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Potential environmental impacts of CO₂ leakage from the study of natural analogue sites in Europe

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Abstract

Sites of natural CO₂ leakage provide opportunities to study the potential environmental impacts of such leakage on near-surface ecosystems. As part of the FP7 RISCS (Research into Impacts and Safety in CO₂ Storage) project a geochemical, botanical and microbiological study have been conducted on a natural CO₂ vent in Florina, Greece and the findings are compared with the results drawn from Latera, Italy and Laacher See, Germany. Plant and microbial communities appear to have adapted to long-term CO₂ exposure. Therefore the findings may not be representative of the effects of potential leakage from man made storage sites.

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1. Introduction

The United Nations Intergovernmental Panel on Climate Change (IPCC) has concluded that 50% to 80% cuts in global CO₂ emissions by 2050, compared to 2000 levels, will be needed to limit the long-term global mean temperature rise to 2.0°C to 2.4°C [1]. The development of carbon capture and storage

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(CCS) technologies is considered as a potential option in the portfolio of required measures to stabilise atmospheric greenhouse gas concentrations at a time of rapidly growing energy needs. Additionally, public acceptance of CCS is widely recognised to be an important factor affecting the feasibility of commercial deployment of this climate change mitigation option. Consequently, the health, safety, and environmental risks associated with potential CO₂ leaks from storage reservoirs need to be evaluated and appropriate monitoring programmes established to provide confidence in the technology.

The effects of naturally occurring CO₂ leaks on vegetation and soil inhabiting organisms have been addressed in relatively few studies, including Mammoth Mountain (California, USA) [2] and Stavesinci (NE Slovenia) [3], [4]; however, considerable work has now been undertaken in Europe at sites in Italy (Latera) [5] and Germany (Laacher See) [6]. The current study assesses the impacts of a natural CO₂ vent on a pasture field located within the Florina basin in Greece, the first such study in a European mountainous environment. Similar methods to those used at Latera and Laacher See were used at Florina and the results from all three sites are compared in this paper. The goal of this study is to integrate the botanical, geochemical and microbial results from these natural sites to better understand how local differences can influence and potentially modify the impact of a CO₂ leak.

The Florina basin is a NNW-SSE trending graben in the NW part of Greece formed by Neogene coarse clastic sediments. The overlying, 800-1000 m thick lacustrine deposits consist of conglomerates, marls, sandstones, and marl limestones with clayey caprocks. Natural CO₂ accumulations alternate with lignite seams in these sediments at various levels. Carbonate rich springs and CO₂-rich gas vents occur throughout the Florina basin, resulting from a slow upwelling of magmatic-hydrothermal CO₂ along faults and fractures. In contrast, the other two sites both occur in extinct volcanic calderas. Here, CO₂ is produced via upper mantle degassing (Laacher See, [6]) or thermo-metamorphic reactions (Latera, [7]), after which it leaks towards the surface along fault and fracture systems and is released to the atmosphere from spatially restricted gas vents.

Research into the impact of CO₂ leakage on the Florina pasture ecosystem consisted of two phases. First, a regional grid of soil gas and CO₂ flux measurements was conducted in July 2011 to define the leakage distribution and orientation. Second, in September 2011, a detailed, 25m-long transect was chosen based on these results and measured every 0.5 m for soil gas concentrations, CO₂ flux, botany, and soil microbiology. Methods used were similar to those applied at both Latera [5] and Laacher See [6], thereby facilitating comparison.

2. Methods

The CO₂ flux measurements were performed using the closed loop accumulation chamber method [5]. This method involves the accumulation of CO₂ that is migrating from the soil to the atmosphere within an open-faced chamber; the increasing concentration over time within this chamber (measured using a non-destructive infrared analyser) combined with the physical dimensions of the chamber itself are used to calculate the flux of CO₂ for that point. Soil gas samples were collected using a 6 mm diameter stainless steel probe that was pounded to the desired depth using a co-axial hammer. On the regional grid samples were collected only at 70 cm depth, whereas those along the transect were sampled at 20, 50 and 70 cm depth. All points were analysed directly in the field for CO₂ using a hand-held infrared analyser, all points along the transect were analysed in the field for O₂, CO, and H₂S, while all 70 cm deep samples along the transect and a portion of the regional grid samples were analysed in the laboratory for CO₂, O₂+Ar, N₂, light hydrocarbon gases using a gas chromatograph and total helium using a mass spectrometer.

The botanical survey was conducted along the entire length of the transect, visually defining the percentage cover of identified plant groups at 0.5 m intervals using a 0.5 m x 0.5 m quadrat. Field flora books were used to identify plant species [8] and digital photographs were taken of each quadrat for a complete visual record. To support these results, water content measurements were also conducted in soil collected at depths of 15-30, 45-60 and 65-70 cm, once every metre.

To assess the CO₂ induced effects on microbial metabolic pathways and populations, samples were collected at 65-70 cm depth at three transect points having CO₂ concentrations that were high (100% at 1.25 m within the vent core), moderate (8-10% at 11.25 m within the transitions zone) and low (<0.9% at 20.25 m within background conditions). The aim is to investigate potential metabolic rates of e.g. anaerobic methane production in soil incubations and the microbial community composition using quantitative real time polymerase chain reaction analyses (qPCR). The qPCR analyses first involved the extraction of high molecular weight DNA from frozen soil samples via cell lysis and DNA purification. Specific fluorescent probes were used targeting the ubiquitous 16S rRNA genes of bacterial or archaeal organisms, which were then quantified by real-time qPCR. The anaerobic methane production activity was determined using a gas chromatograph equipped with a flame ionization detector, as described in [9], [12].

3. Results

3.1 Soil gas concentrations and CO₂ flux

The regional survey results clearly defined NE-SW and NW-SE anomaly trends that align parallel to the main faults of the Florina Basin. Soil gas CO₂ anomalies defined wide features, whereas soil gas CH₄ and He were spatially more restricted in correspondence with CO₂ flux anomalies. The anomalies consist of individual gas vents formed by a core with high flux and CO₂ concentrations near 100% surrounded by a transition zone of gently decreasing CO₂ concentrations but rapidly decreasing CO₂ flux rates. Reactive soil gas species like CH₄ are only found within the essentially anoxic core of the vents. Based on this distribution, anomalies of soil gas CO₂ tend to merge because of their wider extent whereas those of CO₂ flux and soil gas CH₄ tend to be more isolated. These results were used to choose an appropriate gas vent on which to conduct the transect.

The gas geochemistry results from the transect are presented in Figure 1. The CO₂ flux data are presented with both linear and log scales (Fig. 1a), illustrating three distinct zones: 0-3.25 m in the vent core with very high values from 2,000 to 10,000 gm⁻²d⁻¹; 3.25-8.25 m where values drop rapidly from 700 to about 40 gm⁻²d⁻¹; and 8.25-25 m where values remain relatively stable from 40 to 20 gm⁻²d⁻¹. Whereas soil gas CO₂ at 20 cm follows a similar pattern, in that it reaches near background values by about 10 m (Fig. 1b), concentrations at 50 and 70 cm do not reach background levels (c. 1%) until about 20 m from the start of the profile. CO₂ concentrations in the core of the gas vent for the two deeper samples are near 100% while those at 20 cm depth are often over 90% (Note that laboratory analyses of the 70cm deep samples gave slightly lower CO₂ concentrations in this interval, likely due to the qualitative nature of field-based IR detectors in the high concentration range). The O₂ concentration values, not shown, are the exact opposite to those of the CO₂, with conditions being essentially anoxic in the vent core. Finally, both CH₄ and He (Fig. 1c) sampled at 70 cm depth exhibit highly anomalous values in the vent core (from 0-7m), followed by a very rapid drop to background values over less than a metre. This abrupt transition interval corresponds with CO₂ concentrations <50%, CO₂ flux values <500 gm⁻²d⁻¹, and O₂ concentrations

>9%. All parameters show small-scale spatial variations, illustrating the spot nature of gas release as controlled by local, near-surface variability in soil permeability.

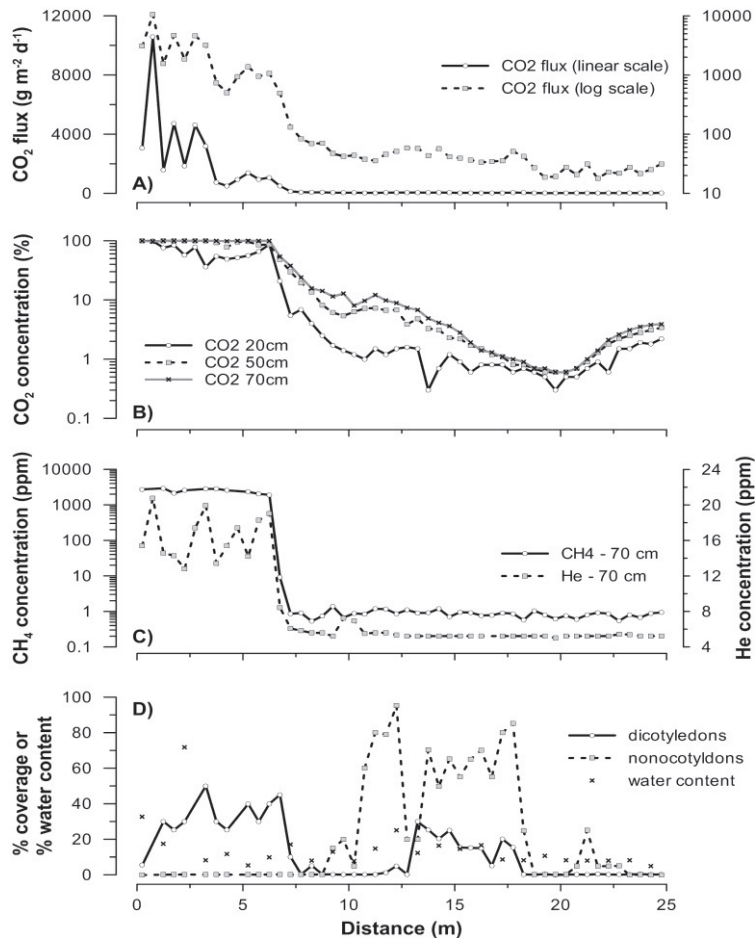


Fig. 1. Results obtained from the 25 m long transect at the Florina study site: A) CO₂ flux plotted on both linear and logarithmic scales; B) soil gas CO₂ concentrations at 20, 50, and 70 cm depths measured with a hand-held infrared analyser; C) soil gas CH₄ and He at 70 cm depth; and D) plant cover and soil water content.

3.2 Botanical surveys

The botanical survey showed a long-term adaptation of some plant species to the hypoxic or anoxic conditions of the naturally CO₂-enriched environment (Fig. 1d). From the centre of the vent to 8.5 m along the transect, where CO₂ concentrations range between 99.7-4 % at 20 cm depth and 99.9-13 % at 50 cm depth, only dicotyledonous plants *Minuartia glomerata* and *Polygonum aviculare* were observed and thus can be considered as bioindicators of high CO₂ in soil. The CO₂ flux range here varied from around 70 to 10.600 g/m²/d with a slight correlation coefficient of -0.63 with the soil CO₂ concentration. The soil oxygen decrease below 10% in this interval (strong negative relationship between soil [CO₂] and [O₂]

with a Pearson correlation coefficient of -0.99) indicates the ability of these plant species to adjust respiration (metabolism) to oxygen deficiency and high CO₂ conditions [3]. At a distance of 9 m from the centre of the vent, concentrations of CO₂ reach the background values of <2% at 20 cm depth. At this point, monocotyledonous plants (such as *Cynodon dactylon*) predominate and *Minuartia glomerata* is not observed although other dicotyledonous plants (e.g. *Astragalus*) are present.

Soil water content (Fig. 1d) at the depth of 15-30 cm is relatively constant along the transect indicating limited effect of elevated soil CO₂ concentrations on soil moisture at the depth of 15-30 cm. Many studies have shown that at sites where elevated CO₂ concentrations limited plant growth, the moisture of the upper soil layers remain higher compared to soils with normally developed vegetation due to the lack of transpirational water loss [10]. Other studies suggested that soil moisture can decrease due to increased plant growth and root biomass depending on the plant species [5].

3.3 Microbiological analyses

Analyses of the CH₄ production rates revealed the influence of different CO₂ concentrations on the methane production rates. As it can be noticed in Fig. 2 these rates were with 0,016 μmol gdw⁻¹ d⁻¹ at the CO₂ vent (100% CO₂) three times higher than at the medium site with 8-10% CO₂ while no methane production could be detected at the reference site (<0,9% CO₂).

In addition to the activity analyses, cell numbers of *Bacteria* and *Archaea* were determined using qPCR. Trends of increasing copy numbers along the transect corresponding with decreasing CO₂ concentrations (Fig. 2) could be observed for both, *Bacteria* and *Archaea*. The numbers of *Bacteria* increased from 3,0*10⁹ to 3,7*10⁹ gene copies gdw⁻¹ of soil from the reference site and vent centre and those for *Archaea* increased from 7,6*10⁸ to 1,5*10⁹ gene copies gdw⁻¹. Similar trends have been observed for both, activity and community analyses at Laacher See [6] and Latera [12].

The results so far show changes in the activity and community composition as a consequence of continuous CO₂ release. The successive replacement of O₂ with CO₂ in the soil along possibly leads to an increase of important anaerobic metabolic pathways like methane production, sulfate reduction and the anaerobic CO₂ production (mineralization). In fact, increasing CH₄ production rates with increasing CO₂ concentrations indicate high abundance of anaerobic methanogens [12], [6]. In contrast, the gene copy numbers of *Archaea* showed no increase with high CO₂ concentrations. Copy numbers of *Archaea* were highest at the reference site with background CO₂ concentrations and were lower at the vent centre. Similar results could be observed for *Bacteria*. One explanation for the detected results might be a CO₂ induced shift in the microbial community composition for the benefit of anaerobic microorganisms which successively replace aerobic microorganisms. The cell numbers won't be necessarily affected by changes within the populations.

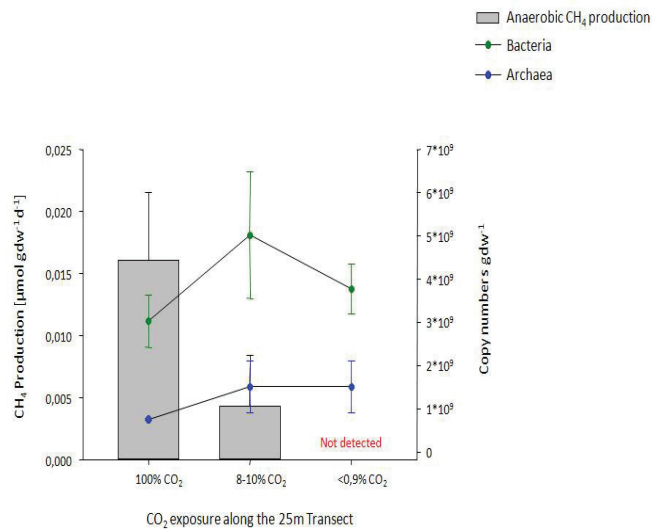


Fig. 2: The variation of the microbial activity and community composition in the soil samples of the CO₂ vent, the transition zone and the reference site of the transect in Florina basin

4. Discussion

Near surface gas geochemistry results along the Florina transect are in many respects very similar to those observed at the Latera site, including a gas vent core with high CO₂ flux, high CO₂, CH₄, C₂H₆, and He concentrations, and very low O₂ levels that is surrounded by a transition zone where flux rates decrease rapidly, concentrations of CH₄, C₂H₆, and He decrease very abruptly over an interval of less than 1 metre, and CO₂ concentrations (at least at the deeper levels) decrease relatively slowly. As discussed in [5], the wider CO₂ concentration distribution is probably due to the dense nature of this gas relative to air, the very narrow distribution of He is due to its very low density, whereas that of CH₄ is likely related to the minimum CO₂ flux rate needed to maintain O₂ concentrations low enough to prevent microbial oxidation. The one principle difference observed between these two sites is the high levels of H₂S in the gas vent core at Latera, in contrast to its almost complete absence at the Florina site. This fact removes one extra variable that could potentially impact on plant growth, thus focusing this work on the impact of elevated CO₂ values alone.

At the Laacher See, soil gas and CO₂ flux measurements present similar findings. Three main zones of higher CO₂ concentrations and fluxes were identified along the traverse: Between locations 11–17 m, 21–28 m, and 30–42 m, the latter representing the so-called centre of the vent where concentration values were greater than 90 vol% CO₂ in 60 cm depth. As the extension of natural CO₂ vents is limited, in less than 15 m from the centre of the vent CO₂ concentrations and fluxes decrease sharply to background values. [9]. Helium anomalies are narrower compared to CO₂ ones, similar to gas geochemistry results along the Florina and Latera transects. The results of the gas surveys conducted at Laacher See highlight also the occurrence of several CO₂ releases along an N-S axis, which is consistent with the known striking thrust of the basement [13], [9].

The observation that dicotyledonous species appear to be more tolerant to very high CO₂ concentrations than monocotyledonous plants in this adapted CO₂ rich environment is similar to the findings from the CO₂ vent at the Laacher See where *Polygonatum arenastrum* was the only observed dicotyledonous plant between 25 and 50 m along the transect when CO₂ concentrations were between 10–35% at 15 cm depth [6]. However, the botanical survey at Latera showed that the greatest impact on vegetation was observed within 6 m of the vent centre, where CO₂ concentrations at 20 cm depth were over 95%, trace reduced gases (CH₄, H₂S, and H₂) present and CO₂ flux rates exceeded 2000–3000 g m⁻² d⁻¹. However, at this site, monocotyledonous plants (e.g. grasses) appear to tolerate elevated CO₂ concentrations [5].

As part of the RISCs project similar observations at the controlled injection sites Asgard Field Site, UK and Grimsrud, Norway suggest that monocotyledonous plants (e.g. grasses) are inherently more tolerant to elevated CO₂ [11].

The microbial activity measurements revealed differences in their spatial distribution for CO₂-rich and reference soils. The reduced aeration of the soil due to elevated CO₂ concentrations and fluxes in CO₂ vents, and the consequently anaerobic and more reduced conditions in the soil lead at the CO₂ vent in Latera to strongly decreasing microbial cell numbers with increasing CO₂ concentrations [12]. On the other hand, anaerobic microbial activities and communities, like sulphate-reduction or methanogenesis, were positively correlated with higher CO₂ levels. The same was observed for the Florina soils, where methane production was strongly stimulated. However in contrast to the studies at Latera, the quantification of 16S rRNA genes in the surface soils at the Florina as well as the Laacher See site showed only minor differences in the overall distribution of archaeal and bacterial gene copy numbers between the sites. In summary, these results indicated a CO₂-induced shift in the environmental conditions leading to more anaerobic and potentially acidic microenvironments.

5. Conclusion

Impacts of CO₂ gas vents on the near-surface ecosystem of natural analogues appear to be highly localized under the prevailing extreme CO₂ regime. The spatial variations of soil CO₂ concentration and fluxes are generally controlled by a complex interplay of factors such as the migration pathway at depth, the physical and chemical properties of the vadoze zone, the features and hydrogeology of the near surface deposits etc. Therefore, soil gas analyses and flux measurements will be required for the baseline characterization of the migration pathways for soil gases and for the design of any near-surface gas geochemistry monitoring program of CO₂ geological storage sites.

The botanical results indicate species-specific response to high CO₂ concentrations depending on soil properties, mineralogical composition and temporal and spatial soil [CO₂] patterns. However the speed of ecosystem's reactivity to short-term leaks in non-adapted systems such as a CO₂ geological storage site is not well investigated up to now.

The outcome from the microbiological analyses at the terrestrial CO₂ vent sites suggest changes in the soil microbial community and significant environmental changes caused by high CO₂ concentrations. Elevated CO₂ concentrations will lead to a shift in the microbial community composition towards anaerobic and acid tolerant microorganisms as well as an ecosystem adaption to the CO₂ induced soil biogeochemistry. This included significant effects on metabolic rates but less pronounced effects on the bacterial and archaeal cell numbers. Further activities will focus on the diversity of anaerobic Bacteria

and Archaea as well as the detection and identification of important soil microorganisms like ammonia oxidizing microorganisms and fungi at the different CO₂ concentrations [12], [6].

Naturally occurring CO₂ deposits provide unique natural analogues that can be used to assess the resulting impacts of the gas release on human health and safety, ecology, surface water, groundwater and the effectiveness of remedial measures. However, at these environments the existing ecosystems have been exposed to elevated CO₂ for considerably long periods, thus the plant/tree species may have adapted. Therefore, the findings from these sites may be either typical of and transposable to various different sites or they may not exhibit characteristics similar to the conditions that would occur with potential leaks from anthropogenic CO₂ storage [11].

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