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CO₂ enrichment can enhance the nutritional and health benefits of parsley (Petroselinum crispum L.) and dill (Anethum graveolens L.)

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eCO₂ improves the nutritional and health benefits of herbs

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Abstract

The functional food value of herbal plants is greatly related to their contents of valuable phytochemicals. Regarding its impact on primary and secondary plant metabolism, CO₂ enrichment could be a candidate strategy to modulate the levels of nutritionally and medicinally interesting phytochemicals in herbal plants. Herein, the concentrations of 98 metabolites and minerals were evaluated in shoot tissues of parsley and dill grown under two levels of CO₂, ambient (378±25 µmole CO₂ mole⁻¹ air, aCO₂) and elevated (627±24 µmole CO₂ mole⁻¹ air, eCO₂). Regardless of the plant species, eCO₂ improved the levels of soluble sugars, starch, organic acids, some EAAs, most of USFA, total phenolics, total flavonoids and vitamins A and E. However, notable variations in the metabolites responsiveness to eCO₂ were recorded among the tested plant species. Moreover, considerable improvements in the total antioxidant capacity, antiprotozoal, antibacterial and anticancer activities were recorded for parsley and dill in response to eCO2.

1. Introduction

Herbs are plants that are utilized both for their nutritional and health-promoting values (Liu, 2010; Opara & Chohan, 2014). Different parts of herbal plants are used in cooking to improve the flavor and aroma of foods (Peter, 2006). Moreover, herbs are used as natural alternatives to synthetic food preservatives (Badei, Faheid, El-Akel, & Mahmoud, 2002). Regarding their use in traditional medicine, herbs are well recognized for their antioxidant, antimicrobial, antiprotozoal, anti-inflammatory, antidiabetic and anticancer properties (Kurian, 2012). Generally, the nutritional and medicinal properties of medicinal herbs are ascribed to their content of minerals, vitamins, unsaturated fatty acids (USFA), essential amino acids (EAAs) and secondary metabolites such as phenolic acids and flavonoids (Bidlack, Omaye, Meskin, & Topham, 2000). Therefore, much attention has been devoted to improve the levels of these valuable phytochemicals in herbs and other medicinal plants. Indeed, plant tissue chemistry is greatly influenced by the growing conditions and agronomic practices (Björkman et al., 2011). For example, environmental stress and climate change have been reported to affect the accumulation of phytochemicals in plants (Oh, Trick, & Rajashekar, 2009). A similar effect is achieved by application of elicitors, which are substances or growth factors that can trigger metabolic events in plants (Baenas, García-Viguera, & Moreno, 2014). Under these circumstances, plants modulate their metabolism in order to accumulate specific compounds that help in coping with the imposed stress, climate change or elicitor. Although the accumulation of these bioactive compounds could improve nutritional and health-promoting values of plants, however it may be accompanied by reduced growth due to shift in allocation of resources, such as non-structural carbohydrates (NSC) and other intermediate compounds, to secondary metabolism (Matyssek et al., 2014). Therefore, improvement of carbon (C) metabolism could

help in adjusting the resource allocation balance between growth and synthesis of secondary metabolites by improving the availability of precursor compounds and metabolic energy (Ballare, 2009).

Elevated CO₂ (eCO₂) has been investigated as an inducer for growth and as a modulator of metabolism in crop and model plants (Azam, Khan, Mahmood, & Hameed, 2013; Noguchi, Watanabe, & Terashima, 2015). As a substrate of photosynthesis, elevated CO₂, upto a physiological limit, has been reported to enhance the photosynthetic C assimilation, therefore improve the accumulation of NSC and their breakdown via dark respiration (Leakey et al., 2009; Li et al., 2017). This can provide the required precursors and metabolic energy for the synthesis of the different classes of phytochemicals. Therefore, CO₂ fertilization can be used as a strategy to improve the nutritional and health-promoting values of herbal and medicinal plants. In this context, there are some evidences for the fertilization effect of eCO₂ treatment on the growth of several medicinal herbs such as Isatis indigotica (Li et al., 2017) and Hymenocallis littoralis (Idso et al., 2000). Moreover, some investigations have pointed to the positive impact of eCO₂ on the accumulation of bioactive phytochemicals in herbal and medicinal plants with a subsequent influence on their biological activities (Ghasemzadeh & Jaafar, 2011; Ghasemzadeh, Jaafar, & Rahmat, 2010; Ibrahim & Jaafar, 2011; Idso et al., 2000; Jaafar, Ibrahim, & Karimi, 2012; Wang, Bunce, & Maas, 2003). However, the current understanding for eCO₂-induced changes in the metabolism of herbal and medicinal plants is still fragmentary, as compared with crop or model plants. Accordingly, more efforts are needed to explore the global metabolic changes triggered by eCO₂ in this important group of plants and to assess the accompanying change in their nutritional and medicinal properties.

Among the commonly utilized culinary and medicinal herbs worldwide, both as fresh materials or essential oils, are the Umbelliferae members parsley (Petroselinum crispum L.) and dill (Anethum graveolens L.). Parsley and dill are valuable sources of minerals, vitamins, USFA, phenolic acids, flavonoids and other bioactive compounds that support their nutritional and medicinal benefits (Karkleliene et al., 2014; Lisiewska, Kmiecik, & Korus, 2006; Parry et al., 2006; Słupski, Lisiewska, & Kmiecik, 2005). They are recognized for their antioxidant, antibacterial, antifungal, antiinflammatory, antidiabetic and anticancer activities (Chahal, Kumar, Bhardwaj, & Kaur, 2017; Farshori, Al-Sheddi, Al-Oqail, & Siddiqui, 2013; Wong & Kitts, 2006). Growth, phytochemicals accumulation and oil composition of these two herbs are known to be influenced by several stress agents and conditions (Fraszczak, 2009; Hashem & Sahab, 1999; Petropoulos, Daferera, Polissiou, & Passam, 2009). However, to our knowledge, the impact of eCO₂ on the nutritional and health-promoting values of parsley and dill has not been investigated. Therefore, the current study was conducted to elucidate the impact of eCO₂ treatment (627±24 µmole CO₂ mole⁻¹ air) on the concentrations of metabolites (sugars, amino acids, fatty acids, vitamins, phenolic acids and flavonoids) and minerals in parsley and dill as well as the associated changes in their biological activities. Moreover, to explore the correlation between different classes of phytochemicals and to understand the global metabolic response of the two herbal plants to eCO₂, Hierarchical Clustering Analysis (HCA) was performed.

2. Materials and methods

2.1. Plant materials, growth microclimate and eCO₂ treatment

Parsley (Petroselinum crispum L.) and dill (Anethum graveolens L.) seeds, purchased from online supplier (https://www.seeds4garden.com/; codes SL2150 and SL2070, respectively), were sown in plastic pots filled with a mixture of loamy soil and organic compost (1:1, v/v). Pots were maintained in controlled-growth cabinet under 21/18 °C air temperature, 16/8 h day/night photoperiod and two mole fractions of atmospheric CO₂, ambient (378±25 µmole CO₂ mole⁻¹ air, aCO₂) and elevated (627±24 µmole CO₂ mole⁻¹ air, eCO₂). eCO₂ scenario was chosen according to the IPCC-SRES B2-scenario prediction of moderate change for the year 2100 (Murray & Ebi, 2012). Two independent growth cabinets were used for the two climate scenarios. For each treatment, twenty pots, six plants each, were distributed in the growth cabinet. All pots were irrigated to maintain soil humidity of 65%. After four weeks, some shoots were harvested, thoroughly washed with double distilled water, gently blotted, dried at 75 °C for 72 h and then used for dry mass determination and elemental analysis. The residual shoot material was immediately frozen in liquid nitrogen and stored at -20 °C until processing. In order to minimize any bias among the cabinets, the experiment had been replicated once again with swapping the two CO₂ levels among the cabinets. CO₂ was supplied in the airflow of the cabinet and its concentration was continuously monitored with a CO₂ analyser (WMA-4, PP Systems, Hitchin, UK).

2.2. Identification and quantification of individual metabolites and their totals

Chromatographic techniques, HPLC or GC/MS, were employed to determine the levels of individual sugars, organic acids, amino acids, fatty acids, phenolic acids, flavonoids and vitamins in plant samples (Hamad et al., 2015; Torras-Claveria, Berkov, Codina, Viladomat, & Bastida, 2014). Fatty acids were identified using NIST 05 database and Golm Metabolome

Database (http://gmd.mpimp-golm.mpg.de). The rest of compounds were identified by comparing their relative retention time with those of the standard mixture. The concentration of each compound was calculated based on peak area and comparison with a calibration curve of the corresponding standard. Nelson's, Folin–Ciocalteu and aluminum chloride colorimetric assays were employed to determine the total content of soluble sugars, total phenolics and flavonoids, respectively (Saleh, Madany, & González, 2015).

2.3. Determination of individual Minerals

For extraction of minerals, 200 mg of powdered-oven dried shoot materials were digested in 10 ml of 30% nitric acid for 48 h. The concentrations of macro and micro-elements were determined using inductively coupled plasma mass spectrometry (ICP-MS, Finnigan Element XR, Scientific, Bremen, Germany).

2.4. Biological activity assessments

Total antioxidant capacity was determined according to the commonly utilized ferric reducing antioxidant power (FRAP) and diphenylpicrylhydrazyl (DPPH) assays and expressed as µmole Trolox equivalent g⁻¹ dry weight or as % inhibition of DPPH radical, respectively. Antibacterial activity of the plant extracts was determined using the disc diffusion method (7.5 mg/disc) on Muller Hinton agar (MHA) and expressed as diameter of inhibition zone, mm (Selim, Aziz, Mashait, & Warrad, 2013). In vitro assessment of the antiprotozoal activity of the plant extracts (5 mg/ml) against *Trypanosoma cruzi* was conducted according to the protocol described by Räz, Iten, Grether-Bühler, Kaminsky and Brun (1997) and expressed as percentage reduction in parasite load relative to the control. Moreover, plant extracts (2 mg/ml) were tested for their anticancer activities against four human cell lines, hepatocellular carcinoma (HepG2), colon carcinoma (Colo205), embryonic kidney adenocarcinoma (293) and urinary bladder

carcinoma (T24P). Cell viability was determined by CellTiter-Blue reagent (Promega, Madison, WI, USA) as described by Solowey et al (2014) and the results were expressed as % dead cells.

2.5. Statistical analyses

Experiments were carried out following a completely randomized block design. Data analyses were performed using the Statistical Analysis System (SPSS Inc., Chicago, IL, USA). Data normality and the homogeneity of variances were checked using the Kolmogorov– Smirnov test and Levene's test, respectively. All the data were subjected to one-way analysis of variance (ANOVA). Tukey's Test ($p \le 0.05$) was conducted as the post hoc test for separation of means. Number of replicates for each experiment were three (n = 3). Cluster analysis was performed by using Pearson distance metric of the MultiExperiment Viewer (MeV)TM 4 software package (version 4.5, Dana-Farber Cancer Institute, Boston, MA, USA).

3. Results

3.1. Biomass, sugars, organic acids and EAAs are enhanced in response to eCO2

Elevated CO₂ treatment (627±24 μmole CO₂ mole⁻¹ air) resulted in significant improvements in the biomass production in parsley and dill, about 1.5 fold relative to aCO₂ conditions (378±25 μmole CO₂ mole⁻¹ air) (Figure 1). The accumulation of individual soluble and insoluble sugars and organic acids were assessed in both plants under the two CO₂ scenarios. Under aCO₂ conditions, parsley shoot tissues contained higher levels of glucose, fructose, sucrose, total soluble sugars, starch and total carbohydrates as compared to dill (Table 1). However, both plants contained comparable levels of organic acids, except for succinic acid that was significantly higher in parsley. eCO₂ treatment improved the accumulation of all sugar fractions in both plants, whereas glucose was the most accumulated soluble sugar in parsley

(0.68) and fructose was the most accumulated one in dill (0.61), as indicated by fold change calculations (Figure 2). Starch showed comparable fold changes (0.29) in both plants as affected by eCO₂. Regarding organic acids, citric, isobutyric, fumaric and malic acids showed similar response to eCO₂ in both plants, whereas the levels of citric and isobutyric were significantly increased and those for fumaric and malic were not significantly affected (Table 1). On the other hand, the level of succinic acid significantly improved in parsley but not affected in dill, and that for oxalic acid significantly reduced in dill but improved in parsley, as affected by eCO₂. On fold change basis, citric, isobutyric and succinic acids showed higher improvement in parsley than in dill, in response to eCO₂ (Figure 2).

Essential (EAAs) and non-essential amino acids (NEAAs) were quantified in shoots of parsley and dill grown under both aCO₂ and eCO₂ scenarios (Table 1). In both plants, lysine was the major EAAs followed by histidine, while glutamine was the abundant NEAAs followed by glutamate. The levels of all the EAAs and most of the detected NEAAs were higher in parsley as compared with their respective levels in dill. Moreover, parsley shoot contained about 4.75 and 1.13 fold amounts of the total essential and non-essential amino acids, respectively, as compared with dill. Under eCO₂, the levels of isoleucine, methionine, cystine, glutamine, ornithine and proline were significantly improved in both plants, when compared with their corresponding levels in aCO₂ plants. Some amino acids (leucine, lysine, phenylalanine and serine) were significantly induced in parsley but not dill, while others (valine and aspartate) were significantly induced in dill only, as affected by eCO₂. Moreover, total content of EAAs showed 0.51 and 0.10 fold increase in parsley and dill, respectively, in response to eCO₂ (Figure 2). The highest fold changes in amino acids levels as affected by eCO₂ were recorded for phenylalanine (1.36) and

lysine (0.66) in parsley and for methionine (1.50) aspartate (0.96), isoleucine (0.93) and cystine (0.90) in dill.

3.2. Levels and composition of fatty acids are altered by eCO₂ treatment

Parsley and dill contained comparable levels of most of the detected saturated fatty acids (SFA), while the contents of the majority of the detected USFA and their total were significantly higher in parsley (Table 1). Hexadecanoic (C16:0) was the dominant SFA, where it accounted for about 25 and 22% of the total SFA in parsley and dill, respectively. However, in both plants, octadecatrienoic and hexadecadienoic acids were the most abundant USFA. CO₂ enrichment did not significantly affect the accumulation of individual SFA or their total content in parsley, but significantly increased their levels in dill. On the other hand, eCO₂ treatment resulted in 0.24 and 0.45 fold increase in the total content of USFA in parsley and dill, respectively, (Figure 2). Such accumulation in total USFA was mainly due to accumulation of hexadecadienoic (C16:2), hexadecatrienoic (C16:3) and octadecatrienoic (C18:3) in parsley and most of the detected USFA in dill. The higher fold changes in the accumulation of USFA was observed for octadecatrienoic (C18:3) (0.60) in parsley and for hexadecanoic (C16:1) (0.91) in dill. Moreover, in both plants, eCO₂ resulted in lower SFA/USFA ratios as compared with aCO₂ conditions, whereas the lowest value (0.83) was detected in the treated parsley.

3.3. Levels of antioxidant metabolites as affected by eCO₂

Parsley and dill possessed similar phenolic acids, flavonoids and vitamins profiles with comparable levels of the detected compounds except for phylloquinone (vitamin K1) that was significantly higher in parsley (Table 2). In both plants, gallic acid, quercetin and α -carotene (vitamin A) were the most abundant phenolic acid, flavonoid and vitamin, respectively. eCO₂

treatment significantly increased the levels of most of the detected individual phenolic acids and flavonoids in parsley but not in dill. However, the total contents of flavonoids and phenolics were significantly enhanced in both plants as affected by eCO₂. Moreover, eCO₂ treatment resulted in significant increase in the levels of β -carotene, β -tocopherol and γ -tocopherol in parsley but not in dill and those for α -tocopherol and δ -tocopherol in dill only, meanwhile the β -cryptoxanthin and total tocopherol content was improved in both plants. On fold change basis, resorcinol was the most enhanced phenolic acid (15.17) in parsley and quercetrin was the most accumulated flavonoid (7.93) in dill, as affected by eCO₂ (Figure 2). Moreover, β -tocopherol and γ -tocopherol showed the highest fold changes (2.38 and 3.71, respectively) in parsley, while δ -tocopherol accounted for the highest fold change (0.64) in dill, in response to eCO₂ treatment.

3.4. Accumulation of minerals in parsley and dill is not affected by eCO₂

The levels of four macronutrients (K, Ca, Mg and P) and four micronutrients (Zn, Cu, Fe and Mn) were assessed in shoot tissues of parsley and dill under both aCO₂ and eCO₂ conditions. Parsley contained higher levels of K, Mg, Zn and Mn, however dill possessed higher values of Ca, P and Fe. The eCO₂ treatment did not exert a significant impact on concentrations of minerals in both plants, but for Ca and Zn there were significantly improvements in response to CO₂ enrichment.

3.5. Elevated CO₂ improves the biological activities of parsley and dill

Under aCO₂ conditions, parsley and dill showed comparable values of total antioxidant capacity, both FRAP and DPPH, antiprotozoal (*Trypanosoma cruzi*), anti-bacterial (*Escherichia coli* and *Streptococcus* sp.) and anticancer (HepG2 and Colo205) activities (Table2). On the other hand, parsley extract showed higher activities against embryonic kidney adenocarcinoma

(293) and urinary bladder carcinoma (T24P) than that of dill. CO₂ enrichment significantly improved all the measured biological activities for both plants, except for antihepatocellular carcinoma (HepG2) in dill. The fold change calculations revealed that the eCO₂-induced improvement in FRAP and DPPH scavenging activities was more obvious in dill than in parsley, however similar fold change was observed for antiprotozoal activity in both plants (Figure 2). Moreover, the highest fold change in antibacterial and anticancer activities was accounted for *Streptococcus* sp. (0.74) and HepG2 cell line (0.54), respectively, in parsley.

4. Discussion

The nutritional and medicinal values of herbs are strongly correlated with their content of minerals, vitamins, USFA, EAAs and secondary metabolites such as phenolic acids and flavonoids. In the current research, the impact of eCO₂ (627±24 µmole CO₂ mole⁻¹ air) on the concentrations of 98 metabolites and minerals in shoot of parsley and dill were evaluated as compared to their aCO₂ conditions. Hierarchical clustering analysis revealed that, under aCO₂ atmosphere, the two plants had qualitatively similar but somewhat quantitatively different metabolic profiles (Figure 3). Moreover, notable variations in the metabolites responsiveness to eCO₂ were recorded among the tested plant species.

CO₂ enrichment is known to improve the rate of photosynthesis by suppressing the oxygenation reaction of rubisco, the key enzyme in photosynthetic C fixation, where CO₂ and O₂ act competitively for this enzyme (Pérez-López et al., 2009). Accordingly, such upregulation of photosynthesis will improve the accumulation of sugars and also enhance its breakdown through dark respiration leading to accumulation of respiration intermediates such as organic acids (Li et al., 2013; Watanabe et al., 2014). Beside their diverse role in plant growth and metabolism, the significant accumulation of soluble sugars, starch and organic acids, such as citric and isobutyric, reported herein under eCO₂ treatment could improve the culinary value of parsley and dill as sugars and organic acids are determinants for taste and flavor (Malundo, Shewfelt, Ware, & Baldwin, 2001). Due to its antimicrobial properties, some organic acids are known to improve preservation of fruits and vegetables and to restrict the harmful microflora in human's digestive system (Nawirska-Olszanska, Biesiada, Soktowska, & Kucharska, 2014). Similar to our results, Moore, Palmquist, and Seemann (1997) had reported significant increases in the levels of sucrose and starch in parsley leaves subjected to 1000 μmole CO₂ mole⁻¹ air. Moreover,

accumulation of soluble sugars and starch in response to eCO₂ had been reported in crop plants such as tomato and wheat (Aranjuelo et al., 2011; Li et al., 2013). Organic acids such as fumarate and malate had also been reported to increase in the model plant *Arabidopsis thaliana* subjected to 780 µmole CO₂ mole⁻¹ air (Watanabe et al., 2014).

Among the factors that determine the nutritional value of plants are EAAs, specially lysine, tryptophan, and methionine that are constrained in cereals and legumes (Ufaz & Galili, 2008). Thus, from commercial and health value perspectives, it is important to enrich the content of EAAs in plant-based food. In the current study, the eCO₂-induced accumulation of isoleucine, leucine, lysine, methionine, phenylalanine, valine and their total in parsley and/or dill suggests a positive impact of eCO₂ on the nutritional value of herbal plants. In this regard, eCO₂ has been reported to affect the amino acid profile in model and crop plant species by improving, through its positive impact on the processes of photosynthesis and dark respiration, the availability of C skeleton and metabolic energy required for biosynthesis of amino acids (Noguchi et al., 2015; Nunes-Nesi, Fernie, & Stitt, 2010).

Fatty acid composition of plant-based foods is of a great health value importance where higher proportions of SFA are linked to cardiovascular diseases (Livingstone, Lovegrove, & Givens, 2012). By its positive impact on dark respiration, eCO₂ is expected to alter fatty acids profile as the precursor of their biosynthesis, malonyl-CoA, is produced from ATP-dependent carboxylation of acetyl-CoA that is an intermediate of the glycolytic pathway (Brown, Slabas, & Rafferty, 2010). Interestingly, the present results revealed that eCO₂ treatment resulted in higher levels of USFA accompanied with lower SFA/USFA ratios in both plants. Thus, parsley and dill produced in eCO₂ atmosphere are healthier than those grown under aCO₂. Unfortunately, little is known about the impact of eCO₂ on fatty acids metabolism, however, dissimilar to the present

results, two earlier investigations had reported a negative impact of eCO₂ on the biosynthesis of fatty acids in silver birch seedlings (Huang, Eglinton, Ineson, Bol, & Harkness, 1999) and root tubers of carrot, radish and turnip (Azam et al., 2013). Therefore, the impact of eCO₂ on the fatty acids biosynthesis and composition seems to be species dependent.

Biological activities of medicinal plants rely on the presence of bioactive phytochemicals, such as vitamins and secondary metabolites, that exert antioxidant, antibacterial, antiprotozoal and anticancer activities (Idso et al., 2002; Kumar, Tirpude, Maheshwari, Bansal, & Misra, 2013; Mohamed, Saleh, Abdel-Farid, & El-Naggar, 2017; Roleira et al., 2015; Sun et al., 2016). Moreover, the role of some micro-elements such as Cu, Zn, Se and Mn in free radical scavenging mechanisms has been reported (Fedor et al., 2017). In the current study, eCO₂ resulted in improved levels of total phenolics and flavonoids, β-Cryptoxanthin (vitamin A) and total tocopherol (vitamin E) in both parsley and dill, but mostly did not affect their mineral composition, except for Ca and Zn. Moreover, some individual flavonoids such as quercetrin and ellagic acid were accumulated in one plant species but not in the other. Such eCO₂-induced accumulation of these vitamins and secondary metabolites could be ascribed to the excess availability of C and N intermediates that are diverted for the biosynthesis of these bioactive phytochemicals (Herms & Mattson, 1992). In this regard, eCO₂ has been reported to affect C and N metabolism as well as nutrient uptake (Asif, Yilmaz, & Ozturk, 2017; Noguchi et al., 2015; Watanabe et al., 2014). As a logical consequence for its impact on the accumulation of bioactive compounds, eCO₂ treatment improved total antioxidant, antiprotozoal, anti-bacterial and anticancer activities of parsley and dill. Similarly, improved antioxidant activity accompanied with accumulation of anthocyanins, pelargonidin, quercetin and kaempferol, has been reported in fruits of strawberry treated with eCO₂ (Wang et al., 2003). Moreover, Ghasemzadeh et al. (2010)

had ascribed the eCO₂-induced improvement in the antioxidant capacity of ginger to the enhanced levels of quercetin, kaempferol, catechin and fisetin. Not only flavonoids, but also some phenolic acids (gallic and caffeic acids) and vitamin C (ascorbic acid) were reported to accumulate in *Labisia pumila* and *Citrus aurantium*, respectively, in response to eCO₂ (Idso et al., 2002; Jaafar et al., 2012). In addition, a similar improvement in the antioxidant and anticancer, against human cell lines MCF–7 and MDA–MB–231, activities has been observed in two varieties of ginger grown under 800 μmol/mol CO₂ (Ghasemzadeh & Jaafar, 2011).

5. Conclusion

Based on the results presented in this research it could be concluded that manipulation of eCO₂ as a strategy to improve the growth and the nutritional and health benefits of herbal plants is worthwhile. eCO₂ (627±24 µmole CO₂ mole ¹ air) improved the accumulation of nutritionally and medicinally valuable phytochemicals in both parsley and dill. However, there was obvious difference between the two plants in the degree by which their individual metabolites respond to eCO₂ treatment. Regardless of the plant species, eCO₂ triggered improvements in the levels of soluble sugars, starch, organic acids, some EAAs, most of USFA, total phenolics, total flavonoids and vitamins A and E. Moreover, parallel improvements in the total antioxidant capacity, antiprotozoal, antibacterial and anticancer activities were recorded for parsley and dill in response to eCO₂.

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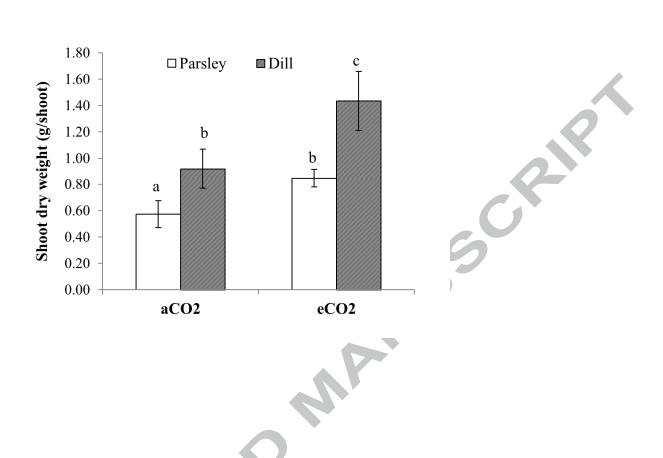
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Figure Captions

- **Figure 1.** Shoot biomass of parsley and dill grown under two levels of CO_2 , ambient (378±25 μ mole CO_2 mole⁻¹ air, a CO_2) and elevated (627±24 μ mole CO_2 mole⁻¹ air, e CO_2). Values are mean \pm standard error of three independent replicates. Same lower-case letters on bars indicate no significant difference at the 0.05 probability level.
- **Figure 2.** Heatmap of fold change ratios in metabolite, vitamins, minerals and biological activities of parsley and dill in response to eCO_2 (627 \pm 24 μ mole CO_2 mole⁻¹ air). Fold change in each metabolite was calculated relative to its corresponding mean (n=3) in the aCO₂ (378 \pm 25 μ mole CO_2 mole⁻¹ air) grown plant. As shown in the color scale, red indicates inhibition, white no change and blue improvement in metabolite levels as affected by eCO₂.

Figure 3. Heatmap of metabolite accumulation in shoot tissues of parsley and dill grown under two levels of CO₂, ambient (aCO₂, 378±25 µmole CO₂ mole⁻¹ air) and elevated (eCO₂, 627±24 µmole CO₂ mole⁻¹ air). The relative accumulation patterns are shown in the heatmap based on the average value (n=3) for each metabolite. Red and blue colors indicate lower and higher ACCEPTED MARINGSCRI concentrations, respectively.



		Parsley Dill		Parsley Dill		Parsley Dill		Parsley Dill
	Sugars				Phenolic acids			
	Glucose	0.69 0.56	Ornithine	0.47 0.60	Caffeic acid	0.33 0.33	γ-tocopherol	3.71 0.06
	Fructose	0.48 0.61	Proline	0.43 0.33	Ferulic acid	0.55 0.34	δ-tocopherol	-0.08 0.64
	Sucrose Total soluble sugars	0.44 0.33 0.49 0.82	Serine Tyrosine	0.28 0.26 0.23 -0.06	Protocatechuic acid Catechin	0.42 0.29 0.54 0.26	Total tocopherol Vitamin K1	1.35 0.25
	Starch	0.29 0.29	alanine	0.12 0.05	Gallic acid	0.58 0.03	Phylloquinone	-0.38 0.18
	Total carbohydrate	0.51 0.60	Asparagine	0.40 0.04	p-Coumaric acid	0.52 -0.19	Macro-elements	0.00
	Organic acids		Total NEAAs	0.24 0.15	Resorcinol	15.17 0.17	K	0.04 -0.08
	Citric acid	0.45 0.27	Saturated Fatty acids (S		Chlorogenic acid	0.44 -0.05	Ca	0.88 0.61
	Fumaric acid	-0.04 0.13	Tetradecanoic (C14:0)	-0.01 0.44	Syringic acid	0.61 0.23	Mg	0.38 -0.12
	Isobutyric acid	0.45 0.28	Hexadecanoic (C16:0)	-0.14 0.23	Flavonoids		Р	0.04 -0.06
	Malic acid	0.11 0.01	Heptadecanoic (C17:0)	-0.01 0.31	Quercetin	0.67 0.16	Micro-elements	
	Oxalic acid	0.31 -0.21	Octadecanoic (C18:0)	-0.02 0.39	Quercetrin	0.42 7.93	Zn	1.18 0.57
	Succinic acid	0.45 0.14	Eicosanoic (C20:0)	-0.11 0.42	Luteolin	0.63 0.00	Cu	0.00 1.00
	Essential amino acid		Docosanoic (C22:0)	0.39 0.04	Apigenin	0.68 -0.06	Fe	0.33 -0.03
	Histidine Isoleucine	0.06 0.08 0.28 0.93	Tricosanoic (C23:0) Tetracosanoic (C24:0)	-0.09 0.41 0.18 0.47	lsoquercetrin Rutin	0.28 0.07 0.92 0.08	Mn Total antioxidant a	0.07 0.19
	Leucine	0.43 0.00	Pentacosanoic (C25:0)	0.00 0.31	Ellagic acid	1.23 0.02	FRAP	0.46 0.59
	Lysine	0.66 0.09	Total SFA	0.01 0.32	Velutin	0.53 0.04	DPPH	0.37 0.60
	Methionine	0.50 1.50	Unsaturated fatty acids (Total flavonods	0.37 0.60	Antiprotozoal	0.07
	Phenylalanine	1.36 -0.08	Hexadecanoic (C16:1)	-0.12 0.91	Total Phenols	0.96 0.32	Trypanosom a cruzi	0.53 0.58
	Valine	-0.03 0.45	Octadecenoic (C18:1)	0.07 0.43	Vitamin A		Antibacterial	
	Threonine	-0.02 0.10	Tetracosenoic (C24:1)	0.02 0.87	α-Carotene	-0.27 -0.10	Streptococcus sp.	0.74 0.24
	Total EAAs	0.51 0.10	Hexadecadienoic (C16:2)	0.31 0.52	β-Carotene	0.56 -0.13	Escherichia coli	0.32 0.42
	Non-essential amino		Octadecadienoic (C18:2)	-0.01 0.37	β-Cryptoxanthin	0.58 0.00	Anticancer	0.54
	Aspartate Cystine	0.38 0.96	Eicosadienoic (C20:2) Hexadecatrienoic (C16:3)	-0.02 0.44	Vitamin B	0.26 0.40	HepG2	0.54 0.00
	Glutamlic acid	0.33 0.90 0.44 0.32	Octadecatrienoic (C18:3)	0.23 0.03 0.60 0.05	Thiamine Vitamin E	-0.36 -0.10	Colo205 293	0.20 0.17 0.06 0.16
	Glutamine	0.19 0.00	Total USFA	0.24 0.45	α-tocopherol	-0.07 -0.14	T24P	0.13 0.28
	Glycine	0.13 -0.07	10101 001 71	0.24	β-tocopherol	2.38 0.06		0.10
			≤-1.5	0.0		≥1.5		
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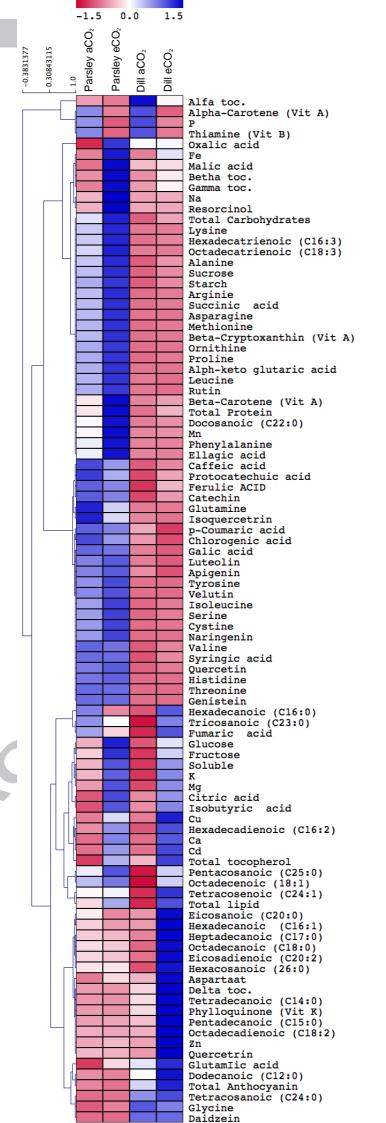


Table 1. Levels of sugars, organic acids (mg g⁻¹ dry weight), amino acids and fatty acids (μg g⁻¹ dry weight) in shoot tissues of parsley and dill grown under two concentrations of CO₂, ambient (378±25 μmole CO₂ mole⁻¹ air, aCO₂) and elevated (627±24 μmole CO₂ mole⁻¹ air, eCO₂). Values are mean ± standard error of three independent replicates. Means followed by the same lower-case letter in a row do not differ significantly at the 0.05 probability level.

_	aCO2		eCO2		_	aCO2		eCO2	
Metabolite	Parsley	Dill	Parsley	Dill	Metabolite	Parsley	Dill	Parsley	Dill
Sugars									
Glucose	$0.62 \pm 0.01 b$	$0.50\pm0.00a$	1.05±0.02d	$0.78\pm0.00c$	Asparagine	4.34±0.78b	1.05±0.11a	6.06±1.17b	1.09±0.06a
Fructose	$0.81 \pm 0.05 b$	$0.59\pm0.02a$	1.20±0.08c	$0.95 \pm 0.03 b$	Glycine	0.64±0.06a	1.41±0.24b	$0.72 \pm 0.03a$	1.31±0.17b
Sucrose	$0.77 \pm 0.03c$	$0.27 \pm 0.00a$	1.11±0.04d	$0.36 \pm 0.00 b$	Ornithine	1.05±0.10c	$0.05\pm0.01a$	1.54±0.15d	$0.08 \pm 0.01 b$
Total soluble sugars	1.74±0.18b	$1.30\pm0.10a$	2.59±0.27c	2.36±0.18c	Proline	3.53±0.37c	$0.58 \pm 0.06a$	$5.04 \pm 0.53 d$	$0.77 \pm 0.08 b$
Starch	12.28±0.45c	4.81±0.02a	15.80±0.58d	6.21±0.02b	Serine	$0.83\pm0.08b$	$0.19\pm0.02a$	1.06±0.11c	$0.24 \pm 0.02a$
Total sugars	24.58±1.02c	10.73±0.14a	37.13±1.55d	17.19±0.23b	Tyrosine	4.94±4.94b	$0.81 \pm 0.12a$	6.10±0.69b	0.76±0.13a
Organic acids					Total NEAAs	267.48±13.4a	236.52±13.1a	331.34±12.6b	271.71±17.1a
Citric acid	1.98±0.10a	2.11±0.11a	2.87±0.15b	2.68±0.14b	Saturated Fatty acids (SI	FA)			
Fumaric acid	0.25±0.01a	$0.23\pm0.01a$	$0.24 \pm 0.02a$	$0.26\pm0.02a$	Tetradecanoic (C14:0)	112.0±3.7a	125.7±4.6a	111.3±2.8a	181.0±7.3b
Isobutyric acid	$1.70\pm0.09a$	1.83±0.09a	2.46±0.13b	2.35±0.12b	Hexadecanoic (C16:0)	319.3±37.3ab	264.9±17.6a	273.6±32.5ab	326.7±15.2b
Malic acid	$7.39 \pm 0.45 a$	$7.54 \pm 0.46a$	8.17±0.37a	7.60±0.29a	Heptadecanoic (C17:0)	245.9±15.4a	231.3±6.3a	242.4±5.0a	303.2±22.9b
Oxalic acid	1.29±0.08ab	1.50±0.08b	1.69±0.10c	1.19±0.03a	Octadecanoic (C18:0)	37.1±1.7a	33.2±1.6a	36.5±1.8a	46.3±1.9b
Succinic acid	3.99±0.24b	$1.24 \pm 0.06a$	5.79±0.35c	1.41±0.07a	Eicosanoic (C20:0)	134.7±8.7a	121.9±8.2a	119.9±4.0a	173.6±17.7b
Essential amino aci	ds (EAAs)				Docosanoic (C22:0)	72.1±1.9b	55.5±0.5a	100.4±4.0c	57.9±3.5a
Histidine	10.21±1.20b	1.61±0.25a	10.80±1.18b	1.74±0.10a	Tricosanoic (C23:0)	8.2±0.3b	5.9±0.5a	7.5±0.3ab	8.3±0.7b
Isoleucine	$0.67 \pm 0.07 b$	$0.15 \pm 0.01a$	0.86±0.09c	0.29±0.02b	Tetracosanoic (C24:0)	228.9±0.7b	160.8±13.5a	270.0±47.4b	236.4±17.7b
Leucine	$0.14\pm0.01b$	$0.01 \pm 0.00a$	0.20±0.02c	0.01±0.00a	Pentacosanoic (C25:0)	22.4±0.7a	23.2±1.7a	22.5±2.0a	30.4±1.2b
Lysine	23.14±0.22b	5.03±0.53a	38.44±4.52c	5.46±0.37a	Total SFA	1270.2±35.6a	1186.1±116.1a	1278.1±56.3a	1562.6±150.2b
Methionine	$0.12\pm0.01b$	$0.02\pm0.00a$	0.18±0.02c	$0.05 \pm 0.01 b$	Unsaturated fatty acids (USFA)			
Phenylalanine	4.79±0.54b	0.53±0.08a	11.32±1.26c	$0.49\pm0.02a$	Hexadecanoic (C16:1)	121.0±0.75a	105.9±2.3a	107.0±1.0a	201.9±19.0b
Valine	5.09±0.44b	1.94±0.20a	4.95±0.43b	2.82±0.35b	Octadecenoic (C18:1)	41.6±1.80b	28.7±1.7a	44.4±4.6b	40.9±1.6b
Threonine	$0.48 \pm 0.01 b$	0.10±0.01a	0.47±0.04b	0.11±0.01a	Tetracosenoic (C24:1)	146.5±0.43b	102.4±0.7a	150.1±10.0b	191.7±7.7c
Total EAAs	44.62±1.31b	9.40±0.61a	67.22±5.8c	10.35±0.51a	Hexadecadienoic (C16:2)	279.4±50.5ab	257.9±31.8a	366.2±21.7c	392.4±26.1c
Non-essential amine	o acids (NEAA	As)			Octadecadienoic (C18:2)	16.3±0.18a	16.3±0.2a	16.2±1.9a	22.3±0.9b

Aspartate	0.21±0.02a	0.26±0.07a	$0.29\pm0.03a$	0.51±0.03b	Eicosadienoic (C20:2)	153.4x12.07b	126.1±13.9a	150.4±6.0b	181.4±8.1c
Cystine	4.07±0.41c	0.20±0.01a	$5.43 \pm 0.55 d$	$0.38\pm0.03b$	Hexadecatrienoic (C16:3)	164.1±3.70b	117.7±7.0a	201.8±8.1c	121.77±9.4a
GlutamIic acid	65.31±4.0a	106.1±16.5b	94.23±9.03b	139.6±16.55	Octadecatrienoic (C18:3)	310.3±11.47b	143.9±20.5a	495.1±19.9c	150.5±14.0a
Glutamine	150.4±4.8c	99.2±2.52a	178.44±5.10d	98.89±5.85b	Total USFA	1232.7±42.2b	898.9±33.a	1530.9±48.2c	1302.9±32.6b
Alanine	32.16±5.34a	26.67±5.11a	36.03±4.09a	28.08±7.30a	SFA/USFA	1.03	1.32	0.83	1.20

Table 2. Concentrations of individual phenolic acids, flavonoids, their totals, vitamins, minerals (mg g⁻¹ dry weight) and biological activities of parsley and dill grown under two concentrations of CO₂, ambient (378±25 μmole CO₂ mole⁻¹ air, aCO₂) and elevated (627±24 μmole CO₂ mole⁻¹ air, eCO₂). Values are mean ± standard error of three independent replicates. Means followed by the same lower-case letter in a row do not differ significantly at the 0.05 probability level.

	aCO2		eCO2			aCO2		eCC)2
Metabolite	Parsley	Dill	Parsley	Dill	Metabolite	Parsley	Dill	Parsley	Dill
Phenolic acids									
Caffeic acid	$0.03\pm0.00a$	$0.03\pm0.00a$	$0.04 \pm 0.00 b$	$0.04\pm0.00a$	γ-tocopherol	$0.024\pm0.00a$	0.031 ± 0.00	0.113±0.01	0.033 ± 0.00
Ferulic acid	3.26±0.46a	3.58±0.39a	$5.04 \pm 0.50 b$	4.79±0.47a	δ -tocopherol	$0.012 \pm 0.00a$	0.022 ± 0.01	0.011 ± 0.00	0.036 ± 0.00
Protocatechuic acid	$0.26\pm0.04a$	$0.28\pm0.03a$	$0.37 \pm 0.04a$	0.36±0.04a	Total tocopherol	0.211±0.09a	0.338 ± 0.06	0.496 ± 0.02	0.421 ± 0.02
Catechin	$0.98 \pm 0.14a$	1.06±0.12a	1.51±0.15b	1.34±0.13a	Vitamin K1				
Gallic acid	25.99±3.70a	28.02±3.0a	40.94±4.08b	28.73±2.83	Phylloquinone	0.121±0.01a	0.496 ± 0.04	0.075 ± 0.03	0.585 ± 0.00
p-Coumaric acid	5.59±0.38a	6.02±0.66a	8.51±0.85b	4.87±0.48a	Macronutrients				
Resorcinol	0.06±0.01a	0.06±0.01a	0.97±0.86b	0.07±0.01a	K	31.59±1.22a	29.20±2.30	32.7±1.30a	26.88±1.12
Chlorogenic acid	$0.34 \pm 0.05a$	$0.37 \pm 0.04a$	0.49±0.05a	0.35±0.03a	Ca	7.75±0.29a	11.94±0.80	14.25±0.52	19.17±0.95
Syringic acid	1.53±0.22a	1.65±0.18a	2.46±0.24b	2.03±0.20a	Mg	4.94±0.52b	2.93±0.26a	6.81±0.53b	2.57±0.06a
Flavonoids					P	5.69±0.22a	$7.40 \pm 0.58a$	$5.89 \pm 0.34a$	$6.97 \pm 0.48a$
Quercetin	2.30±0.33a	$2.48 \pm 0.27a$	3.85±0.38b	2.88±0.28a	Micronutrients				
Quercetrin	$0.26\pm0.04a$	0.28±0.03a	0.37±0.04a	2.50±0.25b	Zn	$0.50\pm0.09b$	$0.21 \pm 0.05a$	1.09±0.17c	$0.33 \pm 0.02b$
Luteolin	$0.08\pm0.01a$	0.08±0.01a	0.13±0.01b	$0.08 \pm 0.01a$	Cu	$0.01 \pm 0.00a$	$0.01 \pm 0.00a$	$0.01 \pm 0.00a$	$0.02\pm0.00a$
Apigenin	$0.50\pm0.07a$	0.53±0.06a	0.84±0.08b	$0.50\pm0.05a$	Fe	0.21±0.01a	$0.34 \pm 0.03a$	$0.28 \pm 0.02a$	$0.33 \pm 0.03a$
Isoquercetrin	1.24±0.18a	1.34±0.15a	1.59±0.16a	$1.44 \pm 0.14a$	Mn	$0.042 \pm 0.00 b$	0.026±0.00	0.045 ± 0.00	0.031 ± 0.00
Rutin	1.61±0.23a	1.74±0.19a	3.09±0.31b	1.88±0.19a	Total antioxidant	activity (FRAP	, μmole Trolox	g ⁻¹ DW; DPP	Н, %

Ellagic acid Velutin	0.40±0.06a 0.47±0.07a	0.40±0.02a 0.47±0.02a	0.89±0.04b 0.72±0.03b	0.41±0.02a 0.49±0.02a	FRAP DPPH	4.01±0.04a 18.56±0.16a	4.20±0.67a 21.56±3.56	5.85±0.07b 25.41±0.33	6.67±0.44b 34.53±3.24
Total flavonods	$72.75 \pm 0.67a$	64.74±2.33	99.58±2.78b	103.49±6.0	Antiprotozoal (%			23.1120.33	31.33±3.21
Total Phenols	55.02±1.29a	53.10±4.89	107.70±4.40	70.29±2.52	Trypanosoma	21.56±0.20a	20.41±3.04	33.07±0.73	32.18±4.34
Vitamin A	20102211294				Antibacterial (dia				
α-Carotene	0.473±0.03a	0.505 ± 0.03	0.346±0.07a	0.452 ± 0.03	Streptococcus sp.	6.17±0.26a	12.22±1.17	10.74±0.14	15.17±1.65
β-Carotene	0.393±0.02b	0.313±0.03a	0.613±0.04c	0.271±0.00	Escherichia coli	18.22±1.48a	17.69±1.24	23.99±0.93	25.08±1.45
β-Cryptoxanthin	0.310±0.02a	0.026 ± 0.00	0.490±0.22b	0.026±0.00	Anticancer (% dea				
Vitamin B					HepG2	47.34±1.11a	56.38±6.82	72.92±4.12	56.53±1.91
Thiamine	$0.348 \pm 0.02a$	0.373 ± 0.03	0.221±0.09a	0.335 ± 0.01	Colo205	66.23±0.61a	60.75±3.00	80.06±2.24	70.93±1.75
Vitamin E					293	73.69±0.65c	53.97±2.59	77.99±1.01	62.56±1.15
α -tocopherol	$0.150\pm0.01a$	0.264 ± 0.10	0.139±0.01a	0.228±0.09	T24P	76.96±0.73b	55.15±4.49	87.21±2.69	70.40±3.90
β -tocopherol	0.032±0.00a	0.032±0.00	0.108±0.02b	0.034±0.00					
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Highlights

- CO₂ enrichment improves the levels of flavor-related compounds such as sugars and organic acids
- Elevated CO₂ enhances the nutritive value through higher EAAs content and lower SFA/USFA ratio
- ullet Levels of antioxidant metabolites and antioxidant activities are improved by elevated CO_2
- CO₂ enrichment supports the value of parsley and dill as functional foods