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Abstract

Changes in the isotopic composition of oxygen associated with phosphate can provide information on the impact of phosphatase activity on soil P dynamics, whereas the use of radioactive P delivers information on P fluxes within soil systems. Although these two tracers may provide complementary data, they have rarely been used together to study soil P cycling. We conducted a dual isotopic soil incubation study of one month with soils originating from four sites of a climatic gradient (Kohala, Hawaii), which provides well-controlled geochemical and biological variations on soils derived from the same parent material. Three groups of soils were incubated in parallel, the first group labelled with ^{32}P radioisotopes, the second group labelled with ^{18}O enriched water and the third group not labelled and used for CO_2 emission measurements. The dual labelling study informed about three processes controlling P dynamics in soils: those that maintain the bond between P and O and transfer phosphate from one pool to another (category I processes), those that involve the cleavage of the P-O bond and transfer phosphate from one pool to another (category IIa processes), and those that involve the cleavage of the P-O bond but do not transfer phosphate from one pool to another (category IIb processes). The use of ^{32}P showed that the studied soils contained a large amount of P that was isotopically exchangeable with the resin P pool (category I process) and that microorganisms had taken up P, but in much lower amounts, from the resin P pool (category I process). ^{18}O added with water was incorporated into microbial and resin P, but not into the other pools obtained from the modified Hedley extraction. Thus, the turnover of O associated with P within microbial cells (category IIb process) and/or enzymatic hydrolysis of organic P (category IIa process) had occurred and had affected active microbes, which passed the ^{18}O labelled phosphate to the resin pool (category I process). The dual isotopic approach thus provided complementary insights on P cycling processes.

1. Introduction

Phosphorus (P) transformations in soils are controlled by sorption/desorption, precipitation/dissolution, microbial mineralisation/immobilisation, plant uptake and death, and organic matter dynamics (Frossard *et al.*, 2000; Oberson and Joner, 2005; McLaren *et al.*, 2019). These transformations are most often studied by using sequential chemical extractions, such as the one developed by Hedley *et al.* (1982). However, information on P distribution among chemical extracts provides a static picture of what is present at a given time; it does not provide information on P cycling processes (Helfenstein *et al.*, 2018). Abiotic and biotic P cycling processes are constantly occurring in the soil even if extracted amounts of P stay constant. Approaches that integrate tracers are therefore required to better understand soil P fluxes and transformations.

In this paper, we evaluate processes controlling P dynamics in soils by targeting the isotopic composition of phosphate molecules, in which one P atom is bound to four oxygen (O) atoms. Diverse process categories are differentiated (Figure 1): those that involve entire phosphate molecules moving as intact units from one pool to another (category I processes), and therefore, the P-O bonds remain untouched; and those that cleave P-O bonds (category II processes). Within the category II processes, we consider those that lead to a phosphate transfer from one pool to another (category IIa processes) and those that do not lead to a phosphate transfer from one pool to another (category IIb processes). Sorption/desorption, precipitation/dissolution as well as P uptake by organisms and release upon cell lysis involve the transfer of phosphate molecules without the cleavage of a P-O bond and therefore can be classified as category I processes. Enzymatic processes mediated by phosphomono- and phosphodiesterase can be assigned to the category IIa processes as they lead to the cleavage of P-O bonds and the release of phosphate from phosphomono- or phosphodiesteres to the soil solution (Blake *et al.*, 2005). Finally, the exchange of O atoms on a phosphate mediated by pyrophosphatase within a cell can be assigned to a category IIb process as the phosphate stays in the same pool (Tamburini *et al.*, 2014).

Location Figure 1

Radioactive P isotopes (^{32}P , ^{33}P) are used to trace the fate of P in soil-plant systems. More specifically, P radioisotopes are used to measure phosphate exchange kinetics between the soil solution and the soil solid phase for assessment of the within a given time isotopically exchangeable phosphate, to estimate the rate of organic P mineralisation, or to quantify the P uptake by a plant or the P recovery from an external source (e.g. a plant residue or a fertilizer) in a specific soil pool (Frossard *et al.*, 2011). Some of these P fluxes are controlled by category I processes (sorption/desorption assessed by isotopic exchange, release of phosphate from dead microbes to the soil solution pool), while other P fluxes belong to category IIa (organic P hydrolysis with P-O bond cleavage and release of phosphate to the soil solution).

In the absence of biological activity and at earth surface temperatures, the P-O bond in phosphate is stable and the isotopic composition of O associated with P ($\delta^{18}\text{O-P}$) remains constant (Blake *et al.*, 2005; Tamburini *et al.*, 2014). Small changes in $\delta^{18}\text{O-P}$ can occur without cleavage of P-O bonds, for instance during apatite dissolution (Liang and Blake, 2007; Jaisi *et al.*, 2010), during short-term sorption (Jaisi *et al.*, 2010; Melby *et al.*, 2013b), or during P uptake by microorganisms (Blake *et al.*, 2005). Enzyme-mediated reactions have on the contrary a strong impact on $\delta^{18}\text{O-P}$. They lead to P-O bond cleavage and O atom exchange between phosphate and water (category II processes), which either results in the equilibration of $\delta^{18}\text{O-P}$ with ^{18}O of water or causes kinetic isotope fractionation effects. Temperature-dependent equilibration can lead to complete O exchange over time and is considered to be mainly intra-cellular and mediated by pyrophosphatase (Chang and Blake, 2015; von Sperber *et al.*, 2017b). Intra- and extra-cellular hydrolysis of organic P cleave one (monoester) or two (diester) P-O bonds. Oxygen atoms from water are integrated into the newly formed phosphate (Liang and Blake, 2006b; Liang and Blake, 2009; von Sperber *et al.*, 2014) with an enzyme-dependent kinetic fractionation.

Phosphorus radioisotopes and stable O isotopes in phosphate have rarely been considered together. An early study conducted by Larsen *et al.* (1989) showed that when phosphate

labelled with both ^{18}O and ^{32}P was added to a soil on which ryegrass was grown, the plant recovered much less ^{18}O than ^{32}P compared to what had been added. The authors concluded that the phosphate had lost its ^{18}O before plant uptake. Furthermore, these authors showed that the phosphate extracted by a resin (plant-available P) at the end of the pot experiment had a lower $^{18}\text{O}/^{32}\text{P}$ ratio compared to the added source. They suggested that the loss of the ^{18}O label associated with P was related to soil biological activity. These results were subsequently confirmed by Saaby Johansen *et al.* (1991) and Melby *et al.* (2013a) who showed that the loss of ^{18}O from ^{18}O labelled phosphate added to soils was related to biological activity. Scheerer *et al.* (2019) supplied beech roots with ^{18}O labelled phosphate and with ^{33}P labelled phosphate via addition to the nutrient solution. They could not observe a correlation between root phosphate uptake determined by incorporation of ^{33}P and by incorporation of ^{18}O . Thus, the phosphate which had been taken up had lost its ^{18}O signature due to a rapid turnover of P within roots. Pfahler *et al.* (2017) supplied soybeans with ^{33}P labelled phosphate and measured the changes in $\delta^{18}\text{O}-\text{P}$ after having stopped or not stopped P inputs. With the ^{33}P tracing, the authors quantified the transfer of P from older plant leaves to younger plant organs when P inputs to the plants were stopped. Also, they showed the occurrence of organic P hydrolysis in older leaves based on the changes in $\delta^{18}\text{O}-\text{P}$. Further, Helfenstein *et al.* (2018) assessed the rate of phosphate exchangeability by use of ^{33}P and measured $\delta^{18}\text{O}-\text{P}$ in P pools extracted sequentially from soils of the Kohala climatic gradient on the island of Hawaii. The use of ^{33}P showed that the soil present at the wettest site had the lowest amount of very slowly exchangeable P, and therefore probably a low concentration of P present in crystallized forms. The high $\delta^{18}\text{O}-\text{P}$ value observed in the HCl extractable P from the sequential extraction measured on this soil showed that the P recovered in this fraction could not be the original apatite derived from the parent material because the O associated with P had been biologically cycled either by plants and/or by microorganisms. The results obtained in the above-mentioned studies show that the use of P radioisotopes and ^{18}O can provide complementary information related to category I and II processes. However, up to now, no experiment has attempted to simultaneously assess the importance of these processes in controlling soil P dynamics.

The objective of this study was to assess whether the addition of water labelled with ^{18}O and of radioactive phosphate to soils would allow us to better understand the importance of category I and II processes during a short-term incubation and how this would improve our understanding of P transformations. This objective was reached by determining P transfer between different P pools with radioactive P (^{32}P), by assessing the incorporation of O from ^{18}O enriched water into phosphate present in different soil pools, and by relating these patterns to soil properties. The incubation was conducted for 34 days with four soils originating from the Kohala climatic gradient on the Big Island of Hawaii. These soils were chosen as they provide a strong gradient in edaphic properties while deriving from the same parent material (Chadwick *et al.*, 2003; Vitousek and Chadwick, 2013; Peay *et al.*, 2017; Helfenstein *et al.*, 2018).

2. Materials and methods

2.1. Sampling sites and soil sample collection and preparation

Soils from four sites on the Kohala climatic gradient located on the leeward slope of the Kohala Peninsula on the Big Island of Hawaii (USA) were studied (Table SM1). Soils are classified as volcanically derived andosols (Chadwick *et al.*, 2003). The soils developed on a Hawi lava flow produced 150,000 years ago (Spengler and Garcia, 1988). The lava is an alkali basalt and contains large amounts of apatite (Chadwick *et al.*, 2003). Mean annual rainfall of the sites 1 to 4 covers a wide range (275, 1578, 2163, and 3123 mm y^{-1}), whereas mean annual temperature varies in comparison on a smaller range (16.2, 17.8, 19.1, and 23.6 $^{\circ}\text{C}$) (Giambelluca *et al.*, 2013). The main vegetation is grassland dominated by either buffel grass (*Pennisetum ciliare* L.) or kikuyu grass (*Pennisetum clandestinum* Hochst. ex Chiov.) (von Sperber *et al.*, 2017a).

Soil samples were taken in February 2016 and in February 2017 from the A horizon of the four sites. The soil samples were kept at room temperature during shipment to Switzerland (one week) and were then stored immediately at -20°C until processing (13 months for February 2016 samples and one month for February 2017 samples, respectively). Soil samples of both

sampling times were unfrozen, sieved moist at 2 mm, and combined to form one composite sample per site in March 2017. Afterwards, soils were kept again at -20 °C until usage (up to two weeks). The combination of the soil samples from the two sampling times was performed to meet the soil dry weight needs of the experiment, this was required due to imposed restrictions on soil sampling and shipment. It was assumed that the soil sampling time and therefore the sample combination does not affect the strong gradient in soil properties, which was the underlying reason for the choice of the sites. However, an impact on microorganisms and available nutrient pools was expected but considered negligible in comparison to changes due to the applied conditions during the incubation experiment.

The water content of field moist soil was determined by drying for 20 hours at 105 °C. The water holding capacity (WHC) of field moist soils was determined by first weighing the soils into tared cylinders to a defined height, then saturating the soils with water by putting them in a water bath for 3 hours, and lastly allowing gravitational water to drain by placing the cylinders on a sand bath for 3.75 hours. For the pre-incubation, the water content of the soils was adjusted to 60% max. WHC by drying the soils at 28 °C (soils of sites 3 and 4) and water addition (soils of all four sites).

2.2. Experimental design

Separate soil subsamples were used for the two tracer (^{32}P and ^{18}O) and the respiration experiments. The incubation of labelled soils lasted 34 days, the respiration measurements lasted 36 days. All soils were incubated in the dark at 25 °C and 70% air humidity. After 23 days of pre-incubation at initially 60% WHC, soils were watered to adjust for evaporation losses and mixed before being assigned to either the labelling or the respiration experiments. Equivalentents of 15 g of dry soil were weighed into jars to measure soil respiration, while equivalentents of 30 g of dry soil were weighed into 250 ml plastic bottles for the tracer (^{32}P and ^{18}O) experiments (Table 1). After adding water and labelling solution, all soils were again at 60% WHC.

Location Table 1

For the ^{32}P experiment, each sample was labelled with carrier-free ^{32}P added in the form of phosphoric acid (Hartmann Analytic GmbH, Braunschweig, Germany) at a rate of 0.034 MBq g^{-1} for samples of sites 1 and 2, and 0.045 MBq g^{-1} for samples of sites 3 and 4, respectively. The rates of ^{32}P applied were based on pre-tests conducted on test samples in 2015. ^{32}P was selected due to a delivery shortage of ^{33}P . The same experiment could have been carried out with ^{33}P , as there is no significant isotopic fractionation between ^{31}P , ^{32}P , and ^{33}P (Frossard *et al.*, 2011). Four replicates were prepared for each time point and site combination by mixing the soil with the radioactive solution for two minutes in the plastic bottle. During the incubation, the samples were arranged according to a random complete block design with one replicate of each time point and site combination per block. Soils were sampled destructively at five time points (1, 4, 14, 26 and 34 days after labelling). Resin and microbial P contents and radioactive counts were measured at each time point to capture the dynamic isotopic exchange. Further explanatory analyses (total and chloroform-labile C and N, potential acid phosphatase activity, water-extractable P (P_w)) were conducted on these samples at selected time points (Table 1). This selection occurred as total and chloroform-labile C and N and potential acid phosphatase activity were expected to stay constant over the incubation time and as P_{resin} was considered a more reliable indicator for available P than P_w throughout the incubation.

For the ^{18}O experiment, half of the soil samples were destructively sampled at the incubation start (day 0) without adding an ^{18}O label. The other half of the soil samples were labelled with 2 ml of ^{18}O enriched water ($\delta^{18}\text{O}$ in labelling water = 300‰, prepared by dilution of 98% ^{18}O -enriched water from Sercon Limited, Crewe, UK) by mixing the soil with the enriched water for two minutes in the plastic bottle. This second set of soil samples was then incubated for 34 days before destructive sampling. During the incubation, the ^{18}O labelled soil samples were kept next to the ^{32}P labelled soil samples, but in a separate randomized block. Determination of $\delta^{18}\text{O}$ in soil water ($\delta^{18}\text{O}-w_s$) and in phosphate ($\delta^{18}\text{O}-P$) of sequentially extracted P pools (resin P, microbial P, inorganic and organic NaOH-EDTA P, and HCl P) was conducted at both time points (day 0 and day 34). Per time point and site combination, four replicates were prepared in plastic bottles. For the analysis of $\delta^{18}\text{O}-P$ two of the four bottles were pooled into

one combined sample to reach P amounts suitable for Ag_3PO_4 purification and analysis, resulting in two experimental replicates per site and time point combination. By using the same (volume and brand) plastic bottles and the same amounts of soil in each bottle for ^{32}P and ^{18}O soil samples, equal conditions were provided, and the results of the ^{32}P and ^{18}O experiments are comparable.

2.3. Analytical methods

2.3.1. Total and microbial carbon and nitrogen

Total soil C and N contents were measured on finely ground samples dried at 40 °C on an elemental analyser (vario PYRO, Elementar Analysensysteme GmbH, Langenselbold, Germany; temperature of combustion 950 °C). Microbial C and N were estimated by the chloroform fumigation-extraction method of Vance *et al.* (1987). Fresh samples with a dry weight equivalent of 6 g dry soil were fumigated for 24 h. Fumigated and non-fumigated subsamples were extracted with 0.5 M K_2SO_4 in a 1:4 soil:solution (mass:mass) ratio. The extracts were frozen and then analysed with a TOC/TN analyser for liquid samples (TOC-L, Shimadzu, Kyoto, Japan). The difference between fumigated and non-fumigated subsamples was calculated to estimate microbial C and N. No conversion factors (k_C and k_N) were applied to correct for inefficient fumigation by chloroform, therefore the terms chloroform-labile C (C_{chl}) and N (N_{chl}) are used instead of microbial C and N.

2.3.2. Soil respiration

Soil respiration was quantified by trapping the released CO_2 in 0.2 M NaOH and back-titrating with 0.2 M HCl (Alef, 1995). Measurements were performed at 8, 13, 20, 28 and 36 days after the day of labelling (Table 1).

2.3.3. Phosphatase activity

Potential acid phosphatase activity in soils was determined by using a 4-methylumbelliferon (MUF) substrate in a microplate assay with 6 analytical replicates following the method developed by Marx *et al.* (2001) and modified by Poll *et al.* (2006). The assay was conducted

for acidic (pH 6.1) phosphatase activity. Fluorescence was measured on a fluorescence plate reader (Biotek FLx800, Fisher Scientific GmbH, Schwerte, Germany). This analysis was performed on a formerly frozen equivalent of 1 g dry soil sampled in the ^{32}P incubation experiment at 14 days (Table 1).

2.3.4. Soil phosphorus pool concentrations

Soil P pool concentrations were assessed on moist soil samples using a modified Hedley sequential extraction (Moir and Tiessen, 2007). The following steps were performed. First three subsamples were treated in parallel: one subsample was extracted with an anion exchange resin (P_{resin}), the other was fumigated with liquid hexanol and extracted with an anion exchange resin (P_{hex}), and the last was spiked with ^{31}P and extracted with an anion exchange resin (P_{spike}). Resins were eluted in 0.1 M NaCl/0.1M HCl by shaking horizontally for 2 hours at 150 rpm. The resin extraction provides an estimation of plant-available P, the difference in P concentration between the resin extract following fumigation and without fumigation provides an estimate of microbial P, while the last sample allows quantifying the sorption of P derived from cell lysis following fumigation (Kouno *et al.*, 1995; Bünemann *et al.*, 2004). The subsample that had been fumigated with hexanol and extracted with a resin was subsequently extracted with 0.25 M NaOH/0.05 M EDTA and the extraction residue was then extracted with 1 M HCl. NaOH/EDTA is considered to extract inorganic P bound to Fe and Al oxides and organic P, while the HCl is considered to extract insoluble calcium phosphates, such as apatite (Hedley *et al.*, 1982). Each extraction step was conducted by shaking the respective soil solution horizontally for 16 hours at 150 rpm. This sequential extraction was performed before the start of the incubation on each soil with three replicates using a 1:15 soil:solution ratio (Pistocchi *et al.*, 2018).

The protocol of Tamburini *et al.* (2018) was used to analyse the $\delta^{18}\text{O}-\text{P}$ in the above-mentioned P pools. This protocol uses a 1:10 soil:solution ratio to collect sufficient P for the measurement of the $\delta^{18}\text{O}-\text{P}$. These analyses were performed before labelling and 34 days after labelling with ^{18}O enriched water in the ^{18}O experiment.

Additional extractions for P_{resin} , P_{hex} and P_{spike} and ^{32}P within these pools were performed at 1, 4, 14, 26 and 34 days after labelling with ^{32}P on the equivalent of 2 g dry soil with a 1:15 soil:solution ratio on four replicates per site in the ^{32}P experiment following the protocol of Pistocchi *et al.* (2018). Added P spikes ranged from 5 to 50 mg P kg⁻¹ during the ^{32}P experiment depending on soil and sampling time. Water extractable P (P_w) was determined on moist equivalents of 10 g dry soil from the incubation 14 days after labelling with ^{32}P for four replicates per soil. The extraction was performed by adding 100 ml H₂O (accounting for water contained in the soil), shaking overhead for 16 hours at 15 rpm, stirring for 30 min at 200 rpm and then filtering the soil solution (0.2 µm cellulose acetate membrane filter, Whatman, Merck, Darmstadt, Germany). Total P was assessed on four replicates per site on soils that had been incubated for 34 days and then air-dried and finely milled. The extraction was conducted by adding 4 ml of 15.6 M HNO₃ to 0.2 g dry soil at 250 °C for 20 min in a microwave (Turbowave, MLS GmbH, Heerbrugg, Switzerland).

The concentrations of resin extractable P (P_{resin}), inorganic NaOH-EDTA P, HCl P, total soil P, and water (P_w) were determined colourimetrically (UV-1800, Shimadzu, Canby, USA) with the malachite green method of Ohno and Zibilske (1991). Total P in NaOH-EDTA extracts was measured by ICP-OES (ICPE-9820, Shimadzu, Kyoto, Japan). Organic NaOH-EDTA P was calculated as the difference between total and inorganic P measured in the NaOH-EDTA extract. Microbial P (P_{mic} , mg P kg⁻¹ dry soil) was calculated as the difference between P_{hex} and P_{resin} subsamples including the correction for sorption (Bünemann *et al.*, 2016):

$$P_{mic} = (P_{hex} - P_{resin}) / rec \quad (1)$$

where P_{hex} (mg P kg⁻¹ dry soil) is the resin extractable inorganic P in the fumigated subsample and P_{resin} (mg P kg⁻¹) is the resin extractable inorganic P in the non-fumigated subsample, and rec is the recovery of added P spike calculated as:

$$rec = (P_{spike} - P_{resin}) / \text{added P} \quad (2)$$

with P_{spike} (mg P kg⁻¹ dry soil) being the resin extractable inorganic P in the subsample to which the P spike had been added. If recovery of added P was above 1, which happened five times

during the ^{32}P experiment (four times for site 1, once for site 2), *rec* was set to 1. For the sequential P extraction before the start of the incubation *rec* was set to 1 for all sites due to amounts of spike P added being not close to actual P_{mic} values resulting in unrealistically low or high *rec* values. No conversion factor (k_P) was applied to correct microbial P for inefficient fumigation by hexanol, as k_P is soil-specific (Oberson and Joner, 2005) and was not determined for the studied soils. Thus, microbial P might be underestimated. Microbial P was found three times to be negative (each time in soils from site 1), in such cases P_{mic} was set to zero.

2.3.5. Radioisotope tracing in soil phosphorus pools

2.3.5.1. Radioactivity in phosphorus pools

Radioactivity in the resin extractable pool (r_{resin}) was directly measured by scintillation counting on a beta-emission counter (Tri-carb 2500 TR, Packard Instruments, Meriden, USA) using 1 ml of sample and 5 ml Ultima GoldTM (Perkin Elmer, USA).

The determination of radioactivity in microbial biomass (r_{mic}) was done after having corrected the release of ^{32}P from the solid phase of the soil caused by the release of ^{31}P from the microbial P pool (Bünemann *et al.*, 2016; Schneider *et al.*, 2017). First, a linear correlation was built between P_{hex} and r_{spike} (the amount of radioactivity recovered in the sample spiked with ^{31}P in Bq g^{-1} dry soil) which is the corrected radioactivity in spiked subsamples

$$r_{spike} = a * P_{hex} + b \quad (3)$$

where *a* and *b* are the slope and intercept of the linear regression.

And then r_{spike} was introduced in equation 4 to obtain r_{mic} (Bq g^{-1} dry soil)

$$r_{mic} = (r_{hex} - r_{spike}) / rec \quad (4)$$

This approach assumes that the recovery of a ^{31}P spike (*rec*) is similar to the recovery of a ^{32}P spike as shown by Bünemann *et al.* (2012), and therefore that it is not necessary to measure this last parameter.

Negative values were observed 18 times for r_{mic} , 12 of these negative values were recorded for the soils of site 1. These values were probably caused by low microbial P concentrations (Table 2) and lower soil microbial activity (Figure 2) combined with propagated measurement uncertainties. Thus, negative values were considered as missing values. As a large fraction of the r_{mic} values of site 1 were missing, we do not discuss r_{mic} in the soils of site 1 and focus on the more reliable complementary data.

2.3.5.2. Radioactivity recovery and specific activity of phosphorus pools

Radioactivity recovery (r/R_{pool} , %) was calculated for each analysed pool by relating the measured radioactivity r in the pool at a given sampling time (r_{pool} Bq g⁻¹ dry soil) to the decay-adjusted introduced radioactivity R (Bq g⁻¹ dry soil, half-life $t_{1/2} = 14.28$ days for ³²P):

$$r/R_{pool} = (r_{pool} / R) * 100 \quad (5)$$

Instead of the commonly used specific activity (SA , Bq mg⁻¹ P), the proportional specific activity in P pools (pSA_{pool} , % kg dry soil mg⁻¹ P) was calculated as follows:

$$pSA_{pool} = (r/R_{pool} / P_{pool}) \quad (6)$$

where r/R_{pool} is the fraction of recovered radioactivity in the P pool (%) and P_{pool} is the P concentration of the P pool (mg P kg⁻¹ dry soil). This modification of the calculation was performed to account for the variable amount of introduced ³²P (R) and thus to enable the comparison between sites.

2.3.5.3. Isotopically exchangeable phosphate

Isotopically exchangeable phosphate was calculated to assess the fluxes of P between the solid phase of the soil and the solution. Since the measurements of P_w were very variable and below the detection limit in the soils of the wetter site 4, we decided to use the resin P pool for the calculation of the isotopically exchangeable phosphate as proposed by Maertens *et al.* (2004). The amount of isotopically exchangeable resin P (E_{resin} , mg P kg⁻¹ dry soil) was calculated according to the following equation:

$$E_{resin} = 100/pSA_{resin} \quad (7)$$

where pSA_{resin} is the proportional specific activity in the resin extract.

2.3.5.4. Estimation of gross phosphorus immobilisation

If we consider that microorganisms take up most of their P from the resin P pool then it becomes possible, using pSA_{resin} and r/R_{mic} , to calculate the amount of microbial P that is derived from the resin P pool (P_{immo} , mg P kg⁻¹ dry soil) using the following equation:

$$P_{immo} = (r/R_{mic}) / pSA_{resin} \quad (8)$$

Equation 8 provides an estimation of the gross P immobilisation (Bünemann *et al.*, 2012). The soils from site 1 were not considered for this calculation.

2.3.6. Oxygen isotopic signatures

2.3.6.1. Oxygen isotopic signatures in soil water and phosphate

Soil water was extracted by cryogenic vacuum extraction (Orlowski *et al.*, 2013) from one replicate per soil of each site before and 34 days after labelling with ¹⁸O enriched water.

Phosphate in the resin, hexanol and HCl extracts was purified following the protocol given by Tamburini *et al.* (2010). For NaOH-EDTA extracts, separation into organic and inorganic P fractions and their respective purification were performed following Tamburini *et al.* (2018). Inorganic NaOH-EDTA extracts were processed for sites 1, 2 and 4, while organic ones were processed only for sites 2 and 4. These samples were selected because we expected differences to occur between the extreme sites 1 and 4. As site 1 had not enough P in the organic NaOH-EDTA extract, site 2 was also considered. The $\delta^{18}O-P$ of inorganic NaOH-EDTA P of site 4 was considerably lower at 34 days compared to before labelling with ¹⁸O enriched water, suggesting a processing error. Because the analysis could not be repeated due to limited soil availability, the results for site 4 are not shown.

Purified Ag₃PO₄ was analysed in one to three analytical replicates for $\delta^{18}O$ on a thermal conversion elemental analyser (vario PYRO cube, Elementar Analysensysteme GmbH, Langenselbold, Germany) in pyrolysis mode (temperature set at 1450 °C), coupled in a

continuous flow to an IsoPrime 100 isotopic ratio mass spectrometer (IRMS). An internal Ag_3PO_4 standard (Acros Organics, Geel, Belgium, $\delta^{18}\text{O} = 14.2\text{‰}$), two benzoic acid standards distributed by the International Atomic Energy Agency (IAEA) (IAEA 601: $\delta^{18}\text{O} = 23.1\text{‰}$, IAEA 602 $\delta^{18}\text{O} = 71.3\text{‰}$) and in-house made standards were used for instrumental drift correction and calibration. Oxygen isotopic signatures are presented in the $\delta^{18}\text{O}$ notation as ‰ deviation values related to the VSMOW (Vienna Standard Mean Oceanic Water).

2.3.6.2. Calculation of equilibrium oxygen isotopic signature

The expected $\delta^{18}\text{O}$ signature of phosphate in equilibrium with soil water ($Eq-w_s$) can be calculated with the following equation (Chang and Blake, 2015):

$$Eq-w_s = e^{(14.43/T-26.54/1000)} * (\delta^{18}\text{O}-w_s + 1000) - 1000 \quad (9)$$

where T is the temperature in K during the incubation and $\delta^{18}\text{O}-w_s$ (‰) is the O isotopic signature of soil extracted water.

Since temperature-dependent equilibration is mainly occurring intracellularly, the proportion of water, produced during metabolic reactions, and its $\delta^{18}\text{O}$ should be considered for the equilibrium estimation. The contribution of metabolic water to intracellular water depends on the metabolic state of the organism. It reached up to 70% for *Escherichia coli* during active growth, whereas the contribution in a stationary phase was considerably smaller (Kreuzer-Martin et al., 2005).

Our estimation of the equilibrium considering soil water and metabolic water ($Eq-w_{s+m}$) was based on two assumptions. First, that intracellular water ($\delta^{18}\text{O}-w_{s+m}$) consists of 59% soil water and 41% metabolic water, as determined by Li et al. (2016) due to the fractionation between $\delta^{18}\text{O}$ of microbial biomass and extracellular water. Second, that $\delta^{18}\text{O}$ of metabolic water is -3.5‰ (Kreuzer-Martin et al., 2005). This results in the following equation:

$$\delta^{18}\text{O}-w_{s+m} = 0.59 * \delta^{18}\text{O}-w_s + 0.41 * (-3.5) \quad (10)$$

where $\delta^{18}\text{O}-w_s$ (‰) is the measured O isotopic signature of soil water, which is accounting for 59% of water within biomass, and -3.5 ‰ is the $\delta^{18}\text{O}$ of metabolic water accounting for 41% of water within biomass (Kreuzer-Martin *et al.*, 2005; Li *et al.*, 2016).

2.3.6.3. Calculation of oxygen isotopic signature in phosphate ($\delta^{18}\text{O}-P$)

Most $\delta^{18}\text{O}-P$ values per individual sample were calculated from three analytical replicates of purified Ag_3PO_4 measured from this sample. For the organic NaOH-EDTA P extracts, the incorporation of ^{18}O during UV-digestion was negligible and therefore the subsamples (non-labelled and spiked with ^{18}O enriched water during UV hydrolysis, see Tamburini *et al.* (2018)) were considered as individual samples for the calculations of mean and corrected sample standard deviation per pool.

The $\delta^{18}\text{O}-P_{mic}$, the O isotopic composition of phosphate in the microbial P pool, was calculated with a mass balance equation (Tamburini *et al.*, 2012):

$$\delta^{18}\text{O}-P_{mic} = (\delta^{18}\text{O}-P_{hex} * P_{hex} - \delta^{18}\text{O}-P_{resin} * P_{resin}) / (P_{hex} - P_{resin}) \quad (11)$$

where $\delta^{18}\text{O}-P_{hex}$ (‰) and $\delta^{18}\text{O}-P_{resin}$ (‰) are the O isotopic compositions of phosphate in the hexanol and resin P pools, respectively and P_{hex} (mg P kg⁻¹ dry soil) and P_{resin} (mg P kg⁻¹ dry soil) are the P concentrations in the hexanol and resin P pools of the respective sample. For sites 1 and 2, $\delta^{18}\text{O}-P$ of the hexanol and resin P pool were too close to each other to allow a reliable mass balance. Thus, $\delta^{18}\text{O}-P_{hex}$ was reported instead.

During the HCl extraction, we used ^{18}O labelled and unlabelled reagents to track possible inorganic hydrolysis related for instance to the presence of pyro- or polyphosphates (Tamburini *et al.*, 2010). If hydrolysis was observed, then the actual $\delta^{18}\text{O}-P_{HCl}$ was calculated according to Pistocchi *et al.* (2017):

$$\delta^{18}\text{O}-P_{HCl} = (\delta^{18}\text{O}-P_l * \delta^{18}\text{O}-w_{nl} - \delta^{18}\text{O}-P_{nl} * \delta^{18}\text{O}-w_l) / (\delta^{18}\text{O}-P_l - \delta^{18}\text{O}-P_{nl} - \delta^{18}\text{O}-w_l + \delta^{18}\text{O}-w_{nl}) \quad (12)$$

where $\delta^{18}\text{O}-P_l$ (‰) and $\delta^{18}\text{O}-P_{nl}$ (‰) are the O isotopic composition of phosphate for the labelled and unlabelled HCl samples, respectively, and accordingly $\delta^{18}\text{O}-W_l$ (‰) and $\delta^{18}\text{O}-W_{nl}$ (‰) the O isotopic signature of water used to prepare labelled and non-labelled reagents.

$\delta^{18}\text{O}-P_{mic}$ and $\delta^{18}\text{O}-P_{HCl}$ per individual sample were obtained with the above-mentioned equations 11 and 12. The means and standard deviations of $\delta^{18}\text{O}-P$ per pool and time point were calculated based on the values of the two individual samples. If no replicate was available ($\delta^{18}\text{O}-P_{resin}$ and $\delta^{18}\text{O}-P_{mic}$ site 4 day 0, $\delta^{18}\text{O}-P$ of inorganic NaOH-EDTA P sites 1 and 2 at both time points, and $\delta^{18}\text{O}-P_{HCl}$ site 1 day 0), the value of the individual sample was reported and no standard deviation was determined.

2.3.6.4. Incorporation of ^{18}O into phosphate

Incorporation of ^{18}O into phosphate of a specific P pool (%) was calculated based on the differences in $\delta^{18}\text{O}$ between time points for soil water and phosphate as following (Liang and Blake, 2006a):

$$^{18}\text{O} \text{ incorporation} = (\delta^{18}\text{O}-P_{end} - \delta^{18}\text{O}-P_{start}) / (\delta^{18}\text{O}-W_{s-end} - \delta^{18}\text{O}-W_{s-start}) * 100 \quad (13)$$

Where $\delta^{18}\text{O}-W_s$ is O isotopic signature of soil water and $\delta^{18}\text{O}-P$ is O isotopic signature of phosphate in the P pool at the start (day 0) and at the end of the incubation experiment (day 34). The calculations were conducted by Monte Carlo error propagation (Anderson, 1976) implementing 10 million iterations varying the $\delta^{18}\text{O}$ signatures according to mean and corrected sample standard deviation. For $\delta^{18}\text{O}-W_{s-end}$ and $\delta^{18}\text{O}-W_{s-start}$ standard deviations of 0.01‰ were assumed (Tamburini *et al.*, 2018).

2.4. Statistical analyses

Statistical analyses were conducted in R version 3.6.0 (R Core Team, 2019). Data residuals were tested for normal distribution and homogenous variance. Some variables were log (chloroform-labile C and total N concentrations, r/R_{resin} of site 2, E_{resin} at day 34, pSA_{resin} of site 4), square root (chloroform-labile N, P_w , P_{mic} within Hedley and total C concentrations, r/R_{mic} , P_{immo}) or inverse (total soil respiration, pSA_{resin}) transformed to meet assumptions of normality

and homogeneity of variances. Analysis of variance (ANOVA) was conducted using *aov* (package *stats* version 3.6.0, R Core Team (2019)). To test site differences, the ANOVA model included site and block effect (data~site+block). The block effect refers to the randomized complete block design sample set-up. To test temporal trends, analyses were conducted separately per site with sampling time point and block effect included in the ANOVA model (data~time.point+block). Afterwards, Tukey's HSD (honest significant difference) post hoc tests for confidence levels 0.95 were performed with *HSD.test* (package *agricolae* version 1.3-1, de Mendiburu (2019)). Figures were produced using the package *ggplot2* (version 3.2.0, Wickham (2009)).

3. Results

3.1. Soil properties

The soil properties varied significantly between sites (Table 2). Water holding capacity, total C and N, chloroform-labile C and N, P_{mic} , organic NaOH-EDTA P and acid phosphatase activity increased from site 1 to 4. In contrast, P_w , P_{resin} and inorganic NaOH-EDTA P decreased from site 1 to 4.

Location Table 2

3.2. Soil respiration

Total soil respiration over 36 days was ten times larger for site 4 (2500 mg C kg⁻¹) compared to site 1 (259 mg C kg⁻¹), with intermediate sites ranging in between and all values being significantly different from each other (Figure 2b). The proportion of total C that was lost by respiration reached 1.6% in the soil of site 1, 1% in site 4, 0.9% in site 2 and 0.8% in site 3. Daily soil respiration rates decreased during the incubation period, most notably for the soil of site 4 with a decrease from 99.0 to 72.8 mg C kg⁻¹ d⁻¹ (Figure 2a).

Location Figure 2

3.3. Phosphorus pools and the fate of radioactive phosphate

3.3.1. Resin phosphorus

Resin P (Figure 3a) concentrations (P_{resin} , mg P kg⁻¹ dry soil) stayed constant in the soil of sites 3 and 4, while they varied without clear trend for soils of sites 1 and 2. Radioactivity recovery in the resin pool (r/R_{resin} , %) on the contrary, strongly decreased with time in all soils. r/R_{resin} varied between 9% (soil of site 3) and 37% (soil of site 1) after 1 day of incubation and between 3% (soils of sites 3 and 4) and 15% (soil of site 1) after 34 days (Figure 3b). r/R_{resin} was always smaller in the soils from the wet compared to the dry sites.

Location Figure 3

Largest amounts of isotopically exchangeable P in the resin extract (E_{resin} , Figure 4) were observed for sites 3 and 2, intermediate amounts for site 2, and the lowest amounts for site 4. The amounts increased over time for all sites. Each of these curves could be described with the equation $E_{resin(t)} = E_{resin(1)} * t^n$ (with t being the incubation time expressed in days, $E_{resin(1)}$ the amount of P isotopically exchangeable after one day, $E_{resin(t)}$ the amount of P isotopically exchangeable after t days and n an empirical parameter) reaching r^2 values higher than 0.95.

Location Figure 4

3.3.2. Microbial phosphorus

Microbial P concentrations (Figure 5a, P_{mic} , mg P kg⁻¹ dry soil) remained constant in the soil of sites 3 and 4 and varied without clear trend for sites 1 and 2. The lowest P_{mic} concentrations were observed in the soil of site 1, and the highest in the soil of site 4. Radioactivity recovery in the microbial biomass (r/R_{mic} , %) (Figure 5b) was low (<10%) and variable in the soils of all sites. It was often not possible to measure r/R_{mic} in the soil of site 1 due to the very small size of the microbial P pool. The radioactive data of microbial P of the soil from site 1 are therefore not discussed. A significant decrease over time was observed in the soil of site 2 with r/R_{mic} ranging from 1 to 0.2% of the added ³²P.

A high proportion of introduced ^{32}P was not detected in either the resin or the microbial P pool already after 1 day of incubation (from 62.0% for site 1 to 89.3% for site 3), indicating a fast disappearance of this tracer towards other P pools.

Location Figure 5

Assuming that microorganisms took up their P from the resin P pool and knowing the specific activity of the resin and microbial pools, we could calculate the amount of P in the microbial biomass derived from the resin pool, (Figure 6, P_{imm} , mg P kg⁻¹ dry soil), which is an estimation for the gross P immobilisation. This amount remained low and variable in the soil of site 2 (below 5 mg P kg⁻¹ dry soil), while it increased significantly with time in the soils of sites 3 and 4 reaching 20 mg P kg⁻¹ dry soil at the last sampling point of the soil of site 4. Gross P immobilisation ranged between 2% of microbial P for the first sampling time point of site 4 (2.0 mg P kg⁻¹ for P_{imm} compared to 109.5 mg P kg⁻¹ for P_{mic}) and 27% for the sampling time point at 14 days of site 3 (15.6 mg P kg⁻¹ for P_{imm} compared to 58.5 mg P kg⁻¹ for P_{mic}).

Location Figure 6

3.4. Oxygen isotopes

3.4.1. Oxygen isotopic signatures

The addition of ^{18}O enriched water resulted in increased O isotopic signatures of soil water ($\delta^{18}\text{O}-w_s$) after 34 days of incubation for all sites (Table 3). Consequently, the calculated equilibrium values between ^{18}O in water and ^{18}O in phosphate were between 1.4 to 4 times higher at the final sampling point compared to the value calculated at day 0 of the incubation. Taking the metabolic water into consideration led to lower equilibrium values.

Location Table 3

The $\delta^{18}\text{O}-P_{resin}$ and $\delta^{18}\text{O}-P_{mic}$ of the soils of each site increased by several permil during the incubation, while the other P pools did not show any increase in $\delta^{18}\text{O}-P$ (Table 3). The $\delta^{18}\text{O}-P$ of none of the P pools reached equilibrium with $\delta^{18}\text{O}-w_s$ or with $\delta^{18}\text{O}-w_{s+m}$ during the incubation.

3.4.2. Incorporation of ^{18}O into soil phosphorus pools

Incorporation of ^{18}O from enriched water into phosphate was observable in the resin and microbial P pools, but not in inorganic or organic NaOH-EDTA P or HCl P (Table 4). In the resin and microbial P pools, approximately 1 out of 4 O atoms per phosphate was exchanged during the incubation in the soils of sites 1 to 3. The soil of the wettest site showed higher exchange rates than the other sites, particularly in resin P (41% ^{18}O incorporation).

Location Table 4

4. Discussion

In the introduction, we considered three types of processes controlling P dynamics in soils: those that maintain the bond between P and O and transfer phosphate from one pool to another (category I processes), those that involve cleavage of the P-O bond and transfer phosphate from one pool to another (category IIa processes), and those that involve cleavage of the P-O bond but do not transfer phosphate from one pool to another (category IIb processes). We discuss how far the use of P and O isotopes allows to assess these three types of processes and their roles in soil P dynamics.

4.1. Category I pathways: phosphates move as intact units

The use of ^{32}P allowed assessing the amount of soil P that was isotopically exchanged over the time of incubation (E_{resin}). The fact that the resin P pool size did not vary significantly with time in each soil, while the fraction of radioactive P in the resin extracts significantly decreased, confirms that the main mechanism controlling ^{32}P losses from the resin P pool was isotopic exchange rather than a net change in pool size. Soil organic P mineralisation could have contributed to the dilution of the added ^{32}P , but it was probably not a dominant mechanism. If we assume that C and P are stoichiometrically mineralised, the amount of organic P mineralised during the incubation would be equal to the ratio between the amount of C mineralised during the incubation to total C multiplied by organic P (Achat *et al.*, 2009). The measured organic NaOH-EDTA P was set as organic P. The amount of mineralised C was

derived from C emitted as CO₂ corrected by the microbial C assimilation efficiency, which was assumed to be 0.4 according to Murphy *et al.* (2003). This calculation suggests that the amount of mineralised organic P would range between 14 for the soil of site 1 and 41 mg P kg⁻¹ dry soil for the soil of site 4, ergo would reach values smaller or equal to the standard deviations observed for E_{resin} after 34 days of incubation. Moreover, the changes in E_{resin} with time can be described for each soil with a simple log/log relation ($E_{resin(t)} = E_{resin(1)} * t^n$), which is similar to the equation used by Fardeau and Zapata (2002) to describe the changes in soil isotopically exchangeable P with time. Isotopic exchange is controlled both by sorption and desorption and by transfer of P from the charged surface to the bulk solution by diffusion and can, therefore, be assigned to the category I processes. The large amounts of soil P isotopically exchangeable with resin P show the importance of category I processes in these soils. Maertens *et al.* (2004) observed the same type of relationship between E_{resin} and incubation time. However, in their work, the increase of E_{resin} with time was several-fold smaller compared to our results. This difference is probably due to the difference in soil types and the difference in total P content as Maertens *et al.* (2004) presented results on a lixisol and a ferralsol from Kenya, which contained less amorphous minerals and less total P than our andosols. The high rate of P exchangeability found in our andosols confirms the results of Helfenstein *et al.* (2018) and is consistent with the presence of high amounts of amorphous minerals in these soils (Chadwick *et al.*, 2003).

The use of ³²P allowed the quantification of the amount of P derived from the resin pool that was taken up by microorganisms (gross P immobilisation, P_{immo}). These microbial processes probably involved pathways in which phosphate was taken up as a unit (category I processes) and processes during which an O-P bond was cleaved (category IIa and IIb processes), for example, if P was taken up after extracellular enzymatic hydrolysis. If we do not consider the soil from site 1 on which a substantial amount of data was missing due to the very low microbial P content, the amount of P derived from the resin incorporated in the microbial biomass was significantly larger for the soils of sites 3 and 4 than for the soil of site 2, and this amount increased with time for the soils of sites 3 and 4. The highest P uptake rate observed in the

soil of site 4 was in agreement with the highest rate of CO₂ emission from this soil and the high microbial biomass. Bünemann *et al.* (2012) and Pistocchi *et al.* (2018) also used P radioisotopes for the estimation of gross P immobilisation rates during incubation experiments. Contrary to our approach, their calculations are based on the changes in specific activity in water extracts and not the changes in specific activity in resin extracts. Nevertheless, they observed higher gross P immobilisation (between 40 and 50 mg P kg⁻¹ soil after 11 days in Pistocchi *et al.* (2018) and between 36 and 49 mg P kg⁻¹ soil after 32 days in Bünemann *et al.* (2012)) compared to our study (between 5 and 20 mg P kg⁻¹ soil after 34 days). Since we have not used a correction factor (k_P) for calculating microbial P content, the fluxes of P through the microbial biomass are underestimated. Nevertheless, even if we applied a k_P factor of 0.8 as in Olander and Vitousek (2005), the resulting values for gross P immobilisation after 34 days of incubation would remain well below the E_{resin} values. For example, for the soil of site 4 k_P corrected gross P immobilisation would only reach 9% of the respective E_{resin} .

Our results contrast with those presented by Olander and Vitousek (2005), who showed that microbial control on P uptake from the solution was at least as strong as the sorption control in a Hawaiian chronosequence after 48 hours of incubation. But, these results are difficult to compare to ours because i) they estimated available P with the Bray I extractant while we used anion exchange resins, and ii) they added 10 µg P g⁻¹ soil which was a significant input compared to the low amounts of P extracted by the Bray I method from the non-treated soils (smaller than 4.5 µg P g⁻¹). We added no ³¹P with the ³²P to minimise changes in chemical equilibrium between the soil and the solution. However, based on the assumption that P extracted by the Bray I method and the resin should target a similar P pool (van Raij *et al.*, 2009), we assume that the soils studied in our work (27.6 to 69.3 mg resin P kg⁻¹, Table 2) had a higher P availability than those studied by Olander and Vitousek (2005) (0.27 to 4.32 mg Bray I P kg⁻¹). Our results together with those presented by Olander and Vitousek (2005), by Schneider *et al.* (2017), and by Pistocchi *et al.* (2017) suggest that the proportion of P cycling through the microbial biomass increases with decreasing P availability.

4.2. Category II pathways: the bond between oxygen and phosphate atoms is cleaved

The $\delta^{18}\text{O}-P$ of the different P pools before labelling with ^{18}O enriched water (Table 3) were resembling those found for these soils by Helfenstein *et al.* (2018).

Although the $\delta^{18}\text{O}-P_{mic}$ had increased during the 34 days of incubation, in none of the soils equilibrium with the $\delta^{18}\text{O}-w_s$ was reached, even when the metabolic water was considered in the calculation of the equilibrium values. This increase shows that ^{18}O had been incorporated into microbial P and that the P-O bond of phosphate had been broken. Thus, phosphate had undergone processes of categories IIa and/or IIb. But the fact that the equilibrium was not reached shows also that the O bound to P in the microbial biomass had not yet been entirely turned over (category IIb process). A complete turnover of O in phosphate in the presence of pyrophosphatase and Mg was observed within a few hours in sterile systems (Chang and Blake, 2015; von Sperber *et al.*, 2017b). However, it is probable that in a heterogeneous medium like soil, within which different types of microorganisms cohabit (Peay *et al.*, 2017), a longer time will be required for O atoms of phosphate to equilibrate with the O atoms of water present within the soil microorganisms. At the end of incubation, in our soils, the mean incorporation of ^{18}O into microbial P was non-linearly correlated with the mean amount of C emitted as CO_2 ($r^2 = 0.919$). This agrees with Middelboe and Saaby Johansen (1992) and Melby *et al.* (2013a), who showed that the de-labelling of ^{18}O from ^{18}O labelled PO_4 was correlated to soil microbial activity.

The resin and microbial P pools had similar $\delta^{18}\text{O}-P$ signatures in the soils of the sites 1, 2 and 3 at day 0 and at the end of the incubation, whereas in the soil of site 4 the $\delta^{18}\text{O}-P_{mic}$ was lower than $\delta^{18}\text{O}-P_{resin}$ at both time points. This suggests a constant and rapid turnover of P between the resin and the microbial pool for the soils of sites 1, 2, and 3. The resin and inorganic NaOH-EDTA P pools were expected to show similar $\delta^{18}\text{O}-P$ signatures, as NaOH P exchanges relatively rapidly with the P in solution, except in systems that contain P in crystallized minerals such as vivianite (Frossard *et al.*, 1996; Helfenstein *et al.*, 2018). However, this was not the

case for the soils of sites 1 and 2. The differences can be possibly explained by i) the fact that 34 days were probably not sufficient for complete equilibration between the different inorganic P pools, and ii) the presence of different forms of inorganic P species extracted by NaOH-EDTA. The first explanation agrees with the fact that only 21 to 56% of the inorganic NaOH-EDTA P was exchangeable after 34 days of incubation. In accordance, the estimated mean residence time of P in the NaOH P pool of Andosols is around six months, substantially longer than the duration of our incubation experiment (Helfenstein *et al.*, 2020). More information on the inorganic P species extracted by NaOH-EDTA would be necessary to provide support for the second explanation.

The soil of site 4 behaved differently compared to the other soils as its $\delta^{18}\text{O}-P_{mic}$ was much lower than its $\delta^{18}\text{O}-P_{resin}$ and a higher proportion of ^{18}O was incorporated in the phosphates of the microbial and of the resin pools. The high rate of incorporation of ^{18}O in the microbial pool reflects the high microbial activity (shown also by the high CO_2 emission and high P uptake by microbes from the resin pool). The lower and far from equilibrium $\delta^{18}\text{O}-P_{mic}$ compared to $\delta^{18}\text{O}-P_{resin}$ seems to suggest the presence of different groups of microorganisms, all of them extractable by hexanol. The first group of organisms is active in uptake and release of phosphate to the soil solution and exchanges O between phosphate and water. The second group, on the other hand, is probably in a more dormant status and characterised by a lower $\delta^{18}\text{O}-P$. This group interacts less with the soil solution and does not actively exchange O between phosphate and water (Chen *et al.*, 2019).

5. Conclusions

Radioactive P is an established tracer to assess the P fluxes from one soil compartment to another, whereas changes in the isotopic composition of O associated with phosphate are investigated to provide information on the impact of phosphatase activity on soil P dynamics. In this paper, we showed that the use of ^{32}P and ^{18}O in parallel experiments generated results that reinforced each other. For instance, the fact that at the end of the incubation only 21 to 56% of the NaOH-EDTA P was measured as isotopically exchangeable agreed with the fact

that the $\delta^{18}\text{O}-\text{P}$ of the resin pool was not equal to the $\delta^{18}\text{O}-\text{P}$ of the NaOH-EDTA pool. With a longer incubation time, we might reach 100% exchangeability in the NaOH-EDTA P pool and thus, the $\delta^{18}\text{O}-\text{P}$ of this pool and of the resin pool would have approached similar values. Furthermore, the use of ^{32}P allowed to quantify the fraction of P derived from the resin pool and taken up by microorganisms, while changes of the $\delta^{18}\text{O}-\text{P}$ in microorganisms allowed to detect the incorporation of O in phosphate within microorganisms, both aspects being linked to microbial activity. In two cases the $\delta^{18}\text{O}-\text{P}$ allowed to follow processes that were not observable with ^{32}P : i) the incorporation of ^{18}O in the resin P pool suggested that either some of this P had been released by enzymatic hydrolysis from organic P or that it has been released from microorganisms in which the O associated with P had been turned over, and ii) the difference between $\delta^{18}\text{O}-\text{P}$ of the resin and the microbial pool at the end of the incubation in the soil of site 4 suggested that this soil could contain groups of microorganisms exhibiting different levels of activities. Further microbiological analyses would be needed to support this last hypothesis. In any case, the results of this study confirm preceding research which showed that $\delta^{18}\text{O}-\text{P}$ cannot be used to quantify the transfer of P from one pool to the next. Nevertheless, we conclude that the two tracers provide complementary information that helps to better understand the soil P dynamics.

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Tables

Table 1: Overview of the experimental schedule during the incubation.

	fresh	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36		
Soil labelling		x																																						
Total and chloroform-labile C and N						x										x																							x	
Soil respiration										x					x						x								x										x	
Phosphatase activity																x																								
P_{conc} of P_w																x																								
P_{conc} and radioactivity of P_{resin} and P_{mic}				x		x										x												x											x	
P_{conc} of Hedley P pools		x																																						
$\delta^{18}O$ of Hedley P pools and soil water		x																																						x

Table 2: Selected properties of studied soils: water holding capacity (WHC); nutrient (carbon, nitrogen and phosphorus) concentrations; pH and acid potential phosphatase activities. Means and corrected sample standard deviations per site, different letters denote significant differences ($P \leq 0.05$) among sites.

	Unit	Site 1	Site 2	Site 3	Site 4
Maximal water holding capacity (WHC) [#]	g H ₂ O g ⁻¹	0.62	1.18	2.24	3.36
Chloroform-labile C (<i>C_{chl}</i>) ^{###}	mg C kg ⁻¹	90 ± 24 ^d	228 ± 40 ^c	416 ± 85 ^b	769 ± 101 ^a
Total C ^{##}	g C kg ⁻¹	16 ± 1 ^d	63 ± 1 ^c	170 ± 3 ^b	239 ± 3 ^a
Chloroform-labile N (<i>N_{chl}</i>) ^{###}	mg N kg ⁻¹	10 ± 5 ^c	43 ± 19 ^b	40 ± 21 ^b	107 ± 41 ^a
Total N ^{##}	g N kg ⁻¹	1.5 ± 0.0 ^d	6.1 ± 0.1 ^c	12.8 ± 0.2 ^b	18.6 ± 0.3 ^a
Water extractable P (<i>P_w</i>) ^{#####}	mg P kg ⁻¹	4.72 ± 0.29 ^a	1.74 ± 0.43 ^b	0.10 ± 0.03 ^c	0.23 ± 0.22 ^c
Resin P (<i>P_{resin}</i>) ^{###}	mg P kg ⁻¹	69.3 ± 3.7 ^a	44.9 ± 1.5 ^b	26.1 ± 0.6 ^c	27.6 ± 2.8 ^c
Microbial P (<i>P_{mic}</i>) ^{####}	mg P kg ⁻¹	0.9 ± 1.0 ^d	13.8 ± 2.2 ^c	50.2 ± 6.4 ^b	93.7 ± 15.3 ^a
Inorganic NaOH-EDTA P ^{###}	g P kg ⁻¹	2.23 ± 0.09 ^a	2.64 ± 0.27 ^a	1.37 ± 0.05 ^b	0.56 ± 0.05 ^c
Organic NaOH-EDTA P ^{###}	g P kg ⁻¹	0.52 ± 0.12 ^c	1.86 ± 0.23 ^b	2.64 ± 0.23 ^b	2.37 ± 0.20 ^{ab}
HCl P ^{###}	g P kg ⁻¹	1.95 ± 0.08 ^a	0.49 ± 0.12 ^c	1.76 ± 0.05 ^b	0.64 ± 0.05 ^c
Total P ^{#####}	g P kg ⁻¹	5.34 ± 0.67 ^a	4.92 ± 0.35 ^a	4.99 ± 0.40 ^a	3.52 ± 0.12 ^b
pH [§]		6.7	5.9	5.4	4.2
Potential acid phosphatase activity ^{#####}	μmol g ⁻¹ h ⁻¹	2.2 ± 0.1 ^d	9.0 ± 0.1 ^c	16.0 ± 0.5 ^b	19.7 ± 1.8 ^a

n=1, ## n=12, ### n=3, #### n=4 replicates per site

§ from Helfenstein *et al.* (2018) analysed on soils sampled in February 2016

Table 3: Oxygen isotopic signatures of soil water ($\delta^{18}\text{O}-w_s$) and of phosphate of soil phosphorus pools ($\delta^{18}\text{O}-P$ of resin, microbial, organic and inorganic NaOH-EDTA and HCl P pools) as well as calculated equilibrium values ($Eq-w_s$ and $Eq-w_{s+m}$) before (day 0) and 34 days after labelling with ^{18}O enriched water. Means and corrected sample standard deviations per site and time point.

	Time [days]	$\delta^{18}\text{O}$ [‰]			
		Site 1	Site 2	Site 3	Site 4
Soil water	0	-3.5 [#]	-2.2 [#]	-1.3 [#]	0.6 [#]
	34	50.9 [#]	26.7 [#]	19.5 [#]	13.2 [#]
Resin	0	22.8 ± 0.2	19.6 ± 0.9	15.5 ± 3.6	15.7 [#]
	34	33.8 ± 0.7	26.5 ± 2.1	19.3 ± 0.4	20.9 ± 1.5
Microbial	0	22.1 ± 0.6 ^H	18.6 ± 0.2 ^H	16.6 ± 2.0	11.5 [#]
	34	32.7 ± 2.6 ^H	25.5 ± 1.4 ^H	20.9 ± 0.4	15.6 ± 2.4
Organic NaOH-EDTA	0	NA	18.6 ± 0.5	NA	21.4 ± 0.5
	34	NA	20.3 ± 0.1	NA	21.5 ± 0.6
Inorganic NaOH-EDTA	0	26.4 [#]	23.7 [#]	NA	NA
	34	27.6 [#]	23.9 [#]	NA	NA
HCl	0	15.6 [#]	25.7 ± 0.6	24.6 ± 1.0	23.4 ± 0.7
	34	14.7 ± 0.3	23.8 ± 2.8	23.1 ± 1.1	23.2 ± 0.4
Equilibrium $Eq-w_s$ (soil water)	0	18.5 [#]	19.8 [#]	20.7 [#]	22.7 [#]
	34	74.1 [#]	49.4 [#]	42.0 [#]	35.6 [#]
Equilibrium $Eq-w_{s+m}$ (soil water and metabolic water) [§]	0	18.5 [#]	19.3 [#]	19.8 [#]	21.0 [#]
	34	51.3 [#]	36.7 [#]	32.4 [#]	28.6 [#]

[#] n=1 (standard deviation not determinable), otherwise n=2
^H $\delta^{18}\text{O}-P_{hex}$ values reported instead of $\delta^{18}\text{O}-P_{mic}$
[§] assumes 41% of metabolic water with given $\delta^{18}\text{O}$ of -3.5‰ in microorganisms
NA not analysed

Table 4: Incorporation of ^{18}O isotopes from soil water into phosphate (%) of resin, microbial, organic and inorganic NaOH-EDTA, and HCl soil phosphorus pools during the incubation (34 days) per site. Means and corrected sample standard deviations per site and time point.

P pool	Incorporation of ^{18}O in phosphate during incubation (34 days) (%)			
	Site 1	Site 2	Site 3	Site 4
Resin	20 ± 1	24 ± 8	18 ± 17	41 [#]
Microbial	19 ± 5 ^H	24 ± 5 ^H	20 ± 10	33 [#]
Organic NaOH-EDTA	NA	6 ± 2	NA	1 ± 6
Inorganic NaOH-EDTA	2 [#]	1 [#]	NA	NA
HCl	-2 [#]	-7 ± 10	-7 ± 7	-1 ± 6

[#] n=1 (standard deviation not determinable), otherwise n=2
^H calculated based on $\delta^{18}\text{O}-P_{hex}$ values instead of $\delta^{18}\text{O}-P_{mic}$

Figures

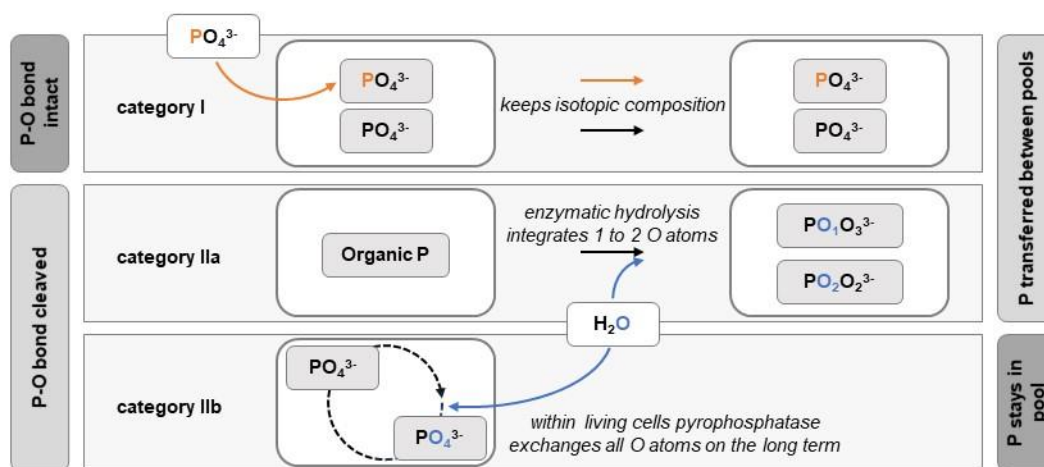


Figure 1: Conceptual visualisation of P cycling process categories and tracer (^{32}P and ^{18}O) movement within.

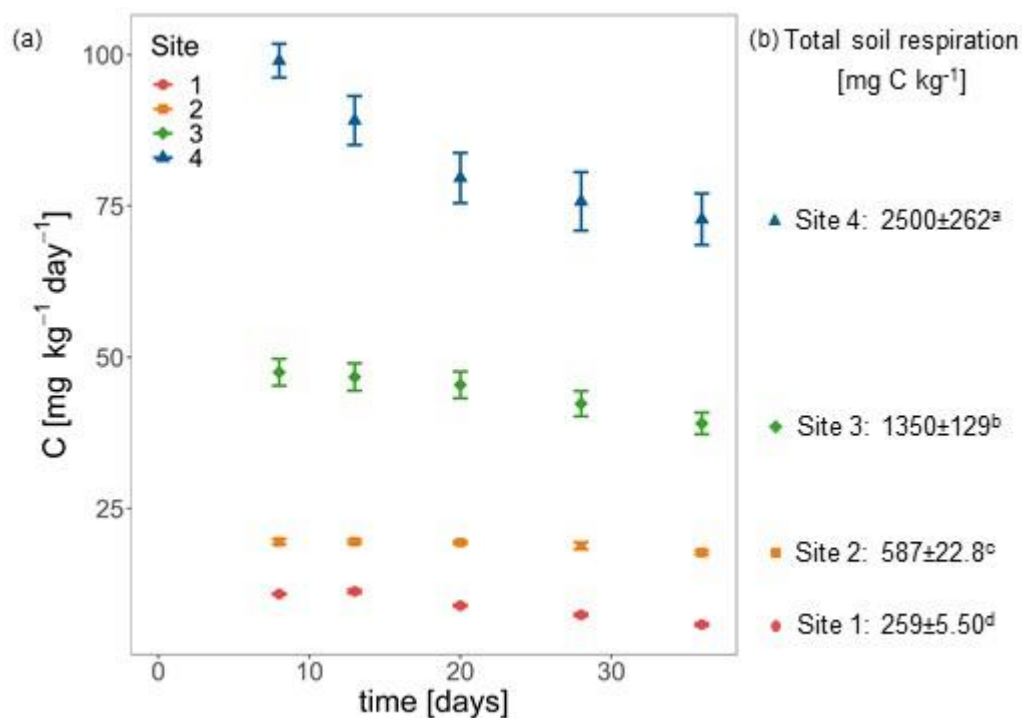


Figure 2: (a) daily ($\text{mg C kg}^{-1} \text{ dry soil day}^{-1}$) and (b) total ($\text{mg C kg}^{-1} \text{ dry soil}$) soil respiration during the incubation period (36 days) measured 8, 13, 20, 28 and 36 days after incubation start. Means and corrected sample standard deviations per site and time point ($n=4$), different letters in (b) denote significant differences ($P \leq 0.05$) among sites.

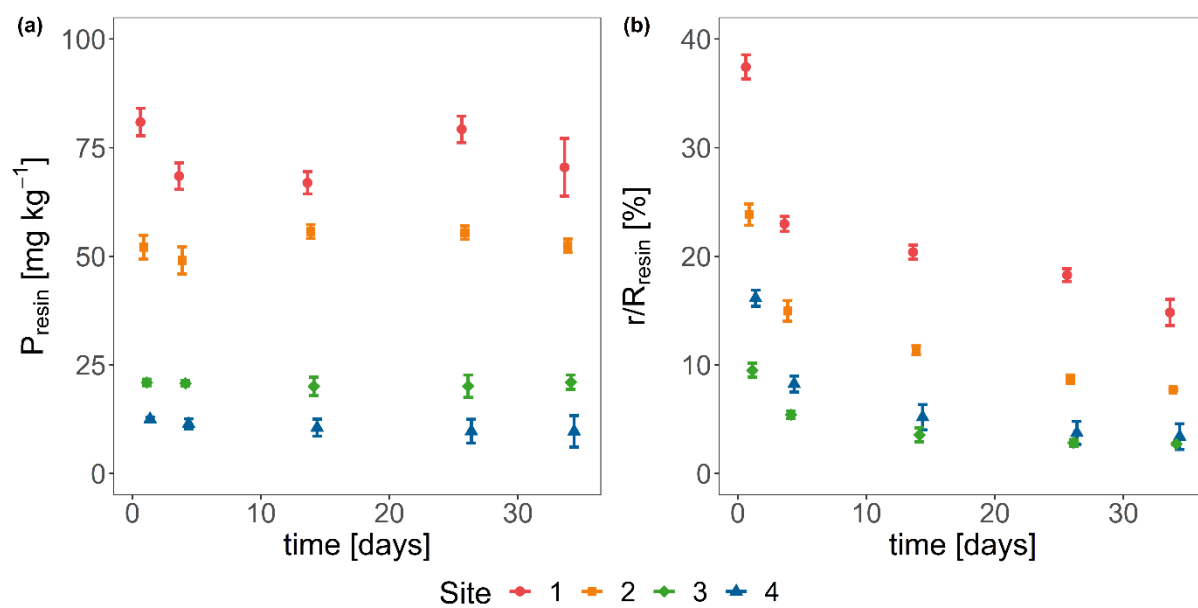


Figure 3: Resin (a) phosphorus concentration (mg P kg^{-1} dry soil) and (b) radioactivity recovery (r/R ; %) at 1, 4, 14, 26 and 34 days after incubation start per site. Means and corrected sample standard deviations per site and time point ($n=4$).

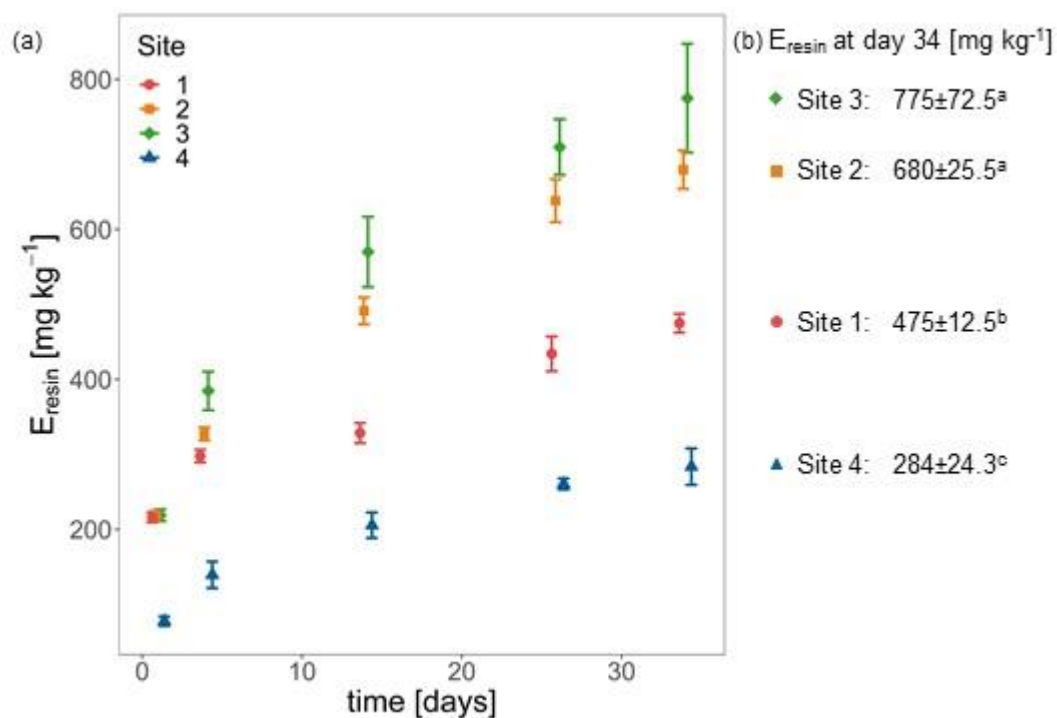


Figure 4: Amount of P isotopically exchangeable with P in resin pool (E_{resin}) in mg P kg^{-1} dry soil at (a) 1, 4, 14, 26 and 34 days and at (b) 34 days after incubation start. Means and corrected sample standard deviations and time point ($n=4$), different letters in (b) denote significant differences ($P \leq 0.05$) among sites.

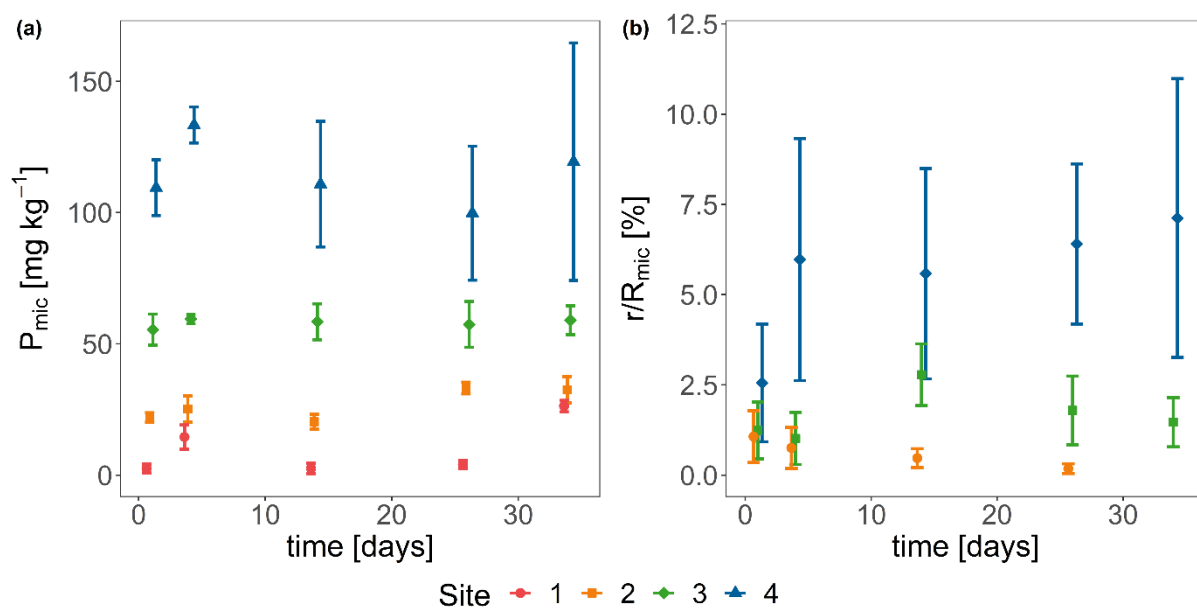


Figure 5: Microbial (a) phosphorus concentration (mg P kg⁻¹ dry soil) and (b) radioactivity recovery (r/R ; %) at 1, 4, 14, 26 and 34 days after incubation start per site. Means and corrected sample standard deviations per site and time point ($n=4$). r/R_{mic} of site 1 is not shown, 4 negative values obtained for r/R_{mic} of site 2 at 34 days were removed from the dataset.

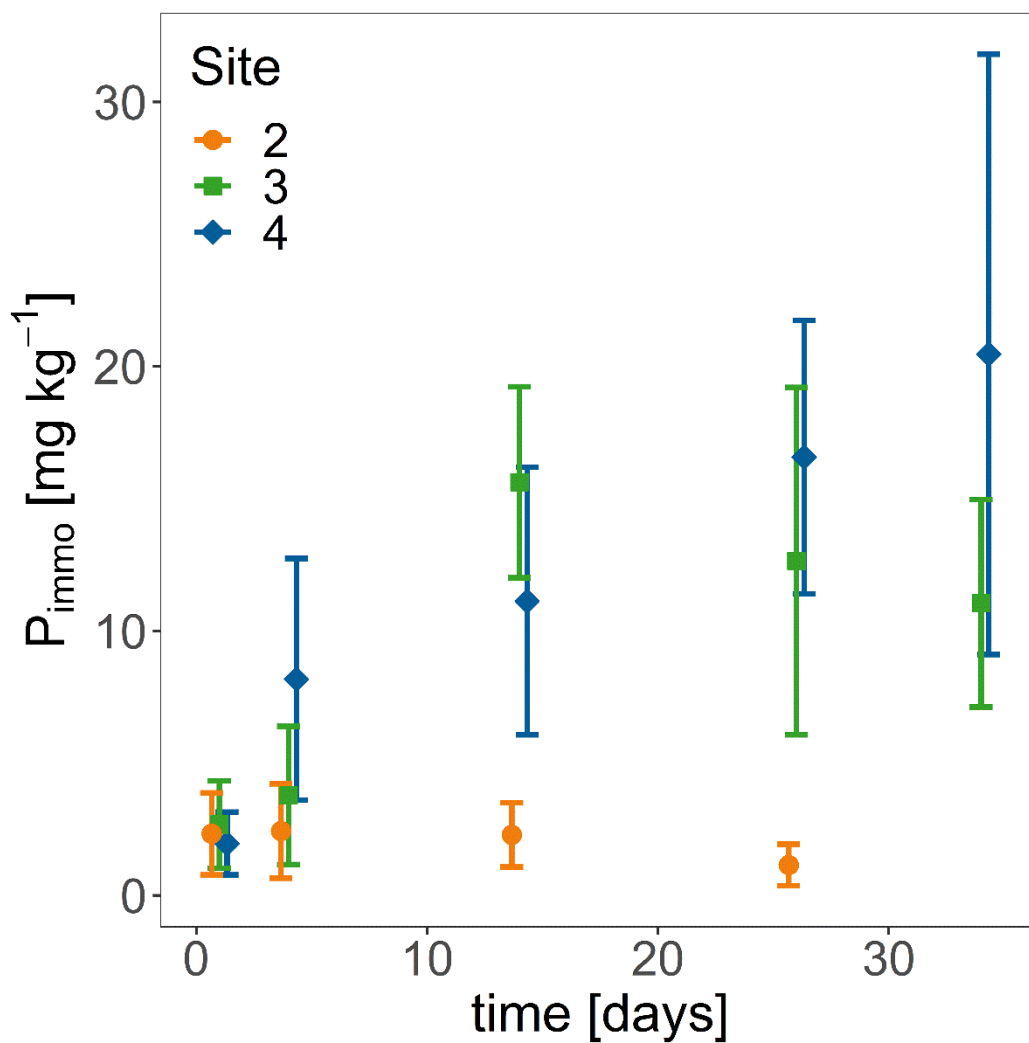


Figure 6: Fraction of microbial phosphorus derived from the resin phosphorus pool (gross P immobilisation, P_{immo}) in mg P kg⁻¹ dry soil at 1, 4, 14, 26 and 34 days after incubation start. Means and corrected sample standard deviations per site and time point (n=4). P_{immo} of site 1 is not shown, 4 negative values obtained for site 2 were removed from the dataset.

Supplementary

Sampling sites

Table SM1: Sampling site description: position (latitude, longitude), elevation, mean annual precipitation (MAP), mean annual temperature (MAT), parent material, soil type, and dominant and additional plant species per site.

	Site 1	Site 2	Site 3	Site 4
Latitude (°N)	20.113	20.151	20.156	20.135
Longitude (°W)	-155.880	-155.821	-155.797	-155.748
Elevation (m a.s.l.)	70	735	860	1059
MAP (mm yr ⁻¹)	275	1578	2163	3195
MAT (°C)	23.6	19.1	17.8	16.2
Parent material	Hawi basalt	Hawi basalt	Hawi basalt	Hawi basalt
Soil type	medial-skeletal, ferrihydritic, isohyperthermic Typic Haplotorrands	medial, amorphic, isomesic Humic Haplustands	medial, amorphic, isomesic Hydric Fulvudands	medial, amorphic, isomesic Hydric Fulvudands
Dominant plant species	buffel grass <i>Pennisetum ciliare</i>	kikuyu grass <i>Pennisetum clandestinum</i>	kikuyu grass <i>Pennisetum clandestinum</i>	kikuyu grass <i>Pennisetum clandestinum</i>
Additional plant species		forage soybean <i>Neonotonia wightii</i> (Wight and Arn.) J.A. Lackey		ohia <i>Metrosideros polymorpha</i> Gaudich.

Specific activity in resin and microbial phosphorus pools

Proportional specific activity in resin P (pSA_{resin} , Figure 5a) showed a clear decrease over time for all sites, indicating large P transfer from P_{resin} to other P pools. The extremest sites showed larger pSA_{resin} than the intermediate sites. Proportional specific activity in microbial P (pSA_{mic} , Figure 5b) varied among time points for sites 2, 3 and 4.

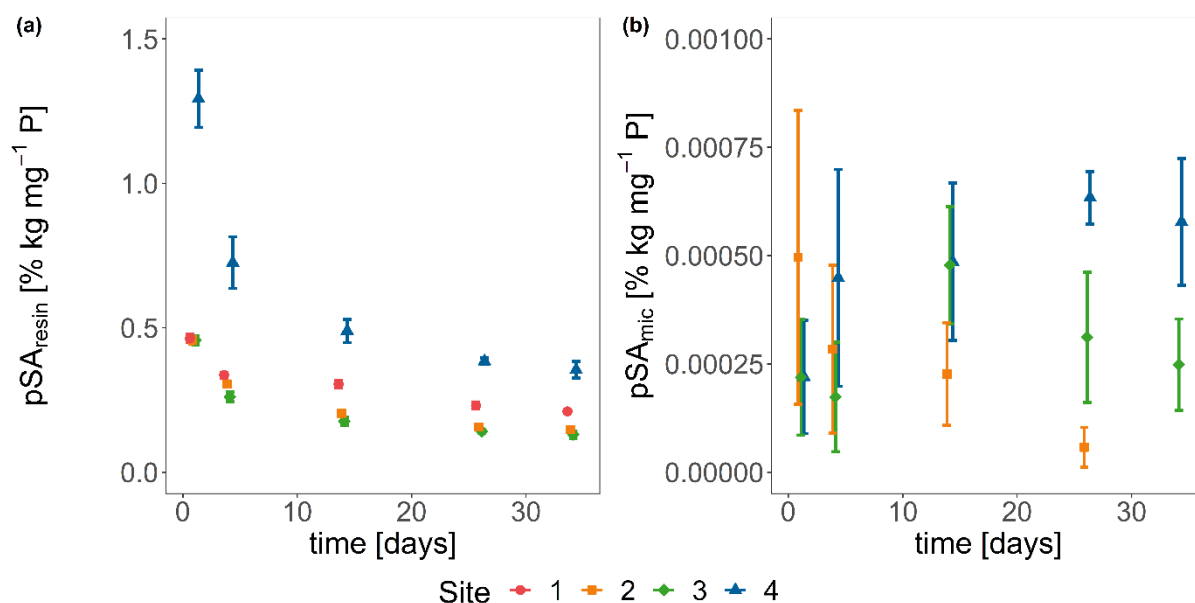


Figure SM1: Proportional specific activity (pSA , % kg dry soil mg^{-1} P) of (a) resin and (b) microbial P pools at 1, 4, 14, 26 and 34 days after incubation start per site ($n=4$). Means and standard deviations per site and time point. pSA_{mic} of site 1 is not shown, 4 negative values obtained for pSA_{mic} of site 2 were removed from the dataset.

Phosphorus pool concentration of Hedley extracts processed for the oxygen isotopic signatures

Table SM2: Phosphorus pool concentrations of the Hedley extracts processed for the oxygen isotopic signatures. Mean and standard deviations per time point and site, n=2.

		P concentrations [mg P kg ⁻¹]			
	Time [days]	Soils of site			
		1	2	3	4
Resin	0	62.3 ± 0.8	37.5 ± 0.7	10.6 ± 0.5	8.3 ± 0.8
	34	44.3 ± 3.0	28.3 ± 1.8	6.5 ± 0.4	3.5 ± 2.6
Microbial	0	1.9 ± 2.4	17.6 ± 0.7	27.1 ± 3.7	87.7 ± 1.5
	34	6.3 ± 2.6	11.4 ± 4.2	21.1 ± 3.1	46.8 ± 28.4
Organic NaOH-EDTA	0	279.9 ± 421.2	1745.5 ± 63.1	1427.6 ± 286.9	2161.2 ± 82.2
	34	675.8 ± 453.2	765.7 ± 228.7	2066.6 ± 862.4	1627.4 ± 261.6
Inorganic NaOH-EDTA	0	2225.8 ± 142.1	2028.5 ± 44.4	1136.8 ± 74.4	548.8 ± 60.7
	34	1979.6 ± 63.2	1891.6 ± 114.8	1146.6 ± 8.4	495.9 ± 14.8
HCl	0	865.6 ± 12.7	202.2 ± 0.1	697.5 ± 27.3	305.8 ± 4.2
	34	958.8 ± 170.7	210.0 ± 57.4	837.3 ± 16.0	284.2 ± 48.7