

Technische Universität München TUM School of Life Sciences

The role of stores in recycling sucrose and supplying respiration of a perennial C₃ grass: on the effects of Last Glacial Maximum to projected end-of-21st-century

atmospheric CO₂ concentration

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Contents

ABSTRACTI
ZUSAMMENFASSUNGIII
LIST OF FIGURES V
LIST OF TABLES X
CHAPTER 1. INTRODUCTION 1
CHAPTER 2. MATERIALS AND METHODS17
CHAPTER 3. RESULTS AND DISCUSSION
3.1 Accuracy and precision of the near-natural abundance ¹³ CO ₂ / ¹² CO ₂ labelling system30
3.2 The role of stores in recycling sucrose40
3.3 The role of stores in supplying respiratory substrate52
3.4 A connection between carbohydrate and respiratory substrate supply system59
CHAPTER 4. CONCLUSION AND OUTLOOK 64
REFERENCE
ACKNOWLEDGEMENTS 75
LEBENSLAUF
APPENDIX77
Publication 1. Atmospheric CO_2 and VPD alter the diel oscillation of leaf elongation in
perennial ryegrass: compensation of hydraulic limitation by stored-growth
Publication 2. ¹⁸ O enrichment of leaf cellulose correlated with ¹⁸ O enrichment of leaf sucrose
but not bulk leaf water in a C ₃ grass across contrasts of atmospheric CO ₂ concentration and
air humidity

Abstract

Aims: Organic carbon stores provide substrate for plant growth and respiration and are mainly formed as carbohydrates (e.g. fructan and sucrose). However, little is known about the effect of past and future carbon dioxide concentration ([CO₂]) on the role of organic carbon stores in the recycling of sucrose and supplying substrate to respiration. This thesis aimed to quantitatively assess the ¹³C tracer dynamics in the main non-structural carbohydrate fractions at the whole shoot level and in plant respiration at the whole plant level of perennial ryegrass (*Lolium perenne* L.) - a perennial C₃ forage grass - and their responses to [CO₂] during plant growth. Of particular interest was (1) to assess the performance of the ¹³CO₂/¹²CO₂ labelling system used to trace C incorporation, turnover and use in respiration, (2) to determine the effects of [CO₂] on the fluxes in central carbohydrate metabolism (fructan, sucrose, glucose, and fructose) in the shoot, including fructan cycling and sucrose recycling, (3) to investigate the [CO₂] effect on the respiratory substrate supply system at a whole-plant level including the numbers, half-lives, size and importance of kinetically distinct pools, and (4) to explore the potential relationship between carbohydrate fractions and respiratory substrate use.

Materials and Methods: Stands of perennial ryegrass were grown in $[CO_2]$ of 200, 400 or 800 µmol mol⁻¹ with a limiting nitrogen fertilizer supply. Well-developed, same-age stands were dynamically labelled with near-natural abundance ${}^{13}CO_2/{}^{12}CO_2$ mixtures, followed by measurements of the ${}^{13}C/{}^{12}C$ ratio (defined as $\delta^{13}C$) of plant respired CO₂ in the dark period and $\delta^{13}C$ of shoot carbohydrate fractions (sucrose, fructan, glucose, and fructose). Finally, relevant compartmental models were parametrized for the tracer kinetics of respired CO₂ at the whole-plant scale and carbohydrate fractions at the shoot level.

Results and Discussion: The results showed that the ${}^{13}\text{CO}_2/{}^{12}\text{CO}_2$ labelling system was prone to inevitable but minor artefacts (mainly due to opening the growth chambers for system maintenance and plant handling and sampling purposes), so that max. 4% error in source CO₂ occurred through air contamination with extraneous CO₂ across the three [CO₂] levels, and a max. 2.6% inter-contamination when a preparative HPLC system separated different carbohydrates. The work highlights the value of a two-chamber approach with CO₂ sources of distinct $\delta^{13}C_{CO2}$ to determine the isotopic end-members of mixing models for quantitative analysis of the tracer kinetics in respired CO₂, biomass, and carbohydrate fractions. Carbohydrate accumulation was enhanced by increasing [CO₂] from 200 to 800 µmol mol⁻¹,

mainly due to a significant increase in fructan mass. A four-pool compartmental model of carbohydrate metabolism indicated that the most significant impact of elevating [CO₂] on central carbohydrate metabolism in the shoot was to increase carbon cycling through the fructan pool. Sucrose re-synthesis from breakdown products of fructan (fructose) accounted for an increasing proportion of total sucrose synthesis and was similar in magnitude to sucrose neo-synthesis at elevated [CO₂]. Independent of the [CO₂] effect, respired C was supplied by a short-term and a long-term pool, and the importance of the long-term pool was substantial (*c*. 58%). Increasing [CO₂] from 200 μ mol mol⁻¹ to 800 μ mol mol⁻¹ increased the size of the long-term pool by 197% and its half-life by 148%. Comparing tracer kinetics of respired CO₂ and individual carbohydrates showed that the long-term pool was likely composed of fructan while the short-term pool probably mainly consisted of a mixture of transport and vacuolar sucrose.

Conclusion: This thesis provides, firstly, a methodical assessment of a quantitative ${}^{13}CO_2/{}^{12}CO_2$ labelling system and demonstrates that a two-chamber approach is a powerful means to analyze the C tracer kinetics in biomass or carbohydrate fractions. Secondly, the analysis of tracer kinetics and compartmental modelling confirmed a constitutive role of the fructan store in recycling sucrose and supplying substrate to respiration. In addition, it allowed the identification of carbohydrate substrates in the respiratory supply system. Finally, my results demonstrate the [CO₂] effect on carbon fluxes in carbohydrate metabolism at the whole shoot level and in the respiratory substrate at the whole plant scale, providing information on how an important perennial C₃ grass may have responded to past Glacial Maximum low [CO₂] and may respond to future high [CO₂].

Zusammenfassung

Zielsetzung: Organische C Speicher liefern C für Pflanzenwachstum und Respiration v.a in Form von Kohlenhydraten (z.B. Fructane und Saccharose). C-Flüsse in Speichern liefern Informationen über den Kohlenhydrat-Stoffwechsel und das respiratorische Versorgungssystem der Pflanzen, und deren Reaktion auf Umweltbedingungen. Das Ziel dieser Arbeit war die Untersuchung der Rolle von organischen C-Speichern und des respiratorischen Versorgungssystems bei unterschiedlicher atmosphärischer CO₂-Konzentration in Deutsch Weidelgras (Lolium perenne, C₃). Von besonderem Interesse war (1) die Beurteilung der Funktionsfähigkeit einer ${}^{13}CO_2/{}^{12}CO_2$ -Markierungsanlage, (2) die Anwendung von Tracer-Kinetiken zur Untersuchung des Einflusses verschiedener atmosphärischer CO₂-Konzentrationen auf C-Flüsse im zentralen Kohlenhydrat-Stoffwechsel (Fructane, Saccharose, Glukose und Fructose) im Sproß, (3) die Untersuchung von C-Flüssen im respiratorischen Versorgungssystem auf der Ebene ganzer Pflanzen incl. Anzahl, Halbwertszeit, Größe und Bedeutung kinetisch unterschiedlicher Pools, und (4) die Untersuchung des Zusammenhangs zwischen verschiedenen Kohlenhydrat-Fraktionen und dem respiratorischen Substrat.

Materialien und Methoden: Im Markierungsversuch wurden Bestände von Deutsch Weidelgras bei atmosphärischen CO₂-Konzentrationen von 200, 400 oder 800 µmol mol⁻¹ und limitierender N-Versorgung angezogen. Geschlossene Bestände gleichen Alters wurden mit CO₂ aus zwei isotopisch unterschiedlichen, natürlichen Quellen markiert, begleitet von Messungen des δ^{13} C in der Dunkelperiode respirierten CO₂ sowie des δ^{13} C der wichtigsten Kohlenhydrat-Fraktionen im Sproß. Mithilfe dieser Daten wurden kompartimentelle Modelle parametrisiert, die die Tracer-Kinetiken im respirierten CO₂ auf der Ganz-Pflanzen-Ebene sowie in den Kohlenhydrat-Fraktionen im Sproß beschreiben.

Ergebnisse und Diskussion: Beim Betrieb der ¹³CO₂/¹²CO₂-Markierungsanalge wurden unvermeidliche aber geringfügige Fehler festgestellt, welche u.a. beim Öffnen der Klimakammern für Wartungs- und Probenahmezwecke auftreten; 4 % des von den Pflanzen aufgenommene C stammte von CO₂ aus der Umgebungsluft (Kontamination) und nicht von den beiden zur Markierung verwendeten CO₂-Quellen. Zudem war die Trennung der verschiedenen Kohlenhydrat-Fraktionen mittels präparativer HPLC mit einem geringen Fehler behaftet (max. 2,6 %). Andererseits konnte die Leistungsfähigkeit des Zwei-Kammer-Ansatzes der ¹³CO₂/¹²CO₂-Markierung demonstriert werden, v.a. für die Bestimmung der beiden

Endglieder des Tracer Mischungsmodels zur Bestimmung der Anteile alten und neuen C in den untersuchten Pools und Flüssen. Die Gehalte der Nicht-Struktur-Kohlenhydrate nahmen mit zunehmender CO₂-Konzentration von 200 auf 800 µmol mol⁻¹ zu, v.a. hinsichtlich der Gesamtmasse der Fructane. Ein Vier-Pool-Model des Kohlenhydrat-Stoffwechsels implizierte, dass der stärkste CO₂-Effekt auf der Umsetzung des Fructan-Pools beruhte. Saccharose-Resynthese aus den Abbauprodukten von Fructanen erklärte einen zunehmenden Anteil der gesamten Saccharose-Synthese und war in der erhöhten CO₂-Konzentration von ähnlicher Größenordnung wie die Saccharose-Neusynthese. Unabhängig von der CO₂-Konzentration wurden der Substratkohlenstoff für die Respiration aus einem kurzfristigen und einem Speicherpool bereitgestellt. Die Bedeutung des Speicherpools für die Respiration war beträchtlich (c. 58 %). Die Erhöhung der CO₂-Konzentration führte zu einer Vergrößerung des Speicher-Pools um 197 % und zu einer Verlängerung seiner Halbwertszeit um 148 %. Der Vergleich der Tracer-Kinetik von respiratorischem CO₂ und einzelnen Kohlenhydraten zeigte, dass der Speicherpool vermutlich mit Fructan identisch war, während der kurzfristige Pool wahrscheinlich hauptsächlich aus einer Mischung aus Transport- und vakuolärer Saccharose bestand.

Schlussfolgerungen: Diese Arbeit diente der Einschätzung der Funktionalität einer ${}^{13}CO_2/{}^{12}CO_2$ Markierungsanlage und demonstrierte die Nützlichkeit eines Zwei-Kammer-Ansatzes mit unterschiedlichen ${}^{13}CO_2/{}^{12}CO_2$ -Quellen für die Analyse von Tracer-Kinetiken in verschiedenen C-Pools und -Flüssen. Tracer-Kinetiken und kompartimentelle Modellierung demonstrierten die grundlegende Rolle von C-Speichern im Saccharose-Stoffwechsel und als Substrat für die Respiration. Zusätzlich ermöglichten die Ergebnisse die Identifizierung von Kohlenhydrat-Fraktionen als Substrat für die Respiration. Schliesslich zeigten die Untersuchungen zu den Auswirkungen unterschiedlicher CO₂-Konzentrationen reagierten und erlauben die Einschätzung wie sie sich an zukünftige, erhöhte CO₂-Konzentration anpassen.

List of Figures

Fig. 1.4 The fraction of unlabelled C ($f_{unlabelled}$) in fructan (filled squares), sucrose (open circles), glucose (filled triangles), and fructose (open triangles) with labelling duration in the youngest

List of Figures

Fig. 2.4 Schematic representation of 13CO2/12CO2 gas exchange and labelling system (A: from Schnyder et al., 2003) and growth chamber picture (B). SC, screw compressor (S40; Boge, Bielefeld, Germany); AD, adsorption dryer (KEN 3100; Zander, Essen, Germany; and molecular sieve: activated aluminium oxide F200; Alcoa, Houston, TX, USA); AR, air receiver (1 m3) (Magnet Kft, Magocs, Hungary); F1, oil and water condensate drain (CSP005; Hiross, Mönchengladbach, Germany), F2 oil, water and particle filter (\geq 0.01 µm; G12XD and filter element: 2030X, Zander), F3, universal filter (\geq 1 µm; G12ZHD and filter element: 2030Z,

List of Figures

Fig. 3.1.2 The δ 13C-difference between δ 13CCO2 outlet and δ 13CCO2 inlet with time (δ 13CCO2 outlet - δ 13CCO2 inlet). CO2 concentration at chamber outlet ([CO2]outlet) was maintained near target [CO2]: 200 (a, b), 400 (c, d) and 800 (e, f) µmol mol-1. Growth chambers were supplied with either 13C-organic (δ 13CCO2 -43.5‰; left) or 13C-mineral CO2 (δ 13CCO2 -5.6‰; right). The criterion of data evaluation was that measurements in the first 45 min of a light period or following the opening of the chamber were removed, and values over $1.5 \times IQR$ (Interquartile Range) away from the mean were removed as outliers. Data points and error bars represent daily means \pm SD (n = 9-23). Note that the dataset included the 14-d

List of Tables

Table 3.1.1 Isotopic difference (termed 'spread') between paired plant growth chambers receiving 13C-organic and -mineral CO2 for different gas exchange, biomass and watersoluble carbohydrate components. Spread ($d\delta 13CX$) was calculated as $d\delta 13CX = \delta 13CX 13C$ mineral – $\delta 13CX 13C$ -organic), with X designating the parameter of interest, e.g. bulk biomass or sucrose. Spread was calculated for $\delta 13CCO2$ at the inlet ($d\delta 13CCO2$ inlet) and outlet $(d\delta 13CCO2 \text{ outlet})$ of paired growth chambers in the light and dark period of days 61 to 64, daytime canopy CO2 exchange (dol3CNd) and respiration in the dark (dol3CRn) for days 61 to 64, bulk shoot (d δ 13Cshoot) and root C (d δ 13Croot), and fructan (d δ 13Cfructan), sucrose (dδ13Csucrose), glucose (dδ13Cglucose) and fructose (dδ13Cfructose) extracted and purified from shoot biomass sampled at the beginning of the light period on day 65. Growth chambers were maintained at near target [CO2] of 200, 400 or 800 µmol mol-1 using one of two CO2 sources, a 13C-organic (δ 13C -43.5‰) or a 13C-mineral source (δ 13C -5.6‰). δ 13CNd and δ13CRn were determined during periods of steady-state gas exchange of chambers, measurements in the first 45 min of a light period or following the opening of the chamber were removed, and values over $1.5 \times IQR$ (Interquartile Range) away from the mean were removed as outliers. Except for [CO2] and δ 13CCO2, all conditions were kept the same in all chambers (see Materials and Methods). Daily means (\pm SD) for n = 2 to 10......35

Table 3.1.2 δ13C of reference carbohydrates measured pure or after HPLC separation of an 80:5:5:4 (wt:wt:wt) mixture of analytical grade inulin-fructan, sucrose, glucose and fructose (all from Merck, Darmstadt, Germany). The concentration of the individual carbohydrates in the mixture corresponded to typical concentrations in shoot extracts of plants grown at [CO2]

Table 3.2.4 Parameters of the four-pool compartmental model of central carbohydrate metabolism in the shoot of L. perenne. Parameters were numerically optimized as described in chapter 2, based on the tracer kinetics data shown in Fig. 3.2.1. Fluxes are expressed as g C m-2 leaf area h-1 and calculated by the differential equations given in chapter 2, using the carbohydrate contents per leaf area as shown in Table 3.2.3. The results of the fluxes predicted by the optimized rate constants are given in bold. Numbers in a row or column with different

superscript	letters	indicate	a	statistically	significant	difference	at	95%	confidence
intervals			••••						49

Table 3.2.5 Results of one-way ANOVA, testing the effect of [CO2]......51

1. Introduction

1.1 Plant responses under varying atmospheric CO₂ concentration

In recent decades, there has been a considerable research interest into the impact of increasing atmospheric CO₂ concentrations ([CO₂]) on various aspects of plant biology, including photosynthesis, growth, nutrition, and morphology (Franks et al., 2012). Studies have explored the effects of [CO₂] across a range of scales, spanning from those present during the Last Glacial Maximum (LGM; 18000–20000 years ago) to projected end-of- 21^{st} -century levels (Sage & Coleman, 2001; Lüscher et al., 2004; Ainsworth and Long, 2005).

An early expectation has been that [CO₂] would enhance photosynthetic capacity, which is typically associated with the rate of carboxylation by Rubisco in C₃ plants, such as perennial ryegrass (Lolium perenne), the object of this study. In C₃ plants, the Rubisco carboxylation reaction is not saturated at current [CO₂] levels, and an increase in atmospheric [CO₂] was generally expected to lead to a boost in photosynthesis (Drake et al., 1997). Conversely, low $[CO_2]$ levels have been found to strongly limit C_3 photosynthesis (Sage & Coleman, 2001), reducing the capacity of plants to assimilate nutrients (Tissue et al., 1995). Additionally, reductions in carbon (C) availability have been observed to alter biomass allocation patterns, with plants allocating more biomass to their leaves at the expense of their roots (Sage & Cowling, 1999). Furthermore, constrained biomass production caused by low [CO₂] levels may have constrained the ability of plants to complete their life cycle (Dippery et al., 1995). Studies have also shown that short-term elevated $[CO_2]$ can significantly stimulate leaf photosynthesis (Ainsworth et al., 2003), increase leaf concentration of total nonstructural carbohydrates and alter the carbon : nitrogen ratio of litter (Körner, 2000), but long-term standand ecosystem-scale responses were generally much smaller and sometimes absent (Long et al., 2006). In particular, the availability of growth resources such as nitrogen (N) and phosphorus (P) were found to strongly influence a plant's response to increasing [CO₂] (Terrer et al., 2016; Terrer et al., 2019). For example, Lüscher et al. (2004) reported that the availability of mineral N in the soil was a main factor limiting the growth response of L. perenne to elevated [CO₂].

Organic C stores represent a significant functional component of plant biomass under rising CO_2 (Körner, 2006), but their role and integration in whole plant C metabolism has not been studied in any detail, as I explain below. Therefore, this thesis utilized experimental and

modelling approaches to explore the interplay of organic C stores in sucrose recycling and respiratory substrate supply to improve understanding of how organic carbon store respond to varying $[CO_2]$ levels, as experienced in the past and as projected for the end of this century. In that I used perennial ryegrass as a model C₃ grass, as it has a very high economic value as a forage crop (Chapman et al., 2017) and its physiology has been studied in relative detail also in response to $[CO_2]$ (Lüscher et al., 2004).

1.2 Ecophysiological functions of organic C stores in C₃ grasses

Virtually all CO₂ fixed in photosynthesis is either deposited in plant structural biomass or returned to the atmosphere through (autotrophic) respiration (Trumbore, 2006; Lehmeier et al., 2008). Thus, respiration and deposition in structural biomass represent the terminal fates for fixed carbon (C) within plants. Interestingly, the balance between being respired and used in growth, denoted by the so-called carbon-use efficiency (CUE, defined as CUE = (photosynthesis – respiration) / photosynthesis), tends to be relatively constant in herbaceous plants with a CUE of ~0.6-0.7, meaning that a relatively constant fraction of 0.3-0.4 of photosynthetically fixed C is respired in diverse environmental contexts (Thornley, 2011; Gong et al., 2017).

However, between being fixed and respired or deposited in structural biomass, C compounds may cycle through diverse metabolic pathways in different cellular compartments in multiple tissues and organs (Gebbing et al., 1998; Lehmeier et al., 2008; Schnyder et al., 2012; Verbančič et al., 2018). The temporal delay between being captured in photosynthesis and respired or deposited as structural C can be extended dramatically by temporary storage as reserves (Chapin et al., 1990; Smith & Stitt, 2007). These stores can represent a substantial fraction of total vegetative plant biomass and are thought to serve primarily as buffers that accommodate or balance fluctuating disparities between photosynthetic activity and assimilate demands for growth, defense and maintenance activities (Pollock & Cairns, 1991; Schnyder, 1993; Chapin et al., 1990). In most C₃ and C₄ plant families, vacuolar sucrose in leaf blades and other photosynthetically active tissues is the main diurnal carbohydrate store in both vegetative and reproductive grass (Jenner & Rathjen, 1972; Schnyder, 1993; Lunn et al., 1999; Barro et al., 2020). This includes the grasses which comprise many of the globally most important crops and a wide variety of cultivated forage species (Farrar & Farrar, 1986; Sicher et al., 1986; Schnyder, 1993; Isopp et al., 2000), such as perennial ryegrass (*L. perenne*, C₃),

the model species used in this study. Starch built up in chloroplasts during the day and degraded during the night is also known as an essential diurnal carbohydrate store, sometimes complementary with vacuolar sucrose, in many plants, e.g. *Arabidopsis* (Sulpice et al., 2009; Graf & Smith, 2011; Stitt & Zeeman, 2012), including in grasses (Gordon et al., 1980; Cairns et al., 2002).

Nevertheless, in C₃ grasses, much greater amounts and concentrations of carbohydrates can be stored longer-term (several days to months) as fructans (Kühbauch & Thome, 1989; Pollock & Cairns, 1991). For example, Wagner et al. (1983) observed a fructan concentration of up to 75% of the dry weight of illuminated barley leaves when photosynthate export was slowed down or blocked by cool conditions or detachment of leaves. Fructan molecules are water-soluble oligomers or polymers of fructose, generally containing one glucose residue, and are synthesized from sucrose in vacuoles (Wagner et al., 1983, 1986; Sprenger et al., 1995; Pollock & Cairns, 1991). Fructan synthesis can occur in virtually all immature growing and mature fully-differentiated tissues of C₃ grasses, including leaf blades, leaf sheaths, leaf growth zones, stems, roots, floral and reproductive tissues (Nelson & Spollen, 1987; Pollock & Cairns, 1991; Schnyder et al., 1993; Pavis et al., 2001). Fructan synthesis in vacuoles occurs by different fructosyl transferase enzymes that catalyse the transfer of the fructosyl residue of a donor sucrose molecule onto an acceptor sucrose molecule, thus forming a trisaccharide or onto an (acceptor) fructan molecule, causing fructan chain elongation by one fructosyl residue (Chalmers et al., 2005). These fructosyl transfers cause liberation of the glucosyl residue of sucrose, which is then used for sucrose (re)synthesis in the cytosol (Nguyen-Quoc et al., 2001). Fructan degradation or depolymerization is catalyzed by fructan exohydrolase enzymes releasing fructose, which is subsequently used for sucrose recycling in the cytosol (Lattanzi et al., 2012). Given that fructan synthesis and hydrolysis may coincide in the same compartment (Wagner et al., 1983; Pollock & Cairns, 1991), the fructan pool in a vacuole can be conceived as an extension of the sucrose pool, which permits the maintenance of a very high concentration of stored carbohydrate without the adverse osmotic consequences that would arise from storage of a similar quantity of sucrose (Nelson & Spollen, 1987).

Fructan storage depends on the source-sink relationships in plants, with an overabundance of photo-assimilates relative to growth and maintenance needs stimulating fructan deposition (Kühbauch & Thome, 1989; Pollock & Cairns, 1991). Such relationships are naturally provided by cool conditions (such as are prevailing in the fall; Pollock & Jones, 1979), nutrient limitation (particularly of nitrogen; McGrath et al., 1997; Lattanzi et al., 2012),

Chapter 1 Introduction

mild drought (Chaves et al., 2003; Muller et al., 2011), high irradiance (Savitch et al., 2000) and elevated atmospheric CO₂ concentration ([CO₂]) (Stitt, 1991; Smart et al., 1994; Isopp et al., 2000; Rogers & Ainsworth, 2006). However, whether or not low [CO₂] – such as the ~200 μ mol mol⁻¹ that existed near the Last Glacial Maximum (LGM), *c*. 20'000 years ago – could have compromised the ability of grasses to store fructan has not been studied to my knowledge.

Generally, it is held that long-term stores of fructan are mobilized during the senescence phase of graminaceous crops that occurs in parallel with grain development and growth (Schnyder, 1993; Smouter & Simpson, 1991; Gebbing et al., 1998) or perennial grasses following the dormant period (Noël et al., 2000), during periods of stress (e.g. drought, Amiard et al., 2003), and after disturbance by fire (de Moraes et al., 2016; Martínez-Vilalta et al., 2016) or defoliation by grazing or mowing (Morvan-Bertrand et al., 2001). Importantly, however, stores also play a constitutive role in supplying substrate for growth and autotrophic respiration of grassland vegetation during undisturbed growth, as indicated by C tracer studies in the field (Gamnitzer et al., 2009; Ostler et al., 2016). Specifically, evidence also supports a constitutive role of fructan in central metabolism, with fructan turnover contributing very significantly to sucrose recycling (regeneration) in leaves even in steady-state conditions when photosynthesis is active and fructan levels remain virtually constant in perennial ryegrass (Lattanzi et al., 2012, see also chapter 1.3, below). Although sucrose is a primary photosynthetic product (Stitt et al., 1987; Goldschmidt & Huber, 1992; Farrar et al., 2000), foliar fructan turnover may participate constitutively in its (re-)cycling, which is vital for understanding the relationship between photosynthetic activity and sucrose availability. The importance is further enhanced by the fact that carbohydrate translocation in grasses (and most other species) occurs in the form of sucrose virtually exclusively (Giaquinta, 1983; Ward et al., 1997). In such a system, sucrose recycled via fructan turnover would become available for growth and respiratory metabolism centers in the heterotrophic plant parts.

Besides carbohydrate stores, soluble proteins have also been discussed as an additional putative source of 'stored' substrate for respiration in connection with protein turnover, particularly under stressful conditions (Araújo et al., 2011; Lehmeier et al., 2012).

As respiration produces 'only' CO_2 , there is no metabolic tag in the respired gas that could inform us of its' metabolic origin, that is the specific metabolic pathways visited by the precursors of the respired CO_2 inside intact plants. However, it is known that – in the strict sense – respired CO_2 derives directly from only a few compounds, mainly malate, pyruvate, isocitrate, α -ketoglutarate, or gluconate-6-P (Heldt, 2005; Tcherkez et al., 2012), which together comprise only a tiny fraction of whole plant biomass (Lehmeier et al., 2008). These compounds must be turned over very rapidly, given the commonly observed respiration rates. However, Lehmeier et al. (2008) demonstrated the presence of a storage pool (with a half-life of ~33 h) that comprised approximately 11.5% of total plant biomass-C feeding whole plant respiration (see chapter 1.3, below). If it is true that stored substrate can be used with a similar CUE as primary photosynthate, then the total size of the store must have been in the order of approximately 35% of shoot biomass-C in the investigations of Lehmeier et al. (2008). That estimate is close to the total metabolic biomass supporting (structural) biomass synthesis of perennial ryegrass in tracer studies in the field (Ostler et al., 2016). At present, a full understanding of the chemical identity of the metabolic pools supporting respiration of perennial ryegrass is still lacking, although it is hypothesized here that fructan plays a central role. In particular, it is presently unknown how the low level of [CO₂] in the LGM and future predicted [CO₂] affect the role of the fructan pool in sucrose recycling and the role of stores in supplying substrate for respiration in grasses.

1.3 Principles and application of carbon isotopes in carbohydrate and respiratory CO₂ measurement

Temporal variation in the transport of C metabolites through metabolic networks and stores of intact plants has been explored by labelling with radioactive (¹¹C- or ¹⁴C-CO₂) or stable C isotope (¹³C/¹²C-CO₂) tracers. Specifically, C tracers have been used to partition CO₂ fluxes, estimate transport velocity, and assess C allocation to different plant parts and partitioning between different biochemical compounds (Schnyder et al., 2012; Kölling et al., 2013). This has included tracing of C through the different metabolites of the reductive pentose phosphate cycle (Bassham et al., 1950), disentangling of photorespiration and gross photosynthesis (Ludwig & Canvin, 1971), distinguishing between dark respiration and net photosynthesis during the light period of the day (Schnyder et al., 2003), and determination of the importance of storage-derived pre-anthesis C to carbohydrate and protein synthesis in wheat grains (Gebbing & Schnyder, 1999). The different isotopes are not equally well suited to studying different physiological questions. For instance, radioactive isotopes (¹¹C, ¹⁴C) are a practical tool for studying translocation velocity *in vivo* by using emission/decay detectors/sensors placed along the translocation path (Geiger & Swanson, 1965; Jahnke et al., 1981). Conversely,

the physiology or biochemistry of longer-term storage cannot be studied with ¹¹C due to its short half-life (20 min). Stable isotope mixtures of CO₂ (i.e. ¹³CO₂/¹²CO₂) are preferable, particularly for applications in the field or for large-scale and long-duration (>1 d) experiments in controlled environments, as they avoid health hazards originating from the use of radioactive isotopes and, by using natural or industrial sources of different origin, are relatively inexpensive.

Two different strategies of applying isotopically labelled CO₂ have been employed in the past: pulse- and dynamic (or steady-state) labelling (Schnyder et al., 2012, 2017). Typically, pulse-labelling experiments have employed either the radioactive isotopes (11 C or 14 C) or 13 C at high isotopic enrichment. Pulse-labelling experiments consist of three phases: (1) a prelabelling period in which the plants are grown in normal, i.e. non-labelled, usually ambient CO_2 , (2) a labelling pulse of short duration (normally seconds to hours depending on the research question), followed by (3) a so-called 'chase' period in which the photosynthetically fixed tracer becomes diluted by new, 'non-labelled' C. Commonly, the distribution of the tracer is followed during the chase period. In contrast, dynamic labelling experiments have only two phases: a pre-labelling period and a labelling period in which tracer distribution is monitored. The labelling period may have a variable duration that may extend from seconds to months or longer, depending on the research question (Schnyder et al., 2012, 2017). For the question at hand here, which concerns the role of stores, dynamic labelling experiments are considered more suitable, as long-term stores are labelled only weakly in short-term pulse-labelling experiments and, consequently, the redistribution of the storage derived C compounds during the chase period is associated with a strongly weakened isotopic signal that may easily escape detection against the background noise in the isotopic baseline (Geiger & Swanson, 1965a, b; Schnyder et al., 2012). Here, I present data from 7-day long dynamic labelling experiments with the near-natural abundance ${}^{13}CO_2/{}^{12}CO_2$ labelling and gas exchange mesocosm described by Schnyder et al. (2003). This system has been used previously to study the role of stores in central carbohydrate metabolism (Lattanzi et al., 2012) or respiration (Lehmeier et al., 2008; 2010a, b) of L. perenne. This technique uses natural and relatively inexpensive sources of CO₂: mineral CO₂ and fossil (or organic) CO₂ derived from industrial processes (Schnyder, 1992). The abundance of ${}^{13}C$ ($\delta^{13}C$) is conventionally expressed as the deviation of the isotopic composition of a sample from that of the universal standard, which is carbon in carbon dioxide generated from a fossil belemnite from the Pee Dee Formation, denoted PDB (Farquhar et al., 1989) or nowadays IAEA Vienna-PDB (V-PDB), defined thus: $\delta^{13}C = R_P/R_S - 1$, with R the

molar abundance ratio ${}^{13}C/{}^{12}C$ as measured by the mass spectrometer, *P* referring to the sample and *S* to the V-PDB standard. As variations of $\delta^{13}C$ are extremely small in natural systems, the $\delta^{13}C$ values are commonly expressed in permil (‰).

The δ^{13} C of the CO₂ sources (δ^{13} C_{CO2}) used in experiments have ranged preferably in the region of -3‰ (mineral) or -47‰ (organic) (Schnyder et al., 2003), similar to this thesis. The precision (standard deviation of repeated measurements) of the ¹³C analysis of plant material or CO₂ in air is typically better than 0.3‰ (e.g. Ostler et al., 2016), providing an isotopic 'signal-to-noise ratio' (SNR) of the labelling system of >100 when SNR is defined as the ratio of the isotopic discrepancy between the contrasting CO₂ sources and the measurement precision for repeated measurements (e.g. SNR is calculated by 44‰ / 0.3‰, for the example cited above).



Fig. 1.1 Scheme illustrating the 'switching CO₂ sources' strategy for dynamic ${}^{13}CO_2/{}^{12}CO_2$ labelling of plant stands in growth chambers at the laboratory of Grünlandlehre of TUM (from Schnyder et al., 2017). Identical plant stands are established in two growth chambers with the same environmental conditions, but different $\delta^{13}C$ in the CO₂ ($\delta^{13}C_{CO2}$) provided to the chambers during the pre-labelling period (see figure insets). The CO₂ sources are switched at a given time, starting the dynamic labelling in both chambers. Individual plants can be sampled prior to and at intervals during the dynamic labelling period to analyze the tracer kinetics in plant parts or compounds of interest (see text). In parallel, the flux of air through the chambers, the CO₂ concentration and $\delta^{13}C_{CO2}$ entering and leaving the chamber are controlled and monitored, enabling the determination of the $\delta^{13}C$ of CO₂ exchanged in the light or respired in the dark.

For labelling plant stands, the laboratory of Grünlandlehre at TUM has typically applied a CO₂ source-switching strategy (Fig. 1.1), where identical plant stands are grown with the same environmental conditions in two growth chambers supplied with different $\delta^{13}C_{CO2}$ (Schnyder et al., 2003). Labelling is initiated by switching the isotopically contrasting CO₂ sources ($\delta^{13}C_{CO2 \text{ mineral}} \Leftrightarrow \delta^{13}C_{CO2 \text{ organic}}$) in parallel chambers. Thus, the chamber receiving mineral CO₂ is switched to organic CO₂, and *vice versa*, at the beginning of labelling. Plants are sampled at the time of (or just prior to) the switch and at intervals during the dynamic labelling for subsequent C isotope analysis of plant parts or compounds of interest (e.g. sucrose; Lattanzi et al., 2012). At the same time, the flux of air through the chambers (controlled by mass flow controllers) and the CO₂ concentration (measured by an infrared gas analyzer) and $\delta^{13}C_{CO2}$ (determined by on-line isotope ratio mass spectrometry) entering and leaving the chamber are used to calculate CO₂ exchange rates and the $\delta^{13}C$ of CO₂ exchanged during the light and dark periods (Schnyder et al., 2003).

The fraction of unlabelled C ($f_{unlabelled}$, with $f_{unlabelled} = 1 - f_{labelled}$, the labelled fraction) in a given entity of interest (C pool or CO₂ flux component, e.g. respiration) is obtained by a two-member mixing model as:

$$f_{\text{unlabelled}} = \left(\delta^{13} C_{\text{sample}} - \delta^{13} C_{\text{new}}\right) / \left(\delta^{13} C_{\text{old}} - \delta^{13} C_{\text{new}}\right)$$
Eqn 1.1

where $\delta^{13}C_{sample}$ equals the $\delta^{13}C$ of a specific sample, and $\delta^{13}C_{new}$ and $\delta^{13}C_{old}$ are the endmembers of the equation given by the C fixed in the presence of the 'new' (labelling) and 'old' (pre-labelling) CO₂ for the same sample type, respectively (Schnyder et al., 2003). $\delta^{13}C_{new}$ and $\delta^{13}C_{old}$ are typically determined in samples from plants constantly maintained in the presence of the 'new' or 'old' CO₂. However, experimenters often have only one labelling chamber. In that case, the 'new' end-member is estimated from: (1) knowledge of ¹³C discrimination of a specific sample ($\Delta^{13}C$) as observed in the presence of the 'old' CO₂ (Farquhar et al., 1989):

$$\Delta^{13}C = (\delta^{13}C_{CO2 \text{ old}} - \delta^{13}C_{old}) / (1 + \delta^{13}C_{old})$$
 Eqn 1.2

and (2) knowledge of the $\delta^{13}C_{CO2}$ of the new, i.e. labelling CO₂ ($\delta^{13}C_{CO2 new}$) as per Schnyder, (1992):

$$\delta^{13}C_{\text{new}} = \left(\delta^{13}C_{\text{CO2 new}} - \Delta^{13}C\right) / (1 + \Delta^{13}C)$$
Eqn 1.3

The one-chamber approach uses the fact that Δ^{13} C is independent of δ^{13} C_{CO2}, as is true based on theory (Farquhar et al., 1989), but also requires that δ^{13} C_{CO2 new} and δ^{13} C_{CO2 old} (as measured for the tank CO₂ gases) are not negatively affected by experimental artefacts in the labelling compartments (e.g. the plant growth chamber) such as by contamination with extraneous CO₂. A critical condition for a quantitative evaluation of tracer data is the virtual constancy of the isotopic composition of CO₂ in the chamber air in the different phases of the experiment. For that reason, it is useful to monitor and control these parameters during the experiment (Schnyder et al., 2003). If experimenters have one parallel growth chamber (two-chamber approach), $\delta^{13}C_{new}$ is taken directly from $\delta^{13}C_{old}$, where $\delta^{13}C_{old}$ in one chamber serves as $\delta^{13}C_{new}$ in its counterpart and *vice versa* (Lehmeier et al., 2010a; Lattanzi et al., 2012). The estimate of $f_{unlabelled}$ via the two-chamber approach is independent of experimental artefacts with the assumption that contamination effects on $\delta^{13}C_{sample}$ and $\delta^{13}C_{old}$ are the same in the parallel chambers with the same growth conditions and practical operation.

Notably, there are systematic but often unknown differences between the δ^{13} C of different C pools or metabolites, even if $\delta^{13}C_{CO2}$ of the CO₂ provided to a chamber is kept constant. This is due to variation of photosynthetic (Farquhar et al., 1989) and post-photosynthetic C isotope discrimination (Cernusak et al., 2009). Generally, however, plant C is variably depleted in ¹³C due to photosynthetic ¹³C discrimination, Δ^{13} C, the primary determinant of overall Δ^{13} C, with a variation of Δ^{13} C as a function of growth conditions and genotype- or functional group-specific biological makeup (Farquhar et al., 1989).

Instantaneous Δ^{13} C – the Δ^{13} C of C fixed on a given moment in time – can be determined from on-line 13 CO₂/ 12 CO₂ gas exchange measurements (Evans et al., 1986) as:

$$\Delta^{13}C = \left[\xi \left(\delta^{13}C_{CO2 \text{ outlet}} - \delta^{13}C_{CO2 \text{ inlet}}\right)\right] / \left[1 + \delta^{13}C_{CO2 \text{ outlet}} - \xi \left(\delta^{13}C_{CO2 \text{ outlet}} - \delta^{13}C_{CO2 \text{ inlet}}\right)\right]$$
Eqn 1.4

with $\delta^{13}C_{CO2 \text{ outlet}}$ and $\delta^{13}C_{CO2 \text{ inlet}}$ the $\delta^{13}C_{CO2}$ at the outlet and inlet of the growth chamber, and

$$\xi = C_{\text{inlet}} / (C_{\text{inlet}} - C_{\text{outlet}})$$
 Eqn 1.5

with C_{inlet} and C_{outlet} the CO₂ concentration at the inlet and outlet of the chamber, respectively, corrected to standard humidity (i.e. 0). For the plant growth chambers of the Lehrstuhl für Grünlandlehre, C_{outlet} and $\delta^{13}C_{\text{CO2 outlet}}$ are very near to the conditions experienced by plants (Schnyder et al., 2003) due to the strong ventilation of air inside the growth chambers.

Importantly, true Δ^{13} C is independent of δ^{13} C_{CO2} (Farquhar et al., 1989; Schnyder, 1992). However, (true) Δ^{13} C may vary by several thousandths between environments (Cernusak et al., 2013), genotypes (Farquhar & Richards, 1984; Yang et al., 2016), plant parts (Schnyder, 1992) and biochemical compounds (Gleixneret al., 1993; Tcherkez et al., 2011). Environmental and genotypic variation of Δ^{13} C is valuable physiological information for C₃ plants, as it correlates closely with intrinsic water-use efficiency (iWUE), the ratio of photosynthesis (*A*) to stomatal conductance (g_s): iWUE = A / g_s (Farquhar et al., 1989; Yang et al., 2016). However, differences in Δ^{13} C between plant parts or biochemical compounds are still not fully understood (Cernusak et al., 2009). Whatever the cause, variation of Δ^{13} C, including via experimental artefacts, must be considered in labelling data evaluation, as it directly affects the isotopic end-members (δ^{13} C_{new} and δ^{13} C_{old}), as seen above. Therefore, organ- and compound-specific Δ^{13} C values (Schnyder, 1992) should be used to calculate *f*_{unlabelled} (Eqn 1.1).

Contamination by extraneous ${}^{13}CO_2/{}^{12}CO_2$ (Gong et al., 2015) is a paramount concern, as it may affect all members of the equation, i.e. $\delta^{13}C_{\text{sample}}$, $\delta^{13}C_{\text{new}}$, and $\delta^{13}C_{\text{old}}$. Contamination by extraneous ${}^{13}\text{CO}_2/{}^{12}\text{CO}_2$ may occur through gas leaks (Gong et al., 2015) in the growth chamber or during the opening of the chambers for plant sampling or other works requiring access to the growth compartment. Such an effect can be explained by the incursion of a mixture of CO₂ from the free atmosphere (δ^{13} C ~ -10‰) and CO₂ exhaled (δ^{13} C ~ -23‰; Epstein & Zeiri, 1988) by people (e.g. experimenters) in the room housing the plant growth chambers. Contamination with extraneous ${}^{13}CO_2/{}^{12}CO_2$ can be minimized by maintaining a slightly positive atmospheric pressure inside relative to outside of the chambers (Schnyder et al., 2003) and by placing air-locks in chamber doors (Lehmeier et al., 2008). Eventual contamination with extraneous (organic) C sources during the handling and processing of samples is another factor that should be considered in labelling experiments. Given that all plant samples are handled and processed in identical ways, any contamination would reduce the isotopic spread between $\delta^{13}C_{new}$ and $\delta^{13}C_{old}$. In a theoretical, although extreme scenario, where the contaminated samples are composed entirely of the contaminant, the (artefactual) estimate of $\delta^{13}C_{new}$ would equal that of $\delta^{13}C_{old}$. Such extreme cases are most likely for tiny samples, where contamination would cause an extreme 'dilution' of the true (original) C isotope signal in the sample. Again, such artefacts would cause an (erroneous) effect on $\delta^{13}C_{CO2}$ on Δ^{13} C, generating errors in tracer data evaluation. In principle, one should expect that tracer experiments at different [CO₂] would be particularly susceptible to such artefacts. Except for the investigation of leak problems with clamp-on leaf chamber ${}^{13}CO_2/{}^{12}CO_2$ exchange measurements by Gong et al. (2015), however, this problem (termed 'isotopic end-member problem' in the following) has not been examined systematically, to the best of my knowledge.

1.4 Compartmental modelling of the respiratory substrate supply system and central carbohydrate metabolism

Compartmental modelling is a mathematical tool for interpreting the tracer time course in a particular entity, e.g. sucrose or respired CO₂, in terms of the metabolic system that generates it (Atkins, 1969; Farrar & Farrar, 1986; Jacquez, 1996; Lattanzi et al., 2012; Schnyder et al., 2012, 2017). Basically, the compartmental model characterises the system, for instance, the respiratory substrate supply system, by the number of kinetically distinct pools, the connections between the pools, the residence time of the tracer in each pool, the size of each pool, the fluxes of material (tracer) through each pool and the environment (including any parts exterior to the system under investigation). Commonly, it is assumed that pools are well-mixed, thus obeying first-order kinetics, with practical support from many systems (e.g. Lattanzi et al. 2005). This means that the time course of the tracer in the parts (pools) of the system can be described by a set of differential equations. Another assumption is that the system is in a steady state, namely that pool sizes and fluxes are constant. In the strict sense, that condition is rarely met, even if plants are grown in controlled conditions. However, the condition may be approximate sufficiently closely, even with plants grown in day-night cycles, when analysing data on a dayby-day timescale over limited periods of time, as shown by Lattanzi et al. (2005, 2012), Lehmeier et al. (2010a), and Gong et al. (2017).

Compartmental models should provide a biologically meaningful description of the system under investigation (Schnyder et al., 2012). At the same time, a compartmental model should be parsimonious, that is, it should not be more complex than is necessary for an unbiased representation of the tracer data. Lehmeier et al. (2008) analysed the tracer time courses in respired CO₂ (i.e. the temporal changes of $f_{unlabelled}$ over time) of shoots of perennial ryegrass grown in continuous light and labelled for periods of 1 h to 25 d and detected three kinetically distinct phases that indicated the existence of a respiratory substrate supply system composed of three pools, with different half-lives (<0.2 h, ~3 h, and 33 h) and sizes (equivalent to 0.2, 10 and 70 mg g⁻¹ biomass-C). Interestingly, the tracer kinetics of the roots were (1) very similar to that for the shoots, but (2) exhibited a temporal delay in tracer appearance that was consistent with assumptions for the time needed for transport of the substrate of respiration from the shoot to the root (~0.8 h), and (3) indicated a total size of the respiratory substrate supply system supporting root respiration that was several times larger than the total quantity of non-structural, i.e. metabolizable C present in the roots. Together, these features (1-3) indicated that at least

Chapter 1 Introduction

the largest and most slowly turning over pool supporting root respiration must have been located outside the root system, i.e. in the shoot. These results revealed new biological insight into the biological system supporting respiration obtained by combining quantitative tracer application and compartmental modelling. A simplified version of that model, that is, a twopool model (inset in Fig. 1.2a) that ignored the presence of the most rapidly turned-over and smallest pool in the system, was subsequently used to study the role of long-term stores in the respiratory substrate supply system of perennial ryegrass grown in continuous light or regular day-night cycles (Lehmeier et al., 2010a; Fig. 1.2) and of $[CO_2]$ in sunflower (Gong et al., 2017).



Fig. 1.2 Time course of the fraction of unlabelled C ($f_{unlabelled}$) in dark-respired CO₂ of shoots (a) and roots (b) during dynamic ¹³CO₂/¹²CO₂ labelling of *L. perenne* grown in continuous light (closed symbols) or a 16:8 h day:night regime (open symbols) (from Lehmeier et al., 2010a). Lines denote the predictions of the two-pool model shown as an inset in (a) for plants grown in continuous light (solid lines) or in the 16:8 h day:night cycle (dashed lines). For details, see Lehmeier et al. (2010a).

Lattanzi et al. (2012) devised a four-pool compartmental model of central carbohydrate metabolism for a photosynthetically active leaf of a C_3 grass (Fig. 1.3). Among other things, that model was used to study the effect of nitrogen nutrition on the role of fructan metabolism in the recycling (re-synthesis) of sucrose. In that model, (1) current photosynthesis supplied central carbohydrate metabolism *via* the sucrose pool, (2) sucrose was consumed in three types of processes: export, fructan synthesis (with the fructosyl moiety of a donor sucrose molecule transferred to an acceptor sucrose or fructan molecule, and simultaneous release of one glucose

Chapter 1 Introduction

molecule), and sucrose hydrolysis by invertase-like processes (generating equimolar amounts of fructose and glucose), (3) fructan hydrolysis produced fructose, (4) fructan turnover in a steady-state yielded equimolar amounts of fructose (released during fructan hydrolysis) and glucose (released during the synthesis step, see above), and (5) all fructose and glucose was (re)utilized in sucrose (re)synthesis. This model provided a virtually perfect fit (R^2 =0.97) to the combined tracer kinetics in sucrose, fructan, glucose, and fructose (Fig. 1.4), supporting the basic assumptions of compartmental analysis, as outlined above. In particular, the model fit to the data also supported stoichiometric constraints, i.e. that equimolar amounts of glucose were produced per fructosyl unit transferred to the fructan pool, and equimolar use of fructose and glucose in sucrose re-synthesis related to invertase activity and fructan metabolism. In other words, in a statistical sense, the stoichiometrically-constrained four-pool model (Fig. 1.3) performed equally well or better than more complex models (Lattanzi et al., 2012). To the best of my knowledge, the approach of Lattanzi et al. (2012) (or equivalent) has not been applied to other research questions, such as scaling up to the whole plant level or investigations of the effect of [CO₂] on the role of fructan metabolism in sucrose recycling.



Fig. 1.3 Four-pool compartmental model of central carbohydrate metabolism in source leaves of *L. perenne* (from Lattanzi et al., 2012). Suc, sucrose; Glc, glucose; Fru, fructose; F_{in} , tracer flux into the system. Q_i represents the size of the carbohydrate pool *i* (Q_{1-4} mean sucrose, fructan, glucose, and fructose pools, respectively), and k_{ij} is the rate constant for the flux from pool *i* to pool *j*. Thus, k_{10} denotes the export of sucrose from the system, k_{12} the fructosyl transferase-catalysed transfer of the fructosyl residue of donor sucrose to a fructan (or sucrose) acceptor molecule, k_{13} glucose production by fructosyl transferase *plus* invertase(-like) activities, k_{14} fructose production by invertase(-like) activity, k_{24} cleavage of fructose from fructan by fructan exohydrolase, k_{31} glucose use in sucrose resynthesis, and k_{41} fructose use in sucrose resynthesis. For mathematical details of the model, including the differential equations describing the fluxes, see Lattanzi et al. (2012).



Fig. 1.4 The fraction of unlabelled C (funlabelled) in fructan (filled squares), sucrose (open circles), glucose (filled triangles), and fructose (open triangles) with labelling duration in the youngest fully expanded leaf blade of *L. perenne* grown at low N (upper panel) and high N supply (lower panel) (from Lattanzi et al., 2012). The insets expand the first 24 h of the dynamic labelling. The curves represent the fits of the four-pool model shown in Fig. 1.3. Carbohydrates were extracted from the blade of the youngest fully expanded leaf of perennial ryegrass grown with either 1 mM nitrate (low N) or 7.5 mM nitrate (high N) in the nutrient solution. Plants were labelled during normal growth in swards kept in ¹³CO₂/¹²CO₂ mesocosms. In both treatments, swards grew in continuous light with a photosynthetic photon flux density of 275 µmol mol⁻¹at 20 °C air temperature and 85% relative humidity. For details on carbohydrate extraction, separation and isotope analysis, see Lattanzi et al. (2012).

1.5 Aim and outline of the thesis

This thesis aims to explore the effects of $[CO_2]$ on the role of stores for recycling sucrose in whole shoots and supplying substrate to respiration of intact plants or stands of perennial ryegrass (*L. perenne*, cv Acento) by use of a modernized version of the mesocosm-scale nearnatural abundance ${}^{13}CO_2/{}^{12}CO_2$ dynamic labelling technique developed at the former TUM Lehrstuhl für Grünlandlehre and described by Schnyder et al. (2003). Perennial ryegrass was chosen as a model perennial C₃ grass of high economic value (Chapman et al., 2017), whose biology, breeding, and agronomy have been studied comparatively intensively in the past. The cultivar Acento was selected, as it had already been the subject of previous physiological studies at the TUM Lehrstuhl für Grünlandlehre (e.g. Lehmeier et al., 2008). My work was based on mesocosm-scale tracer experiments that were performed subsequent to extensive experimental works with the same plant stands during the (pre-labelling) establishment phase (see chapter 3.2 and Baca et al., 2020). As these works might have caused inadvertent ¹³CO₂/¹²CO₂ contamination issues, my work also included investigations on basic methodical assumptions of tracer data evaluation.

Thus, my work addressed the following specific questions and objectives (addressed in chapter 3):

(1) Do ${}^{13}CO_2/{}^{12}CO_2$ artefacts (e.g. through gas leaks or contamination during carbohydrate extraction and separation by HPLC) affect the isotopic end-member of the twomember mixing model used to analyze the labelling kinetics of gas exchange-, biomass- and carbohydrate-components? If so, were such artefacts sensitive to [CO₂] treatments? (chapter 3.1)

(2) Does the four-pool compartmental model of leaf-level central carbohydrate metabolism of a C_3 grass proposed by Lattanzi et al. (2012) also apply to the whole shoot level? Moreover, if so, does [CO₂] affect the role of the fructan store in sucrose recycling? (chapter 3.2)

(3) Does [CO₂] affect the importance of the store in supplying substrate for whole plant respiration? (chapter 3.3)

(4) Does a comparative analysis of the tracer kinetics of carbohydrate components (at shoot level) and whole plant respiration provide clues to the identity of substrates fueling dark respiration? (chapter 3.4).

Perennial ryegrass plants (*L. perenne*) were grown in dense swards in growth chambers at three different [CO₂] levels (treatments): 'half ambient' (approximately equivalent to the [CO₂] at the Last Glacial Maximum), current 'ambient', and 'double ambient' (as projected for the end of this century, IPCC, 2007). Stands in parallel (paired) plant growth chambers were grown in the presence of fossil-organic (industrial) CO₂ (δ^{13} C -43.5‰ as determined for the CO₂ at chamber inlet) and CO₂ from a mineral source (δ^{13} C -5.6‰). The swards were grown using a hydroponic culture system with optimal water supply but reduced nitrogen concentration in the standard nutrient solution (Kavanová et al., 2008; Lehmeier et al., 2008; 2013; Baca et al., 2020; see chapter 2).

In addition, in this thesis, I have included as Appendices two publications in which I am a coauthor. These formed part of the publication-based thesis of Juan C Baca Cabrera ("The key role of stomatal conductance in controlling the grassland vegetation response to a changing environment" passed with distinction in 2022) and were performed in the same experiments as

Chapter 1 Introduction

my studies. I contributed experimentally to these works by performing leaf-level gas exchange measurements (photosynthesis, stomatal conductance, and transpiration), assisting in the sampling of plant material, and carbohydrate analysis, and maintaining the experimental facility. Also, I contributed to the discussion of these works, as acknowledged in the author's contributions of these papers. These include one paper published in *New Phytologist* (Baca et al., 2020) and a preprint published in *ResearchSquare* (Baca et al., 2021). However, the specific topics of these papers are outside the scope of my thesis. Therefore, I did not include these papers in my thesis, although I cite them for some methodical details, which were common also to my work, and findings of interest to the interpretation of my results.

2. Materials and Methods

2.1 Experiment design

The treatments presented in this thesis formed part of a 3×2 factorial experiment with three atmospheric CO₂ concentrations ([CO₂]: 200, 400, and 800 µmol mol⁻¹) corresponding to the Last Glacial Maximum (LGM; sub-ambient), current ambient, and end of the century projections (double-ambient), and two air relative humidity (RH) levels of 50% and 75% during day-time. Night-time RH was 75% in all treatments. A total of five sequential experimental runs were performed using four growth chambers (Table 2.1). My thesis research was limited to the (day-time) 50% RH treatment in the first two runs. A day-time RH of 50% is a typical scenario for local temperate grassland (Hirl et al., 2019). Treatment replicates (*n*=3-5) were allocated to different chambers between sequential runs to minimize chamber effects. Plant management details were the same in all experimental runs. However, I observed no chamber effect on the parameters reported in this study (see also Baca et al., 2020).

Table 2.1 Experiment with a 3 x 2 experimental design: three $[CO_2]$: 200, 400 or 800 µmol mol⁻¹ (C1, C2 or C3) and two RH levels: 50% or 75% during daytime. Nighttime RH was 75%, the same in all treatments. The treatment replicates were distributed between four growth chambers (no. 1-4). The corresponding treatments in this thesis are given in bold.

			Trea	atment		
Run	C1 (50%)	C2 (50%)	C3 (50%)	C1 (75%)	C2 (75%)	C3 (75%)
Chamber no.						
1 st	1/2	3 / 4	-	-	-	-
2 nd	-	1/2	3 / 4	-	-	-
3 rd	-	-	-	3/4	1/2	-
4 th	-	3	-	-	4	1/2
5 th	3	-	1	2	-	4

Plants of *L. perenne* were established and grown singly in individual plastic pot (350 mm height, 50 mm diameter) filled with 800 g of washed quartz sand (0.3-0.8 mm grain size). Pots were arranged at a density of 383 plants m⁻² in plastic containers ($770 \times 560 \times 300$ mm), and two of such containers were placed in each growth chamber (PGR15, Conviron, Winnipeg,

Canada). Plants were supplied with a modified 5 mM nitrate-N Hoagland nutrient solution with a hydroponic culture system (Fig. 2.1). This nitrate-nitrogen (N) concentration corresponded to a nutrient solution with a 33% reduced nitrate-N concentration relative to normal and nominal concentrations of the other nutrients (compared to Kavanová et al., 2008). Thus, the nutrient solution contained 1.7 mM Ca(NO₃)₂, 1.7 mM KNO₃, 1.0 mM MgSO₄, 0.5 mM KH₂PO₄, 0.5 mM NaCl, 0.125 mM iron as EDTA, 46.1 μ M H₃BO₃, 9.1 μ M MnSO₄, 0.8 μ M ZnSO₄, 0.3 μ M CuSO₄, and 0.15 μ M Na₂MoO₄. A nutrient solution with reduced nitrate-N concentration was chosen as most terrestrial ecosystems, including temperate grassland, are typically N-limited (LeBauer & Treseder, 2008; Terrer et al., 2019; Baca et al., 2020).



Fig. 2.1 Schematic representation of a flood-tide hydroponic culture system used in the present study. Nutrient solution (for composition, see text) was contained in a reservoir (size: $1770 \times 690 \times 520$ mm; CEMO, Rems-Murr, Germany) close to the growth chamber (PGR15, Conviron, Winnipeg, Canada), a water pump (JP5BBCVBP, Grundfos, Bjerringbro, Denmark), and a timer (AX-801A, Duwi, Hagen, Germany) which activated the irrigation cycle every six hours. After the pot/container was irrigated for nine minutes, gravity drained the solution back down into the reservoir for *c.* 45 minutes.

Growth chambers were operated with day/night cycles, with a 16 h-long photoperiod and an 8 h-long dark period. The light was supplied by cool-white fluorescent tubes and warmwhite LED bulbs. Irradiance was measured at regular intervals with a quantum sensor (Li-190R, LI-COR Inc., Lincoln, NE, USA), and the distance between the light source and the canopy was adjusted so that a photosynthetic photon flux density of 800 μ mol m⁻² s⁻¹ was maintained at canopy height. The temperature of chamber air was held at 20 °C during the photoperiod and at 16 °C during darkness. RH of the chamber air was kept near the nominal level (SD ± 0.9%) by using a high-pressure water vapour generator (FINESTFOG, Ottobrunn, Germany) that added a known amount of water vapour to the chamber air whenever the nominal RH dropped below a specified threshold (1-2% RH lower than the chamber setpoint). The chamber-based humidification system was inactivated during that period. Air temperature and RH in the chamber were measured continuously with the chamber control system (CMP6050, Conviron, Winnipeg, Canada).

Chapter 2 Materials and Methods

The target $[CO_2]$ inside the chamber was obtained by mixing dry CO₂-free air and tank CO₂ of known carbon isotope composition ($\delta^{13}C$, with $\delta^{13}C = [({}^{13}C/{}^{12}C)_{sample} / ({}^{13}C/{}^{12}C)_{VPDB}$ standard] - 1) (Linde AG, Unterschleißheim, Germany or CARBO Kohlensäurewerke, Bad Hönningen, Germany) using mass flow controllers. Each [CO₂] treatment had two growth chamber replicates, one replicate chamber was supplied with ¹³C-organic CO₂ (δ^{13} C, -43.5‰) while the other replicate chamber received ¹³C-mineral CO₂ (δ^{13} C, -5.6‰). The δ^{13} C of supplied CO₂ was determined at the chamber inlet with a precision (SD) of <0.17‰ for repeated measurements. The airflow and CO₂ concentration of the chamber inlet air were periodically adjusted. The rate of CO₂ supply to the chamber exceeded the CO₂ exchange rate of the stands in light by a factor of 10. This minimised the effects of photosynthesis, respiration, and recycling of respiratory CO₂ on the δ^{13} C of chamber CO₂, so that the growing stands always experienced closely similar δ^{13} C and target [CO₂] in the growth chamber air during the photoperiod. One tank typically contained c. 30 kg of CO₂, so one experiment employed two ¹³C-organic CO₂ tanks and two ¹³C-mineral CO₂ tanks. The δ^{13} C of tank CO₂ was tested before each experimental run and was typically very similar (<0.27‰ SD) between tanks of the same type of CO₂ source (mineral or fossil-organic). The CO₂ concentrations and $\delta^{13}C_{CO2}$ measured in the chamber outlet air represented the growth chamber atmosphere (Schnyder et al., 2003).

Disturbance of air conditions in the chambers during handling of plants was minimized by maintaining *c*. 200 Pa positive atmospheric pressure inside relative to the outside of the chambers (Schnyder et al., 2003) and installing air-locks in chamber doors (Lehmeier et al., 2008) as shown in Fig. 2.2. Experimental operations, which included chamber maintenance activities and plant sampling, carried out with opening growth chamber doors, may have nevertheless resulted in some disturbance of growth chamber air CO₂ and $\delta^{13}C_{CO2}$. The total time that the chamber was open for one entire experiment during daylight was approximately 110 minutes. This time included a total of 20 minutes spent changing the broken light and 90 minutes spent sampling and handling the plants approximately 20 times (see next paragraph).



Fig. 2.2 Picture of air-lock. One air-lock consists of two PVC plates (770 \times 460 mm) and two plastic sheets (570 \times 400 mm). Two PVC plates were installed above and below the door. The middle area was left for the operation window, covered with two plastic sheets. The air-lock covered the growth chamber door completely.

2.2 Experiment timeline

The experiment was set up based on the plant growth stage, as shown in Table 2.2. The first irrigation of the sown seeds, which signifies the initiation of seed imbibition, was considered the start of the experiment (Day 0). The seedling development (Day 0 - 12), plant canopy expansion (Day 13 - 48), and full canopy (Day 49 - 72) are designated the three phases of a complete experiment (10 weeks).

Growth chambers were supplied with ambient air during the seedling development phase (until day 12). RH was maintained at 80% in both the light and dark periods. The air temperature was kept at target levels. Plants were irrigated with nutrient solution for the whole day. [CO₂] and RH treatments were implemented on day 13, in the canopy expansion phase. Plants were irrigated with the nutrient solution every six hours. In that: nutrient solution entered the plastic containers holding the plants for nine minutes and then drained back down into the reservoir over *c*. 45 minutes by gravity. The nutrient solution was renewed every three weeks throughout the duration of each experiment, in all experimental runs. For gas measurements in the chamber inlet or outlet, the concentration of CO₂ was measured from day 13 and δ^{13} C from day 20. All experiment operations were conducted in the full canopy phase. This experimental phase included two 2-d long intensive plant sampling periods, starting on day 49 and day 63, leaf-scale gas exchange measurements from day 49 to day 60, plant leaf temperature measurements from day 49 to day 63, leaf-elongation measurements from day 49 to day 62, and sampling of leaf morphometry measurements on day 64. The labelling experiments (and associated sampling activities) reported in this thesis were started between day 65 and day 72.

Chapter 2 Materials and Methods

Table. 2.2 Timeline of one experiment. Day *X*, designates time during an experimental run, with time 0 defined as when the sown seed was first imbibed. The experiment was started on day 0 (refers to seed imbibition) and ended on day 72. The period consisted of the seedling development phrase, canopy expansion phase, and full canopy phase. The labelling experiment was conducted during the full canopy phase.

Experiment timeline			
Time	Stage description	Details	
Day 0-12	Seedling	Three seeds were sown per plastic tube;	
	development	One week after seed imbibition, plants were reduced to one	
		per tube (Fig. 2.3 A).	
Day 13-48	Plant canopy	On day 13, canopy-scale gas exchange measurement was	
	expansion	started;	
		On day 20, carbon isotopes measurements on chamber	
		gas exchange were started.	
Day 49-72	Full plant canopy	A series of measurements were performed during this	
		period, including two plant sampling phases starting on day	
		49 and day 63, leaf-scale gas exchange, leaf temperature,	
		leaf-elongation (Baca et al., 2020) and leaf morphometry	
		measurements (see also Fig. 2.3 B).	
Day 65-72	Labelling	Dynamic ${}^{13}CO_2/{}^{12}CO_2$ labelling of stand-scale dark respired	
	experiment	CO2 and whole-shoot water-soluble carbohydrate fractions	
		(fructan, sucrose, glucose and fructose).	



Fig. 2.3 Typical stages of plant and stand development during the experiment. A: Arrangement of plants in the growth chamber on day 18. B: Dense canopies had developed by day 49. C: Individual plants removed from the stands on day 65. Treatments did not exhibit differences in development rate or leaf dimensions (Baca et al., 2020). Also, all canopies were closed with a leaf area index >5.5 at the time of harvests (Baca et al., 2020).

2.3 Gas exchange measurement and isotope analysis

The experiment used a new version of the experimental infrastructure described by Schnyder et al. (2003; see also Baca et al., 2020) shown in Fig. 2.3, which consisted of four plant growth chambers. Sample air was collected at the inlet and outlet of each growth chamber and passed on to an infrared gas analyser (Li-840, LI-COR Inc., Lincoln, NE, USA) for measuring CO₂ and H₂O concentrations and to an isotope ratio mass spectrometer (IRMS, Delta plus, Advantage equipped with GasBench II, ThermoFinnigan, Bremen, Germany) for measuring "on-line" δ^{13} C of the sample air. Air for mass spectrometric analysis was pumped continuously via a steel capillary from the gas sampling system to a 0.25 mL sample loop which was attached to an eight-port Valco valve of the GasBench II. Sample air in the loop was introduced into the IRMS via an open split after the passage of a dryer (Nafion) and a GC column (25 m × 0.32 mm Poraplot Q; Chrompack, Middleburg, the Netherlands). After every second sample, a VPDB-gauged CO₂ reference gas was injected into the IRMS via the open split. The precision (SD) of repeated measurements was <0.2‰. The entire measurement cycle of four growth chambers was completed within 30 minutes, including measurement of CO₂ and H₂O concentration and δ^{13} C of the sample air on the inlet and outlet of each growth chamber.


Fig. 2.4 Schematic representation of ¹³CO₂/¹²CO₂ gas exchange and labelling system (A; from Schnyder et al., 2003) and growth chamber picture (B). SC, screw compressor (S40; Boge, Bielefeld, Germany); AD, adsorption dryer (KEN 3100; Zander, Essen, Germany; and molecular sieve: activated aluminium oxide F200; Alcoa, Houston, TX, USA); AR, air receiver (1 m³) (Magnet Kft, Magocs, Hungary); F1, oil and water condensate drain (CSP005; Hiross, Mönchengladbach, Germany), F2 oil, water and particle filter (≥0.01 µm; G12XD and filter element: 2030X, Zander), F3, universal filter (≥1 µm; G12ZHD and filter element: 2030Z, Zander); GMS, gas mixing system consisting of CO₂ flow controller (Red-y, Vögtlin, Muttenz, Switzerland) and air flow controller (EL-FLOW, Bronkhorst, Veenendaal, Netherlands); CO₂, tank containing CO₂ of mineral or fossilorganic origin (Linde AG, Unterschleißheim, Germany or CARBO Kohlensäurewerke, Bad Hönningen, Germany); Growth chamber (PGR15, Conviron, Winnipeg, Canada); SAS, sample air selector (Type 6012, Bürkert, Ingelfingen, Germany); IRGA, CO₂ and H₂O infrared gas analyser (Li-840, LI-COR Inc., Lincoln, NE, USA); IRMS, ¹³CO₂/¹²CO₂ isotope ratio mass spectrometer (IRMS, Delta plus, Advantage equipped with GasBench II, ThermoFinnigan, Bremen, Germany); PC, central control and data acquisition systems (DMP). For more details on the design and its operation, see Schnyder et al. (2003).

2.4¹³C-labelling and sampling

On day 66, in each experiment, ¹³C-labelling of the total photosynthetic carbon flux of all canopies was initiated by switching the CO₂ sources: a growth chamber that had received the ¹³C-rich CO₂ henceforth received ¹³C-depleted CO₂ and *vice versa* (see Fig. 1.1, chapter 1). The switch occurred during the night of day 65, at approximately three hours before the onset of the light period of the next day. This procedure ensured that virtually all of the isotopically "old" (i.e. "pre-switch") CO₂ had been entirely replaced by the "new" CO₂ at the beginning of the 1st labelling light period. The labelling period lasted for a total duration of 8 days.

Chapter 2 Materials and Methods

Plants were collected at the beginning of the dark period (Fig. 2.3 C), just before the switch of the CO₂ source that initiated the ${}^{13}CO_2/{}^{12}CO_2$ labelling, and then at the beginning of the dark periods following the 1st, 2nd, 3rd, 4th and 7th photoperiod after the CO₂ switch. Two replicate samples from each growth chamber were collected at each harvest, with one replicate consisting of three randomly selected plants. Plants were removed from the pots, their roots washed to free them of sand, and dissected into their shoot and root parts. The plant parts were weighed to determine their fresh weight, then frozen in liquid nitrogen and stored at -18 °C before freeze-drying for 72 h. Dry weights were subsequently determined. After that, plant material was ground to a fine powder in a ball mill and stored again at -18 °C until further use.

2.5 Elemental and carbon isotopes analysis

The carbon (C) and nitrogen (N) contents and δ^{13} C of biomass samples were determined for all shoot and root replicates as in Lattanzi et al. (2005). The stored samples were thawed, re-dried at 40 °C for 24 h and stored in desiccator vessels. Aliquots of 0.70 ± 0.05 mg of the shoot and root materials were weighed and packed into tin cups (3.3 x 5 mm, IVA Analysentechnik, Meerbusch, Germany). They were then combusted in an elemental analyzer (NA 1110, Carlo Erba Instruments, Milan, Italy) and interfaced (Conflo III, Finnigan MAT, Bremen, Germany) to a continuous-flow isotope-ratio mass spectrometer (CF-IRMS, Delta Plus, Finnigan MAT, Bremen, Germany), which measured concentrations of carbon and nitrogen, and δ^{13} C of the sample. A solid internal laboratory standard (SILS, fine ground wheat flour) was measured as a reference after every tenth sample to correct for possible instrument drift. All samples and SILS were measured against a laboratory working standard CO₂ gas, which was previously calibrated against a secondary isotope standard (IAEA-CH6; calibration accuracy ± 0.06‰ SD). The precision given as the SD of repeated measurements of the SILS was better than 0.2‰.

The C and N masses of plants and plant parts were obtained as the product of elemental contents (g per g dry mass) of plants and plant parts times their corresponding dry mass. C content of shoot tissue was closely similar for all tissues (44.7% \pm 0.54% SD), meaning that sample dry mass was virtually proportional to C mass. For that reason and to avoid redundancy I do not present dry mass data in this work. Fresh and dry masses of root samples were somewhat contaminated by sand, as reflected in low (but variable) root C contents (average: 40.8%; range 30.1% to 45.5%). This artefact was eliminated by calculating root C mass which corrected the error of the erroneously high dry mass by the correspondingly lower C content, as determined by elemental analysis.

2.6 Water-soluble carbohydrate extraction, separation, quantification, and ¹³C analysis

Water-soluble carbohydrate (WSC) fractions (fructan, sucrose, glucose, and fructose) were extracted from shoot and root samples and separated using the procedures described by Gebbing & Schnyder (2001). Briefly, aliquots of 200 mg of milled sample material were weighed into 2-mL capped Eppendorf tubes and topped off with 1.8 mL of deionized water. Samples were briefly vortexed (Vortex-Genie 2, Scientific Industries, New York, USA), held in a water bath at 93 °C for 10 min, shaken for 45 min (Shaker, Heidolph Instruments, Schwabach, Germany) at room temperature, and centrifuged at 9500 *g* for 15 min (Universal 320, Merck, Tuttlingen, Germany). The supernatant, which contained the dissolved WSC, was passed through nylonmembrane filters with a pore size of 0.45 μ m and then stored in clean 2-mL capped Eppendorf tubes at -18 °C.

WSC fractions (fructan, sucrose, glucose and fructose) were separated, quantified and collected using a high-performance liquid chromatography (HPLC) system similar to Gebbing & Schnyder (2001). Briefly, 0.2 mL aliquots of the filtered supernatant were passed through a guard column (Shodex KS-LG, Showa Denko, Tokyo, Japan) and a preparative column (Shodex KS, Showa Denko, Tokyo, Japan) held at 50 °C, with HPLC-grade water (Carl Roth, Karlsruhe, Germany) as the eluent, at a flow rate of 0.75 mL min⁻¹ and a system pressure of approximately 21 bar. The WSC was detected by refractive index measurement (Shodex RI-101, Showa Denko, Tokyo, Japan) and concentrations quantified by comparing sample peak area against reference calibration curves of pure and mixed standards of analytical grade inulin, sucrose, glucose and fructose (all from Merck, Darmstadt, Germany). Knowing when the individual carbohydrates eluted from the preparative column (Fig. 2.4), I collected fractions of fructan, sucrose, glucose, and fructose individually in test tubes. Aliquots of approximately 0.7 mg of the different carbohydrates were then transferred to tin cups (Capsules, Lüdi Swiss AG, Flawil, Switzerland), dried at 60 °C for 24 h, and then analysed for their δ^{13} C using a continuous-flow isotope ratio mass spectrometer (CF-IRMS) as described above.

The accuracy of HPLC system separation for specific water-soluble carbohydrate was tested as by González et al. (1999): individual stand materials of fructan, sucrose, glucose, and fructose (inulin from chicory, sucrose, D-glucose and D-fructose) were prepared and measured for their δ^{13} C (Pure group). I prepared a 10 ml mixed standard solution containing inulin (413.96 mg), sucrose (25.02 mg), glucose (20.41 mg) and fructose (20.71 mg), which were similar to the concentration of WSCs in the plant, and injected 0.2 mL aliquots of mixed solution into the

HPLC system, recollected the individual carbohydrates after separation, and measured their δ^{13} C as described above (Mixture group).

I used the iodine-potassium iodide solution test to detect starch in plants. Shoot and root powders were taken from plant material of the 800 μ mol mol⁻¹ [CO₂] treatment. I placed a 0.70 mg sample on a glass slide, added two drops of iodine-potassium iodide solution, and watched the colour change with a microscope (475110, Carl Zeiss, Wetzlar, Germany) after 1 minute.



Fig. 2.5 Typical HLPC chromatogram of carbohydrates in the shoot of *L. perenne*. The fractions corresponding to the fructan, sucrose, glucose and fructose peak were separated and detected separately. The thin grey line represents the baseline. The total elution time was about 90 minutes following sample injection. The sample was taken from a plant grown in 800 µmol mol⁻¹ [CO₂].

2.7 Compartmental modelling

Four-pool model for carbohydrates (see chapter 3.2)

I used the same four-pool compartmental model (Fig. 1.2) as Lattanzi et al. (2012) based on the same set of differential equations and implemented this in ModelMaker to analyse the time course of $f_{\text{unlabelled}}$ in the different carbohydrate pools at the whole-shoot scale. Accordingly, for any carbohydrate pool Q_i (with Q_1 , Q_2 , Q_3 and Q_4 representing the size of the sucrose, fructan, glucose and fructose pools, individually), it is assumed that $dQ_i/dt = 0$. This implies

that the sum of all fluxes of substrate carbon into and out of Q_i have the same magnitude, i.e. the system was in steady-state with respect to pool sizes and fluxes at the day-by-day time-scale. The assumptions also imply that a given flux out of a donor pool *i* to a receiving pool *j* (F_{ij}) is the product of a rate constant (k_{ij}) and the size of the donor pool (Q_i) ($F_{ij} = k_{ij} Q_i$). *j*=0 denotes a flux leaving the four-pool system and reflects sucrose consumption in all growth and maintenance processes in the entire shoot. The half-life of a pool ($T_{1/2} Q_i$) is $T_{1/2} Q_i = \ln(2) / k_{In}$, with k_{In} the sum of the rate constants for all fluxes leaving pool Q_i .

The change in the amount of unlabelled carbon in pool i with time t was given by the sum of all fluxes of unlabelled carbon into that pool minus the sum of all fluxes of unlabelled carbon leaving the pool. Accordingly, for the four pools in the system (Lattanzi et al., 2012):

$$df_{\text{unlabelled Q1}}\left(t\right) / dt (t) \cdot Q_{1} = f_{\text{unlabelled In}}\left(t\right) \cdot F_{\text{In}} + f_{\text{unlabelled Q3}}\left(t\right) \cdot k_{31} \cdot Q_{3} + f_{\text{unlabelled Q4}}\left(t\right) \cdot k_{41} \cdot Q_{4} - f_{\text{unlabelled Q1}}\left(t\right) \cdot \left(k_{10} + k_{12} + k_{13} + k_{14}\right) \cdot Q_{1}$$

 $df_{\text{unlabelled Q2}}(t) / dt (t) \cdot Q_2 = f_{\text{unlabelled Q1}}(t) \cdot k_{12} \cdot Q_1 - f_{\text{unlabelled Q2}}(t) \cdot k_{24} \cdot Q_2$

Eqn 2.2

Eqn 2.1

 $df_{\text{unlabelled Q3}}(t) / dt(t) \cdot Q_3 = f_{\text{unlabelled Q1}}(t) \cdot k_{13} \cdot Q_1 - f_{\text{unlabelled Q3}}(t) \cdot k_{31} \cdot Q_3$

Eqn 2.3

 $df_{\text{unlabelled Q4}}(t) / dt(t) \cdot Q_4 = f_{\text{unlabelled Q1}}(t) \cdot k_{14} \cdot Q_1 + f_{\text{unlabelled Q2}}(t) \cdot k_{24} \cdot Q_2$

$$-f_{\text{unlabelled Q4}}(t) \cdot k_{41} \cdot Q_4$$
 Eqn 2.4

where $f_{\text{unlabelled }Qi}(t)$ denotes the fraction of unlabelled C in pool *i* at time *t*, F_{In} the flux into the system (which is equal to the rate of photosynthetic sucrose synthesis in the plant), and $f_{\text{unlabelled}}$ In (*t*) the fraction of unlabelled C in F_{In} with $f_{\text{unlabelled In}}(t) = 1$ before the switch of $\delta^{13}C_{\text{CO2}}$ (i.e. at t < 0) and $f_{\text{unlabelled In}}(t) = 0$ after the switch ($t \ge 0$).

Based on Eqns 2.1 - 2.4, the rate constants were optimized in such a way that $f_{\text{unlabelled }Q_i}(t)$ fitted best the observed $f_{\text{unlabelled }Q_i}(t)$ for the tracer kinetics of sucrose (Q_1) , fructan (Q_2) , glucose (Q_3) , and fructose (Q_4) pools. In this, I constrained the model with the pool size ratios Q_i / Q_j as observed in the experiments. The mean residence time (τ) of carbon in the system was calculated as $\tau = (Q_1 + Q_2 + Q_3 + Q_4) / F_{01}$, with F_{01} the influx of carbon into the sucrose pool, the entry point of the system.

Two-pool model of the respiratory substrate supply system (see chapter 3.3)

The rate of respiration in darkness (R_n , µmol m⁻² s⁻¹) was obtained as per (Schnyder et al., 2003):

$$R_n = (F_{inlet} - F_{outlet}) / A$$
 Eqn 2.5

 F_{inlet} and F_{outlet} are the fluxes of CO₂ (µmol s⁻¹) entering and leaving the chamber, and A is the chamber ground area (m²).

The δ^{13} C of the sample air in the light and dark period were obtained as:

$$\delta^{13}C_{\text{sample}} = \left(\delta^{13}C_{\text{inlet}} \cdot F_{\text{inlet}} - \delta^{13}C_{\text{outlet}} \cdot F_{\text{outlet}}\right) / (F_{\text{inlet}} - F_{\text{outlet}})$$
Eqn 2.6

with $\delta^{13}C_{inlet}$ and $\delta^{13}C_{outlet}$ denoting the $\delta^{13}C$ of CO₂ entering and leaving the growth chamber, respectively.

I used the same two-pool compartmental model (Fig. 1.4) as Lehmeier et al. (2010a) based on the set of differential equations and implemented this in ModelMaker to analyse the time course of $f_{\text{unlabelled}}$ in the respiratory CO₂ of plant stands. $f_{\text{unlabelled}}$ of the respiratory CO₂ was calculated using Eqn 1.1 (chapter 1).

For the two pools in the system:

 $df_{\text{unlabelled }Q1}(t) / dt(t) \cdot Q_1 = f_{\text{unlabelled In}}(t) \cdot F_{01} + f_{\text{unlabelled }Q2}(t) \cdot k_{21} \cdot Q_2$

 $-f_{\text{unlabelled }Q1}(t) \cdot (k_{10} + k_{12}) \cdot Q_1$

Eqn 2.7

 $df_{\text{unlabelled }Q2}(t) / dt (t) \cdot Q_2 = f_{\text{unlabelled }Q1}(t) \cdot k_{12} \cdot Q_1 - f_{\text{unlabelled }Q2}(t) \cdot k_{21} \cdot Q_2$

Eqn 2.8

where $f_{\text{unlabelled }Qi}(t)$ denotes the fraction of unlabelled C in pool *i* at time *t*, F_{01} the flux into the system (which is equal to export flux; the respiratory rate) and $f_{\text{unlabled In}}(t)$ the fraction of unlabelled C in F_{01} with $f_{\text{unlabelled In}}(t) = 1$ before the switch of $\delta^{13}C_{\text{CO2}}(i.e.$ at t < 0), and $f_{\text{unlabelled In}}(t) = 0$ after the switch $(t \ge 0)$.

The pool sizes were given as by Lehmeier et al. (2010a):

$$Q_1 = F_{01}/k_{10}$$
 Eqn 2.9

$$Q_2 = F_{01} / k_{10} \cdot k_{12} / k_{21}$$
 Eqn 2.10

Based on optimized fluxes and pool sizes, the half-life $t_{1/2}$ (Q_i) was obtained by

 $t_{1/2}$ (Q_i) = ln(2)/ k_i

where k_i is the sum of all rate constants leaving the pool Q_i .

The contribution of pool Q_i to respiratory carbon released is defined as the probability that tracer leaves the system without cycling through Q_2 (i.e. current assimilate that was not stored), while the fractional contribution of stores to respiration is the probability that the tracer cycles through Q_2 at least once before it is respired (i.e. that the tracer underwent storage). These fractional contributions are given by:

Contribution of current assimilates =
$$k_{10}/(k_{10}+k_{12})$$
 Eqn 2.12

Contribution of temporary stores =
$$k_{12}/(k_{10}+k_{12})$$
 Eqn 2.13

The mean residence time of carbon in the respiratory supply system (τ) was calculated as:

$$\tau (h) = Q_{\text{total}} / r_{\text{plant}}$$
 Eqn 2.14

where Q_{total} is the total size of all respiratory substrate pools and r_{plant} is the respiration rate of the whole plant.

2.8 Statistical analysis

For the dataset from gas exchange measurements, the measurements in the first 45 minutes of one light and dark period and following the opening of the chamber were removed. In addition, measurements more than $1.5 \times IQR$ (Interquartile Range) away from the mean were removed as outliers. All statistical analyses were conducted in R version 3.6.1 (R Core Team, 2019) and Rstudio version 1.1.383 (Rstudio Team, 2016). The R-packages ggplot2 (Wickhan, 2016) and Agricolae (Mendiburu, 2017) were used for data plotting and ANOVA analysis, respectively. Type III Wald test was used to determine the significance of the CO₂ effects, and the post hoc Tukey's HSD test was performed for pairwise comparison among treatments. T-tests were used to test the δ^{13} C differences of reference carbohydrates between pure and mixture samples The number of replicates varied between measured parameters and treatments and is indicated in the Figure legends and Table captions.

3. Results and Discussion

3.1 Accuracy and precision of the near-natural abundance ¹³CO₂/¹²CO₂ labelling system

Constancy and consistency of the concentration and isotopic composition of CO_2 inside the plant growth chambers

One key methodical criterion of quantitative labelling experiments at different [CO₂] is the accuracy and precision of maintaining [CO₂] near target levels and the constancy of the $\delta^{13}C_{CO2}$ inside the growth chambers during the different phases of the labelling experiment. This includes the pre-labelling phase studied here that determines the so-called end-members of the isotopic mixing model (objective 1, chapter 1). The [CO₂] and $\delta^{13}C_{CO2}$ inside the growth chamber are closely reflected by $[CO_2]$ and $\delta^{13}C_{CO2}$ at the chamber outlet ($[CO_2]_{outlet}$; $\delta^{13}C_{CO2}$ outlet) in these highly ventilated chambers (Schnyder et al., 2003). Besides the technical capabilities and limitations (for specifications, see chapter 2), there are two biological (and hence inescapable) drivers that oppose the constancy of $[CO_2]_{outlet}$ and $\delta^{13}C_{CO2}$ outlet in plant growth chambers: the photosynthetic drawdown of [CO₂] which intensifies over time due to the expansion of the plant canopy and resulting canopy photosynthetic activity, and photosynthetic ¹³C discrimination (Δ^{13} C) which causes a ¹³C-enrichment of the chamber CO₂ that is mainly proportional to the photosynthetic CO_2 flux. In principle, the effect of these drivers on the constancy of [CO₂] close to the target value and maintenance of a stable $\delta^{13}C_{CO2}$ in the chambers can be counterbalanced by adjustments of [CO₂] at the chamber inlet ([CO₂]_{inlet}) and/or increases in the rates of air injection into the chambers. Here, I balanced these variables in such a way that the [CO₂] and $\delta^{13}C_{CO2}$ differences between the chamber inlet and outlet also permitted quasi-continuous measurements of the CO₂ exchange rate (net photosynthesis in the light period and respiration in the dark) and photosynthetic ¹³C discrimination (Δ^{13} C). Accordingly, as soon as the drawdown of [CO₂] between the chamber inlet and outlet started to exceed 10% of the inlet value, decreasing the ξ -value in Eqn 1.3 to <10 (Evans et al., 1986), I adjusted airflow rates in such a way that the drawdown of CO₂ would be kept near-constant while $[CO_2]$ at the chamber outlet remained close to the target value, so that the resulting ξ values were 5.12 (±0.09 SD), 5.87 (±0.24 SD) and 7.66 (±0.22 SD) for 200, 400 and 800 µmol mol⁻¹ [CO₂] respectively, after approximately Day 30. The result of these adjustments is shown in Fig. 3.1.1, which presents the deviation of the daytime mean [CO₂]_{outlet} from the target

values (200, 400 or 800 µmol mol⁻¹ depending on treatment) during the pre-labelling period between day 20 and day 65 of the different experimental runs. These data revealed that [CO₂]_{outlet} actually averaged somewhat higher than the target value in all [CO₂] treatments: +4.0 (±4.3 SD), +7.2 (±6.2 SD) and +13.9 (±8.3 SD) µmol mol⁻¹ for target [CO₂] of 200, 400 and 800 µmol mol⁻¹, respectively. These deviations of [CO₂]_{outlet} from target values were virtually identical (*P*<0.05) in the presence of the ¹³C-organic and -mineral CO₂ sources, and corresponded to mean %-deviations from target [CO₂] of <2% in all treatments. Meanwhile, the enhancement of $\delta^{13}C_{CO2 \text{ outlet}}$ relative to the inlet increased by several ‰ during the first 30 to 35 days of an experimental run, when ξ first attained a value of ≤10, and thereafter was kept relatively stable (<0.31‰ SD) until day 65, the end of the pre-labelling period (Fig. 3.1.2). That temporal pattern was very similar in the chambers receiving ¹³C-organic and -mineral CO₂.



Fig. 3.1.1 CO₂ concentration difference between chamber outlet ([CO₂]_{outlet}) and target [CO₂] ([CO₂]_{target}) with time ([CO₂]_{outlet} - [CO₂]_{target}). Target [CO₂]: (a, b) 200, (c, d) 400 and (e, f) 800 μ mol mol⁻¹. Growth chambers were supplied with either ¹³C-organic ($\delta^{13}C_{CO2}$ -43.5‰; left) or ¹³C-mineral CO₂ ($\delta^{13}C_{CO2}$ -5.6‰; right). The criterion of data evaluation was that measurements in the first 45 min of a light period or following the opening of the chamber were removed, and values over 1.5 × IQR (Interquartile Range) away from the mean were removed as outliers. Data points and error bars represent daily means ± SD (*n* = 9-23). Note that the dataset included the 14-d long period (day 49 to day 63) in which intensive sampling and leaf elongation measurements were performed.



Fig. 3.1.2 The δ^{13} C-difference between $\delta^{13}C_{CO2 \text{ outlet}}$ and $\delta^{13}C_{CO2 \text{ inlet}}$ with time ($\delta^{13}C_{CO2 \text{ outlet}}$ - $\delta^{13}C_{CO2 \text{ inlet}}$). CO₂ concentration at chamber outlet ([CO₂]_{outlet}) was maintained near target [CO₂]: 200 (a, b), 400 (c, d) and 800 (e, f) µmol mol⁻¹. Growth chambers were supplied with either ¹³C-organic ($\delta^{13}C_{CO2}$ -43.5‰; left) or ¹³C-mineral CO₂ ($\delta^{13}C_{CO2}$ -5.6‰; right). The criterion of data evaluation was that measurements in the first 45 min of a light period or following the opening of the chamber were removed, and values over 1.5 × IQR (Interquartile Range) away from the mean were removed as outliers. Data points and error bars represent daily means ± SD (*n* = 9-23). Note that the dataset included the 14-d long periods (day 49 to day 63) in which intensive sampling and leaf elongation measurements were performed.

Isotopic distance (spread) between chambers receiving ${}^{13}C$ -depleted and -enriched CO₂ as observed for components of CO₂ exchange, biomass and water-soluble carbohydrates

The isotopic distance ($d\delta^{13}C$, termed 'spread') between two parallel chambers run with the same target [CO₂] but receiving CO₂ with a different $\delta^{13}C_{CO2}$ (i.e. either ¹³C-organic or -mineral CO₂)

was highly similar for measurements at the inlet ($d\delta^{13}C_{CO2 \text{ inlet}}$, Table 3.1.1). The same was true for measurements at the outlet ($d\delta^{13}C_{CO2 \text{ outlet}}$, Table 3.1.1). Meanwhile, the spread for the $\delta^{13}C$ of net CO₂ exchange rates in daylight (net photosynthesis, $d\delta^{13}C_{Nd}$) and at night (dark respiration, $d\delta^{13}C_{Rn}$) appeared to be slightly smaller than that of CO₂ at the chamber inlet ($d\delta^{13}C_{inlet}$). Thus, $d\delta^{13}C_{Nd}$ was 97 - 99% and $d\delta^{13}C_{Rn}$ 93 - 97% of $d\delta^{13}C_{inlet}$, pointing to a small, and not always significant, contamination (e.g. leak) effect on the $\delta^{13}C$ of gas exchange components, particularly of dark respiration.

Minor CO₂ contamination effects on the $\delta^{13}C$ were also evident for the shoot and root biomass based on their $d\delta^{13}C_{shoot}$ and $d\delta^{13}C_{root}$, although the effects were not always significant. $d\delta^{13}C_{shoot}$ and $d\delta^{13}C_{root}$ averaged 96% of $d\delta^{13}C_{inlet}$, with no apparent significant difference between [CO₂] levels or plant parts. The $d\delta^{13}C$ for water-soluble carbohydrate components $(d\delta^{13}C_{fructan}, d\delta^{13}C_{sucrose}, d\delta^{13}C_{glucose}, and d\delta^{13}C_{fructose})$ was virtually the same as that of $d\delta^{13}C_{shoot}$, indicating that protocols for extraction and separation of the carbohydrates did not introduce any additional contamination that would have further adulterated the original ¹³C signal of the carbohydrates.

The consistently slightly smaller $d\delta^{13}C$ of biomass and water-soluble carbohydrate components relative to $d\delta^{13}C_{Nd}$ must be related to the fact that determination of $d\delta^{13}C_{Nd}$ only considered periods of near isotopic steady-state, that eliminated measurements made during the first 45 min after 'lights-on' during daytime and periods during which chambers had to be opened for operations within the chamber (such as for sampling or making leaf elongation rate measurements). The fact that $d\delta^{13}C$ of biomass and water-soluble carbohydrate components averaged ~96% of $d\delta^{13}C_{inlet}$, means that the integrated effect of all artefacts, including disturbances of the chamber atmosphere during daylight opening times of chambers, accounted for an average error (bias) of ~4% in all [CO₂] treatments. In other words, on average, 4% of the total C in plant biomass was not derived from the ¹³C-organic or -mineral CO₂ sources supplied to the chambers.

Cross-contamination of different carbohydrates due to imperfect separation on the preparative HPLC column (see chapter 2, Fig. 2.2) appeared to be a minor factor in determinations of the δ^{13} C of sugars. This assessment was supported by comparing $\delta^{13}C_{glucose}$ with $\delta^{13}C_{sucrose}$ or $\delta^{13}C_{fructose}$ in pure reference material and after HPLC passage of a reference mixture of fructan, sucrose, glucose and fructose (Table 3.1.3). This is illustrated by the deviation between pure and mixture-derived $\delta^{13}C_{glucose}$ (0.39‰), which was 2.6% of the $\delta^{13}C_{distance}$ distance between glucose and sucrose or fructose. Given the comparatively short separation

(elution) times differences between glucose and sucrose or fructose compared to fructan, and the low concentration of glucose in the reference mixture, the observed 2.6% cross-contamination likely represents near-maximum contamination.

Table 3.1.1 Isotopic difference (termed 'spread') between paired plant growth chambers receiving ¹³C-organic and -mineral CO₂ for different gas exchange, biomass and water-soluble carbohydrate components. Spread ($d\delta^{13}C_X$) was calculated as $d\delta^{13}C_X = \delta^{13}C_{X13C-mineral} - \delta^{13}C_{X13C-organic}$), with X designating the parameter of interest, e.g. bulk biomass or sucrose. Spread was calculated for $\delta^{13}C_{CO2}$ at the inlet ($d\delta^{13}C_{CO2 \text{ inlet}}$) and outlet ($d\delta^{13}C_{CO2 \text{ outlet}}$) of paired growth chambers in the light and dark period of days 61 to 64, daytime canopy CO₂ exchange ($d\delta^{13}C_{Nd}$) and respiration in the dark $(d\delta^{13}C_{Rn})$ for days 61 to 64, bulk shoot $(d\delta^{13}C_{shoot})$ and root C $(d\delta^{13}C_{root})$, and fructan $(d\delta^{13}C_{fructan})$, sucrose $(d\delta^{13}C_{sucrose})$, glucose $(d\delta^{13}C_{glucose})$ and fructose $(d\delta^{13}C_{fructose})$ extracted and purified from shoot biomass sampled at the beginning of the light period on day 65. Growth chambers were maintained at near target [CO₂] of 200, 400 or 800 µmol mol⁻¹ using one of two CO₂ sources, a ¹³C-organic (δ^{13} C -43.5‰) or a ¹³C-mineral source (δ^{13} C -5.6‰). δ^{13} C_{Nd} and δ^{13} C_{Rn} were determined during periods of steady-state gas exchange of chambers, measurements in the first 45 min of a light period or following the opening of the chamber were removed, and values over 1.5 x IQR (Interguartile Range) away from the mean were removed as outliers. Except for $[CO_2]$ and $\delta^{13}C_{CO_2}$, all conditions were kept the same in all chambers (see Materials and Methods). Daily means (\pm SD) for n = 2 to 10.

Spread	CO ₂ concentration (µmol mol ⁻¹)			
-	200	400	800	
		(‰)		
$d\delta^{13}C_{inlet}$	37.7 (0.1)	37.6 (0.1)	37.4 (0.0)	
$d\delta^{13}C_{\text{outlet day}}$	37.7 (0.2)	37.5 (0.2)	37.8 (0.1)	
$d\delta^{13}C_{\text{outlet night}}$	37.3 (0.1)	37.0 (0.2)	36.9 (0.1)	
$d\delta^{13}C_{Nd}$	37.2 (0.1)	36.9 (0.3)	36.2 (0.4)	
dδ ¹³ C _{Rn}	35.0 (1.2)	35.5 (1.0)	36.4 (0.3)	
$d\delta^{13}C_{shoot}$	35.9 (0.1)	35.7 (0.8)	36.2 (1.0)	
$d\delta^{13}C_{root}$	35.8 (0.2)	35.5 (0.5)	36.5 (0.5)	
$d\delta^{13}C_{fructan}$	35.9 (0.3)	36.4 (0.6)	35.4 (1.1)	
$d\delta^{13}C_{sucrose}$	36.1 (1.6)	36.2 (1.0)	36.0 (1.9)	
$d\delta^{13}C_{glucose}$	36.2 (1.6)	35.4 (0.9)	36.0 (1.9)	
$d\delta^{13}C_{fructose}$	36.0 (1.2)	35.5 (0.5)	36.5 (2.8)	

Table 3.1.2 δ^{13} C of reference carbohydrates measured pure or after HPLC separation of an 80:5:5:4 (wt:wt:wt) mixture of analytical grade inulin-fructan, sucrose, glucose and fructose (all from Merck, Darmstadt, Germany). The concentration of the individual carbohydrates in the mixture corresponded to typical concentrations in shoot extracts of plants grown at [CO₂] of 400 µmol mol⁻¹. Data shown are means ± SD (*n* = 2-3). "Pure" represents the pure individual component before being mixed. The δ^{13} C of individual carbohydrates was not significantly different between pure and mixture groups (*P* > 0.05).

	$\delta^{13}C_{fructan}$ (‰)	$\delta^{13}C_{sucrose}$ (‰)	$\delta^{13}C_{glucose}$ (‰)	δ ¹³ C _{fructose} (‰)
Pure	-25.14 ± 0.03	-25.32 ± 0.04	-10.61± 0.14	-25.25 ± 0.09
Mixture	-25.20 ± 0.20	-24.94 ± 0.37	-11.00± 0.87	-25.02 ± 0.4

Post-photosynthetic C isotope discrimination

Although the (isotopic) spread was very similar for dark respired CO₂, shoot biomass-C, and water-soluble carbohydrate-C extracted and isolated from shoot biomass, I did detect some systematic differences between their δ^{13} C values. These differences are shown as deviations from the δ^{13} C of shoot bulk biomass ($\delta^{13}C_X - \delta^{13}C_{shoot}$) in Fig. 3.1.3 and were independent of the isotopic composition of the δ^{13} C of source CO₂ supplied to the plant growth chambers. Of these differences, the deviation between $\delta^{13}C_{sucrose}$ and $\delta^{13}C_{shoot}$ was the most consistent, as it was significant and manifest in all [CO₂] treatments. Overall, sucrose was ¹³C-enriched by ~1‰ relative to shoot biomass and by ~1.5‰ relative to dark respired CO₂, glucose, fructose and structural biomass. Conversely, the relative ¹³C-enrichment of fructan (i.e. $\delta^{13}C_{fructan} - \delta^{13}C_{shoot}$) was only about half that for sucrose. In part, this must have been due to the very high fructan concentration in biomass: fructan accounted for >80% of all water-soluble carbohydrates and up to 60% of total shoot biomass in all [CO₂] treatments (see chapter 3.2). Therefore, its $\delta^{13}C_{fructan}$ must have had a strong effect on $\delta^{13}C_{shoot}$, of which it formed the main component. This also largely explains why the $\delta^{13}C$ of water-soluble carbohydrate-free shoot biomass, a proxy of shoot structural biomass ($\delta^{13}C_{structure}$), was ¹³C-depleted relative to $\delta^{13}C_{shoot}$.

 δ^{13} C-differences among biochemical or flux components have been previously discussed in terms of developmental (temporal), compartmental and true biochemical ¹³C fractionation effects (Bowling et al., 2008; Cernusak et al., 2009; Ghashghaie & Badeck, 2014), but are still not fully understood. However, the relatively close match of the δ^{13} C of whole-plant respiration ($\delta^{13}C_{Rn}$) with $\delta^{13}C_{shoot}$ does agree with observations of Klumpp et al. (2005) and Schnyder & Lattanzi (2005), who found only a tiny ¹³C discrimination in whole plant dark

respiration, although shoot respired CO₂ was systematically ¹³C-enriched and root respiration ¹³C-depleted in two herbaceous dicot species and in *L. perenne*. Also, the ¹³C-enrichment of sucrose and fructan relative to shoot structural biomass agrees with observations of relative ¹³C-enrichment of leaf non-structural carbohydrates (sugars and starch) in comparison with whole-leaf biomass (Bowling et al., 2008), although comparisons of the leaf- and whole shoot- ¹³C may not be entirely straightforward, due to the presence of both ¹³C-enriched autotrophic (leaf) and ¹³C-depleted heterotrophic (i.e. non-photosynthetic) tissues (Cernusak et al., 2009) in whole shoot biomass samples. The stronger relative ¹³C-enrichment of sucrose compared to all other biomass components would also agree with the observed ¹³C-enrichment of leaf sugars, as sucrose concentration in leaf blades was much higher than in any other plant part in the present investigations (Dorn, 2019). In the same work, Dorn (2019) observed the highest fructose and glucose concentrations in the immature growing, fully-heterotrophic shoot tissues, a factor that could be related to their relative ¹³C-depletion.

Finally, differences in residence time of C within a pool (e.g. of sucrose relative to shoot structural biomass) could also affect the relationship between their δ^{13} C. Sucrose is turned over rapidly (in a few hours; chapter 3.2), while C in structural shoot biomass resides in that pool throughout the life span of the plant part (several months). If it is true that ¹³C discrimination and post-photosynthetic fractionation remained unchanged throughout the experiment, some relatively ¹³C-depleted C must have been incorporated into the structural pool during the early phase (e.g. prior to 30 d) of an experimental run when the δ^{13} C of photosynthate was more depleted in ¹³C than in later phases (e.g. past 40 d) due to the increase of $\delta^{13}C_{CO2}$ in the chamber atmosphere (Fig. 3.1.2). By contrast, the δ^{13} C of sucrose obtained from samples collected on day 65 would be determined only by the more ¹³C-enriched CO₂ in the chamber near that sampling time.



Fig. 3.1.3 The δ^{13} C-difference between parameters X and shoot bulk biomass ($\delta^{13}C_X$ $- \delta^{13}C_{\text{shoot}}$), with X referring to the $\delta^{13}C$ of fructan, sucrose, glucose, fructose, structural biomass or dark respiratory CO₂. The δ^{13} C of shoot bulk biomass ($\delta^{13}C_{shoot}$), structural biomass, and the different carbohydrate fractions refer to samples collected on day 65 of the different experimental runs, i.e. before labelling. The $\delta^{13}C$ of structural biomass was determined as the difference between $\delta^{13}C_{shoot}$ and $\delta^{13}C$ of water-soluble carbohydrates and determined by isotopic mass balance. The $\delta^{13}C$ of dark respired CO_2 ($\delta^{13}C_{Rn}$) was determined during dark periods between days 61 to 65. Data (means \pm SE, n = 4-8) represent the average of equinumerous samples derived from chambers supplied with ¹³C-mineral and organic CO₂ and maintained near target [CO₂] of 200 (a), 400 (b), or 800 (c) µmol mol⁻ ¹. Stars mark a significant difference (P <0.05) between a parameter X and shoot bulk biomass ($\delta^{13}C_X - \delta^{13}C_{shoot}$).

The estimation of end-members for isotopic mixing models in labelling experiments

The fraction of unlabelled C ($f_{unlabelled}$) in C pools and CO₂ flux components (e.g. respiration) was quantified by the use of a two-member isotopic mixing model (Eqn 1.1, chapter 1). Estimates of the isotopic composition of the two end-members, i.e. of $\delta^{13}C_{new}$ and $\delta^{13}C_{old}$, can be obtained in two different ways: a one-chamber approach, which is employed when only one growth and labelling facility is available, or a two-chamber approach, which can be used in facilities with more than one labelling unit, as in this study. In both approaches, $\delta^{13}C_{old}$ is determined by measuring the $\delta^{13}C$ of the parameter of interest immediately before the switch of the CO₂ source. This $\delta^{13}C_{old}$ should be flux-, component- or compound specific to account for possible post-photosynthetic fractionation (Fig. 3.1.3). In the one-chamber approach, 38

 $s\delta^{13}C_{new}$ is calculated using the parameter-specific $\Delta^{13}C$ (calculated with $\delta^{13}C_{CO2 old}$) and the $\delta^{13}C$ of the labelling CO₂, i.e. $\delta^{13}C_{CO2 new}$ (Eqn 1.2 and 1.3, chapter 1). However, data in Table 3.1.1 show that approximately 4% of the C fixed was not derived from tank CO₂ but from contaminating CO₂. Hence, the "true" source CO₂ must have been slightly different from $\delta^{13}C_{outlet day}$, which was calculated after eliminating data when chambers were not in isotopic equilibrium. This applies to both the pre-labelling and the labelling phases. Consequently, $\Delta^{13}C$ is associated with some errors (although that error was comparatively small here). Also, the estimation of $\delta^{13}C_{new}$ is subject to a further error, as $\delta^{13}C_{CO2 new}$ for the labelling phase is not precisely reflected by $\delta^{13}C_{outlet day}$.

In the two-chamber approach, $\delta^{13}C_{old}$ in one chamber can be taken as $\delta^{13}C_{new}$ of the parallel chamber (following the change of the CO₂ source) and *vice versa*. Hence, any uncertainties concerning the pre-labelling and labelling of CO₂ sources are avoided. A prerequisite for this approach is that the extent of contamination is identical before and after the switch of the source CO₂. This assumption seems warranted because (1) experimental operations were virtually the same during the pre-labelling and labelling phases, and (2) there was no difference in the isotopic spread in parameters in temporal integration (Table 3.1.1).

In conclusion, the present work offered the unique opportunity to compare the analyses of tracer kinetics using a one-chamber and a two-chamber approach. It should be noted that possible artefacts associated with the one-chamber approach, namely the uncertainties in determining the exact value of the δ^{13} C of the source CO₂, became noticeable and quantifiable only by the two-chamber approach. Still, the analysis revealed a maximum 4% error due to contamination of the source CO₂ in the labelling experiments. Moreover, this work indicated that in investigations of photosynthetic and post-photosynthetic Δ^{13} C, contamination with extraneous CO₂ must be avoided by additional tests (Lehmeier et al., 2008), as those studies must be capable of capturing minute differences in Δ^{13} C. The present study also demonstrated that water-soluble carbohydrates in plant material can be extracted and separated via the HPLC system without cross contamination and adulteration of their δ^{13} C. This is very important for studying the turnover of carbohydrate pools (subsequent chapter) and for studying postphotosynthetic fractionation. As it allowed the quantification of (although minor) artefacts, I used the two-chamber approach to evaluate tracer data in the following chapters (3.2, 3.3, and 3.4).

3.2 The role of stores in recycling sucrose

Plant biomass, leaf area, canopy nitrogen status and water-soluble carbohydrates

Elevating atmospheric CO₂ concentration from 200 to 400 and 800 μ mol mol⁻¹ CO₂ increased the C mass of whole plants by 34% and 59%, and decreased the N mass of whole plants by 12% and 23%, respectively (both *P*<0.001; Table 3.2.1). These effects of rising [CO₂] were very similar for shoots and roots, as the shoot : root C mass ratio was not affected significantly by [CO₂] (*P*>0.05). Meanwhile, leaf area per plant was reduced by 6% and 14%, respectively, a statistically significant reduction (*P*<0.05) for the high [CO₂] level (Table 3.2.1). The nitrogen nutrition index (NNI) of the plant canopies averaged 0.79 at 200 µmol mol⁻¹ CO₂ and decreased by 38% and 52% (both *P*<0.001) by elevating CO₂ concentration to 400 and 800 µmol mol⁻¹ respectively (Table 3.2.1).

Elevating [CO₂] from 200 to 400 and 800 μ mol mol⁻¹ increased the total mass of watersoluble carbohydrates-C (WSCs-C) per shoot by 65% and 116%, respectively (both at *P*<0.001), in association with increases in the concentration of WSCs-C (mg g⁻¹ biomass-C) in shoot biomass of 23% and 37%, respectively (*P*<0.001; Table 3.2.2). Since leaf area per plant decreased with increasing [CO₂], these relationships corresponded to 66% and 147% enhancements of whole-shoot WSCs-C per unit leaf area resulting from the elevation of [CO₂] from 200 to 400 and 800 µmol mol⁻¹, respectively.

Of all WSC-fractions (Table 3.2.3), fructan contents showed the strongest response to the [CO₂] treatments. They accounted for 83-87% of total WSCs in the different treatments and increased by 74% and 161% on a leaf area basis when [CO₂] was increased from 200 to 400 and 800 μ mol mol⁻¹, respectively. This CO₂ response of fructan was almost paralleled by that of fructose which increased by 37% and 121% from 200 to 400 and 800 μ mol mol⁻¹ CO₂ in the air, respectively. Sucrose and glucose contents also increased with increasing [CO₂], but those increases were distinctly smaller. These relationships were stable during the 7 d-long labelling period. No starch was detected in the samples.

Chapter 3 Results and Discussion

Table 3.2.1 Growth parameters of *L. perenne* grown in the presence of $[CO_2]$ of 200, 400 or 800 μ mol mol⁻¹ in the growth chamber atmosphere. Means (±SE) are of biomass parameters and leaf area measurements based on a total of 24 (200 and 800 μ mol mol⁻¹ CO₂) or 48 (400 μ mol mol⁻¹ CO₂) replicates sampled at the beginning of the dark period on the first day after the switch of $\delta^{13}C_{CO2}$ (day 66) and on the following 2nd, 3rd, 4th and 7th days. Determinations of canopy nitrogen nutrition index (NNI, according to Lemaire et al., 2008) are based on 16 (200 and 800 μ mol mol⁻¹ CO₂) or 32 (400 μ mol mol⁻¹ CO₂) replicates sampled on day 63. Shoot and root growth rates were calculated based on biomass change over three weeks (day 49 to day 72). Different superscript letters indicate statistically significant differences among CO₂ treatments, see Table 3.2.5.

	Atmospheric CO ₂ concentration (μ mol mol ⁻¹)		
-	200	400	800
Dry plant biomass (g plant ⁻¹)	11.53 ^c (0.48)	14.75 ^b (0.42)	17.33 ^a (0.41)
Dry shoot biomass t (g plant ⁻¹)	9.21 ^c (0.34)	12.35 ^b (0.40)	14.45 ^a (0.38)
Dry root biomass (g plant-1)	2.33 ^b (0.19)	2.40 ^b (0.11)	2.88ª (0.13)
Whole plant C (g plant ⁻¹)	1.61ª (0.06)	2.15 ^b (0.06)	2.57 ^c (0.06)
Shoot C (g plant ⁻¹)	1.37ª (0.05)	1.84 ^b (0.06)	2.17 ^c (0.06)
Root C (g plant ⁻¹)	0.24ª (0.02)	0.31 ^b (0.01)	0.40 ^c (0.01)
Shoot C : root C ratio	6.17ª (0.44)	5.94ª (0.33)	5.44ª (0.17)
Whole plant N (mg plant ⁻¹)	74 ^a (7)	65 ^{ab} (3)	58 ^b (2)
NNI	0.79 ^a (0.02)	0.49 ^b (0.03)	0.38 ^c (0.01)
Leaf area (dm² plant-1)	10.2ª (0.4)	9.6ª (0.2)	8.8 ^b (0.2)
Growth rate			
shoot (mg C plant ⁻¹ d ⁻¹)	39 ^a (5)	51 ^b (4)	59 ^b (4)
root (mg C plant ⁻¹ d ⁻¹)	2ª (2)	2ª (2)	3ª (3)

Table 3.2.2 Total water-soluble carbohydrates (WSCs) in shoot of *L. perenne*. Means (±SE) of WSC are based on a total of 24 (200 and 800 μ mol mol⁻¹ CO₂) or 48 (400 μ mol mol⁻¹ CO₂) replicates sampled at the beginning of the dark period on the first day after the switch of $\delta^{13}C_{CO2}$ (day 66) and on the following 2nd, 3rd, 4th and 7th days. Different superscript letters indicate statistically significant differences among CO₂ treatments at *P* <0.001, see Table 3.2.5.

	Atmospheric CO ₂ concentration (μ mol mol ⁻¹)		
	200	400	800
WSCs-C			
per shoot (mg plant ⁻¹)	573ª (36)	947 ^b (43)	1240° (65)
per shoot biomass-C (mg g ⁻¹ biomass-C)	418 ^a (20)	515 ^b (11)	571° (27)
per leaf area (g m ⁻²)	39.0ª (1.9)	64.8 ^b (1.4)	96.5° (4.6)

Table 3.2.3 Content of individual water-soluble carbohydrates (sucrose, fructan, glucose and fructose) in the shoot of *L. perenne*. Means (\pm SE) of individual WSC are based on a total of 24 (200 and 800 µmol mol⁻¹ CO₂) or 48 (400 µmol mol⁻¹ CO₂) replicates sampled at beginning of the dark period on the first day after the switch of $\delta^{13}C_{CO2}$ (day 66) and on the following 2nd, 3rd, 4th and 7th days. Different superscript letters within a row indicate statistically significant differences among CO₂ treatments at *P* < 0.001, see Table 3.2.5.

	Atmospheric CO ₂ concentration (µmol mol ⁻¹)		
	200	400	800
Sucrose (g m ⁻²)	3.0ª (0.3)	3.9 ^b (0.2)	5.2 ^c (0.2)
Fructan (g m ⁻²)	32.3ª (1.5)	56.3 ^b (1.3)	84.4 ^c (4.4)
Glucose (g m ⁻²)	1.8ª (0.2)	2.0 ^a (0.1)	2.7 ^b (0.1)
Fructose (g m ⁻²)	1.9ª (0.1)	2.6 ^b (0.1)	4.2 ^c (0.2)



Fig. 3.2.1 Time course of the fraction of unlabelled C ($f_{unlabelled}$) in fructan (red), fructose (green), glucose (blue) and sucrose (orange) with labelling duration. Carbohydrates were extracted from the whole shoot of *L. perenne* grown at [CO₂] of 200 (a), 400 (b) or 800 (c) µmol mol⁻¹. The curves represent the fits of the 4-pool compartmental model shown in Fig. 1.2 (chapter 1). Each value is the mean (± SE) of four (200 and 800 µmol mol⁻¹ CO₂) or eight (400 µmol mol⁻¹ CO₂) replicates.

Fructan concentration in whole-shoot biomass was very high (\geq 35% of biomass) even at sub-ambient [CO₂] (Table. 3.2.1 and 3.2.2). That result was likely connected with an overabundance of assimilate availability (Wagner et al., 1983; Pollock & Cairns, 1991) determined by the high photosynthetically active radiation received by the canopies over the 16 h-long photoperiod and a simultaneously limited sink activity of the plants at reduced nitrogen fertilizer supply (Baca et al., 2020). Indeed, the nitrogen nutrition index of the canopies (defined as in Lemaire et al., 2008) was suboptimal even at sub-ambient [CO₂] (Table 3.2.1). In the same experiment, Baca et al. (2020) also observed that leaf elongation rates and epidermal cell lengths in all CO₂ treatments were typical of the effects of nitrogen limitation as observed in the same cultivar of *L. perenne* by Kavanova et al. (2008). Remarkably, the present data do not support a fundamental penalty to fructan storage in this C₃ grass under sub-ambient [CO₂], at least under the conditions of this experiment. Although I do not know of any other study of fructan levels under glacial maximum [CO₂], I assume that nitrogen limitation and cool climate conditions, both conducive to fructan storage, may have prevailed together under glacial maximum [CO₂].

Still, increasing [CO₂] generated an additional increase in fructan accumulation, while the nitrogen nutrition index of the canopies continued to decline even further (compare Table 3.2.1 and 3.2.2), in line with observations in free-air CO₂ enrichment (FACE) studies (Isopp et al., 2000; Kimball et al., 2002; Högy et al., 2013). However, the whole-shoot fructan concentration at double ambient [CO₂] observed here (57%) was much higher than any reported FACE effect on fructan concentration (Högy et al., 2009, 2013). In fact, the whole-shoot fructan concentration observed here is the highest that I am aware of in an entire system of a C₃ grass, and again, likely resulted from several factors combining to enhance fructan accumulation (see above), including the [CO₂] fertilizer effect and the decrease in the nitrogen nutrition status. This [CO₂] effect on fructan storage occurred while non-WSC biomass of shoots increased very little with [CO₂]. Thus, non-WSC biomass at 800 µmol mol⁻¹ CO₂ was only 11% higher than at 400 and 28% higher than at 200 µmol mol⁻¹ CO₂. In comparison, fructan mass in the shoot increased by 74% and 129%, respectively.

Compartmental modelling of central carbohydrate metabolism

Estimates of the half-life of individual carbohydrate pools, the substrate fluxes among pools and the environment, and the partitioning of sucrose between fructan synthesis, hydrolysis or utilization in growth and maintenance activities (including sucrose export to the root system) were obtained based on optimization of the rate constants when fitting the four-pool model to the time courses of $f_{unlabelled}$. The four-pool model of carbohydrate metabolism of a fructan storing grass species (Fig.1.2, chapter 1) provided a virtually perfect fit to the dynamic labelling data of the whole-shoot carbohydrate system at sub-ambient and ambient [CO₂]. The goodness of fit was near-perfect for the sub-ambient ($R^2=0.99$) and ambient ($R^2=0.98$) [CO₂], and somewhat less so for the double-ambient [CO₂] (R^2 =0.89). Fits with the 'free' (that is stoichiometrically unconstrained) four-pool model, or free models, without consideration of fructose-glucose interconversions (Lattanzi et al., 2012) did not improve the quality of the fit in any [CO₂] treatment. Such a good fit of a compartmental model may not be intuitive given the assumptions of compartmental modelling: that the system is in a steady state at the day-byday timescale, i.e. that fluxes into and out of pools are the same and constant, and that all pools are well-mixed, given the compartmental and developmental heterogeneity of the plant parts that are combined in the system. This includes leaves of different developmental stages and a variety of functionally distinct tissues in both the lamina and sheath parts of each leaf. However,

Chapter 3 Results and Discussion

all plants remained vegetative during the experiment, meaning that the shoot was composed virtually entirely of leaves. In support of modelling assumptions, I found no significant day-byday variation of carbohydrate concentrations during the 7-d long dynamic labelling experiment, as I only sampled near the end of light periods. Also, the shoot growth rate was <3% d⁻¹ (Table 3.2.1) in these well-developed closed stands. This slight deviation from the constancy-of-shootsize assumption was deemed acceptable, as it implied, for instance, that the actual rate of fructan synthesis on a given day underestimated the rate of hydrolysis by less than 3% d⁻¹, as whole shoot fructan concentration did not vary significantly with labelling time. An analogous evaluation and qualification of the steady-state assumption was also performed by Lattanzi et al. (2008 and 2012) in their compartmental modeling of C and N substrate fluxes into leaf growth zones and of central carbohydrate analysis in mature leaves of perennial ryegrass. In comparison, errors in model estimates for the other pools (sucrose, glucose, fructose) must have been minor, as they were turned over much faster. Also, the effects of potential physical compartmentation did not affect model features in Lattanzi et al. (2012). Overall, the present observations may be taken to indicate that biochemical (functional) features of central carbohydrate metabolism were much more critical than potential compartmentation (structural) effects, as the four-pool model described only the broad features of the biochemistry of watersoluble carbohydrate metabolism and yielded excellent fits to tracer data at least under halfambient and ambient [CO₂].

Interestingly, the stoichiometrically-constrained model provided the most parsimonious, best fit to the labelling kinetics of the whole shoot carbohydrate system. That is, more complex models that permitted glucose-fructose interconversion activities or non-stoichiometrically-constrained sucrose re-synthesis from free glucose and fructose, did not demonstrate a statistically superior fit (data not shown). This result is also consistent with the idea that sucrose cycling *via* hydrolysis/re-synthesis and passage *via* the fructan pool may principally occur independently of each other. Whether or not the relative activities of the two cycling mechanisms occurred at the same relative rates in leaves of different ages or leaf blades and sheaths, however, cannot be deduced from this analysis, as all shoot tissues were combined in one sample. However, if the relative magnitude of the two activities did change between developmental stages or tissue types, the fact that the activities occurred independently would explain why the model nevertheless provided an excellent fit to the carbohydrate systems' labelling kinetics.

Tracer kinetics and compartmental analysis

The labelling time courses of the different carbohydrates (i.e. the changes of $f_{unlabelled}$ with time) were qualitatively similar in the different treatments (Fig. 3.2.1): initial label incorporation was fastest and similar in sucrose and glucose, and slowest in fructan. In comparison, fructose demonstrated a labelling time course that was intermediate between that of fructan and sucrose. These time courses of $f_{unlabelled}$ in the carbohydrate system (Fig. 3.2.1) of the whole shoot were well fitted with the so-called 'stoichiometrically constrained' four-pool compartmental model presented by Lattanzi et al. (2012).

The results indicated that [CO₂] had no significant effect on the half-life of any carbohydrate (sucrose, fructan, glucose or fructose) (Table 3.2.4). Also, at all [CO₂], the predicted half-lives were relatively short and similar for the sucrose (~2.4 h), glucose (~3 h) and fructose (~4 h) pools, while that of the fructan pool (~180-190 h) was almost two orders of magnitude higher than that of the sucrose pool. The residence time of carbohydrate-C in the system increased from about 8.9 d to 13.6 d between 200 and 800 µmol mol⁻¹ [CO₂], essentially due to greater sucrose partitioning to fructan synthesis. The half-life $(T_{1/2})$ of fructan observed here $(\sim 7.7 \text{ d})$ was much longer than that in mature photosynthesising leaf blades of the same cultivar of *L*. *perenne* grown in continuous light with high ($T_{1/2} = 1.1$ d) or low nitrogen fertilizer supply ($T_{1/2} = 2.6$ d). As the studies addressed different scales and were performed in different environments, it is unclear which factor explained the difference in half-lives. It is possible, however, that fructan contents in the sheath part of leaves were more prominent and possibly turned over more slowly than in leaf blades (Thomas & James, 1999). The fructan half-life observed here was equivalent to a mean residence time of about 11 d, close to the leaf appearance interval observed by Baca et al. (2020) in the same study. Also, the half-life of fructan corresponded to about one-quarter to one-fifth of the average leaf life span of L. perenne observed elsewhere (Schleip et al., 2013, and compilation therein). Although these relationships do not prove that fructan storage was constrained by leaf life span directly, I cannot rule out that perhaps developmental cues associated with leaf senescence were involved in the control of fructan storage and mobilisation.

Although, I know of no other studies of the half-lives of sucrose, glucose or fructose at the whole shoot level, the half-life estimates here appear to be realistic when compared with studies on the scale of mature leaves (Farrar & Farrar 1985, 1986; Borland & Farrar, 1988; Lattanzi et al., 2012) or leaf growth zones (Schnyder et al., 1987), and the short delays in the system that may be connected with sucrose transport times (Windt et al., 2006) between leaves

Chapter 3 Results and Discussion

and sheaths. Clearly, the idea that the sucrose pool is homogeneous is a significant simplification, given the fact that sucrose may be found in several cellular compartments (cytoplasm, vacuole, apoplast) and in various functionally distinct tissue types, including sieve tubes and companion cells (Ruan 2012, 2014; Du et al., 2019). Indeed, the half-life estimate of sucrose must have been dominated by those tissues that contained the bulk of the sucrose, particularly the cytoplasm of lamina and sheath parenchyma cells. Also, I sampled only near the end of the light period, meaning that a certain fraction of the sucrose in the leaf blades may have been vacuolar (Farrar et al., 1989) if plants used a diurnal sucrose storage mechanism. This factor would tend to increase the half-life of the 'average' sucrose molecule. However, the half-life of sucrose as obtained here was close to (or not much longer) than that of the "transport" sucrose pool (Farrar & Farrar, 1985), which was assumed to include all sucrose not stored in the vacuole of barley leaf blades. Additionally, I would not expect a strong effect of carbohydrate metabolism within sink tissue (that is the leaf growth-and-differentiation zones of the vegetative tillers that composed the shoot, Baca et al., 2020) on the tracer dynamics in the shoot as described by the model, since carbohydrate contents of that tissue accounted for only a tiny fraction (i.e. <10%) of the total quantity of water-soluble carbohydrates in the shoot in the different treatments. For that reason, any eventual discrepancy between labelling patterns of carbohydrates in sinks compared with the vastly greater mass of fully-developed remaining plant parts (source and storage tissue) would have a minimal effect at the scale of the whole shoot tracer kinetics. Together, the predominance of sucrose in the cytoplasm of parenchyma in leaf blades and sheaths, and the short transit times in the transport system linking the plant parts, may have combined to yield a half-life of whole-shoot sucrose that was close to that of the "transport" pool of individual leaves as found in several studies (Geiger et al., 1983; Farrar & Farrar, 1986; Rocher & Prioul, 1987; Borland & Farrar, 1988).

For the different treatments, the model results indicated that about equal fractions of sucrose were partitioned between the three alternative functions: (1) supply of growth and maintenance of the shoot and export to roots, or use in (2) fructan synthesis or (3) hydrolysis (Table 3.2.4). The approximately equal partitioning was true, except for the low [CO₂] treatment, where sucrose partitioning to fructan synthesis (0.22) was significantly lower than its use in the other processes, also by comparison with the treatments with higher [CO₂]. In particular, at low [CO₂], sucrose use in hydrolysis was almost double that for fructan synthesis. For a system in steady-state, that [CO₂] effect implied that sucrose recycling from the products of hydrolysis was almost twice as important as that from fructose released during fructan breakdown at 200 μ mol mol⁻¹ CO₂, while at 400 and 800 μ mol mol⁻¹ CO₂, fructan metabolism

and sucrose hydrolysis contributed approximately equally to sucrose recycling. The summed proportion of the gross flux of sucrose cycling through fructan and hydrolysis was insensitive to $[CO_2]$, and, accordingly, the proportion of sucrose used in growth and maintenance (including export to roots) was also practically constant (Table 3.2.4). At the same time, partitioning of the gross flux of sucrose use between hydrolysis and fructan synthesis did change considerably. The rate of sucrose consumption in hydrolysis was about two times that in fructan synthesis at low [CO₂] while partitioning between hydrolysis and fructan synthesis was approximately equal at ambient and high [CO₂]. Although effects of low [CO₂] on sucrose cycling via hydrolysis/resynthesis have not been studied so far, to my best knowledge, it is well known that sucrose cycling via hydrolysis/resynthesis is common and active in photosynthesizing leaves of dicots and grasses (Wagner et al., 1986; Lemoine, 2000; Du et al., 2020), including in *L. perenne* (Lattanzi et al., 2012) as well as in sink tissue, such as tap roots of sugar beet (Giaquinta, 1979) or tomato fruits (Nguyen-Quoc & Foyer, 2001) or other heterotrophic systems (Kruger et al., 2007). The estimates for the rates of cycling via hydrolysis in the present study were comparable in magnitude to estimates for leaves of Arabidopsis by Nägele et al. (2010), soybean and tobacco by Huber (1989) and cotyledons of germinating Ricinus communis by Geigenberger & Stitt (1991). It has been proposed that leaf sucrose cycling through hydrolysis should increase as photosynthesis acclimates to elevated [CO₂] (Moore et al., 1999). However, my results do not support that hypothesis in terms of sucrose partitioning, although I cannot rule out that a possible enhancement of sucrose cycling via hydrolysis in leaf blades was compensated by decreased cycling elsewhere in the shoot (i.e. in sheaths). Indeed, Lattanzi et al. (2012) observed much greater cycling via hydrolysis in leaf blades. Also, my modelling indicated that the gross flux through hydrolysis at the whole shoot level may have increased with [CO₂], although that result was not statistically significant.

In addition, differences in $[CO_2]$ during growth caused significant differences in the scale of fluxes within and out of the system (Table 3.2.4). For instance, the total flux of sucrose out of the system (F_{10}), supplying growth, maintenance and sucrose export to roots, increased 1.7-fold between 200 and 800 µmol mol⁻¹ [CO₂]. The strongest [CO₂] effect on fluxes was connected with fructan synthesis (F_{12}) and associated fructan hydrolysis (F_{24}). Thus, increasing [CO₂] from 200 to 800 µmol mol⁻¹ caused a 2.8-fold increase in the flux of C through the fructan pool. On the other hand, I found no significant effect of [CO₂] on the sucrose flux through hydrolysis.

Chapter 3 Results and Discussion

In conclusion, increasing $[CO_2]$ from 200 to 800 µmol mol⁻¹ increased total shoot biomass 1.6-fold, fructan mass 2.3-fold and its concentration in shoot biomass from 35% to 51%. Compartmental modelling indicated that the biggest impact of increasing $[CO_2]$ on central carbohydrate metabolism was to increase carbon cycling through the fructan pool. By contrast, the half-life of the fructan pool (~7.7 d) was insensitive to $[CO_2]$ change, as were the half-lives of sucrose, glucose and fructose (2.3-4.5 h). Sucrose re-synthesis from breakdown products of fructan (fructose) accounted for an increasing proportion of total sucrose synthesis and was similar in magnitude to sucrose neo-synthesis at double-ambient $[CO_2]$. The data indicate fructan as the main factor explaining the turnover time of the metabolic pool serving growth and respiration of a temperate perennial grassland species during undisturbed growth.

Table 3.2.4 Parameters of the four-pool compartmental model of central carbohydrate metabolism in the shoot of *L. perenne*. Parameters were numerically optimized as described in chapter 2, based on the tracer kinetics data shown in Fig. 3.2.1. Fluxes are expressed as g C m⁻² leaf area h⁻¹ and calculated by the differential equations given in chapter 2, using the carbohydrate contents per leaf area as shown in Table 3.2.3. The results of the fluxes predicted by the optimized rate constants are given in bold. Numbers in a row or column with different superscript letters indicate a statistically significant difference at 95% confidence intervals.

	Atmospheric CO ₂ concentration (µmol mol ⁻¹)		
	200	400	800
Pool half-life			
Sucrose $(T_{1/2 Q1}, h)$	2.4ª (0.2)	2.5ª (0.2)	2.3ª (0.4)
Fructan (T _{1/2 Q2} , h)	190.5ª (14.5)	181.6ª (16.4)	178.7 ^a (39.1)
Glucose (T _{1/2 Q3} , h)	3.7 ^a (0.3)	3.3ª (0.3)	2.9 ^a (0.5)
Fructose (T _{1/2 Q4} , h)	3.8 ^a (0.3)	4.3 ^a (0.3)	4.5 ^a (0.8)
C flux			
<i>F</i> ₁₀ (g m ⁻² h ⁻¹)	0.18ª (0.02)	0.25 ^b (0.02)	0.30 ^b (0.04)
<i>F</i> ₁₂ (g m ⁻² h ⁻¹)	0.12 ^a (0.01)	0.21 ^b (0.02)	0.33 ^b (0.07)
<i>F</i> ₁₃ (g m ⁻² h ⁻¹)	0.34ª (0.04)	0.42 ^{ab} (0.04)	0.65 ^b (0.12)
<i>F</i> ₁₄ (g m ⁻² h ⁻¹)	0.23ª (0.03)	0.21ª (0.03)	0.33ª (0.10)
<i>F</i> ₂₄ (g m ⁻² h ⁻¹)	0.12ª (0.01)	0.21 ^b (0.02)	0.33 ^b (0.07)
<i>F</i> ₃₁ (g m ⁻² h ⁻¹)	0.34ª (0.05)	0.42 ^{ab} (0.04)	0.65 ^b (0.12)
<i>F</i> ₄₁ (g m ⁻² h ⁻¹)	0.34ª (0.04)	0.42 ^{ab} (0.04)	0.65 ^b (0.13)
Sucrose partitioning			

Chapter 3 Results and Discussion

To growth, maintenance, and	0.34 ^{Aa} (0.04)	0.38 ^{Aa} (0.03)	0.31 ^{Aa} (0.06)
export To fructan synthesis	0.22 ^{Ba} (0.03)	0.32 ^{Ab} (0.04)	0.35 ^{Ab} (0.09)
To hydrolysis	0.43 ^{Aa} (0.07)	0.31 ^{Aa} (0.04)	0.34 ^{Aa} (0.11)
Residence time in the system (h)	213ª (23)	255 ^{ab} (17)	327 ^b (45)
Goodness of model fit (<i>R</i> ²)	0.99	0.98	0.89

Parameter	<i>F</i> -value	P-value
Dry plant biomass (g plant ⁻¹)	30.39	<0.001
Dry shoot biomass (g plant-1)	30.59	<0.001
Dry root biomass (g plant ⁻¹)	3.93	0.23
Plant C (g plant ⁻¹)	39.88	<0.001
Plant N (mg plant ⁻¹)	4.31	0.08
Shoot C (g plant ⁻¹)	32.67	<0.001
Root C (g plant ⁻¹)	33.24	<0.001
Shoot C : root C ratio	1.99	0.14
NNI	109.7	<0.001
Leaf area (dm ² plant ⁻¹)	4.67	0.01
WSC mass (g plant ⁻¹)	34.47	<0.001
WSC concentration (mg g ⁻¹)	15.34	<0.001
WSC content (g m ⁻²)	104.3	<0.001
Fructan (g m ⁻²)	96.53	<0.001
Sucrose (g m ⁻²)	24.91	<0.001
Glucose (g m ⁻²)	12.14	<0.001
Fructose (g m ⁻²)	55.3	<0.001

Table 3.2.5 Results of one-way ANOVA, testing the effect of [CO₂].

3.3 The role of stores in supplying respiratory substrate

Respiration rate

In this section, I report the analysis of the response of plant respiration rate to a change of $[CO_2]$ during growth (object 3, chapter 1). Canopy dark respiration (R_n), that is dark respiration rate per unit chamber ground area, was calculated as specified in Eqn 2.5 (chapter 2). Relative to $[CO_2]$ of 400 µmol mol⁻¹, R_n was 19% lower (P < 0.05) at $[CO_2]$ of 200 µmol mol⁻¹. However, increasing $[CO_2]$ from 400 to 800 µmol mol⁻¹ had no significant effect (Table 3.3.1 and 3.3.2). The latter result aligns with the study of Ryle et al. (1992), that found no significant response of R_n on a ground area basis when *L. perenne* was grown at $[CO_2]$ of 680 µmol mol⁻¹ relative to 340 µmol mol⁻¹. Also, the fact that area-based R_n was reduced under low $[CO_2]$ is consistent with the study of Ayub et al. (2011), who reported that area-based R_n of *Eucalyptus saligna* decreased under $[CO_2]$ of 280 µmol mol⁻¹ relative to 400 µmol mol⁻¹.

Meanwhile, R_n per unit plant-C (which is also termed specific respiration rate, e.g. Lötscher et al., 2004) did not differ significantly between the [CO₂] treatments (Table 3.3.1 and 3.3.2). This indicated that the significant [CO₂] effect on canopy respiration rate (i.e. R_n per unit ground area) observed when increasing [CO₂] above 200 µmol mol⁻¹ was primarily a function of [CO₂] effects on canopy biomass.

In the following compartmental modelling, I used R_n per unit of chamber ground area, as this is also the normal basis for respiration measurements at the plant community level (Amthor, 1991).

Table 3.3.1 Canopy dark respiration rate of *L. perenne*. Respiration was expressed per unit chamber area or per unit plant-C. Plants were grown in the growth chamber with $[CO_2]$ of 200, 400, or 800 µmol mol⁻¹ CO₂, 20/16 °C air temperature and 50%/75% relative humidity during the 16/8 h light/dark period. Photosynthetic photon flux density (PPFD) was 800 µmol m⁻² s⁻¹.at canopy height. Measurements were conducted from day 66 to day 72. Shown are means ± SE of 2 - 4 replicates. Different superscript letters within a row indicate statistically significant differences between CO₂ treatments.

	Atmospheric CO_2 concentration (µmol mol ⁻¹)		
	200	400	800
Dark respiration rate			
per unit ground area (µmol C m ⁻² s ⁻¹)	-4.26 ^a (0.18)	-5.23 ^b (0.08)	-4.99 ^b (0.21)
per unit plant-C (μmol C mol ⁻¹ plant C s ⁻¹)	-0.19ª (0.03)	-0.18 ^a (0.02)	-0.14 ^a (0.005)

Table 3.3.2 Results of one-way ANOVA testing effect of CO₂ on plant dark respiration rate.

Parameter	<i>F</i> -value	P-value
Dark respiration rate		
per unit ground area (µmol C m ⁻² s ⁻¹)	154.8	<0.05
per unit plant-C (µmol C mol ⁻¹ plant C s ⁻¹)	2.4	0.19

Tracer kinetics and compartmental analysis

The tracer kinetics in respired CO₂ were very similar between all [CO₂] treatments, each consisting of two major phases (Fig. 3.3.1): first, the fraction of unlabeled C ($f_{unlabelled}$) decreased rapidly by ~0.25 during the first day of labelling, and then it continued to decrease at a distinctly slower rate during the following phase of continuous labelling. After 7 days of continuous labelling, the value of $f_{unlabelled}$ had decreased to ~0.42. The change of $f_{unlabelled}$ over time appeared to occur only during the 16 h-long photoperiods, as hourly $f_{unlabelled}$ in respired CO₂ did not change significantly during the night, as indicated by the data for the 1st and 2nd night following the start of labelling (Fig. 3.3.2).

In each $[CO_2]$ treatment, fitting two-exponential functions to the tracer kinetics of respired CO₂ gave the best fit, indicating that the respiratory substrate supply system could be

separated into two pools. This compared well with the single-shoot, two-pool compartmental model used by Lehmeier et al. (2010a; see Fig. 1.4, chapter 1) in investigations of day-length effects on the compartmental structure of the system supplying substrate for respiration. In my work, the goodness-of-fit of the two-pool model, given by the R^2 , was >0.99 in all [CO₂] treatments. A one-pool model exhibited a significant lack of fit to the data, while a three-pool model did not improve fit quality compared to the two-pool model, as in Lehmeier et al. (2010a) (data not shown).

In the two-pool compartmental model, the R_n on a ground area basis represents the efflux of the model, which equals the respiratory substrate influx into the system in the steady state. The half-life ($T_{1/2}$) and size (Q) of each pool did not differ significantly between [CO₂] treatments (Table 3.3.3). The $T_{1/2}$ of Q_1 , the fast pool, was in the order of ~11 - 15 h, and that of Q_2 , the slow pool, ranged between 3 - 5 d. Relative to the pool size of Q_2 in ambient [CO₂], Q_2 was 34% smaller at low [CO₂] and 32% larger at high [CO₂]. The proportional contribution of Q_2 to total respiratory substrate supply was very large and highly similar between all [CO₂] levels (58-60%). The carbon residence time in the entire respiratory supply system ranged between 194 h (200 µmol mol⁻¹ CO₂) and 291 h (800 µmol mol⁻¹ CO₂).

The fact that $f_{unlabelled}$ did not change significantly during the course of dark periods (Fig. 3.3.2) is consistent with results by Schnyder et al. (2003), who also reported a constant hourly $f_{unlabelled}$ during the dark period. This finding supports the idea that the respiratory substrate pool is well mixed with "labelled" and "unlabelled" carbon at the whole plant (and community) level.

Within a continuous labelling experiment (as practiced here), unlabelled carbon in the respiratory substrate is gradually replaced by labelled carbon until $f_{unlabelled} = 0$. In a study by Lehmeier et al. (2010a), they found that $f_{unlabelled}$ of shoot or root respired CO₂ decreased to < 0.2 on the 7th day of continuous labelling when *L. perenne* was grown at [CO₂] of 360 µmol mol⁻¹. This progression of labelling was faster than that I observed at all [CO₂] levels. This difference between Lehmeier et al. (2010a) and my investigations may be related to plant growth conditions. Compared with the study of Lehmeier et al. (2010a), in my study, plants were grown at a much higher light intensity (PPFD: 800 compared to 425 µmol m⁻²s⁻¹) and with a lower N supply (N concentration: 5.1 compared to 7.5 mM). The higher photosynthetically active radiation may have led to a higher assimilate availability (Wagner et al., 1983; Pollock & Cairns, 1991), while, simultaneously, the lower nitrogen supply could

have caused sink limitation (Baca et al., 2020). Both factors would lead to a greater accumulation of respiratory substrate in the form of water-soluble carbohydrates (mainly fructan, see chapter 3.2). The slower rate of change of $f_{\text{unlabelled}}$ in my work may be because the turnover of a much larger respiratory substrate store would take much longer, even if the absolute rate of tracer addition to that pool were the same.

In all [CO₂] treatments, the tracer kinetics in respired CO₂ could be fitted with a twoexponential curve. This finding is consistent with the study by Lehmeier et al. (2010a), who reported a tracer time course with two distinct phases in respired CO₂ of shoot and root in *L*. *perenne* when plants grown in [CO₂] of 360 µmol mol⁻¹ were labelled continuously over 14 days. Gong et al. (2017) observed a similar tracer kinetic pattern in respired CO₂ of sunflower stands grown in [CO₂] of 200 or 1000 µmol mol⁻¹ over a 5-d labelling period. Conversely, Lehmeier et al. (2018, 2010b) also found a third, very fast phase immediately at the beginning of labelling. I could not observe this first initial labelling phase, as my first determination of *f*_{unlabelled} only occurred at 16 h after the start of labelling.

The $T_{1/2}$ of Q_1 (12.7 h), the fastest labelled pool detected here, agreed reasonably well with that of the transport and metabolic pool (2 to 8 h), discussed by Lehmeier et al. (2010a). However, I cannot discount the possibility that Q_1 included some contribution from a shortterm or diurnal store (Lehmeier et al., 2010b), which may have included vacuolar sucrose (Farrar & Farrar, 1986; Borland & Farrar; 1988).

In contrast, Q_2 is a long-term storage pool, forming the primary carbon source to feed plant respiration (Table 3.3.3). This finding was in agreement with observations by Lehmeier et al. (2010a, b), who reported that the respiratory storage pool (Q_2) contributed *c*. 60% to total respired C of *L. perenne* when grown in continuous light, regular day-night cycles, or under nitrogen-limited conditions. Nogués et al. (2004) found that the products of current photosynthesis supplied less than 50% of the substrate feeding leaf respiration of French beans. Together, these results highlight that the long-term storage pool formed a central part of the integral C supply system to feed plant respiration. I also found that the carbon stored in Q_2 exceeded the total demand for respiration during a dark period by comparing the size of Q_2 to the rate of F_{10} (Table 3.3.3), indicating this pool may act as a buffer accommodating fluctuating respiratory substrate supply in day-night cycles. This result differs from a study with a starchstoring plant by Gong et al. (2017). Who reported that the long-term storage pool contributed <36% of carbon respired by sunflowers, an amount that was insufficient to meet the demand of dark respiration in one night.

The residence time of respiratory substrate in the plant (τ) is determined by respiratory substrate pool size and efflux rate (F_{10}) (Table 3.3.3). I observed no difference in τ between [CO₂] treatments, despite F_{10} being significantly smaller at [CO₂] of 200 µmol mol⁻¹. τ was more than two times longer than the $T_{1/2}$ of C in the storage pool (Q_2) (234 h vs 93 h) (Table 3.3.3). This result also agreed with the observation that the tracer entered the long-term storage compartment more than once before being respired (Lehmeier et al., 2010b).

In conclusion, R_n was enhanced with increasing [CO₂] from 200 to 400 µmol mol⁻¹ and was insensitive to increasing [CO₂] from 400 to 800 µmol mol⁻¹. Compartmental modelling indicated that an increase of [CO₂] from 200 µmol mol⁻¹ to 800 µmol mol⁻¹ increased the size of the long-term storage pool by 197% and its $T_{1/2}$ by 148%. This highlights the great importance of Q_2 in supplying respired substrate.



Fig. 3.3.1 Time course of the fraction of unlabelled carbon ($f_{unlabelled}$) in respired CO₂ of stands of *L. perenne*. Plants were grown at [CO₂] of 200 (black), 400 (red) or 800 (green) µmol mol⁻¹. The labelling experiment started at day 65. Each value is the mean (± SE) of two (200 and 800 µmol mol⁻¹ CO₂) or four (400 µmol mol⁻¹ CO₂) chamber-scale replicates. The dashed line represents the two-pool model prediction.



Fig. 3.3.2 Time course of the fraction of unlabelled carbon ($f_{unlabelled}$) in respired CO₂ by stands of *L. perenne* during the 1st (a) and 2nd night (b) after the beginning of labelling. Plants were grown at [CO₂] of 200 (black), 400 (red) or 800 (green) µmol mol⁻¹. The labelling experiment started on day 65. Shown are means (± SE) of hourly values of stands during the dark period on the first and second labelling day. n = 2-4.

Table 3.3.3 Effect of atmospheric CO₂ concentration during growth on the parameters of the twopool compartmental model of respiratory substrates in stand *L. perenne*. Parameters were optimized as described in chapter 2, based on the tracer kinetics data shown in Fig. 3.3.1. Fluxes are expressed as g C m⁻² h⁻¹, which is dark respiration rate expressed on a chamber ground area basis. The results of the fluxes predicted by the optimized rate constants are given in bold. Different superscript letters indicate a statistically significant difference at 95% confidence intervals.

	Atmospheric CO ₂ concentration (µmol mol ⁻¹)		
-	200	400	800
C flux			
<i>F</i> ₁₀ (g C m ⁻² h ⁻¹)	0.183ª	0.223 ^b	0.214 ^b
<i>F</i> ₁₂ (g C m ⁻² h ⁻¹)	0.242ª (0.059)	0.309 ^a (0.029)	0.326 ^a (0.078)
<i>F</i> ₂₁ (g C m ⁻² h ⁻¹)	0.242ª (0.059)	0.309 ^a (0.029)	0.326ª (0.078)
Pool half-life			
T _{1/2} (Q ₁) (h)	15 ^a (2)	11 ^a (1)	12ª (2)
T _{1/2} (Q ₂) (h)	76 ^a (17)	90 ^a (7)	112 ^a (28)
Pool sizes			
Q ₁ (g C m ⁻²)	9.1 ^a (0.9)	8.6 ^a (0.4)	9.5 ^a (1.0)
Q ₂ (g C m ⁻²)	26.5 ^a (8.7)	40.0 ^a (4.9)	52.7ª (18.2)
Residence time in the system (h)	194 ^a (48)	218 ^a (22)	291ª (85)
Fractional contribution			
Current assimilation	0.43 ^a (0.07)	0.42ª (0.03)	0.40ª (0.07)
Store	0.57ª (0.15)	0.58ª (0.06)	0.60ª (0.15)
Goodness of model fit (r ²)	0.998	0.999	0.996
3.4 A connection between carbohydrate and respiratory substrate supply systems

Comparison of tracer kinetics in non-structural carbohydrates and dark respired CO2

In this section, I explore the relationships between the tracer kinetics in the dark respired CO₂ at the whole plant level and in the water-soluble carbohydrates (WSCs) of the shoot to seek further clues as to the likely chemical identity of the main mediator carbohydrate substrate feeding respiration (object 4, chapter 1). I first assessed $\Delta f_{\text{unlabelled } X}$, the deviation between the fraction of unlabelled C in a given carbohydrate species X and that in respired CO₂ ($\Delta f_{unlabelled}$ $x = f_{\text{unlabelled }Rn} - f_{\text{unlabelled }X}$) and its temporal pattern during the labelling period (Fig. 3.4.1). This analysis demonstrated that for any given carbohydrate, the magnitude of $\Delta f_{\text{unlabelled } X}$ and its temporal pattern was highly similar for all [CO₂] treatments. $\Delta f_{unlabelled}$ of sucrose ($\Delta f_{unlabelled suc}$) was generally positive throughout labelling. In addition, for glucose, the magnitude and temporal pattern of $\Delta f_{unlabelled}$ ($\Delta f_{unlabelled glc}$) closely resembled that of sucrose. Similar relationships did not exist for fructan ($\Delta f_{unlabelled fructan}$) and fructose ($\Delta f_{unlabelled fru}$), which exhibited consistently negative values. Moreover, $\Delta f_{unlabelled fructan}$ and $\Delta f_{unlabelled fru}$, tended to increase with labelling time, a pattern that was most evident for fructan (Fig. 3.4.1). On average across the three [CO₂] treatments, $\Delta f_{unlabelled suc}$ decreased with the duration of labelling from approximately 0.13 after the first labelling photoperiod to approximately 0.06 on day 7 (Fig. 3.4.1a, b and c). The time lag for dark respired CO_2 to reach the same degree of labelling as whole-shoot sucrose was around 6 h throughout labelling, an estimate associated with an uncertainty (SE) of around 2 h. This pattern was not evident for the other sugars for which $\Delta f_{\text{unlabelled } X}$ stayed constant (glucose) or even increased with the labelling duration (fructan and fructose).

The positive $\Delta f_{unlabelled suc}$ observed throughout the labelling period means sucrose was labelled faster than whole plant (or stand-scale) respiration. This observation supported Schnyder et al. (2012) that sucrose in a leaf blade was labelled faster than the whole shoot respired CO₂ in *L. perenne*. Assuming all substrates feeding respiration are derived from sucrose, the 6 h delay represents the time span between exiting the shoot sucrose pool and passage through the different metabolic events associated with, or preceding, the release of respired CO₂. These results may support the idea that sucrose is indeed the main mediator carbohydrate substrate supplying the respiratory machinery (see below). The fact that $\Delta f_{unlabelled}$ glc was similar to $\Delta f_{unlabelled suc}$ may be unsurprising as both sugars followed similar labelling kinetics, and glucose was turned over rapidly, most likely in connection with sucrose recycling

Chapter 3 Results and Discussion

(see chapter 3.2). Conversely, the results of $\Delta f_{unlabelled fructan}$ and $\Delta f_{unlabelled fru}$ demonstrate that fructan and fructose were labelled much more slowly than the whole plant respired CO₂, a disparity that even increased with labelling duration. This result is consistent with the idea that fructan and fructose did not serve as *immediate* precursors of respiratory substrates at the whole-plant level, although they had a critical mediator role via their involvement in the recycling of sucrose, as discussed in chapter 3.2.

Parameter	<i>F</i> -value	P-value		
[CO ₂] 200 µmol mol ⁻¹				
$\Delta f_{ m unlabelled}$ fructan	12.36	0.008		
$\Delta f_{ m unlabelled}$ suc	2.14	0.21		
$\Delta f_{ m unlabelled}$ glu	2.78	0.15		
$\Delta f_{ m unlabelled}$ fru	1.49	0.33		
[CO ₂] 400 µmol mol ⁻¹				
$\Delta f_{ m unlabelled}$ fructan	5.07	0.009		
$\Delta f_{ m unlabelled}$ suc	0.53	0.71		
$\Delta f_{ m unlabelled}$ glu	4.13	0.02		
$\Delta f_{ m unlabelled}$ fru	4.27	0.02		
[CO ₂] 800 μmol mol ⁻¹				
$\Delta f_{ m unlabelled}$ fructan	11.7	0.009		
$\Delta f_{ m unlabelled}$ suc	8.82	0.02		
$\Delta f_{ m unlabelled}$ glu	97.06	<0.001		
$\Delta f_{ m unlabelled}$ fru	30.38	0.001		

Table 3.4.1 Results of a one-way ANOVA, testing the differences of $\Delta f_{\text{unlabelled X}}$ among different labelling days.



Fig. 3.4.1 Deviation between fractions of unlabelled C between respired CO₂ and individual WSC of *L. perenne*. The deviation ($\Delta f_{unlabelled} \times$) between the fractions of unlabelled C in a given carbohydrate *X* and that in respired CO₂ was expressed by $\Delta f_{unlabelled} \times = f_{unlabelled} R_n - f_{unlabelled} \times$. Individual carbohydrates include sucrose (a, b, and c), fructan (d, e, and f), glucose (g, h, and i), and fructose (j, k, and l). Plants grown at [CO₂] of 200 (left), 400 (middle) and 800 (right) µmol mol⁻¹. Shown are means ± SE of chamber replicates during the whole labelling period from day 65 and the following 7 days, (*n*= 2-4). Different letters indicate significant differences at *P* <0.05.

Further comparative features of the respiratory substrate system and carbohydrate pools

Respiratory substrate-C accounted for *c*. 14% of total plant C across all [CO₂] treatments. By comparison, the average percentage of C in total WSCs in the shoot was 50% across all [CO₂] treatments (Table 3.4.2). The size of the fructan pool (in units of C) was more than two times bigger than the total size of the respiratory substrate pools ($Q_1 + Q_2$). On average, the size of the sucrose pool was close to that of Q_1 , the fast pool, and the size of sum of sucrose, glucose and fructose pools was more than two times greater than that of Q_1 , but two times smaller than

 Q_2 . Again, in comparison, the $T_{1/2}$ of Q_1 was around 5 times longer than that of the sucrose pool (12.7 h vs 2.4 h), but Q_2 had a shorter $T_{1/2}$ than the fructan pool (92.6 h vs 183.6 h).

Table 3.4.2 Plant respiratory substrate pool size and water-soluble carbohydrate (WSC) concentration in the shoot of *L. perenne*. The respiratory substrate system included a fast pool (Q_1) and a slow pool (Q_2), with their sizes predicted from the two-pool compartmental model. Contents of sucrose, fructan, glucose, and fructose were based on a total of 24 (200 and 800 µmol mol⁻¹ CO₂) or 48 (400 µmol mol⁻¹ CO₂) replicates sampled on days 66 to 69 and 72 after seed imbibition during the 7-dlong labelling experiments. Shown are means with SEs indicated in brackets. Different superscript letters within a row indicate statistically significant differences among CO₂ treatments at *P* <0.001.

Size	Atmospheric CO ₂ concentration (µmol mol ⁻¹)			
(mg C g ⁻¹ biomass-C)	200	400	800	
Respiratory substrate				
Q ₁	34ª (4)	25 ^b (4)	22 ^b (3)	
Q2	100ª (43)	116ª (48)	121ª (45)	
WSC				
Sucrose	31ª (3)	30 ^a (1)	30ª (1)	
Fructan	351ª (17)	451 ^b (10)	504° (26)	
Glucose	18ª (2)	15 ^b (1)	15 ^b (1)	
Fructose	19ª (1)	19.0ª (1)	23 ^b (1)	
Total WSC	418ª (20)	515 ^b (11)	571° (27)	

Thus, no simple match between carbohydrate fractions and respiratory pool sizes or half-lives were clearly apparent. However, carbohydrates serve not only as a substrate for respiration but also contribute to C skeletons for structural biomass synthesis (i.e. biomass growth). Therefore, the proportion of carbohydrates in biomass should be much greater than that of the respiratory substrate estimated from the tracer kinetics. Thus, assuming that water-soluble carbohydrates (WSCs) were the sole substrate for respiration, only approximately 28% of the WSCs-C was allocated to respiration. Accordingly, the remainder (72%) would be allocated to new (structural) biomass, indicating a carbon use efficiency (CUE) of 72% for the WSCs. This estimate of the CUE of WSCs was greater than that (53%) in the study of Lehmeier et al. (2008). Nevertheless, the CUE reported here was close to the estimate of CUE (~0.6-0.7) calculated by comparing the current photosynthesis and respiration of herbaceous plants in diverse environmental contexts (Thornley, 2011; Gong et al., 2017). In addition, a very high

CUE was estimated for stored carbohydrates in the vegetative plant tissues of wheat during grain filling (Gebbing et al., 1999).

The $T_{1/2}$ of Q_1 (12.7 h) was in a range that corresponds closely to that of the vacuole sucrose pool of C₃ grasses (12 to 24 h; Farrar & Farrar, 1985), indicating that perhaps vacuolar sucrose could have served an essential function in supplying substrate to respiration. In most C₃ and C₄ plant families, vacuolar sucrose in leaf blades and other photosynthetically active tissues is the primary source of assimilation exported to other plant parts during the dark period (Jenner & Rathjen, 1972; Schnyder, 1993; Lunn et al., 1999; Barro et al., 2020). Also, in parallel studies in the same experiment, Dorn (2019) found the highest sucrose concentration in leaf blades at the end of the light period and substantial decreases in that concentration until the end of the dark period. This pattern was not observed in other tissue types, highlighting some carbohydrate-related structural-functional heterogeneity within the whole shoot. At the whole shoot scale, I found no evidence for a vacuole sucrose pool based on the estimate of the $T_{1/2}$ of sucrose (~2.4 h) which was close to that of the cytoplasmic and transport pool of sucrose (see chapter 3.2). Therefore, I can only speculate that perhaps functional-structural heterogeneity within plants may have obscured a better agreement between carbohydrate and respiratory substrate pool features.

Nevertheless, the $T_{1/2}$ and size of Q_2 suggested a long-term organic carbon storage pool. Considering the size of Q_2 , fructan was probably the main source of carbon for Q_2 , as only its size was larger than Q_2 (Table 3.4.1). This interpretation aligns with that of Lehmeier et al. (2010a), although they only analyzed the pool size of fructan but did not study its labelling kinetics. As already discussed, fructan may supply carbon for the respiratory substrate system via its involvement in the recycling of sucrose (chapter 3.2.). The fact that the half-life of the fructan pool was longer than that of Q_2 (183.6 h compared to 92.6 h), could also be associated with functional structural heterogeneity within plants, or the inclusion and participation of other substrates (non-WSC) to Q_2 . It also needs to be acknowledged that tracer kinetics are often unable to resolve pools with half-lives that differ by less than 3-fold, or sometimes even within one order of magnitude (Cheesman, 1986; Lehmeier et al., 2008). For instance, proteins could form a part of Q_2 , as their half-lives in the range of 84 to 192 h (Simpson et al., 1981; Dungey & Davies, 1982) are close to that of Q_2 . Protein metabolism is intimately connected with respiratory activity (Lea & Ireland, 1999) and products of protein degradation could perhaps serve as an alternative respiratory substrate (Araújo et al., 2011).

Chapter 4 Conclusion and Outlook

This work assessed the performance of the isotope technique in dynamic ${}^{13}\text{CO}_2/{}^{12}\text{CO}_2$ labelling of plant respired CO₂ and carbohydrate fractions under sub-, current-, or double-ambient [CO₂] conditions. The data show that the continuous ${}^{13}\text{CO}_2/{}^{12}\text{CO}_2$ labelling system works well with high accuracy and precision in terms of gas exchange rate and isotope measurements over long experimental periods (up to more than 10 weeks). I demonstrated that contamination of the growth chamber atmosphere, the labelling vessel, was not greater than 4% at all [CO₂] levels analysed, despite intensive sampling and maintenance operations requiring regular access to the chamber interior. In part that precision must have been related to the maintenance of a slight atmospheric overpressure inside the chambers relative to the outside conditions, and the instalment of air-locks that reduced air exchange with the chamber interior when chambers had to be opened. Also, I observed a minimal (max. 2.6%) contamination of carbohydrates during separation with a preparative HPLC system. These minor errors were corrected by using a socalled two-chamber approach to determine the isotopic end-members for the mixing model used to quantify the proportion of labelled and unlabeled C in a sample (gas exchange component, biomass or carbohydrate). Growth conditions inside the growth chambers led to very high fructan concentrations in shoot biomass even at sub-ambient [CO₂], likely due to high photosynthetic activity at high irradiance over long days, and restricted assimilate use in growth processes due to reduced nitrogen availability in the nutrient solution of the hydroponic system used here. A stoichiometrically constrained four-pool compartmental model provided an accurate fit to the whole-shoot carbohydrate (sucrose, fructan, glucose, and fructose) tracer data and yielded an accurate quantitative description of the fundamental features of central carbohydrate metabolism at the shoot scale. This demonstrated that [CO₂] had no significant effect on the half-lives of the different carbohydrate pools. Sucrose re-synthesis from breakdown products of fructan (fructose) was similar in magnitude to sucrose neo-synthesis, but accounted for an increasing proportion of total sucrose synthesis at double-ambient [CO₂]. This result revealed an important constitutive role of fructan in central carbohydrate metabolism, a role that is likely to become even more important at future elevated [CO₂]. It is likely that fructan serves a central and constitutive role in supplying substrate for growth and respiration of C₃ grasses even during undisturbed growth. Dark respiration rate on a ground area basis (canopy respiration) was reduced at sub-ambient [CO₂], but insensitive to ambient or double-ambient [CO₂], broadly aligning with [CO₂] effects on plant biomass production. The interpretation was tentatively supported by the observation that [CO₂] had no significant effect on the specific respiration rate, i.e. respiration per unit biomass. The tracer kinetics of stand-scale dark respiration were well fitted by a compartmental model that indicated the existence of two pools (Q_1 and Q_2) that orchestrate the respiratory substrate supply system. The sizes and half-lives of both pools were independent of [CO₂]. Pool Q_1 was small in size (equivalent to ~3% of total shoot-C) and had a half-life of approximately half a day. Pool Q_2 , termed the respiratory substrate storage pool, had a (specific) size that was equivalent to ~10-12% of total shoot-C, and featured a half-life of approximately four days. Compartmental modelling indicated that ~60% of all respiratory substrate-C cycled through Q_2 at least once independently of [CO₂], meaning that Q_2 was the main source of substrate for whole plant respiration at all [CO₂] levels. A comparative analysis of the tracer kinetics of carbohydrate fractions (at the shoot level) with canopy-scale respiration suggested that sucrose was the mediator substrate that fed metabolic activity connected with respiration, while the role of the fructan pool was indirect, connected through recycling of sucrose. The analysis additionally indicated that the CUE of stored carbohydrates was comparatively high, meaning that <30% of the total water-soluble carbohydrates were lost in respiration.

Based on the methodical and physiological findings of this work, I conclude that the following research topics should be addressed in greater detail in future research. First, my methodological analyses highlight the usefulness of the continuous ${}^{13}CO_2/{}^{12}CO_2$ labelling strategy in conjunction with compartmental modelling to quantify the importance of carbon stores in whole plant metabolism in response to environmental drivers, such as CO₂ as studied here. This approach could be expanded in the future to address the physiological role and importance of stores in response to interactive environmental drivers, such as the interaction of nutrient availability and/or temperature with [CO₂]. Second, although a biochemically-based four-pool compartmental model of central carbohydrate metabolism provided a near-perfect fit to the whole-shoot carbohydrate data, the comparison with a compartmental model of the tracer kinetics in whole plant respired CO₂ was not entirely conclusive. This may have been a consequence of not accounting for alternative mediator substrates for respiration (such as protein) or functional-structural heterogeneity (e.g. differences between the locations of greatest carbohydrate storage and metabolic activity). The latter could likely be addressed by tracer kinetic studies at the organ level, with making distinctions between different developmental stages of different plant organs. Thirdly, it could be interesting to combine such studies with investigations of phloem loading and unloading in the source and sink tissues of plants.

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Atmospheric CO_2 and VPD alter the diel oscillation of leaf elongation in perennial ryegrass: compensation of hydraulic limitation by stored-growth

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Summary

• We explored the effects of atmospheric CO_2 concentration (C_a) and vapor pressure deficit (VPD) on putative mechanisms controlling leaf elongation in perennial ryegrass.

• Plants were grown in stands at a C_a of 200, 400 or 800 μ mol mol⁻¹ combined with high (1.17 kPa) or low (0.59 kPa) VPD during the 16 h-day in well-watered conditions with reduced nitrogen supply. We measured day : night-variation of leaf elongation rate (LER_{day}: LER_{night}), final leaf length and width, epidermal cell number and length, stomatal conductance, transpiration, leaf water potential and water-soluble carbohydrates and osmotic potential in the leaf growth-and-differentiation zone (LGDZ).

· Daily mean LER or morphometric parameters did not differ between treatments, but LER_{night} strongly exceeded LER_{day} , particularly at low C_a and high VPD. Across treatments LER_{day} was negatively related to transpiration ($R^2 = 0.75$) and leaf water potential ($R^2 = 0.81$), while LER_{night} was independent of leaf water potential or turgor. Enhancement of LER_{night} over LER_{day} was proportional to the turgor-change between day and night ($R^2 = 0.93$). LGDZ sugar concentration was high throughout diel cycles, providing no evidence of source limitation in any treatment.

 Our data indicate a mechanism of diel cycling between daytime hydraulic and night-time stored-growth controls of LER, buffering C_a and daytime VPD effects on leaf elongation.

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Introduction

Leaf growth is an integrating plant process (Van Volkenburgh, 1999): leaves intercept light, transpire H_2O and assimilate CO_2 in photosynthesis, supporting the growth and maintenance requirements of all parts of the plant. Conversely, leaf growth is sensitive to plant water status, which is influenced by transpiration, and depends on adequate supplies of assimilates and nutrients. As photosynthesis and transpiration respond to changing atmospheric CO_2 concentration (C_a) and humidity, one may expect fundamental effects on leaf growth (Fig. 1). Surprisingly, however, the scientific literature does not report general strong effects of atmospheric CO_2 on leaf growth (see paragraph 3 in this section), or other morphological features, particularly in conditions with growth-limiting nitrogen fertilizer availability (Reich et al., 2014), a typical situation for many terrestrial ecosystems, including grassland (LeBauer & Treseder, 2008).

Grasses provide a convenient model for studies of leaf growth, as the leaf growth-and-differentiation zone (LGDZ) is physically separated from the photosynthesizing and transpiring leaf tissues. The LGDZ is non-transpiring and entirely dependent on assimilate import, as it is enclosed within the whorl of sheaths of older fully-expanded leaves (Fig. 1). Continuous cell production,

expansion and differentiation in the LGDZ generate an efflux of near-fully mature and photosynthetically competent tissue from the enclosing sheath (Fig. 1). Leaf elongation rate is equivalent to the integral of (axial) cell expansion rates within a cell file spanning the length of the leaf growth zone (Schnyder et al., 1990). Cell expansion depends on a close coordination of several processes and state variables, including (1) cell turgor pressure that causes a tensional stress in the primary cell wall, which stretches irreversibly when a given yield threshold is exceeded, (2) passive water flow into the expanding cell driven by the water potential gradient between the cell and the water source, (3) maintenance of this gradient by continuous adjustment of the cell osmoticum, which - in turn - generates turgor pressure, and (4) metabolic processes involved in cell wall deposition and loosening, as well as synthesis of intracellular constituents (e.g. Lockhart, 1965; Ray et al., 1972; Barlow, 1986; Passioura & Fry, 1992; Martre et al., 1999; Tardieu et al., 1999; Pantin et al., 2012; Cosgrove, 2018).

Photosynthesis of C₃ plants is not saturated with CO₂ at subambient and present-day Ca (Bowes, 1993; Ainsworth & Rogers, 2007), raising the question if leaf growth can be limited, in principle, by the availability of assimilates (Ainsworth & Rogers, 2007), at least at the $C_{\rm a}$ of the Last Glacial Maximum ($\approx 200 \,\mu\text{mol mol}^{-1}$; Lüthi *et al.*, 2008). In fact, increased carbon

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Fig. 1 Schematic illustration of source and hydraulic effects of atmospheric CO_2 concentration and vapor pressure deficit (VPD) on photosynthesis– transpiration–leaf growth physiological relationships in a vegetative grass plant (adapted from Fig. 1 in Liu *et al.*, 2017). The leaf growth-anddifferentiation zone (LGDZ) is fully enclosed within the sheaths of fully-expanded leaves and comprises successive zones of cell division, expansion and differentiation arranged along the longitudinal axis of the leaf. Leaf elongation is determined by (axial) cell expansion, enabled by turgor pressure-driven water uptake. Turgor pressure is the difference between osmotic potential and water potential. Stomatal conductance is sensitive to VPD and CO_2 and represents the physiological control of transpiration that affects water potential in the growth zone. Photosynthesis is influenced directly by CO_2 and indirectly by the CO_2 effect on stomatal conductance, and may act on assimilate supply to the leaf growth zone (source-limitation) as well as on the contribution of sugars (water-soluble carbohydrates) to the osmotic potential of expanding cells.

supply at elevated C_a tends to enhance leaf elongation via the stimulation of cell expansion, cell production or both (Ferris & Taylor, 1994; Ranasinghe & Taylor, 1996; Gamage et al., 2018). However, this response is much weaker than the effect of elevated $C_{\rm a}$ on leaf photosynthesis, stomatal conductance or water-use efficiency (Ainsworth & Long, 2005), with variability in the response linked to interactions with factors such as nutrient availability (Seneweera & Conroy, 2005), vernalization (Masle, 2000), growing season (Ferris et al., 1996), developmental stage (Seneweera et al., 1995; Masle, 2000) or genotype (Masle, 2000; Thilakarathne et al., 2015). Additionally, very high concentrations of carbohydrates in leaf growth zones were observed even when plants were grown at low irradiance at contemporary C_a (Schnyder & Nelson, 1989), providing no evidence of carbohydrate substrate limitation in tall fescue (Festuca arundinacea) in those conditions. Yet, the concentration of monosaccharides, sucrose and low-molecular weight fructans contributed up to > 0.4 MPa to the osmotic potential of leaf growth zone tissue water (Schnyder & Nelson, 1987), suggesting a possibly important role of carbohydrates in osmotic regulation in the leaf growth zone. Whether or not carbohydrate availability in the leaf growth zone can be growth-limiting (as a substrate or osmoticum) over a range of C_a remains unresolved.

Possibly, variation in leaf hydraulics is the most important factor controlling variation of leaf elongation when C_a and atmospheric vapor pressure deficit (VPD) change. A strong and rapid response of leaf elongation rate (LER) to changes in evaporative demand and leaf water status, caused by alterations in temperature, air humidity or light intensity, has been observed in several grasses (Volenec & Nelson, 1982; Parrish & Wolf, 1983; Schnyder & Nelson, 1988; Ben Haj Salah & Tardieu, 1996; Clifton-Brown & Jones, 1999; Bouchabke et al., 2006) in addition to effects of edaphic conditions, including drought (e.g. Passioura, 1988; Passioura, 2002). CO2 and VPD could also indirectly influence the leaf water status of plants through their effect on stomatal conductance (gs) (Sionit et al., 1981; Morrison, 1993; Tyree & Alexander, 1993; Ainsworth & Rogers, 2007; Kimball, 2016; Manderscheid et al., 2016; Buckley, 2019). Results from free-air CO₂ enrichment (FACE) and chamber experiments under various environmental conditions show a systematic, significant decrease of gs in C3 plants at elevated CO2, producing an equivalent decrease in transpiration (Leakey et al., 2009). Such changes affected water potential, turgor pressure, and osmotic potential in the growing leaves of Phaseolus vulgaris (Ranasinghe & Taylor, 1996). However, the combined effect of atmospheric CO₂ and VPD in the growth environment on the mechanisms linking photosynthesis, transpiration and leaf growth at constant temperature have not been investigated in any detail.

One particularly conspicuous feature of leaf elongation rate is its generally strong diurnal variation, with lower rates during the day than during the night when stomata are closed and VPD is low (Bouchabke *et al.*, 2006). That variation is clearly linked to hydraulic effects in the growth zone (Tardieu *et al.*, 2018), but likely also involves metabolic controls, such as diurnal variation

of wall rheological properties or root hydraulic conductance connected with circadian oscillations (Ben Haj Salah & Tardieu, 1996; Pantin et al., 2011; Pantin et al., 2012; Caldeira et al., 2014b). Possibly, such factors can give rise to a 'stored growth' effect (Pantin et al., 2012), that has been defined as 'the ability of a cell to store up a potential for extension during periods of reduced turgor which can be converted into extra extension upon restoration of normal turgor' (Cleland & Rayle, 1972) and was observed for example by Hsiao et al. (1970) and investigated more recently by Proseus & Boyer (2008). One may predict or hypothesize that stored growth would occur at night when turgor is increased due to stomatal closure and decreased VPD. Such an effect would compensate (at least partially) the day-time depression of LER that may be caused by the effect of transpiration on plant and growth zone water potential. It is unknown, at present, if C_a modifies diurnal variation of LER and, if it does, if such variation could be explained by a stored-growth phenomenon.

To address these unknowns, we performed mesocosm experiments with three different CO₂ concentrations: 'half ambient', equivalent to the CO₂ concentration at the Last Glacial Maximum, current 'ambient', and 'double ambient', as projected for the end of this century (IPCC, 2015), combined with high or low VPD during day-time hours, to assess the effects of these environmental drivers on the photosynthesis - transpiration leaf growth relationships in perennial ryegrass (Lolium perenne L.), a major forage crop in temperate climates. Specifically, we asked: (1) Do CO₂ and VPD influence daily-total LER, final leaf length and width, and epidermal cell production and expansion? (2) Does elevated CO_2 decrease the diurnal variation of LER, consistent with predictions of CO₂ effects on stomatal conductance and transpiration? In other words: are relationships between day-time transpiration and day-time LER consistent for CO₂ and VPD effects? (3) Are CO₂ and VPD effects on daytime LER consistent with the effects of these environmental drivers on leaf water potential, osmotic potential and turgor pressure in the LGDZ? (4) What is the effect of CO₂ (and VPD) on the contribution of sugars to (diurnal adjustment of) osmotic potential in the LGDZ? And, (5) do we find evidence for night-time storedgrowth effects on LER compensating daytime CO2 and VPD effects on LER? We studied these relationships with plants growing in sward-like conditions in a culture system with optimal water supply using a nutrient solution with reduced nitrogen concentration, employing previous protocols and experience (Kavanová et al., 2008; Lehmeier et al., 2008, 2013).

Materials and Methods

Experimental design, treatments and growth conditions

The study had a 3×2 factorial design with C_a and daytime VPD as factors, and air temperature controlled at 20 °C : 16 °C during the 16 h : 8 h, day : night cycle in all treatments. Three constant CO₂ concentrations (200, 400 or 800 µmol mol⁻¹), corresponding to Last Glacial Maximum (half-ambient), present-day (ambient) and end of the century projections (double-ambient, i.e. elevated) were combined with high (1.17 kPa, 50% relative

humidity, RH) or low (0.59 kPa, 75% RH) VPD during the day, corresponding to dry or damp summer days in Central Europe. Night-time VPD was held constant at 0.46 kPa (75% RH) in all treatments. Light was supplied by cool-white fluorescent tubes and warm-white light-emitting diode (LED) bulbs with a photosynthetic photon flux density (PPFD) of 800 μ mol m⁻² s⁻¹ at plant height. Treatments were applied 13 d after seed imbibition. Disturbance of atmospheric conditions in the chambers during handling of plants were minimized by installing air-locks in chamber doors (similar to Lehmeier *et al.*, 2008), maintenance of a small overpressure in the chambers relative to the outside atmosphere, and minimizing operations during daylight hours.

Protocols for plant growth followed closely those described by Lehmeier *et al.* (2008). In brief, individual plants of perennial ryegrass (cv. 'Acento') were grown singly in plastic tubes (350 mm height, 50 mm diameter) filled with washed quartz sand (0.3–0.8 mm grain size). Pots were arranged in plastic containers (770 mm × 560 mm × 300 mm) at a density of 383 plants m⁻². The close packing of pots (Supporting Information Fig. S1) resulted in a stand-like situation at harvest (leaf area index > 5.5 after canopy closure, in all treatments).

Two containers were placed in each of four growth chambers (see next paragraph). Plants were supplied with a modified 5 mM nitrate-N Hoagland nutrient solution every 6 h by briefly flooding the containers for 9 min followed by draining by gravity (Lehmeier *et al.*, 2008). The composition of the nutrient solution was the following: 1.67 mM KNO₃, 1.67 mM Ca(NO₃)₂, 1.0 mM MgSO₄, 0.5 mM KH₂PO₄, 0.5 mM NaCl, 134 μ M FeEDTA, 46 μ M H₃BO₃, 9 μ M MnSO₄, 0.8 μ M ZnSO₄, 0.3 μ M CuSO₄, 0.1 μ M Na₂MoO₄. This composition corresponded to a nutrient solution with 33% reduced, i.e. two-thirds-strength, nitrate-N concentration relative to normal and nominal concentrations of the other nutrients (compare with Kavanová *et al.*, 2008; Lehmeier *et al.*, 2013).

The experiments were performed inside the four plant growth chambers (PGR15; Conviron, Winnipeg, Canada) that formed part of a modernized version of the gas exchange mesocosm system described by Schnyder et al. (2003). Air supply to the chambers was performed by mixing dry CO₂-free air and tank CO₂ (from Linde AG, Unterschleißheim, Germany or Carbo Kohlensäurewerke, Bad Hönningen, Germany) using mass flow controllers. CO₂ and water vapor concentration in each growth chambers were measured every 30 min by an infrared gas analyzer (IRGA, Li-840; Li-Cor, Lincoln, NE, USA). Air temperature and RH in the chambers were measured continuously with the chamber control system (CMP6050, Conviron), which was calibrated before each experimental run (see next paragraph) using external sensors. Light intensity at canopy height was measured with a quantum sensor (LI-190R; Li-Cor) and adjusted periodically. RH gradients between the top and bottom of fully-developed canopies were < 5% and neglected in further analysis.

We performed five sequential experimental runs of 10 to 12 wk duration with four growth chambers, with different treatments in each run, and allocation of treatments to different chambers between sequential runs, as in Liu *et al.* (2016) (Supporting Information Table S1). We observed no chamber effect on the parameters reported in this study. Measurements of leaf growth, leaf gas exchange and carbohydrate concentration were performed in experimental runs 1–4 and water status measurements in the last run.

Leaf elongation rate (LER)

LER was determined as in Schnyder & Nelson (1988) and Schnyder et al. (1990). All plants were in the vegetative stage and had 7 to 11 tillers. Briefly, leaf length was measured on the main tiller of eight randomly selected plants per chamber (n = 16-40per treatment), during 14 d after canopy closure (days 49-62 after imbibition of seed). Measurements were performed every day at the end of the light period on all simultaneously growing leaves (one or two leaves) per main tiller, by recording the distance between the tip of the elongating leaf and the ligule of the youngest fully expanded leaf using a ruler. The daily rate of change in blade length was taken as the measure of the mean daily rate of leaf elongation (LER, in mm h^{-1}). For comparisons among treatments the values obtained during the phase of nearmaximum, near-steady growth were used. This corresponded to the phase when the elongating leaf blade had reached $\sim 40-65\%$ of its final length (Fig. S2) and all leaf elongation was due to blade elongation. In addition, leaf length was measured at the end of the day and the end of the following night on two successive days, to obtain the LER during day (LER_{dav}) and night (LER_{night}).

At the end of the 2 wk-long measurement period plants were sampled for the estimation of additional morphological parameters (leaf length, leaf area) using a similar protocol as Liu *et al.* (2016). The IMAGE J software (Schneider *et al.*, 2012) was used for digital analysis.

Epidermal cell length and number

Two fully developed leaf blades from each of four plants per treatment were sampled at 66 d after the start of the experiment. The selected leaves were cut near the ligule and a 3 cm-long replica of the abaxial epidermis was taken in the basal region of the blade as in Schnyder *et al.* (1990) using a thin layer of 4% (w/w) polyvinylformaldehyde (Formvar 1595 E; Merck Darmstadt, Germany).

Digital images of representative sections of each replica (0.7 mm \times 4 mm) were obtained with a fluorescence microscope (BX 61; Olympus Corp., Tokyo, Japan) operated with the CELLSENSE DIMENSION software of the Centre for Advanced Light Microscopy (CALM, Technical University of Munich, Germany), at \times 10 magnification. The IMAGEJ software (Schneider *et al.*, 2012) was used to measure cell length in cell files without stomata. Replicas with fewer than 50 identifiable cells were excluded from the analysis.

Epidermal cell number (in a cell file running from the base to the tip of the leaf blade) was estimated as final leaf blade length divided by epidermal cell length. Former work with perennial ryegrass demonstrated epidermal cell length was virtually constant along the leaf blade (Schnyder *et al.*, 1990). Also, we found no differences in cell length densities near the proximal and distal margins of epidermal replicas taken from the region that corresponded to the cells/tissue that expanded during the LER measurements (Fig. S3).

Stomatal conductance and leaf- and canopy-scale transpiration

Chamber-scale canopy transpiration (E_{canopy}) was measured continuously during the 2-wk interval in which LER was determined. For each chamber, canopy transpiration (in mmol H₂O m⁻² s⁻¹) was calculated every 30 min as the difference between the H₂O fluxes at the inlet (F_{in} , in mmol H₂O s⁻¹) and outlet (F_{out} , in mmol H₂O s⁻¹) of the growth chamber, divided by the chamber ground area (s, 1.5 m²):

$$E_{\text{canopy}} = (F_{\text{in}} - F_{\text{out}})/s \qquad \qquad \text{Eqn 1}$$

These measurements were made while the RH of the chamber atmosphere was kept near the nominal level (SD \pm 0.9%) by using a high-pressure water vapor generator (FINESTFOG, Ottobrunn, Germany) that added a known amount of water vapor to the chamber air whenever the nominal RH dropped below a specified threshold (1–2% RH lower than the chamber setpoint). The chamber-based humidification system was inactivated during that period. Vapor addition rate by the vapor generator was equated with $F_{\rm in}$, as the fresh air supplied to the chambers was dry (dewpoint < –70 °C). Before each experimental run, a calibration was performed to obtain the water addition rate of the nozzles. Water vapor losses due to condensation inside the chambers were quantified by collection and weighing of the condensate and accounted for in the calculations.

In parallel, leaf-level measurements of stomatal conductance and transpiration were performed with a LI-6400 (Li-Cor) portable CO_2/H_2O gas exchange system with a clamp-on leaf cuvette on 6–12 plants per treatment. That measurement system was installed in a separate plant growth chamber (E15, Conviron). For measurements, individual plants were removed from their growth chamber, and the midsection of the youngest fully developed leaf blades of four tillers was enclosed in the 2 cm × 3 cm leaf cuvette. Stomatal conductance to water vapor (g_s , in mol $H_2O m^{-2} s^{-1}$) and leaf transpiration (in mmol $H_2O m^{-2} s^{-1}$) were measured at a leaf temperature of 21 °C and a PPFD of 800 µmol m⁻² s⁻¹. CO₂ concentration and RH in the leaf cuvette were set equal to the conditions in the growth environment. Measurements were logged once steadystate conditions for stomatal conductance and water vapor concentration were reached.

Osmotic potential, leaf water potential and turgor

Total osmotic potential in the LGDZ, leaf water potential, and turgor pressure were estimated in the treatments with C_a of 200 and 800 µmol mol⁻¹ at both high and low VPD. Samples for osmotic potential measurements were collected 2 h before the end of the day and night, on two sampling days between days 61

and 74. Six plants were randomly selected from each chamber and the LGDZ of two fully developed tillers pooled into one sample, sealed in paper bags, frozen in liquid nitrogen and stored at -18° C. The frozen samples (n=4 per treatment) were that at room temperature and sap was extracted under mechanical pressure. Osmotic potential was then measured with a vapor pressure osmometer (5100C; Wescor Inc., Logan, UT, USA). On day 76, leaf water potential of eight plants per treatment was determined with the pressure chamber technique (Scholander et al., 1965): 2 h before the end of the day individual plants were taken from a growth chamber, the youngest fully expanded leaf blade of a major tiller was cut near the ligule and immediately placed in a pressure chamber (Model 1002; PMS Instrument Company, Albany, OR, USA), following the recommendations by Turner (1981). Plants were returned to the same chamber and the protocol repeated on another tiller 2 h before the end of the night. Turgor was estimated as the difference between osmotic potential and leaf water potential, neglecting (1) the water potential gradient between the youngest fully expanded leaf and the LGDZ and (2) the fraction of apoplastic water, that were assumed to be sufficiently small or similar between treatments (Passioura, 1980; Martre et al., 2001). As they comprised the entire LGDZ and youngest fully-expanded leaf blade, our measurements ignored the turgor gradients between expanding and fully-expanded tissue in the LGDZ, and the water potential gradient between the expanding tissue and the water source (e.g. Nonami & Boyer, 1993; Martre et al., 1999; Fricke & Peters, 2002). In detailed investigations of hydraulic conductivities in vegetative plants of tall fescue - a closely related species - Martre et al. (2001, Table 1) found a water potential-gradient of 0.19 MPa between the transpiring, youngest fully-expanded leaf and the leaf growth zone, a difference that corresponded to 10% to 16% of the leaf water potential of the youngest fully-expanded leaf blade in light, in our investigations.

Water-soluble carbohydrates

LGDZ tissue was excised from two mature tillers of six plants per chamber on day 62 after imbibition and pooled together into one sample, both at the end of the day and night period, during experimental runs 1–4 (n=4 per treatment). Fresh weight of the samples was determined and samples were frozen in liquid nitrogen and stored in a freezer at -18 °C until freeze drying. Dry samples were ball-milled to a fine powder and stored again at -18 °C until extraction of water-soluble carbohydrates as in

Table 1 Results of a linear mixed model, testing the response of daily mean leaf elongation rate of *Lolium perenne* to atmospheric CO₂ concentration, daytime vapor pressure deficit (VPD) and their interaction (n = 16-40).

Factor	df	F-value	<i>P</i> -value
CO ₂	9	1.47	0.26
VPD	9	1.12	0.32
$CO_2 \times VPD$	9	1.87	0.21

Schnyder & de Visser (1999). The water-soluble carbohydrates components fructose, glucose, sucrose and fructan were separated by high-performance liquid chromatography (HPLC, Shodex Sugar KS 801 and 802; Showa Denko, Tokyo, Japan) and carbohydrates detected with a refractive index detector (Shodex RI-101). Analytical grade fructose, glucose, sucrose and inulin (all from Merck) were used as standards.

To obtain the mean degree of polymerization of fructans, fructans were separated from other water-soluble carbohydrates components by HPLC, hydrolyzed by heating in 0.02 M hydrochloric acid (HCl) for 1 h (Wolf & Ellmore, 1975), and fructose and glucose in the hydrolysate separated by HPLC and quantified as earlier. The mean degree of polymerization of fructan was obtained as fructose : glucose + 1, and used to calculate the mean molecular weight of fructan in the LGZD.

The osmotic potential of water-soluble carbohydrates in the LGDZ was estimated from the molar concentration of the individual water-soluble carbohydrates per unit tissue water in the LGDZ, assuming 40 mM = 0.1 MPa (Schnyder & Nelson, 1987).

Statistics

Linear mixed models were fitted to analyze the effect of CO₂, VPD, diel period (day or night) and their interactions on LER, morphometric traits, stomatal conductance, transpiration and carbohydrate concentration. Growth chamber, experimental run and multiple measurements on individual plants were included in the models as random effects. Type III Wald test was used for determining the significance of the fixed effects and the post hoc Tukey's HSD test was performed for pairwise comparisons among treatments. Additionally, linear models were used to test the effect of the explanatory variables on leaf water potential, osmotic potential and turgor. Finally, treatment averages were calculated and linear regression analysis was performed to determine the relationship between the different target variables. All statistical analyses were conducted in R v.3.6.1 (R Core Team, 2019). The R-packages NLME (Pinheiro et al., 2019), emmeans (Lenth, 2018) and ggplot2 (Wickham, 2016) were used for fitting linear mixed models, performing the post hoc tests and for data plotting, respectively. The number of replicates varied between measured parameters and treatments and is indicated in the figure legends and table captions.

Results

LER, final leaf length and width, and epidermal cell length and number

Daily mean LER showed no statistically significant responses to CO₂, daytime VPD or their interaction (Fig. 2a; Table 1). Likewise, morphometric traits associated with the leaf growth process, i.e. final leaf blade length and width and epidermal cell length and number (Fig. 2e–h), leaf appearance interval (Fig. S4) and the time from leaf appearance to cessation of leaf elongation (not shown) revealed no significant differences between treatments.

New Phytologist



Fig. 2 Morphometric traits of *Lolium perenne* as influenced by atmospheric CO₂ concentration at low (0.59 kPa; blue color) and high daytime vapor pressure deficit (VPD) (1.17 kPa; red): daily mean leaf elongation rate (LER) (a), LER during day (b), LER during night (c), the relationship between LER during day (LER_{day}) and during night (LER_{night}) (d), final leaf length (e), mean leaf width (f), epidermal cell length (g), and epidermal cell number (h). VPD at night was kept the same in all treatments (0.46 kPa). Symbols: circles, 200 µmol mol⁻¹ CO₂; triangles, 400 µmol mol⁻¹ CO₂; squares, 800 µmol mol⁻¹ CO₂. Significance level of the linear regression in (d): *, P < 0.05. Data points and error bars represent the mean ± SE. For details, see the Materials and Methods section.

Conversely, LER exhibited pronounced diel variation, with higher rates at night in all treatments (LER_{night} > LER_{day}, P < 0.001). The amplitude of the diel variation differed between treatments, due to a significant interaction of LER with daytime VPD (night-time VPD was the same in all treatments) and CO₂ (Table 2, P<0.001; Figs 2b,c, S5). LER_{dav} increased and LER_{night} decreased exponentially with C_a . Even though daytime VPD had no statistically significant effect (Tables 2, S2), a systematic difference in LER between VPD levels was evident. High (relative to low) daytime VPD reduced LER_{day} (-0.04 mm h⁻¹) and enhanced LER_{night} (+0.07 mm h⁻¹, averaged over all treatments) throughout the range of CO2 levels. The greatest divergence between LER_{day} and LER_{night} occurred at low C_a and high daytime VPD (LER_{night}: LER_{day} = 2.0) and the smallest at high C_a and low daytime VPD (1.3). The net result of CO₂ and daytime VPD effects was a close negative and virtually-fully compensating relationship between $\text{LER}_{\text{night}}$ and LER_{day} across all six treatments (Fig. 2d, $R^2 = 0.79$, P < 0.05).

Stomatal conductance, transpiration and relationship with $\mathsf{LER}_{\mathsf{day}}$

The g_s and transpiration were measured on the youngest fully-expanded leaf blade (Fig. 3a,b) under the same conditions of light intensity, C_a and daytime VPD as in the growth environment. The g_s strongly reacted to CO₂ concentration, VPD and their

interaction (Table 3): g_s significantly decreased with both C_a (according to a negative exponential function, P < 0.001) and VPD (P < 0.01), with a VPD-sensitivity that decreased with C_a (Fig. 3a). Accordingly, g_s was highest when VPD and C_a were low, and lowest when VPD and C_a were high; but the enhancement of g_s by low VPD (relative to high VPD) was greater at a C_a of 200 µmol mol⁻¹ (2.6-fold) than at 800 µmol mol⁻¹ (1.5-fold) (Table S3).

Leaf transpiration decreased exponentially with increasing C_a , dropping by more than 55% between 200 and 800 µmol mol⁻¹ (P<0.001; Table 3) at both VPD levels. We also noted a greater leaf transpiration at high relative to low VPD level (> 10% for all

Table 2 Results of a linear mixed model, testing the response of leaf elongation rate of *Lolium perenne* to diel period (leaf elongation during day or night), atmospheric CO₂ concentration (exponential function; see Fig. 2), daytime vapor pressure deficit (VPD) and two-way interactions (n = 16-40).

Factor	df	F-value	P-value
Day/night	100	47.2	< 0.001
CO ₂	9	5.6	0.04
Daytime VPD	9	2.2	0.17
$Day/night \times CO_2$	100	60.5	< 0.001
Day/night × daytime VPD	100	12.9	< 0.001



 CO_2 levels); but this effect was not statistically significant due to the variability of leaf transpiration (Fig. 3c; Table S3). Simultaneously, a significant curvilinear relationship ($R^2 = 0.75$; P < 0.05) existed between leaf transpiration and LER_{day}. LER_{day} decreased



sharply with leaf transpiration when leaf transpiration increased above approximately 2.5–3.0 mmol m⁻² s⁻¹ in plants grown at a $C_{\rm a}$ of 200 µmol mol⁻¹ (Fig. 3c).

Water-soluble carbohydrates

The concentration of the water-soluble carbohydrates components fructose, glucose, sucrose and fructan in the LGDZ was determined at the end of the day and of the night, in all treatments. Very high water-soluble carbohydrate concentrations (with total water-soluble carbohydrate > 53% of dry matter content) were observed in all treatments at the end of both day and night. Differences between treatments and throughout the diurnal cycle were generally small, however treatment effects were observed for the molar concentrations of the individual carbohydrates. This was the case for: (1) sucrose, which showed systematic diurnal variation with concentration decreasing by 25% on average between the end of the day and the end of the night (Fig. 4c, P < 0.001) and strongest relative decreases at low C_a ; (2) a greater fructose concentration in the low C_a treatments relative to the other C_a treatments (+21% on average); (3) a greater fructan concentration at high VPD (+43% in average, due mainly to a lower degree of polymerization relative to low VPD), which determined a greater total water-soluble carbohydrate concentration at high VPD (+16%); (4) a lower fructan concentration at 200 μ mol⁻¹ compared with the other CO₂ levels (-25% on average) and (5) a small decrease of total water-soluble carbohydrate concentration during the night (-5%) on average of all treatments), due mainly to decreases of sucrose and fructan concentration (Tables S4, S5). Conversely, glucose and fructose concentrations did not show signs of depletion during the night (Fig. 4a,b). Together, the monosaccharides accounted for 64 to 77% of the total contribution of all water-soluble carbohydrates to the osmotic potential in the LGDZ. The latter was 26-34% of the total osmotic potential, and did not differ systematically between treatments or diel periods, although the relative contribution was, on average, slightly higher at the end of the night (Fig. 4f).

Leaf water potential, osmotic potential, turgor and relationships with $\mathsf{LER}_{\mathsf{day}}$ and $\mathsf{LER}_{\mathsf{night}}$

The water potential of the youngest fully-expanded leaf blade and the osmotic potential of tissue water in the LGDZ were **Table 3** Results of a linear mixed model, testing the response of stomatal conductance (g_s) and leaf transpiration of *Lolium perenne* to atmospheric CO₂ concentration (exponential function; see Fig. 3), daytime vapor pressure deficit (VPD) and their interaction (n = 6-12).

	gs		Leaf transpiration	
Factor	F-value	P-value	F-value	P-value
CO ₂	75.7	< 0.001	39.8	< 0.001
Daytime VPD	15.6	< 0.01	0.42	0.53
$CO_2 \times daytime VPD$	13.4	< 0.01	0.04	0.84

Conditions in the leaf cuvette during measurement of g_s and leaf transpiration were the same as in the growth chamber of origin of the respective plants.

determined at the end of the day and at the end of the night in the extreme C_a treatments (200 and 800 µmol mol⁻¹ CO₂) with low or high daytime VPD (Fig. 5a,b). Turgor pressure in the LGDZ (Fig. 5c) was estimated as the difference between leaf water potential and osmotic potential, neglecting possible water potential-gradients between the LGDZ and the youngest fully expanded leaf blade.

Treatments had no effect on leaf water potential, or osmotic potential or turgor at the end of the night. However, leaf water potential (P < 0.001), osmotic potential (P < 0.01) and turgor (P < 0.001) changed significantly between the end of the night and the end of the day in each of the four treatments (Fig. 5a–c; Table 4). These changes were inversely related to CO₂ concentration (P < 0.001); that is, changes of leaf water potential, osmotic potential and turgor between end of the night and end of the day

were greater at low than at high C_a (Fig. 5a–c; Table 4). On the contrary, we found a statistically significant effect of VPD on leaf water potential (P < 0.001), but not on osmotic potential or turgor at the end of the day (Table 4).

Leaf water potential was negatively related with canopy transpiration, following a virtually identical relationship across treatments when day and night measurements were pooled (Fig. 5d; $R^2 = 0.98$; P < 0.001). Leaf water potential and osmotic potential exhibited a close proportionality (Fig. 5e; $R^2 = 0.89$; P < 0.001) that also implied a linear (and proportional) increase of turgor with leaf water potential, represented as the difference between the 1 : 1 line and the data points in Fig. 5(e). LER_{day} responded linearly to leaf water potential measured at the end of the day (Fig. 5f, $R^2 = 0.81$). That relationship was determined primarily by the effect of CO₂ on LER_{day} (P < 0.05).

We found no relationship between LER_{night} and leaf water potential or osmotic potential or turgor estimates obtained at the end of the night (Table S6, P > 0.05). But, we observed a tight relationship between the enhancement of LER_{night} relative to LER_{day} and the increase in turgor between the end of the day and the end of the night ($R^2 = 0.93$, P < 0.05; Fig. 6), with the intercept of this relationship not being significantly different from zero (P > 0.05).

Discussion

Our work demonstrates a strong diurnal oscillation of LER in *L. perenne* that was determined by (1) a variable hydraulic limitation of daytime LER driven by atmospheric CO_2 level (C_a) and



Fig. 4 Molar concentration of fructose (a), glucose (b), sucrose (c) and fructan (d) in the leaf growth-and-differentiation zone (LGDZ) of *Lolium perenne*, and osmotic potential of the total water-soluble carbohydrate (WSC) (e), and its contribution to the total osmotic potential measured in the LGDZ (f) as influenced by atmospheric CO₂ concentration at low (0.59 kPa, blue color) and high daytime vapor pressure deficit (VPD) (1.17 kPa, red), for measurements at the end of the day (light colored bars) and at the end of the night (dark colored bars). VPD at night was kept the same in all treatments (0.46 kPa). Data points and error bars represent the mean \pm SE (*n* = 4). Notice the different scales in plots (a)–(d).

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Fig. 5 Water potential of the youngest fully expanded leaf (Ψ_L) of *Lolium perenne* (n = 8) (a), osmotic potential of tissue water in the leaf growth-anddifferentiation zone, LGDZ (π_{LGDZ}) (n = 4) (b), turgor pressure (*P*) in the LGDZ, estimated as the difference between π_{LGDZ} and Ψ_L (n = 4) (c), and relationship between Ψ_L and canopy transpiration (E_{canopy}) (d), between Ψ_L and π_{LGDZ} (e) and between Ψ_L and leaf elongation rate (LER) (f) for measurements at the end of the day (light colored bars or symbols) and at the end of the night (dark colored bars or symbols). In (e) the difference between the 1 : 1 line and the data points represents the estimated turgor. Plants were grown in the presence of half-ambient (200 µmol mol⁻¹, circles) and double-ambient CO₂ (800 µmol mol⁻¹, squares) with low or high daytime vapor pressure deficit (VPD) (low VPD, 0.59 kPa, blue color; high VPD, 1.17 kPa, red) in their growth environment. Night-time VPD in the growth environment was kept the same in all treatments (0.46 kPa). Significance level of the linear regression in (d) and (e): ***, P < 0.001. Data points and error bars represent the mean \pm SE. For details, see the Materials and Methods section.

Table 4 Results of a linear model (t-values) testing the response of leaf
water potential (Ψ_L), osmotic potential in the leaf growth-and-differentia-
tion zone, LGDZ (π_{LGDZ}) and turgor pressure of <i>Lolium perenne</i> to growth
during day or night (a); and effect of atmospheric CO_2 concentration, day-
time vapor pressure deficit (VPD) and their interaction on Ψ_{L} , π_{LGDZ} and P
during growth in the day (b).

Factor	Ψ_{L}	π_{LGDZ}	Turgor
(a) Day/night	12.4***	3.0**	7.4***
	126.8***	7.9*	5.6*
(b) Daytime VPD	20.1***	2.2 ^{ns}	0.11 ^{ns}
$CO_2 \times daytime VPD$	0.3 ^{ns}	0.08 ^{ns}	0.0 ^{ns}

No significant effects were detected during night growth (P > 0.05). Significance levels: ns, not significant (P > 0.05); *, P < 0.05; **, P < 0.01; ***, P < 0.001.

(daytime) VPD, and (2) a full compensation of the daytime hydraulic limitation of elongation by a night-time enhancement of LER, that was consistent with a stored-growth effect. As a result, at daily timescale, LER and a suite of leaf morphogenetic parameters (leaf blade length and width, epidermal cell length and number) were insensitive to C_a (and daytime VPD), explaining, in part, the less-than expected growth response towards elevated CO₂. However, we did note a decrease of specific leaf area



Fig. 6 Enhancement of nocturnal leaf elongation rate (LER_{night}) over diurnal leaf elongation rate (LER_{day}) (Δ LER = LER_{night} – LER_{day}) of *Lolium perenne* as related to the turgor change (Δ P) between end of night and end of day. Symbols: circles, 200 µmol mol⁻¹ CO₂; squares, 800 µmol mol⁻¹ CO₂; blue, low vapor pressure deficit; red, high vapor pressure deficit. Significance level of the linear regression: *, *P* < 0.05. Data points and error bars represent the mean ± SE.

(SLA) with increasing CO₂ level (Fig. S4), a typical CO₂ growthresponse (Poorter & Navas, 2003; Ainsworth & Long, 2005). Furthermore, notably, epidermal cell length was close to that observed by Kavanová *et al.* (2008) in nitrogen-limited conditions with the same cultivar of perennial ryegrass.

The presence of a strong hydraulic limitation of LER_{dav} in all treatments in this work was supported by (1) the strong decreases of leaf water potential between the end of the night and the end of day, and low leaf water potential (<-1.0 MPa) at the end of the day, that was combined with (2) strong decreases of osmotic potential and turgor towards the end of the day, and (3) a significantly lower LER_{day} than LER_{night}, despite of the 4 °C lower air temperature at night. These relationships were unrelated to soil drying during the day as stands were watered frequently, resulting in a near-constant volumetric water content in the soil (water content > 20%, data not shown) and a virtually constant canopy transpiration throughout the light periods in all treatments (Fig. S6). Also, source-limitation was highly unlikely, as watersoluble carbohydrate concentration was greater than 53% of dry mass in the LGDZ in all treatments throughout the day-night cycle. Hydraulic limitation of LER_{dav} occurring independently of soil water deficit has been observed repeatedly (Tardieu et al., 2010; Pantin et al., 2012; Caldeira et al., 2014a) and was related to high irradiance (Gallagher & Biscoe, 1979), high air VPD (e.g. Parrish & Wolf, 1983; Table 2) or low nitrogen nutritional status (Radin & Boyer, 1982). Radin & Boyer (1982) observed greater reductions of daytime relative to night-time leaf expansion in sunflower, when nitrogen nutrition was limiting. We used a Hoagland-type nutrient solution with reduced nitrogen concentration (-33%) relative to the standard solution and nominal concentrations of other elements) which could have contributed to the observed reduction of LER_{day}: LER_{night}. But, betweentreatment variation of LER_{dav} was unrelated to the effect observed by Radin & Boyer (1982) as nitrogen concentration of the LGDZ was highest (Table S7) in the treatment with the greatest disparity between LER_{dav} and LER_{night}. However, we did grow plants at a high irradiance (16 h of 800 μ mol m⁻² s⁻¹ PPFD at canopy height), causing a high daytime evaporative demand particularly in the treatments with high daytime VPD. The effect of high irradiance (and high air VPD) on LER_{dav} is generally associated with strong decreases of leaf water potential and turgor in the leaf growth zone (Tardieu et al., 2010; Pantin et al., 2012).

The C_a or its interaction with VPD strongly modified the hydraulic limitation of LER_{day}, as shown by (1) the negative relationship between LER_{day} and transpiration, principally caused by a negative response of transpiration to C_a , (2) the close negative relationship between leaf water potential and transpiration that depended mostly on variation of C_a , (3) the covariation of leaf water potential and osmotic potential that was primarily driven by variation of C_a , and finally (4) the drastic decrease in turgor and LER during daytime compared to night-time. Accordingly, the effect of C_a on LER_{day} conformed with that expected for hydraulic limitation, predicted based on its effect on stomatal conductance and transpiration during daytime (Fig. 1), although the effect did not persist at the daily level due to compensating

night-time effects on leaf elongation (see 'stored growth' in the following paragraphs). Although hydraulic limitation of LER_{dav} caused by low C_a has not been reported before, a strong negative relationship between (ABA mediated) stomatal conductance and leaf elongation during daytime was demonstrated earlier (Tardieu et al., 2010). It is worth noting that the applied treatments did not alter the water relations in our system. The relationship between canopy transpiration and leaf water potential was extremely tight ($R^2 = 0.98$), suggesting a constant hydraulic conductivity and a lack of acclimation of this parameter to the different C_a and VPD environments. Also, the treatments followed a very similar leaf water potential vs osmotic potential relationship which remained unaltered throughout the day : night cycle. Similarly, sugar (and by difference, non-sugar) contributions to osmotic adjustment remained relatively constant in the diel cycle, and accounted for 26 to 34% (and 66 to 74%) of osmotic potential, independently of C_a or VPD.

Remarkably, none of the treatments showed diel variations in the concentration of fructose and glucose, the most important sugar osmoticum (64-77% of the total osmotic potential attributable to water-soluble carbohydrates). Night-time depletion of sucrose may have resulted (at least in part) from enhanced sucrose hydrolysis by invertase to generate fructose and glucose in the LGDZ (Koch, 2004; Lunn, 2008). Also, the night-time reduction of sucrose concentration in the LGDZ may have contributed to sustain carbohydrate import into the LGDZ (Schnyder & Nelson, 1988) as total water-soluble carbohydrate concentration did not decrease markedly during the night, despite of the greatly enhanced LER_{night} (particularly at low C_a) and associated enhanced growth-related water deposition (nocturnal decreases in water content of the LGDZ were not observed, data not shown). Again, these observations support the view that neither LER_{day} nor LER_{night} were source limited.

In contrast with daytime LER, nocturnal LER showed no evidence of hydraulic limitation, as LER_{night} was not significantly related with any of the hydraulic parameters assessed at the end of the dark period. But, we did observe a negative correlation between atmospheric CO2 concentration and LERnight that was however - non-causal, as reciprocal transfer of plants between CO_2 environments at the beginning of the night period did not alter their subsequent LERnight (Fig. S7; Table S8), effectively demonstrating that LER_{night} was insensitive to night-time CO₂. Clearly, therefore, differences between treatments in LER_{night} were a consequence (carryover) of daytime atmospheric conditions of CO₂ and daytime VPD. Indeed, the enhancement of $\text{LER}_{\text{night}}$ over LER_{day} was closely related to the nocturnal recovery of turgor. This LERnight response conforms with the 'stored growth' phenomenon that is reflected in above-normal growth when turgor recovers after a period of turgor loss and inhibited growth (Serpe & Matthews, 1994; Proseus & Boyer, 2008; Pantin et al., 2012). The mechanism of stored growth is not fully understood, but the phenomenon has been associated with adjustments in wall yielding properties (e.g. Serpe & Matthews, 1994). Studies with Chara corallina cells demonstrated an accumulation of unused cell wall precursors in the cytoplasm during the phase of low turgor, which were used after turgor recovery

and led to an enhancement in cell growth (Proseus & Boyer, 2006). The stored growth effect, however, may not be ubiquitous among grasses (see e.g. the independent response of LER_{night} and LER_{day} to VPD in maize, Bouchabke *et al.*, 2006) or may perhaps interact with other factors, such as source or nutritional limitation.

Although we did not analyze the kinetics of LER during the night, it is well known that abrupt changes from full light to darkness or from darkness to light (as used here and often employed in diurnal 'cycles' in growth chamber experiments), typically produce transient (< 1 h-long) enhancements or inhibitions of LER (Durand *et al.*, 1995), that are indicative of an elastic component of expansion. These short-lived transients are often followed by near constant LER during the remainder of the day or night periods, with the transient enhancements or inhibitions accounting for a relatively small portion of the overall differences between daytime and night-time elongation in well-watered conditions and constant temperature at the growth zone (Schnyder & Nelson, 1988; Durand *et al.*, 1995).

Clearly, the most remarkable result of this work was the virtually complete compensation of reduced daytime leaf expansion by night-time leaf expansion, as LER averaged over a 24 h period did not differ significantly between treatments. As the photoperiod was 16 h, a unit decrease of LER_{dav} actually required a two-unit increase of $\text{LER}_{\text{night}}$ for full compensation. These results are also consistent and indicative of a mechanism governing the daily mean or integral rate of cell wall loosening independently of the variable daytime depressions of turgor that were controlled by atmospheric conditions of CO₂ and VPD. The nocturnal enhancement of cell wall expansion must have been closely proportional to the nocturnal turgor increase, with a very similar proportionality factor (cell length increment per unit increase of turgor) for all treatments, as the enhancement of LER_{night} over LER_{dav} followed the same proportionality with the nocturnal turgor increase. Although we did not observe the processes of cell division and elongation directly, our observations of virtually constant final leaf length, leaf appearance interval, leaf elongation duration post emergence, and final epidermal cell length and number must have some implications for the underlying patterns of cell division and elongation, as LER is a function of cell division (production) and elongation rate along contiguous cell files (Schnyder et al., 1990; Kavanová et al., 2006, 2008), also with respect to CO2 effects (Masle, 2000). It is well established for perennial ryegrass (and related species such as tall fescue), grown in a wide range of environmental conditions, that the duration of individual epidermal cell expansion, from the emergence from the cell division zone to the time when it attains its' final length, is about 3 to 4 d in similar thermal environments (MacAdam et al., 1989; Schnyder et al., 1990; Kavanová et al., 2006, 2008). This would imply that individual elongating epidermal cells experienced several diurnal perturbations of cell expansion rate resulting from the observed treatment-dependent diurnal variations of LER. The simplest, most parsimonious hypothesis that can explain the relationship between the diurnal LER dynamics and diurnal epidermal cell elongation dynamics is, therefore, that epidermal cell

elongation rate exhibited proportionally the same diurnal perturbations as LER. We cannot rule out, however, based on our data alone, that there is perhaps a more complex mechanism, e.g. some complementary compensation between cell elongation rate and duration dynamics that could modify to some extent the exact relationship between LER and epidermal cell elongation rate dynamics (Masle, 2000). In addition, other works have found some CO₂ effect on cell division and expansion (e.g. Ferris et al., 1996; Masle, 2000), and it is not clear what mechanism(s) caused divergence of those results from the present. Certainly, the diel elongation responses found in this work warrant more detailed mechanistic studies with spatio-temporal detail at the level of expanding tissue, including the kinematics of growth, cell division, expansion and associated metabolic processes (e.g. Green et al., 1971; Nonami & Boyer, 1993; Martre et al., 1999; Fricke & Peters, 2002; Moulia et al., 2019).

In conclusion, this work demonstrated a close integration of daytime and night-time leaf elongation in an important forage grass under contrasting atmospheric CO₂ and VPD conditions in a controlled environment mesocosm. LER_{day} was under hydraulic control, in agreement with expected effects of atmospheric CO₂ and VPD on stomatal control, and ensuing effects on transpiration and hydraulic effects on cell wall expansion. Conversely, a compensatory growth mechanism (stored growth) controlled $\text{LER}_{\text{night}}$ in such a way that daily LER remained unaffected by atmospheric conditions of CO₂ and VPD. Source limitation was not a factor under the conditions of this experiment as water-soluble carbohydrate levels in the LGDZ were high even when atmospheric CO₂ concentration was only half-ambient. Absence of source limitation may have resulted, in part, from a relatively limiting nitrogen fertilizer supply and the high radiation received by the stands. Water supply was unlimiting, thus putative feedbacks of eventual soil drying on the leaf elongation process - that would occur more readily with high transpiration at low C_a – had no effect, but are important factors to be considered. So, building on the present findings, studies of the diel oscillation of leaf elongation and underlying mechanism should be expanded to a greater range of plant functional groups, environmental conditions (including photoperiod length, irradiance, and thermal and nutritional conditions) to further improve our understanding of the plant physiology of climate change adaptation.

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Author contributions

JCBC and HS designed the experiment. JCBC performed the research, analyzed the data and wrote the first draft. RTH, JZ and RS helped with gas exchange measurements. All authors contributed to the revision of the manuscript.

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Supporting Information

Additional Supporting Information may be found online in the Supporting Information section at the end of the article.

Fig. S1 Pot arrangement in the growth chamber.

Fig. S2 Daily leaf elongation rate of *Lolium perenne* at different leaf developmental stages as influenced by atmospheric CO_2 concentration at low and high daytime VPD.

Fig. S3 Epidermal cell length density at the margins of an individual leaf blade replica in *Lolium perenne* plants grown at different atmospheric CO_2 concentrations at low or high daytime VPD.

Fig. S4 Leaf appearance interval and specific leaf area of *Lolium* perenne as influenced by atmospheric CO_2 concentration and VPD.

Fig. S5 Time course of leaf elongation rate of *Lolium perenne* as influenced by atmospheric CO₂ concentration and VPD.

Fig. S6 Canopy transpiration of *Lolium perenne* during day-light hours as influenced by atmospheric CO_2 concentration, VPD and irrigation events.

Fig. S7 Nocturnal leaf elongation rate of *Lolium perenne* as influenced by nocturnal transfers to contrasting atmospheric CO_2 concentration.

Table S1 Experimental design.

Table S2 Response of daytime and night-time leaf elongation rate of *Lolium perenne* to atmospheric CO_2 concentration, daytime VPD and their interaction.

Table S3 Stomatal conductance and leaf transpiration of *Lolium* perenne during the day at different atmospheric CO_2 concentration and daytime VPD.

Table S4 Response of water-soluble carbohydrate concentration in the leaf growth-and-differentiation zone of *Lolium perenne* to atmospheric CO_2 concentration, daytime VPD and diel period.

Table S5 Concentration of water-soluble carbohydrates in the leaf growth-and-differentiation zone of *Lolium perenne* at the end of the day and night at different atmospheric CO_2 concentration and daytime VPD.

Table S6 Response of nocturnal leaf elongation rate of *Lolium perenne* to water potential of the youngest fully-expanded leaf, osmotic potential of the leaf growth-and-differentiation zone and turgor.

Table S7 Nitrogen concentration in the leaf growth-and-differ-
entiation zone of *Lolium perenne* for combinations of atmo-
spheric CO_2 concentration and daytime VPD levels.

Table S8 Response of nocturnal leaf elongation rate of Loliumperenne to transfer/swapping of plants between contrasting CO_2 environments.

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180 enrichment of leaf cellulose correlated with 180 enrichment of leaf sucrose but not bulk leaf water in a C3 grass across contrasts of atmospheric CO2 concentration and air humidity

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¹⁸O enrichment of leaf cellulose correlated with ¹⁸O enrichment of leaf sucrose but not bulk leaf water in a C₃ grass across contrasts of atmospheric CO₂ concentration and air humidity

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1 Summary

- The ¹⁸O composition of plant cellulose is often used to reconstruct past climate and plant function. However, uncertainty remains regarding the estimation of the leaf sucrose ¹⁸O signal and its subsequent attenuation by ¹⁸O exchange with source water during cellulose synthesis.
- We grew *Lolium perenne* at three CO₂ concentrations (200, 400 or 800 μmol mol⁻¹) and two relative humidity (RH) levels (50% or 75%), and determined ¹⁸O enrichment of leaf sucrose (Δ¹⁸O_{Sucrose}), bulk leaf water (Δ¹⁸O_{LW}), leaf cellulose (Δ¹⁸O_{Cellulose}) and water at the site of cellulose synthesis (Δ¹⁸O_{CelSynW}).
- $\Delta^{18}O_{Cellulose}$ correlated with $\Delta^{18}O_{Sucrose}$ ($R^2=0.87$) but not with $\Delta^{18}O_{LW}$ ($R^2=0.04$), due to a variable ¹⁸O discrepancy (range 2.0-9.0‰) between sucrose synthesis water ($\Delta^{18}O_{SucSynW}$, estimated from $\Delta^{18}O_{Sucrose}$) and bulk leaf water. The discrepancy resulted mainly from an RH effect. The proportion of oxygen in cellulose that exchanged with medium water during cellulose formation (p_{ex}), was near-constant when referenced to $\Delta^{18}O_{SucSynW}$ ($p_{ex-SucSynW} = 0.52\pm 0.02$ SE), but varied when related to bulk leaf water ($p_{ex-LW} = -0.01$ to 0.46).
- We conclude that previously reported RH-dependent variations of p_{ex-LW} in grasses are related to a discrepancy between $\Delta^{18}O_{SucSynW}$ and $\Delta^{18}O_{LW}$ that may result from spatial heterogeneity in ¹⁸O gradients of leaf water and photosynthetic sucrose synthesis.
- 20

Key words: *Lolium perenne* (perennial ryegrass), Barbour-Farquhar model of ¹⁸O-enrichment
 in cellulose, ¹⁸O in leaf water, sucrose and cellulose, atmospheric CO₂ concentration, relative
 humidity of air.

24 Introduction

The oxygen isotope ¹⁸O/¹⁶O ratio of plant cellulose ($\delta^{18}O_{Cellulose}$) and its enrichment above 25 source water ($\Delta^{18}O_{Cellulose}$, with $\Delta^{18}O_{Cellulose} \approx \delta^{18}O_{Cellulose} - \delta^{18}O_{Source}$) contain important 26 27 environmental and physiological information (see e.g. Roden et al., 2000; Barbour, 2007; 28 Werner et al., 2012; Gessler et al., 2014 for reviews). This is based on the fact that all oxygen 29 in cellulose ultimately originates from water (DeNiro & Epstein, 1979; Liu et al., 2016), and that evaporative ¹⁸O enrichment of water in leaves (Dongmann et al., 1974; Flanagan et al., 30 31 1991; Roden & Ehleringer, 1999; Farquhar & Cernusak, 2005; Cernusak et al., 2016) imprints an ¹⁸O signal onto photosynthetic products (Sternberg & DeNiro, 1983; Sternberg *et al.*, 1986; 32 33 Farquhar et al., 1998) used for cellulose synthesis in growing sink tissues (Barbour et al., 2000; 34 Helliker & Ehleringer, 2002; Cernusak et al., 2005). However, the exact isotopic identity of the water that dictates the ¹⁸O signal of primary photosynthetic products is still uncertain (Lehmann 35 et al., 2017) and a variable proportion ($p_{ex}p_x$, see below) of the original ¹⁸O signal in 36 photosynthetic products appears to be subsequently lost by exchange with source water 37 38 (Helliker & Ehleringer, 2002; Lehmann et al., 2017; Hirl et al., 2021), so that the relationship 39 between the ¹⁸O signal in cellulose and evaporative events determining the ¹⁸O signal in 40 photosynthetic products is still unsettled. The present paper is addressing these uncertainties, 41 and explores the underlying mechanisms, using perennial ryegrass (Lolium perenne, C₃) grown 42 in contrasting CO₂ and atmospheric humidity levels as a model plant.

43 Current mechanistic understanding of the relationship between evaporative ¹⁸O 44 enrichment of water at the site of sucrose synthesis ($\Delta^{18}O_{SucSynW}$) – the most ubiquitous primary 45 photosynthetic product and translocated sugar – and $\Delta^{18}O_{Cellulose}$ can be summarized 46 quantitatively for steady-state conditions (Barbour & Farquhar, 2000) as:

47
$$\Delta^{18}O_{Cellulose} = \Delta^{18}O_{SucSynW}(1 - p_{ex}p_x) + \varepsilon_{bio}, \qquad Eqn 1$$

where p_x denotes the proportion of unenriched source water at the site of cellulose synthesis, p_{ex} is the proportion of oxygen atoms in cellulose that have exchanged with medium water during cellulose formation at that site, and ε_{bio} is the average biochemical fractionation between carbonyl oxygen and water. The term $\Delta^{18}O_{SucSynW} + \varepsilon_{bio}$ represents the ¹⁸O enrichment of leaf sucrose above source water. In field conditions, Eqn 1 requires consideration of non-steadystates and necessitates computation of flux-weighted signals (Cernusak *et al.*, 2005; Barbour, 2007).

55 When applying Eqn. 1, it is generally assumed that p_x , p_{ex} and ε_{bio} are constant 56 parameters: p_x is often set to unity while p_{ex} is often assumed to vary within a narrow range
between 0.4 and 0.5 (Barbour, 2007; Liu et al., 2016) and ε_{bio} is equal to 27‰ (Sternberg & 57 58 DeNiro, 1983; Yakir & DeNiro, 1990). Another, almost general, assumption of previous works has been that $\Delta^{18}O_{SucSynW}$ equals the average ¹⁸O enrichment of bulk or lamina leaf water 59 ($\Delta^{18}O_{LW}$), so that (assimilation-weighted) $\Delta^{18}O_{SucSynW}$ can be replaced by $\Delta^{18}O_{LW}$ in Eqn 1. 60 This assumption is practical, as measurements (and modelling) of $\Delta^{18}O_{LW}$ are relatively 61 62 straightforward in comparison to $\Delta^{18}O_{SucSynW}$. However, this assumption is often hard to validate from cellulose ¹⁸O data because p_{ex} cannot be measured directly, and can only be 63 estimated as a fitted parameter in Eqn 1. Values of p_{ex} estimated in this way therefore absorb 64 all the uncertainty in the other parameters of the equation, including any possible error in the 65 $\Delta^{18}O_{LW} \approx \Delta^{18}O_{SucSynW}$ assumption. 66

The assumption that $\Delta^{18}O_{LW}\approx\Delta^{18}O_{SucSynW}$ has received direct support in only two 67 68 studies that compared ¹⁸O enrichment in phloem sap dry organic matter and assimilation-69 weighted bulk leaf water: one on Ricinus communis during steady-state leaf cuvette 70 measurements (Cernusak et al., 2003) and another on Eucalyptus globulus in the field 71 (Cernusak et al., 2005). Both studies found good agreement between the two signals, provided 72 that the biochemical fractionation ε_{bio} was set at 27%. However, phloem sap is not only 73 composed of sucrose and recent work by Lehmann et al. (2017) with two C₃ grasses in 74 controlled environments found that sucrose extracted from leaves was substantially more ¹⁸O enriched than 27% relative to bulk leaf water, questioning the universal validity of the $\Delta^{18}O_{LW}$ 75 $\approx \Delta^{18} O_{SucSynW}$ assumption. 76

77 Several other recent studies seem to agree that most simplifying assumptions often applied to the Barbour-Farquhar model (i.e. $\Delta^{18}O_{LW} = \Delta^{18}O_{SucSynW}$; $\varepsilon_{bio} = 27\%$; $p_x \approx 1$; $p_{ex} \approx$ 78 79 0.4-0.5) should be questioned. First, there are good indications that the biochemical fractionation ε_{bio} decreases with increasing temperature, with a virtually identical temperature-80 81 dependence in aquatic plants and in heterotrophically grown wheat seedlings (Sternberg & 82 Ellsworth, 2011) and a value of *ca*. 26.7‰ at 20°C. This temperature dependence of ε_{bio} was also required to explain interannual and seasonal variations of leaf $\delta^{18}O_{cellulose}$ in a temperate 83 84 grassland ecosystem (Hirl *et al.*, 2021). In addition, although p_x has been shown to be close to unity in trees (Cernusak et al., 2005) and in the leaf growth-and-differentiation zone of grasses 85 86 (Liu *et al.*, 2017) that is protected from evaporation, p_x is less certain in dicot species because 87 the leaves are directly exposed to evaporative conditions during their growth. Several recent 88 studies (Song et al., 2014; Liu et al., 2016; Cheesman & Cernusak, 2017; Szejner et al., 2020; 89 Hirl *et al.*, 2021) also indicated large variations in p_{ex} , when p_{ex} was estimated using Eqn 1 with $\Delta^{18}O_{SucSynW}$ replaced by $\Delta^{18}O_{LW}$, measured (or well-constrained) estimates of p_x and a 90

91 temperature-dependent ε_{bio} from Sternberg & Ellsworth (2011). Thus far, variations of p_{ex} have 92 been mainly attributed to (1) hexose phosphates going through a futile cycle with triose phosphates before cellulose synthesis (Hill et al., 1995) and an increased probability for an 93 94 oxygen atom derived from sucrose going through an exchangeable carbonyl group with each 95 turn of the futile cycle (Barbour & Farquhar, 2000; Barbour, 2007), (2) unaccounted 96 participation of non-structural carbohydrate stores in cellulose synthesis (Pfanz et al., 2002; 97 Cernusak & Cheesman, 2015) and (3) changes in turnover of non-structural carbohydrate pools 98 (Song et al., 2014).

99 Estimates of p_x are also affected by any error in the $\Delta^{18}O_{SucSynW} = \Delta^{18}O_{LW}$ assumption. 100 This is because true p_x is calculated from determinations of ¹⁸O-enrichment of water at the site 101 of cellulose synthesis ($\Delta^{18}O_{CelSynW}$), source water ($\Delta^{18}O_{Source}$, with $\Delta^{18}O_{Source} = 0$ by definition) 102 and $\Delta^{18}O_{SucSynW}$, using a two-member mixing model that has $\Delta^{18}O_{Source}$ and $\Delta^{18}O_{SucSynW}$ as its 103 endmember:

$$p_{\rm x} = 1 - \Delta^{18} O_{\rm CelSynW} / \Delta^{18} O_{\rm SucSynW}.$$
 Eqn 2

105 Importantly, reported variation of p_{ex} seems to follow environmental patterns across 106 plant functional groups, particularly with respect to relative humidity of air (RH) (Offermann 107 et al., 2011; Liu et al., 2016; Hirl et al., 2021). Relative humidity is known to generally affect 108 the ¹⁸O enrichment of bulk leaf water but also its spatial variations in leaf blades (Cernusak et al., 2016). In particular, very large variations of ¹⁸O enrichment have been found in several 109 110 monocot leaves, from base to tip and center to edge (Helliker & Ehleringer, 2000; Gan et al., 111 2002; Helliker & Ehleringer, 2002; Gan *et al.*, 2003), that may underlie variation of the $\Delta^{18}O_{LW}$ 112 versus $\Delta^{18}O_{SucSynW}$ relationship (Lehmann *et al.*, 2017).

113 Another environmental factor that deserves attention when applying Eqn 1 to biological 114 archives is atmospheric CO₂ concentration, because its rise over the last century may have 115 affected the relationship between $\Delta^{18}O_{LW}$ and $\Delta^{18}O_{SucSynW}$ and resultant estimates of p_{ex} and p_x based on the use of $\Delta^{18}O_{LW}$ (termed p_{ex-LW} in the following) instead of $\Delta^{18}O_{SucSynW}$ ($p_{ex-SucSynW}$) 116 117 in Eqn 1. Atmospheric CO₂ concentration has been shown to have a strong effect on stomatal 118 conductance (Ainsworth & Rogers, 2007; Franks et al., 2013) and consequently on transpiration 119 (Leakey et al., 2009), storage of non-structural carbohydrates (Poorter & Navas, 2003) and on the diurnal oscillation of leaf elongation (Baca Cabrera et al., 2020). All these factors can affect, 120 directly or indirectly, $\Delta^{18}O_{LW}$, $\Delta^{18}O_{SucSynW}$, $\Delta^{18}O_{Cellulose}$, and p_{xx} , estimated with either 121 $\Delta^{18}O_{LW}$ or $\Delta^{18}O_{SucSynW}$. 122

In this study, we explored the combined effects of atmospheric CO_2 concentration (200, 400 or 800 µmol mol⁻¹), relative humidity (RH, 50% or 75% during daytime) and their

interactions on: $\Delta^{18}O_{Cellulose}$, $\Delta^{18}O_{LW}$, $\Delta^{18}O_{CelSynW}$ (estimated as the $\Delta^{18}O$ of water in the leaf 125 growth-and-differentiation zone, $\Delta^{18}O_{LGDZ}$, Liu *et al.*, 2017), $\Delta^{18}O_{SucSynW}$ (estimated as 126 Δ^{18} O_{Sucrose} – ε_{bio}) and p_{ex} and p_x referenced to average leaf water (p_{ex-LW} and p_{x-LW}) and sucrose 127 128 synthesis water ($p_{ex-SucSynW}$ and $p_{x-SucSynW}$). In this, we asked specifically: (1) Do atmospheric CO₂ concentration and relative humidity or their interactions affect $\Delta^{18}O_{SucSvnW}$ and its 129 130 relationship with $\Delta^{18}O_{LW}$? (2) Do these environmental factors influence $\Delta^{18}O_{CelSynW}$? (3) How do $\Delta^{18}O_{SucSynW}$ - and $\Delta^{18}O_{LW}$ -based p_{ex} and p_x respond to these environmental factors? Finally, 131 (4) do we find diurnal variation in these parameters, i.e. between light and dark periods? 132

133

134 Materials and Methods

135 Plant material and growth conditions

136 Perennial ryegrass (cv. 'Acento') plants were grown in four plant growth chambers (PGR15, 137 Conviron, Winnipeg, Canada) in a 16 h : 8 h, day : night cycle (temperature 20/16°C), under a 138 3×2 factorial design: three atmospheric CO₂ concentration levels ('half-ambient' = 200, 'ambient' = 400 or 'double-ambient' = 800 μ mol mol⁻¹) and two daytime relative humidity 139 140 levels (low RH = 50%, high RH = 75%; nighttime RH was 75% for all treatments), as 141 previously described in Baca Cabrera et al. (2020). In brief, L. perenne plants were grown 142 individually in plastic tubes (350 mm height, 50 mm diameter) filled with washed quartz sand (0.3–0.8 mm grain size) and arranged in plastic containers ($770 \times 560 \times 300$ mm) at a density 143 of 383 plants m⁻². Plants were supplied 4 times a day with a Hoagland type nutrient solution 144 145 with reduced nitrate-N content (Baca Cabrera et al., 2020). Light was supplied by cool-white 146 fluorescent tubes and warm-white LED bulbs with a constant photosynthetic photon flux density (PPFD) of 800 μ mol m⁻² s⁻¹ at plant height during the 16 h-long light period. A total of 147 148 five sequential experimental runs were performed, resulting in five chamber scale replicates for 149 the so-called 'reference treatment' (400 µmol mol⁻¹ CO₂ / 50% RH) and three replicate 150 mesocosm-scale runs for the other treatments.

151 CO₂ and RH treatments were installed on the 13th day after seed imbibition. For this, the 152 air supplied to the chambers was mixed from dry CO₂-free air and tank CO₂ (from Linde AG, 153 Unterschleißheim, Germany or CARBO Kohlensäurewerke, Bad Hönningen, Germany), using 154 mass flow controllers. RH and temperature were controlled by the chamber control system 155 (CMP6050, Conviron, Winnipeg, Canada). CO₂ concentration and RH were measured every 156 30 min by an infrared gas analyzer (IRGA; Li-840; Li-Cor) and never deviated more than ± 5 157 µmol mol⁻¹ and $\pm 2.0\%$ relative to the set nominal value, respectively.

158

159 Sampling design and extraction of tissue water, cellulose and sucrose

Plants from each chamber scale replicate were sampled when plant canopies were closed (leaf area index >5.5, at 7-9 weeks after the beginning of the experiment). Sampling took place at c. 2 h before the end of the light and dark periods. Each time, 12 plants were randomly selected, dissected and the sampled plant material of six plants pooled in one subsample (providing two subsamples per chamber and per sampling occasion).

For tissue water extraction, the two youngest fully expanded leaf blades and the leaf growth-and-differentiation zone (LGDZ, see Fig. 1 in Baca Cabrera *et al.*, 2020) of three mature tillers per plant were excised, sealed in 12 mL Exetainer vials (Labco, High Wycombe, UK), capped, wrapped with Parafilm and stored at -18° C until water extraction. Tissue water was extracted for 2 h using cryogenic vacuum distillation as in Liu *et al.* (2016).

170 For cellulose and sucrose extraction, the two youngest fully expanded leaf blades of 171 another two mature tillers from the same plants were excised, placed into paper bags, frozen in 172 liquid nitrogen, stored at -18 °C until freeze-drying, milled and stored again at -18 °C until 173 cellulose and sucrose extraction. a-cellulose was extracted from 50 mg of dry sample material 174 by following the Brendel et al. (2000) protocol as modified by Gaudinski et al. (2005). Water-175 soluble carbohydrates were extracted from 50 mg aliquots of dry material from to the youngest 176 fully-expanded leaf blade and sucrose separated from other compounds using a preparative 177 HPLC technique similar to that described by Gebbing & Schnyder (2001).

178

179 Isotope analysis

180 Oxygen isotope composition was expressed in per mil (‰) as:

181
$$\delta^{18}O = \left(\frac{R_{\text{sample}}}{R_{\text{standard}}} - 1\right) \times 1000,$$
 Eqn 3

182 with R_{sample} the ¹⁸O/¹⁶O ratio of the sample and R_{standard} that in the international standard (Vienna 183 Standard Mean Ocean Water, V-SMOW). δ^{18} O was measured in the following compartments: 184 tissue water of leaf blades ($\delta^{18}O_{LW}$) and of the LGDZ (designated $\delta^{18}O_{CelSynW}$); and cellulose 185 and sucrose of leaf blades ($\delta^{18}O_{Cellulose}$ and $\delta^{18}O_{Sucrose}$). Furthermore, the nutrient solution (the 186 source water for plants, $\delta^{18}O_{Source}$) was sampled 1-2 times per week. $\delta^{18}O_{Source}$ was near constant 187 throughout the experiment (-9.7 ± 0.2‰ standard deviation). $\delta^{18}O_{Source}$ was used to calculate 188 ¹⁸O-enrichment above source water ($\Delta^{18}O_X$) of the different samples (X) as:

189
$$\Delta^{18}O_{\chi} = \frac{\delta^{18}O_{\chi} - \delta^{18}O_{\text{Source}}}{1 + \delta^{18}O_{\text{Source}}/1000},$$
 Eqn 4

Water samples were analyzed by cavity ring-down spectroscopy as described in Liu *et* al. (2016). 1 µL of water sample was injected into a A0211 high precision vaporizer coupled to a L2110-i-CRDS (both Picarro Inc., Sunnyvale, Ca, USA). Each sample was measured five to twelve times depending on memory effects. After every 15–25 samples, heavy and light laboratory water standards, spanning the range of δ^{18} O values in the dataset and previously calibrated against V-SMOW, V-GISP and V-SLAP, were measured for SMOW-scaling and possible drift correction. Analytical uncertainty was <0.2‰.

197 Cellulose and sucrose samples were measured by isotope ratio mass spectrometry 198 (IRMS) as in Baca Cabrera *et al.* (2021). Each sample (sucrose or cellulose) was measured 199 against a laboratory working standard carbon monoxide gas, previously calibrated against a 200 secondary isotope standard (IAEA-601, accuracy of calibration $\pm 0.25\%$ standard deviation). 201 Solid internal laboratory standards (cotton powder) were run each time after the measurement 202 of four samples for possible drift correction and for SMOW-scaling. The precision for the 203 laboratory standard was <0.3‰.

Additionally, δ^{18} O of water vapor in the growth chambers (δ^{18} O_{Vapor}) was measured by 204 cavity ring-down spectroscopy as described in Liu *et al.* (2016). Here, we measured $\delta^{18}O_{Vapor}$ 205 206 continuously during two weeks when canopies were closed, both during the light and the dark periods. $\delta^{18}O_{Vapor}$ was constant across experimental runs and treatments, but was c. 1% more 207 208 enriched during the dark period ($-14.2\% \pm 0.5\%$ standard deviation) than during the light period (-15.2% \pm 0.6% standard deviation). Interestingly, the $\delta^{18}O_{Vapor}$ and $\delta^{18}O_{Source}$ in the 209 210 chambers were quite similar to the multi-season average observed in a nearby grassland 211 ecosystem study (Hirl et al., 2019).

212

213 Statistics

In a first step, linear mixed models were fitted to test the effect of the diel period (day vs. night) 214 215 on $\Delta^{18}O_{CelSynW}$ (n=80), $\Delta^{18}O_{LW}$ (n=160), $\Delta^{18}O_{Sucrose}$ (n=70) and $\Delta^{18}O_{Cellulose}$ (n=76). All 216 available subsamples (pseudo-replicates) were included in the analysis, with growth chamber 217 and experimental run defined as the random factors. As a significant diel trend was only detected for $\Delta^{18}O_{LW}$, day and night data of $\Delta^{18}O_{CelSynW}$, $\Delta^{18}O_{Sucrose}$ and $\Delta^{18}O_{Cellulose}$ were pooled 218 219 for further analysis. In the case of $\Delta^{18}O_{LW}$, only end of day data were used in further calculations, i.e. to estimate p_{x-LW} , p_{ex-LW} or the discrepancy between $\Delta^{18}O_{SucSynW}$ and $\Delta^{18}O_{LW}$. 220 Data from individual chamber scale replications were pooled and two-way ANOVA tests used 221 to assess the effects of CO₂, RH and their interaction on $\Delta^{18}O_{CelSynW}$, $\Delta^{18}O_{LW}$, $\Delta^{18}O_{Sucrose}$, 222 $\Delta^{18}O_{Cellulose}$, p_{x-LW} , $p_{x-SucSynW}$, p_{ex-LW} , $p_{ex-SucSynW}$, and the discrepancy between $\Delta^{18}O_{SucSynW}$ and 223

224 $\Delta^{18}O_{LW}$ ($\Delta^{18}O_{SucSynW} - \Delta^{18}O_{LW}$). All statistical analyses were conducted in R v.4.0.2 (R Core 225 Team, 2020). The R packages nlme (Pinheiro *et al.*, 2019) and ggplot2 (Wickham, 2016) were 226 used for fitting linear mixed models and data plotting, respectively.

227

228 **Results**

¹⁸O enrichment of sucrose, bulk leaf water and the discrepancy between sucrose synthesis- and bulk leaf-water

¹⁸O enrichment of sucrose in leaf blades ($\Delta^{18}O_{Sucrose}$) differed significantly between RH levels, with low RH resulting in a higher $\Delta^{18}O_{Sucrose}$ (+6.4‰ on average) (P < 0.001, Fig. 1a, Table 1). An effect of atmospheric CO₂ concentration was also detected: $\Delta^{18}O_{Sucrose}$ decreased significantly with increasing CO₂ from 45.7‰ to 42.9‰ at low RH and from 39.0‰ to 37.0‰ at high RH (P = 0.03). Across treatments, $\Delta^{18}O_{Sucrose}$ did not differ significantly between samples collected near the end of the day and the end of the night (P > 0.05, Fig. 1a).

Unexpectedly, the ¹⁸O enrichment of bulk leaf water ($\Delta^{18}O_{LW}$) was not affected by RH 237 238 or the interaction between atmospheric CO₂ concentration and RH (Fig. 1b and Table 1). However, we did observe an effect of atmospheric CO₂ concentration on $\Delta^{18}O_{LW}$. That effect 239 involved a decrease of $\Delta^{18}O_{LW}$ with increasing atmospheric CO₂ concentration both when 240 measured near the end of the day (P < 0.01) and end of the night (P < 0.001). On average, 241 $\Delta^{18}O_{LW}$ decreased by 1.7‰ between the 'half ambient' and 'double ambient' CO₂ 242 concentrations. Besides, we observed a significant diurnal trend for $\Delta^{18}O_{LW}$ (P < 0.001): 243 $\Delta^{18}O_{LW}$ was higher at the end of the day (7.9-10.2%) than at the end of the night (6.7-8.8%). 244 That diurnal trend was similar for all treatments. 245

To estimate ¹⁸O enrichment of sucrose synthesis water ($\Delta^{18}O_{SucSynW}$) from $\Delta^{18}O_{Sucrose}$ ($\Delta^{18}O_{SucSynW} = \Delta^{18}O_{Sucrose} - \varepsilon_{bio}$; Barbour 2007) a constant ε_{bio} was assumed (26.7‰, as estimated from Sternberg & Ellsworth, 2011, at 20°C) for all samples collected near the end of the light period. These data indicated that sucrose synthesis water was always more ¹⁸Oenriched than bulk leaf water (Fig.1c). The discrepancy, that is $\Delta^{18}O_{SucSynW} - \Delta^{18}O_{LW}$, seemed unaffected by CO₂ (P > 0.05), but was much higher at low than at high RH (8.5‰ vs. 2.2‰; P< 0.001).

253



255 The ¹⁸O enrichment of water in the leaf growth-and-differentiation zone ($\Delta^{18}O_{LGDZ}$ taken here 256 as a proxy for $\Delta^{18}O_{CelSynW}$) was small for all treatments (0.1-0.9‰) (Fig. 1b). It decreased

200 as a promy for Z Occasynw) was small for an abauments (off obs/ob) (Fig. 10). It abereased

slightly with increasing atmospheric CO₂ (P = 0.04, 0.4‰ decrease between 200 and 800 µmol

258 mol⁻¹, on average), but did not respond to RH or the interaction of CO₂ and RH (Table 1). Also,

- 259 we observed no significant differences in $\Delta^{18}O_{CelSynW}$ between day and night (P > 0.05).
- 260

261	Table 1	Results of	`a two-way	ANOVA	testing t	he effect of	of atmospheric	CO ₂ concentration
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262 RH and their interaction on: Δ^{18} O of sucrose (Δ^{18} O_{Sucrose}), Δ^{18} O of bulk water in the leaf

263 blades ($\Delta^{18}O_{LW}$) and in the leaf growth-and-differentiation zone ($\Delta^{18}O_{CelSynW}$), the discrepancy

264 between $\Delta^{18}O_{SucSynW}$ and $\Delta^{18}O_{LW}$ ($\Delta^{18}O_{SucSynW} - \Delta^{18}O_{LW}$), $\Delta^{18}O$ of cellulose in leaf blades

265 (Δ^{18} O_{cellulose}) and p_{ex} and p_x referenced to average leaf water (p_{ex-LW} and p_{x-LW}) and sucrose

synthesis water ($p_{ex-SucSynW}$ and $p_{x-SucSynW}$), determined for closed canopies of *L. perenne*.

	CO_2		RH		CO ₂ : RH	
Parameter	<i>F</i> -value	<i>P</i> -value	<i>F</i> -value	<i>P</i> -value	<i>F</i> -value	<i>P</i> -value
$\Delta^{18}O_{\text{Sucrose}}(n=18)$	5.6	0.03	63.3	<0.001	0.2	0.68
$\Delta^{18}O_{LW}$ (day) (<i>n</i> =20)	10.7	<0.01	0.4	0.55	0.6	0.47
$\Delta^{18}O_{LW}$ (night) (<i>n</i> =20)	21.2	<0.001	3.2	0.09	1.0	0.34
$\Delta^{18}O_{SucSynW} - \Delta^{18}O_{LW}$ (<i>n</i> =18)	0.4	0.56	74.0	<0.001	1.0	0.34
$\Delta^{18}O_{CelSynW}$ (n=20)	4.7	0.04	1.7	0.22	0.0	0.95
<i>p</i> _{x-LW} (<i>n</i> =20)	3.6	0.07	2.4	0.14	0.0	0.97
$p_{x-SucSynW}(n=18)$	3.4	0.09	11.6	<0.01	0.4	0.55
$\Delta^{18}O_{\text{Cellulose}}(n=19)$	0.0	0.88	79.4	<0.001	0.5	0.47
<i>p</i> _{ex-LW} (<i>n</i> =19)	8.3	0.01	55.4	<0.001	0.1	0.72
$p_{\text{ex-SucSynW}}(n=18)$	3.3	0.09	0.1	0.71	0.2	0.70

267 The number of total canopy scale replicates (*n*) is presented for each parameter, individually.

268 Significant *P*-values are highlighted in bold print.



Fig. 1: Δ^{18} O of leaf blade sucrose ($\Delta^{18}O_{Sucrose}$) (a) Δ^{18} O of bulk water of leaf blades ($\Delta^{18}O_{LW}$, circles) or Δ^{18} O of water at the site of cellulose synthesis ($\Delta^{18}O_{CelSynW}$, diamonds) (b), and difference between Δ^{18} O of sucrose synthesis water and Δ^{18} O of bulk leaf water ($\Delta^{18}O_{SucSynW} - \Delta^{18}O_{LW}$) in the light period (c), as influenced by atmospheric CO₂ concentration at low RH and high RH. Full symbols represent values near the end of the dark period and empty symbols near the end of the light period. Measurements were performed in closed canopies of *L. perenne*. Data points and error bars represent the mean ± SE (n = 3-5).

269

The proportion of source water in the leaf growth-and-differentiation zone (p_x) was calculated using Eqn 2, with either $\Delta^{18}O_{Sucrose} - \varepsilon_{bio} (p_{x-SucSynW})$ or $\Delta^{18}O_{LW} (p_{x-LW})$ as alternative proxies for $\Delta^{18}O_{SucSynW}$. p_{x-LW} varied in a narrow range between 0.92-0.98, but was not significantly affected by CO₂, RH or their interaction (Table 1). In comparison, $p_{x-SucSynW}$ was slightly higher, but also varied in a narrow range (0.93-0.99) that was also not affected by CO₂ or its interaction with RH, but was slightly smaller at low RH compared to high RH (0.95 vs. 0.98, P < 0.01, Table 1).

277

¹⁸O enrichment of leaf cellulose and p_{ex}

¹⁸O enrichment of cellulose in leaf blades ($\Delta^{18}O_{Cellulose}$) was significantly affected by RH (+3.1‰ at low RH relative to high RH, *P* < 0.001), but effects of CO₂ concentration or the interaction of CO₂ concentration and RH were not significant (Fig. 2a, Table 1).

Using the data presented above and Eqn 1, we calculated p_{ex} alternatively as p_{ex} 282 referenced to sucrose synthesis water ($p_{ex-SucSynW}$) or leaf water (p_{ex-LW}). This showed that $p_{ex-Particle}$ 283 284 sucSynW was not significantly affected by CO₂ concentration, RH or their interaction and 285 averaged 0.52 (± 0.02 SE) (Fig. 2b). In contrast, p_{ex-LW} varied strongly between treatments from -0.01 at 800 µmol CO₂ mol⁻¹ and 50% RH to 0.46 at 200 µmol CO₂ mol⁻¹ and 75% RH. pex-LW 286 was significantly affected by both RH (P < 0.001) and CO₂ concentration (P = 0.01) (Fig. 2c). 287 Across all treatments, Δ^{18} O_{cellulose} was closely related to Δ^{18} O_{SucSynW} ($R^2 = 0.87, P \le 0$ 288 0.01, Fig. 3), but a relationship with $\Delta^{18}O_{LW}$ was not evident ($R^2 = 0.04$, P > 0.05). 289 290



Fig. 2: Δ^{18} O of leaf blade cellulose (Δ^{18} O_{Cellulose}) (a) and *p*_{ex}, calculated based on Δ^{18} O of sucrose synthesis water (*p*_{ex-} sucSynW) (b) or Δ^{18} O of bulk leaf water (*p*_{ex-} LW) (c), as influenced by atmospheric CO₂ concentration at low and high relative humidity. Δ^{18} O measurements were performed in closed canopies of *L*. *perenne*. Data points and error bars represent the mean ± SE (*n* = 3-5).



292

293 **Fig. 3** Relationship between Δ^{18} O of sucrose synthesis water ($\Delta^{18}O_{SucSynW}$) and $^{18}O_{-}$

294 enrichment of cellulose ($\Delta^{18}O_{Cellulose}$) as influenced by atmospheric CO₂ concentration

295 (circles, 200 µmol mol⁻¹; squares, 400 µmol mol⁻¹; triangles, 800 µmol mol⁻¹), at high (blue

symbols) and low relative humidity (red symbols). The dashed line and the shadowed area

indicate the values predicted with the Barbour-Farquhar model with $p_{ex}p_x = 0.5$ ($p_{x-SucSynW} =$

298 0.96 and $p_{\text{ex-SucSynW}} = 0.52$) and ε_{bio} at 18 °C (upper limit, $\varepsilon_{\text{bio}} = 27.0\%$), 20°C (dashed line, 299 $\varepsilon_{\text{bio}} = 26.7\%$) or 22°C (lower limit, $\varepsilon_{\text{bio}} = 26.4\%$). Data points and error bars represent the

- 300 mean \pm SE.
- 301

302 **Discussion**

303 Isotopic discrepancy between average leaf water and sucrose synthesis304 water

This work found no negative effect of RH on ¹⁸O enrichment of bulk leaf water ($\Delta^{18}O_{LW}$), which 305 306 was unexpected (Helliker & Ehleringer, 2002; Gan et al., 2003; Xiao et al., 2012; Cernusak et 307 al., 2016; Liu et al., 2016; Liu et al., 2017; Hirl et al., 2019). However, that result was highly 308 reproducible in replicate (n=3-5) mesocosm-scale experiments with different CO₂ 309 concentrations. Also, the result was not a peculiarity of the experimental equipment, as we 310 previously found more typical, negative RH-effects on $\Delta^{18}O_{LW}$ in a range of C₃ and C₄ grasses, 311 including a Lolium sp., in the same system (Liu et al., 2016; Liu et al., 2017). Although we are not aware of previous reports noting complete absence of an RH effect on $\Delta^{18}O_{LW}$, the effect is 312 313 notoriously variable, with significant variation between plant species and stands, including in 314 grasses (Helliker & Ehleringer, 2002; Xiao et al., 2012; Liu et al., 2017). Also, we note that the 315 treatments affected canopy and leaf morphophysiological properties, that may have indirectly influenced $\Delta^{18}O_{LW}$, affecting the apparent RH sensitivity of $\Delta^{18}O_{LW}$. For instance, the high RH 316

317 treatments led to a significantly smaller leaf area index (LAI) and lower nitrogen content per 318 unit leaf area (both P<0.01; Table S1). Both these differences could reduce the apparent RH sensitivity of $\Delta^{18}O_{LW}$ as noted in previous investigations with an isotope-enabled, process-319 320 based soil-plant-atmosphere model of a grassland ecosystem (Hirl et al., 2019). In those 321 investigations, sensitivity analysis indicated that both a decrease of photosynthetic capacity -322 which correlates with nitrogen content per unit leaf area (Kattge et al., 2009) – and LAI generate 323 an increase of $\Delta^{18}O_{LW}$, elevating $\Delta^{18}O_{LW}$ of the stands grown at high RH relative to low RH (Hirl et al., 2019). Although we cannot prove that these indirect mechanisms explained the 324 absence of an RH effect on $\Delta^{18}O_{LW}$ observed here, we note that such an absence was not a 325 necessary condition for the discrepancy between $\Delta^{18}O_{SucSynW}$ and $\Delta^{18}O_{LW}$ as a similar 326 discrepancy was also noted by Lehmann et al. (2017) in conditions with a more common 327 328 (negative) RH response of $\Delta^{18}O_{LW}$.

The negative effect of atmospheric CO₂ on $\Delta^{18}O_{LW}$ was similarly non-intuitive, since 329 330 leaf transpiration decreased with increasing CO₂ (Baca Cabrera et al., 2020), a factor that could drive an increase of $\Delta^{18}O_{LW}$ due notably to a Péclet effect (Farguhar & Lloyd, 1993; Barbour 331 332 et al., 2000; Farquhar et al., 2007). However, Hirl et al. (2019) found no evidence of a Péclet 333 effect in mixed species leaf samples from a temperate grassland ecosystem and in L. perenne 334 and Dactylis glomerata in controlled conditions. Also, Cooper & Norby (1994) did not find consistent effects of atmospheric CO₂ on $\Delta^{18}O_{LW}$ of two deciduous tree species. Besides, we 335 336 know of no other studies of the effect of growth under different atmospheric CO₂ levels on $\Delta^{18}O_{LW}$, which also limits opportunities for discussion. Importantly, effects of atmospheric CO₂ 337 on $\Delta^{18}O_{LW}$ and associated mechanisms at stand scale, including interactive effects of nutrient 338 limitation (as observed here; Table S1), have not been investigated in any detail. 339

In contrast to $\Delta^{18}O_{LW}$, ¹⁸O enrichment of sucrose ($\Delta^{18}O_{Sucrose}$) reflected very closely the 340 341 anticipated negative RH effect on ¹⁸O enrichment of water at the site of photosynthetic sucrose synthesis, i.e. $\Delta^{18}O_{SucSynW}$. That RH sensitivity was -0.25‰ per %RH on average of all 342 343 treatments. The effect appeared to be stable throughout diurnal cycles as we found no significant difference between $\Delta^{18}O_{Sucrose}$ sampled near the end of the light and dark periods. 344 345 Near-constancy of $\Delta^{18}O_{Sucrose}$ and assimilation-weighted $\Delta^{18}O_{SucSynW}$ was likely related to (1) 346 the constant environmental conditions that led to virtually constant daytime stand-scale CO₂ 347 assimilation (Fig. S1) and transpiration rates (Fig. S6 in Baca et al., 2020) and (2) the small day-night variation of $\Delta^{18}O_{LW}$ in all treatment combinations. Additionally, (3) we observed 348 349 diurnal variation of sucrose contents in leaf blades (Fig. S2), suggesting presence of a diurnal sucrose store (Sicher *et al.*, 1984; Schnyder, 1993) but no starch, which may have also helped to maintain a near-constant $\Delta^{18}O_{Sucrose}$.

The fact that $\Delta^{18}O_{SucSynW}$ was significantly higher than bulk leaf $\Delta^{18}O_{LW}$ must have 352 353 resulted from sucrose synthesis being closer to the evaporative sites or a greater proportion of 354 sucrose synthesis in the distal half of the leaf blades, where ¹⁸O enrichment of leaf water is 355 much greater (Helliker & Ehleringer, 2000; Helliker & Ehleringer, 2002; Gan et al., 2003; 356 Affek et al., 2006; Ogée et al., 2007). Indeed, all plants grew in a dense canopy situation (with 357 a LAI >5.5), which must have determined a significant decrease of incident radiation and 358 probably also of photosynthetic sucrose synthesis rate between the tip and the base of leaf 359 blades.

360

p_x , the proportion of source water at the site of cellulose synthesis, was close to 1

When expressed relative to irrigation water, i.e. nutrient solution, ¹⁸O enrichment of water at 363 the site of cellulose synthesis in the leaf growth-and-differentiation zone ($\Delta^{18}O_{CelSynW}$) was very 364 365 low in all treatments. This implied that p_x , the proportion of source water at the site of cellulose 366 synthesis, was close to 1, consistent with prior findings of Liu et al. (2017) for several C₃ and C₄ grasses. Referencing p_x to $\Delta^{18}O_{LW}$ (p_{x-LW}) instead of $\Delta^{18}O_{SucSynW}$ ($p_{x-SucSynW}$) caused only a 367 368 small underestimation of p_{x-LW} (-0.013 ±0.004 SE), due to the small leverage effect of any discrepancy between $\Delta^{18}O_{SucSynW}$ and $\Delta^{18}O_{LW}$ on estimates of p_x when $\Delta^{18}O_{CelSynW}$ is small. 369 370 However, if $\Delta^{18}O_{CelSynW}$ were higher, as may be expected for leaves of dicot species (Kahmen 371 *et al.*, 2011; Song *et al.*, 2014), any difference between $\Delta^{18}O_{SucSynW}$ and $\Delta^{18}O_{LW}$ should exert a 372 greater effect on the difference between $p_{x-SucSynW}$ and p_{x-LW} .

373

374 Is true p_{ex} a constant?

375 This work found a near-constant $p_{ex-SucSynW}$ of 0.52 (± 0.02 SE) across contrasting 376 environmental conditions. This near-constancy of $p_{ex-SucSynW}$ was also conserved when we 377 altered temperature-dependent ε_{bio} (Sternberg & Ellsworth, 2011) within the limits of 378 uncertainty for leaf temperature in our controlled environment experiments (Table S2) and 379 contrasted sharply with estimates of p_{ex-LW} in the different treatments which varied between -0.01 and 0.46. Clearly, the error made in replacing $\Delta^{18}O_{SucSynW}$ with $\Delta^{18}O_{LW}$ in Eqn 1 was the 380 381 principal (if not the only) cause of variation of p_{ex-LW} . This indicates that the treatment-related 382 variation of p_{ex-LW} was virtually fully-independent of actual variation of substrate-oxygen 383 exchange with medium water during transport to and at the site of cellulose synthesis as its

variation was eliminated almost entirely when the discrepancy between $\Delta^{18}O_{SucSynW}$ and $\Delta^{18}O_{LW}$ 384 385 was accounted for in the analysis. So, the principal mechanism underlying variation of p_{ex-LW} 386 resided in the (source) leaf and not in the growing sink tissue. The primary data of Lehmann et 387 al. (2017) are also consistent with that conclusion. However, Lehmann et al. (2017) made a 388 mistake in the estimation of $p_{ex}p_x$ (their p_{sc}), due to an error in Eqn 2 of their paper (compare 389 Eqn 1 with their Eqn 2). If we calculate $p_{\text{ex-SucSynW}}$ from their primary data using our Eqn 1, 390 taking tap water instead of crown water as the source water $\delta^{18}O$ (-10.9‰), a temperature-391 dependent Ebio (Sternberg & Ellsworth, 2011) of 26‰ for 28 °C, the temperature in their growth 392 chamber, and a $p_{x-SucSynW}$ of 0.96 as observed here – Lehmann *et al.* (2017) did not determine 393 p_{x-LW} or $p_{x-SucSynW}$ –, then we obtain a mean $p_{ex-SucSynW}$ for Dactylis glomerata and Lolium 394 perenne of ca. 0.55, close to our observation. Interestingly, our estimate of pex-SucSynW also 395 matches closely the mean p_{ex} estimate (0.53) calculated by Barbour & Farquhar (2000) from 396 the data of Hill et al. (1995). That estimate was based on an alternative approach, that is 397 measurements of randomization of ¹⁴C-labelled hexose phosphates during cellulose synthesis 398 in oak stem tissue.

399 The virtual constancy of $p_{ex-SucSynW}$ in contrasting environmental conditions is also 400 remarkable given its theoretical range of 0.2-1.0 (Barbour & Farquhar, 2000). Clearly, the pex- $_{LW}$ <0.2 observed at 400 and 800 µmol mol⁻¹ CO₂ at a RH of 50% are outside that theoretical 401 expectation. Although many studies have converged to a p_{ex} estimate of 0.4-0.5, if the original 402 403 substrate for cellulose synthesis is carbohydrates (see compilation in Cernusak et al., 2005), 404 Song *et al.* (2014) suggested that true p_{ex} may vary significantly depending on turnover time of 405 non-structural carbohydrates. When using the same approach as Song *et al.* (2014), we found 406 only minor variation of turnover time of non-structural carbohydrates in our data set (Fig. S3), 407 perhaps also contributing to the near constancy of $p_{ex-SucSynW}$. Moreover, L. perenne uses 408 different fructan series, including mixed-linkage fructans, as the primary non-structural 409 carbohydrate store (Pavis et al., 2001) and all plants had very high fructan contents (>35% of 410 dry wt) in both the leaf growth-and-differentiation zone (Baca Cabrera et al., 2020) and leaf 411 blades of fully-expanded leaves in all treatments (Fig. S2). Futile cycling of sucrose appears to 412 be very active in L. perenne (Lattanzi et al., 2012), and a high fraction of the substrate used for 413 leaf structural biomass synthesis likely first passes through the fructan pool in the growth-and-414 differentiation zones of leaves (Schnyder et al., 1988). These factors may have also contributed 415 to the magnitude and relative constancy of $p_{ex-SucSynW}$ in this study.

416 Both RH and atmospheric CO_2 concentration were strong determinants of p_{ex-LW} 417 variation in our experiments. While effects of atmospheric CO_2 concentration during 418 plant/stand growth have not been studied previously, a very similar effect of RH on pex-LW was 419 also observed in a multi-seasonal, ecosystem-scale study of modelled and observed $\Delta^{18}O_{LW}$ and Δ^{18} O_{Cellulose} in a temperate grassland (Hirl *et al.*, 2021), showing that the same effect can also 420 421 occur in natural conditions. A similar RH effect on pex-LW in the C₄ grass *Cleistogenes squarrosa* 422 (Liu et al., 2016) and in several C₃ and C₄ species (Helliker & Ehleringer, 2002) was discussed 423 by Liu *et al.* (2017). Moreover, a tendency for a similar RH effect on p_{ex-LW} is also apparent in 424 the data from R. communis presented by Song et al. (2014). It is tempting to also interpret these effects in terms of a disagreement between $\Delta^{18}O_{LW}$ and $\Delta^{18}O_{SucSynW}$. The present analysis shows 425 that the divergence between p_{ex-LW} and $p_{ex-SucSynW}$ was essentially a direct result of the 426 discrepancy between $\Delta^{18}O_{LW}$ and $\Delta^{18}O_{SucSynW}$, with $\Delta^{18}O_{SucSynW}$ well approximated by: 427

- 428 $\Delta^{18}O_{SucSynW} \approx \Delta^{18}O_{LW} (1 p_{ex-LW})/(1 0.52)$, since Eqn 5a
- 429 $\Delta^{18}O_{SucSynW}(1 p_{ex-SucSynW}p_{x-SucSynW}) = \Delta^{18}O_{LW}(1 p_{ex-LW}p_{x-LW}).$ Eqn 5b

430 Such a rough calculation only requires knowledge of (assimilation-weighted) $\Delta^{18}O_{LW}$, $\varepsilon_{\text{bio}}, p_{\text{x}}, \Delta^{18}O_{\text{Cellulose}}$ (to estimate $p_{\text{ex-LW}}$) and a theoretically- or empirically-based estimate of $p_{\text{ex-}}$ 431 432 sucSynW and can provide a quantitative guess for the magnitude of the discrepancy between $\Delta^{18}O_{LW}$ and $\Delta^{18}O_{SucSynW}$. We suggest that this hypothetical interpretation (Eqn 5a, b) should be 433 434 tested more widely across plant functional groups and environmental conditions to evaluate the magnitude of eventual discrepancies between $\Delta^{18}O_{LW}$ and $\Delta^{18}O_{SucSynW}$ and on their implication 435 in the interpretation of the relationship between (assimilation-weighted) leaf water ¹⁸O 436 enrichment and ¹⁸O enrichment of cellulose. Most certainly, a better understanding of the 437 relationship between $\Delta^{18}O_{LW}$ and $\Delta^{18}O_{SucSynW}$ will require a better knowledge of the spatio-438 439 temporal dynamics of convective and diffusive water fluxes and associated patterns of ¹⁸O-440 enrichment in leaves – both at subcellular and tissue level – and corresponding spatio-temporal 441 patterns of (photosynthetic) sucrose synthesis rates in the different parts of leaves.

442

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450

451 **Author contributions**

- 452 HS, JCBC and RTH designed the study. JCBC, RTH and JZ performed the experiments,
- 453 sampling, and sample processing, with technical assistance (see above). RS performed the
- 454 isotope analyses. JCBC analyzed the data and wrote the first draft. JCBC, HS, JO, RTH, RS,
- 455 JZ and HL contributed to the discussion and revision of the manuscript.
- 456

457 **Competing interests**

458 The authors declare that they have no competing interests

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