

# Climate change impact on legume nutrition and nutritional quality

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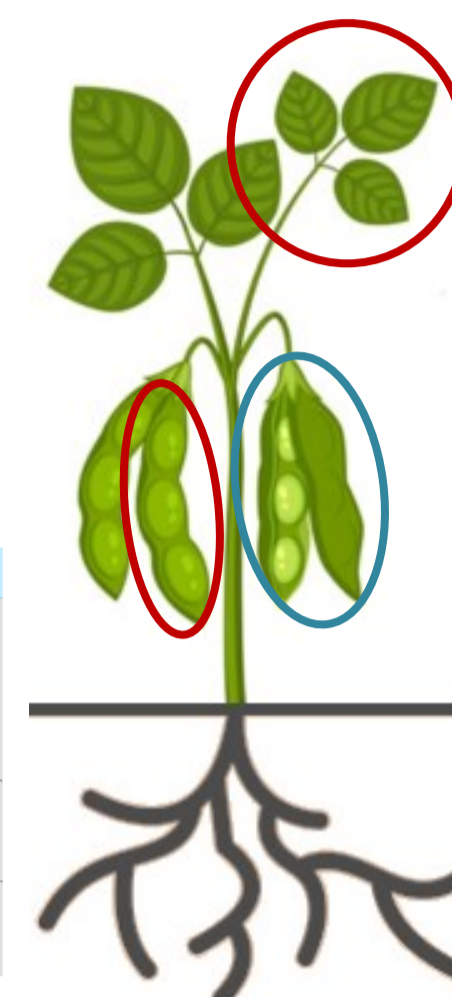
## Introduction

Climate change is a multilayered type of stress, with one of the focal points being the continuous increase of carbon dioxide (CO<sub>2</sub>) presence in the atmosphere, as it is expected to reach 550 parts per million (ppm) by 2050 (Soares et al., 2019). Therefore, understanding the impact of these elevated [CO<sub>2</sub>] (e[CO<sub>2</sub>]) in legumes, key contributors of essential nutrients for human health, iron (Fe) and zinc (Zn), is crucial. In that way, this work relied on free air CO<sub>2</sub> enriched (FACE) experiments, a unique platform to study the increasing levels of atmospheric CO<sub>2</sub> in plants, focusing three genotypes of common bean (*Phaseolus vulgaris* L.) (G1-G2-G3). The genotypes were grown permanently in ambient [CO<sub>2</sub>] (control, a[CO<sub>2</sub>]) or were exposed to a month of e[CO<sub>2</sub>] (600 ppm) during pod-filling stage, to unravel the effects underlying e[CO<sub>2</sub>] responses on biomass yield and nutritional value.

## Methods

BreedFACE  
Free Air CO<sub>2</sub>  
Enrichment

One month  
exposure



### 1. Biomass assessment (G1-G3)

Above ground biomass representative samples

### 2. Mineral analysis (G1-G3)

Mineral analysis of collected samples, was assessed using ICP-OES methodology (Santos et al., 2015)

### 3. Phenolic content and antioxidant capacity (G1/2)

Determined by Folin-Ciocalteu method (Ramos et al., 2019) and DPPH scavenging assay methodology (Gonçalves et al., 2009), respectively.

## Results and Discussion

### 1. Biomass

Genotype	Treatment	Overall biomass (g)				Seed analysis	
		Shoot	Grains	Empty pods	Seed weight (g)	Seed size (mm)	
G1	a[CO <sub>2</sub> ]	Average	106.67	30.41	28.52	0.21	56.42
		SD	36.82	4.67	2.49	0.09	8.83
	e[CO <sub>2</sub> ]	Average	146.67	64.37	48.50	0.28	63.84
		SD	9.43	32.33	17.20	0.10	10.75
% of change		27.27	52.77	41.20	25.00***	11.63***	
G2	a[CO <sub>2</sub> ]	Average	80.00	75.02	51.76	0.41	70.08
		SD	20.00	11.33	2.37	0.10	10.35
	e[CO <sub>2</sub> ]	Average	165.00	109.14	76.50	0.38	81.52
		SD	25.00	23.10	9.55	0.08	11.01
% of change		51.52*	31.26	32.34*	7.89	14.03**	
G3	a[CO <sub>2</sub> ]	Average	130.00	69.14	40.10	0.45	75.71
		SD	40.00	15.00	6.75	0.12	20.00
	e[CO <sub>2</sub> ]	Average	175.00	87.90	53.48	0.49	73.24
		SD	35.00	19.54	14.10	0.14	9.98
% of change		25.71	21.34	25.02	8.16	3.37	

\*\*\* = p<0.01; \*\* = p<0.05; \* = p<0.10

Figure 1- Average shoot, grains, empty pods, seed biomass (g), and average seed size (mm) of *P. vulgaris* genotypes.

Overall biomass and seed weight and size statistically significant increase at e[CO<sub>2</sub>]

- Differential genotypic responses: G1 and G2 could be more susceptible to the e[CO<sub>2</sub>] than G3

### 2. Mineral fluctuations

Mineral (µg g <sup>-1</sup> )	Genotype 1		Genotype 2		Genotype 3		Mineral (µg g <sup>-1</sup> )	Genotype 1		Genotype 2		Genotype 3	
	a[CO <sub>2</sub> ]	e[CO <sub>2</sub> ]	a[CO <sub>2</sub> ]	e[CO <sub>2</sub> ]	a[CO <sub>2</sub> ]	e[CO <sub>2</sub> ]		a[CO <sub>2</sub> ]	e[CO <sub>2</sub> ]	a[CO <sub>2</sub> ]	e[CO <sub>2</sub> ]	a[CO <sub>2</sub> ]	e[CO <sub>2</sub> ]
Mn	62 ± 8 <sup>a</sup>	51 ± 7 <sup>a</sup>	56 ± 2 <sup>a</sup>	81 ± 6 <sup>a</sup>	69 ± 1 <sup>a</sup>	56 ± 7 <sup>a</sup>	Mn	8 ± 1 <sup>a</sup>	7 ± 1 <sup>a</sup>	17 ± 2 <sup>a</sup>	19 ± 2 <sup>a</sup>	27 ± 7 <sup>a</sup>	23 ± 4 <sup>a</sup>
Zn	9 ± 1 <sup>a</sup>	4 ± 1 <sup>a</sup>	18 ± 1 <sup>a</sup>	20 ± 2 <sup>a</sup>	14 ± 1 <sup>a</sup>	18 ± 2 <sup>a</sup>	Zn	50 ± 2 <sup>a</sup>	25 ± 3 <sup>b</sup>	52 ± 5 <sup>a</sup>	59 ± 2 <sup>a</sup>	67 ± 9 <sup>a</sup>	44 ± 3 <sup>b</sup>
Fe	438 ± 18 <sup>a</sup>	889 ± 79 <sup>b</sup>	427 ± 87 <sup>a</sup>	922 ± 16 <sup>b</sup>	809 ± 151 <sup>a</sup>	513 ± 141 <sup>a</sup>	Fe	74 ± 5 <sup>a</sup>	31 ± 1 <sup>b</sup>	82 ± 10 <sup>a</sup>	107 ± 3 <sup>b</sup>	168 ± 3 <sup>a</sup>	120 ± 8 <sup>b</sup>
B	28 ± 5 <sup>a</sup>	19 ± 2 <sup>a</sup>	36 ± 2 <sup>a</sup>	26 ± 5 <sup>a</sup>	25 ± 1 <sup>a</sup>	26 ± 4 <sup>a</sup>	B	8 ± 1 <sup>a</sup>	7 ± 1 <sup>a</sup>	15 ± 1 <sup>a</sup>	18 ± 1 <sup>a</sup>	22 ± 1 <sup>a</sup>	19 ± 1 <sup>a</sup>
Cu	4 ± 0.4 <sup>a</sup>	5 ± 1 <sup>a</sup>	5 ± 0.4 <sup>a</sup>	6 ± 1 <sup>a</sup>	5 ± 0.1 <sup>a</sup>	5 ± 0.2 <sup>a</sup>	Cu	9 ± 1 <sup>a</sup>	8 ± 1 <sup>a</sup>	7 ± 1 <sup>a</sup>	10 ± 1 <sup>a</sup>	13 ± 0.2 <sup>a</sup>	12 ± 3 <sup>a</sup>
Ni	2 ± 0.2 <sup>a</sup>	1 ± 0.5 <sup>a</sup>	2 ± 0.3 <sup>a</sup>	4 ± 0.4 <sup>b</sup>	2 ± 0.8 <sup>a</sup>	3 ± 0.4 <sup>a</sup>	Ni	8 ± 2 <sup>a</sup>	5 ± 0.5 <sup>a</sup>	8 ± 0.6 <sup>a</sup>	12 ± 1 <sup>b</sup>	7 ± 0.1 <sup>a</sup>	7 ± 0.2 <sup>a</sup>
Mg	3066 ± 300 <sup>a</sup>	3749 ± 133 <sup>a</sup>	2567 ± 41 <sup>a</sup>	2620 ± 33 <sup>a</sup>	3197 ± 162 <sup>a</sup>	2573 ± 116 <sup>a</sup>	Mg	1159 ± 42 <sup>a</sup>	1216 ± 29 <sup>a</sup>	2559 ± 54 <sup>a</sup>	3133 ± 188 <sup>a</sup>	3322 ± 277 <sup>a</sup>	2769 ± 334 <sup>a</sup>
K	6674 ± 762 <sup>a</sup>	6095 ± 398 <sup>a</sup>	10854 ± 910 <sup>a</sup>	15602 ± 1258 <sup>b</sup>	9950 ± 407 <sup>a</sup>	13879 ± 1396 <sup>b</sup>	K	10375 ± 481 <sup>a</sup>	10524 ± 306 <sup>a</sup>	23053 ± 982 <sup>a</sup>	25174 ± 1089 <sup>a</sup>	29326 ± 1334 <sup>a</sup>	26126 ± 3972 <sup>a</sup>
Ca	46994 ± 6268 <sup>a</sup>	38004 ± 6141 <sup>a</sup>	36529 ± 3111 <sup>a</sup>	29774 ± 3290 <sup>a</sup>	33108 ± 3218 <sup>a</sup>	26853 ± 2026 <sup>a</sup>	Ca	976 ± 44 <sup>a</sup>	729 ± 71 <sup>a</sup>	1970 ± 276 <sup>a</sup>	2280 ± 378 <sup>a</sup>	4250 ± 671 <sup>a</sup>	4131 ± 686 <sup>a</sup>
P	1128 ± 104 <sup>a</sup>	1728 ± 268 <sup>a</sup>	1722 ± 75 <sup>a</sup>	1675 ± 91 <sup>a</sup>	1465 ± 160 <sup>a</sup>	1837 ± 198 <sup>a</sup>	P	3657 ± 272 <sup>a</sup>	3527 ± 166 <sup>a</sup>	5364 ± 85 <sup>a</sup>	6723 ± 518 <sup>b</sup>	7846 ± 21 <sup>a</sup>	5294 ± 792 <sup>b</sup>

Figure 2- Leaves and grains mineral accumulation, respectively, of *P. vulgaris* three genotypes (G1-G3) grown under a[CO<sub>2</sub>] and e[CO<sub>2</sub>] conditions.

## Conclusions

- Overall biomass increase in e[CO<sub>2</sub>] conditions, with different responses regarding the genotype.
  - G2 presenting the higher increase for the vegetative tissue
  - G1 the most yield-responsive biomass accumulation in grains
- Mineral fluctuations detected in all three genotypes
  - Suggests a response that may operate on a molecular level instead of biomass dilution-effect.
- Decreased phenolic content (G1 and G2) and antioxidant activity (G1), impacting their nutritional value.

These findings elucidate common bean, *P. vulgaris*, responses to a short-term e[CO<sub>2</sub>] exposure, providing a “root to grain” standpoint, while clarifying the importance of genotypic variability of this crop.

## Acknowledgements

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### 3. Phenolic compounds and antioxidant activity

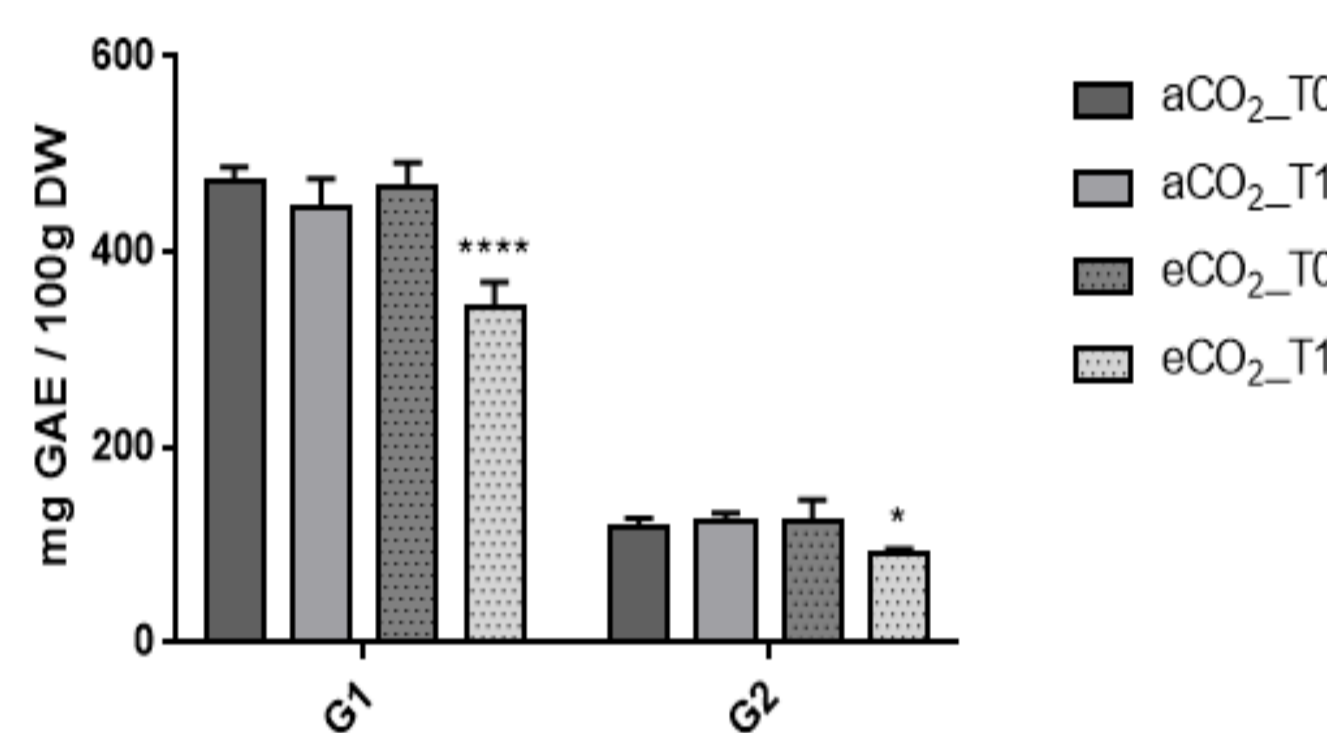


Figure 3- Comparison, through Folin-Ciocalteu method, of total phenolic of *P. vulgaris* grains, G1 and G2, grown at either a[CO<sub>2</sub>] and e[CO<sub>2</sub>] conditions

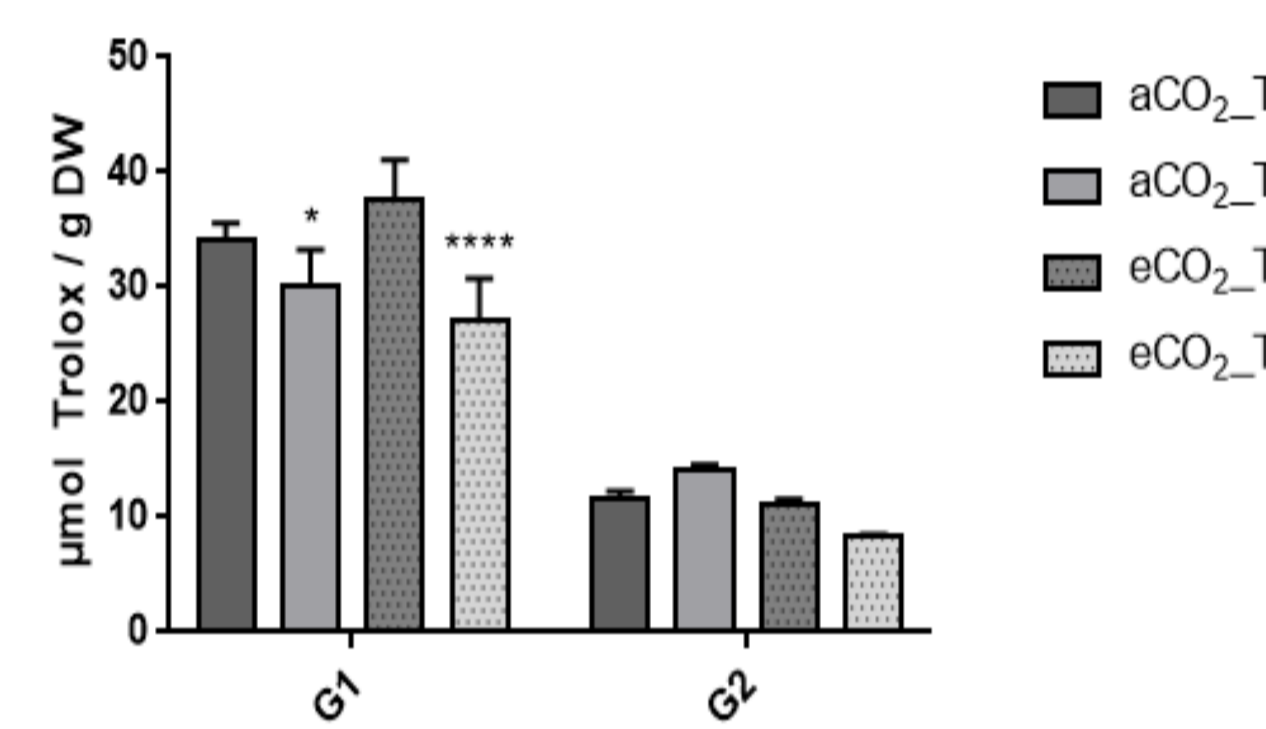


Figure 4- Comparison, through DPPH scavenging effect, of antioxidant activity of *P. vulgaris* grains, G1 and G2, grown at either a[CO<sub>2</sub>] and e[CO<sub>2</sub>] conditions

Significant reductions (27-28 %) in TPC of G1 and G2 under e[CO<sub>2</sub>] conditions.

G2 TPC clearly lower than G1.

28% reduction for G1 considering the DPPH assay under e[CO<sub>2</sub>] conditions.

Lower antioxidant activity is observed for G2.

Besides, the effect of e[CO<sub>2</sub>] exposure, it is again perceivable a **clear contrast between the genotypes tested.**

### Leaves

- Decrease in [Zn] concentration for G1
- Increase on [Fe] concentration for G1 and decreased in G3.

### Grains

- Decreased [Zn] and [Fe] for G1 and G3
- Increase in [Fe] for G2 was also reported.

Highly dependent on the genotype, mineral and tissue tested

## References

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