Topical Editor Decision: Revision by Peter de Ruiter

Comments to the Author:

Dear Dr. Comeau:

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I have seen the comments provided by the reviewers on your manuscript, and your response to these comments. I understand that you are willing to address all comments in a new revised version of the manuscript. I kindly ask you to do so in order to accept the article for SOIL.

Kind regards, Peter de Ruiter

We would like to say thank you to the referees for the comments to improve our manuscript. We appreciate the valuable suggestions from this review process. The co-authors and I have revised the manuscript and carefully addressed all the suggestions. We believe the clarity and quality of the manuscript has been enhanced with the changes.

Please find below our detailed response to all comments in bold font; the comments are in italics. Included in the revised submission is a document (below) with changes visible in Track Changes.

Anonymous Referee #1

15 Received and published: 1 February 2018

It is not an easy task to separate autotrophic and heterotrophic soil respiration. Five methods were compared and none was superior. Accuracy was defined by the difference with the ÉU° 13C-CO2 natural abundance. This is a risky approach because this method has a low precision. Fortunately an second low precision method (root regression) gave a similar estimate of the ratio between heterotrophic and total soil respiration. Strengths and weaknesses of each method are discussed and it is concluded that a combination of methods is needed. This is a thorough investigation on an important subject. The manuscript is well written and easy to read. The methods section is relatively long, but this is inherent to a methodical paper.

We appreciate the positive comments from the reviewer. To address the concern related to "accuracy", the text and table 6 were modified to refer to "relative accuracy" because accuracy was measured in relation to the δ^{13} C-CO₂ natural abundance method.

Non accessible PhD and BSc theses, such as Farmer 2013 and Tong 2015, can be removed from the literature references.

Regarding citations from Farmer (2013) and Tong (2015), the latter was removed from the literature references as suggested and the former was modified in the reference section to allow reader to retrieve it. "Farmer, J.A.: Measuring and modeling soil carbon and carbon dioxide emissions from Indonesian peatlands under land-use change, Ph D, University of Aberdeen, UK, 2013. <u>Available from</u> University of Aberdeen library collections,

https://aulib.abdn.ac.uk/F?func=direct&local_base=ABN01&doc_number=001582866" see lines 395-397.

I found some typos:

At line number

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13 still many uncertainties and unknown. Should be unknowns.

Addressed: "... still many uncertainties and unknowns ..." see line 13.

40 53 Review have been made. Should be reviews.

Addressed: "Reviews of Rh-Rs segregation ..." see line 53.

120 to quantify and CO2. Something missing?

Addressed: "The GC system was equipped with a flame ionization detector and an electron capture detector to quantify CO_2 concentration." see lines 118-119.

45 172 AND 173 time inversion. What is an inversion of time?

Addressed: "The time needed to set up the experiment was assessed by counting the number of working hours (eight hours equal one day) required prior to the start of the measurements." See lines 186-187.

223 potential air contaminations have to be considering. Considered.

Addressed: "... potential air contaminations have to be considered ..." see lines 299-300.

239 could contains some portions of the roots respiration. Contain.

Addressed: "... this technique did not contain..." see lines 252-253.

248 was statistically equal than. Equal to.

Addressed: "...was statistically equal to ..." see lines 278-279.

55 *377 asindependent. Add space.*

Addressed: "...temperature as independent..." see lines 408.

387 deciduousforest. Add space.

Addressed: "... deciduous forest ..." line 419.

433 for 13 C Analysis. 13C.

60 Addressed: "... for 13C ..." line 467.

J. van Leeuwen (Referee)

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Received and published: 6 February 2018

The paper presents an interesting comparison of different respiration measures. I like the paper in general, and it's generally well written. I do however have a number of points on which the paper can be improved for better understanding and readability.

To begin with, please have a good look at the order of paragraphs in methods, results and discussion. Currently the used respiration methods are discussed in random order. I would suggest to follow a consistent order, for example following tables 6 and 7 in all three sections, so starting with regression, followed by lab incubation, root exclusion bags and finish with d13 C. This would improve the readability of the paper.

Addressed: we have corrected this inconsistence. The different sections, paragraphs and sentences related to the five methods used were rearranged to follow a consistent order. Specifically, the following order was standardized throughout the text and tables: root regression, soil cores incubation, hand-sorted root exclusion bags (HS), intact root exclusion bags (IB) soil δ 13C-CO2 natural abundance.

The five used respiration measures are mentioned in the abstract (lines 15-17), but the introduction ends with simply stating that five methods were compared without further elaboration. Please explain the used methods here as well (for example in line 56), with the underlying thoughts on why these specific methods were selected.

Addressed: More detailed information about the specific methods selected were added at the end of the introduction; "The goal of our study was to compare partitioning methods to separate CO2 efflux into its Rs and Rh component in a subtropical secondary forest in Hong Kong. Five methods were selected based on their suitability in the studied ecosystem. Three methods were traditional techniques (i.e. regression between root mass and root derived CO2; root exclusion bags with hand-sorted roots; and soil δ 13C-CO2 natural abundance) and two were innovative variation of existing methods (i.e. root exclusion bags with intact soil blocks; and lab incubations with minimally disturbed soil microcosm cores). The influence of soil moisture and temperature on CO2 efflux was also analyzed." See lines 54-60.

At the start of the methods section, after describing the site selection, please start with soil characterization (lines 151-160).

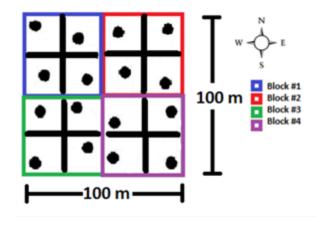
Addressed: section 2.1 is now "Soil general characterization" see line 73.

Also move the soil classification (line 177) to this section.

We recognize that stating the soil type name in the method section would beforehand help the readers to understand the ecosystem were the study was performed. However, we consider that the soil classification we did with the World Reference Base (WRB) system an integral part of the results because it had not been done before in this area; and accordingly, further investigations in this region will refer to our determination/classification.

It could be helpful for understanding the sampling procedure followed in the lab incubation method if a sampling schema is added as figure, as currently it's a bit difficult to follow what happens with the various groups of samples (starting with 16 from the field, and in line 196 there are 22 samples?).

Addressed: a figure showing the sampling procedure followed was added as new Fig 1. Also, clarification was made in the text "... four groups of four soil cores (stratified random design) were collected ..." Lines 105-106.



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Fig. 1. Field sampling for the lab incubations, stratified random design in the 1 ha study area.

In addition, if I read it correctly, the same samples went sequentially from a low incubation temperature to a high temperature. In the ideal scenario, the samples would be divided over the different incubation temperatures and studied independently. In the current setting dependency of the samples could be argued (repeated measurements from the same samples), but more importantly it adds uncertainty in terms of C present in the soils and potential shifts in the microbial community during the incubation (i.e. adapting to lower resource availability). Please include these limitations in the discussion of the used method.

Addressed: The section 4.2 was extended and improved to clarify this potential issue; "to prevent potential shifts in the microbial community during the incubations (i.e. adapting to lower resource availability), prior to the beginning of the experiment the microcosms were left stabilize for two weeks. Accordingly, the fresh and labile organic residues that would in the other segregation methods contribute to the soil respiration had already decomposed before the beginning of the incubations."

120 lines 248-250.

Please remove Table 3, as it contains only information that is fully described in the text already (lines 197-198).

Addressed: Table 3 was removed.

Also, please use consistent units for respiration throughout the paper, so either Mg C ha-1 y-1 or g m2 h-125 1 to improve comparability of the various measures.

Addressed: To be consistent with gas flux units, now everything is presented as Mg C ha⁻¹ y⁻¹.

Some more specific comments:

L124 (equation 1): please specify what f(x,y) represents (CO2 efflux).

Addressed: "where f(x, y) is the CO_2 efflux function ..." see line 125.

130 L150: In the site selection you describe only forest vegetation. What do you mean with different landforms here?

Addressed: "... different landforms (i.e. back-slope and foot-slope) present at the site ..." see line 74.

L177: Please move this sentence to the method section on soil characterization.

See response to "Also move the soil classification (line 177) to this section." comment above.

L195-203: Please split this section according to the respiration measures that you discuss, as regression and incubation are presented as separate methods in tables and in the rest of the paper.

Addressed: this section was divided into 3.2 and 3.3, see lines 201-208.

L199-200: please check carefully, I assume that you mean that Rh represents 54% of Rs, so I think the first time Rs in the sentence should be Rh.

Addressed: This sentence was corrected "The Rh measured when the root regression technique was performed (October 2016) was 6.0 ±2.4 Mg C ha-1 y-1, equivalent to 54% of the Rs (Table 5)." Lines 206-207.

L201-202: are these CO2 efflux values for all moisture groups combined?

Addressed: "... the average (moisture levels combined) CO₂ efflux ... " see line 209.

145 L208: The notably large respiration components. Please remove this sentence as it's trivial information.

Addressed: This sentence was removed.

L245: When I look at figure 1b I don't see an equilibrium in any of the groups. There seems to be the variation in respiration following the changes in temperature, but it does not show stability. Perhaps use the wording careful here.

Addressed: section 4.3 was amended and improved to clarify what we meant with the term "equilibrium"; "After the three months of soil stabilization period, both bag methods for partitioning total soil respiration and root-free soil respiration components successfully produced Rs>Rh in every sampling dates indicating that efflux rates within the bags had reached an apparent post disturbance state (Fig. 1)." lines 273-276.

L211-212: It is unclear where these values are coming from, as I can't find or derive them from table 6?

Addressed: This sentence was re-written to improve clarity "Using the $\delta 13C$ -CO2 method as base-line, the percent increase/decrease of the root regression, lab incubation, hand-sorted and intact block (IB) root exclusion techniques were -11%, -72-87%, +30% above and -20%, respectively (Table 5)." Lines 227-229.

L323: I find "minimally disturbed microcosms" a bit doubtful here, as soils were sorted and repacked in different vials in the lab. Keeping the soil and the larger aggregates intact during such a procedure is hardly possible, I assume that you refer here to undisturbed microaggregates?

Addressed: section 2.3 was amended to specify it is the micro-aggregate that should not be disturbed/broken. "special care to not destroy the micro-aggregates" Lines 107-108.

Minor textual comments:

L64: surveyed -> found

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Addressed: "the following genera were found ..." see lines 67-68.

L102: report -> analysis

170 Addressed: "A linear regression analysis ..." see line 100

L105: soil cores with a volume of

Addressed: "undisturbed soil cores with a volume of ..." see line 104.

L120: the boxes were opened to vent

Addressed: "each measurement session, the boxes were opened to vent ..." see line 120.

175 L140: previous to -> before

Addressed: "... 90 minutes before ..." see line 164.

L180: sub-superficial -> sub-surface or sub-soil

Addressed: "The sub-surface soil ..." see line 185.

L184: based on previous results

180 Addressed: "...status based on previous results ..." see line 195.

L185: type of parent material and level of mineralization of the bedrock

Addressed: "... type parent material and level of mineralization of the bedrock ..." see lines 199-200.

L187:Split sentence in two, the first discussing climate, the second discussing respiration.

Addressed: "During the root exclusion bags measurements period (Feb-Apr 2017), the average air temperature was 16°C and the total rainfall 107 mm. During that period the Rs averaged 6.1 Mg C ha-1 y-1 (Fig. 1)" see lines 213-214.

L252: rephrase for readability: Although not statistically significant, the maximum Rh in the HS and IB bags was found at a relatively low moisture content (9.5 and 21.4%, respectively, Table 4).

Addressed: "... supported by the fact that in the regression fit the maximum Rh was when moisture content was relatively low ..." see lines 283-284.

L262: please remove "In this study the regression had ten points (45%) outside the confidence interval but". The rest of the sentence is a repetition of L197-198, so either generalize for discussion or remove as well.

Addressed: section 3.2 was shortened to remove the redundancies.

195 *L295: variance -> variation*

Addressed: "Fourth, the large variation ..." see line 319.

L298: in this regard

Addressed: "In this regard, the δ^{13} C-CO₂ ..." see line 323.

L300: to standardize

200 Addressed: "... by roots and to standardize ..." see line 324.

L338: variance of -> variation in

Addressed: "...results showed large variation in ..." see lines 364-365.

Separation of soil respiration; a site-specific comparison of partition methods

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Abstract. Without accurate data on soil heterotrophic respiration (Rh), assessments of soil carbon (C) sequestration rate and C balance are challenging to produce. Accordingly, it is essential to determine the contribution of the different sources of the total soil CO₂ efflux (Rs) in different ecosystems, but to date, there are still many uncertainties and unknowns regarding the soil respiration partitioning procedures currently available. This study compared the suitability and relative accuracy of five different Rh/Rs partitioning methods in a subtropical forest: (1) regression between root biomass and root derived CO₂; efflux; (2) lab incubations with minimally disturbed soil microcosm cores; (3) root exclusion bags withhand-sorted roots; (4) root exclusion bags with intact soil blocks; (3) root exclusion bags withhand-sorted roots; (4) lab incubations with minimally disturbed soil microcosm cores; and (5) soil δ¹³C-CO₂ natural abundance. The relationship between Rh and soil moisture and temperature was also investigated. A qualitative evaluation table of the partition methods with five performance parameters was produced. The Rs was measured weekly from February 3rd to April 19th 2017 and found to average 6.1 ±0.3 Mg C ha⁻¹ y⁻¹. During this period, the Rh measured with the in-situ mesh bags with intact soil blocks and hand-sorted roots were estimated to contribute 49 ±7% and 79 ±3% of Rs respectively. The Rh percentage estimated with the root biomass regression, microcosm incubation and δ¹³C-CO₂ natural abundance were 54 ±41%, 8-17% and 61 ±39% respectively. Overall, no systematically superior or inferior Rh/Rs partition method was

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found. The paper discusses the strengths and weaknesses of each technique with the conclusion that combining two or more methods optimizes Rh assessment reliability.

1 Introduction

230 During the 2016 Convention of Parties (COP21) of the United Nations Framework Convention on Climate Change (UNFCCC) in Paris, the goal of increasing global soil organic carbon (SOC) stocks by 0.4 percent per year was set, with the aimof mitigating global anthropogenic greenhouse gas emissions (Minasny et al., 2017). This ambitious target was set based on the concept that the SOC in the top soil layer is sensitive and responsive to management changes and may offer opportunities to mitigate the current increases in atmospheric CO₂ concentration (McConkey et 235 al., 2007). Of the carbon (C) that enters into ecosystems via photosynthesis, a fraction is directly respired by the roots and above ground plant parts (autotrophic respiration) to produce energy (i.e. adenosine-5'-triphosphate), with the other fraction synthesized into organic molecules. Some of these C-containing compounds are harvested or consumed by herbivores and the remainder is added to the soil as plant residues (Janzen et al., 1998). Subsequently, a portion of these fresh organic compounds are respired by organisms (heterotrophic respiration) and the other portion is converted into 240 SOC by the genesis of soil organic matter (SOM) (Janzen, 2006; Lal, 2005). If the amount of new organic residues added to the soil is greater than the Clost by SOC decomposition, SOC content increases (Ellert and Bettany, 1995). Typically, many years (up to decades) are needed to assess SOC stock changes over time in order to evaluate which management practices are beneficial for SOC sequestration (Harmon et al., 2011; Wood et al., 2012). This timeframe is impractical for policy makers to evaluate the mitigation potential of different land management practices, in particular 245 with the pressing need of the UNFCCC goal of increasing the global SOC stocks by 0.4 percent per year. An alternative approach that allows a more rapid evaluation of these long term impacts is to combine the SOC stock change procedure (e.g. VandenBygaart et al., 2008) with the soil C efflux balance approach (i.e. Hergoualc'h and Verchot, 2011), which although demanding and with some uncertainties can provide results on soil dynamics over an annual basis. The soil Cefflux balance approach involves calculating the rate of Centry and exit in the soil. However, 250 the total CO₂ efflux (Rs) from soil does not provide the necessary information to estimate whether the soil is a net source or net sink for atmospheric CO₂ (Kuzyakov and Larionova, 2005). Total soil efflux is a combination of root based respiration (autotrophic (Ra)) and heterotrophic respiration (Rh). Autotrophic respiration does not contribute to

net Closses to the atmosphere as it is cycled within the ecosystem, whereas Rh represent net Closses. However, the

boundary between Ra and Rh is not easy to distinguish (i.e. the rhizo-microbial respiration is linked to both) and realistic Rh assessments are difficult to produce (Braig and Tupek, 2010).

Reviews of Rh-Rs segregation methods have been made (e.g. Kuzyakov, 2006) but no site specific study has been made analysing several different partition techniques simultaneously. The goal of our study was to compare five different partitioning methods to separate CO₂ efflux into its Rs and Rh component in a subtropical secondary forest in Hong Kong. The influence of soil moisture and temperature on CO₂ efflux was also analyzed. The goal of our study was to compare partitioning methods to separate CO₂ efflux into its Rs and Rh component in a subtropical secondary forest in Hong Kong. Five methods were selected based on their suitability in the studied ecosystem. Three methods were traditional techniques (i.e. regression between root biomass and CO₂ efflux; root exclusion bags with hand-sorted roots; and soil δ ¹³C-CO₂ natural abundance) and two were innovative variation of existing methods (i.e. root exclusion bags with intact soil blocks; and lab incubations with minimally disturbed soil microcosm cores). The influence of soil moisture and temperature on CO₂ efflux was also analyzed.

2 Methodology

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The research was conducted in a subtropical secondary forest of Hong Kong (Tai Po Kau Nature Reserve; 22° 27′N, 114° 11′E). The landscape is typical of the escarpment of the Tai Mo Shan mountain range, the system formed by volcanic activities in the Late Jurassic epoch (Langford et al., 1989). The rocks are mainly rhyodacite to rhyolite from the Tsuen Wan Volcanic Group (Davis et al., 1997). The study site was approximatively 600 m. above sealevel and the slope surfaces were stable and vegetated. The forest was approximatively 50 years old and was covered with continuous canopy. More than 100 plant species were registered in the Nature Reserve. The following genera were surveyedfound in the study area: *Machilus sp, Meliosma sp, Garcinia sp, Engelhardia sp, Psychotria sp, Ilex sp, Eurya sp and Lithocarpus sp* (Tong, 2015). The mean annual temperature was 23.3°C and annual precipitation 2400 mm with a hot-humid season (April–September) and a cool-dry season (October–March) (Hong Kong Observatory). The study area was 0.51 ha and was located inside a long-term research site belonging to the Chinese University of Hong Kong. The canopy was closed in the area with an average solar radiation at 2 m high of 13.8 W/m² (non-published data).

2.51 Soil general characterization

Four soil profiles were dug in the study area, characterizing the different landforms (i.e. back-slope and foot-slope) present at the site. Morphological description was done according to Jahn (2006) and the soil was classified with the

World Reference Base (IUSS-Working-Group-WRB, 2014). Soil pH was determined with a glass—calomel electrode pH meter (McLean, 1982). Rainfall and air temperature were recorded hourly with a HOBO Weather station (rain gauge, S-RGB-M002; air temperature/RH, sensor S-THB-M008, Onset Computer Corp., USA). Water holding capacity was assessed by saturating the soils, allowing them to freely drain for 24 h and determining gravimetric water content after oven-drying at 105 °C following Arcand et al. (2016). Root biomass was measured by collecting soil cores (inner diameter 5 cm, height 5 cm) and determined using the approach of Tufekcioglu et al. (1999). The soil was dried, finely ground, and subsequently analyzed for total C and N content using a CNS Analyzer System (Perkin Elmer 2400 Series II CHNS/O Analyzer, USA) at the Earth System Science Laboratory of the Chinese University of Hong Kong (ESSL-CUHK).

2.2 Partitioning soil respiration

To produce estimates of % Rh, five different approaches were designated based on their suitability in the study area. We decided to use three customary methods often employed in this type of ecosystem (i.e. regression between root biomass and CO_2 efflux; root exclusion bags with hand-sorted roots; and soil δ^{13} C- CO_2 natural abundance) and two innovative variation of existing techniques (i.e. root exclusion bags with intact soil blocks; and lab incubations with minimally disturbed soil microcosm cores).

2.2.1 Regression between root biomass and CO₂ efflux method

The root biomass regression technique is based on the relationship between the CO₂ emitted by the root-rhizosphere and root biomass and the CO₂ efflux slope fitted from SOM decomposition (i.e. Rh), corresponding to the intercept of the linear regression line (Kucera and Kirkham, 1971). This method was made following Farmer (2013) with 22 sampling spots. Each spot was a square of 20 x 20 cm randomly distributed in the study area. In each spot, Rs was determined per triplicate using a portable IRGA as described above. Concurrently with CO₂ efflux measurements, air and soil (10 cm depth) temperatures and soil volumetric moisture content were measured at each sampling spot. Immediately after the Rs measurement, the 20 x 20 cm squares were excavated to 25 cm depth. All the visible roots (diameter larger than 0.1 cm) from the excavated soil were collected. In the lab, the roots were washed and then oven dried at 60°C until a steady dry weight was attained, which was then recorded. A linear regression analysis between root quantity and CO₂ efflux was performed using the program R Foundation for Statistical Computing version 2.8.1 (R Development Core Team, 2008).

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2.2.2 Lab incubations

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Software, San Jose, CA).

For the lab incubations, undisturbed soil cores with a volume of 98 cm³ (inner diameter 5 cm, height 5 cm) were collected using a stainless-steel core soil sampler from the upper part of the soil profile (0-5 cm). In the study area, four groups of four soil cores (stratified random design) were collected, then pooled per group and brought to the lab (Fig. 1. complementary fig.1). Subsequently all visible roots were removed but with special care to not destroy the microaggregates. The soil was then repacked to original bulk density in minimally disturbed soil microcosm cores of 45 cm³ (inner diameter 3.5 cm, height 5 cm). The soil cores were separated in four groups of different volumetric moisture content (i.e. 15, 25, 35 and 45). These moisture levels corresponded to the natural annual fluctuation in the field (i.e. from dry to moist season) (Cui and Lai, 2016). After moisturizing the samples with distilled water, each individual soil core was placed into a hermetically sealed 2.9 dm³ plastic container and left to stabilize in dark for two weeks at 25°C. After that, the experiment lasted four weeks and had four different incubation temperature levels (one per week; 14°C, 20°C, 26°C and 32°C) corresponding to the minimum, intermediate and maximum soil temperature values in the field based on preliminary studies (Cui and Lai, 2016). At the beginning of each week, the soil cores were pre-incubated in their incubation box to their corresponding weekly temperature (i.e week #1, 14°C ... week #4, 32°C) for 3 days and then opened and vented for one minute. From all the boxes gas samples were collected (20 ml) with an air-tight syringe (t=0, 24, 72 hour) after box closure. The CO₂ concentrations were analyzed within 48 hours with a gas chromatograph (GC system 7890A, Agilent Technologies). The GC system was equipped with a flame ionization detector and an electron capture detector to quantify -CO₂ concentration. Between each measurement session, the boxes were opened to vent and the moisture of the soil cores was re-adjusted if needed. Gaussian 3D regression fitted curve was derived as shown in equation 1, using SigmaPlot version 10.0 (Systat

$$f(x,y) = a \times \exp\left[-0.5 \times \left(\frac{x - xo}{b}\right)^2 + \left(\frac{y - yo}{c}\right)^2\right]$$
 (1)

where f(x,y) is the CO₂ efflux function; a, b and c are constant coefficients; x is the soil temperature (°C); y is the soil moisture content (%); x_0 is the average temperature; y_0 is the average soil moisture.

2.2.32.1 Root exclusion bag methods

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To partition the CO₂ efflux in-situ into Rs and Rh using mesh bags, two different approaches were followed: 1) the traditional dug soil with hand-sorted root removal and refilling method (HS) (Fenn et al., 2010; Hinko-Najera, 2015) and 2) a variant of it with intact soil blocks (IB). The HS method consisted of digging a pit for each bag with a size matching the bag dimensions (20×20 cm, depth: 25 cm) where the soil is-was excavated in layers (to maintain soil horizons) and visible roots are were removed before repacking the bag inside the pit with the removed soil. The IB variant of this technique consisted in-of extracting a cube as intact as possible from the soil (20×20 cm, depth: 25 cm). This was Then, tightly placeding the soil block into the micromesh bag and inserteding it back into its original pit. For both methods, the same type of micromesh bags (38µmnylon mesh), closed at the bottom but open at the top were used. This mesh size was used to impede roots from entering inside the bags, but allowed mycorrhiza to penetrate (Moyano et al., 2007). In all the pits excavated, no root below 25 cm depth was observed. Collars measuring 10 cm diameter were installed on the soil in the center of each bag to a depth of 8 cm, for heterotrophic emissions sampling.

Seven plots were randomly distributed inside the study area. In each plot, an IB bag was paired with a HS bag with space of 150 cm between them. The root exclusion bags were installed during the month of October 2016 and were let to stabilize for three months. At 1 m distance from each root exclusion bag, a collar was inserted into non-disturbed soil to measure Rs. To assess Rs and Rh without the influence of litterfall decomposition, the collars were cleared of leaves and flowers on a weekly basis.

From February 3rd to April 19th 2017 the collars were measured weekly with an IRGA (Environmental Gas Monitor, EGM-4, PP Systems, UK) attached to a soil respiration chamber (SRC-1, PP Systems, UK). Soil temperature and soil moisture were measured in the area located between the collar and the edge of the bag (to 10cm depth, HH2, Delta-T Devices, Cambridge-England). At the end of the study all the root exclusion bags were removed from the soil and inspected to ensure that no root had penetrated inside. The soil inside the measurement collars was then collected to assess bulk density (van Reeuwijk, 1992). Mathematical calculation and descriptive statistical analyses were done with Microsoft Excel XP[®].

2.2 Root and carbon dioxide efflux regression method

The regression technique is based on the relationship between the CO₂ emitted by the root rhizosphere and root biomass and the CO₂ efflux derived from SOM decomposition (i.e. Rh), corresponding to the intercept of the linear

regression line (Kucera and Kirkham, 1971). This method was made following Farmer (2013) with 22 sampling spots. Each spot was a square of 20 x 20 cm randomly distributed in the study area. In each spot, Rs was determined per triplicate using a portable IRGA as described above. Concurrently with CO₂ efflux measurements, air and soil (10 cm depth) temperatures and soil volumetric moisture content were measured at each sampling spot. Immediately after the Rs measurement, the 20 x 20 cm squares were excavated to 25 cm depth. All the visible roots (diameter larger than 0.1 cm) from the excavated soil were collected. In the lab, the roots were washed and then oven dried at 60°C until a steady dry weight was attained, which was then recorded. A linear regression report between root quantity and CO₂ efflux was

2.3 Lab incubations

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Software, San Jose, CA).

For the lab incubations, undisturbed soil cores of volume 98 cm² (inner diameter 5 cm, height 5 cm) were collected using a stainless-steel core soil sampler from the upper part of the soil profile (0–5 cm). In the study area, four groups of four soil cores were collected then pooled per group and brought to the lab. Subsequently all visible roots were removed but with special care to not destroy the small aggregates. The soil was then repacked to original bulk density in minimally disturbed soil microcosm cores of 45 cm² (inner diameter 3.5 cm, height 5 cm). The soil cores were separated in four groups of different volumetric moisture content (i.e. 15, 25, 35 and 45). These moisture levels corresponded to the natural annual fluctuation in the field (i.e. from dry to moist season) (Cui and Lai, 2016). After moisturizing the samples, each individual soil core was placed into a hermetically sealed 2.9 dm² plastic container. The experiment lasted four weeks and had four different incubation temperature levels (one per week; 14°C, 20°C, 26°C and 32°C) corresponding to the minimum, intermediate and maximum soil temperature values in the field based on preliminary studies (Cui and Lai, 2016). At the beginning of each week, the soil cores were pre-incubated in their incubation box to their corresponding weekly temperature (i.e. week #1, 14°C ... week #4, 32°C) for 3 days and then opened and vented for one minute. From all the boxes gas samples were collected (20 ml) with an air tight syringe (t= 0, 24, 72 hour) after box closure. The CO₂ concentrations were analyzed within 48 hours with a gas chromatograph (CC system 7890A, Agilent Technologies). The CC system was equipped with a flame ionization detector and an electron capture detector to quantify and CO₂. Between each measurement session, the boxes opened to vent and the moisture of the soil cores was re-adjusted if needed.

Gaussian 3D regression fitted curve was derived as shown in equation 1, using SigmaPlot version 10.0 (Systat

$$f(x,y) = a \times \exp\left[-0.5 \times \left(\frac{x-x_0}{h}\right)^2 + \left(\frac{y-y_0}{a}\right)^2\right] \tag{1}$$

where a, b and c are constant coefficients; x is the soil temperature (°C); y is the soil moisture content (%); x_a is the average temperature; y_a is the average soil moisture.

 $\frac{2.2.42.4}{6}$ δ^{13} C natural abundance method

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Millard et al. (2010) have demonstrated that the natural abundance δ^{13} C (‰) of Rs falls between the δ^{13} C values of the Rh and Ra. The δ^{13} C of Rs and ARh respiration was determined following Lin et al. (1999) and Millard et al. (2010). The isotopic partitioning experiment assessed values of the δ^{13} C of the Rs, Ra and Rh. The sampling took place on March 15th 2017. A closed chamber (10 cm diameter, 10 cm high) was positioned on each emissions measurement collar (n=7 as described in section 2.12.2.3). The chambers were flushed for 2 minutes with CO₂-free air to remove all the atmospheric air trapped within the headspace. Chambers were left to incubate for 40 minutes to ensure the concentration of the chamber sample reached above 400 ppm of CO₂ from which a duplicate sample of the gas in the chamber headspace were extracted into evacuated vials to give the δ^{13} C of the Rs. Subsequently, the soil under the chamber was dug and immediately brought to the lab (ESSL-CUHK, less than 30 minutes travel) where the soil and the roots were carefully separated. The roots were gently washed with water to remove adhered soil aggregates and slightly dried afterward with paper towels. Samples of 5 g of root and 10 g of root-free soil per chamber were incubated in CO₂ free air in 250 ml airtight glass bottles to give the δ^{13} C of the Ra and Rh respectively. The bottles were left to incubate for 90 minutes before duplicate extraction into evacuated vials. As recommended by Midwood et al., (2006), previous to before gas sample extraction, the butyl rubber septa used to seal the vials were heated at 105°C for 12 h. The C isotope ratio of the CO₂ in all samples was analyzed using a Gas-bench II connected to a DeltaPlus Advantage isotope ratio mass spectrometer (both Thermo Finnigan, Bremen, Germany) at the James Hutton Institute Scotland UK. The δ¹³Cratios, all expressed relative to Vienna-Pee-Dee Belemnite (VPDB), was calculated with respect to CO₂ reference gases injected with every sample and traceable to International Atomic Energy Agency reference material NBS 19 TS-Limestone. Measurement of the individual signatures of the natural abundance δ^{13} C of the Rs. Rh and Ra allowed partitioning between the different sources using the mass balance mixing model (Lin et al., 1999; Millard et al., 2010):

$$\%Rh = 1 - \frac{\delta Rs - \delta Rh}{\delta Ra - \delta Rh} \times 100 \frac{X - 100}{\delta Ra}$$

(2)

where %Rh is the proportion of Rh from Rs, and δRs , δRh and δRa are the δ^{13} C isotopic signatures.

2.5 Soil general characterization

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Four soil profiles were dug in the study area, characterizing the different landforms present at the site. Morphological description was done according to Jahn (2006) and the soil was classified with the World Reference Base (IUSS-Working Group WRB, 2014). Soil pH was determined with a glass—calomel electrode pH meter (McLean, 1982). Rainfall and air temperature were recorded hourly with a HOBO Weather station (rain gauge, S-RCB-M002; air temperature/RH, sensor S-THB-M008, Onset Computer Corp., USA). Water holding capacity was assessed by saturating the soils, allowing them to freely drain for 24 h and determining gravimetric water content after oven drying at 105 °C following Areandetal. (2016). Root biomass was measured by collecting soil cores (inner diameter 5 cm, height 5 cm) and determined using the approach of Tufekcioglu et al. (1999). The soil was dried, finely ground, and subsequently analyzed for total C and N content using a CNS Analyzer System (Perkin Elmer 2400 Series II CHNS/O Analyzer USA).

2.36 Qualitative comparison of segregation methods

While there is much debate in the literature, and methods are still being developed, Lisotopic partitioning methods are increasingly being recognized as a more accurate approach to segregation of Rh/Rs than non-isotopic techniques (Paterson et al., 2009; Kuzyakov, 2006). Therefore Thus, for comparison purposes we used $_{T}$ the soil δ^{13} C natural abundance method was used as a reference point for segregation relative accuracy. Partition methods that had Rh%: <10, 10-20 and >20 lower or larger than the δ^{13} C-CO₂ natural abundance were categorized as high, intermediate and low relative accuracy, respectively. The level of precision of the segregation methods was determined with the statistical variance associated with the Rh/Rs ratio averages. High, intermediate and low precision were attributed to Rh% standard errors of <10, 10-20 and >20, respectively. The level of complexity was evaluated with the number of steps required to complete each method. For example, the hand-sorted root exclusion bags technique was judged as a four steps method (pit excavation, root removal, bag/pit refiling, and CO₂ efflux measurements). Methods with five steps or less were deemed simple and six steps or more deemed as complex. The time inversion needed to set up the experiment was assessed by counting the number of working hours (eight hours equal one day) required prior to the start of the measurements. The time inversion needed to produce seasonal trends was the number of months of

measurements (in the field or in the lab) required to characterize the Rh at the different temperature and moisture levels of the year.

3 Results

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3.1 Soil characteristics

According to their morphology and diagnostic properties, the soil was classified as Alic Umbris ol (Nechic) and Haplic Alisol (Nechic) (IUSS-Working-Group-WRB, 2014). The difference between the two soil groups was the thickness of humus-containing horizon (between 20 and 30 cm for the Umbrisol; while, 10 to 20 cm for the Alisol). The A horizon had high organic C content (3.2 ±0.2%) and high acidity (pH H2O 4.2) (Table 1). The sub-superficial surface soil was represented by clayey yellow-colored profiles with an argic horizon. Soil texture was heavier in the argic horizon than in the topsoil and parent material. The structure in all the soil profiles was predominantly granular in the upper horizons, whereas the argic horizon was characterized by subangular blocky structure (Table 1). The argic horizon was deemed to be of high-activity clays and low cation base status based toon previous results in the area (Tong, 2015), along with soil acidity, type of parent material and level of mineralization of the bedrock in the soil pits.

3.32 Root regression and Regression between root biomass and CO2 efflux

The 22 quadrats used for the root biomass regression assessment yielded an average Rs of $11.1 \pm 1.0 \,\mathrm{Mg}\,\mathrm{CO}_2$ —Cha¹ y⁻¹. The regression of the CO₂ efflux against root biomass produced a statistically significant slope correlation of 0.08 $\pm 0.04 \,\mathrm{g}\,\mathrm{CO}_2 \,\mathrm{m}^2 \,\mathrm{h}^{-1}$ per mg root cm⁻³ (p=0.03), and set the intercept at $0.25 \pm 0.10 \,\mathrm{g}\,\mathrm{CO}_2 \,\mathrm{m}^2 \,\mathrm{h}^{-1}$ (p=0.02) which represented the basal efflux in the absence of roots i.e. the Rh (Fig. 2 and Table 3). The Rh measured (i.e. slope intercept) when the root biomass regression technique was performed (October 2016) was $6.0 \pm 2.4 \,\mathrm{Mg}\,\mathrm{C}\,\mathrm{ha}^{-1}\,\mathrm{y}^{-1}$, equivalent to 54% of the Rs (Table 6).

3.3 Lab incubation

The 22 quadrats used for the root regression assessment yielded average Rs of 0.46 ± 0.04 g CO₂ m² h⁻¹. The regression of the CO₂ efflux against root density produced a statistically significant slope correlation of 0.08 ± 0.04 g CO₂ m² h⁻¹ per mg root cm⁻² (p=0.03), and set the intercept at 0.25 ± 0.10 g CO₂ m² h⁻¹ (p=0.02) which represented the basal efflux in absence of root i.e. the Rh (Fig. 2 and Table 3). The Rs measured when the root regression technique was performed (October 2016) was 11.1 ± 1 Mg C ha⁻¹ v⁻¹ (Table 6), equivalent to 54% of the Rs.

During the incubation with minimally disturbed soil microcosms, the average (moisture levels combined) CO₂ efflux at 14, 20, 26 and 32°C was 0.015136 +0.50021, 0.670282 +0.38016, 0.05851.40 +0.91038 and 2.240.0938 +1.390.058

Mg CO₂-C ham²¹ yh⁻¹, respectively (Fig. 3). The exponential relationship between CO₂ efflux, soil temperature and moisture is presented in Table 4.

3.24 Root exclusion bag methods

During the root exclusion bags measurements period (Feb-Apr 2017), the average air temperature was 16°C and the total rainfall 107 mm_x and and the During that period the Rs averaged 6.1 Mg C ha⁻¹ y⁻¹ (Fig. 1). One of the requirements for the suitability of root exclusion bag methods to estimate Rh is that soil bulk density, soil temperature and moisture are statistically equal inside and outside of the bags. In this experiment, no significant differences were detected regarding the bulk density and soil temperature (p=0.87 and p=0.15, respectively) but the volumetric soil moisture in the HS bags was on average 17% lower than outside the root exclusion bags (p=0.04) (Table 2). As would be expected, all Rh IB and Rh HS efflux rates were lower than the Rs efflux at each measurement date. Throughout the experiment, the Rh IB was repetitively consistently lower than the Rh HS except on March 31st (Fig. 1b).

2.2 Poot regression and lab incubation

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The 22 quadrats used for the root regression assessment yielded average Rs of 0.46 ±0.04 g CO₂-m²-h². The regression of the CO₂-efflux against root density produced a statistically significant slope correlation of 0.08 ±0.04 g CO₂-m²-h² per mg root cm² (p=0.03), and set the intercept at 0.25 ±0.10 g CO₂-m²-h² (p=0.02) which represented the basal efflux in absence of root i.e. the Rh (Fig. 2 and Table 3). The Rs-measured when the root regression technique was performed (October 2016) was 11.1 ±1 Mg C ha²-y² (Table 6), equivalent to 54% of the Rs-

During the incubation with minimally disturbed soil microcosms, the average CO₂ efflux at 14, 20, 26 and 32°C was 0.0151 ±0.021, 0.0282 ±0.016, 0.0585 ±0.038 and 0.0938 ±0.058 g CO₂ m²-h⁻¹, respectively (Fig. 3). The exponential relationship between CO₂ efflux, soil temperature and moisture is presented in Table 4.

3.54 Soil δ^{13} C-CO₂ natural abundance

On a land-scape basis, \mathbf{T}_1 the δ^{13} C-CO₂ natural abundance determination method reasonably satisfactorily segregated the three respiration components (Table 54). The fact that the δ^{13} C-CO₂ of the Rh HS, Rh IB, and Rh lab were in a very close range (i.e. -16.52 to -16.75), indicated in the field the efflux measured in the root exclusion bags were not contaminated with root respiration. Based on the δ^{13} C-CO₂ of the Rs (-18.21±0.53), the Rh lab (-16.75±0.54), and the

Ra lab (-20.44± 0.65) the average percentage of heterotrophic respiration was 61 ±39% (Table 65). The notably large standard error of the percentage of heterotrophic respiration was due to the large variance in the δ^{12} C CO₂ of the three respiration components.

Using the δ^{13} C-CO₂ method as a base-line, the increase/decrease of Comparing with the Rh from the δ^{13} C-CO₂ method, the Rh from root biomass regression, lab incubation, hand-sorted and intact block (IB) root exclusion techniques were _11% below, _72-87% below, \pm 30% above and _20% below, respectively (Table 6).

4 Discussion

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4.1 Regression between root biomass and CO₂ efflux technique

As demonstrated by Gupta and Singh (1981) the intercept of the regression line between the independent variable (i.e., root biomass) and the dependent variable (i.e., Rs.) corresponds to soil respiration in absence of root (i.e., Rh.) (Fig. 2). In this study the regression had ten points (45%) outside the confidence interval but the intercept and slope -were still statistically significant-. This uncertainty in the regression fit was likely caused in large part by the older roots which are bulkier but respire less than fine and young roots (Behera et al., 1990). However, this method had the closest average Rh/Rs ratio to the δ^{13} C natural abundance technique. Consequently the root biomass regression technique was assessed as high relative accuracy and low precision (Table 7). Previous studies also highlighted large variation of CO₂ efflux and root biomass which causes relatively low coefficient of determinations (Behera et al., 1990; Farmer, 2013). In accordance to Kuzyakov (2005), this method was comparatively simple (Table 7).

4.2 Lab incubation method

Interpreting soil respiration processes in response to seasonal changes is generally challenging because soil temperature and moisture regularly covary (Carbone et al., 2011; Davidson et al., 1998). The lab incubation technique was the only method capable of dividing the effect of soil temperature and moisture on Rh and to produce a significant Gaussian regression fit (Table 3). However, the microcosm incubation produced Rh values notably lower than the other techniques (Table 5). This mayight mainly be due to three different causes. First, the fact that the soil column in the incubation microcosms were 5 cm high thick while the A horizon in the field (i.e. where the Rh assessments from the other techniques were made) was 10 cm thick (Table 2). Second, to prevent potential shifts in the microbial community during the incubations (i.e. adapting to lower resource availability), previous prior to the beginning of the experiment the microcosms were left to stabilize for two weeks. Accordingly, the fresh and labiele organic residues that would in

the other segregation methods contribute to the soil respiration had already decomposed before the beginning of the incubations. Third, the low Rh of the lab incubation method could also be attributed in part to the fact that this technique did not contain any rhizomicrobial respiration and its priming effect (Kuzyakov et al., 2000). That is, this method produced Rh from basal microbial respiration which is considered to be from stabilized SOM with slow turnover rates (Kuzyakov, 2006 Neff et al., 2002). In view of that, with additional field and lab methods development it would be possible to further segregate Rh assessments into percentage of rhizomicrobial respiration, decomposition of plant residues and basal decomposition of SOM. Overall, the lab incubation technique was slightly more complex than the non-isotopic field Rh assessment methods but allowed a prompt determination of Rh whilst simulating year—round field environment conditions (Table 6). Further studies should test the effect of microcosm height on Rh in relation to

4.3 Root exclusion bags methods

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The HS and IB methods had % Rh of 79 ± 3 and 49 ± 7 %, respectively. The se variances around the means (i.e. ±3 and ±7 , respectively) were the lowest of all the field segregation methods tested (Table 5). Comparing the % Rh of the HS and IB with the δ^{13} C natural abundance technique, they resulted 18% above and 12% below, respectively. Thus the root exclusion bags methods were judged of intermediate relative accuracy and high precision. Also, the HS and IB methods were fast and simple to setup (Table 6).

field measurements and determine microbial Cuse efficiency by is othermal microcalorimetry during the incubations.

The micromesh size used in the root exclusion bags was 38µm which was reported to impede root penetration but to allow arbuscular mycorrhizal to spread inside the bags (Moyano et al., 2007; Rühr and Buchmann, 2010). In turn, Fenn et al. (2010) stated that in the mycorrhizal structures the arbuscules exist within roots, and therefore, the CO_2 efflux from these bags could contain some portions of the roots respiration. Contrary to this, the IB and HS air samples analyzed for δ^{13} C had an isotopic signature close and not statistically different from the gas samples collected in the lab airtight glass bottle of fresh soil without roots. This indicates that the root exclusion bags (both IB and HS) did not comprise traces of root respiration that had a significantly larger δ^{13} C- CO_2 signature (Table 4). After the three months period of soil stabilization period, both bag methods for partitioning total soil respiration and root-free soil respiration components successfully produced Rs>Rh in every sampling dates indicating that efflux rates within the bags had reached an apparent post disturbance state (Fig. 1). Also, in both IB and HS, soil temperature and bulk density were statistically equal to the surrounding soil (i.e. Rs) (Table 2). This indicates that the environmental conditions inside and outside of the bags were similar in respect to these two parameters. However, the soil moisture of the IB was

statistically equal to the surrounding soil but for HS it was significantly lower. This was likely caused by the breakdown of the original soil structure at the moment of root removal that increased the drainage inside the HS bags. Moyano et al. (2007) also found that soil moisture can be affected by the hand-sorted root exclusion bag method. Overall, HS had a moisture level 20% lower and an Rh efflux 60% larger than IB (Table 5). The larger HS Rh efflux compared with IB Rh could be due in part to the lower soil moisture in the former. This likelihood is supported by the fact that in the regression fit the maximum Rh was when moisture content was relatively low (i.e. y0, Table 3). In addition, the breakdown of numerous soil aggregates during the root removal likely allowed the soil microorganisms to thrive in previously encrusted SOM domains of the HS soil. It has been shown that the part of the SOM that is located in the interior of the soil aggregates is hardly accessible to microorganisms, and thus not easily mineralized unless the aggregates are shattered (Goebel et al., 2005).

4.14 Soil δ^{13} C natural abundance method

The three respiration components of this method (i.e. δ^{13} C-CO₂ from Rs, Rh and Ra) had large standard errors (Table 54) that produced a high uncertainty value in the Rh/Rs ratio assessment (61 ±39 %, Table 65). This method was accordingly deemed of low precision (Table 76). This, in turn, impeded our ability to produce an Rh/Rs ratio assessment in the individual collars. The low precision of this method indicates that some biases in the assessment of relative accuracy could potentially have been generated. This large δ^{13} C-CO₂ variance was likely caused by variability of δ^{13} C in soil and plants residues and also due to 13 C discrimination by plants that is affected by moisture content and nitrogen availability (Hogh-Jensen and Schjoerring, 1997). In addition, other studies reported the variability of δ^{13} C in soil or plants of at least 1–2‰, which in some cases can limit the capacity to produce soil respiration segregation assessments (Accoe et al., 2002; Cheng, 1996; Farquhar et al., 1989). Because soils are porous mediums, excluding any atmospheric CO₂ that has a different isotopic composition (i.e. δ^{13} C -7.5 to -8.5 ‰) to that of the Rs efflux is challenging and potential air contaminations have to be considereding when analyzing the results (Millard et al., 2010). In our study, the Rh δ^{13} C was measured in the field (IB and HS; potentially air contaminated) and from airtight containers in lab incubations of root free soil (Rs lab; not potentially air contaminated). Both ways produced δ^{13} C in a close range and without statistical differences between them (Table 54). This indicates that the chamber systemused in the field to collect the δ^{13} C efflux samples was adequately effective to prevent air contamination.

Overall, the soil δ^{13} C natural abundance method was fast to setup but was relatively complex to perform with a field and lab component to be accomplished within a short period of time (Table 76). Further studies should use a large amount of sampling points to attempt reducing the respiration components standard errors.

4.2 Root exclusion bags methods

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In the HS and HS methods had % kh of 79 ± 3 and 49 ± 7 %, respectively. These variances around the means (i.e. ± 3 and ± 7 , respectively) were the lowest of all the field segregation methods tested (Table 6). Comparing the % Rh of the HS and IB with the δ^{13} C natural abundance technique, they resulted 18% above and 12% below, respectively. Thus the root exclusion bags methods were judged of intermediate accuracy and high precision. Also, the HS and IB methods were fast and simple to setup (Table 7).

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aggregates during the root removal likely allowed the soil microorganisms to thrive in previously encrusted SOM domains of the HS soil. It has been shown that the part of the SOM that is located in the interior of the soil aggregates is hardly accessible to microorganisms, and thus not easily mineralized unless the aggregates are shattered (Goebel et al., 2005).

4.3 Root and carbon dioxide efflux regression technique

As demonstrated by Gupta and Singh (1981) the intercept of the regression line between the independent variable (i.e. root biomass) and the dependent variable (i.e. Rs.) corresponds to soil respiration in absence of root (i.e. Rh.) (Fig. 2). In this study the regression had ten points (45%) outside the confidence interval but the intercept (0.25 ±0.10 g CO₂ m² h⁻¹ and slope (0.08 ±0.04 g CO₂ X mg root cm²) were still statistically significant (P=0.02 and 0.03, respectively) (Fig. 2, Table 4). These large coefficients of variance caused the largest standard error value in the Rh/Rs assessment (54 ±41 %, Table 6). The uncertainty in the regression fit was likely caused in large part by the older roots which are bulkier but respire less than fine and young roots (Behera et al., 1990). However, this method had the closest average Rh/Rs to the 8 to the 8 to the 70. Previous studies also highlighted large variation of CO₂ efflux and root biomass which causes relatively low coefficient of determinations (Behera et al., 1990; Farmer, 2013). In accordance to Kuzyakov (2005), this method was comparatively simple (Table 7).

4.4 Lab incubation method

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that, with additional field and lab methods development it would be possible to further segregate Rh assessments into percentage of rhizomicrobial respiration, decomposition of plant residues and basal decomposition of SOM. Overall, the lab incubation technique was slightly more complex than the non-isotopic field Rh assessment methods but allowed a prompt determination of Rh whilst simulating year round field environment (Table 7).

The analysis of the five different Rh/Rs partitioning methods examined in this study shows that none of them was fully

satisfactory. That is, each technique had strengths and weaknesses (Table $\frac{76}{10}$). Using δ^{13} C-CO₂ is acknowledged as the

prevent ambient δ^{13} CO₂ contamination) makes that technique more complex than the other methods to assess Rh%.

Third, the ability to perform this technique in remote areas is limited because the δ^{13} C-CO₂ needs to be quickly

assessed with a calibrated and accurate spectrometer (Midwood et al., 2006). Fourth, in our study we found the large

variationnee in δ^{13} C-CO₂ of the respiration components (i.e. Ra, Rh and Rs) that impeded the assessment of Rh% per

individual collar. Accordingly, further studies should analyze the spatial relationships of δ^{13} C-CO₂ with soil properties

and root characteristics. As standalone, the δ^{13} C-CO₂ technique was unable to produce an assessment of soil CO₂

4.5 Comparison of methods and recommendations

preeminent way to segregate Rh/Rs (Cheng, 1996; Kuzyakov, 2006); and accordingly the relative accuracy of the other methods was defined by their difference with this method. However, Wwe recognize this was a precarious approach because the δ^{13} C-CO₂ method had a large variation. Fortunately, the root biomass regression, which is also recognized in the literature as a reliable method (Kuzyakov, 2006), gave a similar %Rh estimate. However, we found several other shortcomings with the δ^{13} C-CO₂ method. First, the conjunction of field and lab procedures makes it difficult to complete this method in one day as needed. Second, the air flushing with CO₂ free gas in the field (to

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efflux; and thus needed to be performed in conjunction with field Rs measurements. In this regards, the δ^{13} C-CO₂ complemented well with the root exclusion bags methods because it allowed <u>us</u> to determine if the buried bags had teared and been invaded by roots and to standardized Rh% determination.

The root biomass regression method had the advantage of being to be simple, to and produced an average Rh% close to the δ^{13} C-CO₂ natural abundance. However it had and the disadvantage to of requiringe a high number of replicates due to low coefficient of determination between CO₂ efflux and root biomass. Another disadvantage of the root biomass regression technique is that in order to produce seasonal trends, the labor intensive procedures (i.e. pit digging, CO₂ measurements and root counting) need to be reinitiated several times during the years. This shortcoming can be particularly impractical in small plot experiments. Complementary studies should assess thresholds of root size to be

included in the regression fit in order to optimize the correlation fit and use the δ^{13} C-CO₂ natural abundance method to determine the effect of root size on the isotopic signature.

The root exclusion bags methods (i.e. HS and IB) had the advantage of being to be easy to monitor throughout the year. That is, because t, capturing temporal variability of the he-% of Rh is unlikely to be constant in time it is important to assess it periodically. The bags methods can be considered as a miniaturization of the traditional soil trenching method. However, contrasting with large trenches (e.g. Comeau et al., 2016; Fisher and Gosz, 1986) the root exclusion bags had the advantage of being to be simpler to establish and to allowed mycorrhiza development inside the mesh bags (Moyano et al., 2007). Conversely, due to the relatively small bag sizes, root webs on the outside edge could potentially contaminate Rh assessment. In this study, the δ^{13} C-CO₂ determination made with the collars located in the center of the bags showed no isotopic signature of root respiration. Similar to ally with the trenching method, the root exclusion bag methods had the disadvantages to of requirings several months of soil stabilization before starting-CO₂ efflux measurements could begin. Compared with the δ^{13} C-CO₂ natural abundance method, the HS and IB overestimated and underestimated %Rh by 18 and 12%, respectively. The divergences were likely caused by soil disturbances, alteration in root demise dynamic and lack of root exudates. Correspondingly, Carbone et al. (2016) found 11% differences in Rh% assessment between an isotopic partition method and the trenching technique. Comparing the HS and IB, the former created more soil disturbances but the latter would not be suitable for soil with a high amount of sand, gravel or rock because the intact blocks would collapse.

The lab incubations with of the minimally disturbed microcosms was the only method that had absolutely no influence of root or mycorrhiza. Thus the results from this method exclusively represented the CO_2 efflux originating from the mineralization of the slow turnover SOC pool (i.e. basal soil respiration) (Pell et al., 2006). Assessment of basal soil respiration in relationship with the total Rh is of great importance in evaluating the dynamic of the stabilized SOC. In this study, the Rh% from the lab incubation was 8-17% while the δ^{13} C- CO_2 natural abundance had an average of 61% Rh. Thus, if the soil incubation results were not affected by the height of the soil columns (as discussed above), basal respiration represented approximately one fifth of the Rh. Because stabilized SOC is a key indicator of soil quality and health (Creamer et al., 2014), further research should study the relationship between basal soil respiration and rhizosphere derived Rh. Also, future studies on soil CO_2 efflux segregation should include other partitioning techniques like the trenching method and the radiocarbon-based assessment (Chiti et al., 2015). Overall, results from field experiments exhibited a range of potential Rh between 2.5 and 6.0 Mg CO_2 -Cha $^{-1}$ y $^{-1}$. With the publication of the total

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annual live biomass growth (i.e. including root and above-grown biomass) at the study site (Tai Po Kau Nature Reserve) assessment of net ecosystem C balance will-would then be possible.

5 Conclusions

Methods for determining ecosystemC fluxes need to be improved and applied to allow a quantitative understanding of the biological processes underlying SOC balance. This study compared five methods to assess Rh and our results showed large variation need fin effluxes and Rh/Rs ratio between the different techniques analyzed. The data revealed that the hand-sorted root exclusion bags and the intact root exclusion bags methods produced similar Rh efflux values and these efflux were slightly lower than the one produced by the root biomass regression method but notably larger than the lab incubation with soil cores We found that methods with higher relative accuracy (soil δ^{13} C-CO₂ natural abundance and root biomass regression) had lower precision (i.e. large variance) and methods with higher precision (root exclusions bags and lab incubation) had lower relative accuracy. Based upon these results, we suggest that when assessing rate of heterotrophic emissions and their contribution to total soil based emissions, two or more methods should be performed to produce more integral assessments.

Acknowledgements

The work described in this paper was supported by two grants from the Research Grants Council of the Hong Kong Special Administrative Region, China (CUHK14302014 and CUHK14305515), as well as a Direct Grant of CUHK (SS16914). Thomas Lui is thanked for assistance with the gas chromatograph analysis. Jack Lin is thanked for the assistance in building the microcosm tubes and for ensuring that the incubation containers were fully airtight. Amelia Amadeo, Christoph Hartmann and Merve Oztoprak are thanked for the assistance during the field measurements. Dr. Barry Thomton from the James Hutton Institute Scotland UK is thanked for the δ ¹³C-CO₂ determination.

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Table 1: Morphological description of the soil profiles at the study site

Horizon depth (cm)	Color (dry)	Color (moist)	Field texture ^a	Structure ^b	Rock fragments volume %	Roots mg root cm ⁻³	pH (H ₂ O)	pH (KCl)	WHC (g H ₂ O g soil ⁻¹)	Soil organic C; N (%)
A 0~10	10 YR 6/2	10 YR 3/2	SL	gr	0	9.3 ± 3.4	4.2	3.3	0.50 ± 0.01	$3.2 \pm 0.2; 0.24 \pm 0.02$
AB 10~25	10 YR 6/3	10 YR 4/2	SL	gr	0	1.6 ± 0.6	4.5	3.9	-	-
Bt 25~60	10 YR 8/8	7.5 YR 6/8	CL	sbk	10-20	0.5 ± 0.2	4.7	4.0	-	-
C 60~100	10 YR 8/8	10 YR 7/8	CL	sbk	>60	0	4.7	4.0	-	-

^a SL, sandy loam; CL, clay loam.
^b gr, granular; sbk, subangular blocky.
WHC, water holding capacity
C, carbon; N, nitrogen

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Table 2: Comparison of environmental parameters inside and outside the root exclusion bags

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Method	Soil temperature	Soil moisture	Bulk density
	(°C)	(vol. %)	$(g cm^{-3})$
Inside hand-sorted root exclusion bags (HS)	$22.4(0.2)\alpha$	20.5 (1.2) β	$1.16(0.04)\alpha$
Inside intact root exclusion bags (IB)	$22.6(0.3)\alpha$	$25.5(1.4)\alpha$	$1.13(0.05)\alpha$
Outside root exclusion bags (Rs)	$22.4(0.2)\alpha$	$24.8(0.8)\alpha$	$1.14(0.03)\alpha$

Values are means and standard error. Values in the same column followed by a different Greek letter (α, β) are significantly different from each other at α =0.05.

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Table 3: Linear regression report between root density and CO efflux

Parameter	Value	SE*	t value	P value
Intercept (g CO ₂ m h)	0.25	0.10	2.50	0.02
Slope (CO₂ X mg root cm⁴)	0.08	0.04	2.31	0.03

Overall r² of the linear regression: 0.21.

*root density in unit of milligram, small (radius between 0.1, 0.5 cm) dried roots (60°C) per cm² of soil.

SE standard error

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Table 34: Parameter values of the Gaussian 3D regression fitted curve (equation 1)

Efflux	Parameter a $(\frac{\text{Mg ha}^{-1} \text{ y}^{-1} \text{g}}{\text{m}^{2} \text{h}^{-1}})$	Parameter xo	Parameter yo	Parameter b	Parameter c
Rh incubation	5.0 0.21 ***	49.2 ***	34.7 ***	15.7 ***	19.2 ***
Rs field	10.3 <mark>0.43</mark> **	24.9 **	18.3 NS	9.6 NS	15.8 NS
Rh IB	5.7 0.24 NS	21.93 NS	21.4 NS	4.8 NS	13.4 NS
Rh HS	8.1 <mark>0.34</mark> NS	21.7 NS	9.5 NS	5.1 NS	14.2 NS

 $\ensuremath{\mathsf{Rh}}$ incubation, heterotrophic respiration from the soil cores incubation

Rs field, total soil respiration from outside of the root exclusion bags

Rh IB, heterotrophic respiration from the intact root exclusion bags

Rh HS, heterotrophic respiration from the hand-sorted root exclusion bags

** and *** significant at p < 0.01 and p < 0.05, respectively; NS non-significant.

Parameters from equation 1. Parameter "a" correspond to the height of the maximum high of the curve (g CO₂ m⁻² h⁻¹); "xo" is the peak soil temperature point (°C) in the curve, "yo" is the peak soil moisture (%) point in the curve, and "b" and "c" are the Gaussian root mean squared widths of the soil temperature and soil moisture, respectively.

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Table <u>45</u>: <u>Average</u> δ^{13} C-CO₂ results

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Method	$\delta^{13}\text{C-CO}_2$
	(‰)
Rs ^a	-18.21 (0.53) αβ
Rh HS ^b	-16.65 (0.44) β
Rh IB ^c	-16.52 (1.07) β
Rh lab ^d	-16.75 (0.54) β
Ra lab ^e	$-20.44(0.65)\alpha$

^a Rs, gas samples collected from the field total soil respiration collars.

Values followed by a different Greek letter (α, β) are significantly different from each other at α =0.05.

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Table <u>56</u>: Comparison of heterotrophic respiration assessment methods

•	Method	Rh efflux ^a	Rs efflux ^b	Rh / Rs
•		Mg CO ₂ -	-C ha ⁻¹ y ⁻¹	%
l	Root biomass regression	6.0(2.4)	11.1 (1.0)	54 (41)
•	Soil cores incubation	$0.4 - 1.9^{c}$	-	8-17 ^d
	Hand-sorted root exclusion bags (HS)	4.8(0.3)	6.1 (0.3)	79(3)
	Intact root exclusion bags (IB)	3.0(0.3)	6.1 (0.3)	49 (7)
	Soil δ ¹³ C-CO ₂ natural abundance	-	-	61 (39)

Values are means and standard error, n = 22 for the root biomass regression, n = 47 for soil incubation,

^b Rh HS, gas samples collected from the field hand-sorted root exclusion bags collars.

^c Rh IB gas samples collected from the field intact blocks root exclusion bags collars.

^d Rh lab, gas samples collected from lab incubations of soil with freshly removed roots.

^e Ra lab, gas samples collected from lab incubations of the roots extracted in Rh lab.

Values are means and standard error, n = 14 for Rs and Ra and n = 7 for HS, IB and Rh lab.

n = 28 for both root exclusion bags techniques.

^a Rh, heterotrophic respiration.

^b Rs, total soil efflux taken alongside the Rh efflux.

^c Efflux range at temperature between 14°C and 26°C.

d Calculated as Rh from incubation at 14°C and 26°C divided by average field Rs at 14°C and 26°C respectively.

Table 67: Comparison of partition methods Qualitative evaluation of the partition methods

G d d d d	D. L. C	D D	C 1 C	m: 1.1.	TD: 1 1 1
Segregation method	<u>Relative</u>	Precision ^b	Complexity of	Time needed to setup	Time needed to produce
	Aaccuracy a		procedures ^c	experiment ^d	seasonal trends
Root biomass regression	High	Low	Simple	2-3 days	6 months to 1 year
Soil cores incubation	Low	High	Complex	5-7 days	<1 to 2 months
Hand-sorted root exclusion bags (HS)	Intermediate	High	Simple	2-3 days	6 months to 1 year
Intact root exclusion bags (IB)	Intermediate	High	Simple	2-3 days	6 months to 1 year
Soil δ ¹³ C-CO ₂ natural abundance	High	Low	Complex	1-2 days	6 months to 1 year

^a Partition methods that had Rh%: <10, 10-20 and >20 lower or larger than the δ^{13} C-CO₂ natural abundance were categorized as high, intermediate and low accuracy, respectively.

b High, intermediate and low precision were attributed to Rh% standard errors of <10, 10-20 and >20, respectively.

c Methods with five steps or less were deemed simple and six steps or more deemed as more complex.
d The time needed to setup experiment was assessed with the number of working hours required prior to be able to start the measurements.

^e The time needed to produce seasonal trends was the number of months of measurements required to characterize the Rh at the different temperature and moisture levels of the

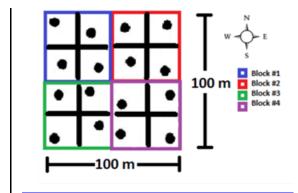


Figure 1. Field sampling for the lab incubations, stratified random design in the 1 ha study area.

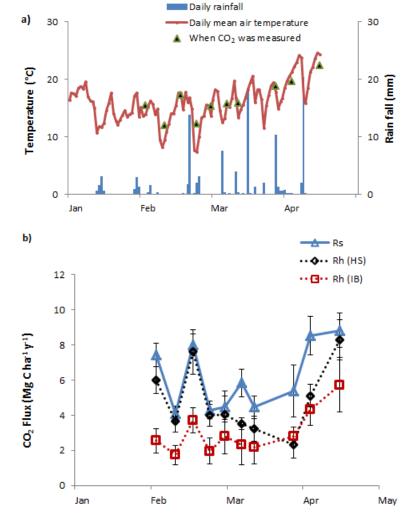


Figure 1: a) Soil and air temperature and daily rainfall over the study period; b) Total soil CO_2 efflux (Rs), heterotrophic CO_2 efflux (Rh) from hand-sorted root exclusion bags (HS), and Rh from intact block root exclusion bags (IB).

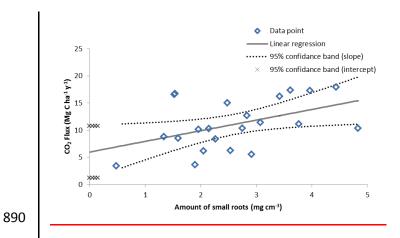
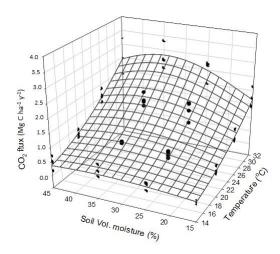


Figure 2: Linear regression between root $\frac{quantity biomass}{quantity}$ and CO_2 efflux.



 $Figure \ 3: Results \ from \ the \ lab \ incubation; regression \ between \ incubation \ temperature, moisture \ and \ CO_2 \ efflux.$