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Diffuse reflectance infrared Fourier transform spectroscopy for a qualitative evaluation of plant leaf pigment extraction

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The extraction and quantification of leaf pigments are easy, fast, and cheap procedures; on the other hand, diffuse reflectance infrared Fourier transform (DRIFT) spectroscopy associated with chemometrics tools could offer new insights into leaf biochemical composition. We aimed to boost the classic leaf pigment quantification, adding leaf biochemical information derived from DRIFT spectroscopy + principal component analysis, using the same leaf pigment extract produced by the classical quantification method. We performed a dose–response experiment using P as the limiting nutrient, and maize (*Zea mays* L.) as a plant-test. After 45 d of growth, we evaluated the effects of P fertilization in total maize shoot biomass, P shoot accumulation, leaf pigment quantification by UV-Vis, and the evaluation of biochemical variations by DRIFT spectroscopy analysis associated with a chemometric approach in the same leaf extract used for pigment quantification. P fertilization raised biomass accumulation (~7.4x), P uptake (~2.3x), and total chlorophyll *a* and *b* contents (~2.1x). DRIFT spectroscopy analysis of extracted pigments revealed an elevated content of proteins and polysaccharides at high P availability. At low P availability, we found a low efficiency of N metabolism suggested by the accumulation of inorganic N forms. DRIFT spectroscopy applied together with the classic leaf pigment extraction and quantification method is a novel and promising tool for plant nutrition studies as a DRIFT spectroscopy metabolic profile protocol.

Introduction

The quantification analysis of pigments has been reported for almost a century.^{1–4} Chlorophylls and carotenoids play a crucial role in the process of photosynthesis, and the concentration of leaf pigments is a parameter recurrently measured

as an indicator of photosynthetic capacity,⁵ N content or whole plant nutrition,^{6,7} and plant health.⁸ At the laboratory scale, there are multitudes of options for pigment extractions using many organic solvents (*e.g.* acetone, methanol, ethanol, and dimethyl sulfoxide), solid–solution ratios, times of extraction and so on. In the last step, the concentration of pigments often follows photometrical determination.^{3,4,9} These procedures are easy, fast, and cheap, which help the leaf pigment quantification in plants to be widely used.

The methods for pigment extraction involve not only pigments but also several leaf compounds soluble in the organic solvent used, which until now has been barely explored by the research. The pigment extraction could give us other qualitative information or eventually semiquantitative data about the biochemical composition status, in the same line as the metabolic profile. The possibility to use the traditional leaf pigments' extraction targeting the soft plant metabolic profile discloses a new analytical approach for a well-established method, providing a deeper level of information on a quick and cheap lab work way.

Currently, the metabolic profile studies use laborious and time and expensive reagent-consuming methods. For example, liquid chromatography–mass spectrometry (LC-MS) analysis shows high sensitivity and can identify and quantify a large number of metabolites, even low concentration secondary metabolites.^{10,11} However, LC-MS¹⁰ requires around 6 days for a complete analysis and shows twelve critical points throughout the process, which increases the chances of failures and discrepancies between samples. All of that together also makes it an expensive analysis, with commercial labs charging around US\$ 100 per sample (*e.g.* Center for Environmental & Human Toxicology, University of Florida). The gas chromatography–mass spectrometry (GC-MS) method offers a good balance of sensitivity and reliability, being considerably more sensitive than nuclear magnetic resonance (NMR) and more robust than LC-MS. However, the GC-MS method¹² requires ten days for final results and shows two critical points as well, including the quantification of low derivatization compounds

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S; 47°36'31.4" W). Samples were air-dried and passed through a 4 mm sieve for experiments and a 2 mm sieve for chemical and physical analyses.²⁵ The soil presented low available P (13 mg dm⁻³, ion exchange resin extractant), 24% of clay,²⁵ a pH of 4.7 (0.01 M CaCl₂ at a ratio of 1:2.5 v/v), low exchangeable Ca²⁺ and Mg²⁺ (41 and 11 mmol_c dm⁻³, respectively, using ion exchange resin extractant), low organic C (27 g kg⁻¹, dichromate oxidation²⁶), and high P buffer capacity (low remaining-P, 22 mg L⁻¹ P²⁷). The soil clay mineralogy is mainly composed of kaolinite and gibbsite, as well as minor contents of Fe oxides such as goethite and hematite.

Soil samples of 2 dm³ were placed in plastic pots. CaCO₃ and MgCO₃ (Ca : Mg ratio of 4 : 1) were used to neutralize soil acidity to reach 50% base saturation. We added water in each pot to raise the soil water content to about 80% of the maximum water retention capacity (MWRC), maintaining the pots under incubation for 14 d. After, the soil samples were air dried and passed through a sieve with a mesh of 4.0 mm, the P fertilizer was mixed with soil samples. The P doses were 0, 30, 60, 120, 180 and 300 mg dm⁻³, using triple-superphosphate (20% P) as the P source, and are defined according to the remaining-P as 0, 0.25, 0.5, 1, 1.5 and 2.5 times the optimal P dose to overcome P soil availability limitation.^{28,29} Lastly on the same day, five maize (*Zea mays* L.) seeds (hybrid Dow SwB585) were sown per pot. After seedlings' emergence (after 4 d), two plants were left per pot.

The addition of other nutrients was carried out by nutrient solution applications according to Barreto *et al.* (2008).²⁹ Briefly, the doses and sources of the other nutrients corresponded to: 320 mg dm⁻³ of N [(NH₄NO₃); 160 mg dm⁻³ of K (KCl); 80 mg dm⁻³ of S [(NH₄)₂SO₄]; 4 mg dm⁻³ of Mn (MnCl₂·4H₂O); 4 mg dm⁻³ of Zn (ZnSO₄·7H₂O); 1.6 mg dm⁻³ of Fe(FeCl₃·6H₂O); 1.5 mg dm⁻³ of Cu (CuSO₄·5H₂O); 0.9 mg dm⁻³ of B (H₃BO₃); and 0.2 mg dm⁻³ of Mo (NaMoO₄·2H₂O). The plants were grown in a greenhouse under natural light conditions, with the temperature varying from 26 to 33 °C (day) to 16–20 °C (night), and a photoperiod (14 h day/10 h night) for 45 days. The pots were kept with the soil moisture close to 80% of the MWRC by irrigation in the upper part of the pots. The experimental design was in randomized blocks, with four replicates.

We aimed to boost the classic leaf pigment quantification, adding leaf biochemical information derived from DRIFT spectroscopy + principal component analysis, using the same leaf pigment extract produced by the classical quantification method. We performed a dose-response experiment revisiting a well-studied concern in plant nutrition about the low phosphate availability in highly weathered soil. Hence, we evaluated the effects of P dose fertilization in maize biomass yield or P accumulation and also presented new insights into biochemical composition from leaf pigment evaluation offered by DRIFT spectroscopy and PCA analysis.

Greenhouse trial

After 45 days of cultivation, we collected the middle third of the last totally expanded leaf, without the central vein and immediately stored at -20°C until subjected to freeze-drying. The total leaf pigments were extracted and measured as described by Lichtenthaler *et al.*⁴ Briefly, 100 mg of dried leaf samples were weighed in polypropylene tubes (15 ml), and 10 ml of cold acetone solution (acetone/deionized water; 80% v/v) were added, gently macerated using a thin plastic pestle (~ 1 min) and left to rest at -8°C for 24 h. The homogenate was centrifuged ($830g$ for 5 min) and the supernatant was collected. The spectrophotometer (UV5, Mettler Toledo) was previously blanked using acetone solution, then the absorbance

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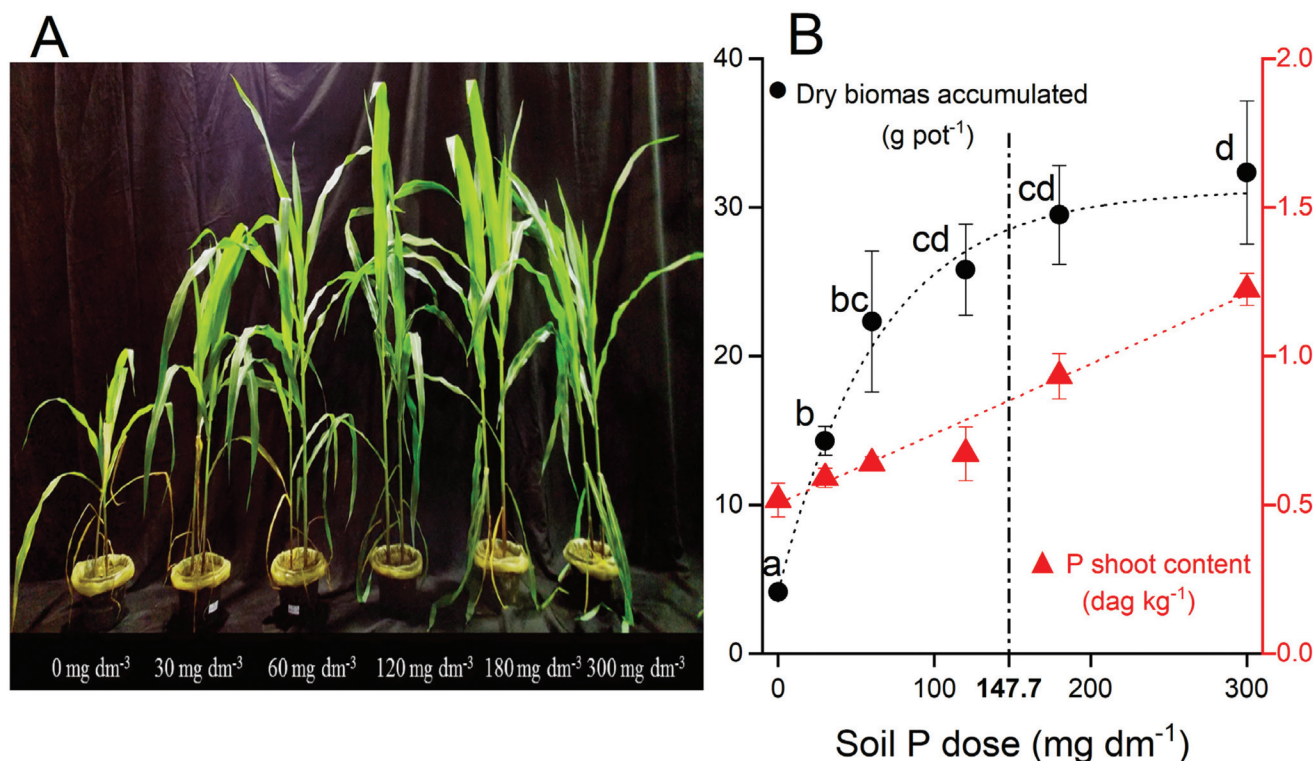


Fig. 1 (A) Maize plants 45 d after seedling emergence at the harvest. (B) The dry biomass accumulated (black circle) and P contents in plant tissue (red circle) over the P fertilizer applied. The different letters above each black circle represent significant differences between treatments at $P < 0.05$ according to Tukey's HSD for one-way ANOVA. Dashed line represents the P dose equivalent of maximum agronomic efficiency (147.7 mg dm⁻³ P). Data are presented as the mean of four replicates followed by its standard error ($\pm$ SE).

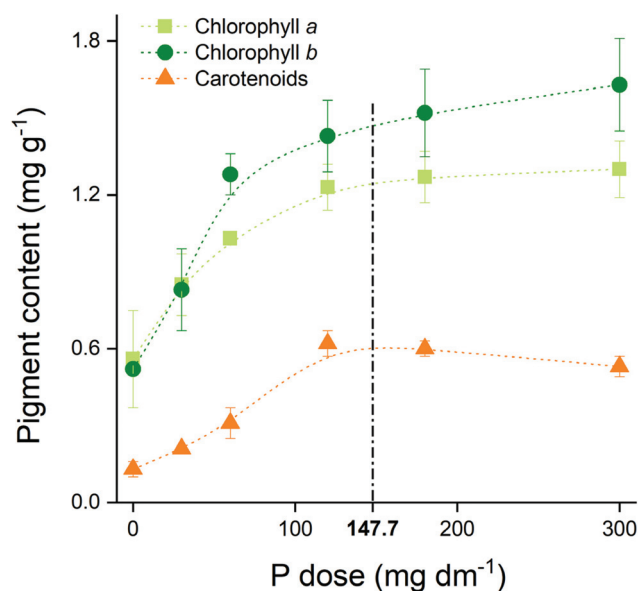


Fig. 2 Pigment content in maize leaf over P doses applied. Dashed line represents the dose equivalent of maximum agronomic efficiency (147.7 mg dm⁻³ P). Data are presented as the mean of four replicates followed by their standard error ($\pm$ SE).

DRIFT spectroscopy investigation

The DRIFT spectra give us some remarkable insights into the acetone extractable compounds in maize leaf (Fig. 3). The use of "acetone extractable" leaf fraction allowed us to investigate the spectra of biochemical compounds that should be hidden above the strong IR absorption band produced by dominant biochemical compounds, such as cellulose and lignin, if we used DRIFT spectroscopy analysis directly on the leaf.

At a lower P supply, the strongest peak was visibly centered at 1400 cm⁻¹, which could be assigned to ionized carboxylic acids (–COO⁻)^{35,36} of organic acids or inorganic N forms as NO₃⁻³⁷ and NH₄⁺.³⁸ The second option is preferential, because the strongest peak of carboxylic acids (1715 cm⁻¹)^{35,36} was not observed, and there is another distinctive peak (3100 cm⁻¹) related to the N–H bond.^{38,39}

The broad FTIR band has a center peak at 1045 cm⁻¹ and its area had a strong increase as a response to P doses (Fig. 3). This region is assigned to the carbohydrate-like fingerprint region,^{21,40,41} such as xyloglucan (1036 cm⁻¹),⁴² starch (~1010 cm⁻¹)⁴³ and sucrose (1056 cm⁻¹).⁴⁴ Also, this band could be associated with the DNA backbone at 1224 (ν_{as} PO₄⁻), 1088 (ν_s PO₄⁻), 1055 (backbone ν C–O), 1021 (furanose vibration),⁴⁵ or even free orthophosphate (–PO₄) groups,¹⁸ because the P content in the leaf was expressive at higher P doses (Fig. 1). The visual inspection of DRIFT spectra



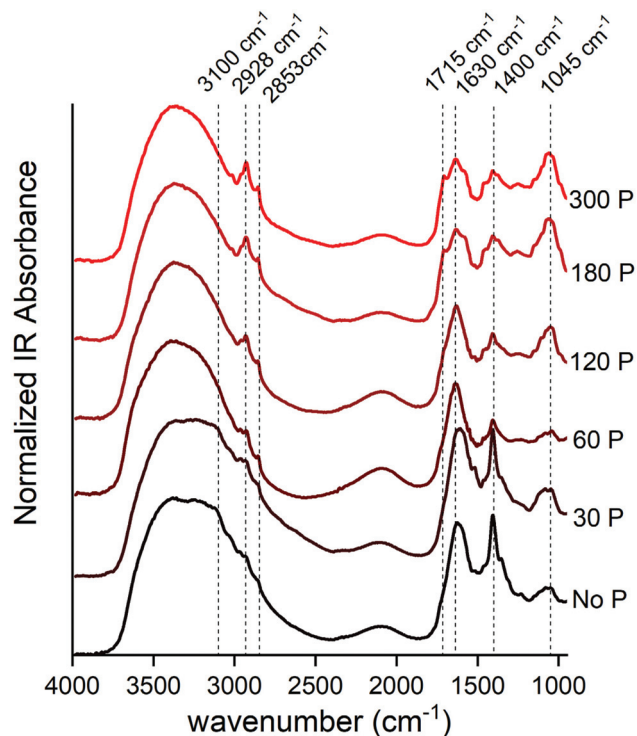


Fig. 3 Normalized DRIFT spectra of maize leaf extract from acetone solution (acetone/deionized water; 80% v/v) dried on KCl salt. The P dose (mg dm^{-3}) applied for rise from bottom (black) to upward (red). The data were stacked to improve visual analyses. The FTIR center peak assignment is: 3100 cm^{-1} = N–H stretch; 2920 cm^{-1} = symmetric methylene ($-\text{CH}_2-$) stretching; 2850 cm^{-1} = symmetric methyl ($-\text{CH}_3$) stretching; 1715 cm^{-1} = carbonyl double bond ($\text{C}=\text{O}$) of carboxylic acids and symmetric $\text{C}=\text{O}$ of ester groups, primarily from lipids and fatty acids; and $\text{C}=\text{O}$ group of chlorophyll molecule; $1650 \pm 50 \text{ cm}^{-1}$ = asymmetric ($\text{C}=\text{O}$) of amides associated with proteins (amide I) or carboxylate from pectin; 1400 cm^{-1} = ionized carboxylic acids ($-\text{COO}^-$) of organic acids and asymmetric N–O stretch of nitrate (NO_3^-) or asymmetric N–H stretching of ammonium (NH_4^+); $1045 \pm 30 \text{ cm}^{-1}$ = polysaccharides.

suggested an increment in both of these peaks, which likely means an increment in both these structural and energetic polysaccharides as well as genetic material, therefore suggesting a increase in whole plant activity linked with the photosynthetic metabolism.

Other peak regions that were boosted following the P dose increment were those around 2928 and 2853 cm^{-1} , which represent the C–H bond, mainly in hydrophobic compounds such as leaf wax and membrane lipid structures.^{46–48} Of note, these peaks are related to the aliphatic methylene chain, such as those found in chlorophylls and carotenoid structures, which is in agreement with the increment of total pigment content shown in Fig. 2.

The peak at 1630 cm^{-1} had a slight decrease from 0 to 300 mg dm^{-3} P. This peak is assigned to $\text{C}=\text{O}$ vibrations of amides associated with proteins (amide I)^{40,42,49} or carboxylate from pectin.⁵⁰ However, the peak concerning amide II (1550 cm^{-1}) was not prominent at lower P doses, and this suggests that the contribution of protein for this band region

began just at moderated P availability and P starvation. Thus, this peak likely is associated with pectin accumulation.

The slight shoulder at 1715 cm^{-1} could be assigned to the carbonyl double bond ($\text{C}=\text{O}$) of carboxylic acids^{35,36} and symmetric stretching $\text{C}=\text{O}$ of ester groups, primarily from lipids and fatty acids,⁴⁰ and the $\text{C}=\text{O}$ group of the chlorophyll molecule.⁵¹ The increase of absorbance derived from these IR bands followed the total improvement of metabolic performance due to P fertilization.

Principal component analysis (PCA)

The plot of the scores of PCs allowed the qualitative recognition of different groups of samples (Fig. 4). The first two principal components (PCs) explained about 93% of the total variability observed in the dataset; meanwhile PC1 did not allow for discrimination between sample groups as observed before,^{52,53} because all samples were located in the negative side of PC1 (data not showed), and since all samples corresponded to a similar biochemical composition of maize leaf regardless of the P dose. However, PC2 (9.49%) and PC3 (4.12%) reached a satisfactory group association, because PC2 segregated a lower P dose at positive score values, and higher P dose at negative ones (Fig. 4A). PC3 segregated the intermediates' P doses (60 and 120 mg dm^{-3}) at negative score values.

The loadings of the PC2 and PC3 plots vs. wavenumber (Fig. 4B) offer information about the link between scores and variables. Negative loadings for PC2, which grouped the higher P dose treatments, showed wavenumber characteristics mostly for methyl and methylene stretching groups (2920 and 2850 cm^{-1}),^{46–48} protein (from 1690 to 1610 cm^{-1}),^{40,42,49} $-\text{CH}_2$ scissoring of lipids ($1430 \pm 10 \text{ cm}^{-1}$) following 2920 cm^{-1} band absorbance improvement, and polysaccharides^{21,40,41} or even nucleic acid⁴⁵ ($1010 \pm 10 \text{ cm}^{-1}$). Of note, the positive PC2 loading at 1380 cm^{-1} , which is associated with inorganic forms of N, reinforces the accumulation of these ions in leaf of P-starved plants.^{54–57}

PC3 segregated in negative score, the sample spectra of the middle P dose (60 and 120 mg dm^{-3}) (Fig. 4A) mainly by negative loadings of amides (1601 cm^{-1}) and $-\text{CH}_2$ scissoring of lipids ($1430 \pm 10 \text{ cm}^{-1}$) and higher (180 and 300 mg dm^{-3}) together with the lower dose (0 and 30 mg dm^{-3}). However, when observing the variation on carotenoids pigments, one can state that there was a maximum at 120 mg dm^{-3} (Fig. 2), which could imply the PC segregation.

Discussion

The P fertilization improved strong biomass accumulation until 120 mg dm^{-3} P dose, and after that, there was no significant increment ($P < 0.05$) (Fig. 1). The maize plants, however, kept accumulating more P (Fig. 1). The maximum agronomic efficiency ($0.9A_{\text{max}}$) was reached with 147.7 mg dm^{-3} P, which is close to the recommendation of about 120 mg dm^{-3} as a suitable P dose for biomass production when related to its soil



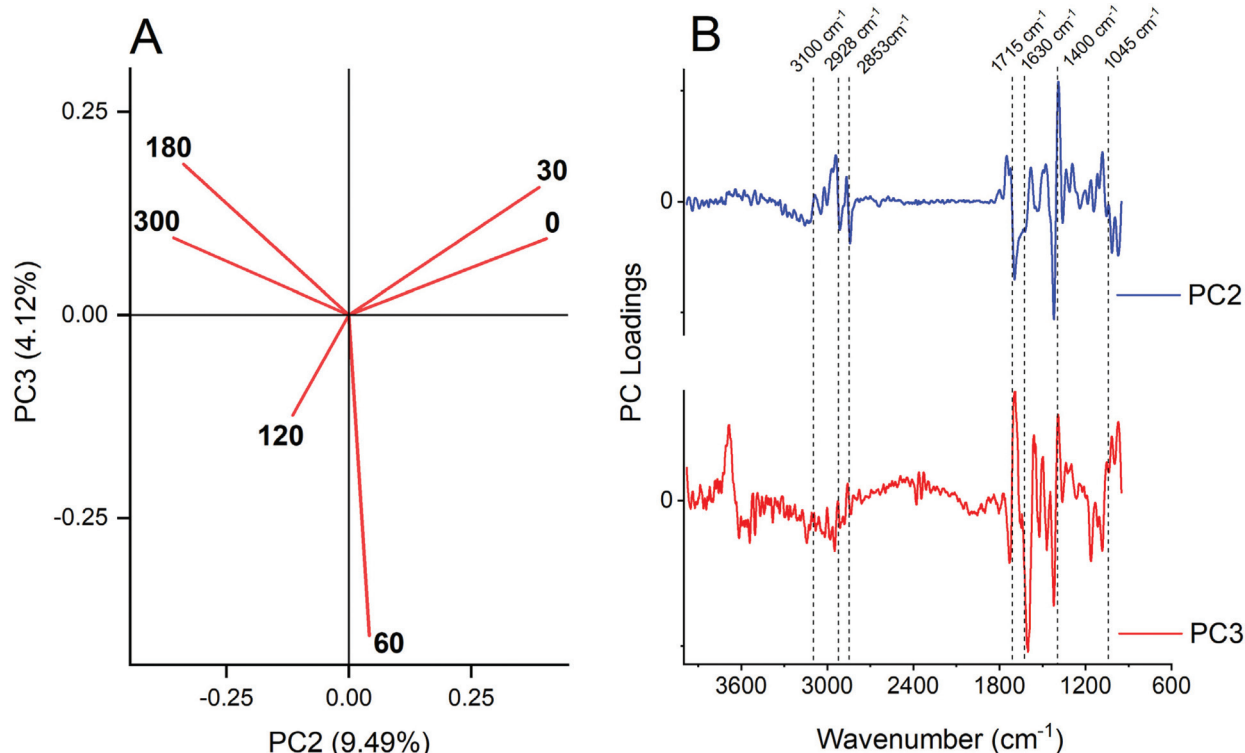


Fig. 4 (A) PCA of the first derivative DRIFT spectra (3900 to 950 cm^{-1} range) of pigment extraction of maize leaf dependent on P dose fertilization. (B) Loading profile of PC2 (blue line) and PC3 (black line). The FTIR center peak assignment is: 3100 cm^{-1} = N–H stretch; 2920 cm^{-1} = symmetric methylene ($-\text{CH}_2-$) stretching; 2850 cm^{-1} = symmetric methyl ($-\text{CH}_3$) stretching; 1715 cm^{-1} = carbonyl double bond ($\text{C}=\text{O}$) of carboxylic acids and symmetric $\text{C}=\text{O}$ of ester groups, primarily from lipids and fatty acids; and $\text{C}=\text{O}$ group of chlorophyll molecule; $1650 \pm 50 \text{ cm}^{-1}$ = asymmetric ($\text{C}=\text{O}$) of amides associated with proteins (amide I) or carboxylate from pectin; 1400 cm^{-1} = ionized carboxylic acids ($-\text{COO}^-$) of organic acids and asymmetric N–O stretch of nitrate (NO_3^-) or asymmetric N–H stretching of ammonium (NH_4^+); $1045 \pm 30 \text{ cm}^{-1}$ = polysaccharides.

P-buffer capacity.^{58–60} The accumulation of P in plant tissue in dose above A_{max} represents a high consumption of P (Fig. 1B), which did not improve the biomass yield.

The P nutrition was strictly associated with the photochemistry process and energy light efficiency for biomass accumulation in our results. The P absorption benefited more the production of Chl *b* than the Chl *a* as observed here and previously.⁶¹ Chl *b* is mostly associated with the light-harvesting complex connected to photosystem II (PSII), and the decrease of the Chl *a/b* ratio is probably indicative of a relative gain of photosystem II (PSII) peripheral antenna, or, alternatively, a change in photosystem stoichiometry in favor of PSII.^{5,61} This increment was followed by the contents of carotenoids, which play a role in the photo-protection of photosynthetic apparatus.^{5,61} Other molecular and physiological responses to P deficiency are well discussed in previous studies.^{62–64}

The higher N absorption and nutrient use efficiency are supported when there is a satisfactory P supply.⁶⁵ The effect of P nutrition in the metabolism of N had been described, and the N:P ratio has strong effects associated with root allocation, nutrient uptake, biomass turnover, and reproductive output.^{54–56} The increment of soluble sugar and decrease of nitrate in cabbage (*Brassica oleracea* L.) leaves were observed when there was proper P supply.⁵⁷

Our results and conclusion described here about P effects in plant metabolism by the DRIFT spectroscopy + PCA approach present a strong agreement with conventional metabolic profile studies. For example, the levels of carbohydrates and glycolysis intermediates in tomato leaves were extremely decreased by P deficiency,⁷ or citrus growing under P-deficient conditions showed increased activity of the arginine biosynthetic due to an increase in the concentration of ammonia in leaves.⁶⁶ The upregulation of protein degradation observed in the P-starvation condition⁶⁷ coincides with the low amide I IR band absorption (Fig. 3, $1650 \pm 50 \text{ cm}^{-1}$). In the end, DRIFT spectroscopy + PCA could not provide a detailed polysaccharide composition (e.g. quantities of saccharose, glucose or starch) as the traditional metabolic profile methods. However, at a very low cost (estimated to be around US\$ 0.15 per sample, including costs for pigment extraction and IR analysis), with a way simpler method that requires few procedure steps, and a relatively faster analysis (2–3 days), we reached conclusions similar to traditional methods for the metabolic profile.

Conclusions

Overall, our study discloses new insights into biochemical compounds and plant nutrition status, using for the first time,



DRIFT spectroscopy together with the classical plant pigment analyses. In summary, the visual inspection (Fig. 3) along with the chemometrics tool (PCA, Fig. 4) gave us considerable information that biochemical compounds are more affected by phosphate nutrition. The low efficiency of N metabolism by accumulating inorganic N forms at P starvation was highlighted as soon as the contents of protein and polysaccharides increased at high P availability. This conclusion is in agreement with a soft “metabolic profile” spectroscopic approach, which represents an upgrade in the conventional analyses for pigment quantification.

There are considerable variations in the scientific literature related to pigment extraction such as: the material used (*e.g.*, fresh, dried, and freeze-dry), solvent (*e.g.*, acetone 80 or 100%, methanol, and dimethyl sulfoxide), sample mix (*e.g.*, grinding and sonication), solvent-sample ratio, contact time, and temperature, even in the equations used for pigment quantification. In this first stage, none of this variation prohibits the DRIFT analysis for the plant leaf extract. The spectroscopic method proposed here could and should be adjusted for each laboratory routine. Other multivariate analyses such as multiple linear regression and hierarchical cluster analysis could be implemented.

The DRIFT spectroscopy + PCA approach described here brings deep level information by an inexpensive and fast workflow. We strongly suggest that this new method should be incorporated in leaf pigment analyses for several plant science studies as an evaluation of other abiotic and biotic plant stresses, as a soft “plant metabolic profile” assessment.

Conflicts of interest

There are no conflicts to declare.

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