

Preparation of uniformly labelled ^{13}C - and ^{15}N -plants using customised growth chambers

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Methodology

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Abstract

Background Stable isotopically labelled organisms found wide application in life science research including plant physiology, plant stress and defense as well as metabolism related sciences. Therefore, the reproducible production of plant material enriched with stable isotopes such as ^{13}C and ^{15}N is of considerable interest. A high degree of enrichment (>96 atom%) with a uniformly distributed isotope (global labelling) is accomplished by a continuous substrate supply during plant growth/cultivation. In the case of plants, ^{13}C -labelling can be achieved by growth in $^{13}\text{CO}_2(\text{g})$ atmosphere while global ^{15}N labelling needs ^{15}N -containing salts in the watering/nutrient solution. Here, we present a method for the preparation of ^{13}C and ^{15}N labelled plants by the use of closed growth chambers and hydroponic nutrient supply. The method is exemplified with durum wheat.

Results In total, 330 g of globally ^{13}C - and 295 g of ^{15}N labelled T. durum wheat was produced during 87 cultivation days. For this, a total of 3.88 mol of $^{13}\text{CO}_2(\text{g})$ and 58 mmol of ^{15}N were consumed. The degree of enrichment was determined by LC-HRMS and ranged between 96-98 atom% for ^{13}C and 95-99 atom% for ^{15}N , respectively. Additionally, the isotopically labelled plant extracts were successfully used for metabolome-wide internal standardisation of native T.durum plants. Application of an isotope-assisted LC-HRMS workflow enabled the detection of 652 truly wheat-derived metabolites out of which 143 contain N.

Conclusion A reproducible cultivation which makes use of climate chambers and hydroponics was successfully adapted to produce highly enriched, uniformly ^{13}C - and ^{15}N -labelled wheat. The obtained plant material is suitable to be used in all kinds of isotope-assisted research. The described technical equipment and protocol can easily be applied to other plants to produce ^{13}C -enriched biological samples when the necessary specific adaptations e.g. temperature and light regime, as well as nutrient supply are considered. Additionally, the ^{15}N -labelling method can also be carried out under regular glasshouse conditions without the need for customised atmosphere.

Background

The use of plants and plant-derived metabolites labelled with heavy stable isotopes offers an interesting perspective in phytochemical research. Some research fields where stable isotopes found application are nutrition (1), evaluation of biosynthetic pathways (2), metabolic engineering (3), flux analysis (2, 4), accurate quantification of plant metabolites (5), studies on plant-microbe interactions (6) and untargeted metabolomics and proteomics (6-12). The increasing popularity of isotopically labelled plant material goes hand in hand with the development of complementary analytical tools e.g. NMR and high throughput GC-MS and LC-HRMS which enable differentiation between native and labelled metabolites (13, 14) as well as the availability of bioinformatics (for automated data processing) and statistical tools (for differential metabolome analysis). Therefore, protocols for the reproducible production of globally labelled biological organisms including higher plants with target stable isotopes are of increasing interest.

Up to now, a variety of *in vivo* stable isotopic labelling methodologies for different organisms were described (e.g. organelles, cells, tissue/organs and whole plants) (3, 15, 16). These approaches are based on the application of isotopically labelled substrates to the organism under investigation which further converts the labelled substrate to various downstream metabolites via endogenous enzymes. Depending on the labelling regime, the organism can be labelled uniformly or non-uniformly with different degrees of enrichment. In tracer (2, 17) or pulse-based methods (time-dependent substrate exposure) (4, 18) a specific submetabolome is labelled. These study types aim to elucidate metabolic networks or determine the rates of metabolic transformations or enzymatic reactions. On the other side, globally labelled organisms can be used for metabolome- or proteome-wide internal standardisation in untargeted -omics approaches which aim to capture the total number of the respective biochemical constituents. In addition, metabolic alterations which may, for example, be caused by various stress factors can be comprehensively investigated.

Global ^{13}C -labelling is used to produce uniformly ^{13}C -labelled plants (here termed as $^{13}\text{C}^{14}\text{N}$ plants) and requires a continuous supply of $^{13}\text{CO}_2$ throughout the whole cultivation. The $^{13}\text{CO}_2$ is applied as a substrate which is *in planta* further converted to all assimilates required to sustain plant metabolism. Therefore, airtight cultivation chambers (labelboxes) with $^{13}\text{CO}_2$ atmosphere become necessary. As the preparation of ^{13}C enriched plants affords long-term cultivations the technical equipment has to enable the regulation of growth parameters such as temperature, light and the atmospheric humidity as well as the levels of CO_2 and O_2 . Global ^{15}N -labelling is accomplished by applying ^{15}N -containing nutrient solution through the root system and ^{15}N distribution takes place by transport via xylem and the initial formation of ^{15}N -labelled glutamine in the plastids and further via transamination reactions. In such a way, ^{15}N -labelled plants (termed $^{12}\text{C}^{15}\text{N}$) can be produced. As ^{15}N is applied via the nutrient solution, hydroponic cultivation with a ^{15}N -labelled ammonium- and/or nitrate solution are prerequisite for defined labelling experiments. If both labelling regimes are used/applied in parallel under the same experimental conditions (cultivation chamber and hydroponics), the obtained plants can be expected to have similar metabolic composition and their metabolomes ($^{13}\text{C}^{14}\text{N}$ and $^{12}\text{C}^{15}\text{N}$) can be correlated. Alternatively, if only ^{15}N labelling is required the ^{15}N salts can be also provided without the use of cultivation chambers.

Up to now, the stable isotopically labelling procedures are implemented in the labs of other research groups who were successful at producing different plants with different degrees of enrichment with ^{13}C (18-22), ^{15}N (23-26) and using both elements in form of dual labelling ($^{13}\text{C}^{15}\text{N}$ plants) or single labelling of plants cultivated in parallel ($^{13}\text{C}^{14}\text{N}$ and $^{12}\text{C}^{15}\text{N}$ plants) (13, 15, 16, 27). Here, we aim to present a fully automated and robust labelling equipment implemented to produce highly ^{13}C and ^{15}N enriched wheat, cultivated in parallel under the same conditions with high N and C metabolome coverage. The presented procedure describes the detailed setup and experimental instructions for the automated long-term cultivation of wheat plants allowing plant manipulation/treatment in closed atmosphere. The procedure enables the production of plants highly enriched with ^{13}C (>96 atom%) via the $^{13}\text{CO}_2$ atmosphere whereas the CO_2 concentration can be controlled during the cultivation. The moisture levels can be

limited and CO₂ or ethylene can be removed by the optional application of a scrubbing device. The included irrigation system allows introduction of nutrient solution into the closed atmosphere. Additionally, ¹⁵N-labelling can be performed by enrichment of the nutrient solution with ¹⁵N containing salts which makes it possible to produce up to 99 atom% of ¹⁵N in ¹²C¹⁵N plants. A dual labelling combining ¹³C and ¹⁵N is also possible. The described setup can easily be extended/adjusted to cultivate other highly ¹³C enriched uniformly labelled plants. Moreover, the presented labelbox setup also allows pulse labelling experiments to study metabolic processes.

Results And Discussion

The presented method describes the cultivation of *T. durum* plants from seedling to flowering stage, for the production of ¹³C¹⁴N plants under controlled ¹³CO₂ atmosphere and ¹²C¹⁵N material by supply with ¹⁵N-labelled NH₄NO₃, Ca(NO₃)₂, and KNO₃ containing nutrient solution. To demonstrate both cultivation and the exemplary application of the plant material in an untargeted metabolomics approach, the paper is structured as follows: (a) cultivation in the labelbox for the production of ¹³C¹⁴N and ¹²C¹⁵N *T.durum* including detailed documentation of the applied and recorded parameters; (b) enrichment of the generated biomass of wheat and the consumption of CO₂ and nutrient solution; (c) assessment of the ¹³C and ¹⁵N enrichment by high resolution mass spectrometry and (d) application of the labelled wheat in an isotope assisted untargeted LC-HRMS approach with the aim to elucidate the ¹³C and/or ¹⁵N-containing metabolome. Additionally, some novel aspects as well as comparison with already presented protocols for stable isotopically labelling will be discussed.

Wheat cultivation in the labelbox

The cultivation of 4 wheat plants in each labelbox lasted 87 days in total. Defined temperature and light regimes were applied depending on the growth stage. The atmospheric parameters inside the labelbox (inner temperature, humidity as well as CO₂ and O₂ levels) were recorded in regular time-intervals (shown in Figure 1) over the whole cultivation period.

The temperature and light regime used during cultivation was adapted to 24 h cycles to simulate natural day and night rhythm for summer wheat. Therefore, the ambient temperature (outside the labelbox) was increased stepwise while the light exposure was extended in the course of cultivation. To simulate day time conditions, the plants were exposed to artificial light during the first 34 days for 12 h, while from day 35 the exposure was extended by 2 h (14 h/10 h at day/night). The ambient temperature setpoints were increased during the cultivation in three steps (Figure 1b). In parallel, the temperature inside the labelbox was recorded with the integrated temperature sensor. The inner temperature reached ambient conditions during night, while it was up to 10°C higher than that of the ambient climate chamber during the day, due to the energy provided by the external light source, regardless of the time point during cultivation (Figure 1c).

The first two monitored growth stages were (Z1) seedling development and (Z2) tillering (Figure 6a, b). During this period, which represents about one third of the cultivation time, the temperature inside the labelbox was 10°C during night and 22°C during the day (Figure 1c) resulting in 3 to 6 tillers per plant. During that stage, the daily CO₂ consumption was between 50 to 100 ml (25 to 50 loops) of CO₂ per day and CO₂ levels did not increase during night (Figure 1c). Further, during the day the O₂ levels were slightly below 20% while the relative humidity was recorded to be 53%. During the night intervals, Peltier cooling was necessary to keep relative humidity at the maximum allowed 70% which depicts the higher humidity through the release of H₂O as plant transpiration product via the stomata.

The first temperature increase initiated the stem elongation stage (Z3) (Figure 6c). The resulting increase in biomass is reflected by an elevated CO₂ consumption during the day and higher CO₂ levels in the dark (up to 1500 ppmv). CO₂ was not removed from the atmosphere in the labelbox as it was quickly assimilated after starting the day time period and this helped to save expensive ¹³CO₂. The O₂ level slightly increased in comparison to the Z1/ Z2 stage but was still below 21%. High O₂ levels would reduce the rate of photosynthesis as shown in (28). It was also observed that the increase in biomass and the associated elevated transpiration and respiration steadily raised relative humidity levels during the day to reach ca. 64% at day 40 (Figure 1c). Similar records were obtained in the ¹⁵N-labelbox.

During growth stages Z4-6 (corresponding to booting, heading and anthesis), the recorded parameter levels were maintained at setpoint levels. O₂ levels and relative humidity reached and stayed at max. values with ~21% and 70%, respectively. With the biomass, the increase in CO₂ consumption was further observed to rise further up to ~2.3 l (1100 loops) at day 66. High demand on CO₂ can also be seen from short intervals between dosages to keep CO₂ at an average level of 400 ppmv (Figure 1c).

The harvest period lasted a total of 14 days. Approximately 6 days (144h) after first anthers appeared, the respective ear and the adjacent stem and leaves were cut, removed from the box and frozen in liquid nitrogen. As expected, the removal of plant material from the labelbox resulted in a decrease of CO₂ consumption, O₂ levels and relative humidity (Figure 1c), starting from day 71.

Summarised, the recorded cultivation conditions are in accordance with the applied temperature- and light regimes and also reflect both the increase in biomass as well as photosynthesis and respiration activity during growth of the plants in the labelbox.

Yield of biomass and consumption of ¹³CO₂ and ¹⁵N-enriched nutrient solution

In total, 330 g of ¹³C¹⁴N- and 295 g of ¹²C¹⁵N-labelled plant fresh weight (FW) were harvested from two labelboxes. The mass fractions obtained for different plant organs was similar for both labelboxes. Roots

represented the greatest mass fraction (50%), followed by stems (ca. 30%), leaves and ears (ca. 10% each) (Figure 2).

The total water content in FW was determined to be between 64% and 70% for different wheat organs. After freeze drying, the rest moisture levels of the plant organs were below 5%. The consumption rate of the nutrient solution in the ^{15}N -labelled wheat was 24.4 ml/g FW and 870 mg of ^{15}N was consumed in total. In the ^{13}C -experiment, 3.81 mol of $^{13}\text{CO}_2$ were consumed which corresponds to a consumption rate of 11.5 mmol $^{13}\text{CO}_2$ /g FW (Additional file 1: equation (S1, S4)). The consumed amounts of ^{15}N or $^{13}\text{CO}_2$ are in good agreement with the literature (29-31).

Determination of the degree of ^{13}C - and ^{15}N enrichment by LC-HRMS

The degree of isotopic enrichment with ^{13}C and ^{15}N , were determined by LC-HRMS in $^{13}\text{C}^{14}\text{N}$ - and $^{12}\text{C}^{15}\text{N}$ wheat ear extracts respectively. For this, the LC-HRMS data was firstly processed by the MetExtractII software (14). Then, ca. 9 highly abundant metabolites from the obtained list that contains truly wheat-derived metabolites were selected and used to determine the degree of enrichment. A detailed annotation or identification was beyond of the scope of this study. However, as ^{13}C - and ^{15}N -labelling was employed the number of C and N atoms contained in the target compounds were known exactly. The considered metabolites contained between 9 and 14 carbon atoms and 1 to 4 nitrogen atoms, respectively. Further, the isotopologue abundance distribution for each metabolite in measured mass spectra were examined manually with Xcalibur 4.0 (Thermo Fisher Scientific, Bremen, Germany) and exemplified with one tested compound (tryptophan) in Figure 3. All isotopologues of the same metabolite show the same retention times as they coelute from the chromatographic column.

For calculation of the degree of enrichment, peak height intensities at three different regions of chromatographic peak (centre, front and tail of peak) were considered. MS peak height intensities $>10^6$ were taken into account to accurately determine the degree of isotopic enrichment (32).

The degree of enrichment (E_X) for native and labelled forms was calculated according to **equation (4)** which is derived from the binominal distribution equation (33). R_X represents the intensity ratio of the inspected isotopologues of a molecule. The count C/N is the number of carbon or nitrogen atoms contained in the molecule. The $I_{M+0.997}$ represents the intensity of the low abundant ^{15}N mass peak in native samples.

[Please see the supplementary files section to view the equations.] (4)

(5)

The degrees of ^{13}C and ^{15}N in *T. durum* cultivars were determined to be (98.4 ± 0.1) atom% and (99 ± 0.2) atom%, respectively. Similar levels were obtained for earlier labelling experiments with *T. aestivum* (Table 1). In comparison to other genotypes, the cultivar Apogee had a lower degree of enrichment for ^{13}C and ^{15}N . This might be explained with the fact that initially native seeds were used. In comparison to other genotypes, the Apogee is a “dwarf wheat”. The amount of produced biological material per seed is ca. 2.8 times lower for Apogee wheat than for Karur plants. Therefore, the fraction of ^{12}C and ^{14}N per seed that originated from the seeds is higher in Apogee than in the other cultivars.

Table 1 Determined degree of enrichment after cultivation of ^{13}C - and ^{15}N -labelled *T. aestivum* and *T. durum* cultivars as well as the enrichment in native analogues.

Higher abundant stable isotope	Sample type	Species	Genotype	Nr. of metabolites	E_x [atom%]	Standard deviation [atom%]	Standard error of the mean [atom%]
^{13}C	Labelled	<i>T. aestivum</i>	Remus	9	98.4	0.1	0.02
			Apogee	9	96.7	0.2	0.05
		<i>T. durum</i>	Karur	9	98.4	0.1	0.02
^{12}C	Native	<i>T. aestivum</i>	Remus	9	98.91	0.75	0.14
			Apogee	9	98.99	0.09	0.02
		<i>T. durum</i>	Karur	9	99.00	0.09	0.02
^{15}N	Labelled	<i>T. aestivum</i>	Remus	8	99.4	0.3	0.05
			Apogee	8	95.6	0.5	0.13
		<i>T. durum</i>	Karur	8	99	0.2	0.04
^{14}N	Native	<i>T. aestivum</i>	Remus	6	99.7	0.1	0.02
			Apogee	7	99.6	0.1	0.02
		<i>T. durum</i>	Karur	8	99.7	0.1	0.02

Annotation of the global metabolome

Extracts of the stable isotopically labelled wheat (^{13}C and ^{15}N separately) were used to internally standardise native wheat extracts. These sample mixtures were measured with LC-HRMS and processing

of the obtained raw data was performed with the in-house developed MetExtract II software (7, 8, 14, 34). The program automatically performs an efficient filtering of all non-wheat derived metabolites by extracting metabolic features based on their chromatographic coelution and isotopic pattern (which is characteristic to the mixture of native and labelled extracts) (14). The LC-HRMS data of the ^{13}C -experiment contained 1467 ions (global metabolome including all N-containing ions) while 367 ions were found in the ^{15}N -experiment (only N-containing ions were considered). In the next step, both feature lists were merged into one matrix by combining ions with same characteristics. For this, m/z values and retention times of native isotopologues containing ^{12}C and/or ^{14}N were considered in both experiments. This combined matrix, encompassed 1621 ions in total. In order to remove false positive ions from the N metabolome (e.g. those containing N but no C), the isotopologue pattern of each metabolite putatively containing N was additionally curated manually by the aid of the MZmine software (35) and the isotopologue pattern was reviewed in all experiments (e.g. in native, ^{13}C and ^{15}N labelling experiment similar isotopologue patterns were expected as those shown in Figure 3).

Finally, the data set contained 1519 ions which were assigned to 652 unique metabolites with a mass range between m/z 116 and 875 (Figure 4). Amongst the detected metabolites, 143 contained nitrogen. As illustrated in the feature plot of Figure 4, metabolites containing nitrogen atoms tended to elute during the first 15 minutes of the chromatographic run. With respect to number, the majority of the detected N-containing metabolites contain 1, 2 or 3 nitrogens.

Applications of the presented method and comparison with existing protocols

Previously, two types of equipment for stable isotopic labelling of plants with ^{13}C were described. The first employs commercially available, smaller sized automatic cultivation chambers with a built-in regulation module for light and temperature regimes (16, 20, 36). The second one makes use of (in-house designed) labelling devices placed in bigger sized climate chambers, climate rooms or glasshouses which provide an external light and temperature regulation (13, 15, 18, 19, 27). Both approaches enable continuous monitoring and regulation of atmospheric parameters. Enrichment with ^{15}N or ^{34}S does not require systems with an enclosed atmosphere but rather hydroponic cultivation systems (23-25, 37). Generally, both equipment types are in principle suitable to perform pulse (16, 18) and uniformly labelling (13, 20, 38) of plants at different enrichment degrees.

The method described here employs a customised labelbox that can be purchased commercially. In contrast to the existing approaches the presented method enables the manipulation of plants during cultivation by the use of four gloves which are tightly attached to the labelbox. Additionally, a gate compartment with an air lock system constitutes a part of the labelbox, enabling the introduction or removal of manipulation and sampling tools as well as sampled plant material from the labelbox without contaminating the inner atmosphere with ambient $^{12}\text{CO}_2$ during cultivation.

The presented setup may be also used to mimic abiotic stress - e.g. low or elevated nutrient supply or targeted control of O₂ or CO₂ levels (16). In combination with the application of stable isotopically labelled ¹³CO₂ the thereby caused molecular alterations may be effectively investigated. Further, the presented equipment is perfectly suited for pulse labelling or - to reduce experimental costs - uniform labelling with lower degrees of enrichment. The latter can be achieved with carbon dioxide showing a lower degree of ¹³C enrichment. In addition, the monitoring of both ¹²CO₂ and ¹³CO₂ can be performed in one single box. For pulse labelling, synthetic air consisting of ¹⁴N₂ and ¹⁶O₂ only might be replaced by CO₂-containing air and ¹³CO₂ can be dosed according to the needs of the respective experiment.

While the costs for ¹⁵N-salts are rather moderate, the ¹³CO₂ labelling is cost demanding but affordable on a long-run. For example, 330 g of wheat plant material were produced from ~100 L ¹³CO₂ in a single labelbox, giving enough material for global internal standardisation of ~10⁵ analytical samples according to the described LC-HRMS workflow. This would correspond to ca. 0.1 euros per sample for global internal standardisation. For making sure that the labelled plant material can be used over a long period, it should be freeze-dried to less than ca. 5% rest moisture and stored at -80°C. To our experience, under such conditions, the plant material may be preserved for several years without a significant change of its metabolic composition.

Conclusion

The method presented in this article enables the reproducible production of stable isotopically labelled plant material under controlled conditions. By the use of a customised growth chamber (labelbox), the composition of the atmosphere inside the labelbox including ¹³CO₂ can be pre-selected and regulated during long-term cultivation. By this, degrees of global ¹³C enrichment as high as 98% can be achieved. Furthermore, with the presented equipment, plants can also be treated during cultivation without interfering with the labelling process. Complementary ¹⁵N-labelling is facilitated by watering with ¹⁵N-enriched nutrient solutions, which can also be done in the greenhouse without the need of the labelbox.

The globally labelled plant material can be used as internal reference in metabolism-related plant research and phytochemical analysis including metabolomics, proteomics and lipidomics (34, 39-42). The generated plants can also be further processed to generate fractions of tailored substance classes or to isolate pure phytochemicals. Moreover, the labelled plants or generated fractions can be used for internal standardisation of experimental native samples to improve accuracy and reliability in both qualitative and quantitative analysis (43, 44).

While here, method performance characteristics have been evaluated for global uniform ¹³C- and ¹⁵N-labelling of cereals, the presented approach can also easily be adapted to culture other plant species or used for short-term labelling, the production of partly enriched plant material or kinetic studies using alternate exposure to ¹²CO₂ and ¹³CO₂ in pulsed time intervals (18).

Methods

This chapter includes (1) a description of the technical equipment used for the global ^{13}C - and ^{15}N -labelling in detail; (2) a presentation of the cultivation procedure adapted to produce globally labelled wheat from seedling to flowering stage and (3) a sample preparation workflow for the applied untargeted LC-HRMS based metabolomics approach.

Description of the labelling equipment

This section is related to the functional description of the cultivation process to produce globally ^{13}C and ^{15}N labelled plants using the PhytolabelBox equipment (ECH, Halle, Germany). All symbols contained in this section are related to Figure 5.

The plants were cultivated in two specially designed cultivation chambers, so called labelboxes \boxtimes . Each of these 214-L Plexiglas labelboxes is equipped with two internal fans as well as a sensor assembly **S** for temperature, relative humidity and overpressure measurement. Each labelbox also contains 4 openings tightly sealed with gloves (Glove Box Glove Jugitec® H, chlorosulfonated polyethylene, JUNG RUBBERTEC) allowing plant manipulation during cultivation. Control and regulation of cultivation parameters (e.g. CO_2 supply, overpressure and atmospheric humidity), which are essential to achieve and maintain adequate conditions for plant growth were implemented by different modules, which are together with the labelbox termed as PhytolabelBox equipment. These modules are navigated by the control unit \boxtimes based on the user predefined parameter setpoints via the software on the PC \boxtimes . Separate from this, the nutrient supply is manually carried over the irrigation delivery module \boxtimes .

Gas supply and control module: CO_2 is supplied from the gas bottles \boxtimes which are connected via stainless steel and copper lines to the labelbox \boxtimes . Since carbon labelling was realised in one of the boxes (^{13}C -labelbox), this is supplied with $^{13}\text{CO}_2$ gas while the ^{15}N -labelbox was operated with native $^{12}\text{CO}_2$. Each gas bottle is equipped with a pressure reducer, which was set to 1 bar. The amount of the introduced CO_2 is balanced by dosage loops $\boxed{2b}$ (ca. 2 ml at 1 bar) located in the control unit \boxtimes . The dosage loops $\boxed{2b}$ are covered with magnet valves $\boxed{2a,c}$ from both sides (in- and outlet). These valves are opened reciprocally in short time intervals during gas dosing operation. The pressure difference between the pressure reducer at

the gas bottle and the interior of the labelbox forces the CO₂ into the loop space and further into the labelbox \boxtimes . The CO₂ [ppmv] measurement was performed by IR absorption using separate cuvettes 2e for ¹³CO₂ and ¹²CO₂. Simultaneously, the O₂ [%] is also measured with an electrochemical sensor \boxtimes 2f, which is located in the control unit. Immediately before measurement, the sensor and cuvettes are rinsed with air from the labelbox to achieve a homogeneous gas sample. The labelbox is always operated at slight overpressure (the difference between ambient and inner pressure) of 10 ± 2 mbar to prevent ambient air from entering the closed and controlled atmosphere.

The overpressure levels are constantly recorded by the integrated sensor S in each labelbox \boxtimes and the pressure signal status is continuously transferred via the control unit \boxtimes to the PC \boxtimes . If the overpressure level decreases below the allowed minimum setpoint (slight gas leaks are expected and occur), the dosing of synthetic air (mixture of N₂ + O₂) is activated. To this end, a magnet valve \boxtimes 2d, which is located in the control unit opens and the synthetic air is dosed into the labelbox \boxtimes until the overpressure level reaches the setpoint again. If the pressure in the cultivation box exceeds the maximum allowed overpressure of 12 mbar, a magnet valve \boxtimes opens to release excess air. This mechanism is controlled by the software and is additionally used for the air exchange in case CO₂ and/or O₂ levels exceed the defined maximum values. To prevent damage to the plexiglas labelboxes by excessive overpressure, a power-independent mechanism preventing too high pressure in the closed labelboxes is installed (overpressure limitation unit \boxtimes). For this, a water-filled bottle connected to the labelbox provides a robust safety tool, as by a simple physical mechanism, overpressure > 15mbar is automatically released.

Humidity control module: To regulate the maximum humidity levels, each cultivation chamber contains an internal relative humidity (polymer-based capacitive) sensor **S**, an external membrane pump \boxtimes 5a and Peltier element \boxtimes 5b as well as a condensate bottle \boxtimes 5c. A permanent gas circulation (30 L/min) is maintained throughout the whole cultivation period passing the Peltier element \boxtimes 5b by use of a membrane pump \boxtimes 5a. In such a way, not only continuous homogenous air is provided to the plants but also relative humidity levels can be limited. The increase of relative humidity above a specified value caused by plant transpiration activates an increase of Peltier current in the drying module. Peltier current forces one side of the block to cool down, which enables water condensation from the air coming from the labelbox. The

condensate is further collected in separate bottles whereas the dried air is guided back to the cultivation chambers.

Irrigation system: The irrigation system \boxtimes consisted of two external bottles with nutrient solutions (for the growing period, Table 2) connected over hoses and pipes to the plant pots in the labelboxes. Depending on the labelling regime the two boxes were supplied with different nutrient solutions $\boxed{6b,c}$. The labelbox used for global ^{15}N -labelling was supplied with salts enriched with ^{15}N $\boxed{6c}$. To prevent contamination of the air in the boxes during watering, the airspace in the nutrient solution bottles was connected with labelboxes to allow pressure compensation with air from inside the boxes.

CO₂ scrubber: At the beginning of the cultivation, before placing the seedlings into the labelbox, CO₂ was removed from air inside the labelbox. This is achieved by the use of CO₂ scrubber which is attached to the gas hose between the membrane pump $\boxed{5a}$ and labelbox \boxtimes . It is filled with CO₂ adsorbent (3-4 mm diameter, Soda Lime Carbon Dioxide adsorbent spherical granules, Spherasorb™, Intersurgical Ltd., Wokingham, UK) which traps CO₂ after navigating the dry air from the labelbox through the CO₂ scrubber. In the presented setup, switching of the valve to direct the gas flow through the CO₂ scrubber has to be done manually.

Chemicals

Methanol (MeOH, LC-MS CHROMASOLV®), acetonitrile (ACN, LC-MS CHROMASOLV®) and formic acid (FA, MS grade, ~ 98% purity) were purchased from Riedel-de Haën, Honeywell (Seelze, Germany). The ultra-pure water was obtained from an ELGA Purelab system Veolia Water (Ultra AN MK2, Vienna, Austria). The salts KOH ($\geq 99.5\%$), NH₄NO₃ ($\geq 99\%$), Na₂MoO₄*2H₂O, KH₂PO₄ ($\geq 99.8\%$), KNO₃ (65%) were obtained from Merck (Darmstadt, Germany) and MgSO₄*7H₂O, ZnSO₄*7H₂O, Ca(NO₃)₂*4H₂O, Ferric sodium - EDTA (C₁₀H₁₂N₂NaFeO₈), MnCl₂*4H₂O, ZnSO₄*7H₂O, CuSO₄*5H₂O (>98%) from Sigma-Aldrich (Steinheim, Germany). NH₄NO₃ (^{15}N , 98 atom%), Ca(NO₃)₂ (^{15}N , 98 atom%), KNO₃ (^{15}N , 98 atom%) and $^{13}\text{CO}_2$ (99 % purity) was purchased from Eurisotop (St-Aubin, France) while CO₂ and synthetic air were obtained from Messer (Gumpoldskirchen, Austria).

Plant material

Wheat genotypes Karur (*T.durum*), Remus (*T.aestivum*) and Apogee (a dwarf cultivar of *T. aestivum*) were generated and used in native and labelled (^{13}C and ^{15}N) form. The native seeds of Karur and Remus were obtained from - and grown in the glasshouse at - the Institute of Biotechnology in Plant Production (University of Natural Resources and Life Sciences, Vienna, Department of Agrobiotechnology, IFA-Tulln, Austria). The native Apogee seeds were obtained from the Department of Applied Genetics and Cell Biology (University of Natural Resources and Life Sciences, Vienna, University Research Center Tulln, Austria) and grown in the labelbox while supplied with non-labelled substrates. The ^{13}C - and ^{15}N -labelled material of each genotype was produced in the labelbox from native seeds.

Experimental description

The cultivation of *T.durum* plants lasted 87 days in total and comprised following steps: 1. preparation of the nutrient solution, wheat seedlings and labelboxes; 2. cultivation and 3. harvest

Preparation of the nutrient solution

To generate globally ^{13}C - and ^{15}N -labelled *T.durum* plants, the seedlings were grown in a perlite substrate using nutrient solutions adapted from Hoagland (1950) (45) and Bugbee, Spanarkel (1994) (46) (Table 2). The salt concentration was adapted to the growing stage of the plants resulting in two nutrient solutions (one for the germination and one for the growing period respectively, Table 2). Both labelling regimes were provided with the same nutrient solutions, with the only difference that the ^{15}N labelling experiment was carried out with highly ^{15}N -enriched salts. From two stock solutions, a total of 4 different nutrient solutions were prepared for two developmental periods, i.e. 2 with labelled ^{15}N (98-99 atom% enriched) and 2 with native ^{14}N salts.

Table 2 Adapted Hoagland solution. *Salts substituted with highly ^{15}N enriched analogues in the ^{15}N -labelling experiment

Salts	Hoagland Stock solutions [g/l]	Germination period [mg/l]	Growing period [mg/l]
KNO ₃ *	202	101	404
Ca(NO ₃) ₂ x4H ₂ O*	236	472	472
C ₁₀ H ₁₂ N ₂ NaFeO ₈ (Ferric sodium – EDTA)	15	3.75	0.975
MgSO ₄ x7H ₂ O	493	123.3	98.6
NH ₄ NO ₃ *	80	8	8
H ₃ BO ₃	2.86	0.143	0.072
MnCl ₂ x4H ₂ O	1.18	0.59	1.18
ZnSO ₄ x7H ₂ O	0.22	0.88	0.33
CuSO ₄	0.051	0.051	0.051
Na ₂ MoO ₄ x2H ₂ O	0.12	0.024	0.0072
KH ₂ PO ₄	136	68	68

All 4 nutrient solution types were prepared from the Hoagland stock solutions (composition in Table 2) in cool (< 10°C) ultra-pure autoclaved water and stirred for ca. 10 min on a magnet stirrer. Low amounts of ¹²C as well as ¹⁴N were present in the nutrient solutions to provide iron uptake in the form of ethylenediamine tetraacetic acid (EDTA) chelate complex.

Preparation of wheat seedlings

Before placement into the labelboxes, wheat seedlings were prepared according to the following three steps: germination, vernalisation and planting.

Germination: Wheat seeds were placed in blocks of rock wool (4x4 cm, Grodan) with the embryo upwards so that each block contained three seeds. The seed-contained blocks were placed in two separate darkly shaded boxes and watered with solutions for the germination period containing either ¹⁵N or ¹⁴N salts

until saturation of the rock wool. Boxes were closed and the seeds germinated at room temperature in the darkness for two days.

Vernalisation: Germinated seedlings were proceeded to vernalisation in the dark for 2 days at 4°C in the cooling room to promote shoot yield and acceleration of the flowering process (47). Under these conditions, previous experiments have shown to result in 4-6 shoots per *T.durum* plant during cultivation.

Planting of seedlings: Plant pots (~1L) were wrapped in aluminum foil as depicted in Figure 6, to prevent access of light to the medium and thus growth of algae. Pots were filled with perlite and nutrition solutions for the growth period (Table 2). For ¹⁵N-labelling nutrient solution containing ¹⁵N salts was used. Germinated and vernalised seedlings were transferred from rock wool blocks into corresponding prepared plant pots. After the placement of seedlings, a layer of rock wool was added on top of the pot, to reduce water evaporation from the nutrient solution and prevent access of light during plant cultivation.

Preparation of the PhytolabelBox equipment

Prior to placement of plants into the labelboxes, the technical performance of the equipment was verified.

First, the tightness of labelboxes was tested. The labelboxes are operated with slight overpressure which is regulated at (10 ± 2) mbar above the ambient atmospheric pressure. The ability to keep the overpressure at the setpoint range can be taken as a measure of tightness of the system including the labelboxes. To check this, the pressure cycle consisting of the pressure drop from 10 mbar to 8 mbar and build up back to 10 mbar by dosage of synthetic air was monitored, resulting in pressure cycle periods of 10 min for the ¹⁵N- and 20 min for the ¹³C-labelbox respectively. These values slightly increased with the progress of the cultivation.

Second, the increase of CO₂ in the labelbox per dosed loop was determined. This is required for the estimation of the CO₂ consumption rate of plants after the cultivation is finished. Tests showed that per dosed loop of CO₂, the concentration of CO₂ in the labelbox increased by (4.6 ± 0.3) ppm ¹²CO₂ in the ¹⁵N-labelbox and (9.7 ± 0.3) ppm in ¹³C-labelbox when the pressure reducing device on the CO₂ bottles was set to 1 bar each.

Cultivation of plants

Start of the cultivation

1. The pots with seedlings were placed in the corresponding ^{15}N - and ^{13}C -labelboxes.
2. The labelbox cover at the back was tightly sealed with screws. Note: Too tightly sealed screws may damage the Plexiglas housing. To promote tightness a silicone paste was applied on the sealing gums of the labelboxes.
3. The atmospheric CO_2 in the ^{13}C -labelbox was removed by passing the air through the CO_2 Further, $^{13}\text{CO}_2$ gas was manually dosed to reach a level of 400 ppm in the labelbox.
4. The setpoint values of CO_2 , O_2 , overpressure and humidity were defined in the software (Table 3).
5. The cultivation was initiated by starting the measurement in the software. All user predefined setpoints were automatically monitored and regulated by the system during the whole cultivation period (Table 3).

Table 3 Setpoint values for the recorded and regulated atmospheric parameters for both labelboxes.

Regulated parameters	Set point values	Allowed parameter range	
		Min.	Max.
CO_2 [ppmv]	400	300	1600
O_2 [%]	20	-	-
Overpressure [mbar]	10	8	12
Humidity [%]	variable	-	70

CO_2 , O_2 and overpressure were maintained at setpoints by the control unit while values below the minimum led to the activation of gas dosing system. The maximum allowed setpoint for relative humidity was defined while levels below depended on the plant biomass and thus was increasing with growth (see Figure 1).

Cultivation conditions

The temperature and light-duration setpoints outside the labelbox for the day/night cycle were realised over an external control unit of the climate chamber, in which the labelboxes were located throughout the cultivation period. These setpoint values are shown in Table 4. The light intensity remained constant during the whole cultivation period. The light intensity at simulated day conditions was estimated with the Li-COR sensor (LI-190SA Quantum Sensor, Li-COR, Germany) and was measured on top of the rock

wool immediately after placing the pots with seedlings into the labelbox to be 206 $\mu\text{mol}/(\text{m}^2\cdot\text{s})$ in the ^{13}C -labelbox and 220 $\mu\text{mol}/(\text{m}^2\cdot\text{s})$ in ^{15}N -labelbox respectively.

Table 4 Light and temperature intervals in the climate chamber during the cultivation process

Cultivation days	Light cycle day [h]/night [h]	Temperature day [°C]/night [°C]	Growth stage (Zadoks scale (48))
0 - 28	12/12	12/10	Z1 and Z2
28 - 31	12/12	14/12	Z2 and Z3
31 - 38	14/10	16/14	
38 - 86	14/10	18/16	Z3, Z4 and Z6

Zadoks scale: Z1- seedling development, Z2 - tillering, Z3 - stem elongation, Z4 - booting and Z6 - anthesis

Irrigation

Irrigation rhythm was adapted to the growth stage of the plants. In the first three weeks of cultivation, the plants were irrigated once a week and from day 28 until harvest twice a week. Irrigation was accomplished manually by an external peristaltic pump (S20, Vederflex® smart, UK) and a flow velocity of 100 ml/min. The volume of nutrient solution applied during watering was recorded. Both labelboxes were watered with the respective nutrient solution by the use of separate hoses.

Harvest

Each ear and the adjacent stems, leaves and roots inside the ^{13}C -labelbox were harvested separately 144 h after the first anthers had appeared on the ear, respectively. The sample material was removed via the gate compartment with an air lock system and immediately shock frozen in liquid nitrogen outside the labelboxes. The time between cutting the samples and freezing was kept as short as possible. The ^{15}N -plants were harvested under similar experimental conditions while the ^{15}N -labelbox was open, as the ambient atmosphere does not disrupt the ^{15}N -labelling process.

Freeze-drying and sample storage

Sample material was freeze-dried (FreeZone 6Plus, Labconco, Kansas City, MO, USA) to <5% rest moisture for long term storage at -80°C. To this end, wheat ears were dried for 6 days, stems for 1 day and leaves for 2 days at -80°C and ~0.4 mbar. Rest moisture was estimated with the infrared moisture analyzer (LC 4800P-00V1, Sartorius, Göttingen, Germany).

Sample preparation

Two set of samples were prepared for the measurement. One is used for the determination of the degree of enrichment and the other for the annotation of the global C and N metabolome. The procedure for milling and extraction as well as the solvent composition of samples at the time point of measurement was the same in both cases. The only difference is in the composition of the sample extracts. For annotation of the global metabolome, wheat ear extracts of freshly sampled native wheat from glasshouse were mixed with either freeze-dried $^{12}\text{C}^{15}\text{N}$ or freeze-dried $^{13}\text{C}^{14}\text{N}$ ears from the labelbox. For the determination of the degree of enrichment, extracts of either native, $^{12}\text{C}^{15}\text{N}$ - or $^{13}\text{C}^{14}\text{N}$ - wheat ears were used.

The wheat ears were milled to a fine powder with a ball mill (MM400, Retsch, Haan, Germany) while being kept in frozen condition. The extraction was performed similarly as reported in (8). In short, 100 mg of fresh native wheat powder and 30 mg of dried wheat powder was extracted separately with 1 ml MeOH/ACN/H₂O (1.5/1.5/1 v/v/v) + 0.1 % formic acid (FA). 70µl of H₂O was added to the extraction solvent of dried wheat in order to compensate for the loss of water during the drying process. For the detection of the global C and N metabolomes, native extracts were mixed 1:1 (v/v) with $^{13}\text{C}^{14}\text{N}$ or $^{12}\text{C}^{15}\text{N}$ extracts and diluted with H₂O + 0.1% FA in order to obtain 1:1 organic: water ratio prior to LC-HRMS (8). For determination of the degree of enrichment, the labelled and native extracts were diluted with H₂O + 0.1% FA individually.

LC-HRMS measurement

The LC-HRMS measurement was performed in positive ionisation mode on an Orbitrap mass spectrometer (QExactiveHF, Thermo Fisher Scientific, Bremen, Germany) coupled to a Vanquish uHPLC (Thermo Fisher Scientific, Bremen, Germany) with the method described in (49).

Data processing

Data processing was performed with the AllExtract module in MetExtract II software (14) LC-HRMS spectra of the sample set for global C and N metabolome annotation were considered. Briefly summarised, the MetExtract II software was set up to search for pairs of chromatographic peaks, which originate from co-eluting native and (^{13}C - or ^{15}N -) labelled metabolite ion forms. Native and labelled metabolite forms must show their distinctive and mirror-symmetric isotopologue patterns. Moreover, their chromatographic peak shapes must be highly similar and show perfect coelution. Such detected metabolite ions were then aligned across all samples of the either ^{13}C - or ^{15}N -labelling experiment.

The measured LC-HRMS raw files were converted into mzXML format using the MSConvertGUI (version 3.0.19166-cc86d1f56) from Proteowizard (50), and further loaded in the MetExtract II. Data processing parameters of MetExtract II were: Intensity threshold for M and M': 10000 counts; Chromatography start and end time: 3-36 minutes; Chromatographic peak scales: 7-21; Maximum allowed deviation for signal pairs: ± 3 ppm; Maximum isotopologue abundance error: $\pm 15\%$; Minimum peak correlation: 0.85 (Pearson correlation); MZ-delta between native and labelled metabolite ion form: 1.00335484 for the ^{13}C -labelling experiment and 0.99703000 for the ^{15}N -labelling experiment; Isotopologue purity: 0.9893 for the native metabolite form, 0.9850 for the ^{13}C -labelled metabolite ion form, 0.9951 for the ^{15}N -labelled metabolite ion form; Maximum MZ deviation of consecutive signal pairs: ± 8 ppm; Number of isotopologues checked: 2; Number of carbon atoms searched for in the ^{13}C -labelling experiment: 3-60; Number of nitrogen atoms searched for in the ^{15}N -labelling experiment: 1-12. Parameters for the combination of the ^{13}C - and the ^{15}N -labelling experiment: Maximum allowed MZ deviation: 5 ppm; Maximum allowed retention time shift: ± 0.15 minutes.

List Of Abbreviations

^{13}C - Stable isotope of carbon containing 6 protons and 7 neutrons

^{13}C - labelbox - cultivation chamber for ^{13}C -labelling

^{15}N – Heavy stable isotope of nitrogen which contains 7 protons and 8 neutrons

^{15}N - labelbox - cultivation chamber for ^{15}N -labelling

DW - Dry weight

FW - Fresh weight

GC-MS - Gas chromatography-mass spectrometry

Labelbox - Cultivation chamber in which stable isotopic labelling is performed

LC-HRMS - Liquid chromatography - high resolution mass spectrometry

MS - Mass spectrometry

mzXML - File format for LC-HRMS data

NMR - Nuclear magnetic resonance

T. aestivum - *Triticum aestivum*

T. durum - *Triticum durum*

Z1 - Growth stage of wheat after Zadoks scale (48) and represents seedling development

Z2 - Growth stage of wheat after Zadoks scale (48) and represents tillering

Z3 - Growth stage of wheat after Zadoks scale (48) and represents stem elongation

Z4 - Growth stage of wheat after Zadoks scale (48) and represents booting

Z5 - Growth stage of wheat after Zadoks scale (48) and represents heading

Z6 - Growth stage of wheat after Zadoks scale (48) and represents anthesis

Declarations

Ethics approval and consent to participate

Not applicable

Consent for publication

Not applicable

Availability of data and materials

The datasets used and/or analysed during the current study are available from the corresponding author on reasonable request.

Competing interests

The authors declare that there is no conflict of interest.

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Author's contributions

AC, MD, and RS contributed scientifically and technically to the development of the cultivation method and the preparation of the manuscript. AK, HB and ML scientifically contributed in adaptation of the cultivation method while AP and KX technically supported during plant cultivation. AC performed the LC-HRMS measurements and data evaluation and CB performed data processing, statistical analysis and data visualisation. All authors contributed to, have critically revised and approved the manuscript.

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Figures

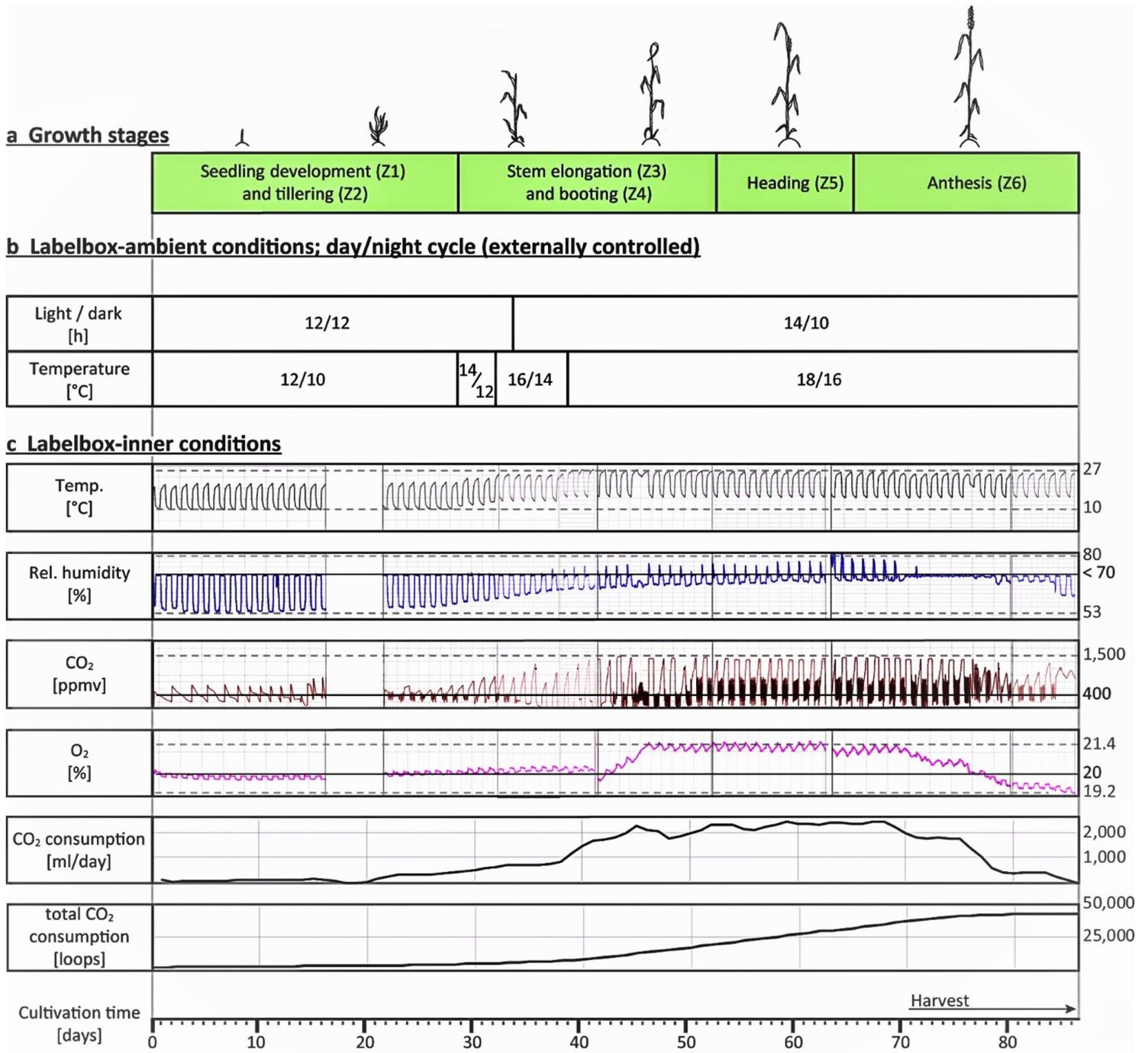
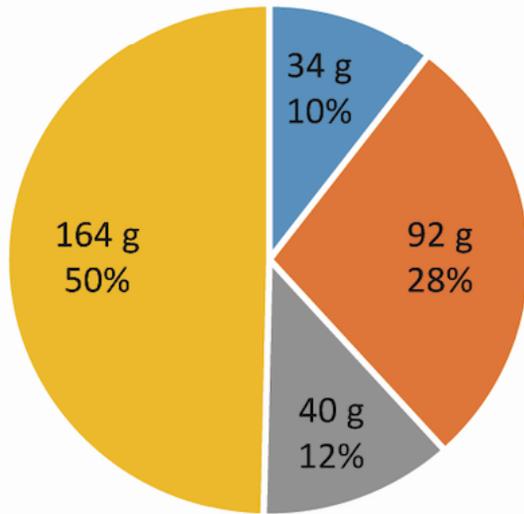


Figure 1

13C-labelbox cultivation record of wheat for the inner atmosphere correlated to the observed growth stages and applied setpoint levels for ambient temperature and light duration. Similar records were obtained for the 15N-labelbox.

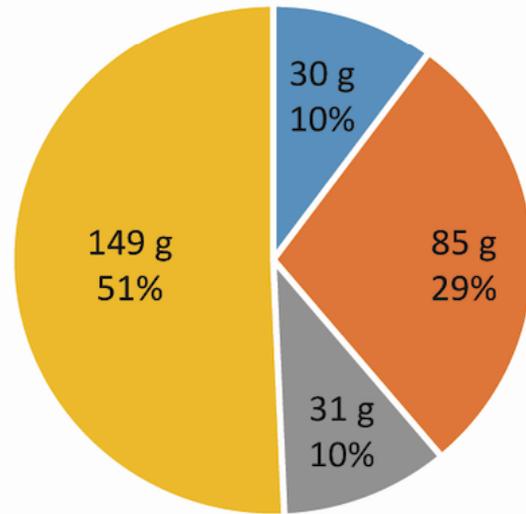
a ^{13}C -labelled *T. durum*

Total weight: 330 g



b ^{15}N -labelled *T. durum*

Total weight: 295 g



■ Ears ■ Stems ■ Leaves ■ Roots

Figure 2

Mass fractions in [%] relative to the whole wheat plant weight.

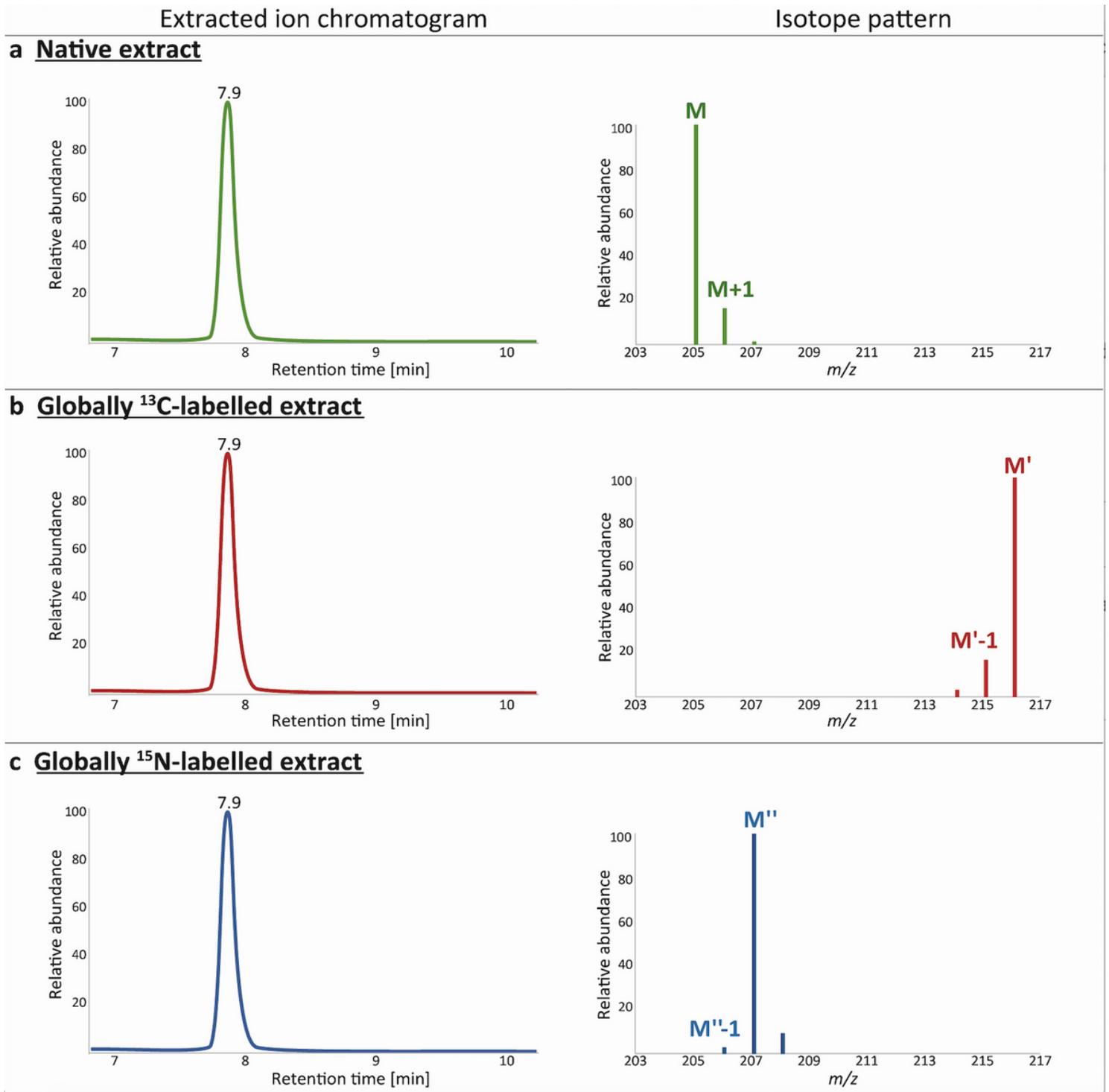
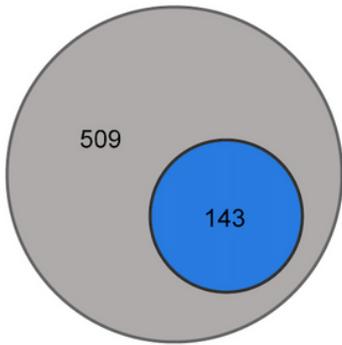


Figure 3

In each experiment high abundant isotopologues of tryptophan were represented as M (all carbons are ¹²C), M' (all carbons are ¹³C) and M'' (all nitrogens are ¹⁵N) and the low abundant isotopologues of the same molecule are M+1 (one ¹³C and rest ¹²C), M'-1 (one ¹²C and rest ¹³C) and M''-1 (one ¹⁴N and rest ¹⁵N). The relative isotopologue intensities were used to determine the degree of enrichment.

a Venn diagram

n = 652



b Feature plots

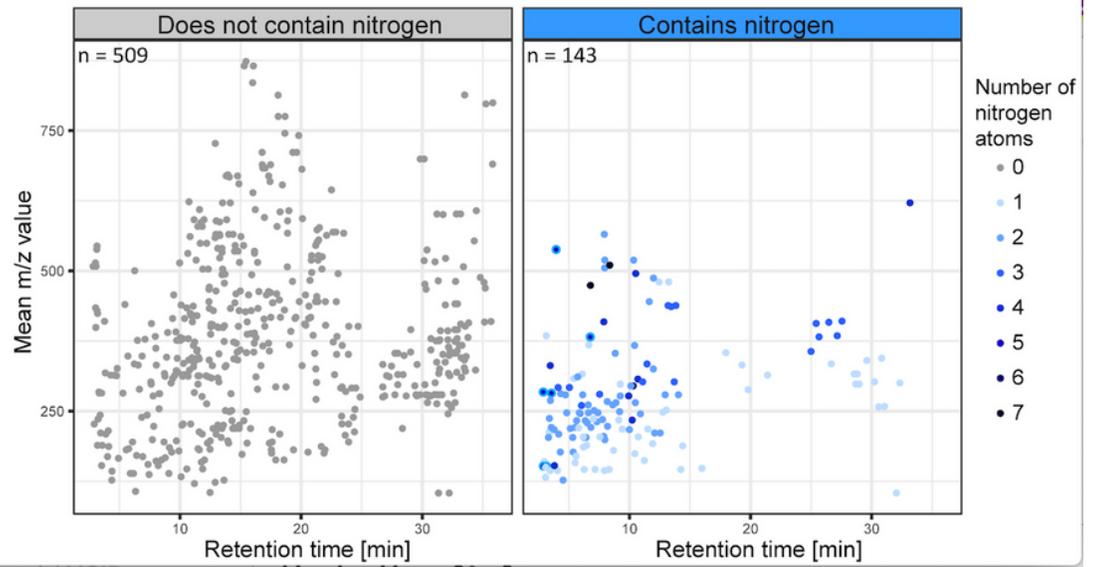


Figure 4

Each dot represents one metabolite found in the ^{13}C and ^{15}N experiment.

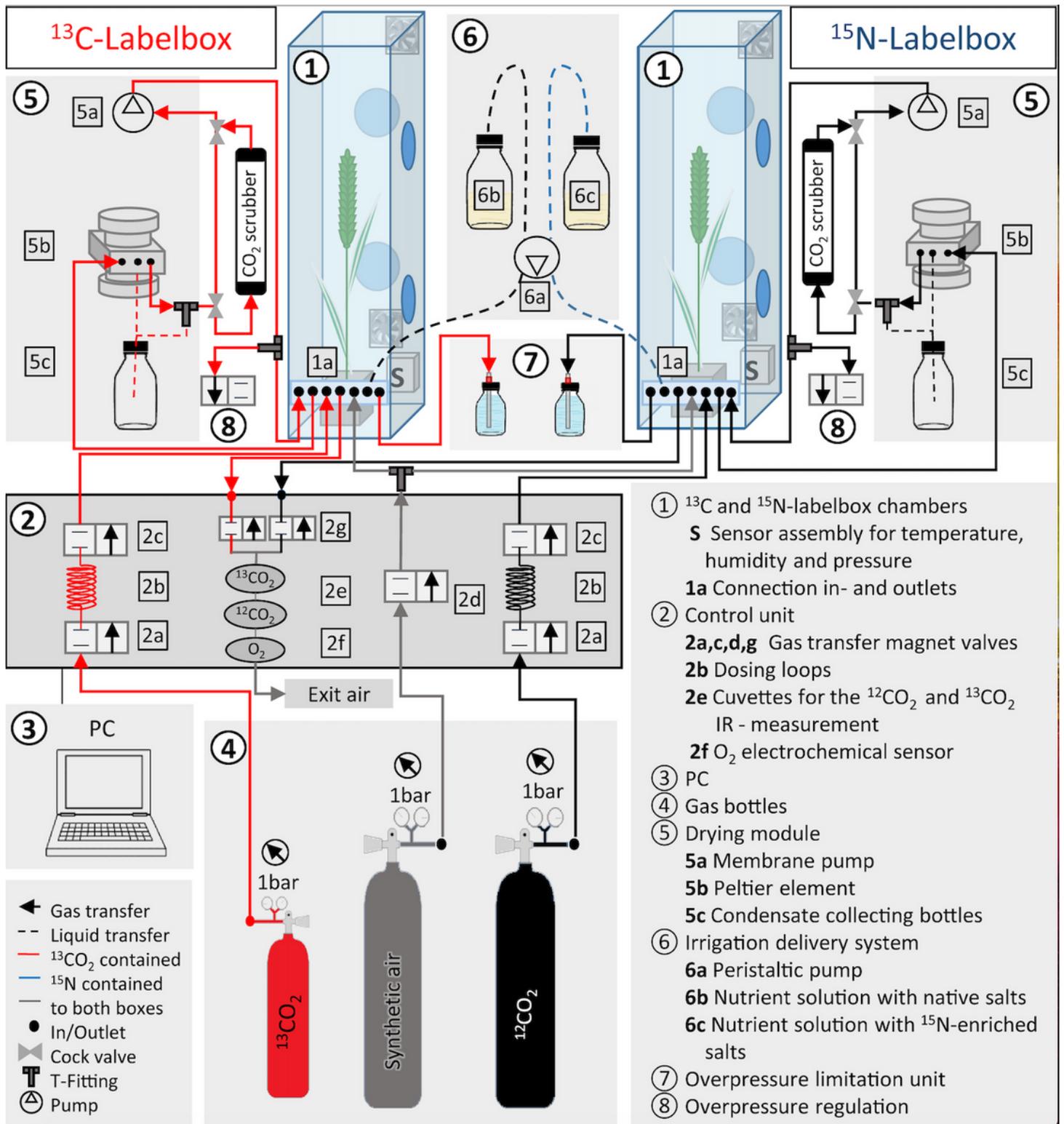


Figure 5

Simplified schematic overview of the labelling process by applying the PhytolabelBox equipment (ECH, Halle Germany).

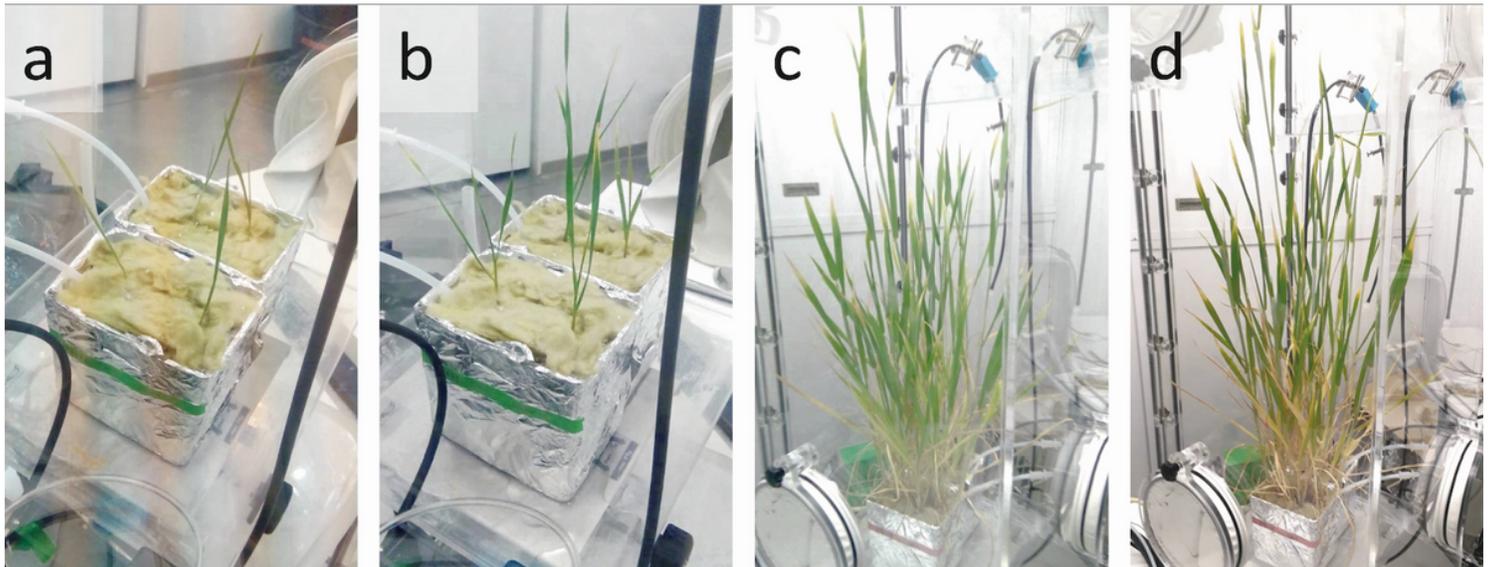


Figure 6

Wheat in the cultivation box at different growth stages: a seedling development (Z1), b tillering (Z2), c stem elongation (Z3), d heading (Z5).

Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [Supplementarymaterial.docx](#)
- [Equations.docx](#)