass yield of over 30 t/ha and a protein content of 8%. However, its cultivation requires 900 mm of water and 300 kg/ha of nitrogen (27). Amaranthus cruentus cv Mexicano has shown adaptability to the local environmental conditions with yields of 21 t/ha, needing 800 mm and 150 kg/ha of water and nitrogen requirements, respectively (39, 40, 41). Therefore, amaranth could be an alternative forage resource for the region given its high biomass production and low management requirements. Seguín et al. (2013) stated that amaranth shows high rumen degradability when used as animal feed, either by direct grazing or as conserved forage. The nutritive value of this forage varies according to the developmental stage, standing out for its high crude protein content (8-29%), low lignin values (1.7-7.3%), and high in vitro digestibility (59-79%) (23, 28). However, high moisture and protein content could negatively affect the ensiling process. According to Borreani et al. (2018), moisture content produces effluents that reduce soluble carbohydrates, whereas high protein values in the forage have a buffering capacity and prevent pH decrease (<5.0) in the silage. A common practice to decrease forage moisture is to wilt the forage in the field for 24-48 hours under good weather conditions (no risk of rain) and with minimal mechanical treatment. This increases dry matter, resulting in a higher concentration of soluble carbohydrates that favors the fermentation process and nutritive value (37). The use of additives, such as bacterial inoculants, could favor a rapid decrease in pH, improving the conservation and nutritive value of silage (4, 32). Among the most commonly used inoculants are lactic acid bacteria (LAB), which ferment carbohydrates into lactic acid, acidifying the silage and inhibiting growth of undesirable bacteria (24), thus improving the fermentation and aerobic stability of the final product. Although silage techniques have been studied for this crop, different results depend on variety, cutting time, processing methodology, and place of origin (16, 20, 28, 29). In South America, there is no information on amaranth as fodder or its conservation in silage. In this study, we hypothesized that wilting and LAB inoculation would improve fermentation quality of amaranth forage. The objective of this study was to evaluate the nutritional quality of amaranth silage in response to previous wilting of forage and addition of lactic acid bacteria.

#### MATERIALS AND METHODS

#### Site location, climatic conditions, and forage production

Field experiments were conducted at the INTA-Estación Experimental Agropecuaria Valle Inferior del Río Negro (40°48' S, 63°05' W; 4 m). This area in Patagonia, Argentina, has an irrigation system that covers 24,000 ha. During the growth period of amaranth in 2019 (November to April), rainfall was 186 mm and the average temperature was 19°C. The physicochemical characteristics of the upper 50 cm of the experimental loam soil were: pH (8.20); electrical conductivity (1.2 mmhos cm<sup>-1</sup>); organic matter (3.8%); total nitrogen (0.18%); N-NO<sub>3</sub> (24.60 mg kg<sup>-1</sup>); P (Olsen, 16.60 mg kg<sup>-1</sup>); S (14.7 mg kg<sup>-1</sup> as a SO<sub>4</sub><sup>=</sup>); Ca (8.230 mg kg<sup>-1</sup>); Mg (1.170 mg kg<sup>-1</sup>); sodium-adsorption ratio (1.83). The INTA laboratory performed the soil characterization.

The cultivar evaluated was *A. cruentus* cv Mexicano, sown in rows with a horticultural seeder at the end of spring (November 20<sup>th</sup>). The sowing area was 70 m<sup>2</sup> (10 furrows 0.7 m wide × 10 m long). Weeding was performed manually when the plants reached 20-30 cm in height; thinning was performed by hand, leaving ten plants m<sup>-2</sup>. Furrow irrigation was applied according to soil moisture retention curve before reaching the permanent wilting point, with a total lamina of 800±50 mm. Fertilization with granulated urea (46% N), with a nitrogen (N) dose of 90 ha<sup>-1</sup>, was carried out in two stages; on plants 60 cm high and at bloom.

#### Treatment of plant material before the silage process

The forage was cut at advanced flowering stage (between milky-pasty grains), corresponding to a chronological time of 123 days or 1627 growing degree days, with dry matter (DM) values above 20%. Amaranth plants were manually cut 50 cm above the soil surface (26) and then wilted in the field for 24 hours before chopping the material. The plants were chopped using a Thomas Willey mill until they reached a size of 1-3 cm. The chopped material was then treated with lactic acid bacteria (LAB) under conditions recommended by the manufacturer using an atomizer. (Bemix Plus® 2% w/v). Four different types of silage were obtained: unwilted, ensiled amaranth (UWAE); unwilted, ensiled amaranth with added lactic acid bacteria (UWAEL); wilted, ensiled amaranth (WAE); and wilted, ensiled amaranth with added lactic acid bacteria (WAEL).

#### **Ensiling process**

The silage was prepared at laboratory scale using 110mm PVC tubes 30 cm long, known as experimental microsilage. The filling of the experimental microsilage was performed by placing the plant material in compacted layers with a hydraulic press (140 kgf) ensuring homogeneous compaction. The tubes were capped and sealed to achieve anaerobic conditions (four replicates for each forage type). The experimental microsilage was maintained under environmental conditions for 60 days.

#### Characterization and chemical analyses of the forage and silage types

Chop size: A fresh sample (100 g) of non-ensiled plant material was measured using a Vernier caliper and grouped according to length. After the ensiling period, the average temperature was determined with an infrared thermometer (AMPROBE IR-710) and a sample (50 g) was collected at a depth of 15 cm from each silo. All samples were frozen at -18°C until use for quality determination. An aqueous extract (1:4) was made and left to rest for 1 h at room temperature, after which pH was measured with a pH meter (Foodcare HI99161) (10). Dry matter content (% DM) was determined according to the AOAC (2000). The samples were ground using a grinder (ARCANO®). Ash content, crude protein (CP%), Neutral Detergent Fiber (NDF %), and Acid Detergent Fiber (ADF %) were determined following the methodology proposed by the AOAC (2000). The dry matter digestibility (% DMD) was estimated using the predictive equation proposed by Rohweder *et al.* (1978). Metabolizable Energy (ME) was determined using the following formula: EM: 3.61\*DMD according to Di Marco (2011). All measurements were performed in triplicate.

#### Statistical analysis

A completely randomized block design was used, with four treatments and four replicates for each treatment. Pre-ensiling quality variables were analyzed using simple ANOVA (WA-C; UWA-C). Silage quality parameters were analyzed by double ANOVA with the following sources of variation: wilting (UWAE; WAE), inoculation (UWAEL; WAEL), and their interaction as main effects ( $2 \times 2$  factorial design). The T° and pH showed no interaction, while DM, CP, ash, NDF, and SDF were analyzed by simple ANOVA to evaluate the effect of wilting and addition of lactic acid bacteria separately. Comparisons of means were made using Fisher's least significant difference (LSD) test at 5%. Statistical analyses were performed using InfoStat (8).

#### **RESULTS AND DISCUSSION**

#### Pre-ensiling characteristics of forage

Visual differences in coloration between UWA-C and WA-C were observed in the chopped plant material (data not shown). The latter had an opaque yellow-olive color, whereas UWA-C was bright olive-green, which could be associated with the loss of water during the wilting process. In addition, drying plant material in the sun results in the formation of indigestible protein-carbohydrate complexes called Maillard products. The DM values are presented in table 1. Knowing the proper DM content of forage at the pre-ensiling stage is essential to achieve efficient compaction and an anaerobic environment while preventing growth of undesirable microorganisms. DM content of 30-35% is usually recommended to obtain high-quality amaranth silage (14). Krawutschke *et al.* (2013) achieved an optimum DM content of 30-35 % when wilting red clover (*Trifolium pratense L.*), which like amaranth is difficult to ensile due to a lower water-soluble carbohydrate concentration (WSC), higher buffering capacity (BC) and lower dry matter (DM) content at harvest. In amaranth, wilting (WA-C) resulted in dry matter values that reached the recommended values for silage; therefore, wilting could favor a higher concentration of soluble carbohydrates and a lower concentration of lactic acid (21).

# **Table 1.** Nutritional parameters of unwilted amaranth control (UWA-C) and wiltedamaranth control (WA-C).

	DM %	CP %	NDF %	ADF %	DMD %	ME Mcal/kg DM
UWA-C	20.3 b	15.8 a	46.1 b	26.4 b	68.3	2.5
WA-C	34.8 a	14.1 b	51.7a	27.6a	67.4	2.4
Standard error	0.36	0.09	0.58	0.27		
p-value	0.0001	0.0007	0.0001	0.0025		

**Tabla 1.** Parámetros nutricionales del forraje de amaranto no oreado control (UWA-C) yforraje de amaranto oreado control (WA-C).

Dry matter (DM); Crude Protein (CP), Neutral Detergent Fiber (NDF); Acid Detergent Fiber (ADF), Dry Matter Digestibility (DMD), Metabolizable Energy (ME) associated with the quality of the forage before silage.

Values of the same variable followed by the same letter are not statistically different according to Fisher's LSD test (p > 0.05).

Materia Seca (MS); Proteína Cruda (PC), Fibra Detergente Neutra (FDN); Fibra Detergente Ácida (FA), Digestibilidad de la Materia Seca (DMS), Energía Metabolizable (EM) asociados a la calidad del forraje antes del ensilaje.

Los valores de las mismas variables seguidos de la misma letra no son estadísticamente diferentes según la prueba LSD de Fisher (p > 0,05).

On the other hand, table 1 (page 120), shows the nutritional parameters of the material before the ensiling process and statistically significant differences (p> 0.001) were observed between the UWA-C and WA-C treatments. Forage total N determines the total amount of N available for animal consumption. In this study, the %CP of UWA-C was higher than that of WA-C, possibly because the drying process favors loosing nonstructural carbohydrates, volatile organic substances, and protein degradation. This results in amino acids with higher solubility that can be removed from plant tissues, and although this process can significantly decrease the %CP, it varies depending on the drying method. N decreased by only 5% in this case because of wilting conditions. WA-C had the highest fiber content (NDF and SDF), possibly because exposure of the material to ultraviolet radiation (wilting) decreased forage soluble components. In this study, NDF content ranged from 46.1% to 51.7%, and FAD content ranged from 26.4% to 27% when amaranth plants were wilted. Fiber components, such as NDF, FAD, and lignin, are generally inert during plant respiration, but their concentrations increase because of a decrease in oxidized soluble compounds. Therefore, slow-drying methods are expected to result in higher proportions of NDF and FAD. The digestibility of UWA-C was significantly higher (p > 0.001) than that of WA-C, mainly given by the lower fiber content of the undried plant material.

On the other hand, the size of the plant material obtained in this study before ensiling was between adequate ranges (1-3 cm), coinciding with the recommendations of Citlak and Kilic (2020).

#### Determination of parameters related to fodder processing and conservation

A double ANOVA determined that the wilting and inoculation treatments did not affect the temperature and pH variables. However, as shown in table 2, significant differences in T° were observed with the wilting effect, where the minimum and maximum values (UWAE and WAE, respectively) were recorded, but there were no differences in this parameter after LAB addition. Temperature affected silage fermentation, and the best results were obtained with a moderate T between 20 and 30°C. For corn silage, Zhou *et al.* (2019), reported that increasing storage temperature from 5°C to 25°C did not affect fermentation profiles of most biochemical parameters or bacterial and fungal populations. In the present study, the low temperatures observed when opening the experimental microsilage could be related to low ambient temperature at the study site.

 Table 2. Parameters related to fodder processing and conservation.

 Tabla 2. Parámetros relacionados con el procesado y conservación del forraje.

Treatment	T° (°C)	рН
UWAE	15.95 ± 1.42 b	4.6 ± 0.13b
WAE	18.10 ± 1.17 a	4.9 ±0.22a
UWAEL	16.05 ± 1.86 ab	4.5 ± 0.04b
WAEL	17.50 ± 0.71 ab	4.6 ± 0.71b
Standard error	0.68	0.05
p-valor	0.6158	0.0595

Temperature (T°), and pH of the experimental microsilage at the moment of opening for unwilted amaranth ensiled (UWAE); unwilted amaranth ensiled with added lactic acid bacteria (UWAEL); wilted amaranth ensiled (WAE); wilted amaranth ensiled with added lactic acid bacteria (WAEL).

Values of the same variable followed by the same letter are not statistically different according to Fisher's LSD test (p > 0.05).

Temperatura (T°), y pH de los microsilos experimentales en el momento de apertura. Amaranto ensilado sin orear (UWAE); amaranto ensilado oreado (WAE); amaranto ensilado sin orear con adición de bacterias lácticas (UWAEL); amaranto ensilado oreado con adición de bacterias lácticas (WAEL).

Los valores de las mismas variables seguidos de la misma letra no son estadísticamente diferentes según la prueba LSD de Fisher (p > 0,05).

When anaerobic fermentation occurs in the silage process, lactic acid bacteria in the forage are desirable because they rapidly lower the pH, achieve rapid stabilization, and maintain the characteristics of the ensiled material (9, 36). The pH (average of 4.65) observed for all silages was within the established values for good-quality silages, which is comparable with the values reported for different amaranth varieties, harvest stages, and place of cultivation (26). Several factors may be responsible for silage having a higher than normal pH (~4.0), including buffering capacity provided by high protein content (17). In UWAEL and WAEL, the use of bacterial inoculants slightly increased acidity of the experimental microsilage. Similar results have been obtained for corn, oats, and amaranth silages (4, 20). In alfalfa, a forage with high buffering capacity, wilting pretreatment, and a pH of 5.19 was reported after 60 days of ensiling (11). In this study, silage with pre-wilting and higher DM (WA-C, table 1, page 120) showed a slight increase in T° and pH; however, the silage obtained with wilted forage and addition of lactic acid bacteria (WAEL) only showed a decrease in pH.

Other authors have reported that lactic acid bacteria in temperate to cold environments favor a lower pH and rapid production of desirable metabolites, such as lactic acid, accelerating fermentation, and better-conserving silage nutrients over a wide range of growth temperatures (4, 13).

#### Nutritional quality of ensiled plant material

For the DM, CP, Ash, NDF, and ADF variables, the double ANOVA detected interactions between sources of variation (wilting and inoculation with LAB); therefore, the results are presented separately (table 3). When DM was determined after the ensiling process, values of approximately 20% were observed in the unwilted plant material (UWAE and UWAEL) and values approached 34% in the wilted material (WAE and WAEL). These values are similar to those obtained before ensiling (table 1, page 120). Good fermentation results in DM losses of less than 10% (3), and the dry matter reduction was <1% in our case.

Without LAB	DM (%)	CP (%)	ASH (%)	NDF (%)	ADF (%)	DMD (%)	ME (Mcal/kg DM)
UWAE	20.24b	11.14b	14.19a	48.58a	26.36a	68.37	2.47
WAE	34.47a	13.43a	14.31a	44.07b	24.30b	69.97	2.53
Standard error	0.58	0.12	0.01	0.06	0.03		
p-value	0.0001	0.0001	0.0603	0.0001	0.0001		
With LAB	DM (0/ )	CP (%)	ASH (%)	NDF (%)	ADF (%)	DMD	ME
	DM (%)	CI (70)	1011 (70)	NDI (70)		(%)	(Mcal/kg DM)
UWAEL	20.66b	13.32a	15.23a	48.80a	26.61a	<b>(%)</b> 68.17	(Mcal/kg DM) 2.46
UWAEL	20.66b 33.59a	13.32a 12.71b	15.23a 12.87b	48.80a 41.11b	26.61a 19.12b	(%) 68.17 74	(Mcal/kg DM) 2.46 2.67
UWAEL WAEL Standard error	20.66b 33.59a 0.25	13.32a 12.71b 0.56	15.23a 12.87b 0.03	48.80a 41.11b 0.04	26.61a 19.12b 0.28	(%) 68.17 74	(Mcal/kg DM) 2.46 2.67

# Table 3. Parameters related to nutritional quality. Tabla 3. Parámetros relacionados con la calidad nutricional.

Dry Matter (DM); Crude Protein (CP), Neutral Detergent Fiber (NDF); Acid Detergent Fiber (ADF), Digestibility of Dry Matter (DMD), Metabolizable Energy (ME) of ensiled plant material. Unwilted amaranth ensiled (UWAE); wilted amaranth ensiled (WAE); unwilted amaranth ensiled with added lactic acid bacteria (UWAEL); wilted amaranth ensiled with added lactic acid bacteria (WAEL).

Values of the same variable followed by the same letter are not statistically different according to Fisher's LSD test (p > 0.05).

Materia Seca (MS); Proteína Cruda (PC), Fibra Detergente Neutra (FDN); Fibra Detergente Ácida (FDA), Digestibilidad de la Materia Seca (DMS), Energía Metabolizable (EM)) del material vegetal ensilado. Ensilado de amaranto sin orear (UWAE); ensilado de amaranto oreado (WAE); ensilado de amaranto sin orear con adición de bacterias lácticas (UWAEL); ensilado de amaranto oreado con adición de bacterias lácticas (WAEL).

Los valores de las mismas variables seguidos de la misma letra no son estadísticamente diferentes según la prueba LSD de Fisher (p > 0,05).

High protein content is generally associated with higher forage quality, and protein content has been reported to vary from 11.5 to 14% in amaranth silage (16). These variations could be associated with on-site environmental conditions and agronomic practices affecting nutritional composition (39). Table 3 (page 122) shows protein content of the different silages used in this study, with values within the mentioned range. Thus, UWAE had the lowest value (11.14%) for this variable, whereas WAE had the highest value (13.43%). A decrease in protein content of the ensiled material was observed when compared with the non-silage forage (WA-C and UWA-C). The greatest decrease in protein content was observed in UWAE (26% and 12% in UWAE and UWAEL, respectively); however, these decreases were smaller in previously wilted forage (7% WAE and 10% WAEL). In general, a higher osmotic pressure results from higher dry matter (DM) content, which affects plant enzymatic activity and reduces proteolytic capacity (16, 30). This leads to a decrease in soluble protein (CP) and an increase in the insoluble protein fraction, although the latter is still potentially degradable in wilted silages (34, 35).

In contrast, the addition of LAB to unwilted forage decreased proteolysis by 54% and CP loss decreased. However, in the case of WAEL, the addition of LAB did not decrease proteolysis, indicating an interaction between wilting and the addition of LAB, reflected in a decrease in CP. These results agree with those published by Abbasi *et al.* (2018), who observed decreases in protein content in relation to fresh material of 9%, 10%, and 7.5% in ensiled forage, silage with lactic acid bacteria, and wilted forage, respectively.

Ash contents of the different experimental microsilages is listed in table 3 (page 122). The silages with BL (UWAEL and WAEL) had the highest and lowest ash values, respectively. However, silages without the addition of lactic acid bacteria (UWAE-WAE) did not show statistically significant differences (p>0.05), and the wilting process did not affect ash content. Amaranth species are characterized by high mineral content that can vary according to variety, harvest time, climate, soil, and crop management (39). The ash values obtained were within the range of those published by other authors for amaranth silage, which provides more minerals for livestock (25, 28). Seguín *et al.* (2013) reported that the major mineral in amaranth is Ca, generally found as calcium carbonate, and a fermentation process can favor the forage buffering capacity to the detriment of its acidification.

Fibers in a feed consist of structural carbohydrates in the cell walls and soluble nonstructural carbohydrates. Concerning fiber content of fresh chopped or forages (UWA-C and WA-C), the ensiling process reduced the NDF and ADF values, in agreement with previous reports (16, 20, 25). Fiber values in the silages obtained (table 3, page 122) were within the ranges described in the literature for this crop (NDF: 28-47.7% and ADF: 26-31%) (14, 21). ADF and NDF showed statistically significant differences due to the effect of wilting, and a decrease was observed as a consequence. However, the addition of LAB further favored this decrease in ADF and NDF. Similar results have been reported for amaranth under similar conditions (wilting and addition of LAB) (1). However, in our study on unwilted silages, no such decrease was observed. The higher moisture content and lower temperature of the experimental microsilage in UWAE and UWAEL may have limited the fermentation process and decreased the action of cellulolytic bacteria responsible for the reduction in the fibrous fraction (14, 38). Fiber content is directly related to digestibility and influences the rate at which feed passes through the digestive tract of an animal (18). Digestibility values for UWAE and UWAEL mirrored those observed for FDN and FAD, and the addition of LAB did not increase this parameter. However, WAE and WAEL showed higher digestibility and metabolizable energy values (table 3, page 122). A synergistic effect was observed between wilting and the addition of LAB, achieving silage with a digestibility of 74.02%. This silage would be classified as good quality according to Di Marco's classification (7), whereas the rest of the silages would be of medium quality. Regarding the ME of WAEL (2.67 Mcal/k DM), an 8% increase was observed with respect to the non-silage plant material (WA-C). This energy value can be expressed as 11.17 MJ/k DM and is within the range determined for silages of other amaranth species (1, 29). Based on the results obtained in this study, we can conclude that WAEL presented the best organoleptic, conservation, and nutritional characteristics among the four silages evaluated. By comparing the quality of this silage with the values described for crops traditionally used in this type of conservation technique (table 4, page 124), we can infer that it is comparable to the quality of corn silage.

# Table 4. Comparison of nutritional quality parameters of the main crops used as fodder.Tabla 4. Comparación de los parámetros de calidad nutricional de los principales cultivosutilizados como forraje.

	DM (%)	рН	CP (%)	Ash (%)	ADF (%)	NDF (%)	DMD (%)	ME	Reference
Corn	36	3.5	8.5	5.5	15.8	36.6	75.5	2.7	(21)
Sorghum	33.7	4	6.9	8.7	30.7	55	66.3	2.4	(6)
Amaranth	33.5	4.6	12.7	12.9	19.1	41.1	74	2.7	This work

Dry Matter (DM); pH; Crude Protein (CP), Ashes; Acid Detergent Fiber (ADF), Neutral Detergent Fiber (NDF); Digestibility of Dry Matter (DMD), Metabolizable Energy (ME).

Materia seca (MS); pH; Proteína Cruda (PC), Cenizas; Fibra Detergente Ácida (FDA);

Fibra Detergente Neutra (FDN); Digestibilidad de la Materia Seca (DMS), Energía Metabolizable (EM).

Likewise, it is evident that amaranth protein content did not affect the ensiling process, and its digestibility was comparable to that of corn silage. Regarding corn biomass production for silage, previously reported values in the Lower Rio Negro Valley vary between 16-34 Tn DM/ha, depending on the hybrid and management (19). In this sense, amaranth yields (7.8-21 Tn DM/ha) were comparable to those of corn (39).

#### CONCLUSION

Wilting and inoculation of amaranth forage with lactic acid bacteria before ensiling resulted in silage with nutritional characteristics (crude protein percentage, fiber content, dry matter digestibility, and metabolizable energy) that can be classified as a high-quality component of animal diets. The practice of ensiling amaranth as an alternative for conserving forage can be considered viable in semi-arid regions such as Patagonia. However, further research under *in vivo* conditions is required, especially regarding animal responses according to category and species, as well as the exploration of combinations with other ingredients to achieve complementarity and a better balance of nutrients and energy.

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# Morphostructural composition and meat quality in local goat kids from the northeastern region of Mexico

### Composición morfoestructural y calidad de la carne en cabritos locales de la región noreste de México

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#### ABSTRACT

Goat farming is an important activity in northern Mexico. In this sense, "cabrito" or kid goat is a typical regional dish with high economic and cultural value. However, information on the morphostructural composition and meat quality of these local specimens is scarce. Given this, the objective was to evaluate morphostructural characteristics, carcass and meat quality in local kids according to sex in the northeastern region of Mexico. For this purpose, 14 kids (7 males and 7 females) 57 days old were slaughtered. Morphostructural composition was evaluated with 22 zoomometric and phenotypic variables. Carcass characteristics were evaluated by considering different body structures, carcass yield and degree of fatness. Meat quality was determined by physicochemical characteristics, nutritional value and fatty acid profile. The sex effect was evaluated by t-test of independent means and Chi-square. Meat physicochemical characteristics, nutritional value and morphostructure of local kids were heterogeneous and showed no differences ( $P \ge 0.05$ ) concerning sex. Carcass, kidneys, head, neck, rib and loin weights were higher in males than in females ( $P \le 0.05$ ). Fatty acids (FA) found in greater proportion were palmitic (C16:0), oleic (C18:1, n-9), stearic (C18:0), and myristic (C14:0). These FA comprised 80.85 % of the lipid profile of male meat and 76.83% of females. These results are the basis for future programs aimed to improve production systems. Differences found could shed light on future efforts on how to differentiate goat meat from this region of Mexico and enter new markets directly benefiting small producers.

#### Keywords

meat • carcass • nutritional quality • fatty acid profile

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#### RESUMEN

La caprinocultura es una actividad muy importante en el norte de México por la producción de leche y carne. En este sentido, el cabrito es un platillo típico de esta región con valor económico y cultural elevado, no obstante, la información que describe la composición morfoestructural y calidad de carne de estos ejemplares locales, ha sido poco documentada. Debido a lo anterior, el objetivo fue evaluar las características morfoestructurales, calidad de canal y carne en cabritos locales de acuerdo con el sexo (machos y hembras) en la región noreste de México, para esto se sacrificaron 14 cabritos (7 machos y 7 hembras) de 57 días de edad. Para evaluar la composición morfoestructural, se consideraron 22 variables zoométricas y fenotípicas. Las características de la canal se evaluaron considerando las distintas estructuras corporales, rendimiento en canal y grado de engrasamiento. Se determinó calidad de carne midiendo las características fisicoquímicas, valor nutricional y perfil de ácidos grasos. Se evaluó el efecto del sexo mediante una prueba de t de medias independientes y Chi cuadrada. La estructura morfoestructural de los cabritos locales es heterogénea, y no mostraron diferencias (P≥0,05) respecto del sexo. El peso de la canal, riñones, cabeza, cuello, costillar y lomo fueron mayores en machos que en hembras (P≤0,05). Las características fisicoquímicas y valor nutricional de la carne no mostraron diferencias entre sexos. Los ácidos grasos (AG) que se encontraron en mayor proporción fueron; palmítico (C16:0), oleico (C18:1, n-9), esteárico (C18:0), y mirístico (C14:0). Estos AG comprendieron el 80,85 % del perfil lipídico de la carne de los machos, mientras que en hembras representaron el 76,83 %. Estos resultados son la base para futuros programas de mejora del sistema productivo y donde las diferencias encontradas podrían arrojar luz sobre esfuerzos futuros, sobre cómo diferenciar la carne de cabrito en esta región de México e ingresar a nuevos mercados que beneficiarían directamente a los pequeños productores.

#### **Palabras clave**

carne • canal • calidad nutricional • perfil de ácidos grasos

#### INTRODUCTION

In Mexico, goat production is focused on meat and milk. During 2022, carcass meat production achieved 77,000 tons, and 160 million liters of milk (32). The predominant production system is extensive, with animals known as "local", with an undefined phenotype derived from mating different breeds such as Alpina, Saanen, Nubia, and Toggenburg (31). Besides feeding suckling kids, the produced milk is used for cheese, cajeta (a milk candy-type), and typical regional candies. Kids are slaughtered and consumed at approximately 30 days old. Their meat is soft and tender, low-fat, pearly white and juicy. As a traditional Mexican dish, particularly in the north of Mexico, it is consumed on special events and can be cooked in various presentations like "al pastor" (shawarma), fried, or roasted. It is also considered a gourmet dish reaching high prices in restaurants (32).

Similar dishes are prepared in other countries such as Spain, where this meat should meet certain characteristics (9), India, China, Pakistan, Nigeria, Bangladesh, and Iran, where goat production is important (13). However, in Mexico, particularly in the northeast, information on body structure, carcass quality, and nutritional properties of kid meat is scarce. Nevertheless, breeds such as Payoya, Gokcead, Maltese, Majorera, Blanca celtiberica, Negra serrana and Moncaica, have been extensively documented (6).

In this context, generating local information on kid meat produced in this region of Mexico is important given that it supplies almost all the consumed kid meat in the northern states of Mexico. Regional goat raising relies on the environmentally best-adapted breeds. In this sense, particular production strategies can set particular qualities, where denominations of origin can trigger added value (26). The aforementioned is particularly important for small-scale, and highly social and economically marginalized producers (31, 35). Therefore, our objective was to evaluate morphostructural traits, carcass and meat quality, and fatty acid profile considering sex in local kids from the northeastern region of Mexico.

#### MATERIAL AND METHODS

Animal management and study protocol was approved by the Bioethics Committee of the Facultad de Medicina Veterinaria y Zootecnia – Universidad Autónoma de Tamaulipas in the pronouncement CBBA\_01\_2023.

#### **Place of study**

This experiment was conducted in a commercial production unit, located in ejido Ignacio Zaragoza, Viesca, Coahuila, Mexico, within the region known as Comarca Lagunera. The climate is desert, semi-warm with cool winters (BWhw), mean annual rainfall of 240 mm, with average temperature of 25°C, ranging from -1°C in winter to 44°C in summer.

#### Animals and feeding

Before the experiment, during the summer, a herd of 150 local empty goats mated naturally during grazing and in housing pens.

Fourteen 57-day-old kids (7 males and 7 females) weighing 7.7 kg (live weight; LW) were housed with their mothers from birth in individual 2 x 3 m pens provided with shade, drinkers and *ad libitum* mineral salts.

Goat feeding was based on grazing from 8:00 a.m. to 1:00 p.m., and from 4:00 p.m. to 8:00 p.m., keeping the goats penned during the hottest hours of the day, and taking advantage of this space of time for kid suckling. Table 1 shows plant nutritional value during grazing.

**Table 1**. Average chemical composition of the main plant species consumed by local goatsin northeastern Mexico.

Gradian	Nutritional Content (Dry basis; g kg DM)								
species	DM	СР	ADF	NDF	ME	NEI			
Cynodon dactilon	95.3	5.9	36.9	61.4	1.7	1.5			
Amaranthus palmeri	93.8	14.4	25.9	33.7	2.1	1.3			
Setaria macrostachia	94.0	13.2	37.4	61.4	1.7	1.1			
Solanum eleagnifolium	93.7	22.1	32.6	37.7	2.5	1.3			
Spharalcea angustifolia	94.1	11.9	39.1	51.4	2.0	1.2			
Enneapogon desvauxii	94.8	5.3	44.3	71.0	1.5	0.8			
Cucumis melo (Vegetative part)	90.7	13.8	33.8	34.1	2.2	1.4			
Cucumis melo (Fruit)	92.9	10.5	30.9	35.7	2.2	1.1			

**Tabla 1.** Composición química promedio de las principales especies de plantasconsumidas por caprinos locales en el noreste de México.

#### Transportation and slaughtering of goat kids

Kids were slaughtered at weaning (57 days of age). Twelve hours before slaughter, they were separated from their mothers and transported to the Municipal slaughterhouse in Matamoros, Coahuila, where they were slaughtered following the Official Mexican Standard NOM-033-SAG/Z00-2014.

#### Zoometric and morphostructural measurements

Before slaughter, zoometric traits were measured: live weight (LW), face width (FW), skull length (SL), ear length (EL), ear width (EW), neck width (NW), neck length (NL), height at withers (HW), chest circumference (CC), barrel circumference (BC), flank depth (FD), lumbosacral height (ASL), leg length (LL), cane perimeter (CP). Morphostructural traits recorded included skin pigmentation (SP), hoof pigmentation (HP), mucous membrane pigmentation (MP), presence of wattles, beard, and horns (1=present, 2=absent) (8, 12, 14).

DM= dry matter; CP= crude protein; ADF= acid detergent fiber; NDF= neutral detergent fiber; ME= metabolizable energy; NEl= net energy for lactation. DM= materia seca; CP= proteína cruda; ADF= fibra detergente ácido: NDF= fibra detergente neutro; ME= energía metabolizable: NEl= energía neta para la lactancia.

#### **Carcass yield**

Before slaughter, the PV was recorded. Subsequently, carcass productive components were sectioned and removed (head without skin, skin, legs, lungs and trachea, liver, heart, rumen, intestine and testicles -in males-). Yield was calculated by dividing cold carcass weight (24 hours postmortem at 4°C) by the initial live weight, expressed as a percentage (4).

#### Nutritional value

From each kid, 200 g of *Longissimus dorsi* muscle meat were grounded to homogenize and determine protein, fat, collagen, and moisture content with a FoodScan<sup>™</sup> Meat Analyzer.

#### Physicochemical characteristics of meat

To measure meat physicochemical characteristics, the *Longissimus Dorsi* muscle was removed with a transverse cut between the 12<sup>th</sup> and 14<sup>th</sup> rib, 24 hours *postmortem* (4).

#### pН

The pH was measured 24 h *post mortem* from a cut of the *Longissimus dorsi* muscle at the 12<sup>th</sup> rib, inserting the electrode of a portable potentiometer (HANNA® instruments, HI99163, Singapore), previously calibrated with pH 4.00 and 7.00 buffer solutions.

#### Color

Color was measured at three different points on the surface of the *Longissimus dorsi* muscle 24 h *post mortem*, using the Hunter method. A colorimeter (Minolta, Mod CR-400/410, Tokyo, Japan) determined L\* (lightness), a\* (red-green) and b\* (yellow-blue) (11).

#### Drip loss

Drip loss was determined after Wang *et al.* (2016) with modifications. Approximately 30 g of the *Longissimus dorsi* muscle meat sample were weighed and placed in Styrofoam cups hanging from a thread without touching the walls of the cup. Subsequently, they were stored at 4°C and weighed 24 h later. Drip loss was expressed as the percentage of weight loss to initial weight (8).

#### Water retention capacity (WRC)

The WRC was analyzed following Guerrero *et al.* (2002), with modifications. Five g of finely minced meat of the *Longissimus dorsi* muscle, 24 h *postmortem*, were weighed and homogenized with 8 ml of sodium chloride for 1 minute using a glass rod. Subsequently, it was left to rest for 30 minutes in an ice bath. The extract was centrifuged for 25 minutes at 35,000 r.p.m. The supernatant was drained and the volume was measured in a graduated cylinder. The amount of ml of solution retained in 100 g of meat was reported.

#### **Cooking yield**

Cooking yield was determined after Liu *et al.* (2012) with modifications. From each meat sample, 50 g of the *Longissimus dorsi* muscle, 24 h *postmortem* were weighed using an analytical scale and placed in Ziploc-type bags. They were then placed in a water bath at 90°C for 15 minutes. Meat internal temperature was measured with a stem thermometer. Subsequently, they were left to rest at room temperature for 30 minutes. After this time, they were re-weighed. Cooking yield was obtained by considering initial *vs.* final weight differences, expressed as percentages.

#### Fatty acid (FA) profile

Fat purification was carried out through FA methylation. We proceeded to oven-dehydrate 30 g of the *Longissimus dorsi* muscle at 60°C. Subsequently, meat samples were purified (15) and methylated according to Jenkins (2010) modified by Granados-Rivera *et al.* (2017). Once the FA methyl esters were obtained, they were determined in a Hewlett Packard 6890 chromatograph with an automatic injector equipped with a silica capillary column (100 m x 0.25 mm x 0.20  $\mu$ m thickness, Sp-2560, Supelco). FA identification was done by comparing retention times of each peak obtained from the chromatogram against a standard of 37 FA methyl ester components, and a specific standard for cis-9, trans-11 and trans-10, cis-12 isomers (Nu-Check).

#### Statistical analysis

Significant differences among quantitative zoometric traits, nutritional value, carcass and meat quality, and fatty acid profile between male and female goat kids were determined by a t-student test for independent means with the SAS version 9.3 program. Given morphostructural variables are frequencies, a Chi-square ( $\chi^2$ ) test was used to assess independence concerning sex.

#### RESULTS

#### Zoometric, morphostructural, and carcass measurements

Morphostructure of local kids was heterogeneous, and no differences ( $P \ge 0.05$ ) were found between sexes (table 2).

# **Table 2.** Absolute (AF) and relative (RF) frequencies for morphostructural traits in localgoat kids.

**Tabla 2.** Frecuencias absolutas (FA) y relativas (FR) para las característicasmorfoestructurales en cabritos locales.

Skin pigmentation		Male	Female	Mucous membrane pigmentation		Male	Female
Drecont	AF (n)	3	3	Drecont	AF (n)	7	6
Present	RF (%)	21.43	21.43	Present	RF (%)	50.0	42.86
	AF (n)	4	4		AF (n)	0	1
Absent	RF (%)	28.57	28.57	Absent	RF (%)	0	7.14
		χ <sup>2</sup>	1.0000			χ <sup>2</sup>	0.2994
Hoof pigmentation		Male	Female	Presence of wattles		Male	Female
Dresset	AF (n)	5	4	Dresset	AF (n)	1	2
Present	RF (%)	28.57	35.71	Present	RF (%)	7.14	14.29
	AF (n)	2	3	Absent	AF (n)	6	5
Absent	RF (%)	21.43	14.29		RF (%)	42.86	35.71
		χ <sup>2</sup>	0.5770			χ <sup>2</sup>	0.5148
Presence of horns		Male	Female	Presence of beard		Male	Female
Dresset	AF (n)	4	5	Dresset	AF (n)	0	0
Present	RF (%)	28.57	35.71	Present	RF (%)	0	0
	AF (n)	3	2		AF (n)	7	7
Absent	RF (%)	21.43	14.29	Absent	RF (%)	50.0	50.0
		χ <sup>2</sup>	0.5770				

Concerning live weight, males were significantly heavier compared to females at 57 days old, with an average difference of 0.940 kg. Regarding other body traits, no differences (P $\ge$ 0.05) were found between sexes (table 3, page 132).

Variable (cm)	Male	Female	SEM	P value
Live weight (kg)	8.23	7.29	0.3298	0.0001
Face width	6.07	5.71	0.3141	0.2842
Skull length	8.57	8.21	0.5487	0.5287
Ear length	9.42	10.07	0.1279	0.5836
Ear width	4.85	4.50	0.3891	0.3798
Neck width	17.35	17.28	1.1628	0.9522
Neck length	14.58	14.78	0.7214	0.9423
Height at withers	34.61	34.57	1.3087	0.9522
Chest circumference	32.08	30.37	0.855	0.0688
Barrel circumference	32.92	31.21	1.4487	0.2634
Flank depth	11.35	11.78	0.5933	0.4842
Lumbosacral height	35.51	34.92	1.3863	0.9600
Leg length	14.42	14.07	0.4862	0.4809
Calf perimeter	6.14	6.07	0.7641	0.7787

Table 3. Live weight and zoometric measurements of local goat kids
Table 3. Peso vivo y medidas zoométricas de cabritos locales.

SEM: Mean standard error. SEM: error estándar de la media.

Regarding yield components and carcass traits, weights of cold carcass, kidneys, head, neck, ribs, and loin were higher in males than females ( $P \le 0.05$ ), showing significant differences between sexes. Other carcass components did not show differences between sexes (table 4).

Variable (kg)	Male	Female	Female SEM		
Head	0.52	0.47	0.0192	0.0313	
Skin	1.02	0.85	0.0905	0.0872	
Blood	0.25	0.26	0.0497	0.7788	
Hooves	0.33	0.30	0.0233	0.1674	
Rumen	0.18	0.15	0.0212	0.1644	
Intestine	0.27	0.23	0.0358	0.2539	
Heart	0.04	0.04	0.0149	0.2577	
Kidneys	0.05	0.03	0.0142	0.0285	
Liver	0.16	0.14	0.0126	0.1389	
Cold carcass weight	5.20	4.22	1.0222	0.0121	
Omental fat	0.05	0.05	0.0136	0.6091	
Mesenteric fat	0.06	0.05	0.0126	0.6095	
Perinephric fat	0.11	0.07	0.0322	0.2075	
Ribs	1.49	1.24	0.1025	0.0296	
Legs	0.84	0.75	0.0911	0.3072	
Loin	0.55	0.42	0.1013	0.0024	

**Table 4**. Yield components of the carcass in local goat kids.**Tabla 4.** Componentes de rendimiento de la canal en cabritos locales.

SEM: Mean standard error. SEM: error estándar de la media.

#### Nutritional value and meat physicochemical characteristics

Meat nutritional value of male and female kids showed no differences ( $P \ge 0.05$ ) (table 5).

Variable (%)	Male	Female	SEM	P value
Dry matter	24.34	25.51	0.6118	0.0923
Moisture	75.65	74.48	0.6118	0.0923
Protein	21.58	22.23	0.3177	0.0648
Fat	2.49	2.18	0.2882	0.1948
Collagen	1.25	1.17	0.0983	0.4597
pH	5.99	6.15	0.2733	0.5627
Drip loss %	2.28	3.66	0.9086	0.0689
Cooking yield %	69.65	65.51	3.8076	0.2987
Water retention capacity (mL/100 g)	38.02	26.34	6.9588	0.1224
L*	48.61	48.17	1.5912	0.7835
a*	17.46	16.99	0.9604	0.6324
b*	10.06	10.18	0.8332	0.8919

**Table 5.** Nutritional value and meat quality of local goat kids.**Tabla 5.** Valor nutricional y calidad de la carne de cabritos locales.

### Fatty acid profile

The FAs found in greater quantity in meat were palmitic (C16:0), oleic (C18:1, n-9), stearic (C18:0), and myristic (C14:0). These, in total, represented an average of 80.85% of the FA that make up male meat and 76.83% of female meat (table 6, page 134).

Meat concentration of caproic, lauric, myristic, and oleic acids showed differences (P $\ge$ 0.05) regarding sex, being higher in male meat.

The concentration of saturated FA was significantly higher in males compared to females with values of 56.45% and 46.72%, respectively, while the amount of monounsaturated (40.53%) and polyunsaturated (9.28%) FA was higher females compared to males (P<0.05).

#### DISCUSSION

#### Zoometric, morphostructural and carcass measurements

Morphostructural characteristics were heterogeneous, without defined traits in terms of sex. In this regard, Maldonado-Jáquez *et al.* (2023) report that, in local kids from northern Mexico, the dominant phenotype corresponds to animals without wattles or beards. This coincides with our study since no animal presented wattles (total frequency of 78.57%), or beard. Furthermore, these same authors mention that local kids present pigmented mucous membranes and horns. In this study, 85.71% of the animals presented pigmented mucous membranes and 94.28% presented horns, reaffirming this information. These results can be attributed to local animals of this region being a cross between different breeds, with varying phenotypic traits.

On the other hand, sex had a significant effect on animal weight, where males were heavier than females. This same result is reported by Maldonado-Jáquez *et al.* (2023) for 30-day-old kids, where males weighed an average of 800 g more than females. This effect is explained by goat growth curves (2), showing shorter growth phases in males (ending up to 4 months before) than females. Moreover, the growth hormone has a marked effect on the early development of males, when the highest growth rates are observed between 20 and 60 days old (27, 29).

Regarding body measurements, our results differ from the reported by Maldonado-Jáquez *et al.* (2023), who indicated differences in neck length and width, body length, chest circumference, and leg length, probably given by age differences. While

L\*: lightness index; a\* red to green index; b\*: yellow to blue index; SEM: Mean standard error. L\*: índice de luminosidad; a\*índice de rojo a verde; b\*: índice de amarillo a azul; SEM: error estándar de la media. Maldonado-Jáquez *et al.* (2023) considered 1 to 30 days, our research measured at 57 days old. Age significantly influences live weight and body conformation (3).

Cold carcass was heavier in males than in females, as found by Todaro *et al.* (2004) who reported differences in carcass weight with respect to sex with values of 5.7 kg and 5.3 in males and females of the Nebrodi breed, respectively, at 47 days old. However, Bonvillani *et al.* (2010) indicated no differences concerning sex, with average weights of 5.34 and 5.48 kg for females and males respectively in local kids from Córdoba, Argentina, at 60 to 90 days old. This could be due to age heterogeneity rather than sex. However, the breed effect could also influence carcass weight of males and females.

Differences in rib and loin weights between males and females can be attributed to males having a higher live weight at 56 days old, and consequently, a higher carcass weight reflected in a higher weight in these structures. This sex difference may assist decisions considering males being directed to the sale of cuts (30), for wholesale sale and/or in restaurants, reaching high prices. This relies on the fact that a single piece is equivalent to 50% of the price paid to the producer for a whole live goat (5). Females not meeting breeding characteristics can be commercialized as meat.

Fatty acid	Common name	Male	Female	SEM	P-value
C4:0	Butyric	0.761	1.309	0.4719	0.2675
C6:0	Caproic	0.511	0.187	0.0588	0.0005
C12:0	Lauric	1.632	0.842	0.1539	0.0003
C14:0	Myristic	9.841	6.920	0.9025	0.0071
C14:1	Myristoleic	0.339	0.511	0.0848	0.0658
C15:0	Pentadecanoic	0.826	0.570	0.1414	0.0958
C16:0	Palmitic	30.55	25.690	2.7086	0.0990
C16:1 cis-9	Palmitoleic	2.797	3.333	0.6165	0.4048
C17:0	Heptadecanoic	1.198	0.956	0.1728	0.1864
C17:1cis-10	Cis-10-Heptadecenoic	0.797	1.222	0.1971	0.0519
C18:0	Stearic	10.686	9.663	1.6219	0.5400
C18:1 cis-9	Elaidic	0.527	0.849	0.2435	0.2184
C18:1, n-9	Oleic	29.780	34.56	1.1955	0.0017
C18:2	Linoleic	3.697	5.353	1.1752	0.1843
C18:3 n-9,12,15	Linolenic	0.525	0.756	0.1613	0.1772
C18:2 cis-9, trans-11	CLA	0.283	0.305	0.0299	0.4821
C20:1 cis-9	Gondoic	0.048	0.044	0.0046	0.3771
C20:3 n-6	Dihomo-γ-linolenic acid	0.1130	0.149	0.0438	0.4172
C20:3 n-3	Cis 11,14,17 Eicosatrienoic	1.663	2.49	0.7037	0.2692
C22:0	Behenic acid	0.431	0.584	0.2197	0.5001
C22:6 n-3	Docosahexaenoic acid (DHA)	0.174	0.225	0.0653	0.4503
SFA		56.45	46.72	1.7464	<.0001
MUFA		34.29	40.53	1.1829	<.0001
PUFA		6.45	9.28	1.1008	<.0001
Undetermined		2.80	3.45		

**Table 6.** Fatty acid profile (g/100 g<sup>-1</sup> of fat) in local goat kids' meat. **Tabla 6**. Perfil de ácidos grasos (g/100 g<sup>-1</sup> de grasa) en carne de cabritos locales.

<sup>ab</sup> Different letters in the same row show statistical differences  $(P \le 0.05); SEM:$ Mean standard error; SFA= saturated fatty acids, MUFA= monounsaturated fatty acids; PUFA= polyunsaturated fatty acids. <sup>ab</sup> Letras diferentes en la misma fila presentan diferencias estadísticas ( $P \le 0,05$ ); SEM: error estándar de la media; SFA= ácidos grasos saturados, MUFA= ácidos grasos monoinsaturados; PUFA= ácidos grasos poliinsaturados.

#### Nutritional value of meat

Sex does not influence meat nutritional value. Other authors concluded that in Nebrodi breed kids, sex did not change the protein, fat, and ash contents of meat (34). Protein content of local kid meat from northeastern Mexico is higher than the 20.79% and 19.72%

reported by Horcada *et al.* (2012), as well as the 2.37% fat reported by Kawęcka *et al.* (2022). Fat percentage in kid meat is low compared to other species, probably because of age, since fat formed in early stages is mesenteric, while intramuscular fat is formed during adulthood (20).

#### Physicochemical characteristics of meat

Sex does not influence meat pH at 24 hours *post mortem*. Values found are close to those reported in Payoya kids at 30 days old (19). However, others report higher pH (23). Regardless, our values are over the recommendations for normal meat considering species for meat production (1). This effect could be due to kids being only fed with milk and muscle glycogen before slaughter is not abundant given goat restless behavior (20) rather than chronic stress before slaughter, a condition that has been documented to cause high pH values in meat (1).

Drip loss, water retention capacity, cooking yield, and color were not modified by sex, as already observed (33, 34). This may be explained by absent differences in pH since low or high final pH will determine the amount of water lost during handling, as well as pale or dark colors. The lightness index and yellow index values obtained in this study are within the reported ranks (10, 19, 34). Conversely, the values obtained for the a\* parameter are higher than other reports (33, 34), probably given by factors like breed and age at the time of slaughter.

#### Fatty acid profile

Fatty acids found in greater quantity are within the reported ranges. Regardless of breed or feeding systems, FAs are palmitic acid (C16:0) with minimum values from 17.32 g/100 g (34) to 25.0 g/100 g (20); stearic acid (C18:0) with values from 7.87 g/100 g (21) to 19.71 g/100 g (7); and oleic acid (C18:1, n-9) with values from 25.38 g/100 g (34) to 51.08 g/100 g (21). The fact that these fatty acids predominate in meat of young and adult animals can be explained by ruminant animals with endogenous synthesis in the adipocyte from acetate, obtained during ruminal fermentation. This determines palmitic acid as the main final product, later elongated into stearic acid or desaturated to oleic acid. Long-chain fatty acids are easily synthesized in adipose tissue (17, 28).

While other studies found no differences regarding fatty acid content concerning sex in Nebrodi and Criollo Cordobes breeds (10, 34), our study found the opposite effect with a greater amount of saturated fatty acids; caproic, lauric, and myristic in male meat. While females have a higher amount of oleic unsaturated fatty acid due to the above, males showed a higher amount of saturated fatty acids and a lower amount of monounsaturated and polyunsaturated fatty acids. These differences could be due to breed or diet, since, even though kids are milk-fed, the lipid profile of the mother's milk will be largely determined by her diet (25).

#### **CONCLUSIONS**

Based on the results, we conclude that kid morphometric characteristics are heterogeneous in females and males in northeastern Mexico. Sex did not affect carcass characteristics, nutritional value, and physicochemical traits of meat. However, sex tended to modify FA profile, favoring a higher concentration of caproic (C6:0), lauric (C12:0), myristic (C14:0), and oleic (C18:1, n-9) acids in males.

This constitutes a pioneer study on morphostructure, carcass, and meat quality characterization of local goat kids from northeastern Mexico and will lay the foundations for future programs to improve the production system. The differences found could shed light on future efforts to differentiate kid meat from this region of Mexico oriented to new markets that would directly benefit small producers.

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# Effect of three different fixed-time AI protocols on follicular dynamics and pregnancy rates in suckled, dual-purpose cows in the Ecuadorian Amazon

## Efecto de tres protocolos diferentes de IA a tiempo fijo sobre la dinámica folicular y las tasas de preñez en vacas de doble propósito amamantando en la Amazonía Ecuatoriana

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#### ABSTRACT

Reproductive performance is crucial for profitability of dual-purpose cow-calf production in the Ecuadorian Amazon. To evaluate three different fixed-time artificial insemination (FTAI) protocols in suckled, dual-purpose cows in the Ecuadorian Amazon. Lactating, Brown Swiss cows (n=301) received 2 mg estradiol benzoate (EB) and an intravaginal device containing 0.5 g of progesterone (P4) on Day 0. They were allocated randomly into the following three treatment groups: Cows in the EB group received 500 µg cloprostenol (PGF2 $\alpha$ ), and P4 devices were removed on Day 7, and 1 mg EB was administered on Day 8. Cows in the ECP group were treated as those in the EB group, except that they received 0.5 mg estradiol cypionate (ECP) at the time of P4 device removal on Day 7 instead of EB on Day 8. Cows in the J-Synch group had the P4 device removed and PGF2 $\alpha$  administered on Day 6. All cows were FTAI on Day 9; cows in the J-Synch group also received 100 µg gonadorelin at that time. Although the diameter of the dominant follicle and the resulting CL were greater (P<0.05) in cows in the J-Synch group, pregnancies per AI did not differ (P>0.2) among groups (EB: 51.0%, ECP: 53.0% and J-Synch: 59.4%). The three protocols tested were applied successfully in suckled, dual-purpose cows with no differences in pregnancies per AI.

#### Keywords

prolonged proestrus • estradiol/progesterone • P4 device • FTAI • pregnancy per AI • lactating dual-purpose cows • tropical

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#### RESUMEN

El desempeño reproductivo es crucial para la rentabilidad de la producción doble propósito en la Amazonía Ecuatoriana. Evaluar tres protocolos para Inseminación Artificial a tiempo fijo (IATF) en vacas en lactancia de doble propósito. Vacas Pardo Suizo en lactancia (n=301) recibieron 2 mg de benzoato de estradiol (EB) y un dispositivo intravaginal con 0,5 g de progesterona (P4) el Día 0 y fueron asignadas aleatoriamente en tres grupos. Las vacas del grupo EB recibieron 500 µg de cloprostenol (PGF2 $\alpha$ ) y se les retiró el dispositivo el Día 7, y recibieron 1 mg de EB el Día 8. Las vacas del grupo ECP fueron tratadas como las del grupo EB, excepto que recibieron 0,5 mg de cipionato de estradiol (ECP) en el día del retiro del dispositivo (Día 7) en lugar de EB el Día 8. En las vacas del grupo J-Synch, el retiro del dispositivo y la administración de PGF2 $\alpha$  se realizaron el Día 6. Todas las vacas fueron IATF el Día 9 y las del grupo J-Synch recibieron 100 µg de gonadorelina en ese momento. Aunque el diámetro del folículo dominante y del CL fueron mayores (P<0,05) en las vacas del grupo J-Synch, las tasas de preñez a la IATF no difirieron entre los grupos (EB: 51,0%, ECP: 53,0% y J-Synch: 59,4%). Los tres protocolos probados se pueden aplicar con éxito en vacas de doble propósito amamantando.

#### Palabras clave

proestro prolongado • estradiol/progesterona • dispositivo de P4 • IATF • preñez por IA • vacas doble propósito en lactancia • tropical

#### INTRODUCTION

Dual-purpose cattle production (i.e., milk and meat) represents a main source of income for small farms in tropical areas around the world, such as the Ecuadorian Amazon. However, there is a need for improvement in terms of reproductive performance and genetics in dual-purpose livestock (31). Less-than-optimal efficiency is due to multiple factors, including the environment (high temperature and humidity), nutrition, health, management, and poor animal welfare, among others (31). One of the technologies that has had the greatest impact on reproductive performance in beef and dairy cattle has been the systematic application of artificial insemination (AI), without estrus detection, known as fixed-time AI (FTAI) (4). Currently, there is a wide range of FTAI protocols used in beef and dairy cattle (4, 15, 47) which requires testing in dual-purpose cattle in the Ecuadorian Amazon. FTAI protocols are classified according to the main hormones used. Ovsynch-type protocols are based on the use of GnRH and prostaglandin F2 $\alpha$  (PGF2 $\alpha$ ; 35, 47) and may be combined with the use of an intravaginal progesterone (P4) releasing device in beef (26, 29) and dairy cattle (41). Estradiol and P4-based protocols (2, 30) have also been commonly used for FTAI in beef and dairy cattle, especially in South America (3, 4, 48). This treatment has been simplified by the administration of estradiol cypionate (ECP) as an ovulation-inducing agent, given at the time of P4 device removal (14) replacing the use of estradiol benzoate (EB) 24 h after P4 device removal in beef (14, 44) and dairy cattle (40, 42). In general, both FTAI protocols (GnRH-based and estradiol/P4-based) result in pregnancies per AI (P/AI) between 40 to 60% in beef (1, 5) and 30 to 50% in dairy cattle (15, 43). In 2012, a new estradiol/P4-based protocol, called J-Synch, was developed (19). In this treatment protocol, the P4-releasing device is inserted for a shorter period of time (6 days) and the administration of ECP at the time of P4 device removal was replaced by the administration of GnRH at the time of AI (72 h after P4 device removal), promoting a longer proestrous period. Subsequent studies have shown that the J-Synch protocol is an efficient treatment to synchronize ovulation in beef (19, 32) and dairy (36) heifers, resulting in greater pregnancy rates than those achieved with the conventional estradiol/P4-based protocols which used ECP or EB to induce ovulation (5, 20, 36). The higher pregnancy rates obtained with the J-Synch protocol were associated with longer exposure to elevated endogenous estradiol concentrations during the proestrus period, which resulted in a more appropriate environment for early embryo development (20).

The current experiment was designed to evaluate three different estradiol/P4-based FTAI protocols, which differ mainly in the duration of the proestrus period, in suckled dual-purpose cows of the Ecuadorian Amazon. The hypothesis to be tested was that the protocol with the prolonged proestrus and utilizing GnRH to induce ovulation, named J-Synch, would result in greater P/AI than conventional estradiol/P4 protocols, which utilized EB or ECP to induce ovulation.

#### MATERIALS AND METHODS

The study was performed in the Ecuadorian Amazon, at the Center of Postgraduate Research and Conservation of the Amazon (CIPCA) of the Amazon State University in the province of Napo, which is located in the Northeast of Ecuador (Latitude: 01°14'32" South, Longitude: 53°77'13" West), from October 2015 to April 2016. The climate is tropical, with 4000 mm of precipitation per year, average relative humidity of 80% and temperatures ranging between 25 and 30°C. The altitude varies from 580 to 990 m above sea level, and although the soils have a very heterogeneous composition, most originated in fluvial sediments from the Andean plateaus.

#### Animals and feeding

Lactating, Brown Swiss cows (n = 301) with suckling calves were used. The study was performed in three replicates of 100 animals each. Cows were grazing a mixed pasture based on *Brachiaria decumbens* (17.585 kg DM/ha/year), *Brachiaria brizantha* (26.970 kg DM/ha/year), *Arachis pinto* (6.212 kg DM/ha/year), *Desmodium ovalifolium* (5.890 kg DM/ha/year) and *Stylosanthes guianensis* (15.237 kg DM/ha/year). They received the standard vaccination and health protocols applied commonly to cattle at the CIPCA. This included the application of vitamins and minerals, deworming, insecticide dips against ticks and flies, vaccinations for Foot and Mouth Disease, Bovine Rabies and Vesicular Stomatitis.

The cows were multiparous (*i.e.*, 2-5 lactations),  $112.0\pm1.0$  (mean±SEM) days postpartum, with a mean body condition score (BCS) of  $2.4\pm0.3$  (1 to 5 scale; 22), and they were producing  $6.7\pm0.6$  Kg of milk per day. Cows were selected for treatment by the presence of a CL, or at least one follicle >10 mm in diameter in their ovaries through an ultrasound examination and without any abnormalities of the reproductive tract. All cows were handled in the same milking facility and were inseminated with frozen-thawed semen from one bull of proven fertility. Animal procedures were approved and supervised by the Animal Care and Use Committee of the CIPCA and the Institutional Committee for the Care and Use of Laboratory Animals (CICUAL) of the University of Villa Maria (UNVM), Argentina.

#### Treatments

Cows were randomly allocated to one of three FTAI protocols. The EB protocol (n = 100): On Day 0, cows received a P4-releasing intravaginal device (DIB 0.5 g, Zoetis, Ecuador) plus 2 mg EB (Gonadiol, Zoetis) by intramuscular (i.m.) injection. On Day 7, the P4 device was removed, and cows received 500  $\mu$ g cloprostenol (PGF2 $\alpha$ , Ciclase DL, Zoetis) i.m. On Day 8, all cows received 1 mg EB i.m. and the tail heads were painted (Divasa - Farmavic, Spain) for the determination of estrus. On Day 9, (48 to 54 h after P4 device removal) cows were FTAI (figure 1, page 141).

The ECP protocol (n = 100): On Day 0, cows received a P4 intravaginal device plus 2 mg of EB i.m. On Day 7, the P4 device was removed, and cows received PGF2 $\alpha$  and 0.5 mg ECP (Cipiosyn, Zoetis) i.m. and the tail heads were painted as in the EB protocol. On Day 9, (48 to 54 h after P4 device removal), cows were FTAI (figure 1, page 141).

The J-Synch protocol (n = 101): On Day 0, cows received a P4 device plus 2 mg of EB i. m. On Day 6, the P4 device was removed, cows received PGF2 $\alpha$  i.m. and the tail heads were painted as in the previous groups. On Day 9, (72 h after P4 device removal), cows received 100 µg gonadorelin acetate (GnRH, Gonasyn gdr, Zoetis) and were FTAI concurrently (figure 1, page 141).


EB group: Day 0, P4 releasing device (0.5 g of P4) plus 2 mg EB; Day 7 device removal plus 500 µg cloprostenol (PGF); Day 8, 1 mg of EB; Day 9, FTAI (48 to 54 h after P4 device removal). ECP group: Day 0, P4 releasing device (0.5 g of P4) plus 2 mg EB; Day 7 device removal plus 500 µg cloprostenol (PGF) and 0.5 mg ECP; Day 9, FTAI (48 to 54 h after P4 device removal). J-Synch group: Day 0, P4 releasing device (0.5 g of P4) plus 2 mg EB; Day 6 device removal plus 500 µg cloprostenol (PGF); Day 9, GnRH and FTAI (72 h after device removal).
EB: Día 0, dispositivo liberador de P4 (0,5 g de P4) más 2 mg de EB; Día 7, retiro del dispositivo más 500 µg de cloprostenol (PGF); Día 8, 1 mg de EB; Día 9, IATF (48 a 54 h después de retirado el dispositivo de P4). ECP: Día 0, dispositivo liberador de P4 (0,5 g de P4) más 2 mg de EB; Día 7, retiro del dispositivo más 500 µg de cloprostenol (PGF); Día 9, IATF (48 a 54 h después de retirado el dispositivo de P4). J-Synch: Día 0, dispositivo más 500 µg de cloprostenol (PGF) y 0,5 mg de ECP; Día 9, IATF (48 a 54 h después de retirado el dispositivo de P4). J-Synch: Día 0, dispositivo más 500 µg de cloprostenol (PGF) y 0,5 mg de ECP; Día 6, retiro del dispositivo más 500 µg de cloprostenol (PGF); Día 9, GnRH y IATF (72 h después de retirado el dispositivo de P4).

#### Figure 1. Treatment Groups.

Figura 1. Grupos de tratamiento.

# Ultrasonography

All cows were examined by real-time ultrasonography (Ibex Pro and Lyte ®, USA), with a linear probe of 5.0 MHzon Days 7, 9 and 10. A map was made to measure and record all follicles >5 mm present in both ovaries, and to identify the dominant follicle which was defined as the follicle with the largest diameter at the time of device removal. To determine the timing of ovulation (disappearance of the follicle of greatest diameter) animals were examined at the time of FTAI (Day 9) and then every 12 h for the next 24 h. If the cows ovulated after Day 10, the ovulation time was defined as an ovulation occurring > 24 h after FTAI. The ovulation was reconfirmed by the presence of a CL 7 days after FTAI. Two measurements of the CL were taken using the equipment's software which included the vertical and horizontal diameter (width and height) of each CL as described previously (25).

## **Estrus detection**

Estrus was recorded visually and by the disappearance of the tail-paint. The animals with >30% of the paint removed were considered to be in estrus (13). The visual observations 3 times a day: morning, noon and afternoon of Days 7, 8 and 9 were initiated after the removal of the P4 device.

# Blood sampling and progesterone analysis

Seven days after FTAI a blood sample was taken for P4 analysis. After cleaning and disinfecting the base of the tail, samples were taken from the coccygeal vein using 10 mL Vacutainer tubes. Samples were stored at the laboratory at 4°C for 4 to 6 h after extraction

and then centrifuged at 3000 RPM for 20 minutes to separate the serum. The serum was extracted and then stored at -20°C until its subsequent analysis (18). The serum P4 concentrations were determined together, in duplicates, using a progesterone ELISA kit (Progesterone Elisa, DiaMetra S.R.L., Italy). The minimum detectable P4 concentration of the kit was 0.05 ng/mL and the low and high intra-assay coefficients of variation were 4.4% and 19.5%, respectively.

### **Pregnancy diagnosis**

The diagnosis of pregnancy was determined by ultrasonography at 35 to 40 days after FTAI.

## Statistical analysis

For each continuous variable studied, the arithmetic mean (x) and the standard error (SEM) were estimated. The data were analyzed using ANOVA and means were compared by the Tukey-Kramer HSD test. Differences were considered significant with a P value of <0.05. The statistical analysis was carried out using the JMP program (JMP R, 2003) in its version 5.0 for Windows. Estrus, ovulation and P/AI rates were compared among groups using logistic regression for binary data (*i.e.*, 1 = success, 0 = failure) and a logit link, using the Infostat program (21).

### RESULTS

### Estrus expression

Overall, 90.4% (272/301) of the cows showed estrus. The percentage of cows in estrus was lower (P < 0.01) in the J-Synch treatment group (78.2%, 79/101), than in the ECP (93%, 93/100) or EB (100%, 100/100) treatment groups.

# **Follicular characteristics**

The diameters of the dominant follicle at the time of P4 device removal and at the time of FTAI are shown in figure 2. Cows in the J-Synch group had the smallest mean diameter follicle at the time of P4 device removal and the largest diameter follicle at the time of FTAI (P < 0.05).



**Figure 2**. Diameter of dominant follicle (means ± SEM) at the time of P4 device removal (orange columns) and at the time of FTAI (yellow columns) in suckled, dual-purpose cows subjected to three different estradiol/P4-based protocols.

**Figura 2.** Diámetro del folículo dominante (media ± SEM) en el momento del retiro del dispositivo P4 (columnas naranjas) y en la IATF (columnas amarillas) en vacas de doble propósito sometidas a tres protocolos que utilizan estradiol/P4.

<sup>abc</sup> Different superscripts indicate significant differences between groups (P<0.05).</p>
<sup>abc</sup> indica diferencias significativas entre grupos (P<0,05).</p>

### **Time of ovulation**

The mean interval from P4 device removal to ovulation was longer (P < 0.01) in the J-Synch group (87.7 ± 0.6 h) than in the ECP (73.7 ± 0.6 h) or EB (75.7 ± 0.6 h) groups. Overall, the percentage of cows that ovulated by 12 h after FTAI was 41.9% (126/301) and it was greater (P < 0.01) in the J-Synch group than in the other two groups (table 1). Conversely, the percentage of cows that ovulated between 12.1 and 24 h after FTAI was less in the J-Synch group than in the EB and ECP treatment groups (P < 0.01). The percentage of cows that ovulated between 12.1 and 24 h after FTAI was less in the J-Synch group than in the EB and ECP treatment groups (P < 0.01). The percentage of cows that ovulated more than 24 h after P4 device removal was 3% (9/301) and all were in the EB treatment group (9%; P > 0.1).

**Table 1.** Ovulatory response up to 12 h and between 12.1 and 24 h after FTAI in suckled dual-purpose cows subjected to three different estradiol/P4-based protocols.

**Tabla 1.** Respuesta ovulatoria hasta 12 y entre 12.1 y 24 h después de la IATF en vacas de doble propósito sometidas a diferentes protocolos que utilizan estradiol/P4.

Treatments	n	Ovulation up to 12 h after FTAI (%)	Ovulation between 12.1 and 24 h after FTAI (%)	Acumulated ovulation* (%)
J-Synch	101	68 (67.3)ª	31 (30.7) <sup>b</sup>	99 (98.0)
ECP	100	35 (35.0) <sup>b</sup>	65 (65.0) ª	100 (100.0)
EB	100	23 (23.0) <sup>b</sup>	68 (68.0) ª	91 (93.0)
Total	301	126 (41.9)	164 (54.5)	290 (97.0)

Percentages with different superscripts differ (abP <0.01). \*Ovulations occurring within 24 h after FTAI. Los porcentajes con

diferentes superíndices indican diferencias significativas (abP <0,01).

\*Ovulaciones que ocurren dentro de las 24 h posteriores a la IATF.

## CL diameter and serum P4 concentration 7 days after FTAI

The CL diameter was greater (P < 0.05) in cows in the J-Synch group compared to the other two groups, which did not differ from one another (table 2). Similarly, serum P4 concentrations were higher (P < 0.05) in the J-Synch group than in the EB group but were intermediate in the ECP group which did not differ from either of the other treatment groups (table 2).

**Table 2.** CL diameter and serum progesterone concentrations 7 days after FTAI (mean ± SEM) in suckled, dual-purpose cows subjected to three different estradiol/P4-based protocols.

**Tabla 2.** Diámetro del CL y concentración sérica de progesterona a los siete días de la IATF (media ± SEM) en vacas de doble propósito sometidas a diferentes protocolos que utilizan estradiol/P4.

Treatments	n	CL diameter (mm)	Progesterone concentration (ng/ml)
J-Synch	101	$25.4 \pm 0.2^{a}$	$11.4 \pm 0.3^{a}$
ECP	100	24.2 ± 0.2 <sup>b</sup>	$10.6 \pm 0.3^{ab}$
EB	100	24.5 ± 0.2 <sup>b</sup>	9.9 ± 0.3 <sup>b</sup>

Means within columns with different superscritps differ (ab P < 0.05). Las medias con diferentes superíndices indican diferencias significativas (ab P < 0,05).

# Pregnancies per FTAI (P/AI)

Overall, 164/301 (54.4%) of the cows were pregnant following FTAI. P/AI were numerically, but not significantly higher (P<0.2) in cows in the J-Synch group (59.4%, 60/101), compared to the EB (51.0%, 51/100) and ECP (53.0%, 53/100) treatment groups.

Another analysis investigated the effects of the diameter of the dominant follicle at the time of FTAI and the diameter of the ensuing CL on P/AI (table 3). Dominant follicle diameter at the time of FTAI was greater (P < 0.01) in cows that became pregnant than in those that did not become pregnant. Furthermore, the diameter of the CL tended (P=0.0861) to be greater in cows that became pregnant than in those that did not become pregnant to FTAI.

**Table 3.** Comparison of the follicular and luteal characteristics in suckled, dual-purpose cows that became pregnant or did not become pregnant following treatment with three different estradiol/P4-based FTAI protocols (Mean ± SEM).

**Tabla 3.** Comparación de las características foliculares y luteales entre vacas que quedaron preñadas o no, en vacas de doble propósito sometidas a tres protocolos diferentes que utilizan estradiol/P4 (media ± SEM).

Variable	Pregnancy Status*	n	Diameter (mm)	P value	
DE et ETAL	Р	164	13.4 ± 0.2	<0.0001	
Dr at r IAI	0	137	12.3±0.3		
	Р	164	24.9±0.8	0.00(1	
CL / days after FTAI	0	137	24.5±0.9	0.0801	

\*Pregnancy Status: (P) pregnant; (O) non-pregnant. \*Estado de gestación: (P) preñadas; (O) no preñadas.

# DISCUSSION

The environment, genetics, nutrition, health and management have been reported to be determinants of the reproductive performance in cattle. Although the prolonged proestrus protocol (J-Synch) resulted in the largest dominant follicle at the time of FTAI and the largest CL 7 days after FTAI, the hypothesis that this treatment results in greater P/AI than those in the conventional estradiol/P4 based protocols currently used in South America was not supported. Nevertheless, the results of this study have important implications because they demonstrate that other estradiol/P4-based FTAI alternatives can be applied successfully in suckled dual-purpose cattle in tropical environments. The P/AI exceeded 50% in the three treatment groups, which agrees with other studies performed on beef cattle in tropical regions (1, 5, 27, 38), but superior to those reported with dairy cattle in the same region (40, 46).

Another interesting factor related to tropical regions is the low estrus expression rate and a tendency to show estrus during the night, greatly affecting the efficiency of AI programs (4). However, in the present study, the overall estrus expression was 90.4%, which is higher than those reported for beef and dairy cattle in tropical regions (15, 33, 38, 39). Several studies have shown that the estrus expression at the time of FTAI, detected by tail-paint or estrus detection patches is associated with greater P/AI (13, 33, 38, 39, 42, 43) suggesting that tail-paint or patches are useful aids to determine that cows are indeed showing estrus at the time of FTAI, which is likely to result in improved pregnancy outcomes following FTAI.

The duration of the proestrus period tended to have an influence on P/AI in this study and has been reported to be positively related to an improved uterine environment and fertility (7, 8, 9, 11). Furthermore, the difference in dominant follicle diameter at P4 removal and at the time of FTAI in this study agrees with a similar study reported by de la Mata *et al.* (2018), where cows receiving the J-Synch treatment had the smallest follicles at P4 device removal, but had follicles of the greatest diameter at the time of FTAI compared with two other conventional (short proestrus) protocols. Although P/AI was only numerically greater in cows in the J-Synch group, the cows that became pregnant, regardless of treatment groups, had a greater follicle diameter at the time of FTAI than those cows that did not become pregnant. This is also consistent with the results reported by Yañez *et al.* (2016) on dairy cows in Ecuador. It has also been reported that *Bos taurus* beef heifers have the maximum probability of pregnancy when the follicular diameter at the time of FTAI is >12.8 mm (34). Furthermore, *Bos indicus* cows with follicles > 15 mm in diameter had the greatest probability of ovulating and becoming pregnant in estradiol/P4 based FTAI programs (38, 39). Thus, efforts to increase ovulatory follicle size would appear to be a worthy endeavor.

When the time of ovulation was evaluated in relation to the time of FTAI, the percentage of cows ovulating up to 12 h after FTAI in J-Synch treatment group was the highest (67.3%) compared to the ECP (35.0%) or EB (23.0%) treatment groups; while the percentage of cows that ovulated between 12.1 and 24 h after FTAI was 30.7% for the J-Synch treatment, and 65.0% and 68.0% for the ECP and EB treatment groups, respectively. The determination of the percentage of cows ovulating 12, 24 and after 24 h from FTAI may have important implications for fertility to FTAI programs. It has been estimated that the maximum sperm viability in the female genital tract is 24 to 30 h (23) and after ovulation occurs, the oocyte has an even shorter fertile lifespan (6). Therefore, achieving greater synchrony between ovulation and sperm arrival in the oviduct is very likely to increase P/AI (37). Although pregnancy rates did not differ among groups, inseminating cows nearer to the time of ovulation may have given an advantage to the J-Synch group; however, results did not differ significantly and increasing P/AI in the ECP and EB groups may have been due to later ovulations within 24 h after FTAI, which is still considered appropriate in cattle (16).

In this study CL size 7 days after FTAI was greater in the J-Synch treatment group than in the other groups, which is in agreement with the results reported by de la Mata *et al.* (2018). Nevertheless, it is important to note that when all cows were considered, regardless of their treatment, those that became pregnant had larger dominant follicles and tended to have larger CL than those that did not become pregnant, which is in agreement with other studies (12, 45). Thus, larger ovulatory follicles should result in larger CL and greater P/AI regardless of FTAI treatment protocol, again making this a worthy endeavor.

Endocrine and uterine environments associated with elevated estradiol concentrations before ovulation have been reported to affect the maintenance of the conceptus (9). Furthermore, preovulatory estradiol concentrations have been reported to have a positive impact on subsequent conceptus development (17). Therefore, the positive relationship between the expression of estrus, follicle diameter, plasma P4 concentration and P/AI may be explained by the effects of estradiol and P4 on the endometrial tissue, as has been shown in other studies (10, 20, 24, 28).

## CONCLUSIONS

Although the prolonged proestrus protocol (J-Synch) resulted in the largest dominant follicle at the time of FTAI and the largest CL 7 days after ovulation, P/AI did not differ from the other estradiol/P4-based treatments in dual-purpose cows. Nevertheless, the present study has shown a positive association between the manifestation of estrus, follicle diameter at the time of FTAI and CL diameter 7 days later in suckled dual-purpose cows synchronized with estradiol/P4-based protocols for FTAI. Finally, it is concluded that the three protocols tested can be successfully applied in lactating dual-purpose cows in the Ecuadorian Amazon.

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### **AUTHOR CONTRIBUTIONS**

GAB: Conceptualization, Writing - review & editing; PRM: Writing - review & editing; JCLP: Supervision, Data curation, Methodology, Writing - original draft. AM: writing-review & editing.

## **CONFLICTS OF INTEREST**

The authors declare no conflicts of interest.

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### A DATA AVAILABILITY STATEMENT

The data that supports this study will be shared upon reasonable request to the corresponding author.

# Green solvents for the recovery of phenolic compounds from strawberry (*Fragaria* x *ananassa* Duch) and apple (*Malus domestica*) agro-industrial bio-wastes

# Uso de solventes verdes para la extracción de compuestos fenólicos a partir de residuos agroindustriales de frutilla (*Fragaria* x *ananassa* Duch) y manzana (*Malus domestica*)

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# ABSTRACT

We aimed to study the obtention of valuable phenolic compounds from tissue by-products of agro-industrial processing of apple (GS) and strawberry (RF) using green solvents and Soxhlet extraction methodology. The effects of solvent type [water (W); 80% ethanol, (EtOH)] and extraction ratio (1:10, 1:20, 1:30, and 1:40 p/v) were determined on total phenolic content (TPC), antioxidant capacity (DPPH), and the profile of phenolic compounds of the GS and RF extracts. The solvent type and the extraction ratio significantly affected TPC and DPPH of GS and RF extracts. Extraction with EtOH and 1:40 ratio produced the highest yields, obtaining an RF extract with 15.8 g GAE/Kg (TPC) and 19 mmol TE/Kg (DPPH). The tetra-galloyl glucose isomer and agrimoniin (0.8-0.4 g/Kg) were the main RF phenolic compounds of the eight identified. GS extract, obtained with EtOH and 1:40 ratio, had 11.9 g GAE/Kg (TPC) and 20.5 mmol TE/Kg (DPPH), having quercetin -3-o-glucuronide (0.43 g/Kg) the highest concentration among the eight phenolic compounds identified. The results highlight the potential of green solvents to obtain valuable compounds from low-cost raw materials, like the high-antioxidant capacity phenolic compound extracts obtained herein.

# Keywords

green extraction  ${\mbox{\circ}}$  bioactive compounds  ${\mbox{\circ}}$  hydrolysable tannins  ${\mbox{\circ}}$  flavonols  ${\mbox{\circ}}$  natural antioxidants

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### RESUMEN

El objetivo fue estudiar la extracción de compuestos fenólicos con alto valor bioactivo, a partir de subproductos del procesamiento de manzana (GS) y frutilla (RF), utilizando solventes amigables con el medio ambiente y la extracción Soxhlet. Se determinaron los efectos del tipo de solvente [agua (W); etanol 80% (EtOH)] y la relación de extracción (1:10, 1:20, 1:30 y 1:40 p/v) sobre el contenido de fenoles totales (TPC), capacidad antioxidante (DPPH) y perfil de compuestos fenólicos. El tipo de solvente y la relación de extracción afectaron significativamente el TPC y la DPPH. La extracción con EtOH a 1:40 produjo los rendimientos más altos, obteniendo un extracto de RF con 15,8 g GAE/Kg (TPC) y 19 mmol TE/Kg (DPPH). El isómero de tetra galoil glucosa y agrimoniin (0,8-0,4 g/Kg) fueron los principales compuestos fenólicos de RF, entre los ocho identificados. El extracto de GS, obtenido con EtOH a 1:40, tuvo 11,9 g GAE/Kg (TPC) y 20,5 mmol TE/Kg (DPPH), con quercetin-3-o-glucuronido (0,43 g/Kg) como el principal compuesto fenólico entre los ocho identificados. Los resultados destacan el potencial de los solventes verdes para obtener compuestos fenólicos de alta capacidad antioxidante potencialmente bioactivos a partir de materias primas de bajo costo.

### **Palabras claves**

extracción verde  ${\scriptstyle \bullet}$  compuestos bioactivos  ${\scriptstyle \bullet}$  taninos hidrolizables  ${\scriptstyle \bullet}$  flavonoles  ${\scriptstyle \bullet}$  antioxidantes naturales

## INTRODUCTION

One-third of fruit and vegetable production is wasted or lost in the production chain, producing avoidable and non-avoidable waste (3, 31). The former includes the losses generated by wrong handling during postharvest, processing, transport, storage, and distribution, and the non-avoidable waste is that part of the vegetable or fruit that must be eliminated for processing and sale (peels, seeds, core, and inedible parts). Therefore, the appropriate disposal of this waste is essential to reduce the environmental and economic impact of agro-industrial activity. The circular economy proposes different solutions to prevent this waste plant tissue from ending up in city landfills (35). Although the composition of fruit and vegetable by-products includes vitamins, minerals, and carbohydrates (27), as well as different bioactive compounds like phenolic compounds, these residual plant tissues are mainly used as ingredients in animal feed or as a source of energy for boilers. Waste fruit and vegetable tissues are sources of phenolic compounds with significant health benefits for consumers and antioxidant properties (24, 33). Phenolic compounds are synthesised in the plant's secondary metabolism during the normal development of plants. Their chemical structure has at least one aromatic ring with one or more hydroxyl groups, showing different biological activities according to their carbon bridges and hydroxyl substitution.

The production, consumption and industrial processing of apples and strawberries continue to increase worldwide, and so do the wasted tissues associated (12). Non-edible or non-processable fruit parts may have better composition and bioactivity than edible tissues (5, 34). The main phenolic compounds of strawberries are ellagitannins, phenolic acids and flavonoids like anthocyanins (32), and the principal apple phenolic compounds are catechins and proanthocyanidins (17), depending on the cultivar, the area, and the type of production of these crops. The extraction method of these compounds significantly impacts the extraction yields and their bioactive potential. Considering the structural diversity of the phenolic compounds, no single extraction system could produce the total recovery of these metabolites of interest from all vegetable tissues.

The solid-liquid extraction is commonly used to obtain phenolic compounds, and the success of this process depends on the plant matrix, kind of solvent, temperature, pH, and extraction technology. Polar protic solvents, like ethanol and methanol, can donate hydrogen bonds (33). Otherwise, polar aprotic solvents (acetone and ethyl acetate) have dipoles but do not possess hydrogen atoms bonded with a high electronegativity atom that can be donated in a hydrogen bond. Finally, apolar solvents (hexane and petroleum ether)

have bonds between atoms with similar electronegativities (33). The extraction technique also affects phenolic compound yield. The Soxhlet extraction technique is widely used for obtaining secondary metabolites from different plant tissues, being a reference method for comparing the performance of other extraction techniques. The Soxhlet extraction has good extraction yields without using large quantities of solvent, with a total extraction time of 1-6h, including multiple extraction cycles. This technique commonly employs flammable, hazardous, and toxic organic solvents. The low production cost of ethanol (by fermentation from renewable sources), low energy requirements for final disposal and moderate chronic toxicity make it a sustainable option to replace traditional solvents, along with water, considered the greenest solvent. Water application is generally limited to low-polarity metabolites; nevertheless, it is possible to broaden the solubilisation spectrum of water with suitable parameters (8). Therefore, green extraction with safe and non-toxic solvents, like water, ethanol, and binary ethanol-water mixtures, must be studied. Both solvents are considered safe and acceptable for use in the food industry by regulatory agencies (FDA). The ethanol-aqueous solutions for polyphenol extraction from plant tissues have better yields due to their azeotropic behaviour (1). Additionally, safe solvents in Soxhlet extraction could also yield higher amounts of phenolic compounds from some agro-industrial waste (7, 8, 26).

Complementary treatments with phenolic compounds that strengthen the immune system or their use as antioxidant agents would promote obtaining these phenolic compounds of interest at a low cost, facilitating the accessibility to a larger population and valuing the wasted agro-industrial tissues using green technologies. Therefore, this work aims to study the extraction of the phenolic compounds from strawberry by-products and 'Granny Smith' apple peel, characterise the obtained extracts using the Soxhlet method, evaluate the impact of two green solvents (water and ethanol 80%), and different solid-liquid ratios on the total phenolic content, phenolic profile, and in-vitro antioxidant activity of the extracts.

#### **MATERIALS AND METHODS**

### Plant material and Experimental design

The by-products of strawberry (RF) (*Fragaria x ananassa Duch*) cv 'Festival', consisting of sepals and stems, with non-processable parts of the fruit (part of the fruit closest to the sepal and peduncle), came from a single field (Coronda, Santa Fe, Argentina) during postharvest preparation for industrial processing. 'Granny Smith' apple peel (GS, 1 mm thickness) was obtained from the minimal processing of apples, according to Rodríguez-Arzuaga and Piagentini (2018). The RF and GS samples (89.2% and 80.4% moisture content, respectively) were weighed, packed in 40  $\mu$ m polyethylene bags, and stored at -20°C until processing. Before extraction assays, both samples were ground to a particle size <1 mm.

The effect of the extraction solvent (S) and the solid-liquid ratio (R) in the phenolic compound extraction of each vegetable tissue (RF and GS) was studied through a factorial experimental design. The two experimental variables of each factorial design were S and R, with two [S: water (W) and ethanol 80% (EtOH)] and four levels [R: 1:10, 1:20, 1:30 and 1:40 w/v], respectively. The total phenolic content (TPC), phenolic compound profile, and antioxidant capacity (DPPH) were determined on each extract of RF and GS. The extraction times were four hours for EtOH extractions and eight hours for W extractions (extraction times determined in preliminary assays). The extracts were cooled (20°C), filtered, and stored at -20°C for further analysis. Each phenolic compound extraction assay was performed in triplicate.

# **Total phenolic content (TPC)**

Each extract TPC was determined in triplicate by the Folin-Ciocalteu method (34). Gallic acid (Sigma-Aldrich, San Luis, Missouri, USA) was used as the standard reagent to perform the calibration curve, measuring the absorbance of the reaction at 760 nm in a spectrophotometer (Genesys 10s UV-Vis, Thermo Scientific<sup>™</sup>, Waltham, Massachusetts, USA). The concentration of TPC was reported as g of gallic acid equivalent (GAE) per kilogram of RF or GS (g GAE/Kg).

### Phenolic compound profile

The phenolic compound profile of the extracts was performed with an LC-20AT HPLC with a photodiode array detector (PAD), with the software Lab Solutions for data processing and control (Shimadzu Co., Kyoto, Japan). The separation was performed with a hybrid reverse phase column C18 Gemini  $5\mu$  110Å of  $250 \times 4.6$  mm, attached to a guard column (Phenomenex Inc, CA, USA). The analysis was performed according to Villamil-Galindo *et al.* (2021) for RF and Villamil-Galindo and Piagentini (2022a) for GS. The quantification of the identified compounds was performed using the following external standards (Sigma-Aldrich Inc. St. Louis, Missouri, USA): Ellagic acid (EA), Kaempferol-3-O-glucoside (K3G), Quercetin-3-O-glucoside (Q3G), Chlorogenic acid (ACI), Procyanidin B2 (PACB2), (-) Epicatechin (EPQN), (+) Catechin (CQN), Floretin (FLN), Gallic acid (GA), Coumaric acid (CUA), and Ferulic acid (FRA). The results were expressed in g per Kg of tissue by-product.

### Antioxidant capacity (DPPH)

The extract DPPH was determined using the 2.2-diphenyl-1-picrylhydrazyl radical (DPPH\*) scavenging assay, performed by triplicate (34). A volume (200  $\mu$ L) of the DPPH\* methanolic solution (0.08 mM) reacts with 25  $\mu$ L of extract or reference reagent 2,5,7,8-tetramethylchroman-2-carboxylic acid (Trolox, Sigma-Aldrich), and the absorbance was measured at 515 nm in a microplate reader (Asus UVM 340, Cambridge, England) after 2 h. The results were expressed as mmol Trolox equivalents/Kg of tissue by-product (mmol TE/Kg).

### Statistical analysis

All assays were carried out in triplicates, and data were presented as the mean ± standard deviation (SD). The effect of the solid-liquid ratio and the extraction solvent on the total phenolic content, phenolic compound profile and antioxidant capacity of RS and GS extracts were determined by the analysis of variance (ANOVA). Tukey's test (5% significance level) was used to determine the significant differences among treatment means. Besides, the correlation between the individually identified phenolic compounds and the antioxidant capacity determined in each extract was determined using Pearson's correlation test. The statistical analysis was performed with STATGRAPHICS Centurion XV software (StatPoint Technologies Inc., Warrenton, VA, USA).

## **RESULTS AND DISCUSSION**

# Phenolic compounds recovery from strawberry agro-industrial by-products (RF)

Both Soxhlet extraction parameters [solid-liquid ratio (R) and type of solvent (S)] affected (p<0.001) the yields of total phenolic content (TPC), the phenolic compounds profile, and the antioxidant capacity (DPPH) of the strawberry agro-industrial by-products extracts. The interaction term between R and S was also significant (p<0.001), meaning that R affected in a different way TPC, DPPH, and the phenolic compounds profile of the extracts depending on the type of solvent used.

The increase of R values improved the TPC yields of RF Soxhlet extraction with water (W), with the highest yield (10.69 g GAE/Kg) obtained at R 1:40, up to 48% higher than those obtained at lower R values (table 1, page 153). The concentration gradient between the solute and solvent was the driving force for the diffusion process. Therefore, better extraction yields were obtained with the highest R-value (1:40). Therefore, strawberry by-products have great potential as a source of bioactive compounds obtained with water.

The RF Soxhlet extraction with EtOH significantly increased the recovery of TPC as compared with W by 47-170% (table 1, page 153). Contrary to W extractions, R did not affect the TPC yields of the EtOH extracts (mean value 15.37 g GAE/Kg) (table 1, page 153). The phenolic compounds of RF could present preferential solvation when using water-ethanol. The compounds would show hydrophobic hydration with the few polar groups they could have in their structure, and a large amount of (-OH) groups would allow more interaction with water (6). Therefore, this phenomenon would produce more effect than the increase in the extraction ratio. However, the use of EtOH on the Soxhlet extraction system significantly

increased the TPC yields compared to RF water extracts, improving extraction yields by up to 170% (table 1). Therefore, the lower the extraction ratio, the smaller the solvent needed, and the lower the production costs of the extracts.

**Table 1.** Total phenolic content (TPC) and antioxidant capacity (DPPH) of the extracts from strawberry by-products (RF) and 'Granny Smith' apple peel (GS).

**Tabla 1.** Contenido de fenoles totales (TPC) y capacidad antioxidante (DPPH) de los extractos de sub-productos de frutilla (RF) y cáscara de manzana 'Granny Smith' (GS).

Sample	R	TPC		DPPH	
		W	EtOH	W	EtOH
RF	1:10	$5.58 \pm 0.08^{CB}$	$15.10 \pm 0.14^{dA}$	$4.82 \pm 0.20^{\text{cB}}$	$9.95 \pm 0.08^{aA}$
	1:20	$6.57 \pm 0.23^{bcB}$	$15.13 \pm 0.18^{dA}$	$10.43 \pm 1.16^{\text{bB}}$	15.44 ± 0.72 <sup>bA</sup>
	1:30	$6.27 \pm 0.02^{\text{bB}}$	$15.41 \pm 0.21^{dA}$	$10.70 \pm 0.21^{\text{bB}}$	19.01 ± 0.86 <sup>cA</sup>
	1:40	$10.69 \pm 0.69^{aB}$	$15.82 \pm 1.76^{dA}$	$15.34 \pm 0.12^{aB}$	18.99 ± 1.16 <sup>cA</sup>
GS	1:10	$4.18 \pm 0.02^{aB}$	$6.33 \pm 0.12^{dA}$	$3.60 \pm 0.14^{dB}$	$7.26 \pm 0.20^{dA}$
	1:20	$3.90 \pm 0.31^{abB}$	7.27 ± 0.001 <sup>cA</sup>	8.22 ± 0.35 <sup>cB</sup>	12.89 ± 0.38 <sup>cA</sup>
	1:30	3.55 ± 0.13 <sup>bB</sup>	$7.92 \pm 0.09^{\text{bA}}$	11.48 ± 0.39 <sup>bB</sup>	13.75 ± 0.15 <sup>bA</sup>
	1:40	$4.35 \pm 0.24^{aB}$	$11.90 \pm 0.01^{aA}$	$14.05 \pm 1.09^{aB}$	$20.51 \pm 0.79^{aB}$

Mean ± standard deviation. R: Solid-liquid ratio. W: water; EtOH: 80% ethanol. Different capital letters and lowercase letters indicate significant differences (p<0.05) by Tukey's test, between solvent and among different solid-liquid ratios, respectively.

Promedio ± desviación estándard. R: relación sólido-líquido. W: agua; EtOH: 80% etanol. Letras mayúscula y minúsculas indican diferencias significativas (p<0,05) por el test de Tukey, entre solvente y entre diferentes relaciones sólido-líquido, respectivamente.

Furthermore, these results confirm that binary ethanol-water mixtures are suitable bio-solvents for obtaining phenolic compounds due to their polar protic properties. The yields obtained for RF with EtOH using Soxhlet extraction were higher than those reported for pandan leaves (6.6 g GAE/Kg), mango by-products (4.5-6.6 g GAE/Kg), asparagus (2.8-3.7 g GAE/Kg), cauliflower (1.1-1.8 g GAE/Kg), and bergamot lemon (4-10 g GAE/Kg) (4, 14, 15, 27).

Regarding the antioxidant capacity of RF water extracts, they were significantly affected by R-value, comparable to TPC. The highest DPPH value was obtained for the 1:40 ratio (15.3 mmol TE/Kg), being up to 69% higher than that obtained at R 1:10. Otherwise, like with the content of phenolic compounds, EtOH improved the antioxidant capacity of the RF extracts obtained compared with W extracts. The EtOH RF extracts with the highest DPPH values (18.9-19.0 mmol TE/Kg) were those obtained with R 1:40 and 1:30 (p>0.05), respectively. The phenolic compounds are excellent antioxidants of natural origin due to their reducing- capacity, shown by the highly significant correlation between the TPC and the antioxidant capacity of the RF extracts.

Eight major phenolic compounds were identified for the strawberry agro-industrial waste tissue (RF), belonging to three main phenolic compound classes: hydrolysable tannins, ellagic acid derivatives, and flavonols (figure 1a and figure 2, page 154). Tetragalloyl-glucose isomer (TGI) and galloyl-bis-HHDP-glucose dimer (agrimoniin) (AGN) were identified among the hydrolysable tannins; ellagic acid pentoxide (EAP) and free ellagic acid (EA) among the ellagic acid derivatives; and finally, the flavonols were represented by quercetin-3-O-glucuronide (Q3G), quercetin hexoxide (QHS), kaempferol-3-O-glucuronide (K3G), and kaempferol hexoxide (KHS).

TGI: Tetragalloyglucose isomer, EAP: Ellagic acid pentoxide, AGN: Dimer of galloyl-bis-HHDPglucose (agrimoniin isomer), EA: Ellagic acid, Q3G: Quercetin-3-0-glucuronide, QHS: Quercetin Hexoxide, K3G: Kaempferol-3-O-glucuronide. (+) CTQN: Catechin, PACB2: Procyanidin B2, (-) EPQN: Epicatechin, PACT: Procyanidin tetramer, QPN: Quercetin pentoxide. TGI: Tetragalloyglucosa isomero, EAP: pentoxidode ácido Ellagico, AGN: Dimero de galloylbis-HHDP-glucosa (agrimoniin isomero), EA: ácido Ellagico, Q3G: Quercetin-3-O-glucuronido, QHS: Quercetin Hexoxido, K3G: Kaempferol-3-0glucuronido. (+) CTQN: Catequina, PACB2: Procianidina B2, (-) EPQN: Epicatequina, PACT: Procianidin tetramero, QPN: Quercetina pentoxido.

TGI: Tetragalloyglucose isomer, EAP: Ellagic acid pentoxide, AGN: Dimer of galloyl-bis-HHDPglucose (agrimoniin isomer), EA: Ellagic acid, Q3G: Quercetin-3-0-glucuronide, OHS: Quercetin Hexoxide, K3G: Kaempferol-3-0-glucuronide, **TPC**<sub>HPLC</sub>: Total phenolic compounds analyzed by HPLC. Different lowercase letters indicate significant differences (p<0.05) by Tukey's test, between different solidliquid ratios.

TGI: Tetragalloyglucosa isomero, EAP: pentoxido de ácido Ellagico, AGN: Dimero de galloylbis-HHDP-glucosa (agrimoniin isomero), EA: ácido Ellagico, Q3G: Quercetin-3-O-glucuronido, QHS: Quercetin Hexoxido, K3G: Kaempferol-3-0glucuronido, **TPC**<sub>HPLC</sub>: Compuestos fenólicos totales analizados por HPLC. Diferentes letras minúsculas indican diferencias significativas (p<0,05) por el test de Tukey, entre diferentes relaciones sólido-líquido.



**Figure 1.** Typical HPLC-UV chromatogram obtained of (a) strawberry by-products at 254 nm and (b) 'Granny Smith' apple peel at 320 nm.

Figura 1. Cromatograma típico HPLC-UV obtenido para (a) sub-productos de frutillas a 254 nm y (b) cáscara de manzanas 'Granny Smith' a 320 nm.



Figure 2. Phenolic compounds from strawberry by-products (RF) extracted with a) water (W) and b) 80% ethanol (EtOH).

Figura 2. Compuestos fenólicos de los sub-productos de frutilla extraídos con a) agua (W) and b) 80% etanol (EtOH).

For the water extracts at R 1:10, hydrolysable tannins represented 38.5% of the total phenolic compounds, similar to flavonols with 39.9%, followed by ellagic acid derivatives with 21.6% (figure 2a, page 154). However, as the extraction ratio increased, hydrolysable tannins represented 64% of the total phenolic compounds in the extracts obtained with R 1:40, showing that the increase in the concentration gradient in the extraction solvent (W) allowed greater recovery of hydrolysable tannins (33).

The highest concentration of tetragalloylglucose isomer (TGI) in W extracts was obtained at R 1:40 (0.18 g/Kg); for the other R-values, the TGI concentration was similar (0.07-0.1 g/Kg) (figure 2a, page 154). EtOH extraction significantly improved the TGI yields for all extraction ratios, with the maximum concentration at R 1:40 (0.83 g/Kg). The Ellagic acid pentoxide (EAP) had the highest concentration, 0.088 g/Kg (p<0.05), in the EtOH extracts (figure 2, page 154). EAP concentrations were higher than those reported for the stem of Sanguisorba Officinalis L (Rosaceae family) (0.038 g/Kg) (20). Agrimoniin (AGN) is a compound derived from hexahydroxydiphenic acid (HHDP), considered a taxonomic marker of the Rosaceae, with great importance in the nutraceutical industry due to its bioactive properties (30). The AGN yield obtained with water was about 0.064-0.084 g/kg for the higher R values. The AGN yields were significantly improved for all R-values (p<0.05) when EtOH was used, with a maximum concentration of 0.44 g/Kg at R 1:40 (figure 2, page 154). This AGN concentration was higher than the reported for whole strawberry fruit extracts (0.12 g/Kg) obtained with 70% methanol (25), showing this residual tissue as a valuable source of AGN. The antioxidant properties of hydrolysable tannins have been reported *in-vitro* and *in-vivo* (16). Simirgiotis et al. (2010) reported that cyanidin glucosides and ellagic acid were the compounds with the highest participation in the antioxidant activity in the edible part of strawberries. For RF, the hydrolysable tannins TGI and AGN had a significant correlation (p<0.01) with the DPPH\* antioxidant capacity. These compounds have polyhydric alcohol in the centre, and their hydroxyl groups could be partially esterified with ellagic acid or HHDP, having the capacity to yield electrons and thus neutralise the free radicals present (14). The ellagic acid (EA) concentrations were lower in W extracts for any extraction ratio (p>0.05). Like AGN, EtOH improved the extraction of EA, with a maximum concentration of 0.10 g/Kg R 1:40 (figure 2, page 154). The consumption of ellagic acid derivatives was associated with numerous health benefits since, in-vivo conditions, different types of urolithins were metabolised by the microbiota, which had powerful antiproliferative and cancer cell apoptosis-inducing activities (21).

The flavonol Quercetin-3-o-glucuronide (Q3G) concentrations in RF water extracts were 0.002-0.04 g/Kg, obtaining the higher at R 1:40 and 1:30 (p>0.05). Nevertheless, using 80% ethanol significantly increased Q3G yields up to 0.11 g/Kg (figure 2, page 154), similar to chokeberry extracts (11). QHS concentrations obtained with EtOH (0.12-0.13 g/Kg) were higher than those obtained with W (0.04-0.05 g/Kg). Kaempferol is one of the most common flavonols in different botanical species. In the cell vacuoles, Kaempferol tends to glycosidate with some carbohydrates to have more stability in the pH of the medium (28). The highest kaempferol-3-o-glucuronide (K3G) concentration obtained with water was 0.043 g/Kg. EtOH improved the K3G extraction yields by about 50% (figure 2, page 154). K3G values in RF were close to those reported for green tea (60% methanol), a popular antioxidant infusion (19). Finally, the Kamepferol Hexoxide (KHS) yield increased by up to 91% in EtOH extractions compared to water extracts (figure 2, page 154).

Therefore, the agro-industrial strawberry by-product showed a large variability of phenolic compounds of interest. The highest recovery of total phenolic compounds ( $TPC_{HPLC}$ ) with W was achieved with R 1:40 (0.40 g/Kg). As expected, the EtOH increased  $TPC_{HPLC}$  up to 425%, obtaining the highest concentration also at R 1:40 (figure 2, page 154). Similar to the results of TPC and DPPH, the extractions with the ethanol-water binary mixture (80:20) had the highest recovery of phenolic compounds with high antioxidant capacity. The maximum  $TPC_{HPLC}$  content (1.70 g/Kg) obtained for RF EtOH extracts was comparable to that reported for strawberry plant leaves (1.95-2.07 g/Kg) obtained with methanol-formic acid (99:1) (30). The phenolic compounds of RF have an excellent antioxidant, anti-inflammatory and anticarcinogenic potential for colorectal cancer (34, 36). Therefore, the high concentration of hydrolysable tannins and ellagic acid derivatives in RF enables the promotion of this kind of agro-industrial by-products as a low-cost source of healthy compounds.

# Phenolic compounds recovery from 'Granny Smith' apple peel (GS).

The solid-liquid ratio (R) and the type of solvent (S) affected (p<0.001) the content of phenolic compounds and the antioxidant capacity of GS extracts. The interaction term between R and S was also significant (p<0.001) for TPC, DPPH, and phenolic compounds profile.

The use of EtOH improved the TPC recovery from GS, like RF (table 1, page 153). The TPC of GS EtOH extracts increased as R increased, obtaining the highest yield (11.9 g GAE/Kg) for R 1:40. Castro-López *et al.* (2017) reported that R-values higher than 1:20 increased the recovery of phenolic compounds. Binary alcohol-water mixtures offer an eco-friendly solvent system for obtaining phenolic compounds from different wasted plant matrices than those using pure ethanol or other organic solvents. The water and ethanol mixture act synergistically and could provide a suitable polarity range for extracting phenolic compounds (medium-high polarity). The former is fundamental as a swelling agent of the plant matrix, allowing the lower viscosity ethanol to diffuse through the material and break the non-covalent interactions between the solute and the matrix, facilitating the preferential solvation sphere transferring the analyte to the dissolution medium (33).

Phenolic compounds are the plant-secondary metabolites with the highest reported antioxidant activity. Each compound antioxidant capacity differs due to its oxidation-reduction reactions, phenyl ring structure resonance, and hydroxyl group substitution pattern (32). The antioxidant capacity of GS extracts at R 1:40 is 47% higher in EtOH extract than in W extract. The DPPH values obtained were comparable to those reported for other plant materials by Soxhlet extraction (sugar beet molasses, rapeseed, and flowers of *Jatropha integerrima*) (10, 13). The use of EtOH enhanced the antioxidant capacity of GS extracts as R increased (table 1, page 153), comparable to TPC, and therefore, showing a strong correlation between TPC and DPPH (p<0.01).

The two main classes of phenolic compounds identified and quantified in 'Granny Smith' apple peel (GS) were the flavan-3-ols with (+) catechin [(+) CTQN], Procyanidin B2 (PACB2), (-) epicatechin [(-) EPQN], and Procyanidin tetramer (PACT); and the flavonols with the Quercetin-3-o-glucuronide (Q3G), Quercetin pentoxide (QPN), Quercetin Hexoxide (QHS), and Kaempferol-3-o-glucuronide (K3G) (figure 1b, page 154, and figure 3, page 157).

For the GS W extracts, the flavan-3-ols and flavanols represented each 50 % of the total phenolic compounds (R 1:10, 1:20 and 1:30), increasing the flavan-3-ols proportion with R 1:40 up to 76 % of the total of the quantified compounds, showing the affinity of this class of phenolic compounds for a polar solvent like water (figure 3a, page 157). Nevertheless, flavonols accounted for more than 50% of the total phenolic compounds in all the EtOH extractions, with a maximum of 84% in the extracts with R 1:40 (figure 3b, page 157).

R did not affect (p>0.05) the (+)CTQN extraction yield with water. EtOH extractions increased (+)CTQN yields up to 0.066 g/kg, 77% higher than W extracts (figure 3, page 157). The (+)CTQN yields obtained were lower than those reported by Almeida et al. (2017) for 'Granny Smith' apple peel extracted with 100% acetone (0.17 g/Kg). PACB2 (epicatechin-epicatechin dimer) is the most common proanthocyanidin determined in high concentrations in fruits like peaches, apples, and plums. The PACB2 content in W extracts increased with R; it was at least 71% higher for R 1:40 than for the other extraction ratios. However, ethanol did not improve the PACB2 extraction yields. Procyanidin B2 in a liquid medium from 90°C onwards starts a degradation process by oxidation and epimerisation, lowering the procyanidin B2 recovery. The highest concentration of EPQN, reported as the main phenolic compound in apples (23), was achieved with W and R 1:40 (0.1 g/Kg), being higher than that reported for apple pomace extracts (0.02 g/Kg) (20). The highest PACT concentration in W extracts was 0.04 g/Kg, obtained at R 1:40. The EtOH improved PACT yields (p<0.05) for all R values (figure 3, page 157). Procyanidins are oligomers composed of catechin and epicatechin; their structure and high molecular weight give them different bioactive and functional properties for the food industry.



Figure 3. Phenolic compounds from 'Granny Smith' apple peel (GS) extracted with a) water (W) and b) 80% ethanol (EtOH).

Figura 3. Compuestos fenólicos de cáscara de manzana compounds from 'Granny Smith' (GS) extraidos con a) agua (W) and b) 80% etanol (EtOH).

Q3G extraction yields with W were affected by R, obtaining the highest one at R 1:40, 82% higher than the yields obtained at lower R values. Nevertheless, the EtOH improved Q3G extraction yields, as expected for a medium polarity compound. The highest Q3G yield with EtOH (0.43 g/Kg) was obtained at R 1:40, 50% higher than those obtained at lower R values (figure 3). Moreover, Q3G was the GS individual compound with the highest correlation with the antioxidant capacity, mainly for its structure and hydroxyl groups that allowed the donation of electrons, neutralising the free radicals present in the medium (32). Contrarily, RF flavonols did not show a highly significant correlation with the antioxidant capacity determined by DPPH. Consequently, the bioactive potential did not depend only on the bioactive compound concentration but also on the interaction with the food matrix. The values obtained with EtOH were similar and even higher than those reported for 'Granny Smith' apple peel acetone extracts (0.18-0.4 g/Kg) (2).

The highest concentration of QPN, the other quercetin glucoside derivative, was obtained with EtOH at R 1:40 (0.08 g/Kg). The yields of QHS in EtOH extracts were higher than in W, obtaining the highest concentration at R 1:40 (0.14 g/Kg). According to previous reports, quercetin and its glycosides have a low bioavailability (16-25%) due to their low water solubility and crystalline structure at body temperatures (16). Like the other flavonols, EtOH enhanced the recovery of K3G, with the highest concentration (0.25 g/Kg) at R 1:40 (figure 3).

Considering the sum of the compounds identified and quantified, the  $TPC_{HPLC}$  obtained for W extracts increased with R, determining the highest concentration in W at R 1:40 (0.55 g/Kg) (figure 3a). Higher  $TPC_{HPLC}$  values were obtained with EtOH and higher R values.  $TPC_{HPLC}$  extracted with EtOH 1:40 (1.07 g/Kg) was higher than that reported for

(+)CTQN: Catechin, PACB2: Procyanidin B2, (-)EPQN: Epicatechin, PACT: Procvanidin tetramer. Q3G: Quercetin-3-0-glucuronide, **QPN:** Quercetin pentoxide. OHS: Ouercetin Hexoxide. K3G: Kaempferol-3-0-glucuronide, TPC<sub>HPLC</sub>: Total phenolic compounds analyzed by HPLC. Different lowercase letters indicate significant differences (p<0.05) by Tukev's test, between different solid-liquid ratios. (+)CTQN: Catequina, PACB2: Procianidina B2, (-)EPQN: Epicatequina, PACT: Procianidina tetramero, Q3G: Quercetina-3-O-glucuronido, QPN: Ouercetina pentoxido. QHS: Quercetina Hexoxido, K3G: Kaempferol-3-0glucuronido, TPC<sub>HPLC</sub>: Compuestos fenólicos totales analizados por HPLC. Diferentes letras minúsculas indican diferencias significativas (p<0,05) por el test de Tukey, entre diferentes relaciones sólidolíquido. the apple pulp (17), showing the bioactive potential of apple peel. The GS  $\text{TPC}_{\text{HPLC}}$  highly correlates with antioxidant activity (R<sup>2</sup> 0.87), mainly due to flavonols. These results encourage the integral use of the apple peel as a source of valuable compounds, focusing on green solvents use with low environmental impact and cost (35).

### CONCLUSIONS

There is a growing demand for nutraceutical products of vegetable origin, as their frequent consumption has been associated with a decreasing risk of having chronic non-transmissible diseases. The market for nutraceutical compounds is booming, and the extraction of bioactive compounds using clean solvents from agro-industrial waste tissues, like the strawberry by-products and apple peel, presents an opportunity to reduce costs and the environmental impact. The conventional Soxhlet extraction technique has good yields, low complexity, and high efficiency, allowing optimal use of natural resources, especially those that are rejected for industrial processing, like the waste tissue produced during the postharvest trimming of the strawberry (about 7-20% of the fruit intended for minimal processing) and Granny Smith apple peel (about 12% of the fruit intended for minimal processing).

This study demonstrates that waste vegetable tissues can be transformed into valuable phenolic compounds with antioxidant properties using eco-friendly solvents such as water and ethanol. The extracts with the highest content of phenolic compounds and antioxidant capacity were obtained for Soxhlet extraction with 80% ethanol and 1:40 extraction ratio for both the strawberry by-products (15.8 g GAE/Kg and 19 mmol TE/Kg) and the 'Granny Smith' apple peel (11.9 g GAE/Kg and 20.5 mmol TE/ Kg). Additionally, eight main phenolic compounds were identified and quantified in both waste tissues. The hydrolysable tannins, like Tetragalloyglucose isomer (TGI: 0.83 g/Kg) and Dimer of galloyl-bis-HHDP-glucose (agrimoniin isomer, AGN: 0.44 g/Kg), were the main phenolic compounds extracted from RF, while flavonols accounted for 83.7% of the total extracted phenolic compounds from GS, obtaining for Quercetin-3-O-glucuronide the highest yield (Q3G: 0.43 g/Kg).

These results demonstrated the importance of by-products as low-cost sources of bioactive compounds with high nutraceutical potential through a circular process approach in the fruit and vegetable industry. Currently, these bio-wastes are disposed of in landfills without any use. The information obtained in this study provides a pathway towards the integral use of strawberry and apple by-products. The challenge is to continue studying the development of a procedure for obtaining bioactive compounds from strawberry by-products and 'Granny Smith' apple peel with higher yields, shorter extraction times and lower energy consumption, using more sustainable and efficient technologies stimulating an integral use of these by-products.

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