

The impact of soil microorganisms on the global budget of $\delta^{18}\text{O}$ in atmospheric CO_2

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Edited by Christopher B. Field, Carnegie Institution of Washington, Stanford, CA, and approved October 22, 2009 (received for review May 13, 2009)

Improved global estimates of terrestrial photosynthesis and respiration are critical for predicting the rate of change in atmospheric CO_2 . The oxygen isotopic composition of atmospheric CO_2 can be used to estimate these fluxes because oxygen isotopic exchange between CO_2 and water creates distinct isotopic flux signatures. The enzyme carbonic anhydrase (CA) is known to accelerate this exchange in leaves, but the possibility of CA activity in soils is commonly neglected. Here, we report widespread accelerated soil CO_2 hydration. Exchange was 10–300 times faster than the uncatalyzed rate, consistent with typical population sizes for CA-containing soil microorganisms. Including accelerated soil hydration in global model simulations modifies contributions from soil and foliage to the global CO^{18}O budget and eliminates persistent discrepancies existing between model and atmospheric observations. This enhanced soil hydration also increases the differences between the isotopic signatures of photosynthesis and respiration, particularly in the tropics, increasing the precision of CO_2 gross fluxes obtained by using the $\delta^{18}\text{O}$ of atmospheric CO_2 by 50%.

carbon cycle | water cycle | carbonic anhydrase | oxygen isotopes | terrestrial biosphere

The Earth's climate system is intimately connected to the movement of water and carbon across the planetary surface. As global warming proceeds, it is expected that photosynthetic CO_2 uptake will increase in colder regions of the world and diminish in those regions that are already warm and dry (1). At the same time, warming is expected to increase microbial activity, at least where water is not limiting, and therefore lead to an enhanced breakdown of organic matter in the soil, producing a large respiratory flux of CO_2 back to the atmosphere (2). Because terrestrial ecosystems presently sequester about a quarter of the CO_2 emissions associated with fossil fuel burning (7.1 GtC y^{-1}) (1), it is critical that we understand how large-scale, climate-driven changes will affect the carbon sequestration of the terrestrial biosphere. Currently, the precise response of terrestrial CO_2 sources and sinks to changes in climate remains uncertain (3) and its understanding requires the ability to quantify the amount of CO_2 taken up during photosynthesis separately from the amount released by respiration.

The oxygen isotope composition of atmospheric CO_2 (δ_a) was shown to be a powerful tracer of photosynthetic and respiratory CO_2 fluxes while at the same time providing information on the intensity of water cycling within terrestrial ecosystems (4–6). This tracing property occurs because the oxygen isotope composition ($\delta^{18}\text{O}$) of leaf and soil water pools is transferred to

atmospheric CO_2 during photosynthetic and respiratory CO_2 exchange, via an isotopic exchange during CO_2 hydration (7): $\text{CO}_{2\text{aq}} + \text{H}_2^{18}\text{O} \rightleftharpoons \text{CO}^{18}\text{O}_{\text{aq}} + \text{H}_2\text{O}$. Despite the short residence time of CO_2 in leaves, CO_2 involved in photosynthesis is nearly completely relabeled by ^{18}O -enriched leaf water because of the enzyme carbonic anhydrase (CA; EC 4.2.1.1), a very efficient catalyst of CO_2 hydration and isotopic exchange (4, 5, 8, 9). Typically the $\delta^{18}\text{O}$ of leaf and soil water pools are very different. There is a tendency for the heavier molecules of water to accumulate more readily in leaves than in soils during evapotranspiration because of the difference in water pool size (10, 11). Because the CO_2 - H_2O exchange in leaves (associated with photosynthesis) or soils (associated with soil respiration) produces such contrasting ^{18}O signals, estimates of the amount of CO_2 exchanged during photosynthesis and respiration can in principle be constrained by using the $\delta^{18}\text{O}$ signal of atmospheric CO_2 (6, 12).

However, our ability to partition gross fluxes of CO_2 may be complicated because the $\delta^{18}\text{O}$ of soil water (δ_{sw}) can often display a strong vertical gradient at the soil surface because soil evaporation also leads to an enrichment of heavy water molecules in the uppermost layers (13–15). Thus, to determine the $\delta^{18}\text{O}$ of CO_2 exchanged between soils and the atmosphere accurately it becomes necessary to know the shallowest depth (z_{eq}) where diffusing CO_2 molecules (from the atmosphere or produced by soil respiration; Fig. 1A) have enough time to fully equilibrate isotopically with soil water. With increasing temperature and moisture, CO_2 hydration increases relative to the diffusion rate so that z_{eq} moves closer to the surface, and toward more enriched $\delta^{18}\text{O}$ values (see *Methods*, Eq. 4). Although we know that CA accelerates the rate of hydration in leaves, the possibility of CA activity in soils is commonly neglected (4, 15),

Author contributions: L.W. and J.O. designed research; L.W., J.O., M.C., B.G., I.R., D.Y., K.M., E.G.P., M.M.B., B.M., R.B., J.M., M.M., J.H.S., J.H., and J.G. performed research; P.P. contributed analytic tools; L.W., J.O., M.C., B.G., I.R., and U.S. analyzed data; and L.W., J.O., U.S., and D.Y. wrote the paper.

The authors declare no conflict of interest.

This article is a PNAS Direct Submission.

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This article contains supporting information online at www.pnas.org/cgi/content/full/0905210106/DCSupplemental.

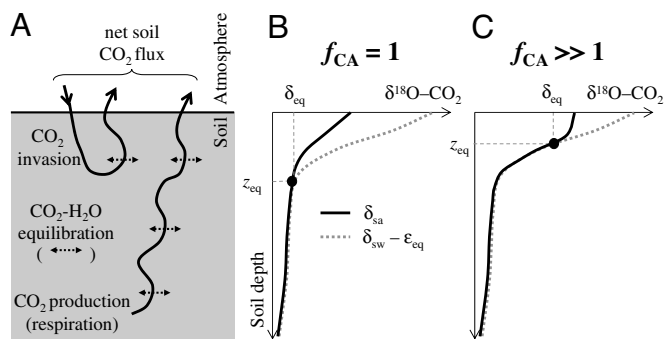


Fig. 1. Schematic showing the influence of CO₂ hydration rates on vertical profiles of δ¹⁸O in soil air CO₂. (A) The net soil-atmosphere CO₂ exchange is composed of CO₂ molecules moving from the atmosphere into the soil and back to the atmosphere (i.e., invasion) and further CO₂ molecules produced during soil respiration. Because of oxygen isotopic exchange between soil CO₂ and water, both invasion and respiration fluxes modify the isotopic composition of atmospheric CO₂, and their ¹⁸O isotopic signature depend on the extent of CA activity in the soil. (B) Typical profile of δ¹⁸O in soil air CO₂ (δ_{sa}) for uncatalyzed CO₂ hydration in soil water (enhancement factor $f_{CA} = 1$). (C) Same as in B but for catalyzed CO₂ hydration (enhancement factor $f_{CA} \gg 1$). In deep soil layers where vertical gradients of δ_{sw} are weak, the residence time of CO₂ is long enough to reach full isotopic equilibrium with soil water (δ_{sa} = δ_{sw} - ε_{eq}), where ε_{eq} denotes the isotopic equilibrium fractionation between CO₂ and water (22). Above a certain depth z_{eq} (where, by definition, δ_{sa} = δ_{eq}), CO₂ molecules diffuse too rapidly to fully equilibrate with local soil water. If CO₂ hydration is enhanced because of CA activity ($f_{CA} \gg 1$), the equilibration becomes faster and z_{eq} shallower, thus δ_{eq} becomes more enriched.

because the abundance and location of CA in soils is still somewhat unclear, with only indirect and isolated indications based on measurements of δ¹⁸O of soil CO₂ or COS fluxes (14, 16, 17). Substantial CA activity in soils would lead to a faster equilibration of CO₂, moving z_{eq} further toward the surface where soil water is more ¹⁸O enriched (Fig. 1 B and C). So far global simulations have assumed uncatalyzed CO₂ hydration in soils (18–20) and equilibration depths below the region of strong evaporative enrichment (5, 21).

Results and Discussion

Evidence for Enhanced Soil CO₂ Hydration Rates. Here, we demonstrate that, in contrast to current assumptions, the observed rate of soil CO₂ hydration is always substantially faster than the uncatalyzed rate. We compared measurements of depth-resolved soil water δ¹⁸O (δ_{sw}) and observed δ¹⁸O signatures of chamber-based soil CO₂ fluxes (δ_{flux}) in seven different ecosystems that encompass most of the major land biomes, providing a global perspective of ¹⁸O exchange in soils (Table S1 and see Tables S5–S7). From the δ_{sw} data, we determined the depth-resolved δ¹⁸O of soil CO₂ in full equilibrium with soil water (δ_{eq}), equal to δ_{sw} - ε_{eq} where ε_{eq} is the temperature-sensitive equilibrium fractionation between CO₂ and water (22). Most sites exhibited strong gradients in δ_{sw} - ε_{eq} at the soil surface, reflecting the evaporative enrichment of soil water (Fig. 2). From the δ_{flux} data, we determined the δ¹⁸O of soil CO₂ at z_{eq} (δ_{eq}, see Fig. 1) for different rates of hydration expressed as an enhancement factor (f_{CA}) with respect to the uncatalyzed CO₂ hydration rate (see Eq. 6 in Methods). Increasing f_{CA} shifts z_{eq} toward surface layers (Fig. 2) and δ_{eq} toward δ_a. The best estimate for f_{CA} would be one in agreement with both soil water and chamber flux measurements. This is obtained when the point (δ_{eq}, z_{eq}) derived from the chamber data intersects the δ_{eq} curve derived from soil water measurements. At all sites, this intersection occurs for values of f_{CA} between 10 and 300, with the lowest f_{CA} in the cooler temperate ecosystems while higher f_{CA} were found at the Mediterranean and subtropical sites (Fig. 2). As a consequence, the equilibration depth z_{eq} was in most cases within the top 5 cm of the soil, the zone containing the strongest δ_{sw} gradients. A reduction in the effective diffusivity of soil CO₂ would also lead to shallower equilibration depths z_{eq} by increasing the residence time of CO₂ in soils, but it would not yield simultaneous solutions for both soil water and CO₂ flux isotope data (14). Thus, an enhanced CO₂ hydration rate is the only plausible mechanism to explain these chamber-based measurements.

Consistency with CA Activities in Soil Microorganisms. The uppermost soil layers host many bacterial, algal, and fungal species that produce intracellular and sometimes extracellular CAs (23–25). Based on a literature survey, we claim that this mixed population

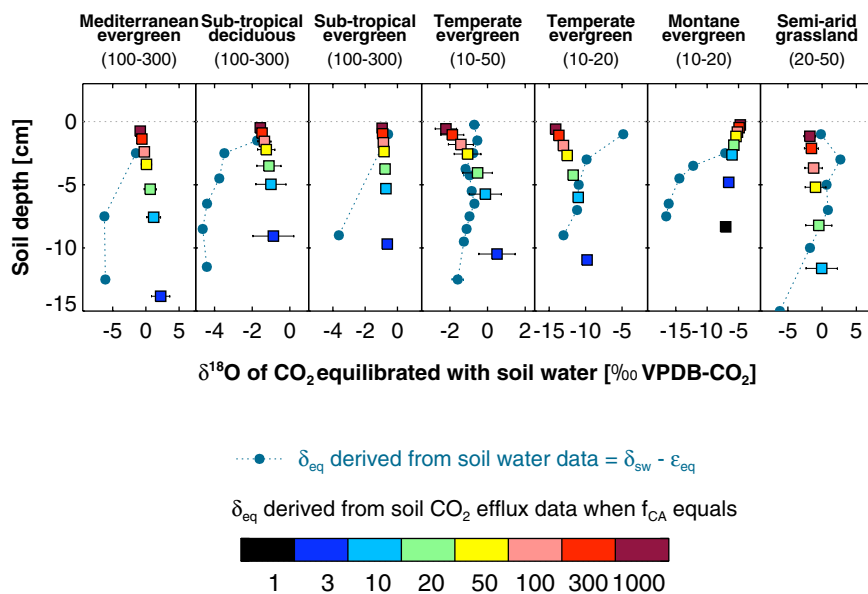


Fig. 2. The δ¹⁸O of soil CO₂ at the depth of full equilibration (δ_{eq}, z_{eq} ; see Fig. 1) estimated from chamber flux measurements for different levels of hydration rates (f_{CA}). Depth-resolved soil water data yields the δ¹⁸O of CO₂ in isotopic equilibrium with soil water (δ_{sw} - ε_{eq}). The point at which the two curves intersect indicates the most likely value for the enhancement factor, f_{CA} , listed below the ecosystem type for each site (see Table S1). The horizontal error bars on the squared symbols represent the standard deviation of δ_{eq} values over the number of δ_{flux} measurements ($n = 1$ –15).

of soil microorganisms is responsible for the accelerated soil CO₂ hydration. Most soils contain 10³ to 10⁶ algae per g of dry soil, but populations can reach 10⁸ algae per g of dry soil (Table S2). Bacterial population sizes are even larger at 10⁸ to 10⁹ cells per g of dry soil (Table S3). At 25 °C, the CO₂ hydration rate in soil algal and cyanobacterial cells can be up to 172,500 times the uncatalyzed rate, comparable to CA activities found in plant chloroplasts (Table S4). With a cell volume of ≈100 μm³ and population of 10⁶ per g of dry soil, algae could explain a significant fraction of our observed soil CA activities. Indeed, we found that the presence of algae developing naturally on the surface of a peat soil dramatically enhanced CA activity (Table S4). Laboratory studies have also reported high CA activities (*f_{CA}* = 50) in bulk soil extracts from subtropical karst forests containing a mixture of bacterial and fungal species (Table S4). Based on a cell volume of ≈1 μm³ and population of 4 × 10⁹ cells per g of dry soil (Table S3), this soil-level *f_{CA}* value would be consistent with soil bacteria operating at a cell-level CO₂ hydration of 8,000 times the uncatalyzed rate (Table S4). These estimates demonstrate that soil microorganisms are likely to be responsible for enhanced soil CO₂ hydration rates of 20–300 times the uncatalyzed rate.

Impact of Soil CA Activity at the Global Scale. The accelerated hydration of CO₂ in soils has been missing in the mass budget of δ¹⁸O in atmospheric CO₂. To explore the impact of soil CA activity on the δ¹⁸O of atmospheric CO₂, and its north–south (N-S) gradient, we incorporated this biological process into the global model of δ¹⁸O in atmospheric CO₂, Meebeth (18, 19) (see *Methods*). Simulations were performed over an average year calculated from the 1990s and compared with observations from the worldwide network of atmospheric stations for the same decade (26, 27). Three scenarios are discussed here: one uncatalyzed (abiotic) scenario (*f_{CA}* = 1) and two globally uniform *f_{CA}* scenarios covering the range of soil chamber estimates (*f_{CA}* = 20 and *f_{CA}* = 300).

The high CA activity scenario (*f_{CA}* = 300) improves the agreement between the modeled and observed N-S gradient in δ_a, particularly when compared with the uncatalyzed, abiotic scenario (Fig. 3). This latitudinal feature in δ_a is largely driven by the N-S gradient in the δ¹⁸O of precipitation that creates depleted leaf and soil water pools toward the northern latitudes. In the uncatalyzed scenario (*f_{CA}* = 1), photosynthesis dominates the N-S gradient. Introducing *f_{CA}* enhances the invasion flux (the number of CO₂ molecules from the atmosphere that equilibrate with soil water and go back to the atmosphere; see Fig. 1*A* and *Methods*, Eq. 5). The contribution of this invasion flux to the N-S gradient increases with *f_{CA}* and at some latitudes becomes larger than the contribution of respiration.

Incorporating high CA activity (*f_{CA}* = 300) also reduces the mean value of δ_a by ≈1‰ relative to the abiotic case (*f_{CA}* = 1), bringing the model closer to atmospheric observations. This reduction is the result of complex interactions between δ_a and the δ¹⁸O signatures of all component fluxes. The influence of each process on δ_a can be represented by using the concept of isoflux (*I_x*) defined as the product of a gross CO₂ flux (*F_x*) and its isotopic composition (δ_x) relative to δ_a: *I_x* = *F_x* (δ_x - δ_a). Photosynthesis tends to enrich the atmosphere (positive isoflux, δ_x > δ_a), whereas respiration and soil invasion usually have the opposite effect (negative isoflux, δ_x < δ_a). Nonbiospheric fluxes (from ocean, fossil fuel, and biomass burning) also tend to deplete δ_a, but to a much lesser extent (19, 28). In the uncatalyzed scenario (*f_{CA}* = 1), photosynthetic (*I_A*) and respiratory (*I_R*) isofluxes balance the nonbiospheric isofluxes globally while the soil invasion isoflux (*I_{inv}*) remains close to zero at all latitudes (Fig. 3). When *f_{CA}* is increased, the isotopic signatures of soil invasion and respiration become progressively enriched as a result of the isotopic gradient in δ_{sw} (Fig. 1), but usually remain

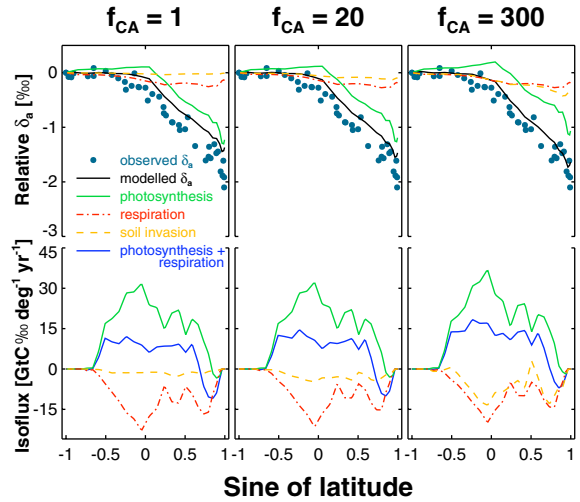


Fig. 3. Simulated contributions of different biospheric processes to the N-S gradient in δ_a for uncatalyzed (*f_{CA}* = 1) or enhanced (*f_{CA}* ≫ 1) CO₂ hydration rates in the soil, compared with measured δ_a. Enhanced hydration increases the corresponding isofluxes of soil invasion and photosynthesis + respiration, i.e., the isotopic imbalance required for gross flux partitioning. Note that δ_a values are always reported relative to the South Pole and thus do not show the absolute changes in δ_a (−0.1‰ and −1.1‰ for *f_{CA}* of 20 and 300 relative to *f_{CA}* = 1, respectively).

below δ_a (*I_R* and *I_{inv}* remain negative). Most importantly, the mass of atmospheric CO₂ molecules that equilibrate with soil water increases from 25 GtC yr^{−1} in the uncatalyzed scenario to 450 GtC yr^{−1} when *f_{CA}* = 300. As a result, *I_{inv}* becomes very large and negative, reaching nearly the same magnitude as soil respiration. The associated depletion in δ_a is partly compensated by an increase in both *I_A* and *I_R*. When *f_{CA}* = 300, the absolute value of *I_{inv}* increases by 571 GtC ‰ yr^{−1}, whereas *I_R* decreases by 226 GtC ‰ yr^{−1} and *I_A* increases by 269 GtC ‰ yr^{−1}. Soil CA activity thus strongly modifies the relative contribution of photosynthesis and respiration to the CO₂ budget in our global model.

Consequences for the Retrieval of CO₂ Sources and Sinks. Measurements of δ_a have been proposed as one of the few tools available to partition net CO₂ fluxes into photosynthesis and soil respiration, but it critically depends on the existence of sufficient imbalance between *I_A* and *I_R* (4–6, 9, 12, 21, 28). In previous studies, where δ_a was prescribed and not dynamically coupled to soil and leaf water pools as in our study, such an imbalance had been restricted to the boreal regions (5, 21, 28). In contrast, we now show a significant isotopic imbalance at nearly all latitudes (photosynthesis + respiration curve in Fig. 3), greatly enhancing the potential of the δ¹⁸O approach for partitioning CO₂ fluxes. At the continental scale, there is a strong isotopic imbalance over Europe and North America coinciding with the peak season of photosynthetic activity in the Northern Hemisphere (Fig. 4). Over the tropics, the isotopic imbalance between *I_A* and *I_R* increases by up to 50% when *f_{CA}* = 300 and is maintained year-round in many areas (Fig. 4). The uncertainties of tropical gross CO₂ fluxes could thus be reduced by an equivalent amount (12), making the δ¹⁸O of atmospheric CO₂ a better tracer for terrestrial gross CO₂ fluxes than previously thought in these regions. From a one-box global mass balance budget and using globally averaged fluxes and isotopic signatures from Meebeth, we calculated that, neglecting “biotic” invasion associated with soil CA activity in inversion studies leads to errors in the isotope-derived estimates of global photosynthesis by up to 30 GtC yr^{−1}, i.e., ≈30% of current estimates (1).

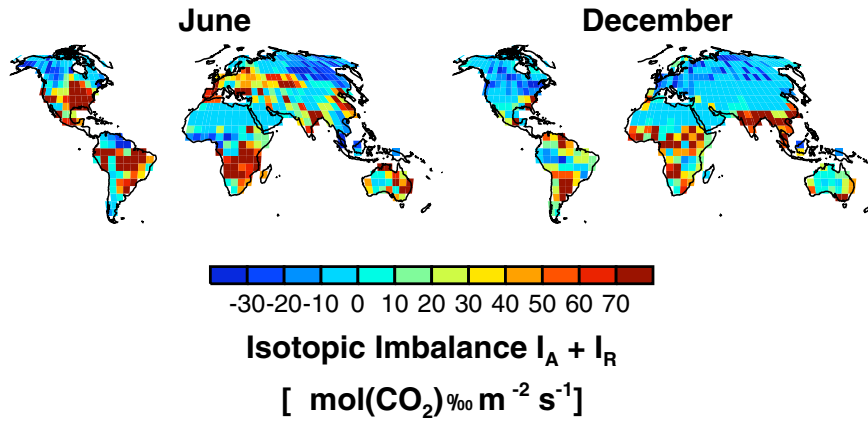


Fig. 4. Global distribution in the extent of isotopic imbalance ($I_A + I_R$) across continental surfaces for June and December simulated by the global model Mecbeth for the most enhanced soil CO_2 hydration scenario ($f_{CA} = 300$). Regions where $I_A + I_R$ is the most different from zero correspond to regions of strong isotopic imbalance where biospheric gross CO_2 fluxes are expected to be the most constrained by $\delta^{18}\text{O}$ data.

Future Directions for Global Isotope-Enabled Models. This study demonstrates that enhanced rates of CO_2 hydration occur at the soil surface and appreciably impact the oxygen isotope composition of atmospheric CO_2 . This enhanced exchange in the soil brings into focus our limited ability to predict the isotopic enrichment of soil water near the surface (18, 29), highlighting a need for future improvements in this research area. Also, although we provided the basic observations and parameterization, more work is now needed to further assess the variability in f_{CA} in different ecosystems, plant functional types, or regions within the global model, including attempts to establish the mechanistic basis to underpin the observed differences in CA activity between ecosystems. Developments on these fronts will greatly enhance our capabilities to use the $\delta^{18}\text{O}$ of atmospheric CO_2 to quantitatively inform us of large-scale changes in the intensity of carbon and water cycling in terrestrial ecosystems.

Methods

Soil CO^{18}O Budget Equation. In a given soil layer, the number of moles of CO^{18}O changes as a result of (i) CO^{18}O production during heterotrophic and autotrophic respiration, (ii) diffusion of these molecules through the soil layer, and (iii) oxygen isotopic exchange with the surrounding soil water (30–32):

$$\theta_t \frac{\partial C\mathcal{R}}{\partial t} = \mathcal{R}_c S_c + \frac{\partial}{\partial z} \left[D_{c,\text{iso}} \frac{\partial C\mathcal{R}}{\partial z} \right] + k_{h,\text{iso}} B \theta_w C (\mathcal{R}_{\text{eq}} - \mathcal{R}), \quad [1]$$

where C [$\text{mol}\cdot\text{mol}^{-1}$] is the CO_2 mole fraction in soil air, \mathcal{R} , \mathcal{R}_c , and \mathcal{R}_{eq} are the $^{18}\text{O}/^{16}\text{O}$ ratios of the CO_2 in soil air, respired CO_2 , and CO_2 in isotopic equilibrium with the surrounding soil water, respectively, S_c ($\text{mol}\cdot\text{m}^{-3}\cdot\text{s}^{-1}$) is the respiration rate density, $D_{c,\text{iso}}$ ($\text{m}^2\cdot\text{s}^{-1}$) is the effective diffusivity of CO^{18}O in soil air, θ_w ($\text{m}^3\cdot\text{m}^{-3}$) is the volumetric soil water content, B is the CO_2 solubility coefficient, and θ_t ($\text{m}^3\cdot\text{m}^{-3}$) is the total CO_2 porosity. Denoting by θ_a the soil air porosity we have (31): $\theta_t = \theta_a + B\theta_w$. The solubility coefficient B depends on soil temperature T_s (K) according to ref. 33: $B = 1.739\exp(-0.039(T_s - 273.15) + 0.000236(T_s - 273.15)^2)$. \mathcal{R}_{eq} is related to the $^{18}\text{O}/^{16}\text{O}$ ratio in soil water \mathcal{R}_{sw} through $\mathcal{R}_{\text{eq}} = (1 + \varepsilon_{\text{eq}})\mathcal{R}_{\text{sw}}$, where $\varepsilon_{\text{eq}} = 17.604/T_s - 0.01793$ is the CO_2 - H_2O equilibrium fractionation (22). Because there are three oxygen atoms present in the bicarbonate intermediate, the isotopic exchange rate during CO_2 hydration equals one-third the hydration rate (7): $k_{h,\text{iso}} = f_{CA}k_{h,\text{uncat}}/3$, where (34) $k_{h,\text{uncat}} = 0.037 \times \exp(0.118(T_s - 298.15))$. In this framework, CA activity is expressed as an enhancement factor (f_{CA}) of the uncatalyzed CO_2 hydration rate ($k_{h,\text{uncat}}$). The effective CO^{18}O diffusivity in soil air is calculated as $D_{c,\text{iso}} = D_{c,\text{eff}} \alpha_d$, where $\alpha_d = 0.9913$ is the isotopic discrimination during molecular diffusion of CO_2 in air and $D_{c,\text{eff}}$ ($\text{m}^2\cdot\text{s}^{-1}$) is the effective CO_2 diffusivity in soil air. Several parameterizations of this effective diffusivity exist in the literature that differ mostly for wet soils (35). Results presented in this study use ref. 31: $D_{c,\text{eff}} = 0.66 \times \theta_a \times 1.4 \cdot 10^{-5} (T_s/298.15)^{1.75}$.

Full Equilibration Depth. The budget equation above contains two time scales. One time scale indicates the half-life of CO_2 molecules before being isotopically equilibrated with the surrounding water:

$$\tau_k = \ln 2 \cdot \left(\frac{\theta_t}{k_{h,\text{iso}} B \theta_w} \right) \quad [2]$$

and another time scale indicates the time required for a plume of C^{18}O molecules to diffuse through the soil over a given distance z :

$$\tau_d(z) = \frac{\theta_t z^2}{2D_{c,\text{iso}}} \quad [3]$$

Full equilibration within a soil layer of thickness z is satisfied when the time scale for isotopic equilibration is smaller than the time scale for diffusion through this layer, i.e., $\tau_k \ll \tau_d(z)$. When $\tau_k = \tau_d(z)$, full equilibration can occur if the soil layer has uniform soil temperature, moisture content, and isotopic composition. However, in the top centimeters of the soil, strong gradients of T_s , θ_w , and \mathcal{R}_w are more likely. The shallowest depth of full equilibration, z_{eq} , must therefore satisfy the inequality: $\tau_k < \tau_d(z_{\text{eq}})$. In the following we will define z_{eq} as: $\tau_k = \tau_d(z_{\text{eq}})/4$, or similarly:

$$z_{\text{eq}} = 2 \sqrt{\frac{2 \ln 2 D_{c,\text{iso}}}{k_{h,\text{iso}} B \theta_w}} \quad [4]$$

The factor 4 was determined by matching the value of f_{CA} deduced in Fig. 2 with that obtained from simulations using the full numerical model (Eq. 1), i.e., $f_{CA} \approx 300$ for the Mediterranean evergreen site (14) and $f_{CA} \approx 20$ for the montane evergreen site (15). Eq. 4 with $f_{CA} = 20$ also provides seasonal variations of z_{eq} at the temperate evergreen site that correspond to the depth where $\delta^{18}\text{O}$ in soil air CO_2 (δ_{sa}) and $\delta_{\text{sw}} - \varepsilon_{\text{eq}}$ (estimated using the full numerical model, Eq. 1) start to diverge by >0.3 ‰ (a threshold chosen for practical purposes to represent the overall precision of soil water isotope measurements).

Other studies (14, 35) use a different formulation for $D_{c,\text{iso}}$, leading to values of this diffusivity 5-fold smaller in saturated soils. Using this other formulation does not fundamentally change the results presented in Fig. 2.

Soil CO_2 Isoflux. In the steady state, and assuming isothermal and uniform soil water conditions, Eq. 1 can also be solved analytically (30–32). In this framework, the isotopic composition of the soil CO_2 flux δ_{flux} is:

$$\delta_{\text{flux}} = \delta_{\text{eq}} + \varepsilon_{d,\text{eff}} + (\delta_{\text{eq}} - \delta_a) v_{\text{inv}} \frac{C_a}{F_R}, \quad [5]$$

where $\varepsilon_{d,\text{eff}}$ is the effective isotopic fractionation during diffusion, F_R is the soil CO_2 efflux, and $v_{\text{inv}} = \sqrt{B\theta_w k_{h,\text{iso}} D_{c,\text{iso}}}$ has the dimensions of a velocity ($\text{m}\cdot\text{s}^{-1}$) that when multiplied by C_a gives the soil invasion flux F_{inv} . The product $(\delta_{\text{flux}} - \delta_a)F_R$ is called the soil CO_2 isoflux. It can be seen as the sum of two isotope fluxes: a respiration isoflux, $I_R = (\delta_{\text{eq}} + \varepsilon_{d,\text{eff}} - \delta_a)F_R$, and an invasion isoflux, $I_{\text{inv}} = (\delta_{\text{eq}} - \delta_a)F_{\text{inv}}$, sometimes defined as abiotic because it is independent of

F_R . Assuming a uniform soil CO_2 production S_c over a soil column of depth z_0 , $\varepsilon_{d,\text{eff}}$ can be estimated as (31): $\varepsilon_{d,\text{eff}} = \varepsilon_d(1 - z_1/z_0(1 - \exp(-z_0/z_1)))$, where $z_1 = (2\sqrt{2\ln 2})^{-1} z_{\text{eq}}$. Eq. 5 can then be inverted to estimate δ_{eq} as a function of δ_{flux} , C_a , δ_a , and F_R measurements:

$$\delta_{\text{eq}} = \frac{\delta_{\text{flux}} - \varepsilon_{d,\text{eff}} + v_{\text{inv}}C_a/F_R\delta_a}{1 + v_{\text{inv}}C_a/F_R} \quad [6]$$

Oxygen Isotope Composition of the Net CO_2 Flux from Soil Chambers. The steady-state oxygen isotope signal of the net soil CO_2 flux during chamber closure (δ_{ch}) was calculated by using a simple isotopic mass balance:

$$\delta_{\text{ch}} = \frac{\delta_{\text{out}}C_{\text{out}} - \delta_{\text{in}}C_{\text{in}}}{C_{\text{out}} - C_{\text{in}}}, \quad [7]$$

where C_{out} , C_{in} and δ_{out} , δ_{in} are the mole fractions and isotopic compositions of CO_2 in the air leaving and entering the chamber, respectively. In the case of the two sites that used closed chambers (subtropical evergreen and semiarid grassland), C_{out} , C_{in} and δ_{out} , δ_{in} are the mole fractions and isotopic compositions of CO_2 at the start and end of a defined chamber closure period, respectively.

To derive δ_{eq} values from soil chamber data, we use Eq. 6, neglect chamber effects, and make the common assumption that the atmosphere inside the chamber is well mixed ($C_a = C_{\text{out}}$ and $\delta_a = \delta_{\text{out}}$).

Oxygen Isotope Composition of Soil Water. Depth-resolved soil samples were collected at each experimental site within proximity of the soil chamber and at approximately the same time as gas exchange measurements. In the case of the Mediterranean evergreen, subtropical evergreen, and both temperate ever-

green sites, soil water was extracted cryogenically from bulk soil samples and $\delta^{18}\text{O}$ analysis of CO_2 equilibrated with the extracted water was completed (14). For the montane evergreen, subtropical deciduous, and semiarid grassland sites CO_2 with a known isotopic composition was equilibrated directly with fresh soil samples and stored in gas-tight containers for 12 h. Equilibrated CO_2 was then sampled from the container and analyzed for its $\delta^{18}\text{O}$ composition (15).

Global Model Simulations. The global model Mecbeth calculates the sources and sinks of CO_2 , water, and their respective isotopes and transports them in the atmosphere (18, 19). It merges a description of the biospheric energy, water, and carbon fluxes with a global climate and water isotope model. The atmosphere and biosphere are dynamically coupled to account for feedbacks of the accelerated equilibration of CO_2 with soil water on δ_a and the isotopic signatures of leaf and other fluxes. The model parameterization of soil water isotopes was improved in this study to provide depth-resolved descriptions of soil water and soil water isotopes (35), a necessary step if CA activity occurs in soils containing strong vertical gradients in δ_{sw} (14). Several soil layers of varying thickness were included in the model. The most important upper layers relevant to this study consisted of a top layer at 0–6 cm and another layer at 6–20 cm.

ACKNOWLEDGMENTS. We thank P. Richard for isotopic analysis of the soil water data from Le Bray, France and W. T. Baisden for contributions to the sample collection at Canterbury, New Zealand. L.W. was supported by the CarboEurope-IP research program funded by the European Union. Soil CA assays were made possible through funding from the Institut National de la Recherche Agronomique Projet Innovant, Centre National de la Recherche Scientifique Ecosystèmes Continentaux et Risques Environnementaux and Région Provence-Alpes-Côte d'Azur.

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Supporting Information

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Table S1. Experimental set-up and description of sites used in this study

Site description	Site name	Location	Soil type	Dominant species	Chamber	Sample date	Ref./ source
Mediterranean evergreen forest	Mitra	Portugal	Sandy loam	<i>Quercus suber</i> L.	Open	9/7/2004	1
Subtropical deciduous forest	Oak Ridge	Tennessee, USA	Silty loam	<i>Quercus</i> : and <i>Acer</i>	Open	7/16/1998	2
Subtropical evergreen forest	Tallahassee	Florida, USA	Sandy silt	<i>Pinus elliotii</i> Engelm.	Closed	6/2/2003	3
Temperate evergreen forest	Bray	France	Sandy podzol	<i>Pinus pinaster</i> Ait.	Open	10/1/2007	4
Temperate evergreen forest	Canterbury	New Zealand	Silty loam	<i>Pinus radiata</i> D. Don.	Open	10/21/2005	This study
Temperate Montane evergreen forest	Niwot	Colorado, USA	Sandy loam	<i>Pseudotsuga menziessii</i> Mirb.	Open	8/4/1997	5
Semiarid grassland	Fort Collins	Colorado, USA	Sandy loam	C3-C4 grasses	Closed	7/26/2001	6

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Table S2. Algal populations in surface soils of different ecosystems

Site description	Location	# g dry soil ⁻¹ ($\times 10^3$)*	Ref.
Temperate montane forest	Central Europe	50 / 4 / 5	1
Temperate montane forest	Central Europe	50 / 0 / 0	1
Temperate montane forest	Central Europe	1 / 0 / 0	1
Mediterranean agricultural field	Italy	1,000 / 300 / 60	2
Mediterranean agricultural pasture	Italy	700 / 300 / 100	2
Mediterranean agricultural vineyard	Italy	100 / 0 / 10	2
Mediterranean agricultural field	Italy	100 / 20 / 5	2
Desert agricultural field (irrigated)	Kazakhstan	100 / 10 / 0	3
Desert	Kazakhstan	2 / 0.4 / 0	3
Antarctic tundra	Antarctica	1–4,000 / 0–350 / 0 [†]	4
Temperate grassland	England	500–4,000 [†]	5
Temperate forest	England	50	5
Temperate grassland and gardens	Europe	40–2,000	6
Temperate forests	Europe and USA	10–793	6
Temperate agricultural fields	Europe and USA	3,100–33,000	6
Semiarid grassland	USA	340–110,000	6

*If available, population sizes are given for green and yellow-green algae/cyanobacteria/diatoms, separately. In general, populations of green and yellow-green algae > cyanobacteria > diatoms, with approximate proportions 20:2:1. This indicates that the algal group displaying the highest f_{CA} (see Table S4) also has the highest abundance.

[†]Assuming a soil bulk density of 1,300 kg·m⁻³.

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Table S3. Bacterial populations in surface soils of different ecosystems

Site description	Location	# g dry soil ⁻¹ (×10 ⁹)	Ref.
Tundra peatland	Siberia	2.3	1
Tundra peatland	Siberia	0.45*	2
Arctic alpine soil	Norway	0.43 - 2.6	3
Boreal deciduous forest	Sweden	2.3	4
Boreal evergreen forest	Sweden	5.9	4
Boreal evergreen forest	Sweden	0.14	4
Boreal evergreen forest	Sweden	2.3 - 7.0	5
Boreal pasture	Sweden	0.08	4
Boreal garden	Sweden	0.85	4
Boreal garden	Sweden	1.4	4
Temperate peatland	Germany	0.17*	2
Temperate deciduous forest	Denmark	1.0	6
Temperate deciduous forest	Denmark	1.0	6
Temperate mixed forest	New York, USA	4.3	7
Temperate evergreen forest	Netherlands	2.0 - 6.5	8
Temperate evergreen forest	Sweden	1.6	4
Temperate evergreen forest	South Carolina, USA	0.1	9
Temperate soils	Germany	12	10
Temperate agricultural field	Japan	6.0	11
Temperate agricultural field	New York, USA	1.9	7
Temperate agricultural field	Denmark	0.2	6
Temperate agricultural field	Germany	1.1	10
Mediterranean oak Litter	France	0.15 - 0.6	12

Only those studies using techniques capable of enumerating viable population sizes are included.

*Population expressed as # cells g⁻¹ (wet weight) of peat.

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Table S4. CA activity of soils, microorganisms, and leaves

	Conditions of growth	Reference compartment	Assay type	Enhancement factor at 25°C	Ref/source
Soil					
Peat from phytotron pot, lower soil layer	Ambient CO ₂	Bulk soil	¹⁸ O	4	This study
Peat from phytotron pot upper soil layer with algae	Ambient CO ₂	Bulk soil	¹⁸ O	44	This study
Subtropical karst forest soils in China (mean value)	Ambient CO ₂	Bulk soil	pH	50 ^a	1
Bacteria					
GLCa103	Agar, CaCO ₃	Cell volume	pH	7,500 ^b	1
GLCa104	Agar, CaCO ₃	Cell volume	pH	6,200 ^b	1
NLCa601	Agar, CaCO ₃	Cell volume	pH	7,900 ^b	1
NLCa602	Agar, CaCO ₃	Cell volume	pH	8,500 ^b	1
<i>Methanosarcina barkeri</i>	Anaerobic, pH6.4	Cell volume	pH	2,600 ^b	2
Fungi					
JFSP303		Cell volume	pH	4,300	1
NLPCa201		Cell volume	pH	7,300	1
Algae*[§] and cyanobacteria[§]					
<i>Chlamydomonas reinhardtii</i>	Ambient CO ₂	Protoplast	¹⁸ O	94,000 ^c	3
<i>Chlamydomonas reinhardtii</i>	5%	Protoplast	¹⁸ O	4,200 ^d	3
<i>Chlorella ellipsoidea</i>	Ambient CO ₂ , pH5.5	Cell volume	pH	8,400 ^e	4
<i>Chlorella ellipsoidea</i>	Ambient CO ₂ , pH7.2	Cell volume	pH	172,500 ^e	4
<i>Synechococcus</i> PCC7942	0.004%	Cell volume	¹⁸ O	1,800 ^f	5
<i>Synechococcus</i> PCC7942	1.0%	Cell volume	¹⁸ O	350 ^f	5
<i>Synechocystis</i> PCC6803	Ambient CO ₂	Cell water	¹⁸ O	770	This study
Plant leaves (as references)					
<i>Arabidopsis thaliana</i> L.	Ambient CO ₂	Chloroplast	¹⁸ O	113,000	This study
		Leaf water		13,000	This study
<i>Commelina communis</i> L.	Ambient CO ₂	Chloroplast	¹⁸ O	239,000 ^g	6
		Leaf water		8,600 ^g	6
<i>Nicotiana tabacum</i> L.	Ambient CO ₂	Chloroplast	¹⁸ O	362,500 ^h	7
		Leaf water		36,700 ⁱ	7

For the pH-type assay, CA activity values are usually expressed in Wilbur-Anderson (WA) unit defined as $x(t_{\text{uncat}}/t_{\text{cat}} - 1)$, where t_{uncat} and t_{cat} are times taken to attain a given drop in pH (usually ≈ 2 pH units) and x is an integer, originally taken equal to 1 (e.g., ref. 2) but also sometimes equal to 10 (e.g., refs. 1 and 7). To compare better with the ¹⁸O-type assay, we used the 2.34 conversion factor proposed by Price et al. (7): $2.34 U = 10/x \text{ WA}_{(x)}$, where U is the deviation of the catalyzed (k_{cat}) to the uncatalysed rate (k_{uncat}) and defined as $U = k_{\text{cat}}/k_{\text{uncat}} - 1$ and $\text{WA}_{(x)}$ is the CA activity given in WA units for a given x . The enhancement factor is then calculated as $f_{\text{CA}} = U + 1$. In studies using the pH-type assay, U is commonly expressed per mass of protein (or chlorophyll or dry soil). The enhancement factor f_{CA} at the reference compartment level (cell, bulk soil, . . .) is then computed as: $f_{\text{CA}} = U \cdot V_{\text{cuvette}} \cdot [\text{P}]_{\text{comp}} + 1$, where V_{cuvette} is the cuvette volume and $[\text{P}]_{\text{comp}}$ is the concentration of protein (or chlorophyll or dry soil, . . .) in the reference compartment. A Q_{10} value of 2 is also used to express all enhancement factor to the same temperature.

*Terrestrial species.

[§]Freshwater species (but strains of these species are resident in soils).

^aGiven a soil bulk density of 1.3 g·cm⁻³.

^bIntracellular activity only. Extracellular activity is usually lower. Given a bacterial protein content of 60 g/L.

^cGiven a low-C_i protoplast specific volume of 130 μL/mg Chl (3).

^dGiven a high-C_i protoplast specific volume of 290 μL/mg Chl (3).

^eGiven a specific cell volume of 77 μL/mg Chl (8).

^fGiven a specific cell volume of 60 μL/mg Chl (5).

^gGiven a leaf chlorophyll concentration of 1.2 g Chl/L, a specific chloroplast volume of 30 μL/mg Chl and a room temperature of 25°C (6).

^hGiven a specific chloroplast volume of 30 μL/mg Chl (7).

ⁱGiven a leaf water content of 90%.

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Table S5. Details of soil chambers used in this study

Site description	Site name	Location	Chamber type	Chamber area, m ²	Chamber height, m	Soil surface porosity, m ³ m ⁻³	Code
Mediterranean evergreen forest	Mitra	Portugal	Open	0.0572	0.03	0.54	MI
Subtropical deciduous forest	Oak Ridge	Tennessee, USA	Open	0.0487	0.27	0.66	TN
Subtropical evergreen forest	Tallahassee	Florida, USA	Closed	0.4225	0.24	0.6	FL
Temperate evergreen forest	Bray	France	Open	0.071	0.1	0.66	BR
Temperate evergreen forest	Canterbury	New Zealand	Open	0.0154	0.24	0.6	NZ
Temperate Montane evergreen forest	Niwot	Colorado, USA	Open	0.0439	0.1	0.5	NI
Semiarid grassland	Fort Collins	Colorado, USA	Closed	0.00785	0.1	0.6	FC

Table S6. Chamber flux data

Date	Air temperature, °C	Soil surface temperature, °C	Flow rate, l min ⁻¹	C _{in,r} , ppm	C _{out,r} , ppm	d _{in} (VPDB-CO ₂)	d _{out} (VPDB-CO ₂)	Soil surface moisture, m ³ m ⁻³	Code
9/6/2004	29.39	24.7	0.966	377.01	499.29	-1.3709	-1.7215	0.051	MI
9/6/2004	29.13	27.96	0.975	374.2	498.1	-1.2073	-1.4169	0.049	MI
9/6/2004	27.03	26.05	0.993	373.9	484.2	-0.6839	-1.084	0.05	MI
9/6/2004	19.81	22.16	1.01	402.2	479.8	-1.1205	-0.6243	0.052	MI
9/7/2004	28.47	20.21	1.003	374.15	451.48	-1.2682	-1.4082	0.054	MI
9/7/2004	30.21	23.9	0.975	374.75	459.08	-0.9145	-1.6917	0.052	MI
9/7/2004	21.71	22.33	1.021	425.51	476.08	-1.0794	-0.27	0.051	MI
7/16/1998	30	22	2.08232	372	434	-0.003	-1.573	0.147	TN
7/16/1998	30	22	3.45512	413	488	-0.633	-1.633	0.147	TN
7/16/1998	30	22	2.49032	359	468	0.027	-1.903	0.147	TN
6/2/2003	34	39	900	450.7	582.8	-0.99	-3.12	0.03	FL
10/1/2007	17.4	14.7	1.996	481.8	718.3	-0.94	-2.07	0.16	BR
10/1/2007	17.5	14.7	1.996	444.3	675.4	-0.98	-2.09	0.18	BR
10/1/2007	18.1	14.8	1.996	452.3	692.7	-0.63	-2.52	0.18	BR
10/1/2007	20.8	15.3	1.996	418.6	694.5	-0.49	-2.83	0.18	BR
10/1/2007	21.7	15.1	1.994	443.6	741.3	0.15	-2.15	0.17	BR
10/1/2007	19.1	15.3	1.995	530.6	838.1	-0.79	-2.1	0.17	BR
10/1/2007	18.5	15.7	1.994	504.6	766.2	-1.29	-2.13	0.17	BR
10/1/2007	18.7	15.5	1.995	459.2	738	-0.76	-2.42	0.17	BR
10/2/2007	17.2	15.5	1.995	462.8	742.6	-0.32	-2.96	0.19	BR
10/2/2007	17.7	15.8	1.995	444.4	736	-1.11	-3.4	0.19	BR
10/2/2007	18.2	15.5	1.995	463	731.7	-1.43	-3.12	0.18	BR
10/2/2007	19.5	15.9	1.996	430.2	733.5	0.09	-2.91	0.18	BR
10/2/2007	22.4	16	1.995	422.2	724	-0.14	-3.45	0.18	BR
10/2/2007	26.3	18.9	1.993	425.2	780.6	0.16	-3.17	0.18	BR
10/2/2007	22.7	18.2	1.994	499.4	830	-0.97	-3.19	0.18	BR
10/21/2005	12.6	11.6	0.444	394.1	514.5	-14.55	-14.58	0.191	NZ
10/21/2005	13.9	11.8	0.444	394.9	504.5	-14.66	-14.57	0.191	NZ
10/21/2005	15.1	12.1	0.438	396.7	496.7	-15.23	-15.01	0.191	NZ
10/4/1997	22.2	22.2	2.2576	366.9	475.8	-0.92	-4.44	0.25	NI
7/26/2001	22.76	26.81	480	417.29	634.61	-2.25	-6.94	0.03	FC
7/26/2001	15.46	22.09	480	517.48	558.77	-2.65	-1.65	0.03	FC
7/27/2001	21.05	17.68	480	393.28	525.02	-1.39	-2.06	0.03	FC

Table S7. Soil water isotope data for each site

Date	Soil depth, cm	d_{sw} (VSMOW)	Code
9/6/2004	2.5	-1.006	MI
9/6/2004	7.5	-6.475	MI
9/6/2004	22.5	-6.435	MI
9/6/2004	26	-5.594	MI
9/6/2004	2.5	-1.891	MI
9/6/2004	7.5	-5.919	MI
9/6/2004	12.5	-6.055	MI
9/6/2004	17.5	-4.667	MI
7/15/1998	1.5	-2.05	TN
7/15/1998	2.5	-3.79	TN
7/15/1998	4.5	-4.06	TN
7/15/1998	6.5	-4.71	TN
7/15/1998	8.5	-4.94	TN
7/15/1998	11.5	-4.72	TN
7/15/1998	16.5	-5.36	TN
7/15/1998	21.5	-5.88	TN
7/15/1998	31.5	-6.35	TN
7/15/1998	41.5	-6.03	TN
7/15/1998	51.5	-6.09	TN
7/15/1998	61.5	-5.76	TN
6/2/2003	1	2.174	FL
6/2/2003	9	-0.886	FL
10/2/2007	0	-2.19	BR
10/2/2007	0.5	-2.55	BR
10/2/2007	2.5	-2.34	BR
10/2/2007	3.5	-2.67	BR
10/2/2007	4.5	-2.24	BR
10/2/2007	5.5	-2.29	BR
10/2/2007	6.5	-2.28	BR
10/2/2007	7.5	-2.54	BR
10/2/2007	8.5	-2.68	BR
10/2/2007	9.5	-2.71	BR
10/2/2007	12.5	-3.31	BR
10/2/2007	17.5	-3.57	BR
10/2/2007	22.5	-3.36	BR
10/2/2007	0	-1.95	BR
10/2/2007	0.5	-2.2	BR
10/2/2007	1.5	-2.06	BR
10/2/2007	2.5	-2.22	BR
10/2/2007	4	-2.7	BR
10/2/2007	5.5	-2.43	BR
10/2/2007	6.5	-2.15	BR
10/2/2007	7.5	-2.41	BR
10/2/2007	8.5	-2.58	BR
10/2/2007	9.5	-2.84	BR
10/2/2007	12.5	-2.89	BR
10/2/2007	17.5	-3.26	BR
10/2/2007	22.5	-3.55	BR
10/21/2005	1	-7.14	NZ
10/21/2005	3	-12.16	NZ
10/21/2005	5	-13.24	NZ
10/21/2005	7	-13.46	NZ
10/21/2005	9	-15.32	NZ
10/4/1997	2.5	-7.4	NI
10/4/1997	3.5	-12.5	NI
10/4/1997	4.5	-14.7	NI
10/4/1997	6.5	-16.4	NI
10/4/1997	7.5	-16.8	NI
7/27/2001	1	-0.45	FC
7/27/2001	3	2.39	FC
7/27/2001	5	0.37	FC
7/27/2001	7	0.57	FC
7/27/2001	10	-2.05	FC
7/27/2001	15	-6.45	FC
7/27/2001	25	-9.39	FC
7/27/2001	50	-9.7	FC