Inorganic carbon fluxes across the vadose zone of planted and unplanted soil mesocosms

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Abstract. The efflux of carbon dioxide (CO₂) from soils influences atmospheric CO₂ concentrations and thereby climate change. The partitioning of inorganic carbon (C) fluxes in the vadose zone between emission to the atmosphere and to the groundwater was investigated to reveal controlling underlying mechanisms. Carbon dioxide partial pressure in the soil gas (pCO₂), alkalinity, soil moisture and temperature were measured over depth and time in unplanted and planted (barley) mesocosms. The dissolved inorganic carbon (DIC) percolation flux was calculated from the pCO₂, alkalinity and the water flux at the mesocosm bottom. Carbon dioxide exchange between the soil surface and the atmosphere was measured at regular intervals. The soil diffusivity was determined from soil radon-222 (²²²Rn) emanation rates and soil air Rn concentration profiles and was used in conjunction with measured pCO₂ gradients to calculate the soil CO₂ production. Carbon dioxide fluxes were modeled using the HP1 module of the Hydrus 1-D software.

The average CO₂ effluxes to the atmosphere from unplanted and planted mesocosm ecosystems during 78 days of experiment were 0.1 ± 0.07 and 4.9 ± 0.07 µmol C m⁻² s⁻¹, respectively, and grossly exceeded the corresponding DIC percolation fluxes of 0.01 ± 0.004 and 0.06 ± 0.03 µmol C m⁻² s⁻¹. Plant biomass was high in the mesocosms as compared to a standard field situation. Post-harvest soil respiration (Rₛ) was only 10 % of the Rₛ during plant growth, while the post-harvest DIC percolation flux was more than one-third of the flux during growth. The Rₛ was controlled by production and diffusivity of CO₂ in the soil. The DIC percolation flux was largely controlled by the pCO₂ and the drainage flux due to low solution pH. Modeling suggested that increasing soil alkalinity during plant growth was due to nutrient buffering during root nitrate uptake.

1 Introduction

The global flux of carbon dioxide (CO₂) from the soil to the atmosphere amounts to 59–76.5 Gt carbon (C) yr⁻¹ and is one of the largest fluxes in the global C budget (Raich and Potter, 1995; Houghton, 2007). Agriculture strongly enhances this flux, accounting for 10–12 % of global anthropogenic emissions (Robertson et al., 2000; Barker et al., 2007; Vermeulen et al., 2012). The flux of CO₂ from the soil to the groundwater as dissolved inorganic carbon (DIC) is estimated at 0.2 Gt C yr⁻¹ and is hence much less than the upward flux of CO₂ (Kessler and Harvey, 2001). Agriculture enhances the DIC percolation flux by up to 4 times compared to undisturbed systems (Barnes and Raymond, 2009). Although numerous measurements have been made of either gas or aqueous phase CO₂ emission from cropland to

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the atmosphere and groundwater, respectively, few studies have investigated the total CO$_2$ emission with regard to its controls. In the light of the climate change induced by the present atmospheric concentration of CO$_2$ of 400 ppm and its increment rate of ∼2 ppm yr$^{-1}$ (IPCC, 2007; Lal, 2011), the magnitudes and underlying mechanisms of the soil CO$_2$ effluxes to the atmosphere and groundwater from agricultural systems are of crucial importance for prediction of the climate forcing. The residence time of DIC in groundwater is at least as long as the residence time of groundwater itself, which may be in the order of hundreds to thousands of years (Kessler and Harvey, 2001). The atmospheric residence time of CO$_2$ may be as low as 5 years (Solomon et al., 2007; Archer and Brovkin, 2008) implying that even small changes in the C emission balance can have important effects on the atmospheric CO$_2$ concentration (Schimel, 1995). The present study explores the total CO$_2$ emission for a cropland mesocosm system and investigates the underlying mechanisms.

The soil partial pressure of CO$_2$($p_{CO_2}$) and the soil CO$_2$ efflux to the atmosphere, also referred to as the soil respiration ($R_s$), are a function of the combined CO$_2$ production from microorganisms and roots, the soil gas diffusivity and a limited contribution from mineral reactions via the carbonate system (Kuzyakov, 2006; Trumbore, 2006). Soil temperature and moisture are the main abiotic factors controlling the biological production of CO$_2$ (Schlesinger, 1973; Maier et al., 2011). Further factors such as the overall soil nutrient content, soil mineralogy, land use history and plant phenology also play an important role for the magnitude of the soil CO$_2$ production (Lohila et al., 2003; Trumbore, 2006).

Diffusion of CO$_2$ in air is about 10$^4$ times faster than in water (Suarez and Šimůnek, 1993). Rain increases the $R_s$ due to a stimulation of microbial respiration and/or due to displacement of CO$_2$-rich soil air by advection (Huxman et al., 2004, Lee et al., 2002). However, frequent and heavy rains eventually result in a high soil water content that lowers the soil gas diffusivity, leading to accumulation of CO$_2$ in air-filled pores (i.e., higher $p_{CO_2}$) and a reduced $R_s$ (Jassal et al., 2005; Zhang et al., 2010).

Dissolved inorganic carbon is the sum of C in carbonic acid, H$_2$CO$_3$ (where H$_2$CO$_3$ = CO$_2$(aq) + H$_2$CO$_3$), bicarbonate, HCO$_3^-$, and carbonate, CO$_3^{2-}$. The concentration of DIC, [DIC], is closely linked to the $p_{CO_2}$ via Henry’s law. In addition, the [DIC] depends on soil solution chemistry because of the pH-dependent solubility of inorganic C species, and is as such largely influenced by processes that increase soil alkalinity, e.g., the weathering of carbonate and silicate minerals (Appelo and Postma, 2005; Walmsley et al., 2011). The DIC percolation flux to the groundwater can be described by multiplying the [DIC] by the recharge flux (Appelo and Postma, 2005; Thaysen et al., 2014a).

Here, we measured upward and downward CO$_2$ transport in both gas and aqueous phases in unplanted and planted mesocosms to quantify the total CO$_2$ emission from ecosystems. The mesocosms, established to simulate the top 0–80 cm soil profile of a barley cropland, were incubated under controlled environmental conditions, allowing a model-based investigation (HP1 module, Hydrus 1-D) of the biogeochemical controls on CO$_2$ production and transport in the soil profile prior to seeding, during growth and after harvest. Mesocosms have been shown to provide useful environments for conducting process-related research in unsaturated soil (Hendry et al., 2001; Thaysen et al., 2014a). Reactive transport modeling may further increase the understanding of the coupled physical, chemical, and biological processes influencing CO$_2$ transport within soils (Steefel et al., 2005). HP1 allows for the complex biogeochemical modeling of CO$_2$ transport in the vadose zone by providing options for simulation of soil water content, root growth, root water and solute uptake, as well as for gas diffusion and geochemical reactions of all possible chemical species, the latter being a novelty amongst vadose zone models.

2 Methodology

2.1 Setup of mesocosms

Emissions of CO$_2$ and DIC from unplanted and planted soil profiles may be directly compared in equally structured and textured soil maintained under controlled environmental conditions. In this study, seven mesocosms were constructed and incubated in a climate chamber. A detailed description of the experimental set-up and filling of mesocosms is given in Thaysen et al. (2014a). Soil was collected from the A and C horizon of an experimental field site located in an agricultural area (Voulund, Denmark, 56°25′35.7 N, 9°8′101.1 E) characterized as a coarse–sandy alluvial sediment (Podzol). The soil was sieved (6 mm) and packed into plexiglass cylinders (length: 85 cm, diameter: 19 cm) (Fig. 1a). The A and C horizons were located at 0–30 and 30–78.5 cm depth, respectively. The bottom plate of the mesocosms at 82–85 cm had an embedded suction disc (thickness of 10 mm) and the hydraulic connection between the C horizon and the suction disc was optimized by means of a thin layer of quartz flour (∼81.5–82 cm) and a layer of quartz flour mixed with C horizon (∼78.5–81.5 cm).

The mesocosms were subjected to different treatments (Table 1). Two mesocosms remained unplanted (referred to as the unplanted treatment). These mesocosms were shown to exhibit low variability and high reproducibility (Thaysen et al., 2014a). To investigate the additional variability introduced by the presence of plants, barley (Hordeum vulgare L. cv Anakin) was sown into three mesocosms. In these mesocosms CO$_2$ fluxes were investigated during growth (days 14 to 58 after sowing) and after harvest (days 58–117) (referred to as the barley growth treatment #1 and the post-harvest treatment, respectively). The agreement between mesocosms remained good (Figs. 3, S1 in the Supplement), also allowing
Table 1. Overview over mesocosm experiments. The post-harvest experiment was the successive of the barley growth experiment #1.

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Duration of experiment (days)</th>
<th>Fertilization</th>
</tr>
</thead>
<tbody>
<tr>
<td>Barley growth #1</td>
<td>58</td>
<td>Addition to soil prior to experiments</td>
</tr>
<tr>
<td>Barley growth #2</td>
<td>78</td>
<td>Nutrient irrigation</td>
</tr>
<tr>
<td>Post-harvest (post growth #1)</td>
<td>60</td>
<td>Addition to soil prior to experiments</td>
</tr>
<tr>
<td>Unplanted</td>
<td>78</td>
<td>Nutrient irrigation</td>
</tr>
</tbody>
</table>

for a reduction to a sample size of two for planted mesocosms. In further two mesocosms the growth period of barley was extended to 100 days with monitoring up to 78 days after sowing (barley growth treatment #2). In each planted mesocosm, 15 barley seeds were sown at 3 cm depth, and upon germination the seedlings were thinned to eight per mesocosm, corresponding to 280 plants m$^{-2}$.

Prior to packing of the mesocosms for the barley growth treatment #1, basal nutrients were mixed into the upper 9 cm of the A horizon in the following amounts (mg kg$^{-1}$ soil): $\text{NH}_4\text{NO}_3$, 30; $\text{K}_2\text{SO}_4$, 75; $\text{CaCl}_2$, 75; $\text{MgSO}_4 \times 7\text{H}_2\text{O}$, 45; $\text{MnSO}_4 \times \text{H}_2\text{O}$, 10.5; $\text{ZnSO}_4 \times 7\text{H}_2\text{O}$, 5.4; $\text{CuSO}_4 \times 5\text{H}_2\text{O}$, 2.1; $\text{Na}_2\text{MoO}_4 \times 7\text{H}_2\text{O}$, 0.2; $\text{CoSO}_4 \times 7\text{H}_2\text{O}$, 0.4. Mesocosms in the barley growth #1 and post-harvest treatment were irrigated with milli-Q water at 1–2 days intervals throughout the experiment. Mesocosms in the barley growth treatment #2 were irrigated in the same manner before day 56 and thereafter daily, however, irrigation was with milli-Q water prior to germination of barley and subsequently with a 50% strength Hoagland nutrient solution (Hoagland and Amon, 1950). The different mode of fertilizer application for the second barley growth treatment was implemented in order to avoid nutrient depletion of the soil during the longer experimental runtime. Unplanted mesocosms were initially irrigated using similar irrigation amounts as for planted mesocosms to compare the magnitude of soil respiration in unplanted and planted soil. Irrigation amounts were decreased after 24 days due to observed water logging at the mesocosm bottoms, but topsoil volumetric water content (VWC) was maintained at approximately the same level as in barley mesocosms (Fig. S1). Average irrigation amounts for the barley growth #1, the barley growth #2, the post-harvest and the unplanted treatments were 5.3, 13.5, 4.6 and 4.4 mm d$^{-1}$, respectively. Irrigation amounts exceeded those in the typical field situation because soil temperatures and plant biomass in mesocosms were elevated, leading to higher evapotranspiration (see Sect. 4.1). Maintenance of mesocosms was as in Thaysen et al. (2014a).

2.2 Sampling and calculations

Due to the limited amount of replicates in this study, we report data ranges instead of averages for all our core measurements. Ranges represent either the smallest and highest measured value for the barley growth #1 and the post-harvest treatment or both values for the barley growth #2 and the unplanted treatment.

2.2.1 Inorganic C speciation calculations and DIC percolation

Soil water alkalinity ($\sim$ sum of $[\text{HCO}_3^-]$ and $2 \times [\text{CO}_3^{2-}]$), soil $\rho\text{CO}_2$, soil temperature and moisture content were determined as function of depth and time, as described in Thaysen et al. (2014a). The speciation of inorganic C was calculated from the $\rho\text{CO}_2$, alkalinity and soil temperature using PHREEQC (Parkhurst and Appelo, 2013). The weekly DIC percolation flux in each mesocosm was estimated from the calculated [DIC] at the lowest sampling depth and the measured drainage flux. When low $\rho\text{CO}_2$ was measured at the...
mesocosm bottoms due to high water content (see Thaysen et al. 2014a), the upper next sampler was used to obtain the $p$CO$_2$ value.

2.2.2 CO$_2$ exchange

The exchange of CO$_2$ between the soil surface of the mesocosms and the atmosphere was measured using the static closed chamber technique (Ambus et al., 2007). A transparent cylindrical chamber ($V = 22.6\,L$) was carefully mounted gastight (Teflon sealant) on the mesocosm tops. Before deployment, the chamber was filled with air of ambient CO$_2$ concentration. During measurement, the $p$CO$_2$ in the chamber space was closed-loop analyzed continuously with an environmental gas monitor (EGM-2, PP-Systems, Amesbury, MA, USA) (Fig. 1b), and the CO$_2$ flux was estimated from the concentration change, volume and measurement time. Carbon dioxide exchange in barley mesocosms was measured under light and in the dark to quantify net ecosystem exchange (NEE) and ecosystem respiration (ER, $R_n$+ canopy respiration), respectively. The barley plant canopy occupied an area that was approx. 4 times the surface area of the mesocosm, which implied careful insertion of shoots into the measurement chamber (Fig. 1). For unplanted mesocosms, only $R_s$ was measured.

2.2.3 Radon profiles

Sedimentary produced radon-222 ($^{222}$Rn) was measured at different VWCs for unplanted mesocosms and mesocosms of the barley growth #1 treatment, in order to determine the soil diffusivity. Evacuated ZnS(Ag)-scintillation cells ($V = 200\,mL$) equipped with a manometer were attached to the mesocosm gas sampling ports. The scintillation cells were removed when their internal pressure had increased to 1013 hPa. Samples were analyzed on a scintillation counter (EDA-200, CAN). The background activity of Rn-222 inside the climate chamber was measured with an Alpha-guard PQ-2000 (Alphaguard, Genitron, DE) and was about 100 Bq m$^{-3}$. Radon emanation rates of each soil horizon were determined in the laboratory from five replicate soil aliquots of 200 g through incubation in a closed container for 2 months and subsequent scintillation counting of an air sample from the container. Experimentally determined Rn profiles were modeled with the RnMod3d software (Andersen, 2001) using measured VWCs and Rn emanation rates and assuming homogeneity within each soil horizon. Resulting bulk diffusivities were then used in the modeling of the CO$_2$ (Sect. 2.3) and to estimate the soil CO$_2$ production rate, $R_{CO_2}$ (µmol m$^{-2}$ s$^{-1}$) from Fick’s first law of diffusion (e.g., Fierer et al., 2005):

$$ R_{CO_2} = D_{soil} \cdot (C_T - C_B)/z, $$

(1)

where $D_{soil}$ (m$^2$/s$^{-1}$) is the bulk CO$_2$ diffusion coefficient in the A horizon soil, $C_T$ is the concentration of CO$_2$ at the mesocosm surface (µmol m$^{-3}$), $C_B$ is the concentration of CO$_2$ at the sampling depth of peak $p$CO$_2$ (µmol m$^{-3}$), and $z$ is the depth interval (m). The variable $D_{soil}$ was determined by dividing the CO$_2$ diffusion coefficient in air with the ratio between the Rn diffusion coefficient in air and the Rn bulk diffusion coefficient for the A horizon. Equation (1) assumes isobaric conditions, no downward flux of CO$_2$ beyond $C_B$, and CO$_2$ transport by gaseous diffusion only. It also neglects the effect of spatial differences in VWC.

2.2.4 Evapotranspiration

Weekly evapotranspiration was estimated from the difference between calculated and measured mesocosm weights. The calculated weight of a given mesocosm was obtained by subtracting the water removal due to effluent and sampling, from the sum of the mesocosm weight and the volume of irrigation water.

2.3 Modeling of CO$_2$ fluxes

In order to determine the controls on gaseous and dissolved CO$_2$ fluxes, results from barley growth experiments were modeled using the HP1 module of the Hydrus software (Šimůnek et al., 2006; Jacques et al., 2008). Due to the similarity between mesocosms (Figs. 3, S1), only simulation results of one mesocosm from the second barley growth experiment are presented herein. Besides the coupling between variably-saturated water flow, multicomponent transport, heat transport and biogeochemistry, particular features of the conceptual model are (i) CO$_2$ production accounting for respiration of both soil organisms and plant roots based on the SOILCO$_2$ model (Šimůnek and Suarez, 1993; Suarez and Šimůnek, 1993), linking explicitly soil respiration to root growth, (ii) cation exchange reactions and geochemical buffering by minerals, and (iii) root growth and root solute uptake.

2.3.1 Model setup

The mesocosm model contained three materials. The first two materials represented the A- and C horizons, respectively. The suction plate at the bottom of the mesocosm, the quartz flour layer and the layer of quartz flour mixed with the C horizon were lumped in the third material at 80–83 cm depth. The model domain was discretized in 157 nodes with the highest node densities at the mesocosm top and at the interfaces of the soil materials. Water flow was described by the Richards equation (Richards, 1931) and the constitutive relations of the van Genuchten–Mualem model without hysteresis (Mualem, 1978; van Genuchten, 1980). Upper boundary conditions for the water flow accounted for temporally variable irrigation rates and potential evapotranspiration, which was modeled using weekly estimates (Sect. 2.2.4). At the domain bottom, a variable pressure head boundary condition was used to account for the applied suction range of $-0.1$
to −0.75 bar relative to atmospheric pressure (Thaysen et al., 2014a). The root depth was fixed at 30 cm throughout the experiment, corresponding to a measured root depth of 30 cm on simulation day 0 (day 15 after sowing) and allocation of ~90% of the root mass in the A horizon at the end of the experiment (as measured in a similar experiment; Thaysen et al., 2014b). A normalized exponential root distribution function was employed to describe the vertical root distribution within the A horizon (Hoffmann and van Genuchten, 1983). Root water uptake was modeled using the root distribution and the S-shaped water uptake model without solute stress and default parameterization (van Genuchten, 1987).

Water retention parameters were obtained from inverse modeling of the water flow in an unplanted mesocosm (data not shown) using Hydrus 1-D (Šimůnek et al., 2013) and a global stochastic optimization algorithm (Vrugt et al., 2009).

Solute transport was modeled with the advection-dispersion equation. Carbon dioxide transport processes included in HP1 are diffusion in the gas phase and advective-hydrodynamic dispersion in the aqueous phase. Carbon dioxide diffusion in the water and gas phase, respectively, was modeled by scaling the CO2 diffusion coefficient in water and air with the Millington and Quirk tortuosity (Millington and Quirk, 1961). The boundary layer height was set at 0.02 m. Equilibrium CO2 distribution between the gas and the water phase is described in HP1 by Henry’s law. More details are available in Šimůnek et al. (2006) and Jacques et al. (2008).

Heat transport was also described with an advection-dispersion equation using thermal conductivity data for sand (Chung and Horton, 1987). Soil temperatures of 22 and 18 °C were chosen for the upper and lower boundary conditions, respectively, corresponding to the measured temperature decline from the top to the bottom of the mesocosm (Fig. S1). Day and night cycles were modeled with temperature amplitudes of 5 °C.

2.3.2 CO2 production and root growth

Soil CO2 production was modeled through implementation of equations and parameters from the SOILCO2 model (Šimůnek and Suarez, 1993; Suarez and Šimůnek, 1993), making a few modifications. The total CO2 production rate, \( S \) (mol m\(^{-2}\) s\(^{-1}\)), is considered as the sum of the production by the soil microorganisms, \( \gamma_s \) (mol m\(^{-2}\) s\(^{-1}\) g\(^{-1}\) DW\(_{\text{root}}\)), and the production by plant roots, \( \gamma_p \) (mol m\(^{-2}\) s\(^{-1}\) g\(^{-1}\) DW\(_{\text{root}}\)), and the production by plant roots, \( \gamma_p \) (mol m\(^{-2}\) s\(^{-1}\) g\(^{-1}\) DW\(_{\text{root}}\)), and the production by plant roots, \( \gamma_p \) (mol m\(^{-2}\) s\(^{-1}\) g\(^{-1}\) DW\(_{\text{root}}\)), according to

\[
S = (\gamma_s + \gamma_p)\text{RMI},
\]

where the subscripts \( s \) and \( p \) refer to soil microorganisms and plant roots, respectively, and RMI is the root mass index in the system (g DW). Both \( \gamma_p \) and \( \gamma_s \) were linked to the root distribution, since microbial respiration is primed by root mass (Kuzuyakov, 2002; Zhu and Cheng, 2011). It is assumed that individual CO2 producing processes are additive and that it is possible to superpose individual mechanisms that reduce production from an optimal value. The production rates for microbial and root respiration at any given point in time and space are described by Eqs. 3–5

\[
\gamma_s = \gamma_{s0} \prod_i f_{s_i},
\]

\[
\gamma_p = \gamma_{p0} \prod_i f_{p_i},
\]

\[
\prod_i f_i = f(h) f(T) f_{CO_2}(c_a) f(h_o) f(z),
\]

where \( \gamma_{s0} \) and \( \gamma_{p0} \) are the optimum respiration rates for microorganisms and roots (mol m\(^{-2}\) s\(^{-1}\) g\(^{-1}\) DW\(_{\text{root}}\)), respectively, and the term \( \prod f_i \) multiplies all reduction coefficients. The coefficient \( f(h) \) is the reduction coefficient as a function of pressure head (unitless), \( f(T) \) is the reduction coefficient as a function of temperature (unitless) and \( f_{CO_2}(c_a) \) is the reduction coefficient as a function of CO2 concentration (reduced oxygen availability) (unitless). These coefficients are assumed to be equal for \( \gamma_{s0} \) and \( \gamma_{p0} \). The coefficient \( f(h_o) \) is the reduction coefficient as a function of osmotic head (unitless) which is set to 1 for \( \gamma_{s0} \) (no reduction). Expressions for all reduction coefficients except for the reduction coefficient as a function of depth in the soil profile, \( f(z) \), are identical to those in SOILCO2, more details are available in Šimůnek and Suarez (1993). The coefficient \( f(z) \) is described differently for \( \gamma_{s0} \) and \( \gamma_{p0} \), as defined below. All reduction terms have values between 0 and 1, except the \( f(T) \) which is higher than 1 above 20 °C and smaller than 1 below 20 °C.

Increasing CO2 production due to the evolving root mass, RMI, is described by a linear biomass increase with time:

\[
\text{RMI} = R_{\text{init}} + (\text{time} \cdot r),
\]

where \( R_{\text{init}} \) is the initial root mass at simulation time zero (g DW), and \( r \) is the root growth rate (g s\(^{-1}\)).

The depth reduction factor for root respiration, \( f_p(z) \), is directly linked to the modeled vertical root distribution. For the microbial respiration, \( f_s(z) \) is described with an exponential function containing an \( a \) parameter (see Šimůnek and Suarez, 1993) which was set to 0.0015 m\(^{-1}\) in our simulations.

In SOILCO2, there is no time dependency of microbial respiration (i.e., \( \gamma_s \) is not linked to RMI), and the Verhulst–Pearl logistic growth function is used to describe root growth from differences in vertical root penetration depth. The use of these assumptions revealed a poor fit to our data set. For our simulations, \( R_{\text{init}} \) and \( r \) were fixed at 2 g DW and 2.4 \times 10\(^{-6}\) g s\(^{-1}\), respectively. Assuming linear root growth, the \( r \) was calculated by dividing the measured root biomass at the end of a similar mesocosm experiment (13.7 g at 70 days after sowing and 64.5 days after plant emergence; Thaysen et al., 2014b) with 64.5 days of plant growth. The \( R_{\text{init}} \) was then calculated from multiplication of the \( r \) with the number
of days after plant emergence at simulation time zero (i.e., 15 days after sowing minus 5.5 days of plant emergence). The $\gamma_{p0}$ and $\gamma_{a0}$ parameters were set to 0.8 $\mu$mol m$^{-2}$ s$^{-1}$ gDW$^{-1}$ root, assuming equal contributions of microbial and root respiration to the total $R_s$ (mean of reported contributions of root respiration to the total $R_s$ of 10–90%; Swinnen, 1994; Hanson et al., 2000; Kocyigit and Rice, 2006; Moyano et al., 2007). When multiplied by the root mass at the end of the experiment, the sum of $\gamma_{p0}$ and $\gamma_{a0}$ resulted in a CO$_2$ production of 16 $\mu$mol m$^{-2}$ s$^{-1}$, which was within the calculated range of the actual average CO$_2$ production of 15–31 $\mu$mol m$^{-2}$ s$^{-1}$ in planted mesocosms (Sect. 3.4). An overview of all applied parameters is given in Table S1 in the Supplement.

The simulated CO$_2$ efflux from the ecosystem to the atmosphere is essentially the soil respiration, $R_s$, because there is no plant module in this conceptual model. Hence, our ER measurements could not be directly compared to the simulated CO$_2$ efflux. Anticipating that (1) canopy respiration was roughly 50% of the total plant respiration (Poorter et al., 1990; Loveys et al., 2002), and (2) root respiration accounted for 50% of the total $R_s$ (see above), we corrected the ER by a factor of 0.67 to get an estimate of the $R_s$.

2.3.3 Root solute uptake

Root solute uptake was simulated based on literature values on the average plant content of major ions. Average plant nutrient contents were (µmol g DW plant$^{-1}$; from Marschner, 1995): K$^+$, 250; Ca$^{2+}$, 125; Mg$^{2+}$, 80; PO$_4^{3-}$, 60; SO$_4^{2-}$, 30; Na$^+$, 0; total N (modeled as NO$_3^-$), 1000. The daily influx per g DW root, $J$, was calculated by multiplying the average plant nutrient content by the rate of total plant biomass increase over the experiment (1.05 g d$^{-1}$) and dividing by the total root mass at the end of the experiment (~13.7 g). The root mass dependent influx, $J_{TIME}$ (mol s$^{-1}$), was then obtained by multiplication of $J$ with the RMI and the normalized vertical root distribution, $f_p(z)$, (Eq. 7).

$$J_{TIME} = J \cdot RMI \cdot f_p(z).$$  (7)

Because the root length was fixed at 30 cm (Sect. 2.3.1), but the root mass within this depth increased by the RMI, root solute uptake was simulated to increase linearly over time within the A horizon.

Root cation uptake was described by simultaneous excretion of protons, except for K$^+$ for which Na$^+$ was assumed to be excreted. Anion uptake was modeled by OH$^-$ excretion. This approach for root solute uptake modeling is in agreement with general knowledge on root nutrient uptake mechanisms (Marschner, 1995).

2.3.4 Geochemistry

Aqueous speciation and complexation were accounted for using PHREEQC (Parkhurst and Appelo, 2013) and the wateq4f.dat database. Cation exchange capacities and initial compositions were as measured from soil extractions (Kjøller et al., 2004). Initial exchanger compositions were equilibrated with a solution composition in contact with a $p$CO$_2$ of 1% and with 0.3 meq L$^{-1}$ alkalinity, as measured on day 15 after sowing. Charge balance of the initial solution was achieved using Li$^+$ and an equilibrium constant for the exchange reaction of log $k = -100$. Nutrient irrigation was modeled using the Hoagland solution composition diluted 1 : 1 by H$_2$O.
Figure 3. Gaseous and dissolved CO₂ fluxes from mesocosms. (a): ecosystem respiration (ER) from barley mesocosms during growth (black) and soil respiration (Rₛ) from barley mesocosms after harvest and unplanted mesocosms (green). (b): net ecosystem exchange (NEE) from mesocosms with barley. (c): dissolved inorganic carbon (DIC) percolation flux recalculated into gaseous efflux unit. (d): cumulative DIC percolation flux as a function cumulative drainage.

3 Results

3.1 Distribution of total inorganic carbon between gaseous and aqueous phase species

The total inorganic C distributed between gaseous and aqueous phase species (i.e., CO₂(g) and DIC) increased during barley growth and decreased after harvest (Fig. 2a and b). Over time, inorganic C was mainly found as CO₂(g) and changed little beyond 20 cm depth. A downward moving front of elevated [DIC] was not visible. The latter was expected in the middle to bottom range of the C horizon due to the downward movement of initially present pore water. In unplanted mesocosms, the amount of inorganic C and distribution between species hardly changed over the course of the experiment (Fig. 2c).

3.2 CO₂ exchange and aboveground biomass

The ER increased with time during plant growth (Fig. 3a). After harvest of the aboveground biomass, the Rₛ declined logarithmically (Fig. 3a). The Rₛ from unplanted mesocosms was much lower than from barley mesocosms and was relatively constant. The negative NEE, i.e., ecosystem CO₂ uptake, increased with plant age up to day 54 after which it decreased to less negative values similar to those observed 20 days after sowing (Fig. 3b). Soil surface temperatures were 20–23 °C during measurements (Fig. S1). The aboveground biomass in mesocosms of the barley growth treatment #2 was 2918 ± 83 g m⁻².

3.3 DIC percolation

The DIC percolation flux varied considerably in all treatments but tended to increase with plant growth (range: 0.02–1 µmol CO₂ m⁻² s⁻¹), decrease after harvest (0.03–0.3 µmol CO₂ m⁻² s⁻¹), and remain fairly constant in unplanted soil (0.01–0.2 µmol CO₂ m⁻² s⁻¹; Fig. 3c).

The cumulative amount of DIC that leached from barley mesocosms in the first growth treatment and post-harvest was 303–311 mmol C m⁻² at a cumulative drainage of 212–234 mm (Fig. 3d). This drainage corresponded to an estimated 1.8–2.0 times the initially water-filled pore volumes, with 0.6–0.7 times the water-filled pore volumes flushed during growth. The average [DIC] during the pre-harvest and post-harvest periods was 1.8–2.0 and 1.1–1.3 mmol L⁻¹, respectively.

In the second barley growth treatment, the cumulative amount of DIC leached was 360–440 mmol C m⁻² and significantly higher than the 80–82 mmol C m⁻² leached from the corresponding unplanted mesocosms (Fig. 3d). The cumulative drainage from barley mesocosms of the second barley treatment and from unplanted mesocosms was 139–168 and 141–158 mm, respectively (Fig. 3d) and corresponded to 1.3–1.5 and 1.1–1.3 times the initially water-filled pore volumes. The average [DIC] in mesocosms of the second barley treatment and in unplanted mesocosms was 2.5 and 0.4–0.5 mmol L⁻¹, respectively. The cumulative DIC percolation fluxes from all mesocosms were highly correlated with the cumulative drainages (Fig. 3d; R² = 0.96–0.99).
Figure 5. Measured (full lines or dots) and simulated (dashed lines) temporal variation (expressed in days after sowing) of soil air \( p\text{CO}_2 \) (a), alkalinity (b), pH (c), \( \text{Al}^{3+} \) (d), \( \text{CaX}_3 \) (e), \( \text{Ca}^{2+} \) (f), \( \text{AlX}_3 \) (g), \( \text{Ca}^{2+} \) root uptake (h), \( R_n \) (i) and cumulative DIC percolation (j) during barley growth. Geochemical reactions in simulations were described by root nutrient uptake and cation exchange (scenario 1). In (i) small fluctuations in the simulated \( R_n \) around the baseline arise from diurnal temperature variations. Large fluctuations are numerical noise caused by the fact that the numerical solution does not fully obey the von Neumann stability criteria.

3.4 CO₂ emission partitioning

The cumulative CO₂ effluxes to the atmosphere from unplanted and planted soil (barley growth #2) during 78 days of experiment, using linear interpolation between measurements, were 0.4–1.1 and 32.8–33.4 mol C m\(^{-2}\), respectively (not shown). The corresponding cumulative DIC percolation fluxes of 0.080–0.082 and 0.36–0.44 mol C m\(^{-2}\), respectively, constituted 7.3–20.5 and 1.6–2.0 % of the aboveground CO₂ emission. The cumulative post-harvest CO₂ efflux to the atmosphere during 60 days of experiment was 2.4–2.7 mol C m\(^{-2}\) at a cumulative DIC percolation flux of 0.15–0.16 mol C m\(^{-2}\), corresponding to a relative magnitude of the cumulative DIC flux compared to the cumulative aboveground CO₂ efflux of 5.6–6.7 %.

3.5 Soil diffusivity and CO₂ production

The Rn emanation rates of the soil A and C horizons were 4.93 ± 0.46 and 2.65 ± 0.13 atoms s\(^{-1}\) kg\(^{-1}\), respectively. At a VWC of ∼20 % in the A horizon, the Rn concentration [Rn] was 1.5 times larger in unplanted mesocosms than in planted mesocosms, with a difference of ∼1000 Bq m\(^{-3}\) (Fig. 4). The bulk soil diffusivity was determined by comparing measured [Rn] profiles at a given VWC with modeled [Rn] at different bulk diffusivities. The code RnMod3d could reproduce the Rn profiles in unplanted mesocosms using bulk diffusivities of 0.6 × 10\(^{-6}\) and 2.0 × 10\(^{-6}\) m\(^2\) s\(^{-1}\) for the A and C horizon, respectively. For barley mesocosms, the bulk diffusivities for the A and C horizon were 2.1 × 10\(^{-6}\) and 1.9 × 10\(^{-6}\) m\(^2\) s\(^{-1}\), respectively (Fig. 4). The confidence intervals in Fig. 4 mark the range of diffusivities at which agreement between measured and modeled [Rn] could be achieved. Modeled soil bulk diffusivities of the unplanted
A and C horizons showed reasonable agreement with bulk diffusivities of $0.44 \times 10^{-6}$ and $0.9 \times 10^{-6}$ m$^2$ s$^{-1}$, respectively, calculated from empirical formulas in Rogers and Nielsen (1991) and Andersen (2001) (see Text S1 in the Supplement).

The CO$_2$ diffusion coefficient in unplanted and barley mesocosms was estimated to 1.2–2.3 $\times 10^{-6}$ and 1.8–6.8 $\times 10^{-6}$ m$^2$ s$^{-1}$, respectively, using the Rn bulk diffusivities (Eq. 1). Average soil CO$_2$ production as calculated from the range for the A horizon bulk diffusivities was 1.0–1.8 and 15.1–31.1 µmol m$^{-2}$ s$^{-1}$ for mesocosms of the unplanted and barley growth #2 treatment, respectively, where CO$_2$ production during barley growth increased from $\sim$5 to 59 µmol m$^{-2}$ s$^{-1}$ (confidence intervals: 3.5–7.4 to 30–87 µmol m$^{-2}$ s$^{-1}$).

### 3.6 Modeling of CO$_2$ fluxes

The CO$_2$ diffusivity and production estimates (Sect. 3.5) were used to simulate the measured CO$_2$ fluxes in one mesocosm of the second barley growth treatment. Different geochemical constraints were applied in three scenarios. In scenario 1, root nutrient uptake was coupled to cation exchange. Scenario 2 had cation exchange and equilibrium (saturation index = 0) for amorphous aluminium hydroxide, Al(OH)$_3(a)$, as the soil solution chemistry in the mesocosms revealed a possible control by Al(OH)$_3(a)$ (Fig. S2 in the Supplement). Scenario 3 was equal to scenario 2 except that root nutrient uptake was included. The fit of scenarios 2 and 3 to measured parameters was poor (not shown) and hence the discussion of our modeling is limited to scenario 1.

Due to the dependency of the $p$CO$_2$ and the $R_5$ on the VWC, a correct description of the VWC in the mesocosms at any given point in time was crucial. The dependency of the DIC percolation flux on the drainage flux implied a need for an exact simulation of the drainage flux. Both water flow and drainage flux were generally well described (Fig. S5 in the Supplement).

The temporal variation of the simulated $p$CO$_2$ and $R_5$ captured the main trends in the observations (Fig. 5a, i). However, in the upper part of the mesocosm, the simulated $p$CO$_2$ was somewhat higher than the measured $p$CO$_2$ up to day 36 despite a good agreement between simulated and observed topsoil VWC (Fig. S5) and the $R_5$ (Fig. 5i). The simulated cumulative DIC percolation provided a good fit to the measured data up to day 46 but underestimated the measured DIC percolation on day 53 (Fig. 5j) due to lower simulated alkalinity towards the mesocosm bottom (Fig. 5b).

A general increase in alkalinity over time was caused by the high root uptake of NO$_3^-$ that exceeded uptake of any other plant nutrients (Fig. S3 in the Supplement) and caused a net excretion of OH$^-$. This counteracted the pH drop caused by CO$_2$ dissolution and buffered the pH around 5.6–6.0 (Fig. 5c). Mismatch between the simulated and the calculated pH were largely caused by differences between measured and simulated alkalinity and $p$CO$_2$. The simulated high near-surface pH (Fig. 5c) was due to low $p$CO$_2$ caused by the diffusional loss of CO$_2$ and high alkalinity caused by evaporation processes (see tracer simulation, Fig. S4 in the Supplement).

Scenario 1 simulated increasing [Al$^{3+}$] and [Ca$^{2+}$] with time that approached the measurement on day 71 (Fig. 5d and f, respectively). Increasing [Al$^{3+}$] resulted from displacement of exchanger-bound Al by cations in the incoming nutrient solution. High [Al$^{3+}$] led to supersaturation of Al(OH)$_3(a)$ (not shown), in accordance with Fig. S3 in the Supplement. Calcium was displaced on the A horizon exchanger as well (Fig. 5e), resulting in increasing [Ca$^{2+}$] with time. However, the largest increase in [Ca$^{2+}$] resulted from irrigation with nutrient solution and subsequent up-concentration due to evaportranspiration (Fig. 5f). Root uptakes of Ca$^{2+}$ (and other nutrients) were small compared to [Ca$^{2+}$] in the soil solution (Figs. 5h, S3) but the accompanying release/uptake of protons had a big impact on the soil alkalinity and pH (Fig. 5b and c). Exchanger-bound Al and Ca were displaced by K (not shown), probably due to higher [K$^+$] in the nutrient solution compared to other ions, and the high ionic strength of the nutrient solution (13.9 mmol kg$^{-1}$ water) (Appelo and Postma, 2005).

### 4 Discussion

In this work, we have quantified the inorganic C dynamics in unplanted and unplanted soil mesocosms representing the vadose zone of a cultivated podzol. The data were generated from a detailed collection of gaseous and aqueous samples and from subsequent modeling enabled by the novel implementation of SOILCO$$_2$$ into HP1. We show that inorganic C dynamics in ecosystems are not well described from measurements of the CO$$_2$$ pools and fluxes alone, and that biogeochemistry has a potentially major impact on the movement of dissolved CO$$_2$$ in the vadose zone. In the following section, our results are discussed in the context of soil type effects on CO$$_2$$ fluxes and differences between mesocosm and field studies are addressed.

#### 4.1 Soil respiration: gaseous CO$$_2$$ efflux

The $R_5$ and the $p$CO$$_2$ (Figs. 3a and S1) in unplanted mesocosms were generally in agreement with field studies on a range of different arable soils (Table 2). In barley mesocosms, the ER and the $p$CO$_2$ were generally higher than in published field studies but were in accordance with pot-grown barley in humic clay (Simojoki et al., 1991) (Table 3). Higher respiration rates in barley mesocosms than in the field were probably a function of a larger plant biomass, as also observed by Simojoki et al. (1991). The above-ground and root biomass was 3–4 times and 2–5 times higher, respectively, than the values reported from field studies.
Table 2. Ranges of soil respiration ($R_s$) and soil partial pressure of CO$_2$ ($p$CO$_2$) in unplanted mesocosms as compared to other studies (one in pots, four in the field). NA represents not available, VWC represents volumetric water content.

<table>
<thead>
<tr>
<th>$R_s$ (µmol m$^{-2}$ s$^{-1}$)</th>
<th>$p$CO$_2$ (%)</th>
<th>Crop</th>
<th>Soil type</th>
<th>Soil texture</th>
<th>Soil surface temperature (°C)</th>
<th>Daytime air temperature (°C)</th>
<th>VWC (%)</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>−3.0–3.2</td>
<td>0.2–1.2</td>
<td>Fallow</td>
<td>Podzol</td>
<td>Coarse sand</td>
<td>22–23</td>
<td>18</td>
<td>20–25</td>
<td>This study (mesocosms)</td>
</tr>
<tr>
<td>NA</td>
<td>NA</td>
<td>Fallow</td>
<td>Gleyic Cambisol</td>
<td>Sandy soil</td>
<td>18–22</td>
<td>NA</td>
<td>NA</td>
<td>Weihermüller et al., 2009 (pot experiment)</td>
</tr>
<tr>
<td>~0.7–4.5</td>
<td>NA</td>
<td>Fallow</td>
<td>Arenosol</td>
<td>Sandy soil</td>
<td>3–20</td>
<td>−10–28</td>
<td>10–25</td>
<td>Herbst et al., 2008</td>
</tr>
<tr>
<td>NA</td>
<td>0.5 ±0.1</td>
<td>Fallow</td>
<td>Montmorillonitic, Mesic Udolic Ochraqualfs</td>
<td>Silt loam</td>
<td>15–20</td>
<td>13–15</td>
<td>27–30</td>
<td>Buyanovsky and Wagner, 1983</td>
</tr>
<tr>
<td>∼0.05–3.7</td>
<td>NA</td>
<td>Fallow</td>
<td>Orthic Humic Gleysol</td>
<td>Loam</td>
<td>10–26</td>
<td>NA</td>
<td>25–40</td>
<td>Rochette et al., 1992</td>
</tr>
<tr>
<td>1.5–8</td>
<td>NA</td>
<td>Fallow</td>
<td>Endocalcari–Epiphygleyic Cambisol</td>
<td>Clay loam</td>
<td>19–27</td>
<td>11–22</td>
<td>NA</td>
<td>Feizene et al., 2008</td>
</tr>
<tr>
<td>−2–10</td>
<td>NA</td>
<td>Fallow</td>
<td>Endocalcari–Epiphygleyic Cambisol</td>
<td>Clay loam</td>
<td>19–27</td>
<td>11–22</td>
<td>NA</td>
<td>Feizene et al., 2008</td>
</tr>
</tbody>
</table>

Table 3. Ranges of ecosystem respiration (ER) and soil partial pressure of CO$_2$ ($p$CO$_2$) during growth in barley mesocosms as compared to other studies (two in mesocosms, seven in the field). NA represents not available, VWC represents volumetric water content.

<table>
<thead>
<tr>
<th>ER (µmol m$^{-2}$ s$^{-1}$)</th>
<th>$p$CO$_2$ (%)</th>
<th>Crop</th>
<th>Soil type</th>
<th>Soil texture</th>
<th>Soil surface temperature (°C)</th>
<th>Daytime air temperature (°C)</th>
<th>VWC (%)</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>0.8–52</td>
<td>0.2–8.8</td>
<td>Barley</td>
<td>Podzol</td>
<td>Coarse sand</td>
<td>20–24</td>
<td>18</td>
<td>20–25</td>
<td>This study (mesocosms)</td>
</tr>
<tr>
<td>NA</td>
<td>~2–10</td>
<td>Barley</td>
<td>Humic clay soil</td>
<td>Clay</td>
<td>NA</td>
<td>NA</td>
<td>15–32</td>
<td>Simejoki et al., 1991 (mesocosms)</td>
</tr>
<tr>
<td>NA</td>
<td>~1–16</td>
<td>Barley</td>
<td>Podzol</td>
<td>Coarse sand</td>
<td>−7 to 29 (mean ~8)</td>
<td>8 (mean)</td>
<td>6–34</td>
<td>Our field site (Lohila et al., 2003)</td>
</tr>
<tr>
<td>NA</td>
<td>0.3–2</td>
<td>Barley</td>
<td>Eutric Cambisol</td>
<td>Sandy soil with high humus content</td>
<td>12–23</td>
<td>18–30</td>
<td>13–23</td>
<td>Lohila et al., 2003</td>
</tr>
<tr>
<td>NA</td>
<td>0.1–0.3</td>
<td>Barley</td>
<td>Haplic Phaeozem</td>
<td>Silty clay loam</td>
<td>NA</td>
<td>~9.3 (mean)</td>
<td>~17–33</td>
<td>Walmsley et al., 2011</td>
</tr>
<tr>
<td>2.8–4.5</td>
<td>0.3–6</td>
<td>Barley</td>
<td>Epiphygleyic Cambisol</td>
<td>Clay loam</td>
<td>19–26</td>
<td>11–22</td>
<td>15–30</td>
<td>Buyanovsky and Wagner, 1983</td>
</tr>
<tr>
<td>2.3–0.75</td>
<td>NA</td>
<td>Barley</td>
<td>Orthic Humic Gleysol</td>
<td>Loam</td>
<td>10–19</td>
<td>NA</td>
<td>10–35</td>
<td>Rochette et al., 1992</td>
</tr>
</tbody>
</table>

4.2 Percolation: DIC fluxes

The DIC percolation flux was calculated from the alkalinity, the $p$CO$_2$ at the mesocosm bottom and the drainage flux but was primarily a function of the latter two variables. The smaller effect of the alkalinity on the downward DIC flux was caused by the low soil pH that shifted DIC primarily towards H$_2$CO$_3^+$ (Fig. 2). The absence of a downward moving DIC front was caused by the fact that H$_2$CO$_3^+$ is a direct function of the $p$CO$_2$ that was constant with depth (Fig. S1).

The higher $p$CO$_2$ in barley mesocosms caused about a 4 times higher DIC percolation flux than in unplanted mesocosms at similar drainage (Fig. 3c and d). Average post-harvest [DIC] were significantly higher than the average [DIC] in the percolation water from unplanted soil but may have partly been influenced by high [DIC] during plant growth due to incomplete exchange of the initial water-filled pore volume of during barley growth in the first barley treatment. The latter implies that high [DIC] arising from high $p$CO$_2$ conditions at the mesocosm top during barley growth were not transported all the way to the mesocosm outlet.

The DIC percolation flux and the average [DIC] under barley were within the range of reported values for cropland soil (Table 4). Further comparison of the [DIC] in the percolation water is impeded by the numerous factors influencing soil $p$CO$_2$ and the [DIC] (see Sects. 1 and 4.1). The DIC fluxes.
percolation flux from unplanted soil was similar to the previously reported percolation fluxes from sandy forest soils (Kindler et al., 2011).

### 4.3 CO₂ emission partitioning and controls on CO₂ fluxes in the vadose zone

Our measurements revealed a pivotal influence of vegetation on CO₂ fluxes in the vadose zone as both upward and downward transport of CO₂ was strongly increased in planted mesocosms (Fig. 3). The cumulative DIC percolation flux was small relative to the cumulative aboveground CO₂ efflux. Results from planted mesocosms (1.6–2.0 %) were higher than the global emission flux partitioning (0.26 %) (Raich and Potter, 1995; Kessler and Harvey, 2001) but lower than a 2.5 % fraction reported for an onion field (Sawamoto et al., 2003). The relatively higher ratio between cumulative DIC percolation flux and the cumulative Rₚ in unplanted mesocosms resulted from lower cumulative Rₚ in unplanted soil that was further decreased by periods of net uptake of CO₂ (Fig. 3a). The potential for manipulating the emission balance by increasing the DIC export to the groundwater via irrigation is substantial after harvest where the absence of plant transpiration facilitates fast percolation of water through the soil and where relatively high soil pCO₂ may transmit to increases in the [DIC].

The interpretation of CO₂ fluxes in mesocosms needs to take into account the high plant biomass density in mesocosms, the constant summer conditions in the climate chamber and the coarse-sanded, acidic soil of this study. Elevated biomass of plants grown at relatively high temperature increased the soil respiration which again increased both upward and downward flux of CO₂. Smaller [Rn] in barley mesocosms (Fig. 4) indicated that root growth and decay increased the soil diffusivity (by $\sim 1.5 \times 10^{-6} \text{ m}^2 \text{ s}^{-1}$). This may have enhanced the Rₚ, implying a decreased net uptake of atmospheric CO₂ (NEE) by the plant-soil ecosystem. Increased diffusivity due to the presence of biopores and cracks has indeed been reported (Holford et al., 1993; Hoff, 1997). However, some reduction in the [Rn] in planted soil may have been caused by Rn removal via plant transpiration (Lewis and MacDonell, 1990; Jayaratne et al., 2011). In any case, the high soil diffusivity of the coarse sand and the relatively low soil pH of $\sim 6$ caused much less downward transport of DIC compared to the CO₂ emission to the atmosphere (Fig. 3).

Modeling of the CO₂ fluxes using different explanatory scenarios was a valuable tool for identifying the key processes behind the observed evolution in the dissolved CO₂ fluxes. Scenario 1 (root nutrient uptake coupled to cation exchange) was in accordance with previous demonstrations of slight soil alkalization when nitrate is the dominating inorganic N species (Marschner et al., 1991; Weligama et al., 2010). However, our modeling of the nutrient uptake remains somewhat uncertain due to lack of data for actual nutrient uptake. A control of Al(OH)₃(a) on the soil solution was unlikely but other buffering processes such as the dissolution of small pieces of lime (Fig. S2) may be possible.

Soil pCO₂ and Rₚ were hardly affected by any buffering process, despite the low subsoil pH that caused partitioning of DIC towards H₂CO₃ and some degassing to CO₂(g) (Fig. 2). This is in accordance with existing knowledge that mineral reactions affecting the carbonate system have only marginal effects on the pCO₂ (Kuzakoy, 2006).

The modified SOILCO₂ model made use of several simplifications of reality, one of them being the linkage of the $yₐ$ to the RMI without inclusion of a bulk soil $yₐ$. However, due to the dense root growth in the A horizon (see Thaysen et al., 2014b, Fig. 1 and Sect. 4.1) plant root independent $yₐ$ was probably negligible in the A horizon. The exponential decline in the $yₐ$ with soil depth by the $fₚ(z)$ (Eqs. 3–5) further implied that the $yₐ$ in the C horizon was small compared to the topsoil. Hence, for the experimental conditions of this study, the omission of a plant-root independent $yₐ$ in our model seems justified.
5 Conclusions

The DIC percolation flux of $\sim 5$ mmol C m$^{-2}$ d$^{-1}$ during barley growth was $\sim 1.6$–$2.0\%$ of the $R_s$ at increased plant biomass and elevated soil $p$CO$_2$ compared to the field situation. After harvest, the magnitude of the DIC percolation flux was lowered to $\sim 2.5$ mmol C m$^{-2}$ d$^{-1}$ but the importance of DIC percolation flux for the overall cropland $CO_2$ emission increased to $\sim 6$–$7\%$ of the $R_s$.

At constant conditions of temperature and water content, the $R_s$ was controlled by the production and diffusivity of $CO_2$ in the soil, both of which were increased by plant growth. The DIC percolation flux was primarily controlled by the recharge flux and the $p$CO$_2$ due to the low soil pH in the acidic soil of our study. Modeling suggested that nutrient buffering during root nitrate uptake dominated any mineral control on the soil solution in planted mesocosms.

This study showed that the integration of experimental and modeling work is a powerful tool in advancing process-understanding of $CO_2$ fluxes in the vadose zone. Our findings are important for improving current base understanding of $CO_2$ partitioning in the vadose zone and may be included in the optimization of climate models. Further research is needed to outline the effect of different crops and soil amendments on the $CO_2$ emission portioning of croplands.

The Supplement related to this article is available online at doi:10.5194/bg-11-7179-2014-supplement.

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