

Climate change impact on legume nutrition and nutritional quality

Rafael D.C. Duarte ^a, Juan Quirós-Vargas ^b, Onno Muller ^b, Marta W. Vasconcelos ^a

^a Universidade Católica Portuguesa, CBQF - Centro de Biotecnologia e Química Fina – Laboratório Associado, Escola Superior de Biotecnologia, Rua Diogo Botelho 1327, 4169-005 Porto, Portugal

^b Institute of Biogeosciences, IBG2: Plant Sciences, Forschungszentrum Jülich GmbH, 52425 Jülich, Germany

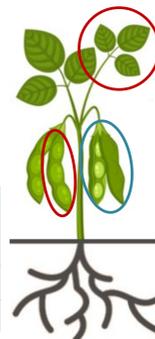
Introduction

Climate change is a multilayered type of stress, with one of the focal points being the continuous increase of carbon dioxide (CO₂) 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 [CO₂] (e[CO₂]) 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 CO₂ enriched (FACE) experiments, a unique platform to study the increasing levels of atmospheric CO₂ in plants, focusing three genotypes of common bean (*Phaseolus vulgaris* L.) (G1-G2-G3). The genotypes were grown permanently in ambient [CO₂] (control, a[CO₂]) or were exposed to a month of e[CO₂] (600 ppm) during pod-filling stage, to unravel the effects underlying e[CO₂] responses on biomass yield and nutritional value.

Methods

BreedFACE
Free Air CO₂
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 ₂] | Average | 106.67 | 30.41 | 28.52 | 0.21 | 56.42 |
| | | SD | 36.82 | 4.67 | 2.49 | 0.09 | 8.83 |
| | e[CO ₂] | 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 ₂] | Average | 80.00 | 75.02 | 51.76 | 0.41 | 70.08 |
| | | SD | 20.00 | 11.33 | 2.37 | 0.10 | 10.35 |
| | e[CO ₂] | 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 ₂] | Average | 130.00 | 69.14 | 40.10 | 0.45 | 75.71 |
| | | SD | 40.00 | 15.00 | 6.75 | 0.12 | 20.00 |
| | e[CO ₂] | 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[CO₂]

- Differential genotypic responses: G1 and G2 could be more susceptible to the e[CO₂] than G3

2. Mineral fluctuations

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

Figure 2- Leaves and grains mineral accumulation, respectively, of *P. vulgaris* three genotypes (G1-G3) grown under a[CO₂] and e[CO₂] conditions.

Conclusions

- Overall biomass increase in e[CO₂] 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[CO₂] 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

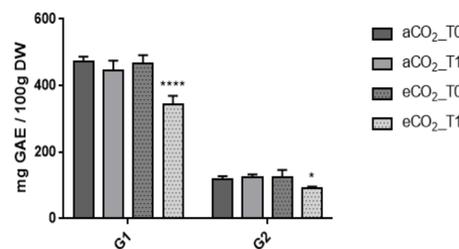


Figure 3- Comparison, through Folin-Ciocalteu method, of total phenolic of *P. vulgaris* grains, G1 and G2, grown at either a[CO₂] and e[CO₂] conditions

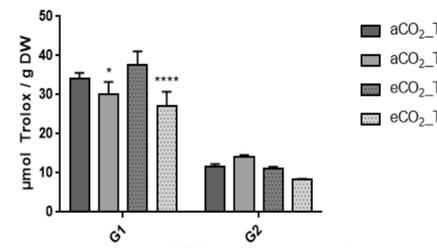


Figure 4- Comparison, through DPPH scavenging effect, of antioxidant activity of *P. vulgaris* grains, G1 and G2, grown at either a[CO₂] and e[CO₂] conditions

Significant reductions (27-28 %) in TPC of G1 and G2 under e[CO₂] conditions.

G2 TPC clearly lower than G1.

28% reduction for G1 considering the DPPH assay under e[CO₂] conditions.

Lower antioxidant activity is observed for G2.

Besides, the effect of e[CO₂] 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

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