

Dear Editor and Reviewers,

Thank you for giving chance for our manuscript with this revision. We appreciate your support and your valuable comments and hope that our manuscript improved well and could reach the standards of BG.

The main changes in the revised version (highlighted by red in the manuscript) are the following:

1. First, since the main concern of the reviewers was the lack of the spatial replication in component measurements, we added a supplementary material to the manuscript dealing with the experimental setup and position changes of the soil chambers. There were replications both for root-exclusion and for root- and mycorrhiza-exclusion and the positions of the chambers were changed in every 2-3 weeks during the study. The figures of the supplementary material (Fig S1-4) represent the schematic map of the experimental area, the position changes of the soil chambers during the study period, the air-flow diagram of the measuring system and the regressions used for estimation of component's isotopic signals.
2. Contributions of the different soil respiration components were calculated (mixing models applied for daily measurements data) for the different days and averaged for the study periods in the former version. Now we removed these daily estimations due to the lack of spatial replication on a single day. The contributions of the components are now estimated by the mixing models for the different periods (30-50 days), therefore at least 2-4 spatial replications were used for component estimations (Fig. 1: position changes indicated by arrows, P5 L17-31, P8 L12-14, Supplementary material Fig. S2).
3. Description of isotopic measurements has been rewritten (P7-8).
4. Other results (cross-correlation between NEE and component contributions) and texts related to the daily estimations of the components were also omitted from the manuscript due to the lack of the spatial replication on a single day.
5. Some results have changed due to the changes in the estimations of the component contributions and the different error propagation (e.g. P12 L19-20, L29-31, P13 L4-15), but the main results- effect of drought on component's CO₂ efflux - remained the same.
6. The text is re-structured and several parts of the introduction and the methods (mainly the introduction concerning P3 L6-11, L14-17, P4 L11-16, L24-30) have been rewritten. The results and discussion sections have also been re-structured according to the comments of Reviewer#2 (the most important changes being the order of the sections and figures) and several parts of them have been rewritten (P13-14, P15 L28-

32). Sections 4.1 and 4.3 have not merged (as it was suggested) to keep the results and discussions comprehensible.

7. We also tried to clarify the terminology following Moyano et al. (2009) and added a few lines to 2.4 section where components were described (P9 L9-15, L23-33).
8. Instead of bootstrapping, gaussian error propagation method was used to calculate uncertainty, as the Editor proposed (P11 L9-12). The propagated uncertainties have increased by a few% due to the different method, but the ratios and tendencies remained roughly similar to the former version. (see Results).

The authors would like to thank the anonymous reviewers for their valuable remarks, and the Editor for her constructive comments, which we hope helped to increase the overall quality of the manuscript.

Gödöllő, 2016.04.14.

Sincerely Yours
On behalf of the authors
János Balogh

Autotrophic component of soil respiration is repressed by drought more than the heterotrophic one in a dry grassland

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Abstract

Summer droughts projected to increase in Central Europe due to climate change strongly influence the carbon cycle of ecosystems. Persistent respiration activities during drought periods are responsible for a significant carbon loss, which may turn the ecosystem from sink to source of carbon. There are still gaps in our knowledge regarding the characteristic changes taking place in the respiration of the different components of the ecosystem respiration in response to drought events.

Here, we combined a physical separation of soil respiration components with continuous measurements of soil CO₂ efflux and its isotopic (¹³C) signal at a dry grassland site in Hungary. The physical separation of soil respiration components was achieved by the use of inox meshes and tubes inserted into the soil. The root-excluded and root- and mycorrhiza excluded treatments served to measure the isotopic signal of the rhizospheric, mycorrhizal fungi and heterotrophic components, respectively.

In the dry grassland investigated in this study the three components of the soil CO₂ efflux decreased at different rates under drought conditions. During drought the contribution made by the heterotrophic components was the highest (54±8%; mean±SE). Rhizospheric component was the most sensitive to soil drying with its relative contribution to the total soil respiration dropping from 66±7% (non-stressed) to 35±17% (mean±SE) under drought conditions. According to our results, the heterotrophic component of soil respiration is the major contributor to the respiration activities during drought events in this dry grassland ecosystem.

1 Introduction

Grassland ecosystems strongly respond to drought events via substantial reduction of primary production (GPP, Hoover et al., 2014; Parton et al., 2012; Reichstein et al., 2013). In contrast, below-ground respiration is not as strongly affected (van der Molen et al., 2011; Yang and Zhou, 2013), but tends to be reduced as well under drought (Balogh et al., 2011; Suseela and Dukes, 2013). **Soil respiration is the second largest component of carbon cycling in grasslands and returns 50–90% of annual GPP back to the atmosphere (Bahn et al., 2008). Thus, the magnitude of soil respiration can turn the carbon budget from a net sink to a net source in dry years (Nagy et al., 2007). Here we address the question whether under drought this is primarily a function of autotrophic respiration which declines as GPP declines, whereas heterotrophic respiration remains largely unaffected.**

According to climate change scenarios the frequency of droughts is expected to increase in Central Europe (Prudhomme et al., 2014), where dry grassland ecosystems represent one of the major land use types. **It is well known that there is a need for better mechanistic models to address the effects of climatic extremes on carbon fluxes (e.g., Blagodatsky and Smith, 2012), but progress has been limited due to the high complexity of responses of different ecosystem respiration components to the climatic drivers.**

Soil organic matter (SOM) and litter derived respiration is considered to belong to the heterotrophic soil respiration component (Moyano et al., 2009). This decomposition is attributed mainly to soil bacteria and fungi and have an about 50% share in the total soil respiration in dry grasslands (Bao et al., 2010; Chen et al., 2009; Gomez-Casanovas et al., 2012). On the other hand, some of the soil fungi using recent photosynthetic assimilates are contributing to the autotrophic respiration component. Arbuscular mycorrhizal fungi (AMF) are obligatory symbiont soil fungi, forming intimate mutualistic associations in 70-90% of the plant species. **About 10-20% of the assimilated C may be allocated to AMF in exchange of acquisition of water and essential nutrients for plant productivity (van der Heijden et al., 2015).** Therefore soil respiration includes components of an autotrophic-heterotrophic continuum from roots through the root-associated (rhizospheric and mycorrhizal) to non-root-associated (heterotrophic) microbial components.

Belowground CO₂ production by the autotrophic and heterotrophic components show large diel and seasonal variability (Fassbinder et al., 2011; Moyes et al., 2010). The drivers behind all this are not fully revealed and the role of soil microbes is still not clear mainly due to the

diversity of soil biota (Bardgett et al., 2008). Moreover, drivers of CO₂ production frequently interact with each other (Balogh et al., 2015; Vargas et al., 2010), hampering the partitioning of the total CO₂ efflux into components. Studies found a stronger effect of photosynthesis than that of temperature on root respiration (Gomez-Casanovas et al., 2012; Heinemeyer et al., 2012; Hopkins et al., 2013). Both autotrophic and heterotrophic components were shown to be sensitive to water shortage (Carbone et al., 2011; Moyano et al., 2013). Autotrophic component was found to be dominant over the heterotrophic one during drought periods in a Mediterranean woodland ecosystem (Casals et al., 2011), but we have limited information from grasslands of shallow rooted herb species regarding the dominant source of carbon during drought periods.

The widely used separation techniques (trenching and girdling) are not considered suitable for grasslands (Epron, 2009), thus the physical separation of the soil CO₂ efflux components via root exclusion is hardly feasible without serious disturbance of soil structure and the root system. A viable option, however, is the use of stable isotopic signatures ($\delta^{13}\text{C}$) of soil respiration for estimating the relative contributions of the main components (Carbone et al., 2011; Hopkins et al., 2013). Although diel patterns in $\delta^{13}\text{C}$ may also be subject to biases in measuring methods (Fassbinder et al., 2011; Midwood and Millard, 2011), seasonal changes are expected to reflect the changes in the contributions of source components rather than the changes in the isotopic signal of the component itself (Knohl et al., 2005). However, SOM $\delta^{13}\text{C}$ can also change during the year: fresh plant material is more depleted in ^{13}C than the older SOM components (Bowling et al., 2002), therefore fresh litter may contribute to the decreasing $\delta^{13}\text{C}$ of the heterotrophic component. Drying of the surface layers can also modify $\delta^{13}\text{C}$ of the heterotrophic component, since heterotrophic respiration could be restricted to the deeper layers of the soil (Moyes et al., 2010). Soil drying can also change the amount of CO₂ produced in the topsoil layer (Balogh et al., 2015) by allowing greater atmospheric invasion, therefore enriched $\delta^{13}\text{C}$ in soil air (Phillips and Nickerson, 2010). The disequilibrium between the measured isotopic composition and the isotopic composition of the respiratory source could be significant, especially in tracer experiments (Gamnitzer et al., 2011), but it is assumed to be less pronounced in open chamber measurements due to the steady-state diffusion (Nickerson et al., 2013).

Uncertainties in estimates about the contributions of soil respiration components could be reduced by a combination of different methodologies (Risk et al., 2012). The question we are asking is: which of the investigated soil respiration components (autotrophic - including

rhizospheric and mycorrhizal fungi - and heterotrophic components) of a dry grassland dominates during drought? Our hypothesis is that autotrophic respiration would be reduced linearly with photosynthesis, whereas heterotrophic respiration might not be affected as strongly, resulting in net C loss from the soil carbon. To achieve our goals, we used an experimental setup of physical separation of soil respiration components combined with measurements of soil CO₂ efflux and its isotopic (¹³C) signal.

2 Methods

2.1 Site description

The vegetation at the Bugac site (46.69° N, 19.6° E, 114 m above sea level) is a dry sandy grassland dominated by *Festuca pseudovina*, *Carex stenophylla* and *Cynodon dactylon* and it has been under extensive management (grazing) for the last 20 years. Ten-year mean annual precipitation (2004-2013) was 575 mm, and the mean annual temperature reached 10.4 °C. The soil is a chernozem type sandy soil with high organic carbon content (Balogh et al., 2015).

2.2 Spatial separation of soil CO₂ efflux components

In 2010, ten soil cores (160 mm in diameter and 800 mm in depth, one of them 600 mm in diameter) were excavated. The roots were removed and the root-free soil was packed back - layer by layer - into PVC tubes with the same dimensions. Four tubes were used to exclude both roots and mycorrhiza. Walls of another 6 tubes were partially removed and replaced by inox mesh (40 µm mesh size) to exclude roots, while ensuring that the mycorrhiza filaments can grow into the tubes (Moyano et al., 2007). These root-free and root- and mycorrhiza-free soil cores were settled at a distance of 6 m from the eddy covariance tower to the south direction (Fig. 1.). The distance between the soil cores/tubes was 50 cm.

Soil CO₂ efflux and its isotopic signal were measured in plots:

- with undisturbed soil (various positions, 36 positions in total within a ~4 m² plot): total soil respiration, R_{soil} , $\delta^{13}\text{C}_{\text{Rsoil}}$,
- without roots and arbuscular mycorrhizal fungi (4 spatial replications) = heterotrophic component only, R_{rme} , $\delta^{13}\text{C}_{\text{Rme}}$,
- with root-excluded soil (6 spatial replications) = without roots, but with arbuscular mycorrhizal fungi, R_{re} , $\delta^{13}\text{C}_{\text{Re}}$.

2.3 Gas exchange measuring systems

Three different gas exchange systems were used in this study: eddy-covariance system (EC), automated soil respiration measuring system (SRS), connected to an isotopic CO₂ analyser (cavity ring-down spectroscopy system, CRDS). The experimental area was in the EC footprint (Supplementary material Fig. S1), but the size of the EC flux footprint area was larger by several orders of magnitude than the area covered by the SRS. Care was taken during the establishment of the experiment to select a plot with the same average soil characteristics and vegetation cover as found in the EC footprint area. Hence, the NEE and ET estimates obtained in this way can be considered representative also for the small-scale SRS and isotope measurements.

Data from 15th May 2013 to 12th November 2013 (182 days) were analysed in this study.

2.3.1 Eddy covariance setup

The EC system at the Bugac site has been measuring the CO₂ and H₂O fluxes continuously since 2002. In dry years, this grassland can turn into a net carbon source (Nagy et al., 2007), but the longer-term annual sums of net ecosystem exchange (NEE) show it to be a net sink, ranging from -171 to +106 g C m⁻² yr⁻¹ (Pintér et al., 2010) with a -100 g C m⁻² yr⁻¹ average. The EC system consists of a CSAT3 sonic anemometer (Campbell Scientific, USA) and a Li-7500 (Licor Inc, USA) open-path infra-red gas analyser (IRGA), both connected to a CR5000 data logger (Campbell Scientific, USA) via an SDM (synchronous device for measurement) interface. Additional measurements used in this study were: air temperature and relative humidity (HMP35AC, Vaisala, Finland), precipitation (ARG 100 rain gauge, Campbell, UK), global radiation (dual pyranometer, Schenk, Austria), incoming and reflected photosynthetically active radiation (SKP215, Campbell, UK), volumetric soil moisture content (CS616, Campbell, UK) and soil temperature (105T, Campbell, UK). These measurements were performed as described in Nagy et al. (2007) and Pintér et al. (2010). Fluxes of sensible and latent heat and CO₂ were processed using an IDL program after Barcza et al. (2003) adopting the CarboEurope IP methodology. For a detailed description of data processing and gap-filling see Nagy et al. (2007) and Farkas et al. (2011).

2.3.2 Soil respiration system

The 10 chamber automated soil respiration system was set up in July 2011. The system is an open dynamic one, consisting of an SBA-4 infrared gas analyser (PPSystems, UK), pumps,

flow meters (D6F-01A1-110, Omron Co., Japan), electro-magnetic valves, and PVC/metal soil chambers. The chambers were 10.4 cm high with a diameter of 5 cm, covering a soil surface area of 19.6 cm². The flow rate through the chambers was 300 ml min⁻¹, replacing the air in the chamber in 40 seconds. The PVC chambers were enclosed in a white metal cylinder with 2 mm airspace in between to stabilize the chamber and to prevent warming by direct radiation. Four vent holes with a total area of 0.95 cm² were drilled in the top of the chambers. Vent holes also served to allow precipitation to drip into the chambers. The system causes minor disturbances in the soil structure and the spatial structure of the vegetation. It is applicable without cutting the leaves/shoots of the plants, so it is not disturbing transport processes taking place within the plant stems and roots. It is suitable for continuous, long-term unattended measurements of soil CO₂ efflux and has been used in previous experiments (Nagy et al. 2011). The soil respiration chambers contained no standing aboveground plant material. Other studies (Nickerson et al., 2013; Risk et al., 2011) also used this chamber size, arguing that these chambers can be placed between the plants in grasslands, while larger chambers might create a non-representative surface due to the cutting necessary for placing the chambers on the ground (Risk et al., 2011).

R_{soil} was measured by 6 SRS chambers.

R_{rme} was measured by 2 SRS chambers.

R_{re} was measured by 2 SRS chambers.

2.3.3 Isotopic (¹³CO₂) measurements

A Picarro G1101-i gas analyser (CRDS, Picarro Inc., CA, USA) was attached to the soil respiration system from May to November in 2013. This CRDS system measured the isotopic composition inside the chambers and in the reference air 10 cm above the surface. The SRS sequentially measured each of the 10 chambers for 3 minutes. Every second chamber was additionally probed for isotopic signature measurements by the CRDS (3 minutes), followed by reference air measurements for another 3 minutes. Thus, 5 chambers for isotopic measurements took 30 minutes to measure in a single cycle. The CRDS integration time was set at 10 seconds, thus the CRDS provided 18 measurement points per chamber per cycle. Although the system response of the CRDS was clearly slower than the response of the SRS, the 3-minute duration was long enough to obtain robust results. Since CRDS followed the 3-minute intervals of SRS measurements, no additional grace time had to be considered for the isotopic measurements.

Although this sampling scheme provides a very good temporal coverage (replication in time), it is not perfectly addressing spatial variability, and hence the position of each of the chambers was moved 11 times to randomly selected locations during the study period (i.e., every 2–3 weeks) to obtain sequential spatial replications for each plot type (undisturbed, root-excluded, root- and mycorrhizal fungi excluded; see Supplementary material Figs S1 and S2). In detail, $\delta^{13}\text{C}_{\text{Rsoil}}$ was measured by 3 chambers at 36 (3 chambers x 12 positions) randomly selected positions within the experimental area (undisturbed soil, Supplementary material Fig. S1). $\delta^{13}\text{C}_{\text{Rre}}$ was measured by 1 chamber which was moved among positions 1, 3, 5, 6, 8, 9 during the study period (Supplementary material Fig. S2). $\delta^{13}\text{C}_{\text{Rme}}$ was measured by 1 chamber which was moved among positions 2, 4, 7, 10 during the study period (Supplementary material Fig. S2).

Since contributions by the different soil CO₂ efflux components were estimated for five different periods within the study period distinguished by NEE, SWC values and isotopic signals (see Results), data for each estimation originated from 2-3 spatial replications.

2.4 Data processing and modelling

Data processing and statistical analysis were done in R (R Core Team 2014). Before calculating daily averages of $\delta^{13}\text{C}$ values, a filtering method was applied to each dataset. Out of each 180 s long measurement on a certain chamber, the first 70 s (to measure a steady state signal) and the last 20 s were cut and the remaining values were used for further calculations. As reference and chamber air were measured sequentially, reference values during chamber measurements were estimated by linear interpolation between the neighbouring reference sequences.

After the interpolation, $\delta^{13}\text{C}$ values of the soil CO₂ efflux were calculated using the isotopic mass balance approach in each plot:

$$\delta^{13}\text{C}_R = \frac{\delta^{13}\text{C}_{\text{out}} \times c_{\text{out}} - \delta^{13}\text{C}_{\text{in}} \times c_{\text{in}}}{c_{\text{out}} - c_{\text{in}}} \quad (1)$$

where $\delta^{13}\text{C}_{\text{out}}$ and $\delta^{13}\text{C}_{\text{in}}$ are the isotopic signature of the outgoing and incoming air of the chamber and c_{out} and c_{in} are the CO₂ concentration of the outgoing and incoming air of the chamber, respectively.

$$\delta^{13}\text{C} = \frac{R_{\text{sample}}}{R_{\text{standard}}} - 1 \quad (2)$$

and R stands for the $^{13}\text{C}:^{12}\text{C}$ isotope ratio of the sample and the international VPDB standard (0.011182), respectively.

Individual measurements were filtered out by using a moving-window procedure if the investigated value (at the window center) was outside the range of the mean \pm median absolute deviation of the values in a 10 days moving window. This filtering procedure left an overall data availability of 68-70%. Daily averages were calculated using the remaining data.

To determine the isotopic signature of the ecosystem respiration (R_{eco}), Keeling plots were constructed by plotting the night-time $\delta^{13}C$ values measured 10 cm over the surface against the inverse of the CO_2 concentration. The extrapolated y-intercept of the linear regression was used as $\delta^{13}C_{Reco}$ values.

Total soil CO_2 efflux was separated isotopically into its components. We defined the components following the terminology presented by Moyano et al. (2009):

Heterotrophic respiration= microbial respiration from litter and SOM decomposition.

Autotrophic respiration= mycorrhizospheric respiration including rhizospheric and mycorrhizal fungi components.

Rhizospheric respiration= respiration of roots and root-associated microorganisms in the rhizosphere, not including mycorrhizal fungi.

Two-source mixing models were used to estimate the fraction (a) of the rhizospheric and (b) mycorrhizospheric components, based on the measured isotopic signals:

$$\delta^{13}C_{R_{soil}} = a \times \delta^{13}C_{R_{rhizo}} + (1 - a) \times \delta^{13}C_{R_{re}} \quad (3)$$

$$\delta^{13}C_{R_{soil}} = b \times \delta^{13}C_{R_{mycrhiz}} + (1 - b) \times \delta^{13}C_{R_{rme}} \quad (4)$$

where $\delta^{13}C_{R_{soil}}$ is the $\delta^{13}C$ of the total soil CO_2 efflux, $\delta^{13}C_{R_{re}}$ is the $\delta^{13}C$ of the root-excluded soil, $\delta^{13}C_{R_{rme}}$ is the $\delta^{13}C$ of the root- and mycorrhiza excluded soil (heterotrophic respiration), a is the fraction of the rhizospheric component (R_{rhizo}) and b is the fraction of the mycorrhizospheric component ($R_{mycrhiz}$) to the total soil efflux. According to these equations $1-b$ represents the ratio of heterotrophic respiration component to the total soil efflux and $b-a$ represents the ratio of mycorrhizal fungi component.

$\delta^{13}C_{R_{rhizo}}$ value was estimated by plotting $\delta^{13}C_{R_{soil}}$ values against the R_{re}/R_{soil} ratio (Supplementary material Fig. S3b). Since R_{re}/R_{soil} is hypothetically zero when only rhizospheric respiration is present, y-intercept of the linear regression was assumed as $\delta^{13}C_{R_{rhizo}}$. $\delta^{13}C_{R_{mycrhiz}}$ was estimated using the same approach (Supplementary material Fig. S3a), $\delta^{13}C_{R_{soil}}$ values were plotted against the R_{rme}/R_{soil} ratio and y-intercept of the linear regression was assumed as $\delta^{13}C_{R_{mycrhiz}}$. Similarly, $\delta^{13}C_{R_{re}}$ values were plotted against the R_{rme}/R_{re} ratio and y-intercept of the linear regression was assumed as $\delta^{13}C_{R_{myc}}$ (Supplementary material Fig. S4c), but this value was not used in further calculations.

Contributions of rhizospheric, mycorrhizal fungi and heterotrophic respirations to total soil respiration were calculated by the mixing models applied on subsets (periods) of the dataset of the total study period.

2.5 Microbial investigations

Soil samples for the microbial investigations were taken after the gas exchange measurements in May 2014 to avoid the disturbance of the measurements by the soil sampling. Sampling date was chosen considering the maximum of the carbon sequestration capacity of the investigated grassland (Nagy et al., 2007). 5-5 samples were taken from 5 soil layers (0-10 cm, 10-20 cm, 20-30 cm, 30-40 cm and 40-50 cm) in each plot.

Determination of AM fungal hyphal length in the soil was based on the methods of Bååth and Söderström (1979) using separation by wet-sieving and centrifugation. The separated fungal hyphae were stained using agar solution (0.75%) containing trypan blue (0.05%) then dried for 24 h at 70°C. The hyphal length was measured in the dried agar film by the intersection method (Tennant, 1975) under a binocular microscope.

The fluorescein diacetate (FDA) hydrolysis assay was used to estimate the total microbial activity in soil samples and expressed as mg fluorescein released kg⁻¹ dry soil (Adam and Duncan, 2001).

2.6 Uncertainty assessment

Isotopic signal of soil respired CO₂ has been studied extensively but several uncertainties related to the different methods have been revealed. Steady-state methods were found to provide more robust estimates than static chambers but still charged with biases (e.g. diffusive fractionation, Nickerson and Risk 2009). Open systems have the advantage of unattended automatic measurement collecting large amount of data but are less sensitive to small isotopic differences (Midwood and Millard, 2011).

In our study $\delta^{13}\text{C}_{\text{Reco}}$ estimates were independent of chamber related biases, using night-time $\delta^{13}\text{CO}_2$ and CO₂ concentration data of the free air over the surface for the calculation (Keeling-plot approach). This approach gave similar results to the chamber-based measurements, providing also partial verification of the latter ones. Moreover, isotopic measurements were independent on soil CO₂ efflux measurements, since IRGA and CRDS systems took different air samples from the same soil chambers. Isotopic data together with

CO₂ efflux rates were collected during 1980 measurement cycles on 182 days in order to have robust estimates of isotopic signals.

A C₄ grass (*Cynodon dactylon*) was also present in the study site potentially modifying the δ¹³C of the respired CO₂. Its cover was about 10% in the pasture (Koncz et al., 2014), but it was less frequent (i.e. less than 5%) in the experimental area. Calculated uncertainties of the relative contributions of each components (rhizospheric, mycorrhizal fungi and heterotrophic) contain the uncertainty due to a possible 5% contribution by the C₄ grass. **Isotopic signal of CO₂ efflux by the C₄ plant was supposed to be -14‰.**

In order to estimate the uncertainty of the measurements and estimated contributions by the different components to the total soil respiration, random errors of each factor (CO₂ concentrations, isotopic compositions, model fit errors and possible C₄ contribution) were propagated by Gaussian error propagation (Lo, 2005).

3 Results

3.1 Meteorological conditions, NEE, ET, soil CO₂ efflux, δ¹³C of CO₂ efflux

The end of May and the beginning of June was the most productive period in the year due to good water availability, the lowest NEE (strongest carbon sink activity) and highest evapotranspiration (ET) values were measured in this period (Fig. 1a). It rained only a few times from the end of June to 19th August (total precip: 10 mm) and the accompanying high temperature resulted in drought. Daily minimum NEE was around zero at the end of July and in August. Rain events after the drought period had significant effects on soil CO₂ effluxes (Fig. 1c). There was a second active period following autumn rains, but CO₂ uptake and ET were smaller than in May or June.

R_{soil} was the highest among the soil CO₂ effluxes, while R_{rme} was the lowest, the average CO₂ effluxes in the whole study period were 5.0±2.1, 3.8±1.6 and 2.6±1.2 μmol CO₂ m⁻² s⁻¹ (mean±SD) in R_{soil}, R_{re} and R_{rme}, respectively. R_{re} was sometimes higher than R_{soil}, especially shortly after rain events. **Lowest daily average total soil CO₂ efflux was measured in 15th August (2.22 μmol CO₂ m⁻² s⁻¹), while the lowest daily average R_{re} and R_{rme} values were observed in 2nd October (1.25 μmol CO₂ m⁻² s⁻¹) and 2nd November (1.04 μmol CO₂ m⁻² s⁻¹), respectively. Highest values of soil CO₂ effluxes were measured in May in all treatments (R_{soil}, R_{re} and R_{rme}). Sudden increases in R_{re} and R_{rme} were observed shortly after rain events, but R_{soil} showed slower (but more persistent) response to precipitation.**

Isotopic signature of R_{eco} was the lowest in May and June, increased in July and August and decreased again in October and November. $\delta^{13}\text{C}_{R_{\text{eco}}}$ showed clear responses to precipitation pulses: sudden declines were observed during the rain events. Chamber-based $\delta^{13}\text{C}_{R_{\text{soil}}}$ showed similar changes during the study period. $\delta^{13}\text{C}_{R_{\text{rme}}}$ and $\delta^{13}\text{C}_{R_{\text{re}}}$ showed large scatter during the whole study period with no clear and detectable trends. Differences between $\delta^{13}\text{C}_{R_{\text{soil}}}$ and $\delta^{13}\text{C}_{R_{\text{rme}}}$ were largest in the active period and smallest under drought conditions.

According to the NEE, SWC values and isotopic signals we distinguished 5 periods within the study period: an active period from 15th May to 20th June, a drying (stress development) period from 21st June to 22nd July, a drought period from 23rd July to 19th August, a wetting (stress release) period from 20th August to 16th September and a re-greening (recovery) period from 17th September to the end of the study period (11th November) (Fig. 1).

3.2 $\delta^{13}\text{C}$ of the respiration components

Fig 2. shows the measured and estimated $\delta^{13}\text{C}$ values of the different soil CO_2 efflux components. $\delta^{13}\text{C}_{R_{\text{rme}}}$ was the highest, while $\delta^{13}\text{C}_{R_{\text{eco}}}$ was the lowest, suggesting that rhizospheric respiration was the most substantially depleted, while heterotrophic respiration was the least depleted in ^{13}C . Mean of $\delta^{13}\text{C}_{R_{\text{eco}}}$, $\delta^{13}\text{C}_{R_{\text{soil}}}$, $\delta^{13}\text{C}_{R_{\text{re}}}$ and $\delta^{13}\text{C}_{R_{\text{rme}}}$ were $-27.9\pm 0.5\text{‰}$, $-26.8\pm 1.3\text{‰}$, $-26.4\pm 1.8\text{‰}$ and $-25.7\pm 2\text{‰}$ (mean \pm SE), respectively. The estimated isotopic signal of the respiration of mycorrhizospheric ($\delta^{13}\text{C}_{R_{\text{mycrhiz}}}$), rhizospheric ($\delta^{13}\text{C}_{R_{\text{rhizo}}}$) and mycorrhizal fungi components ($\delta^{13}\text{C}_{R_{\text{myc}}}$) were $-28.6\pm 1.6\text{‰}$, $-28.9\pm 1.7\text{‰}$ and $-27.2\pm 2.3\text{‰}$ (estimate \pm SE), respectively (Fig. 2).

36% of the variation in $\delta^{13}\text{C}_{R_{\text{soil}}}$ was explained by SWC ($\delta^{13}\text{C}_{R_{\text{soil}}} = -0.1267 \times \text{SWC} - 25.537$, $R^2=0.36$, $P<0.0001$), while only 3% of the variation of $\delta^{13}\text{C}_{R_{\text{rme}}}$ was explained by SWC and there was no correlation between $\delta^{13}\text{C}_{R_{\text{re}}}$ and SWC. Similar results were found between T_s and isotopic signals, but the correlation was weaker ($\delta^{13}\text{C}_{R_{\text{soil}}} = 0.1056 \times T_s - 28.588$, $R^2=0.11$, $P<0.0001$).

3.3 Ratio of the different components in total soil respiration during the vegetation period

Two end-member mixing models (eq. 3 and 4) were used to estimate the relative contributions of rhizospheric, mycorrhizal fungi and heterotrophic components to total soil respiration during the study period. The estimated contributions by the different components were $50\pm 6\%$, $13\pm 8\%$ and $37\pm 6\%$ (mean \pm SE) for the rhizospheric, mycorrhizal fungi and

heterotrophic components, respectively. The autotrophic component (mycorrhizospheric component) of soil respiration showed significant decrease during the drying and drought periods. Rhizospheric component was the most sensitive to drying and drought. Average contribution by the rhizospheric component to total soil CO₂ efflux decreased from 66±7% (mean±SE) in the active period to 35±13% during the drought period (Fig. 3). After drought, rhizospheric contribution increased again and become dominant during the re-greening period in autumn 63±7% (mean±SE). During the transient (drying and wetting) periods the rhizospheric contribution to the total soil CO₂ efflux was 38±11% and 46±8%, respectively. Relative mycorrhizal contribution was between 8-21% during the whole study period, with the highest contribution (21±11%; mean±SE) during the wetting period. Heterotrophic contribution to soil respiration was lowest in the active period (21±7%) and highest under drought (54±13%) (Fig. 3).

Changes in soil CO₂ effluxes showed similar responses to drying and drought conditions as isotopic signals. Average R_{soil} decreased by 60% (referenced to the average during the active period) as a response to drought, while R_{re} and R_{rme} showed declines of 56 and 52% respectively, suggesting declines in root respiration to be substantially larger than that in R_{soil} (60%).

3.4 Microbial biomass and activity

Hyphal length (on dry soil weight basis) was significantly lower in the upper layers of root- and mycorrhiza excluded soil than in undisturbed soil, while it was significantly higher in root-excluded plots at 10-20 cm depth. Hyphal length in the root-excluded soil was similar to undisturbed soil in the other soil layers. Fluorescein values were significantly lower in all soil layers in the root- and mycorrhiza excluded plots than in the undisturbed soil. Fluorescein values in the root-excluded plots were also lower than in undisturbed soil, but this difference was not significant (Fig. 4).

4 Discussion

4.1 Estimated contributions by the different components to the total soil CO₂ efflux and effect of drought on CO₂ effluxes and δ¹³C values

Ratio of the autotrophic component in total soil CO₂ efflux was 63±6% on average (rhizospheric and mycorrhizal fungi components, 50±6% and 13±8%, respectively) and it was

much higher than the ratio of the heterotrophic ($37\pm 6\%$) component, but the contributions by the different components showed significant changes during the growing season. In other studies, conducted in grassland ecosystems, the estimated yearly average ratio of the autotrophic component was found to be lower: 38-52% of the total soil respiration (Bao et al., 2010; Heinemeyer et al., 2012), while reaching 74% during the growing season in a prairie grassland (Gomez-Casanovas et al., 2012) and 60-74% in an arid perennial grassland (Carbone et al., 2008).

Soil CO_2 effluxes decreased in all treatments (R_{soil} , R_{re} , R_{rme}) under dry conditions, the largest decline was observed in total soil respiration (R_{soil}), therefore a strong response of the autotrophic component to drought could be assumed. The observed increase in $\delta^{13}\text{C}_{\text{Reco}}$ and $\delta^{13}\text{C}_{\text{Rsoil}}$ values during the drying period and during the drought also showed the decline of the autotrophic component. The same phenomenon was shown by the modelling results, with the smallest contribution by the rhizospheric component estimated for the drought period ($35\pm 13\%$; mean \pm SE), while the highest for the active period ($66\pm 7\%$; mean \pm SE). Fraction of the heterotrophic respiration were the highest during the drought ($54\pm 13\%$ mean \pm SE) and mycorrhizal fungi respiration showed only a small decrease during drought compared to the active period (from $13\pm 10\%$; to $11\pm 18\%$), suggesting that the non root-associated microbes and mycorrhizal filaments were less sensitive to water shortages than the rhizosphere. Soil aggregates are expected to provide micro-habitats for soil organisms that should be moist enough for those organisms to thrive even under drought (Davidson et al., 2012). Since there was an absence in plant photosynthetic supply during drought period, mycorrhizal fungi component is expected to use stored carbon for respiration (van der Heijden et al., 2008).

Low $\delta^{13}\text{C}_{\text{Rsoil}}$ and $\delta^{13}\text{C}_{\text{Reco}}$ were measured in the wetting and re-greening periods due to the drought-induced fall of the fresh litter to the surface as fresh plant material could be more depleted than the old litter (Bowling et al., 2002). The declines in $\delta^{13}\text{C}_{\text{Rsoil}}$ and $\delta^{13}\text{C}_{\text{Reco}}$ immediately after the rain events during drying and drought periods could also be explained by the wetting of the litter layer, exposing relatively fresh substrate to degradation for short periods. This phenomenon could also cause an overestimation in contributions made by the depleted components (rhizospheric) during rain events. Since the rhizospheric contribution estimated for the re-greening period was high, it is supposed that this result was obtained partly due to the increased amount of fresh litter. Similar results were found in a tallgrass prairie by Gomez-Casanovas et al. (2012), where the autotrophic components were more sensitive to soil drying than the heterotrophic ones. In contrary, Carbone et al. (2008) found

more sensitive response by the heterotrophic component in an arid (<150 mm annual precipitation) perennial grassland. Ratio of autotrophic components were reported to increase in response to drought in a woodland ecosystem, suggesting signature of the recent photosynthetic supply became enriched during drought and that could also explain the enrichment of the soil respired CO₂ (Casals et al., 2011). A drought induced increase in $\delta^{13}\text{C}$ of root respiration of trees was also assumed in a recent study (Risk et al., 2012), suggesting that the isotopic signal of the assimilates, thereby the signal of the autotrophic component might also increased. In our study, $R_{\text{rme}}/R_{\text{soil}}$ showed significant positive correlation with $\delta^{13}\text{C}_{R_{\text{soil}}}$ (the regression was used to estimate $\delta^{13}\text{C}_{R_{\text{rhizo}}}$, **Supplementary material Fig S4**), so $\delta^{13}\text{C}_{R_{\text{soil}}}$ was high if the ratio of heterotrophic CO₂ efflux to the total soil CO₂ efflux was found to be high. According to these studies and to our results we can assume that the different vegetation types may respond differently to drought: woodlands may increase the autotrophic contribution, while in grasslands it may decrease (Casals et al., 2011; Gomez-Casnovas et al., 2012; Risk et al., 2012). Since plants with different rooting habits have different water availability during dry periods (van der Molen et al., 2011), this could explain the differences in the response to drought by the different ecosystems.

4.2 Measured and estimated isotopic signals of the soil respiration components

Measured and calculated $\delta^{13}\text{C}$ values of the different respiration components showed differences similar to the ones as reviewed by Bowling et al. (2008). $\delta^{13}\text{C}_{\text{Reco}}$ (containing also the signal from above ground green biomass) was the most depleted, while $\delta^{13}\text{C}_{R_{\text{rme}}}$ (**heterotrophic components only**) was the least depleted. $\delta^{13}\text{C}$ of the root- and mycorrhiza excluded respiration was similar to SOM $\delta^{13}\text{C}$ measured in a previous study (Denef et al., 2013): -25‰ and -26‰ in the topsoil layers (without the litter layer). CO₂ effluxes from mycorrhizal fungi were expected to be more enriched in ¹³C relative to the total soil respiration (about +3‰, Bowling et al. 2008). Estimated $\delta^{13}\text{C}$ of mycorrhizal fungi component was **-27.2±2.3‰** (estimate±SE), which is 1.7‰ higher than the rhizospheric component (**-28.9±1.7‰**; estimate±SE).

In our study, neither $\delta^{13}\text{C}_{R_{\text{rme}}}$ values (heterotrophic respiration), nor $\delta^{13}\text{C}_{R_{\text{re}}}$ values (heterotrophic+mycorrhizal fungi respiration) showed correlation with SWC, but $\delta^{13}\text{C}_{R_{\text{soil}}}$ (total soil respiration) showed significant negative correlation with SWC. We can assume that $\delta^{13}\text{C}$ of heterotrophic respiration was not influenced by SWC changes during the growing season, as it was found also by other studies (Phillips and Nickerson, 2010; Risk et al., 2012).

Further, the lack of correlation by the present study also suggests that soil moisture induced changes in diffusivity (disequilibrium effect due to changing soil moisture) were not large enough to affect the measured $\delta^{13}\text{C}$ values.

4.3 Microbial investigations

High hyphal density was maintained in R_{re} plots and low, but still significant microbial activities (SOM decomposition) were detected in R_{rme} plots, therefore the measured $\delta^{13}\text{C}$ values characterized the sources of the root-free ($\delta^{13}\text{C}_{Rre}$) and root- and mycorrhiza-free ($\delta^{13}\text{C}_{Rrme}$) soils. The fact that very high amounts of hyphae were found in the root-excluded soil in the 10-20 cm layer proved that, mycorrhizal fungi filaments were able to penetrate through the inox mesh and supported significant microbial activity. Grasses have extensive fibrous root systems with moderate to high levels of mycorrhizal colonization (van der Heijden et al., 2015). The range of AM hyphal lengths found in this study (1.9–8.8 m g⁻¹ soil) are similar to those reported in the literature (e.g. Mummey and Rillig 2008). The higher hyphal densities found in root-free soil might have been related to the higher availability of SOM-derived nutrients and to more space without the roots (i.e. lack of competition). According to our results, significant amount of CO₂ was respired from mycorrhizal filaments in the undisturbed soil, having 12-31% share in the respiration by the autotrophic component.

Values of fluorescein in root-excluded plots were similar to those measured in the undisturbed soil, probably because of the fact that hyphae of AM fungi provide an increased area for interaction with other microorganisms (hyphosphere, Andrade et al. 1997), but were much lower in root- and mycorrhiza excluded soil. These results support the component estimations showing the significant activity of root-associated microorganisms.

5 Conclusions

In the dry grassland investigated in this study all three components of the soil CO₂ effluxes decreased, following different dynamics under drought conditions. Both measured CO₂ effluxes and isotopic signals showed similar results regarding component responses. The strongest decrease in response to drought was seen in rhizospheric respiration (relative contribution to total respiration decreased from 66±7% to 35±13%; mean±SE), while the relative contribution to the total soil respiration by the heterotrophic components increased during soil drying. During drought the contribution of the heterotrophic component was found to be the highest (54±8%; mean±SE). Mycorrhizal fungi respiration had its higher share in

soil respiration ($21 \pm 11\%$; mean \pm SE) in the wetting period after drought. According to these results, the carbon source activities identified by NEE measurements originated from carbon sources already stored, thereby decreasing the carbon content of the soil.

Drought events are expected to be more frequent in Central Europe in the future, and it is expected that the productivity of grassland ecosystems may strongly respond to projected dryness, influencing the carbon cycle of the ecosystems. Since potential productivity is generally linked to soil carbon content, a pronounced decrease in soil organic matter due to the enhanced activity of the heterotrophic component under drought may directly affect the long term productivity of grasslands.

Author contributions

J. Balogh, M. Papp, K. Pintér and Z. Nagy conceived and designed the experiment, M. Papp, K. Pintér and K. Posta performed the experiment, J. Balogh, Sz. Fóti, W. Eugster and Z. Nagy analyzed the data and wrote the paper, but all co-authors contributed to writing.

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Table 1 Measured and estimated CO₂ effluxes and isotopic signals in this study

	CO ₂ efflux	Isotopic signals
measured	$R_{\text{eco}}, R_{\text{soil}}, R_{\text{re}}, R_{\text{rme}}$	$\delta^{13}\text{C}_{\text{Reco}}, \delta^{13}\text{C}_{\text{Rsoil}}, \delta^{13}\text{C}_{\text{Rre}}, \delta^{13}\text{C}_{\text{Rrme}}$
estimated	$R_{\text{rhizo}}, R_{\text{myc}}, R_{\text{het}}$	$\delta^{13}\text{C}_{\text{Rmycrhiz}}, \delta^{13}\text{C}_{\text{Rrhizo}}, \delta^{13}\text{C}_{\text{Rmyc}}$

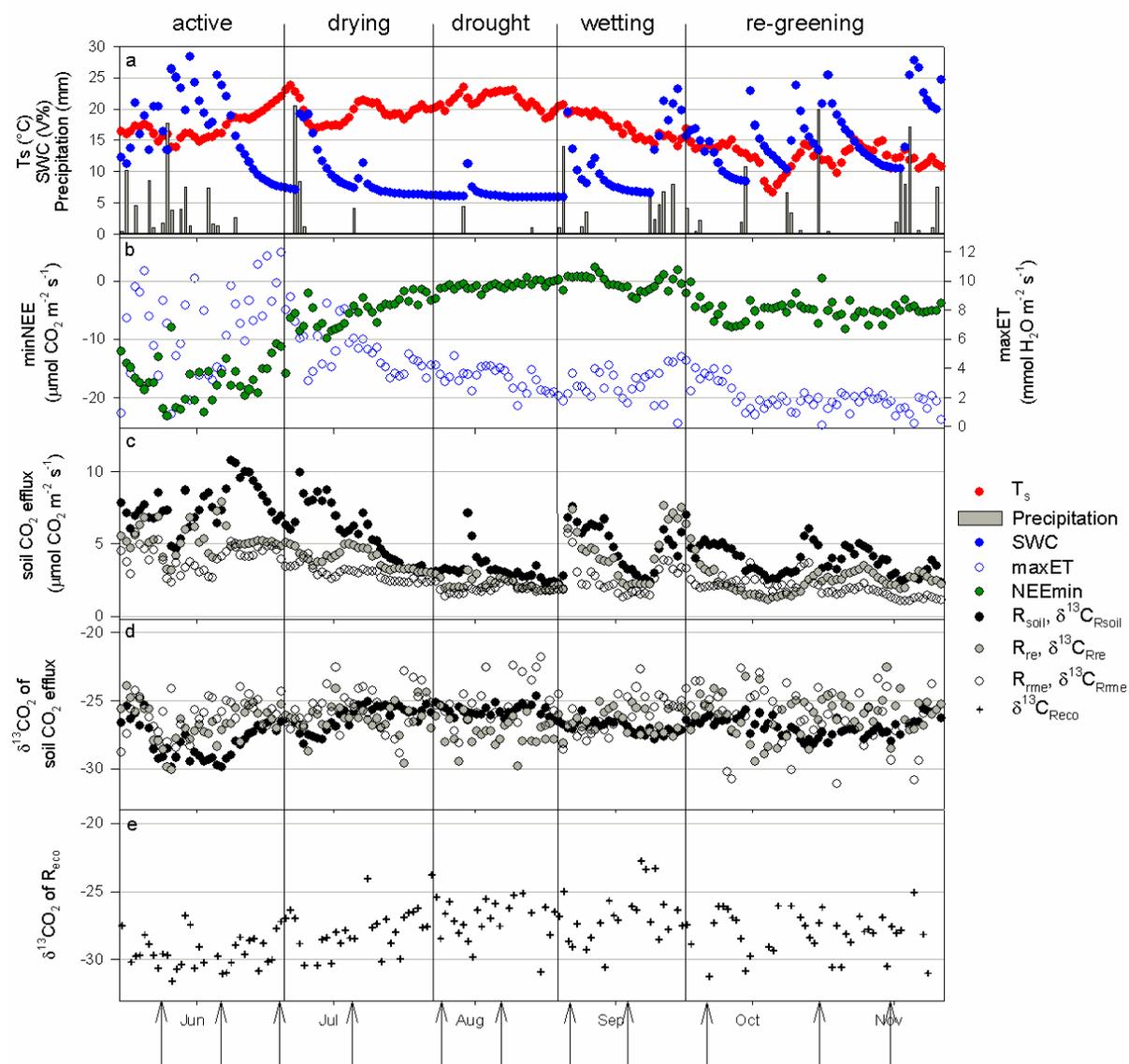


Fig. 1 (a) Daily averages of soil temperature (T_s), soil water content (SWC) at 5 cm depth and daily sum of precipitation, (b) daily minimum half-hourly NEE and maximum half-hourly ET, (c) daily averages of CO_2 efflux in undisturbed soil (R_{soil}), root-excluded soil (R_{re}) and root- and mycorrhizal fungi excluded soil (R_{rme}), (d) daily averages of $\delta^{13}\text{C}$ of soil CO_2 efflux in undisturbed soil ($\delta^{13}\text{C}_{R_{\text{soil}}}$), root-excluded soil ($\delta^{13}\text{C}_{R_{\text{re}}}$) and root- and mycorrhizal fungi excluded soil ($\delta^{13}\text{C}_{R_{\text{rme}}}$) and (e) daily averages of $\delta^{13}\text{C}$ of ecosystem respiration ($\delta^{13}\text{C}_{R_{\text{eco}}}$) during the study period in 2013, at Bugac site. **Arrows indicate the positions changes of the soil chambers.**

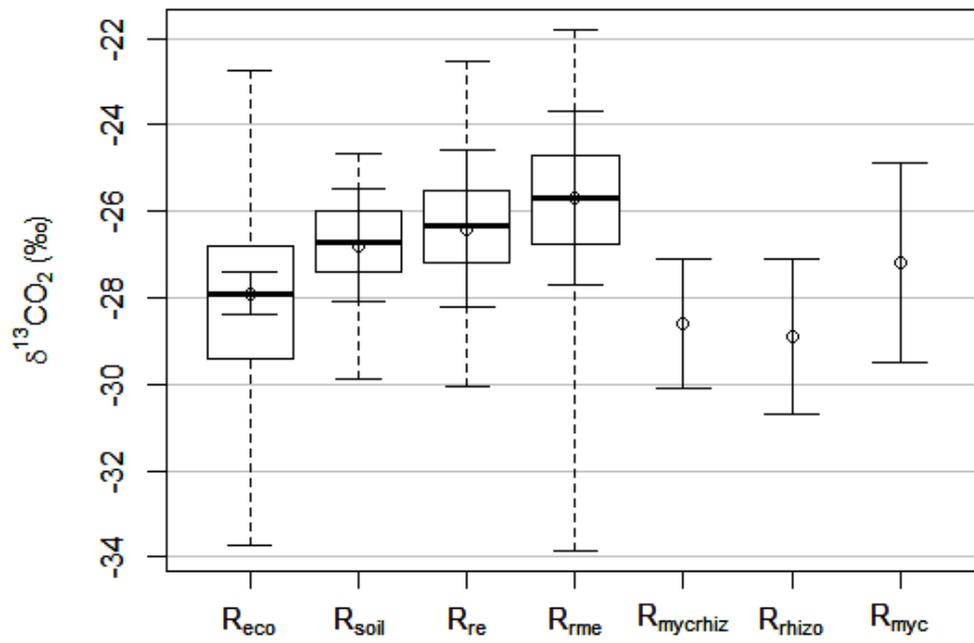


Fig. 2 Measured (R_{eco} , R_{soil} , R_{re} , R_{rme}) and estimated ($R_{mycrhiz}$, R_{rhizo} , R_{myc}) $\delta^{13}C$ values of the respiration components. Horizontal lines in boxes show medians and dashed whiskers show data extremes. Open circles and solid whiskers show means \pm propagated standard errors.

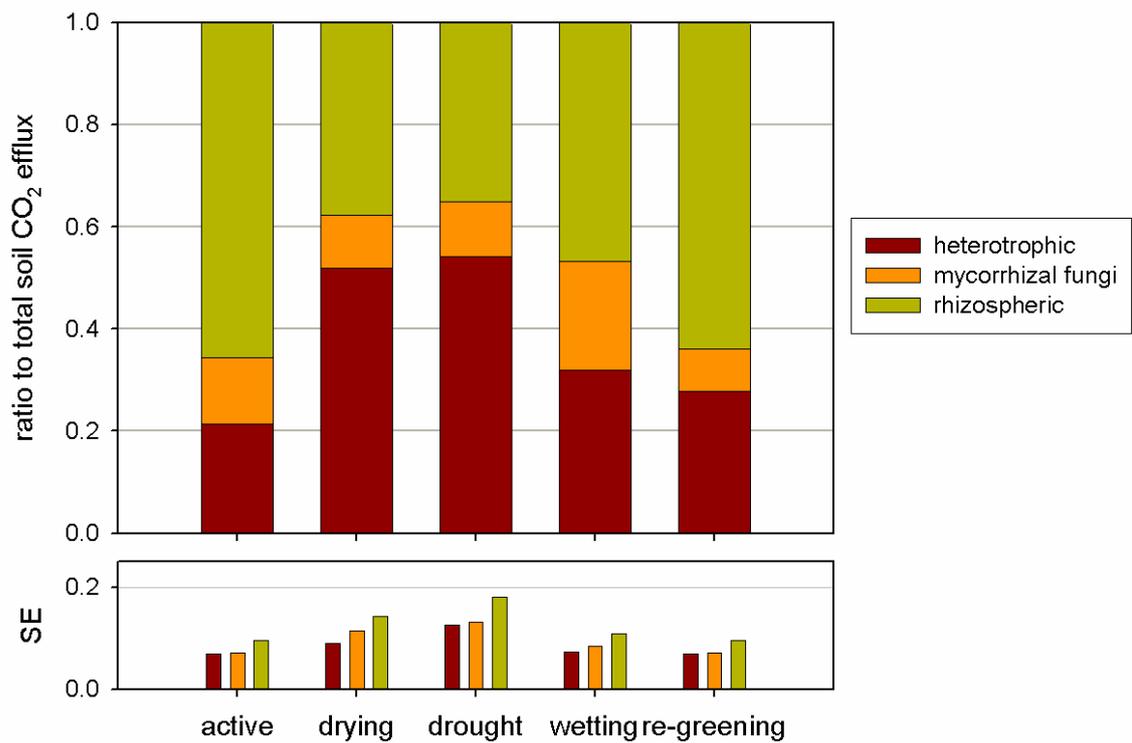


Fig. 3 Relative contributions made by rhizospheric, mycorrhizal fungi and heterotrophic components to the total soil respiration in the different parts of the vegetation period (15/05/2013-12/11/2013) at Bugac site. Propagated uncertainties of each estimate are shown in the lower panel.

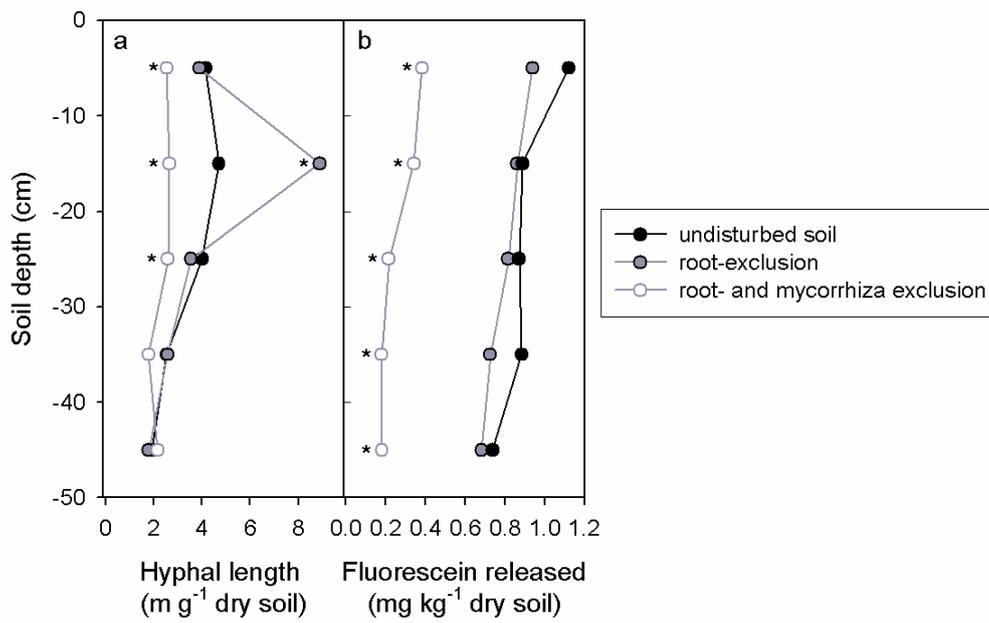


Fig. 4 (a) Mean hyphal length (m g⁻¹ dry soil) and (b) mean microbial activity expressed as fluorescein released (mg kg⁻¹ dry soil) in the undisturbed soil, root-exclusion and root- and mycorrhiza exclusion in different soil depths. Asterisks denote significant differences from undisturbed soil.