



Separating microbial respiration of exudates from root respiration in non-sterile soils: a comparison of four methods

Y. Kuzyakov*

Institute of Soil Science and Land Evaluation (310), University Of Hohenheim, Emil-Wolff-Strasse 27, D-70593 Stuttgart, Germany

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Abstract

Partitioning the root-derived CO₂ efflux from the soil into actual root respiration (RR) and microbial respiration of exudates and root residues is very important for determining the carbon (C) and energy balance of soils. Studies based on artificial root environments like hydroponics or sterile soils give unrealistic figures for C partitioning and are unsuitable for predicting C flows under natural conditions. To date, only four methods have been suggested to separate RR and rhizomicrobial respiration in non-sterile soils: (1) the isotope dilution method, (2) the model rhizodeposition technique, (3) modeling of ¹⁴CO₂ efflux dynamics, and (4) the exudate elution procedure. All four methods are based on the pulse labeling of shoots in a ¹⁴CO₂ atmosphere and subsequent monitoring of ¹⁴CO₂ efflux from the soil. However, the basic assumptions and principles of these methods, as well as the results observed in the original papers, all differ from one another. This study describes the separation of RR of *Lolium perenne* grown on a loamy Haplic Luvisol from microbial respiration of rhizodeposits by means of all four methods under the same experimental conditions.

In spite of alternative principles, the isotope dilution and the ¹⁴CO₂ dynamics methods show a similar level of RR: accordingly, 39 and 45% of total root-derived CO₂ efflux were accounted for by RR. The remainder is rhizomicrobial respiration. The exudate elution method, which underestimates the total rhizodeposition, shows that at least 19% of root-derived CO₂ is produced by exudate decomposition. The microbial respiration of rhizodeposits calculated using the model rhizodeposition technique is also underestimated. The exudate elution method is the only procedure allowing physical separation of both C flows. The assumptions and principles of all four methods are reviewed and the effects of possible shortcomings on the separation results are discussed. In conclusion, RR contributes about 40–50% to the root-derived CO₂ efflux. The remaining 50–60% comprise the microbial decomposition of root exudates and other rhizodeposits. The longer the period of monitoring the CO₂ efflux after the pulse labeling is, the higher the contribution of rhizomicrobial respiration to the total root-derived CO₂ efflux from soil. © 2002 Elsevier Science Ltd. All rights reserved.

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1. Introduction

Differentiating the root-derived CO₂ efflux from the soil into actual root respiration (RR) on the one hand and into microbial respiration of exudates and of root residues on the other hand is very important in quantifying the C and energy balance of the soil. Exudates and root residues are energy-rich; they enhance the underground C stock and are metabolized by soil microflora. These C sources, which are readily available to microorganisms, contribute to fast C turnover in the soil and to higher microbial activity in the rhizosphere when compared with root-free soil. Stimulation of microbial growth and activity around roots increases the mineralization of native soil organic matter (reviewed by

Cheng and Kuzyakov (2002) and Kuzyakov (2002)) and subsequently increases the availability of mineral nutrients. In contrast to the root exudates, CO₂ originating from RR cannot be used by microorganisms for their growth: it is energy-poor and does not affect the turnover of microbial biomass and soil organic matter. Therefore, accurate C and energy budgets of the soil cannot be determined without separately estimating RR and microbial utilization of root exudates. As stated by Killham and Yeomans (2001), “Discriminating between CO₂ which is directly derived from RR and that which is derived from mineralization of the components of C flow is exceptionally difficult and has presented one of the greatest challenges to quantifying rhizosphere C flow”.

The term ‘root-derived CO₂’ is used here to describe the sum of RR and CO₂ evolved by microbial decomposition of

* Tel.: +49-711-459-2327; fax: +49-711-4593-117.

exudates, secretions as well as root residues such as sloughed root cells, root hairs and dead roots. Strictly speaking, the term ‘rhizosphere CO₂’, frequently used in the literature, refers to the location of CO₂ production; from this point of view it must include not only RR and CO₂ evolved by microbial utilization of exudates, but also the CO₂ derived by microbial decomposition of rhizosphere soil organic matter.

Nutrient solution cultures (Helal and Sauerbeck, 1991; Meharg and Killham, 1991; Hodge et al., 1996; Groleau-Renaud et al., 1998), soil sterilization (Barber and Martin, 1976; Martin, 1977; Merbach et al., 1990; Merbach and Ruppel, 1992) and fumigation techniques (Helal and Sauerbeck, 1991) related to ¹⁴C or ¹³C labeling have been used to investigate microbial respiration of root exudates. The results show that investigations based on environments that are artificial for the roots, like hydroponics or sterile soils, give unrealistic figures for C partitioning (Bowen, 1980; Schönwitz and Ziegler, 1988; Merbach et al., 1990; Meharg and Killham, 1991; Schulze et al., 1994). This makes them unsuitable for predicting C flows under natural conditions. Some earlier field studies used the component integration method; this involves manually separating the constituent soil components that contribute to CO₂ efflux (i.e. roots, sieved soil, litter), followed by measurements of the specific respiration activity of each component part (Edwards and Harris, 1977; Blagodatsky et al., 1993; reviewed by Hanson et al. (2000)). This method, however, was never broadly applied, probably because of the strong impact of physical separation on the components and the significant change of rhizosphere conditions along with the imperfect separation procedure (reviewed by Hanson et al. (2000)).

More recently, efforts have been made to divide root-derived CO₂ (as a sum of RR and microbial respiration of root-derived organic C) into CO₂ originating from RR and that from microbial respiration of root-borne substances during plant growth on non-sterile soils. Four methods have been suggested: The first method, known as isotope dilution, is based on the addition of a solution of unlabeled glucose to the soil and simultaneous ¹⁴C pulse labeling of growing plants (Cheng et al., 1993). Using this method, RR of 3-week-old wheat plants was found to account for about 41% of the root-derived CO₂. The second method, based on adding ¹⁴C-labeled model rhizodeposits to the soil (Swinnen, 1994), shows that the contribution of RR of 30-days-old wheat and barley to the total root-derived CO₂ was between 89 and 95%. The third method, based on the dynamics of ¹⁴CO₂ efflux from the soil after ¹⁴C pulse labeling of shoots (Kuzyakov et al., 1999, 2001; Kuzyakov and Domanski, 2002), showed that RR and respiration of rhizosphere microorganisms in growing *Lolium perenne* amounts to 41 and 59% of root-derived CO₂ efflux from the soil, respectively. The contribution of RR varied from 17 to 61% of total CO₂ efflux from the soil, depending on the age of the *Lolium* plants. A detailed description of these three

methods, along with their shortcomings and advantages, is discussed in Kuzyakov (2001) and Kuzyakov and Domanski (2002). The fourth method is based on the elution of ¹⁴C labeled exudates from soil before microorganisms utilize them (Kuzyakov and Siniakina, 2001). Here, the CO₂ efflux from soil is monitored simultaneously to exudates elution, and water movement is driven by air circulation according to the siphon principle. The siphon elution method shows that at least 19% of root-derived CO₂ can be attributed to soluble organic substances exuded by roots.

In summary, the four methods that have been suggested to separate RR and microbial respiration of rhizodeposits in non-sterile soil show different results. Each is based on many assumptions and has certain shortcomings. It remains unclear whether the different results are method-inherent or reflects environmental and experimental conditions, i.e. different plants, soils, equipment, and environmental conditions, etc. As Hanson et al. (2000) conclude, comparative studies of different methods are sorely needed but unfortunately very rare.

The present contribution describes the separation of RR from microbial respiration of rhizodeposits by means of all four methods described earlier under the same experimental conditions. It is designed to compare methods, independently estimate both CO₂ sources, as well as select an appropriate procedure for future studies. A few improvements that help extend the original methods were used in this work.

2. Material and methods

2.1. Soil and test crops

Soil, a loamy Haplic Luvisol, was taken from the top 10 cm (Ap horizon) of the University of Hohenheim’s long-term field experimental station Karlsruh. The soil contains no CaCO₃ and has the following characteristics: pH 6.0, C_t 1.2%, N_t 0.13%, sand 4.4%, clay 23%, silt 73%. Each container was filled with 400 g soil, which was air-dried and sieved on a 2 mm screen prior to the experiment. A polycarbonate filtration device ‘CombiSart’ with a volume of 250 ml (Merck®-Laborkatalog, 2000) was used as a soil container (Fig. 1, ⓐ). The overall volume of the device, including the space under the lid, was about 340 ml. The lid of the CombiSart device contained three inlets. PVC tubes brought air through the three inlets into the CombiSart device from the output of a membrane pump. The soil was separated from the outlet in the bottom of the CombiSart device by a perforated filter support, delivered together with the filtration device, overlaid by two layers of perforated (holes = 0.5 mm) polyethylene. The outlet was connected through a PVC tube to a test tube containing 20 ml of 0.25 M NaOH for CO₂ trapping (Fig. 1, ⓑ).

One pre-germinated seed of *L. perenne* L. was put into each container and grown under 27/22 °C day/night temperature, 14 h photoperiod and 400 μmol m⁻² s⁻¹ light

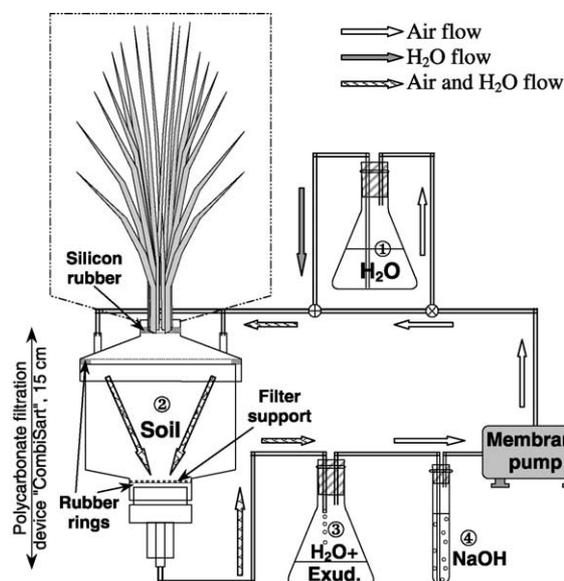


Fig. 1. Experimental setup for separate measurement of RR and exudation. ①—flask with water for elution, ②—polycarbonate filtration device CombiSart (Merck®-Laborkatalog, 2000) with soil and roots, ③—collection flask with eluted exudates, ④—test tube with NaOH solution for CO₂ trapping, ⊗—regulation clamp, ⊕—joint where air and water flows are connected. For all variants except variant 11, the outlet of the membrane pump was connected directly to the inlets of the CombiSart in the lid and its outlet was connected to the test tube with NaOH solution for CO₂ trapping. (Kuzyakov and Siniakina, 2001).

intensity. The soil water content of each chamber was adjusted daily with deionized water to 60% of the available field capacity.

The preparation of soil samples, the main soil characteristics, and the growing conditions are the same as described in detail by Domanski et al. (2001).

2.2. Methods tested

All four methods known from the literature were compared in one experiment. The experimental design of

the experiment is presented schematically in Table 1. Furthermore, in the soil of variants 8 and 10, more glucose was added than recommended in the original contribution of Cheng et al. (1993) and Swinnen (1994). This allowed conclusions to be drawn about applying the method under a wider range of conditions than described in the original papers. Additionally to the pots with planted soil without any glucose addition, the unplanted soil treated with different amounts of ¹⁴C labeled glucose was used as controls.

2.2.1. Isotope dilution method

The isotope dilution method is described in detail by Cheng et al. (1993). Briefly, unlabeled glucose was added to the soil with plants that were pulse labeled in a ¹⁴CO₂ atmosphere in order to dilute the labeled ¹⁴C rhizodeposits. The underlying assumption is that the dilution of ¹⁴C by ¹²C in the CO₂ originating from microbial respiration of rhizodeposits is proportional to the amount of unlabeled glucose added. However, only the microbial respiration of exudates is diluted, but the contribution of RR remains constant. Originally (Cheng et al., 1993), two glucose concentrations were used: 171 and 881 μg C g⁻¹ soil. Besides the recommended glucose amounts (160 and 800 μg C g⁻¹ soil), one lower (32 μg C g⁻¹) and one higher (4000 μg C g⁻¹) value were also used in our study. Two latter variants (32 and 4000 μg C g⁻¹) were tested to check a potential extended use of the method. All four glucose amounts were injected as 20 ml aqueous solution by syringe with a long needle into the soil of variants 5–8. The injection was concomitant with the ¹⁴C pulse labeling of variants 4–8 and 11 in ¹⁴CO₂ atmospheres.

2.2.2. Model rhizodeposition method

This study introduced some improvements of the original model rhizodeposition method described by Swinnen (1994). The original method involved two variants: (1) plants were ¹⁴C pulse labeled and no model rhizodeposits

Table 1
Variants and soil treatments used to separate root and rhizomicrobial respiration by means of four methods

Variant	Method name	Plant	¹⁴ CO ₂ pulse (kBq Pot ⁻¹)	¹² C glucose (μg C g ⁻¹)	¹⁴ C glucose (kBq Pot ⁻¹)	Author/reference
1	Control 0	–	0	0	0	
2	Control 1	–	0	32	119	
3	Control 2	–	0	160	119	
4	¹⁴ CO ₂ dynamics	+	118	0	0	Kuzyakov et al. (1999, 2001)
5	Isotope dilution	+	118	32	0	Cheng et al. (1993)
6	Isotope dilution	+	118	160	0	Cheng et al. (1993)
7	Isotope dilution	+	118	800	0	Cheng et al. (1993)
8	Isotope dilution	+	118	4000	0	Changed: Cheng et al. (1993)
9	Model rhizodep.	+	0	32	119	Swinnen (1994)
10	Model rhizodep.	+	0	160	119	Changed: Swinnen (1994)
11	Exudates elution	+	118	0	0	Kuzyakov and Siniakina (2001)

were added to the soil, (2) ^{14}C labeled model rhizodeposits were added to the soil with unlabeled plants.

Firstly, only ^{12}C -D-glucose was used here instead of model rhizodeposits to enable comparability with the isotope dilution method of Cheng et al. (1993). This substitution is valid because low molecular weight sugars are the main component of root exudates (Jones and Darrah, 1993; Merbach et al., 1999). The other components of rhizodeposits—carboxylic acids, amino acids, etc.—have similar microbial availability as sugars (Kuzyakov and Demin, 1998). Originally (Swinnen, 1994), ^{12}C glucose solution was used as one variant.

Secondly, the same amount of unlabeled glucose ($32\ \mu\text{g C g}^{-1}$) was added to the soil with labeled plants (variant 5) as the amount of ^{14}C labeled glucose added to the soil with unlabeled plants. This permits comparison of these variants. Otherwise, as in the case of the original method, different results might merely reflect different treatment (no glucose addition into the soil with labeled plants). Moreover, one additional glucose concentration ($160\ \mu\text{g C g}^{-1}$) was investigated. This level was used to check whether the results of the model rhizodeposition method depend on the glucose concentration. In the original paper, 1.23, 3.69 and $12.3\ \mu\text{g C glucose g}^{-1}$ soil were investigated. However, these amounts were used in different experiments, making it difficult to compare the effect of glucose amount on the separation results.

^{14}C glucose ($119\ \text{kBq pot}^{-1}$) solution was injected into the soil of variants 2, 3, 9 and 10 in the same way as in the isotope dilution method.

2.2.3. $^{14}\text{CO}_2$ dynamics method

The third method is based on the $^{14}\text{CO}_2$ efflux dynamics from soil after ^{14}C pulse labeling of shoots and subsequent modeling of the C flows in the rhizosphere (Kuzyakov et al., 1999, 2001). The C flow simulation reveals a time delay of CO_2 coming from microbial utilization of rhizodeposits versus the CO_2 coming from RR. Monitoring the $^{14}\text{CO}_2$ efflux from soil after ^{14}C pulse labeling of plants, fitting the model parameter on the measured $^{14}\text{CO}_2$ efflux, and subsequently modeling RR and microbial decomposition of exudates allow the independent estimation of both flows. This study used the second version of the model, in which all model parameters (except the two described above) were estimated in a separate experiment with ^{14}C labeling of *L. perenne* (Kuzyakov and Domanski, 2002). The $^{14}\text{CO}_2$ dynamics from the soil with labeled plants of the fourth variant (without glucose addition) were used to fit the two model parameters responsible for RR rate and exudation rate.

2.2.4. Exudate elution method

This method is based on the elution of exudates from the rhizosphere before soil microorganisms take them up and decompose them. The CO_2 evolved from RR is collected simultaneously to the exudates elution. The soil was

therefore flushed with a continuous flow of air–water mixture (Fig. 1). When this mixture left the soil, the water with eluted exudates was collected in a flask ③ containing Ag^+ solution (Micropur[®]) to suppress microbial decomposition of leaked exudates before analysis (Deubel, 1996; Gransee and Wittenmayer, 2000). The CO_2 evolved from RR was trapped in the NaOH solution (④) placed behind the flask for exudates.

The traps for $^{14}\text{CO}_2$ evolving from RR and those for eluted exudates were started at the beginning of labeling. The supply flask ① was filled up regularly with 400 ml of distilled water and the collecting flask ③ emptied. At the same time, 20 ml of 0.25 M NaOH solution in the test tube ④ was exchanged. To obtain the dynamics of RR and exudation, the exchange of solutions was done five times on the first day after the labeling and only twice on the fourth day. Fresh air was introduced into each container once daily to compensate for the O_2 consumed by soil microorganisms and roots. The method is described in detail by Kuzyakov and Siniakina (2001).

2.3. Labeling

The plants were labeled on day 64 and harvested on day 69. One day before labeling, the soil surface under the hole of the lid was sealed with a 2 mm layer of silicone rubber that overlaid a 2 mm layer of low melting point Paraffin. Three hours before the labeling the containers were flushed with CO_2 -free air to remove CO_2 evolved prior to labeling.

Two kinds of labeling were used in different variants of the experiment (Table 1): pulse labeling of *Lolium* shoots in $^{14}\text{CO}_2$ atmosphere and injection of ^{14}C glucose as aqueous solution directly into the soil.

The ^{14}C pulse labeling has been described in detail (Kuzyakov et al., 1999, 2001; Domanski et al., 2001). Briefly, after sealing with silicone, each container with soil and roots was enclosed separately within a small chamber. *Lolium* shoots were carefully placed into the chamber. ^{14}C as $\text{Na}_2^{14}\text{CO}_3$ (118 kBq) was put into a 2 ml Eppendorf micro test tube in the upper chamber and the chamber was then closed. One ml of 5 M lactic acids was added to the $\text{Na}_2^{14}\text{CO}_3$ solution in the microtest tube through a Teflon pipe. This allowed complete evolution of $^{14}\text{CO}_2$ into the chamber atmosphere. Assimilation took place within 1 h after the pulsing of $^{14}\text{CO}_2$. After the 1 h labeling period, trapping of CO_2 from the upper compartment was started to remove the remaining unassimilated $^{14}\text{CO}_2$. The air of the upper chamber was pumped through 20 ml of 0.5 M NaOH solution to remove $^{14}\text{CO}_2$. Then the top of the chamber was removed. The plants of the variants 4–8 and 11 were labeled in this manner in the $^{14}\text{CO}_2$ atmosphere.

The injection labeling variant involved adding aqueous ^{14}C glucose solution. The labeled D-[U- ^{14}C]-glucose (119 kBq) was added to 20 ml aqueous solution of unlabeled D-glucose containing a total of 32 (variants 2 and 9) or 160 (variants 3 and 10) mg glucose and was

mixed. This was then injected into different parts of the soil through three inlets in the lid of the polycarbonate filtration device (see Section 2.1) as 20 ml aqueous solution using a syringe with a long needle. Soil of variants 2, 3, 9 and 10 were labeled with ^{14}C glucose in this manner. The unlabeled glucose was added in the same way to the soil of variants 5–8. The injection and pulse labeling took place at the same time.

2.4. Sample analysis

During the experiment, the CO_2 evolved from the soil-root compartment was trapped in 20 ml of 0.5 M NaOH solution by continuous pumping ($100\text{ cm}^3\text{ min}^{-1}$) with a membrane pump (Fig. 1). Fig. 1 shows the detailed layout of the CO_2 trapping and exudates collection system for the variants of the exudates elution method (variant 11). The chambers of all other variants were connected in the same way, yet without the flasks ① and ③. Thus, the outlet of the soil–root chamber ② was connected to the NaOH test tube ④ and inlet of membrane pump. The outlet of the membrane pump was connected to the three inlets of the lid of the soil–root chamber ②.

The trap for $^{14}\text{CO}_2$ evolved from the soil was changed every 2 h immediately after labeling, but after 2 days only twice daily. Five days after labeling, the soil–root chamber was opened, each plant cut at the base, and each root–soil column pulled out. Shoots and roots were dried at 60°C . Dry samples of shoots, roots, and soil were mixed and pulverized in a ball mill (Retsch Co.) prior to analyses for radioactivity and for total C and N determination.

$^{14}\text{C-CO}_2$ collected in a NaOH solution was measured with the scintillation cocktail Rothiscint-22x (Roth Company) on 1 ml aliquots of NaOH after the decay of chemiluminescence. The ^{14}C counting efficiency was about $89 \pm 1\%$ and the ^{14}C -activity measurement error did not exceed 2%. The absolute ^{14}C -activity was standardized by addition of NaOH solution as quencher to the scintillation cocktail and using a two-channel ratio method of extended standard (tSIE).

Radioactivity of shoots, roots and soil samples was measured after combustion of 1 g of sample within an oxidizer unit (Canberra Packard Co. Ltd, Model 307) with the scintillation cocktail Permafluor E⁺ (Canberra Packard Co. Ltd) by a Liquid Scintillation Counter Tri-Carb 2000CA (Canberra Packard Co. Ltd).

The total CO_2 content collected in NaOH solution was measured by titration with 0.2 M HCl against phenolphthalein after addition of 2.0 M BaCl_2 solution (Black, 1965). Total C and N content in shoot and root was measured by a C–N–Analyzer (Carlo-Erba). The total C content in shoots was about $41 \pm 2\%$ of dry mass and was accepted as a constant. Total C content in washed roots varied considerably because of different contents of mineral soil particles.

The experiment was conducted with four replicates for the variants listed in Table 1. In a first step, the $^{14}\text{C-CO}_2$ data

were calculated as percentages of total assimilated ^{14}C for each replication. The total assimilated ^{14}C was calculated according to the equation

$$^{14}\text{C}_{\text{ass}} = ^{14}\text{C}_{\text{input}} - ^{14}\text{C}_n - ^{14}\text{C}_r,$$

$^{14}\text{C}_{\text{ass}}$ is activity of total assimilated ^{14}C ; $^{14}\text{C}_{\text{input}}$ is total activity introduced as $\text{Na}_2^{14}\text{CO}_2$ (118 kBq); $^{14}\text{C}_n$ is activity of the NaOH solution after flushing the upper chamber; $^{14}\text{C}_r$ is the ^{14}C not volatilized after addition of lactic acid. In a second step the total $^{14}\text{C-CO}_2$ (root-derived CO_2) evolved from the soil was equated to 100% and the contributions of RR and rhizomicrobial contribution to the total root-derived CO_2 were calculated. Standard deviation (SD) was calculated as a variability parameter.

3. Results

3.1. Isotope dilution method

In the variant without glucose addition, the dynamics of $^{14}\text{CO}_2$ efflux intensity were similar to the results observed in previous studies in our laboratory. Increasing glucose additions to the soil decreased specific ^{14}C activity of the evolved $^{14}\text{CO}_2$ (Fig. 2). Even 16 and $80\text{ }\mu\text{g g}^{-1}$ soil reduced the activity of evolved $^{14}\text{CO}_2$ by about 20%. Further addition of unlabeled glucose diminished the evolved $^{14}\text{CO}_2$ activity by up to 39% of the variant without glucose addition. To calculate the function of evolved $^{14}\text{CO}_2$ from the amount of added glucose ($^{14}\text{CO}_2\%(\text{Glucose})$), the parameter (k) of exponential equation with a constant RR were fitted on the measured data:

$$^{14}\text{CO}_2\%(\text{Glucose}) = (100 - \text{RR})\exp(-k\text{Glucose}) + \text{RR} \quad (1)$$

where RR is root respiration, Glucose is the concentration of added unlabeled glucose, k is a proportionality coefficient of decreasing specific activity.

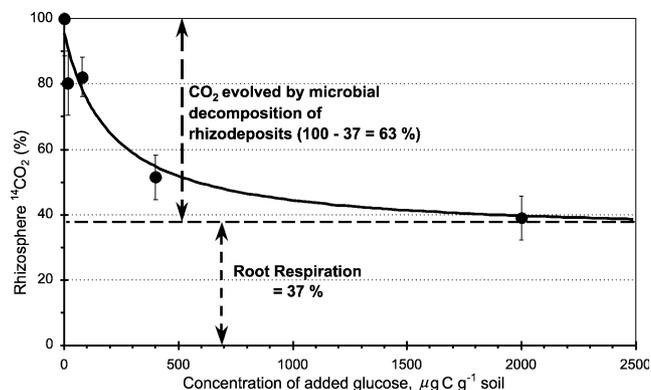


Fig. 2. Separation of RR and rhizomicrobial respiration using the isotope dilution method (Cheng et al., 1993): Dilution of $^{14}\text{CO}_2$ efflux from soil with $^{12}\text{CO}_2$ coming from different amounts of added unlabelled glucose. Vertical bars show standard deviation.

The fitted RR coefficient shows that RR amounts to about 37% of total rhizosphere respiration (Fig. 2). The difference to 100% yields the respiration of microorganisms decomposing rhizodeposits in the soil of the variant without glucose addition. These results are very close to those presented in Cheng et al. (1993) original contribution: 41% for RR and 59% for rhizomicrobial respiration. The slightly lower RR here probably reflects the much higher glucose amount ($4000 \mu\text{g C g}^{-1}$) added in the soil of variant 8.

3.2. Model rhizodeposition method

The calculation of RR and rhizomicrobial respiration by the model rhizodeposition method assumes a constant ratio between microbially respired ^{14}C ($^{14}\text{C-MR}$) and ^{14}C remaining in the soil ($^{14}\text{C-Soil}$) in the variant with natural rhizodeposits and with model rhizodeposits (Swinnen, 1994)

$$^{14}\text{C-MR}_C / ^{14}\text{C-Soil}_C = ^{14}\text{C-MR}_{\text{Glu}} / ^{14}\text{C-Soil}_{\text{Glu}} \quad (2)$$

where $^{14}\text{C-MR}_C$ and $^{14}\text{C-MR}_{\text{Glu}}$ are ^{14}C activity evolved by microbial respiration from the soil of the control variant with labeled plants and from the soil with added ^{14}C glucose, and $^{14}\text{C-Soil}_C$ and $^{14}\text{C-Soil}_{\text{Glu}}$ are ^{14}C activity remaining in soil residue in the variant with labeled plants and in the variant with added ^{14}C glucose.

The two glucose levels tested here (16 and $80 \mu\text{g g}^{-1}$ soil) differ from those of Swinnen (1994) (1.23 and $3.67 \mu\text{g g}^{-1}$ soil). The latter concentrations were close together and investigated in different experiments, making it difficult to estimate the effect of the amount of added glucose. The significantly different concentrations chosen here show that RR depends on the glucose concentration (Table 1): at the lower value, RR was $83 \pm 1\%$ of total root-derived CO_2 , whereas at the higher value it was $71 \pm 6\%$. Thus, adding an additional $64 \mu\text{g glucose g}^{-1}$ soil decreased the contribution of microbial respiration by 12%. According to this method, the microbial respiration of the exudates is only about 17–29% of total root-derived CO_2 . However, these results are 2–3 times higher than in Swinnen (1994), who reported values of only 5–11%. I explain this result by the higher glucose concentration used in my experiment.

3.3. $^{14}\text{CO}_2$ dynamics method

The separation of root and microbial respiration by this method is based on the assumption that these two processes are predominant in total root-derived $^{14}\text{CO}_2$ efflux at different times following ^{14}C pulse labeling of shoots. The most rapid process is the CO_2 efflux from RR, which predominates in the first day after labeling (Fig. 3, phase 1). The CO_2 evolution by microbial respiration of root exudates occurs later than RR because it consists of a chain of

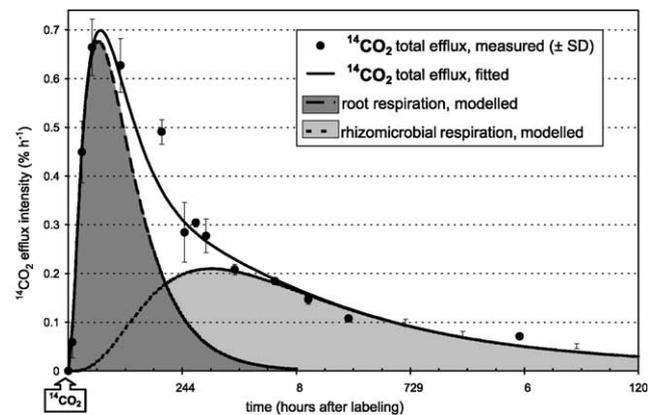


Fig. 3. Separation of RR and rhizomicrobial respiration using the $^{14}\text{CO}_2$ efflux dynamics method (Kuzyakov et al., 1999, 2001; Kuzyakov and Domanski, 2002): $^{14}\text{CO}_2$ efflux evolved by RR appears earlier than $^{14}\text{CO}_2$ respired by rhizomicrobial respiration. Vertical bars show standard deviation.

successive processes: exudation of organic substances from the root, their intake by microorganisms and their use for microbial respiration. It predominates between the second and fifth day after labeling (Fig. 3, phase 2). The microbial respiration of dead roots is very slow (on the time scale used) and therefore contributes only negligibly to the total $^{14}\text{CO}_2$ efflux in these first days. It predominates after day 5 (Kuzyakov et al., 2001; Domanski et al., 2001).

Using the model of below-ground C translocation described earlier (Kuzyakov and Domanski, 2002) allows separate simulation of root and rhizomicrobial respiration (Fig. 3). The fitted curve of the total root-derived CO_2 and the subdivision in (1) RR and (2) microbial respiration of root exudates and dead roots substantiate these assumptions. In order to calculate the total amount of C passed through RR and root exudation separately each process was integrated. The former contributed 45%, the latter about 55% of total root-derived CO_2 (measured as $^{14}\text{CO}_2$). These relationships are close to previously reported ones (Kuzyakov et al., 1999, 2001). Note here that prolonging CO_2 monitoring period for longer than 5 days will increase the contribution of rhizomicrobial respiration and diminish the contribution of RR.

3.4. Exudates elution method

This method of separating the two respiration processes is based on the blowout of $^{14}\text{CO}_2$ coming from RR by continuous air pumping and simultaneous leakage of original root exudates as well as of organic compounds modified by microorganisms by continuous water flow through the rhizosphere soil.

^{14}C activity was found in the $^{14}\text{CO}_2$ coming from the rhizosphere and in the water with eluted organic compounds. Note that the eluted ^{14}C -labeled compounds stem from original root exudates as well as from the organic

compounds metabolized and modified by microorganisms during elution.

The total ^{14}C activity found in $^{14}\text{CO}_2$ coming from the soil corresponded to 8.5% of total assimilated ^{14}C , and the corresponding value in eluted organics was 2.3%. Thus, rhizomicrobial respiration contributed at least 19% to the total root-derived CO_2 . This yields a ratio ($^{14}\text{CO}_2$ respired/ ^{14}C eluted) of about 4.5 over the whole observation period. However, this ratio strongly depends on sampling time, especially during the first 2 days after start of assimilation (labeling). The first samples, taken 1 h 40 min after the start of labeling, revealed nine times more ^{14}C in $^{14}\text{CO}_2$ compared to ^{14}C in eluted root-derived organic compounds. The maximum $^{14}\text{CO}_2$ efflux occurred roughly 12 h after labeling, but there were two ^{14}C maxima in eluted root-derived organic compounds: the first after 5 h and the second between 20 and 24 h. Note that both these maxima correspond with the light phases (Fig. 4), i.e. they occurred during the photosynthesis period. On day 3 (ca. 40 h) after labeling, a smaller third peak appeared also during the light phase. In contrast, both minima measured during the first 2 days occurred at night.

The ratio of $^{14}\text{CO}_2$ to ^{14}C in eluted root-derived organic compounds was about 6–9 at night and dropped to about 2–3 during the assimilation time (Fig. 5). Exudation intensity is thus much higher during the daytime. Two days after assimilation the ratio showed no diurnal dynamics and remained on the level of 5–6.

Exudation intensity decreased after the second maximum and, two days after labeling, was below $0.01\% \text{ }^{14}\text{C h}^{-1}$. This suggests that 2 days after assimilation, most of the ^{14}C activity was located in the non-soluble organic compounds, such as root hairs, sloughed cells and partially in mucigels, microbial biomass, etc. Therefore, the $^{14}\text{CO}_2$ coming from the rhizosphere on day 3 mainly originates from microbial decomposition of root hairs, sloughed cells and turnover of

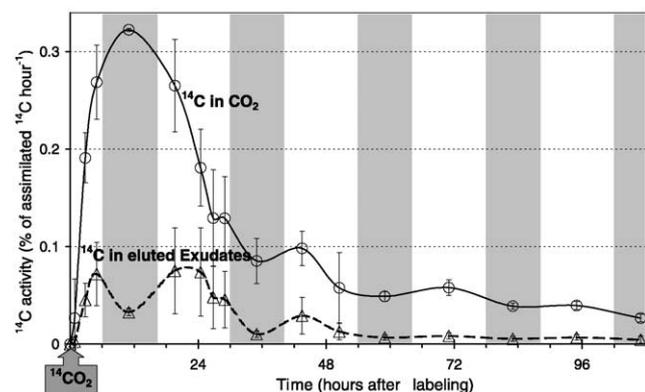


Fig. 4. Dynamics of ^{14}C in RR and exudates separated with the exudates elution method after ^{14}C pulse labeling of shoots: Elution of root exudates occurs simultaneously to blow out of $^{14}\text{CO}_2$ coming from RR. Vertical bars show standard deviation. Grey area shows the nighttime (Kuzyakov and Siniakina, 2001).

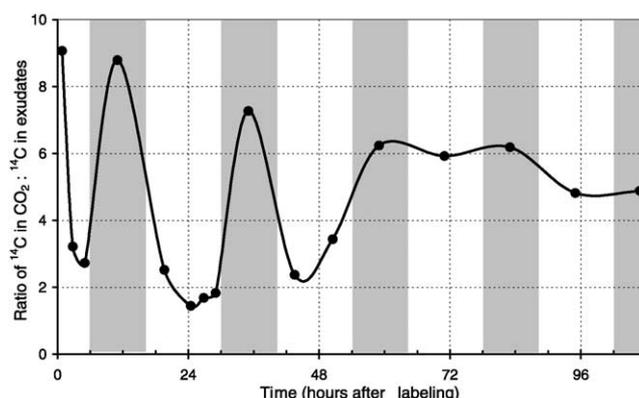


Fig. 5. Quotient between ^{14}C in RR and in exudates during four and a half days after assimilation. Grey area shows the nighttime. (Kuzyakov and Siniakina, 2001).

microorganisms. Strictly speaking, it is no longer connected with the mineralization of actual root exudates (water soluble low molecular organic substances).

4. Discussion

Four different methods allowing separate estimation of RR and rhizomicrobial respiration in non-sterile soil were tested in this experiment under the same environmental and experimental conditions. Thus, the observed differences between investigated methods can be only attributed to the methods themselves and to their assumptions. This makes it important to enumerate the assumptions considered by each method, to expose possible shortcomings, and to evaluate their possible effects on the separation results.

4.1. Isotope dilution method

Some assumptions of this method were discussed by Cheng et al. (1993, 1994, 1996): (1) injection of glucose does not produce short-term effects on plant physiology other than diluting the root exudates; (2) glucose is compatible with root exudates in terms of substrate specificity; (3) adding glucose does not stimulate or suppress the microbial activities in the rhizosphere during the experiment; (4) the dilution of $^{14}\text{CO}_2$ evolved from the soil shows a simple and proportional relationship with the amounts of added glucose. The last assumption enables calculating the ratio of RR to rhizomicrobial respiration. In my opinion, all these assumptions are acceptable because they have no effect on the separation results. However, one very important hidden assumption not discussed in the original paper has also to be considered: the ratio of root to rhizomicrobial respiration is accepted as fixed during the experiment and is extrapolated for the whole period of rhizosphere respiration. Based on the $^{14}\text{CO}_2$ efflux, Warembourg and Billes (1979), Nguyen et al. (1999) and Kuzyakov et al. (1999, 2001) have indirectly demonstrated that this ratio changes during plant development after a ^{14}C

pulse. An additional shortcoming of this method is that the above ratio can be registered only briefly (about 4–5 h) after supplying the soil with glucose. After this lag-period the microorganisms begin to grow and the third and fourth assumptions can no longer be accepted.

4.2. Model rhizodeposition method

This method assumes that the $^{14}\text{CO}_2$ efflux from soil with added labeled model rhizodeposits corresponds to rhizomicrobial respiration in the soil with applied above-ground labeling of plants (Swinnen, 1994). The $^{14}\text{CO}_2$ efflux from the soil with added model rhizodeposits is monitored and compared with the $^{14}\text{CO}_2$ efflux from the soil with pulse labeled plants (Swinnen, 1994). The ratio of respired and remaining ^{14}C in both variants is used to estimate the RR (Eq. (2)). Accordingly, the contribution of RR of 30-day-old wheat and barley to total root-derived CO_2 was between 89 and 95%. These results appear too high considering the high concentration of exudates in the rhizosphere (Cheng et al., 1993) and oppose many other investigations about partitioning of the $^{14}\text{CO}_2$ efflux (Helal and Sauerbeck, 1989; Kuzyakov et al., 1999; reviewed by Kuzyakov and Domanski, 2000). The possible source of the artifacts may be the missing addition of unlabeled model rhizodeposits to the soil with labeled plants. This possible shortcoming was improved in the reported experiment.

The hidden assumption of the method is that the microbial utilization of glucose and its absorption by clay minerals and soil organic matter is the same in the rhizosphere and in the bulk soil. When glucose (in this experiment) or model rhizodeposits (in the original paper) are artificially injected into the soil, only part of the added amount is located in the rhizosphere. The main part is distributed in the root-free soil. The microbial decomposition of glucose and other rhizodeposits to CO_2 in the rhizosphere is higher than in the bulk soil because rhizosphere microorganisms are more strongly limited by N and other nutrients. Moreover, different microbial communities of rhizosphere and root-free soil may be responsible for various yield factors (Y) (Payne, 1970) and conversion to CO_2 . Finally, many rhizosphere microorganisms are located directly on the rhizoplane at the exudation sites. The exuded organic substances can be directly taken up by microorganisms and have no chance to be absorbed by clay minerals and soil organic matter. After the experiment, more ^{14}C therefore remains in the bulk soil with added ^{14}C glucose compared with the ^{14}C remaining in the rhizosphere. This result decreases the contribution of rhizomicrobial respiration according to Eq. (2) and increases the contribution of RR to the total root-derived CO_2 .

4.3. $^{14}\text{CO}_2$ dynamics method

This method is based on the dynamics of $^{14}\text{CO}_2$ efflux from soil after pulse labeling (Kuzyakov et al., 1999; 2001).

It assumes that $^{14}\text{CO}_2$ generated by RR appears earlier than that derived from rhizomicrobial respiration. This delay reflects the time necessary for the synthesis of exudates, for the exudation and secretion processes, and for the uptake and utilization of rhizodeposits by microorganisms (Warembourg and Billes, 1979). However, this time delay is not introduced in the model artificially (no time lags in the model) and the parameters responsible for RR and root exudation are of the same order. Therefore, this delay occurs because of the successive processes.

The following other assumptions are used in the C flow model by the dynamics separation method (Kuzyakov and Domanski, 2002): (1) The plant biomass does not significantly change during the whole period of $^{14}\text{CO}_2$ monitoring until the end of C allocation. (2) The influence of plant growth (reverse transport of ^{14}C -labeled compounds from the roots to the shoots) on partitioning processes was omitted from the model. (3) The model does not consider the diurnal changes in assimilation, translocation and respiration activity. (4) All ^{14}C flows in the model are described by first-order kinetics. All these assumptions are used in developing the model and have no short-time effects (several days after ^{14}C pulse labeling) on the separation results.

4.4. Exudates elution method

The last method is based on the elution of exudates from soil before microorganisms can utilize them. It also includes many assumptions and shortcomings (Kuzyakov and Siniakina, 2001). The first shortcoming involves the limited elution of some mucigels secreted by roots as well as the ^{14}C incorporated in root hairs and sloughed root cells. Merbach et al. (1999), however, showed that up to between 60 and 80% of the root-borne organic compounds was mainly water-soluble. Similarly, Jones and Darrah (1993) found that, depending on the removal of nutrient solution, soluble low molecular weight exudates account for between 48 and 86% of root-derived organic compounds.

The exudates in this method are mainly eluted by preferential flow. Therefore, during exudation, the processes near the preferential flow pathway might differ from those farther away. The mean time for exudate elution by preferential flow amounts to about 5–10 min. The elution time of organic substances exuded far from the main water streams is longer, but it is difficult to estimate it. Thus, microorganisms can decompose the exudates during their transport from the root to the exudate collector. The eluted organics therefore consist not only of the original exudates but also include substances modified by microorganisms during elution. This shortcoming may depend on the soil texture. Sandy soils are probably more suitable for the separation with this method than the clay soils.

Continuous water flow in the microcosm may change the amount and composition of the C released by the roots. Jones and Darrah (1993) reported an up to 98% re-uptake of

maize exudates in a sterile static nutrient solution culture. Using ^{14}C labeled glucose, Paterson and Sim (1999) show a 75% re-uptake of exudates by *L. perenne* roots in a sterile nutrient solution culture. However, it is doubtful whether such re-uptake plays a significant role under non-sterilized soil conditions. Under field conditions, microorganisms on the root surface strongly compete with roots for exudates. In our system the removal of exudates from roots by water flow may be accepted as uptake by microorganisms.

These shortcomings of the exudates elution method may increase the ^{14}C in CO_2 , thereby decreasing it in exudates. Therefore, the ^{14}C measured in eluted organic compounds is probably underestimated and the ^{14}C in CO_2 overestimated. Hence it can be concluded that the method shows only the minimal amount of water-soluble exudates released from roots.

Despite its shortcomings, one key advantage over the other tested methods deserves mention: the exudates elution method is only one technique allowing physical separation of different CO_2 sources. The three other methods are based on calculations and not on physical separation. Therefore their results cannot be verified directly and remain unverifiable hypotheses. The physical separation in the exudates elution method shows that the ratio of CO_2 derived from RR and C respired by microbial utilization of rhizodeposits depends on the diurnal dynamics of photosynthesis (Figs. 4 and 5): At the light-on events the exudation intensity is 2–3 times higher than at night, while RR remains nearly constant, independent of photosynthesis. Thus, the clear diurnal dynamics of total and root-derived CO_2 efflux by growing wheat on a C_4 soil observed by Kuzyakov and Cheng (2001) could be explained by diurnal dynamics of exudation and not by RR.

The advantage of physical separation of root exudates by this method could be used later to study the composition of organic substances exuded by roots in non-sterile soils. Until now such investigations have been conducted only for nutrient solution studies or sterilized soils, in which the root growth conditions are unnatural (Bowen, 1980; Schönwitz and Ziegler, 1988; Merbach et al., 1990; Meharg and Killham, 1991; Schulze et al., 1994).

4.5. General comparison of the four methods and conclusions

In spite of the different approaches and assumptions used by each method, there is some important conformity: All four methods are based on the pulse labeling of shoots in $^{14}\text{CO}_2$ atmosphere and subsequent monitoring of $^{14}\text{CO}_2$ efflux from the soil. Labeling is a mandatory requirement to separate root-derived CO_2 from SOM-derived CO_2 . Continuous labeling is hardly useful for such separations, especially for isotope dilution and $^{14}\text{CO}_2$ dynamics methods. Theoretically ^{13}C could be used instead of ^{14}C . However, the very high sensitivity of ^{14}C isotope

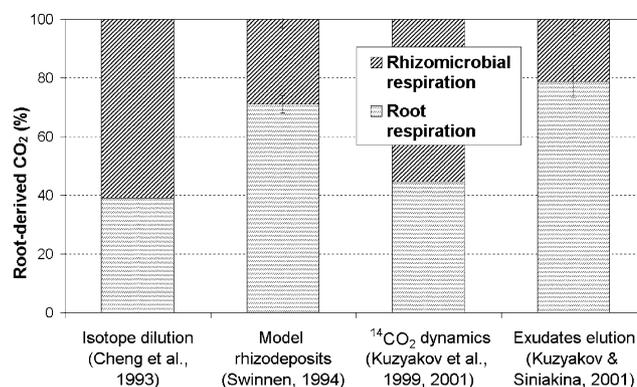


Fig. 6. General comparison of the results obtained by four methods separating RR and rhizomicrobial respiration in non-sterile soils. Mean values \pm SD are shown. SD for isotope dilution and $^{14}\text{CO}_2$ dynamics methods cannot be calculated.

analyses, its easy application and cheap analyses make ^{14}C an unmatched tracer for such studies.

Fig. 6 shows the generalized results observed by each method tested. Two pairs of results were similar: The isotope dilution and the dynamics method show similar amounts of RR, 39 and 45% of total root-derived CO_2 efflux, respectively. The RR estimated using the model rhizodeposition technique and the exudates elution method was 71 and 79% of total root-derived CO_2 efflux, respectively.

Interestingly, similar results were obtained by the isotope dilution and the $^{14}\text{CO}_2$ dynamics method despite alternative underlying assumptions. This mutually exclusive assumption involves the ratio of ^{14}C in CO_2 coming from RR to that derived by microbial respiration of rhizodeposits after the pulse labeling. The former method assumes that this ratio is constant during the observation, whereas the latter method accepts this ratio as variable. The results of the exudates elution method, however, show that this ratio is not constant, that it changes after labeling, and that it depends on photosynthesis cycles (Fig. 4).

As described earlier, the exudates elution method shows only the minimal amount of water-soluble exudates released from roots. Nonetheless, this minimal amount is about two to three times higher than that estimated by the original model rhizodeposition method (Swinnen, 1994) and is similar to the results of the improved model rhizodeposition method used in our experiment. The model rhizodeposition method therefore also underestimates rhizomicrobial respiration and strongly overestimates RR. Thus, the addition of ^{14}C -labeled model rhizodeposits to soil (Swinnen, 1994) is unacceptable as a satisfactory method for separating RR from microbial respiration of exudates.

The amount of eluted root-derived organic substances measured with the exudates elution method is about two times smaller than that measured with the isotope dilution (Cheng et al., 1993) or $^{14}\text{CO}_2$ dynamics method (Kuzyakov et al., 1999, 2001). A similar factor of about two emerged

when comparing the dipping method (almost 100% of cold-water-soluble exudates) with the percolation method (comparable with the presented Siphon method) in extracting exudates from the rhizosphere (Gransee and Wittenmayer, 2000). Thus, we can accept that roughly half of the substances exuded by roots are water-insoluble and that microorganisms will mineralize a part (no more than half) of the exuded organic compounds during the leaching process through the loamy soil. Merbach et al. (1999) also found that the amount of water-insoluble secretions comprises less than 40% of root-borne organic compounds. We can therefore conclude that RR contributes about 40–50% to the root-derived CO₂ efflux. The remaining 50–60% comprise the microbial decomposition of root exudates and other rhizodeposits. We can also conclude that the longer the monitoring period of CO₂ efflux after pulse labeling, the higher the contribution of rhizomicrobial respiration to the total root-derived CO₂ efflux from soil.

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