

Genotype-by-environment interaction and selection of superior *Physalis peruviana* L. genotypes

Interacción genotipo por ambiente y selección de genotipos superiores de *Physalis peruviana* L.



FRANCY LILIANA GARCÍA-ARIAS^{1, 3}
ERIKA SÁNCHEZ-BETANCOURT¹
FRANKLIN GIOVANNY MAYORGA-CUBILLOS¹
DAVID RODRÍGUEZ-PUERTAS²
VÍCTOR MANUEL NÚÑEZ-ZARANTES^{1, 3}

Cape gooseberry fruit obtained in an experimental plot in the department of Cundinamarca.

Photo: E. Sánchez-Betancourt

ABSTRACT

Cape gooseberry (*Physalis peruviana* L.) productivity in Colombia can be increased by developing high-yielding and adaptable varieties identified in multi-environment trials. In this study, the genotype-by-environment interaction (G×E) for fruit yield and fruit weight of 13 cape gooseberry genotypes at seven locations was evaluated using a randomized complete block design. G×E interaction was significant for yield and fruit weight, suggesting a differential response of genotypes across environments. Through the AMMI analysis, similar and contrasting environments were identified, as well as the genotypes that contributed the most to the G×E interaction. Genotypes I, B, D, and H were the high yielding, ranging from 25.2 to 27.3 t ha⁻¹, so they could be recommended for commercial cultivation. Genotypes B and D was stable in yield and widely adapted; while the genotypes I and H showed a specific adaptation for yield and exhibited heavier fruits. Genotype R1 exhibited the greater fruit weight in most locations except Ipiales but showed low fruit yield. The suitable locations for the cultivation of cape gooseberry were Pasto, Puerres, and Ipiales since they presented the highest yields and fruit weight.

Additional key words: cape gooseberry; G×E; AMMI; mega-environments; fruit yield; adaptability.

¹ Corporación Colombiana de Investigación Agropecuaria – Agrosavia, Centro de Investigación Tibaitatá, Bogota (Colombia). ORCID García-Arias, F.L.: <https://orcid.org/0000-0003-3112-9950>; ORCID Sánchez-Betancourt, E.: <https://orcid.org/0000-0002-3024-3180>; ORCID Mayorga-Cubillos, F.G.: <https://orcid.org/0000-0003-1482-0280>; ORCID Núñez-Zarantes, V.M.: <http://orcid.org/0000-0002-5087-9864>

² Corporación Colombiana de Investigación Agropecuaria – Agrosavia, Centro de Investigación Obonuco, San Juan de Pasto (Colombia). ORCID Rodríguez-Puertas, D.: <https://orcid.org/0000-0001-7161-3013>

³ Corresponding authors. flgarcia@agrosavia.co; vnunez@agrosavia.co

RESUMEN

La productividad de la uchuva (*Physalis peruviana* L.) en Colombia puede incrementarse mediante el desarrollo de variedades estables de alto rendimiento, identificadas en ensayos multi-ambientes. En este estudio, se evaluó la interacción genotipo por ambiente (G×E) para el rendimiento y peso de frutos de 13 genotipos de uchuva en siete localidades utilizando un diseño de bloques completos al azar. La interacción G×E fue significativa para rendimiento y peso de fruto, lo que sugiere una respuesta diferencial de los genotipos a través de los ambientes. Mediante el análisis AMMI se identificaron ambientes similares y contrastantes, así como los genotipos que más contribuyeron a la interacción G×E. Los genotipos I, B, D y H presentaron altos rendimientos, que oscilaron entre 25,2 a 27,3 t ha⁻¹, por lo tanto, podrían recomendarse para el cultivo comercial. Los genotipos B y D fueron estables en rendimiento y de adaptación amplia; mientras que los genotipos I y H mostraron una adaptación específica para rendimiento y presentaron frutos de mayor peso. El genotipo R1 mostró frutos de mayor tamaño en la mayoría de las localidades excepto Ipiales, pero presentó bajo rendimiento. Las localidades adecuadas para el cultivo de uchuva fueron Pasto, Puerres e Ipiales por presentar los mayores rendimientos y peso de fruto.

Palabras clave adicionales: uchuva; G×E; AMMI; mega-ambientes; rendimiento de fruto; adaptabilidad.

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INTRODUCTION

Cape gooseberry (*Physalis peruviana* L.) is a native plant from the Andean region that belongs to the Solanaceae family (Fischer and Melgarejo, 2020). The cape gooseberry fruits contain high amounts of vitamins A, B, and C, polyunsaturated fatty acids, phytosterols, essential minerals, antioxidants, withanolides, and physalins. It has medicinal benefits, with antitumor, anti-inflammatory, antiparasitic antimicrobial, and anticancer properties (Demir *et al.*, 2014; El-Beltagi *et al.*, 2019; Huang *et al.*, 2020, Miranda and Fischer, 2021).

About 80 species of *Physalis* have been reported worldwide, but only three ecotypes have been cultivated in Colombia: Kenya, South Africa, and Colombia, whose names are derived from the country where they come from (Peña *et al.*, 2010; Valderrama and Núñez, 2021). The Colombia ecotype is the most important genetic material in the producing areas since it stands out for its color, aroma, and high sugar content in the fruit (Miranda, 2005). However, the commercial plantations of this ecotype have shown high phenotypic variability probably due to the mixture of the different ecotypes grown in the country (Bonilla *et al.*, 2019).

Colombia produces around 20,000 t year⁻¹ of cape gooseberry fruits, mainly in the departments of Cundinamarca, Boyaca, Antioquia, and Nariño. This

fact has positioned Colombia as the world's leading producer of cape gooseberries (Minagricultura, 2020, 2022). However, the national yields have decreased by 30%, from 17.8 t ha⁻¹ in 2009 to 10.4 t ha⁻¹ in 2020 (Agronet, 2020; Minagricultura, 2020) due to the discarding of cracked fruits (up to 50%) (Fischer *et al.*, 2021; Criollo *et al.*, 2014), also due to the use of plant material of unknown origin and identity, and mixture of ecotypes (Rodríguez and Bueno, 2006). Therefore, it is important to develop high-yielding varieties with high-quality fruit to meet national demand and access new export destinations.

Several studies have been carried out to identify genetic resources useful for the cape gooseberry breeding program. Some of these investigations include the study of ploidy level and genetic diversity, agronomic characterization, the development of haploid and dihaploid lines via anther culture, and the generation of F1 populations (Suescún *et al.*, 2011; Liberato *et al.*, 2014; García-Arias *et al.*, 2018; Mayorga-Cubillos *et al.*, 2019; Franco-Florez *et al.*, 2021; Sánchez-Betancourt and Nuñez, 2022). From these studies, nine promising materials with desirable attributes of yield and fruit quality were chosen for evaluation across multiple locations to assess G×E interaction and select genotypes with specific or broad adaptation in Colombia.

Several multivariate methods have been proposed to study G×E interaction, including analysis of variance (ANOVA), principal components analysis (PCA), linear regression, genotype plus genotype by environment (GGE), and additive main effects and multiplicative interactions (AMMI). The AMMI is one of the most used multivariate methods for statistical analysis of yield trials and includes ANOVA and PCA analysis in a unified approach (Zobel *et al.*, 1988; Crossa, 1990). According to Gauch *et al.* (2008), this model has shown results with a clear agronomic meaning and has been effective due to its ability to capture the G×E interaction. At the same time, this model facilitates the identification of mega-environments and allows breeders to interpret the data and make reliable decisions in the selection of superior genotypes for each environment (Zobel *et al.*, 1988; Gauch *et al.*, 2008; Movahedi *et al.*, 2020).

The AMMI model includes more than one model, that is, a family of models, discriminated by the number of components that it retains and are called AMMI0, AMMI1, AMMI2, AMMI3, and so on. AMMI0 being the one that does not contain components, while AMMI1 contains a component, etc. The model AMMI1 is used to estimate the prediction of yield across environments and combines the additive main effect from genotypes and environments with the interaction effects estimated from the principal component 1 (PC1). According to Crossa *et al.* (1991) and Gauch (2013), AMMI1 is a simpler and more suitable model for mega-environment delineation and identification of winning genotypes in each environment compared to the other AMMI family members.

Considering the few reported studies in G×E interaction in cape gooseberry and the need to identify genotypes with differential response across the environments, it is crucial to carry out studies on this topic. Therefore, this research aimed to evaluate the G×E interaction of 13 cape gooseberry genotypes in different locations and identify high-yielding genotypes with broad or specific adaptation to Colombia.

MATERIALS AND METHODS

Genetic material and experimental locations

The plant material included nine elite genotypes labeled “A to I” and four commercial cape gooseberry genotypes (N, R1, R2, and R3) currently in use in the

Agrosavia breeding program. These genotypes were previously evaluated in different studies and selected for their high yield and high sugar content (García-Arias *et al.*, 2018; Mayorga *et al.*, 2019). The elite genotypes came from the Universidad Nacional de Colombia - UNAL and Corporación Colombiana de Investigación Agropecuaria - Agrosavia collections, whereas the commercial genotypes came mostly from farmers in the different producing areas. The genotype N (Colombia ecotype) is widely cultivated across the country and the genotypes R1, R2, and R3 are cultivated in Cundinamarca, Antioquia, and Nariño departments, respectively (Tab. 1).

The evaluation of the genotypes was conducted across seven locations in three departments (Antioquia, Cundinamarca, and Nariño) in Colombia. The locations were La Union (LU), Rionegro (RN), and San Vicente (SV) in Antioquia; Mosquera (MO) in Cundinamarca; and Ipiales (IP), Pasto (PA), and Puerres (PU) in Nariño. The locations are representative of the producing areas of the country, and they are located between 2,117 and 2,856 m a.s.l. with different rainfall regime that range from 700 to 1,800 mm. The average of temperature of the locations ranges from 10.2 to 17.0°C and showed a soil pH between 4.9 to 6.19 (Tab. 2).

Trial establishment and data collection

Genotypes were tested under field conditions during the same year using a randomized complete block design with four replicates in each location. Each experimental plot had five plants per replicate and was planted with a spacing of 2.5 m between rows and 2.0 m between plants, with a total area of 5 m² per plot. Fertilization was carried out according to the soil analysis for each location.

The evaluated traits were fruit yield and fruit weight. Fruit yield (FY) was determined from the accumulated fruit weight collected in 15 harvests from 5.5 to 10 months after transplantation, expressed as t ha⁻¹ per year for each genotype; and fruit weight without calyx (FW) as the average weight of one fruit obtained from weighing 5 fruits per replicate and harvest.

Statistical analysis

Data from yield and fruit weight trials at all locations were summarized and subjected to analysis of variance (ANOVA), with Dunnett's and Tukey's test, at $P < 0.05$, for genotypes and locations, respectively.

Table 1. Passport information of cape gooseberry genotypes included in the study.

Label	Genotype	Other * codes	Country	Department	Town	Germplasm collection	Observation
A	12U336	-	Colombia	--	--	Agrosavia	Experimental genotype derived from anther culture
B	12U346	-	Colombia	--	--	Agrosavia	Experimental genotype derived from anther culture
C	12U372	-	Colombia	--	--	Agrosavia	Experimental genotype derived from anther culture
D	13U406	09U047	Colombia	Boyaca	Tota	UNAL	Experimental genotype
E	13U413	09U279	Colombia	Nariño	Pasto	Agrosavia	Experimental genotype
F	13U416	12U381	Colombia	Boyaca	Pesca	UNAL	Experimental genotype
G	09U089	-	Colombia	Antioquia	San Pedro de Uraba	Agrosavia	Experimental genotype
H	09U278	-	Colombia	Nariño	Pasto	Agrosavia	Experimental genotype
I	12U396	-	Colombia	Antioquia	El Peñol	Agrosavia	Experimental genotype
R1	13U411	09U274	Colombia	Cundinamarca	--	Private company	Cundinamarca Regional control
R2	09U171	-	Colombia	Antioquia	--	Farmer	Antioquia Regional control
R3	13U417	-	Colombia	Nariño	Ipiales	Farmer	Nariño Regional control
N	09U275		Colombia	Cundinamarca	Choconta	Agrosavia	National control

* Garcia-Arias *et al.* (2008); Mayorga *et al.* (2019).

Table 2. Geographic and environmental characteristics of the experimental locations in Colombia.

Department	Location	Latitude (N)	Longitude (W)	Altitude (m a.s.l.)	Soil texture	Soil pH	Average temperature (°C)	Average rainfall (mm)	Topography
Antioquia	La Union (LU)	05°08'01.40"	75°23'40.40"	2,444	Silty	5.68	13.0	1,800	Strongly wavy
Antioquia	Rionegro (RN)	06°08'12.42"	75°25'06.70"	2,117	Silty	5.90	17.0	1,800	Flat
Antioquia	San Vicente (SV)	06°20'48.70"	75°16'14.10"	2,354	Clay loam	4.90	17.0	1,500	Wavy
Cundinamarca	Mosquera (MO)	04°41'50.65"	74°12'17.64"	2,550	Silty loam	6.19	12.9	700	Flat
Nariño	Ipiales (IP)	00°51'52.00"	77°35'10.00"	2,856	Sandy loam	5.30	10.2	743	Slight slope
Nariño	Pasto (PA)	01°12'06.00"	77°18'27.00"	2,760	Silty loam	5.53	11.8	870	Slight slope
Nariño	Puerres (PU)	00°54'58.00"	77°28'55.00"	2,612	Clay loam	5.70	12.5	798	Slight slope

The G×E interaction was analyzed using the AMMI method through SAS Enterprise software, version 8.3 (SAS Institute, 2020).

The AMMI model combines the ANOVA with principal component analysis (PCA):

$$Y_{ijk} = \mu + G_i + E_j + \sum \lambda_n \gamma_{in} \delta_{jn} + R_{ij} + \varepsilon_{ijk}$$

where Y_{ijk} is the observed value of the i^{th} genotype in the j^{th} environment and k^{th} repetition; μ is the grand

mean; G_i and E_j are the deviation of the i^{th} genotype and j^{th} environment from the grand mean, respectively; λ_n is a singular value of the n axis in the PCA; γ_{in} and δ_{jn} are genotype and environment factors, respectively, in the singular vectors associated with λ_n from the interaction matrix; R_{ij} is the part of the interaction that is explained by the PCA, and ε_{ijk} is the deviation of the i^{th} genotype in the j^{th} environment of the k^{th} repetition (Gauch, 1992).

RESULTS AND DISCUSSION

There were significant differences among genotypes, environments, and $G \times E$ interactions for fruit yield and fruit weight, indicating a differential response of the genotypes across the environments, therefore the possibility of selecting superior and stable genotypes. The environment explained 62.0 and 77.3% of the total sum of squares for fruit yield and fruit weight, respectively, whereas genotypes and $G \times E$ interactions each accounted for less than 9% of the variation (Tab. 3). The significantly higher contribution of the environments suggests that the environmental impact on fruit yield and fruit weight is more relevant than the main effect of each genotype, affecting the physiological processes of the plant, and in turn impacts yield (Sánchez *et al.* 2015). In line with this finding, Panayotov *et al.* (2016) also found a statistically significant contribution of environment to cape gooseberry yield, which was greater than the effect of genotypes and $G \times E$ interaction. Similarly, Rosero *et al.* (2021) found highly significant genotype-environment interaction in the evaluation of the combinatory ability of 36 cape gooseberry hybrids, obtained from nine inbred lines in four locations in the department of Nariño: Gualmatan, Puerres, Pasto, and Ipiales. Therefore, substantial variation is generated between genetic materials depending on the environment

in which they develop. Likewise, Gauch and Zobel (1997) reported in *Solanum tuberosum* that the main effect of environments could account for 80% of the total variation, whereas the effects of genotype and $G \times E$ interactions only accounted for 10%, as found in the present study. Ngailo *et al.* (2019) also reported great contribution of environments (64%) and $G \times E$ interactions (25%) for variation in storage root yield compared with the main effect of genotypes (11%) in sweet potato. In the same way, Ceballos-Aguirre *et al.* (2020) also reported a larger contribution of the environment (34.37-75.08%), compared to that of genotypes (1.24-11.83%) and $G \times E$ interaction (23.68-53.80%) for quality traits in cherry tomatoes.

The portioning of $G \times E$ components of fruit weight showed that the two first principal components (PC1 and PC2) of the interaction were significant and captured 50.9 and 19.2% of the variation, respectively. Nevertheless, the AMMI analysis for fruit yield indicated that only the first principal component of the $G \times E$ interaction was significant ($P < 0.005$), and PC1 accounted for more than 45% of the interaction. However, the eigenvalues of the first two principal components were greater than one, explaining the interaction in 67.5%. Thus, the analysis of AMMI with PC1 seems appropriate and can explain the effect of the interaction, as it has been shown in $G \times E$

Table 3. AMMI analysis of variance for fruit yield and fruit weight of 13 cape gooseberry genotypes over seven locations.

Source of variation	Fruit yield (FY)			Fruit weight (FW)		
	df	MS	%SS	df	MS	%SS
Genotype (G)	12	180.1***	6.3	12	2.58***	8.8
Environments (E)	6	3567.5***	62.0	6	45.42***	77.3
Block (Environment)	21	84.3	5.1	14	0.06	0.2
$G \times E$ interactions	72	35.6*	7.4	72	0.35***	7.1
PC1	17	68.1**	45.2	17	0.75***	50.9
PC2	15	38.1 ^{ns}	22.3	15	0.32**	19.2
Residual	40	20.8	32.5	40	0.19	29.9
Error	252	26.35	19.2	168	0.14	6.6
CV		20.7%			7.4%	

df: Degree of freedom; MS: Mean square; %SS: explained percentage of the sum of square; CV: coefficient of variation. Asterisks indicate significant differences at *5%; **0.5%; *** < 0.01%; ^{ns} not significant.

interaction studies in tomato (Ceballos-Aguirre *et al.*, 2020) and sesame (Movahedi *et al.*, 2020). There are no studies on cape gooseberry in this regard.

According to the unadjusted means, six genotypes had high yield in at least one environment, this number of genotypes is elevated, unwieldy, and impractical for planting recommendation in the producing areas. The methodology proposed by Crossa *et al.* (1991) and Gauch (1992; 2013) using the AMMI1 model reduce the number of winning genotypes and identifying mega-environments. The mega-environment is a subregion with similar genotypic responses and better performing cultivars (Yan *et al.*, 2023). The locations of this study were grouped in two mega-environments based on PC1 scores (Tab. 4), which were primarily defined by having similar climatic and soil conditions. Mega-environment 1 (ME1) consisted of Rionegro, Puerres, Ipiales, and Mosquera, with a mean of 24.1 t ha⁻¹, whereas mega-environment 2 (ME2) consisted of La Union, Pasto, and San Vicente, with a mean of 25.7 t ha⁻¹. These mega-environments presented similar conditions between some locations. The locations in ME1 are mainly at >2,500 m a.s.l. and had rainfall between 700 and 900 mm, except

Rionegro, and most had soils with high pH values (>5.7). On the other hand, ME2 locations are located below 2,500 m a.s.l., have soils with pH values below 5.7, and rainfall > 1,500 mm, except Pasto. Higher and well-distributed precipitation throughout the year compared to ME1, favored fruit production. Similar results have been reported by Fischer and Miranda (2012), who reported that well-distributed precipitation of 1,000 to 1,800 mm favors the vegetative and reproductive development of cape gooseberry, especially the fructification and guarantees high yields.

In warmer locations or when there are long periods of high temperature, the yields can be affected due to insufficient fruit set or increased fruit drop (Fischer and Orduz-Rodríguez, 2012), as observed in the locations of Rionegro and San Vicente of Antioquia department, where the temperature can range from 17 to 22°C. The locations of the Nariño department contributed to obtaining high yields, probably due to its natural soil conditions and microclimates suitable for the cape gooseberry production, specifically good availability of water resources, optimal temperatures, solar radiation, appropriate wind regime, and high fertility.

Table 4. AMMI1 estimates (top line) with their rank (in parenthesis) and unadjusted means (bottom line) for fruit yield (t/ha) of 13 cape gooseberry genotypes grown over seven locations.

Environment (E)	Mean E	Genotype (G)													
		A	B	C	D	E	F	G	H	I	N	R1	R2	R3	
ME1	RN	18.0 de 11.8*	9.2 (13) 11.8*	19.7 (7) 24.0	17.0 (9) 17.1	18.8 (8) 14.5	11.3 (11) 12.4*	20.5 (5) 20.9	24.1 (1) 22.9	15.2 (10) 11.6*	23.1 (3) 22.7	23.2 (2) 24.3	11.1 (12) 11.0*	20.6 (4) 20.7	19.9 (6) 20.6
	PU	25.9 bc 16.0	17.7 (13) 16.0	27.6 (6) 23.4	25.0 (9) 23.3	26.7 (8) 28.4	19.7 (12) 20.5	28.2 (4) 26.8	31.2 (1) 32.5	23.7 (10) 27.8	30.5 (3) 31.0	30.6 (2) 29.5	19.9 (11) 19.3	28.1 (5) 31.2	27.5 (7) 26.9
	IP	31.4 b 21.2*+	25.0 (13) 21.2*+	33.0 (4) 33.1	30.8 (10) 29.9	32.2 (8) 37.0	26.4 (12) 23.8*+	33.0 (5) 32.0	34.4 (3) 33.9	30.9 (9) 32.6	34.6 (1) 34.0	34.6 (2) 37.0	27.8 (11) 30.4	32.8 (6) 29.3	32.3 (7) 34.3
	MO	21.0 cd 17.1	15.4 (13) 17.1	22.6 (4) 21.9	20.6 (10) 24.6	21.8 (7) 22.9	16.5 (12) 17.2	22.2 (5) 23.3	23.0 (3) 24.8+	21.3 (9) 19.4	23.6 (1) 23.2	23.5 (2) 21.7	18.6 (11) 17.2	21.9 (6) 20.5	21.5 (8) 19.0
ME2	LU	13.6 e 11.9	9.4 (13) 11.9	15.2 (2) 14.0	13.4 (10) 14.1	14.4 (5) 12.5	10.1 (12) 8.8	14.3 (6) 18.2	13.8 (8) 13.2	15.3 (1) 14.9	15.1 (3) 18.0	14.9 (4) 12.3	13.1 (11) 12.5	13.8 (7) 14.8	13.6 (9) 11.2
	PA	38.2 a 34.5	35.2 (13) 34.5	39.7 (2) 40.9	38.3 (8) 38.5	39.0 (4) 38.5	35.6 (12) 34.7	38.4 (7) 37.0	36.9 (11) 35.8	41.1 (1) 41.4	38.7 (5) 38.1	38.5 (6) 38.4	39.4 (3) 39.4	37.8 (9) 36.9	37.7 (10) 42.4
	SV	25.4 c 22.5	23.0 (13) 22.5	26.9 (3) 27.4	25.6 (5) 23.4	26.2 (4) 25.5	23.1 (12) 25.3	25.4 (7) 23.7	23.4 (11) 24.1	28.8 (1) 28.9	25.5 (6) 24.4	25.2 (8) 27.2	27.4 (2) 27.7	24.7 (9) 26.8	24.6 (10) 22.4
Mean ME1	24.1	16.5	25.6	23.7	25.7	18.5	25.8	28.5	22.9	27.7	28.1	19.5	25.4	25.2	
Mean ME2	25.7	23.0	27.4	25.3	25.5	22.9	26.3	24.4	28.4	26.8	26.0	26.5	26.2	25.3	
Mean G	24.8	19.3*	26.4	24.4	25.6	20.4*	26.0	26.7	25.2	27.3	27.2	22.5*	25.7	25.3	

*Significant according to Dunnett's test at 5% level (control genotype N); +Significant according to Dunnett's test at 5% level (control genotypes R1, R2, and R3 of Cundinamarca, Antioquia, and Nariño departments, respectively). Locations with the same letter are not significantly different according to Tukey test. Locations: RN: Rionegro; PU: Puerres; IP: Ipiales; MO: Mosquera; LU: La Union; PA: Pasto; SV: San Vicente.

The fruit yield reported for most of the genotypes in this study ranged from 19.3 to 27.3 t ha⁻¹, being above the national commercial yield reported in 2020, with a mean of 12.1 t ha⁻¹ (Minagricultura, 2020). Genotypes I, N, G, B, D and H presented the best yields and were above the grand mean, whereas genotypes A and E had lower yield (Tab. 4). The N genotype or Colombia ecotype is cultivated throughout the country despite the lack of uniformity in cultivation and fruit traits; therefore, it was considered a national reference in the analysis.

The AMMI estimates and unadjusted fruit yield means showed different top-ranking genotypes in five of the seven locations tested. Based on unadjusted means, six of 13 genotypes (D, F, G, H, N, and R3) “won” in at least one environment; however, according to AMMI1 estimates, there were three winners:

genotypes G, H, and I (Tab. 4). The genotypes G and I were the best genotypes for ME1, and their performance was better compared to the three regional controls. The genotype H was the genotype with the highest performance for ME2 and superior to the three regional controls. Additionally, genotypes D and B presented the PC1 scores closest to zero, suggesting that they were the most stable genotypes (Fig. 1). However, only genotype B showed a mean yield higher than 26 t ha⁻¹; its yield was similar to that of Colombia ecotype (genotype N) in La Unión, Rionegro, San Vicente, and Mosquera and its yield was higher in Pasto and La Union. This high-yielding genotype was the most stable in all the locations evaluated, which indicates a broad adaptation. Thus, genotype B can be recommended for planting in the locations of the departments of Cundinamarca, Antioquia, and Nariño. The genotype B and genotypes

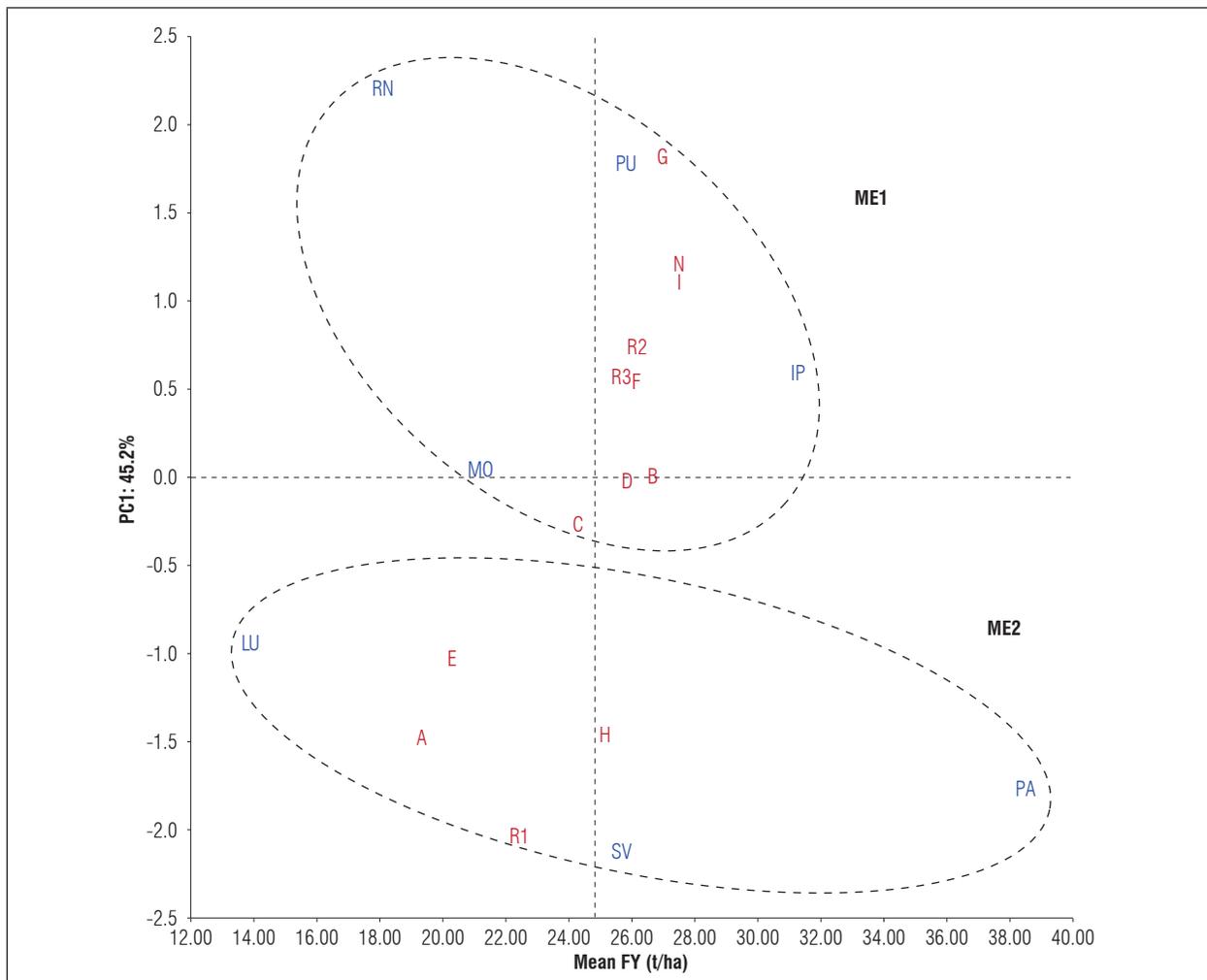


Figure 1. AMMI1 biplot fruit yield (FY) vs. PC1 of 13 cape gooseberry genotypes grown in seven locations.

G, I, and H could contribute significantly to the economic growth of the country's producing regions and maintain Colombia's position as the leading producer of cape gooseberry worldwide and, above all improve the quality of life of small farmers. Also, these could potentially be used in other countries with similar climatic conditions.

The most unstable genotypes were G, H, I, N, A, and R1. The genotypes G, H, I, and N, as mentioned above, exhibited high yield, whereas R1, A, and E were among the lowest yielding ones. The genotype R1, used as a regional control of the Cundinamarca department, presented low yields in ME1 and its mean yield was 19.5 t ha⁻¹. The regional control of Antioquia (genotype R2) and Nariño (genotype R3) had a lower performance than genotype N, except in Pasto. The Cundinamarca regional control (genotype

R1) showed low adaptation to Mosquera environmental conditions in comparison with the other genotypes, although this material has traditionally been cultivated in this department. The regional controls R2 and R3 had good yields, but these were lower than those of genotypes I, N, G, B, and H.

For fruit weight, the AMMI analysis identified only one mega-environment, but the Ipiales locality was far from the other locations that made up the mega-environment (Fig. 2). Ipiales locality showed the higher fruit weight followed by Pasto and Puerres of Nariño department, indicating that Nariño department is suitable environment to fruit weight. In contrast, Rionegro, La Union and San Vicente showed the smallest fruit weights and Mosquera exhibited fruits with intermediate weight. This suggest that locations with altitudes above 2,500 m a.s.l. and

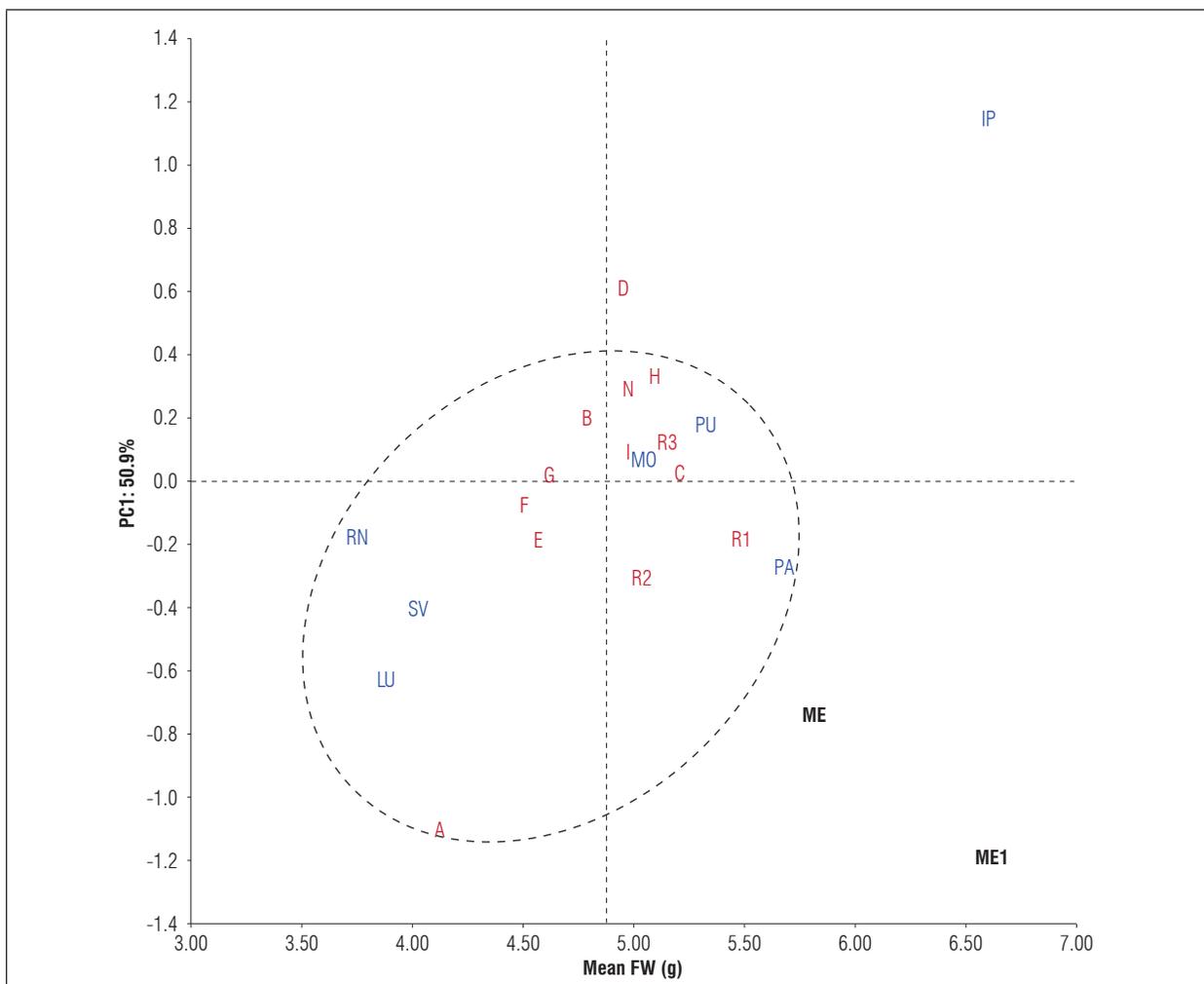


Figure 2. AMMI1 biplot fruit weight (FW) vs. PC1 of 13 cape gooseberry genotypes grown in seven locations.

temperatures between 12 and 15°C stimulated production of heavier fruits, as was the case in the locations of Nariño. Similar results were reported by Angulo (2005), who observed that fruits developed in low altitude regions were lighter than those from higher altitudes, directly affecting fruit yield. Likewise, Mazorra *et al.* (2003) found that to low altitude, the fruits of cape gooseberry were smaller than high altitudes.

In the same way that fruit yield, the selection of winner genotypes for fruit weight was realized based on AMMI estimates. According to this, two genotypes were the winners: genotypes D and R1 with 4.9 and 5.5 g (Tab. 5). The genotype D showed an adaptation for Ipiales while the genotype R1 for the mega-environment conformed by Puerres, Mosquera, Rionegro, Pasto, San Vicente, and La Union. However, the genotype R1 was one of the lowest fruit yielding genotypes with only 22.5 t ha⁻¹. The stable genotypes were G and C; while the unstable were D and A. The genotype A exhibited low fruit weight with 4.2 g, but genotype D exhibited a fruit weight higher of the average. The high yielding genotypes G, and B had a fruit weight below the average, suggesting that the fruit yield was a result of the number of fruits and not their weight. Conversely, the genotypes I,

and H with good fruit yield and adaptation to ME1 and ME2, respectively, showed a fruit weight above average and similar to national and regional controls. Similar results have been reported by Mendoza *et al.* (2012), who found an average of fruit weigh of 4.7 g in different farms of Cauca, demonstrating that the fruit satisfies with the national and international standards.

The fruit yield and fruit weight are critical traits to select genotypes for commercial plantation. Considering both traits, AMMI1 analysis revealed that genotype I can be recommended for planting in ME1 conformed by Rionegro, Puerres, Ipiales and Mosquera, genotype H for ME2 conformed by La Union, Pasto and San Vicente, and genotype B can be planting for both ME1 and ME2 due to stable performance in fruit yield, although it has a slightly smaller fruit that genotypes I and H. The locations from Nariño department were top environments with high yield and fruit weight of cape gooseberry.

CONCLUSION

Significant differences among genotypes, environments, and G×E interactions were found in this study. The AMMI1 analysis simplified and enable the

Table 5. AMMI1 estimates (top line) with their rank (in parenthesis) and unadjusted means (bottom line) for fruit weight (g) of 13 cape gooseberry genotypes grown over seven locations.

Environment (E)	Mean E	Genotype (G)													
		A	B	C	D	E	F	G	H	I	N	R1	R2	R3	
ME1	IP	6.6 a	4.6 (13) 4.6*	6.7 (8) 7.0	7.0 (6) 6.7	7.4 (1) 7.3	6.1 (12) 6.0	6.1 (11) 6.4	6.3 (10) 6.2	7.2 (2) 7.3	6.8 (7) 6.6	7.1 (3) 7.1	7.0 (4) 7.0	6.5 (9) 6.5	7.0 (5) 7.0
	PU	5.3 c	4.4 (13) 4.6	5.3 (9) 4.7	5.6 (2) 5.6	5.5 (7) 6.0+	5.0 (11) 5.3	4.9 (12) 4.7	5.0 (10) 5.5	5.6 (3) 5.3	5.5 (8) 5.5	5.5 (5) 5.4	5.9 (1) 6.0	5.5 (6) 5.0	5.6 (4) 5.5
	MO	5.0 d	4.3 (13) 4.1	5.0 (9) 4.6	5.4 (2) 5.8	5.1 (8) 4.9	4.7 (11) 4.8	4.7 (12) 4.4	4.8 (10) 4.6	5.3 (3) 5.5	5.2 (7) 5.8	5.2 (6) 4.9	5.6 (1) 5.3	5.2 (5) 5.5	5.3 (4) 5.3
	RN	3.7 f	3.2 (13) 3.0*	3.6 (9) 3.6	4.0 (2) 4.0	3.6 (8) 3.4	3.4 (10) 3.2	3.3 (12) 3.3	3.4 (11) 3.4	3.9 (5) 3.8	3.8 (6) 3.9	3.8 (7) 3.9	4.4 (1) 4.6	4.0 (3) 4.2	3.9 (4) 3.7
	PA	5.6 b	5.2 (13) 5.0	5.5 (8) 5.7	5.9 (2) 6.1	5.5 (9) 5.3	5.4 (10) 5.5	5.3 (12) 5.3	5.3 (11) 5.3	5.8 (5) 5.4	5.7 (6) 5.3	5.7 (7) 5.8	6.3 (1) 6.7	5.9 (3) 6.0	5.8 (4) 6.1
	SV	4.0 e	3.7 (11) 3.8	3.8 (8) 4.0	4.3 (3) 4.1	3.7 (10) 3.8	3.8 (9) 3.6	3.6 (13) 3.8	3.7 (12) 3.8	4.1 (5) 4.1	4.0 (6) 3.9	4.0 (7) 3.9	4.7 (1) 4.6*	4.3 (2) 4.1	4.1 (4) 4.2
	LU	3.9 ef	3.8 (7) 3.9	3.6 (10) 3.7	4.1 (3) 3.8	3.5 (13) 3.6	3.7 (9) 3.7	3.5 (12) 3.8	3.5 (11) 3.5	3.8 (6) 4.0	3.9 (5) 3.8	3.8 (8) 3.6	4.6 (1) 4.4*	4.2 (2) 4.3	4.0 (4) 3.9
Mean ME1	4.6	4.1	4.5	4.9	4.5	4.3	4.2	4.3	4.7	4.7	4.6	5.2	4.9	4.8	
Mean G	4.9	4.2	4.8	5.2	4.9	4.6	4.5	4.6	5.1	5.0	5.0	5.5*	5.1	4.9	

*Significant according to Dunnett's test at 5% level (control genotype N); +Significant according to Dunnett's test at 5% level (control genotypes R1, R2, and R3 of Cundinamarca, Antioquia, and Nariño departments, respectively). Locations with the same letter are not significantly different according to Tukey test. Locations: RN: Rionegro; PU: Puerres; IP: Ipiales; MO: Mosquera; LU: La Union; PA: Pasto; SV: San Vicente.

identification and selection of potential genotypes of cape gooseberry with high yield and fruit weight for two mega-environments with specific and abroad adaptation. Besides, the analysis identified suitable environments for cape gooseberry crops according to the evaluated traits.

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