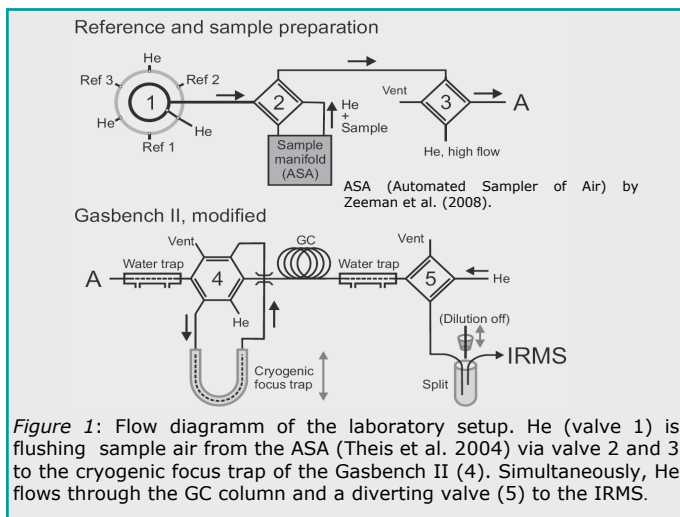


Measurement of $\delta^{13}\text{C}$ of atmospheric CO_2 on a routine basis

Introduction:

The $\delta^{13}\text{C}$ value of CO_2 in canopy air provides information about physiological processes underlying biosphere-atmosphere net CO_2 exchange. Since CO_2 from "background air" and respiration processes have different $\delta^{13}\text{C}$ values, these CO_2 sources can be distinguished, and the coupling of terrestrial and atmospheric carbon fluxes can be addressed. The "Keeling Plot" approach (regression of the $\delta^{13}\text{C}$ of CO_2 to its inverse $[\text{CO}_2]$) can be used to determine the $\delta^{13}\text{C}$ value of ecosystem-respired CO_2 with the possibility to partition net CO_2 exchange into assimilation and respiration. This implies the precise and accurate determination of $\delta^{13}\text{C}$ in CO_2 in large numbers of air samples in order to assess temporal and spatial variability within an ecosystem.

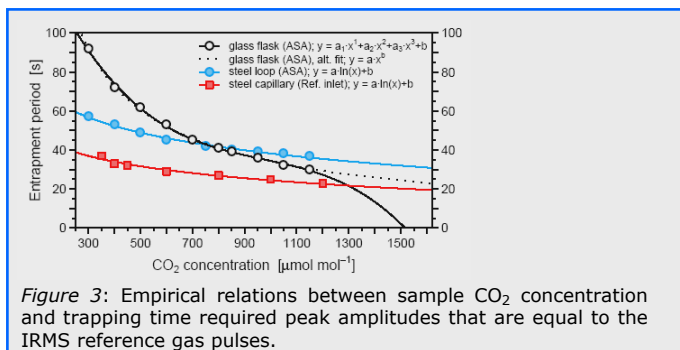
15th WMO/IAEA Meeting of Experts on Carbon Dioxide

Problem:

Linearity tests with gases with different CO_2 mixing ratios have shown a strong relationship between peak amplitude and corresponding δ -values.

Possible reasons:

- Signal-to-noise ratio
- Maximum linearity deviation of standard pulses tolerated by Finnigan MAT is 0.06 ‰/V
- Small memory-effect in the ASA tubing system ?



Results:

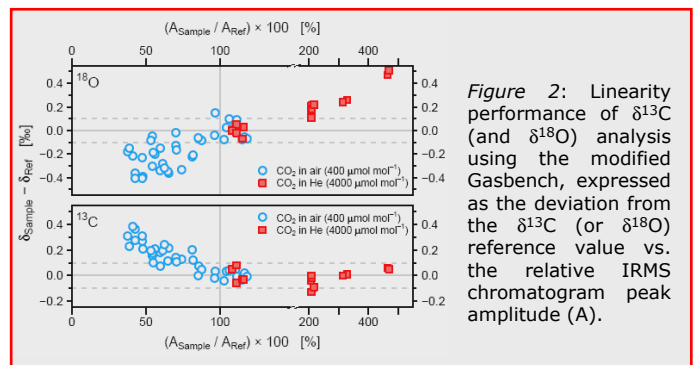
- The overall precision of $\delta^{13}\text{C}$ measurements of CO_2 was determined to be <0.08 ‰ (σ) for samples with standards stored in glass flasks inside an ASA ($n=33$) and <0.06 ‰ (σ) for directly supplied standards ($n=5$), over the course of several measurement campaigns between February 2006 and March 2008.
- The $\delta^{13}\text{C}$ -values of different mixing ratios of one identical source CO_2 ($n=12$) with synthetic air (from 300 to 1800 ppm) can be measured with a total precision of 0.04 ‰ (σ) using normalized peak amplitudes.

Literature:

DE Theis et al. (2004): Rapid Comm. Mass Spectrom. 18, 2106ff
RA Werner, WA Brand (2001): Rapid Comm. Mass Spectrom. 15, 501ff

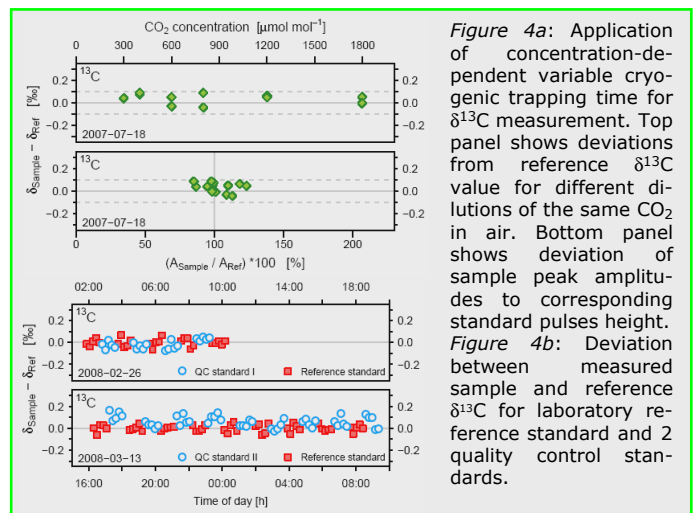
Laboratory setup:

- Sampling system ASA (Theis et al., 2004) modified
- Multiple reference gas inlets, sharing the flow path of the sample gas (Fig. 1, 1 + 2). Referencing after **I**dentical-**T**reatment principle (Werner and Brand, 2001) now possible
- Vents to purge the capillaries, release (over)-pressure (Fig. 1, 3 + 4)
- Modified Gasbench II with ConFlo III split allowing undiluted transfer of sample CO_2 to the IRMS
- Homebuilt cryogenic trap (Ni-wire in steel capillary)
- Automated $\text{N}_2(\text{l})$ refill system for cold trap controlled by ISL scripts
- Software optimisation (automated adaption of trapping time)



Solution:

Optimizing sample peak amplitudes close to the peak heights of the reference gas pulses by freezing the same amount of CO_2 for each sample, independent of $[\text{CO}_2]$. Implementing an ISL script which adapts trapping time relative to $[\text{CO}_2]$.



MJ Zeeman et al. (2008): Rapid Comm. Mass Spectrom. 22, 3883ff

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