

Post-photosynthetic fractionation of stable carbon isotopes between plant organs—a widespread phenomenon[†]

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Discrimination against ¹³C during photosynthesis is a well-characterised phenomenon. It leaves behind distinct signatures in organic matter of plants and in the atmosphere. The former is depleted in ¹³C, the latter is enriched during periods of preponderant photosynthetic activity of terrestrial ecosystems. The intra-annual cycle and latitudinal gradient in atmospheric ¹³C resulting from photosynthetic and respiratory activities of terrestrial plants have been exploited for the reconstruction of sources and sinks through deconvolution by inverse modelling. Here, we compile evidence for widespread post-photosynthetic fractionation that further modifies the isotopic signatures of individual plant organs and consequently leads to consistent differences in $\delta^{13}\text{C}$ between plant organs. Leaves were on average 0.96‰ and 1.91‰ more depleted than roots and woody stems, respectively. This phenomenon is relevant if the isotopic signature of CO₂-exchange fluxes at the ecosystem level is used for the reconstruction of individual sources and sinks. It may also modify the parameterisation of inverse modelling approaches if it leads to different isotopic signatures of organic matter with different residence times within the ecosystems and to a respiratory contribution to the average difference between the isotopic composition of plant organic matter and the atmosphere. We discuss the main hypotheses that can explain the observed inter-organ differences in $\delta^{13}\text{C}$. Copyright © 2005 John Wiley & Sons, Ltd.

Enzymatic and diffusional fractionation processes lead to discrimination against ¹³C during photosynthesis that varies between C₃, C₄, and CAM plants.¹ In C₃ plants, CO₂ is fixed within the Calvin cycle yielding C₃ compounds as first products and an isotopic signature that varies between −21‰ and −35‰. In C₄ plants, CO₂ is first fixed in C₄ organic acids that transport the carbon to separate tissues where sugar synthesis within the Calvin cycle takes place. The isotopic composition ranges between −9‰ and −20‰. CAM plants have an intermediate isotopic signature between C₃ and C₄ with variation depending on the proportion of carbon fixed in the dark into C₄ organic acids or in the light via the Calvin cycle. For instance, under water-deficient conditions, CAM plants open their stomata and fix CO₂ mainly at night time via the C₄ pathway accumulating malic acid. During the day, they close their stomata avoiding dehydration, and then the malic acid is decarboxylated providing CO₂ to the Calvin cycle.

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Photosynthesis leaves enriched CO₂ behind in the atmosphere. Carbon stored in plant materials is released to the atmosphere by respiratory processes. Therefore, the atmospheric CO₂ is relatively more enriched during summer when carbon fixation by photosynthesis is much greater than release by respiration. The degree of limitation of photosynthesis by resistances to diffusion of CO₂ to the sites of carboxylation and by photosynthetic capacities and the corresponding draw down of CO₂ is variable and hence also photosynthetic discrimination (Δ) varies with the relative contribution of enzymatic and diffusional fractionation processes (b and a , respectively). This variation has been shown to be a linear function of the ratio of CO₂ partial pressure in the leaf interior sub-stomatal cavities and air surrounding the leaf (p_i and p_a , respectively) [i.e. $\Delta = a + (b-a) p_i/p_a$].² All the factors changing stomatal conductance and/or photosynthetic capacity (e.g. water deficit, light, vapour pressure deficit), thus changing p_i/p_a , will also change the photosynthetic discrimination. Stomata are apertures in the leaf surface that facilitate gas exchange and can be actively regulated by the plant.

The seasonal signal of $\delta^{13}\text{C}$ in atmospheric CO₂ and the variation of discrimination with the water relations of the plants open up a wide range of applications for isotopic studies. They can be used to partition fluxes in the

ecosystem,³ to determine the fate of carbon of C₃ vegetation in C₄ soils, and vice versa,⁴ to study resource use in parasite-host systems,^{5,6} or to detect the time delay with which C is respired via root and soil respiration.⁷ In applying these methods it needs to be known if the isotopic signature of plant materials derived from photosynthesis can further be altered by post-photosynthetic fractionation processes. Evidence for the operation of (photo)respiratory fractionation in leaves has recently been reviewed.⁸ Results published subsequently have confirmed that leaves and whole shoots respire CO₂ enriched relative to the respiratory substrate for five woody species⁹ and four cultivated herbaceous species,^{10,11} respectively. The isotopic signatures of plant materials can be further modified if fractionation occurs during transport of metabolites or if respiratory fractionation is different in different organs. If this is the case, then the useful simplifying assumption that all plant organs built with substrates of the same photosynthetic origin will have an identical isotopic composition¹² cannot be verified.

Post-photosynthetic fractionation at the biochemical level is a well-documented phenomenon. Fractionation due to equilibrium and kinetic isotopic effects determines differences in isotopic signatures between metabolites and intramolecular positional isotopic effects.^{13,14} Here, we only refer to these differences where they are relevant for the interpretation of isotopic signatures of whole plant organs. Remarkable differences between $\delta^{13}\text{C}$ of different plant organs have been documented for organic matter or specific biochemical substances in many cases. For example, with some exceptions, leaves are generally isotopically lighter than the other plant organs¹⁵ and it has been postulated that this is due to a generic difference between autotrophic (gaining energy from photosynthesis) and heterotrophic (gaining energy from respiration of organic molecules) organs.¹⁶

However, at present no clear-cut conclusions can be drawn from the observation of differences in $\delta^{13}\text{C}$ of different organs. *A priori* it cannot be ruled out that these differences are an occasional product of the use of substrates produced at different times under different environmental and ontogenetic conditions (as, e.g., proposed by Francey and co-workers¹⁷ to explain the differences between leaves and wood). For reasons of parsimony this is a hypothesis to be favoured because it can potentially explain the phenomenon on the grounds of already known sources of variation in photo-

synthetic discrimination in the course of time. However, a metabolic origin of the variation of ^{13}C abundance between organs cannot be ruled out *a priori*. In search of alternative hypotheses explaining the differences between C-isotopic compositions of organs, it can be proposed that: (1) discrimination occurs during the transport of assimilates; (2) the metabolites used for export are enriched in ^{13}C (e.g. sucrose) with respect to the photosynthetic products; (3) respiratory processes discriminate against ^{13}C at different degrees in different organs along with different types of metabolic pathways; (4) the biochemical composition varies from one organ to another along with characteristic signatures of metabolites; and (5) different rates of carboxylation for replenishment of Krebs cycle intermediates (via PEP-carboxylase, PEPc, EC 4.1.1.31) lead to varying rates of incorporation of relatively heavy carbon (some of these hypotheses were formulated earlier^{18–21}).

The aim of the present work is to (a) check the consistency of the difference in $\delta^{13}\text{C}$ of autotrophic and heterotrophic organs with an analysis of published data and (b) to evaluate the alternative hypotheses that can potentially explain these differences. We also report some supplementary unpublished data on French bean plants.

RESULTS

Differences between $\delta^{13}\text{C}$ of organs and metabolites from published sources

Published results on $\delta^{13}\text{C}$ of organic matter or a specific substance class were compiled from studies that reported at least results for leaves and one other organ (see list in the Appendix). Data were not normally distributed. The hypothesis of zero difference between sample means was tested with the Wilcoxon signed rank test for paired samples. Statistical analyses were done with R.²² For 410 reported results on isotopic composition of plant organs, leaves were on average 1.26‰ more depleted in ^{13}C than other plant organs (see Table 1 and Fig. 1). In only 38 cases were leaves found to be isotopically heavier; in 39 cases the difference was smaller than 0.2‰. A total of 333 comparisons resulted in leaves being more than 0.2‰ more depleted in ^{13}C . Nine out of the 39 cases where leaves were ^{13}C -enriched by more than 0.2‰ were measured on *Pinus edulis* branch vs. leaves including the most extreme value. For 38 reported results on isotopic composition of total belowground and aboveground plant

Table 1. Differences in ^{13}C -isotopic signature of organic mass or classes of metabolites of different plant organs (p = error level for rejection of null hypothesis of zero difference)

	Belowground— aboveground	Other organs— leaves	Roots— leaves, C ₃	Roots— leaves, C ₄
Average	1.23	1.26	1.08	0.09
Minimum	-1.4	-2.06	-0.43	-1.8
Maximum	5.64	8.4	3.5	1.5
Standard error of the mean	0.19	0.07	0.09	0.28
p	<0.001	<0.001	<0.001	0.68
n	38	410	78	10
n with diff. < -0.2	1	38	1	2
n with diff. > +0.2	33	333	71	4
n with -0.2 < diff. < 0.2	4	39	6	4

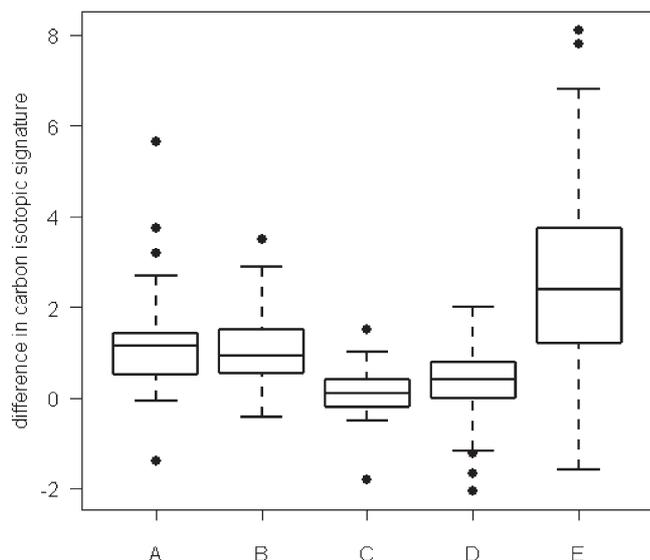


Figure 1. Differences between isotopic carbon signatures of plant organs: A) belowground—aboveground; B) roots—leaves, C_3 plants; C) roots—leaves, C_4 plants; D) non-woody stems—leaves; E) woody stems—leaves. The line in the middle of the box represents the median. The box spans from first to third quartile. The whiskers are at the level of the most distant data point within 1.5 times the interquartile distance from the box.

organs, the aboveground fraction was on average 1.2‰ more depleted in ^{13}C than the belowground fraction. These overall figures include classes of comparison with contrasting results. The 88 comparisons of root and leaf organic mass resulted in an average difference of 0.96‰. This is the only case for which sufficient data ($n > 9$ for individual groups) were available to analyse C_4 and C_3 vegetation separately. Roots of C_3 plants were always ^{13}C -enriched (average 1.08‰) compared with leaves or had a statistically non-distinguishable signature, with one exception of wheat plants at ripeness where the difference was -0.43 ‰. The differences of the means mentioned up to here were all significantly different from zero at $p < 0.001$. As opposed to roots of C_3 plants, C_4 roots did not show a statistically significant difference compared with leaves ($n = 10$). In four cases the difference was less than 0.2‰. In two cases roots were isotopically light-

er, in four cases heavier than leaves. In some cases also stems were found to be ^{13}C -depleted compared with leaves. This was the case for woody stems (average difference 1.91, minimum -2.0 , maximum 8.1, $n = 116$) as well as non-woody stems (average difference 0.32, minimum -2.1 , maximum 2.0, $n = 90$). Thus, especially non-woody stems had isotopic signatures close to leaf tissues.

Cellulose was on average 1.29‰ ^{13}C -enriched compared with the organic mass of the respective organ while lipids and lignins were 4.67‰ and 3.17‰ depleted, respectively (Table 2). Sugars and starch as cellulose were ^{13}C -enriched compared with the organic mass of the organ from which they were extracted.

DISCUSSION

Differences between heterotrophic and autotrophic organs

Our analysis of published data of plant organ isotopic composition shows a consistent difference between leaves and other organs. Roots and woody stems were isotopically heavier than leaves, while non-woody stems showed only a slight similar trend for higher $\delta^{13}C$ than leaves. In 12% of the comparisons for non-woody stems a difference more negative than -0.2 ‰ was found, while in 67% of the cases the differences were above 0.2‰. The experimental data obtained on *Phaseolus vulgaris* (Table 3) fit into this pattern.

These results add to earlier evidence in favour of post-photosynthetic fractionation processes^{15,16,19,20,23} that lead to consistent inter-organ differences in $\delta^{13}C$. The special case of herbaceous stems that bear isotopic signatures closer to those measured in leaves would fit into a general pattern if the observed differences are related to the degree of heterotrophy. Herbaceous stems are mostly green photosynthesising organs and in this respect metabolically closer to leaves than to roots or woody stems.

It is unlikely that changes in photosynthetic discrimination in the course of time can explain the general trends in organ $\delta^{13}C$ differences we observed in the current study. Intrinsic ontogenetic changes generally lead to a closure of stomates and decrease in photosynthetic discrimination in the course of time. Therefore, it can probably not explain differences between leaves and roots.

In a study on rice,²⁴ Scartazza and coworkers investigated whether differences in the isotopic composition of organs

Table 2. Differences in ^{13}C -isotopic signature for the chemical components cellulose, lipids and lignin as compared with total organic matter

	Cellulose— organic mass	Starch— organic mass	Sugars— organic mass	Lipids— organic mass	Lignin— organic mass
Average	1.29	2.02	1.71	-4.68	-3.17
Minimum	-1.3	0.0	-3.8	-8.8	-6.7
Maximum	4.2	4.1	7.7	-0.6	1.6
Standard error of the mean	0.08	0.21	0.3	0.23	0.30
p	<0.001	<0.001	<0.001	<0.001	<0.001
n	107	34	46	84	37
n with diff. < -0.2	5	0	6	84	33
n with diff. $> +0.2$	98	32	36	0	2
n with $-0.2 < \text{diff.} < 0.2$	4	2	4	0	2

Table 3. Isotopic signature of carbon in total organic matter (TOM), starch, heat precipitated proteins (HPP) and sucrose of different organs of *Phaseolus vulgaris* L. cv. Contender. Culture conditions and analyses methods as in previous experiments^{29,33,46} (mean \pm SD, $n = 3$, and $n = 1$ for data without SD)

Organ	TOM	Starch	HPP	Sucrose
Bud	-29.11 \pm 0.46	-27.64 \pm 0.41	-28.61 \pm 0.34	-28.61 \pm 0.03
Cotyledon, leaf blade	-28.49 \pm 0.16	-27.87 \pm 0.49	-27.25 \pm 0.36	-28.30 \pm 0.62
Cotyledon, leaf veins	—	-26.37 \pm 0.28	-27.19 \pm 0.43	-28.46 \pm 0.10
Cotyledon, leaf petiole	—	-26.41 \pm 0.73	-25.88 \pm 0.36	-28.40 \pm 0.17
First leaf blade	-29.1 \pm 0.42	-28.28 \pm 0.39	-28.11 \pm 0.24	-28.45 \pm 1.04
First leaf veins	—	-27.94 \pm 0.58	-27.58 \pm 0.76	-28.32 \pm 0.44
First leaf petiole	—	-27.69 \pm 1.11	-28.54	-28.22 \pm 0.48
Second leaf blade	-29.01 \pm 0.69	-27.51 \pm 0.08	-28.02 \pm 0.26	-28.56 \pm 0.52
Second leaf veins	—	-27.62 \pm 0.28	-28.00 \pm 0.48	-27.45 \pm 0.19
Second leaf petiole	—	-28.92	-27.81	-27.23 \pm 0.39
Third leaf (whole)	-29.14 \pm 0.51	-27.68 \pm 0.51	-28.80 \pm 0.35	-28.53 \pm 0.34
Root	-27.63 \pm 0.13	-26.50 \pm 0.51	-27.98 \pm 0.39	-28.17 \pm 0.70
Stem	-28.97 \pm 0.27	-27.73 \pm 0.28	-26.90 \pm 0.61	-28.67 \pm 0.14

could be explained as a consequence of the use of assimilates that had been produced at different times with different overall photosynthetic discrimination. They found that even plants that were well watered all the time showed differences in $\delta^{13}\text{C}$ between organs and concluded that different photosynthetic discrimination in the course of the time cannot explain all the observed variation in isotopic composition.

In addition, studies of heterotrophic metabolism have provided further evidence for post-photosynthetic fractionation processes. Heterotrophically growing plant organs (tomato and tobacco) exhibit a more positive $\delta^{13}\text{C}$ than the substrates used for growth, irrespective of the use of externally supplied sugars or sugars imported from other plant organs.¹⁹ ^{13}C signatures in young expanding leaves of several tropical tree species were less negative than they were in adjacent older, mature leaves.²⁵ This difference could not be explained by differences in p_i/p_a . The young leaves had higher p_i/p_a ratios; consequently, a higher discrimination can be expected in the younger leaves. Also, the anatomical changes in the course of early leaf ontogeny should lead to increases in the resistance to gas transport in the course of time and thus impose a trend of decreasing discrimination, which in turn cannot explain the observed differences between old and young leaves. The author could convincingly show that seasonal variations in the environmental conditions and air $\delta^{13}\text{C}$ can be ruled out as explanation for the observed phenomenon.

Also, the result that sugars and starch were on average less depleted in ^{13}C than organic mass (Table 2) can be taken as an indication of post-photosynthetic discrimination. Generally, sugars and starch will represent the isotopic signature of recent photosynthetic products.¹ The wide range of variation in the difference between sugars and organic mass $\delta^{13}\text{C}$ can be expected because photosynthetic discrimination varies with stomatal aperture. However, on average, organic mass was more negative than sugars, indicating an overall apparent discrimination between primary photosynthetic products and the bulk organic matter.

The simplest hypothesis to be favoured, as long as it cannot be falsified, states that consistent differences in isotopic composition of plant organs can be explained by varying

isotopic signature of assimilates produced at different times. On the basis of the results discussed so far we conclude that this hypothesis does not have a high explanatory power. There are good reasons to look for alternative hypotheses because the simplest hypothesis (i.e. the one to be favoured according to Occam's razor) cannot explain the main trends in the observations.

$\delta^{13}\text{C}$ of organic matter in organs and fractionation during biosynthesis

If the phenomenon we seek to understand is not due to a simple difference in time of carbon assimilation at different discrimination, then it must be due to additional fractionating processes during the building up of the considered organs. A corollary of this statement is that there exists a difference in the isotopic signature of incoming and outgoing fluxes. The exchange fluxes that can potentially carry different signatures are: respiration, carbon fixation via carboxylases, import and export (including phloem and xylem transport, ablation of surface waxes, exudation, losses due to foraging of selected components by parasites and predators, and root uptake of organic substances).

In addition, other potential causes have frequently been invoked for the explanation of differences in isotopic composition of organs, namely differences in the biochemical composition of the organs.^{24,26–28} Given that classes of biochemical substances do not have the same isotopic signature (Table 2), a high depletion of an organ in ^{13}C can be correlated with high lignin or lipid contents, while high cellulose content is expected to correlate with a relatively low depletion. However, the chemical composition of an organ cannot be the cause of a difference in the isotopic composition of two organs because of mass balance, but only a consequence of another exchange process that induces the overall difference. For example, even a high production rate of, e.g., lipids, leading to a high fraction of the total mass that is strongly depleted in ^{13}C , cannot lead to a depletion of ^{13}C in organic matter, because the heavy carbon atoms not used in lipid synthesis will reside in other organ components that are isotopically heavier than the average organic mass. A correlation of high concentration of depleted compounds and organic matter that is depleted relative to the source can

only result (a) if fractionation occurs during respiration and heavy carbon atoms are respired preferentially, (b) if carbon export of isotopically heavy compounds occurs, or (c) if carboxylation reactions add carbon to the organ with an isotopic signature different from the signature of respiratory substrates. Thus, differences in chemical composition cannot be the cause of correlated differences in isotopic signature but only a corollary of exchange fluxes that enrich or deplete the organ under consideration.

Discrimination during transport

The comparison of sugars in leaf blades, petioles and major veins (Table 3) as well as sugars in blades and phloem (=conduits within plant stems that transport assimilates from source to sink organs) sap of *Phaseolus vulgaris* did not show a consistent isotopic difference so that fractionation during transport leading to preferential export of heavy carbon isotopes from photosynthesizing leaves cannot be proven with these data. Phloem sap was collected with the method reported by Groussol *et al.*²⁹ Leaf petioles were placed in 5 mmol L⁻¹ EDTA at pH 6 for 4 h in the dark. Sugars in second leaf blades and sugars collected from phloem sap just below the second leaves had only slightly different isotopic signatures (sucrose in blade -27.06 and -27.05, sucrose in phloem sap -27.18 and -27.85, fructose plus myo-inositol in phloem sap -27.79 and -26.62 in sample 1 and 2, respectively).

Accordingly, only a small and non-significant difference between total phloem exudates (-30.4‰) and sugars plus organic acids in leaf blades (-29.9‰) of *Triticum aestivum* plants in the ear-forming stage was reported.¹⁵ In the grain-filling stage also only a small and non-significant difference between total phloem exudates (-29.5‰) and sugars plus organic acids in leaf blades (-29.4‰) was found. However, sugars and acids extracted from the ears had an isotopic carbon signature of -26.6‰.

Hobbie and coworkers¹⁶ interpreted findings by Pate and Arthur³⁰ on differences between leaf organic matter and phloem sap sample signatures in eucalypt trees as an indication for fractionation during export. However, as Pate and Arthur pointed out the leaf dry matter signatures were expected to be more negative than phloem exudates in a dry period of the year, because the leaves had been produced during the less stressed winter/spring period prior to the stress conditions. Thus the measurements show the contrast of leaf signatures and phloem signatures of plant materials of different age and cannot be taken as evidence for fractionation during export.

The strongest indication of a possible discrimination associated with the export of carbohydrates from leaf tissues comes from measurements of isotopic signatures of sugars in leaves and current year stem sections of European beech, *Fagus sylvatica*, all throughout the growing season.²⁶ From the end of May to October the leaf sucrose had a 2–3‰ more negative $\delta^{13}\text{C}$ than the sucrose extracted from stems. Similar differences of 2–3‰ were reported for the same species under different environmental conditions when the canopy weighed average of leaf sugar $\delta^{13}\text{C}$ was compared with the sugars sampled from phloem at breast height.²⁷ In contrast to this, in another study on European beech,³¹ a depletion of

phloem exudates of twigs relative to organics in the water-soluble fraction of leaves in the shade crown of beech by 0.5‰ was found in July. For the sun crown a similar but non-significant trend was measured whereas phloem bleeding-sap collected at breast height was enriched by 1.1‰ relative to leaf water-soluble organics. In September no difference was found between all the leaves, twigs and base of stem.

However, these data should be interpreted with caution because stem photosynthesis can provide a fraction of the stem sugars and exchange with storage carbohydrates can occur. Depending on the balance between respiration, gas transport in the stem and resistances to the gas exchange with the air surrounding the stems, the source CO₂ isotopic signature for stem photosynthesis can vary between the signature of CO₂ released by respiration and the atmospheric CO₂ signature and thus the discrimination during stem photosynthesis can vary with the balance of the carbon fluxes.

While the results discussed so far were obtained on aboveground organs, we are aware of only one comparison of leaf and root sugars reported in the literature.¹⁸ Leaf sugars were found to be 2.1‰ more depleted on average of measurements at different times of the day in *Beta vulgaris*. In *Phaseolus vulgaris* a difference with the same sign although non-significant and much smaller (0.1–0.4‰) was found (Fig. 2).

The evidence for differences between $\delta^{13}\text{C}$ of transport metabolites sampled in the source organ and the sink organ or along the transport pathway supports the notion of an apparent discrimination associated with export fluxes. However, it does not allow differentiation between two groups of potential mechanisms: Either a discrimination occurs at a step of the export process itself or the exported substrates come from a pool within the exporting cell that has a signature different from the average signature of this substrate within the cell. Hobbie and Werner²⁰ favoured the latter hypothesis. They proposed a fractionation of photosynthetic products in leaf cells. Carbon isotopically lighter than primary photosynthetic products would be incorporated into lipids and lignin. The remaining enriched

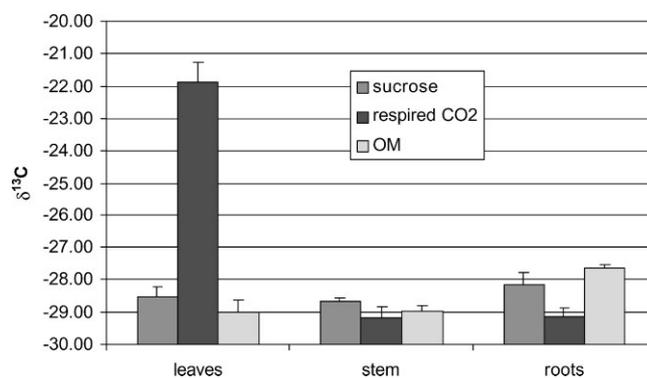


Figure 2. Isotopic signatures ($\delta^{13}\text{C} \pm \text{SE}$) of sucrose, respired CO₂ and dry organic matter in leaves, shoots and roots of *Phaseolus vulgaris*. $\delta^{13}\text{C}$ of root respired CO₂ was independent of the pH (varied between 2.8 and 7) used in the solution for washing the roots, indicating that it is not an artefact due to discrimination during solution/dissolution processes. The root respiration rate during these measurements was $2.11 \pm 0.29 \text{ nmol CO}_2 (\text{g fresh weight})^{-1} \text{ s}^{-1}$.

carbohydrates would be used for cellulose synthesis and export. Thus, exported carbohydrates would be less depleted than the primary photosynthetic products and so are sink organs like roots.

In summary, the hypothesis that fractionation during assimilate transport is the cause of the observed consistent differences between root and leaf carbon isotopic signatures currently cannot be rejected nor confirmed on the basis of the scarce experimental results available. More experiments with sampling of leaf source sugars and phloem sap sugars along the export paths are needed in order to further explore the potential role of fractionation during transport in post-photosynthetic fractionation processes.

The effect of respiration

It has been shown for *Phaseolus vulgaris* leaves that CO₂ respired in the dark is constantly enriched by about 6‰ compared with sucrose,³² a result confirmed by the 6.3‰ enrichment compared with sucrose as reported above for a more recent experiment on the same species. As opposed to this, the apparent discrimination in roots (Fig. 2) indicated net release of lighter carbon isotopes as compared with the root sugars. In both cases, the results on apparent discrimination are consistent with the difference found between sucrose, the potential substrate and bulk organic matter, the average product. These results suggest that release of heavy carbon in leaves leads to an organic matter that is isotopically lighter than the source molecules while in roots organic matter is relatively enriched in ¹³C by preferential net release of lighter carbon isotopes.

A similar opposite apparent respiratory fractionation was found in a study¹⁰ on whole shoots and roots of sunflower (*Helianthus annuus*), perennial ryegrass (*Lolium perenne*) and alfalfa (*Medicago sativa*). Respiratory CO₂ of whole shoots was enriched by 1.06–5.64‰ relative to organic mass, while the respiratory CO₂ of whole roots was depleted by 0.46–5.39‰ relative to organic mass (OM). At the same time roots had higher δ¹³C than shoots, in accordance with the results reported with our review. The slope of a regression of the difference in OM signatures on the difference in respiratory discrimination (forced through the origin) indicates that, for every ‰ difference between shoot and root respiratory CO₂, the difference between OM signatures is about 0.26‰. This relationship is consistent with the view that respiratory discrimination is associated with biosynthesis, and the moles of carbon incorporated into biomass per moles of C consumed are approximately 0.75–0.8.

The isotopic signature of respired CO₂ of a given organ tended to be more negative when the respiratory quotient, RQ (CO₂ released per O₂ consumed), was lower (data not shown). The variation in δ¹³C with RQ corresponds to results obtained on *Phaseolus vulgaris* leaves previously.³³ RQ varies with the oxidation state of the respiratory substrates. Less oxygen per mole of carbon released as CO₂ is required during respiration of highly oxidised substrates (e.g. organic acids) and more when highly reduced substrates (e.g. lipids) are used. At the same respiratory quotient, the CO₂ released by leaf respiration was always several ‰ less depleted in ¹³C than the CO₂ released by respiration of roots and shoots (data not shown) suggesting that the signature of substrates used

for respiration is not the only source of variation of the signature of respired CO₂.

If ¹³C is preferentially released during leaf respiration⁸ how can the opposite discrimination during root respiration be explained? The most promising candidate mechanism is dark-assimilation of carbon via PEPc. When apparent fractionation⁸ during respiratory metabolism is measured, with the traditional gas-exchange methods currently available, the effects of carboxylation reactions cannot be factored out. In consequence, up to now the measured fractionation represents the net result of all processes exchanging CO₂ between organ carbon and the atmosphere. During dark respiration as well as during photosynthetic metabolism in the light, several carboxylating enzymes can potentially fix inorganic carbon and add carbon atoms to the organic matter.³⁴ PEPc carboxylates phosphoenolpyruvate and provides C₄ organic acids for the replenishment of the Krebs cycle. This function is required when organic acids are exported from the Krebs cycle either for regulation of cellular pH or as carbon skeletons for amino acid synthesis. The carbon fixed by PEPc is 5.7‰ enriched³⁴ in ¹³C relative to source CO₂. Qualitatively, the incorporation of inorganic C *in vivo* by PEPc,³⁵ the release of some of the fixed carbon after short residence times,³⁶ and dilution of the label by transport of newly synthesised biomolecules from roots to other plant organs^{37,38} have been demonstrated with ¹⁴C pulse labelling. Use of organic acids in pH regulation and provisioning of carbon skeletons for amino acid synthesis both imply a high variability of PEPc activity³⁴ that is known to change with the type of nitrogen nutrition (NO₃⁻ vs. NH₄⁺),³⁸ availability of other nutrients in the soil solution,³⁹ and soil CO₂ partial pressure.^{40,41} A general quantitative model for the rate of CO₂-fixation by PEPc is not yet available (however, see Raven and Farquhar³⁴ for a characterisation of the quantitative bounds at the whole plant level).

However, the observed rates and *in vitro* measurements of PEPc activity indicate that the PEP carboxylation can be sufficient to explain the differences between the signatures of respired CO₂, sugars and organic mass in roots. In *Phaseolus vulgaris*, for the results shown in Fig. 1, *in vitro* PEPc activity (measured as previously described^{42,43}) was 5.3 and 29.3 nmol s⁻¹ (g fresh weight)⁻¹ in roots (comparable with the rates reported on the same species⁴⁴) and leaves, respectively. Thus, in roots, the *in vitro* rate was higher than the net respiration rate of 2.11 ± 0.29 nmol CO₂ (g fresh weight)⁻¹ s⁻¹ suggesting that PEPc can potentially fix carbon at rates of several 10% of the gross respiration rate.

In the soil-root space most of the CO₂ is derived from respiration of organic molecules. The effect of variable rates of PEPc activity fixing CO₂ with signature of organic mass is illustrated in Fig. 3.

The model calculations show that the results obtained on the differences between the signatures of sugars compared with respired CO₂ and organic mass, respectively, in *Phaseolus vulgaris* roots can be explained by respiration rates and dark carboxylation rates in the range of commonly observed values. More experiments with pulse-chase labelling are needed to study the role of dark carboxylation and effects of export of amino acids and organic acids from the roots towards other organs.

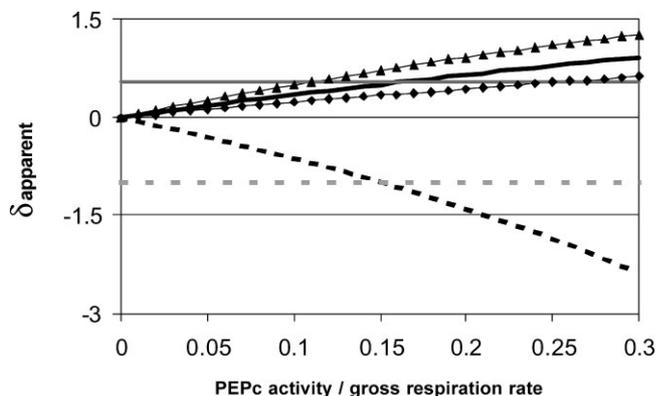


Figure 3. Modelled difference between signatures of respired CO_2 (dark broken line) and signature of organic mass (dark continuous line), respectively, and isotopic signature of source molecules for respiration and synthesis of organic mass for varying rates of PEPc activity and PEPc discriminating by -5.7‰ relative CO_2 in the gas phase. It was assumed that the source signature is constant in time, carbon fixed by PEPc is not subsequently respired within the Krebs cycle, and the use of respiratory substrates proceeds without discrimination. The signature in organic mass depends on the ratio of carbon used for syntheses to carbon respired, u . Three cases are shown for $u = 1.5$ (dark continuous line), $u = 1.0$ (dark continuous line with triangles), and $u = 2.34$ (dark continuous line with diamonds). The grey continuous line represents the difference between organic mass and sugars and the broken grey line the difference between respired CO_2 and sugars shown in Fig. 1 for *P. vulgaris*

Significance of the post-photosynthetic fractionation for ecological and physiological studies

For the application of inverse modelling the implicit or explicit¹² assumption is used that $\delta^{13}\text{C}_{\text{plant}} = \delta^{13}\text{C}_{\text{leaf}} = \delta^{13}\text{C}_{\text{photosynthetic product}}$, i.e. the respiratory flux always carries the signature of the photosynthetic products. We showed above that in general this is not the case in real plants.

However, if organic matter stemming from organs of different isotopic composition is respired during decomposition with an approximately constant mixture, the basic assumption could still be kept up in a modified way. This is the case if (a) the release of carbon on inter-annual time scales proceeds with virtually constant proportions from slowly and rapidly decaying fractions, (b) we are interested in averages over large areas that are relevant for inverse modelling. However, in managed ecosystems, large fractions of biomass are removed from the system (e.g. wood harvesting) and transported to distant places where they are used. A harvest flux that exports a fraction of biomass with an isotopic signature deviating from the average signature of primary photosynthetic products will bring about a horizontal isoflux. This isoflux changes the isotopic mass balance of the local and regional systems exchanging with the atmosphere. The relevance of this potential effect can be checked by an extensive comparison of tissues with low versus tissues with high residence times in the ecosystem.

Fung and coworkers¹² determined the overall discrimination during uptake of carbon by the biosphere (Δ_{bio}) for the net primary production flux from estimates of photosynthetic discrimination. Thus it was assumed that autotrophic respiration releases carbon of the same signature as gross photosynthetic products. If the overall discrimination during autotrophic respiration was of the order of magnitude (0.7‰) as determined by Klumpp and coworkers,¹⁰ this would increase the estimate of Δ_{bio} by 4–6% and change the estimated biospheric sink by about 5%.

When isoflux measurements are used to break down the different respiratory sources within an ecosystem, instantaneous differences between the signature of CO_2 respired by different organs as shown in Fig. 2 need to be known. These differences stem from differences in the isotopic signature of the respiratory substrate and apparent fractionation during respiration, which are both not yet well characterised, especially for roots. Therefore, further studies of the organ-specific respiratory isotopic signatures are required.

The difference between root and leaf isotopic composition can partly explain the observed profiles of $\delta^{13}\text{C}$ in soils.⁴⁵ Often the isotopic signature of organic mass in surface litter layers is close to isotopic composition of aboveground plant organs. Then, a rapid increase of $\delta^{13}\text{C}$ in soil organic matter with the first centimetres of increasing soil depth is observed, followed by a slower rise down to greater depth. Under C_3 vegetation the predominant input of more depleted leaf litter to the surface soil layers and less depleted dead roots to lower soil layers contributes to this observed profile.

Finally, a good understanding of post-photosynthetic fractionation in plants lays the basis for the use of measurements of isotopic signatures in gas-exchange fluxes as a non-invasive tool of studies of metabolism.

CONCLUSIONS

When the variation of carbon isotopic signature of different plant organs due to the seasonal variation in photosynthetic discrimination is accounted for, there still remain differences between organ signatures that cannot be explained only on the grounds of the seasonal variation. Mass balance requires that these differences are associated with fractionation during exchange fluxes across the organ boundaries. The biochemical composition of the organs cannot be the cause of the observed inter-organ differences in isotopic signature but only a corollary of the exchange fluxes described above. The inter-organ differences can be caused either by fractionation associated with transport of metabolites across organ boundaries or by fractionation during heterotrophic metabolism. In the first case, either a selection of metabolites to be exported from a source organ may occur because of compartmentation within the source organ or fractionation at a metabolic branching point or the fractionation may occur during the export process itself. In the second case, fractionation along respiratory metabolic pathways may occur or dark fixation via carboxylases may introduce carbon with a signature distinct from the plant internal source carbon. Currently, there is evidence available in favour of both the two main groups of hypotheses. More studies are needed, especially on carbon import to roots and isotopic signatures of root

respired CO₂, in order to disprove any of the hypotheses. A good understanding of the nature of the post-photosynthetic fractionation between plant organs is required for the use of isoflux measurements in plant physiology, ecosystem studies and studies of the biogeochemistry of the terrestrial biosphere at the global scale.

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APPENDIX

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