Improving accuracy and precision of ice core δD(CH₄) analyses using methane pre-pyrolysis and hydrogen post-pyrolysis trapping and subsequent chromatographic separation

M. Bock, J. Schmitt, J. Beck, R. Schneider, and H. Fischer
Climate and Environmental Physics, Physics Institute and Oeschger Centre for Climate Change Research, University of Bern, Sidlerstrasse 5, 3012 Bern, Switzerland

Correspondence to: M. Bock (bock@climate.unibe.ch)

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Abstract. Firn and polar ice cores offer the only direct palaeoatmospheric archive. Analyses of past greenhouse gas concentrations and their isotopic compositions in air bubbles in the ice can help to constrain changes in global biogeochemical cycles in the past. For the analysis of the hydrogen isotopic composition of methane (δD(CH₄) or δ²H(CH₄)) 0.5 to 1.5 kg of ice was hitherto used. Here we present a method to improve precision and reduce the sample amount for δD(CH₄) measurements in (ice core) air. Pre-concentrated methane is focused in front of a high temperature oven (pre-pyrolysis trapping), and molecular hydrogen formed by pyrolysis is trapped afterwards (post-pyrolysis trapping), both on a carbon-PLOT capillary at −196 °C. Argon, oxygen, nitrogen, carbon monoxide, unpyrolysed methane and krypton are trapped together with H₂ and must be separated using a second short, cooled chromatographic column to ensure accurate results. Pre- and post-pyrolysis trapping largely removes the isotopic fractionation induced during chromatographic separation and results in a narrow peak in the mass spectrometer. Air standards can be measured with a precision better than 1 ‰. For polar ice samples from glacial periods, we estimate a precision of 2.3 ‰ for 350 g of ice (or roughly 30 mL – at standard temperature and pressure (STP) – of air) with 350 ppb of methane. This corresponds to recent tropospheric air samples (about 1900 ppb CH₄) of about 6 mL (STP) or about 500 pmol of pure CH₄.

1 Introduction

Methane (CH₄) is a potent greenhouse gas, which shows increased atmospheric concentrations since the industrial revolution (Intergovernmental Panel on Climate Change, 2007). A recent assessment of the present-day methane budget is presented in Kirschke et al. (2013). However, the atmospheric load of CH₄ has varied on various timescales. A wealth of information has been gained from concentration measurements regarding annual (Dlugokencky et al., 1995), decadal (Mitchell et al., 2011), and millennial up to glacial-interglacial (Loulergue et al., 2008) CH₄ variability. Stable isotope data of methane on recent air samples (e.g. Quay et al., 1999) and on the past atmosphere using ice cores (e.g. Ferretti et al., 2005; Fischer et al., 2008; Sowers, 2010; Sapart et al., 2012; Möller et al., 2013) provide further insight into processes and sources controlling the global methane cycle. For instance, knowledge of the temporal evolution of the hydrogen isotopic composition of methane (δD(CH₄) or δ²H(CH₄)) over the termination of the last ice age (14,000–18,000 years before present) (Sowers, 2006) as well as during rapid warming events between 32,000–42,000 years before present (Bock et al., 2010b) made it possible to reject the “clathrate gun hypothesis” proposed by Kennett et al. (2003) as the trigger for the steep atmospheric methane increases.

However, we are still far from a complete picture of the biogeochemistry of methane in the past. Ice core isotope studies on δD(CH₄) have the potential to improve our understanding of the global CH₄ cycle but are still scarce due to analytical difficulties (e.g. Bock et al., 2010a; Sapart et al., 2011) and the large sample amount needed. To date, the
few published ice core δD(CH₄) studies required from 0.5 kg (Bock et al., 2010b) to more than 1 kg (Sowers, 2006; Mischler et al., 2009) of ice from multi-parameter deep ice cores with a typical precision of around 3 to 4 ‰. This error bar is still large in view of the observed natural variability, which is rather small: about 30 ‰ for glacial–interglacial and 20 ‰ for rapid changes during the last glacial (Sowers, 2006; Bock et al., 2010b). This study presents new developments based on Bock et al. (2010a) to improve precision and accuracy and significantly reduce the sample size for (ice core) δD(CH₄) measurements.

2 Experimental

We present an improved continuous-flow gas chromatography (GC) pyrolysis (P) isotope-ratio monitoring mass spectrometry (irmMS) system (GC/P/irmMS) designed to analyze δD(CH₄) from (ice core) air samples (Fig. 1) with high precision. In the following we give a short summary of our previous instrumentation (Bock et al., 2010a) and new developments concerning the physical system and data processing.

The most important new features presented here are pre- and post-pyrolysis trapping (pre&postPT) of CH₄ and molecular hydrogen (H₂), respectively, and subsequent gas chromatographic separation using a cooled porous layer open tubular (PLOT) column, which improve accuracy and precision and reduce the required sample amount considerably. A systematic dependency of δD(CH₄) on the amount of CH₄ (signal dependency) is observed but can be precisely corrected for. The successful implementation of pre&postPT requires as a prerequisite better purification of helium. A new calibration software tool has also been developed, enabling a one-step correction of system drifts over time and signal dependency (linearity) in an iterative way.

Note that the current status of our set-up has been reached in separate steps over the last few years. The different states are summarized in Table 1 and Fig. 1, named after the year(s) and depicted in different colours. Until 2010 no pre&postPT and subsequent gas chromatographic separation was performed. In the years 2011 and 2012, we took advantage of post-pyrolysis trapping and basic GC separation afterwards. In 2013 we also implemented pre-pyrolysis trapping but still used the same basic second GC. Furthermore, since the beginning of 2014 an enhanced, cooled 2nd GC is additionally operated.

2.1 Instrumentation

The system is fed by helium (He) (Alphagaz I, 99.9990% purity; Carbages, Switzerland), which is purified using a high-capacity gas purifier and an inline gas purifier (both Supelco, Bellefonte, PA, USA). In addition to the description given by Bock et al. (2010a), we further purge the He used for the complete system in a 3 m long 1/4 in. stainless steel (SST) capillary (i.d. 5.3 mm) filled with charcoal (grain size 0.3–0.5 mm, 0.41 g cm⁻³, Fluka 05112; Sigma-Aldrich Chemie, Steinheim, Germany). This trap is immersed in liquid nitrogen (LN) during the week and can be vented at room temperature over the weekend using valve V0 in Fig. 1, a pneumatic six-port, two-position valve (1/16 in. fittings, 0.4 mm port diameter, Valcon M rotor; Valco, VICI AG, Schenkon, Switzerland). The additional He purification cold trap lowers the blank CH₄ contribution considerably compared to Bock et al. (2010a). Extracting remnant gas in the sample cylinder (containing melt water) after an ice sample results in an H₂ peak 0.5 % the area of a small CH₄ peak characteristic for the LGM (Last Glacial Maximum – CH₄ concentration of 350 ppb – about 20 ka before present (BP), where present is defined as 1950; Loulergue et al., 2008; Clark et al., 2009). In comparison, the same procedure in the old system led to a 1 % area of a LGM peak; thus, blank CH₄ contribution appears to be reduced by 50 % for the improved set-up. To factor out the blank contribution from the extracted sample and, for example, the sample cylinder walls, we trapped background gases (from the He carrier gas) on T2, thus bypassing the sample cylinder, which resulted in a peak area of only 0.2 %.

The following sample preparation steps are similar to Bock et al. (2010a): a glass vessel containing an ice core sample is evacuated, and the enclosed air is released upon melting the ice. In a high flow (He, 500 mL min⁻¹), water vapour is removed using a cooled Nafion membrane and a cold trap (T1) while the air sample is transferred to a trap filled with charcoal (T2) immersed in LN.

Contrary to Bock et al. (2010a), T1 is made up of an empty 1/8 in. tube of three coils that enter or leave a dewar maintained at −90 °C. Temperature controlled cooling of the dewar is achieved using LN droplets released into the dewar (Schmitt, 2006; Bock et al., 2010a). Only residual water vapour is removed by T1, while CO₂ is adsorbed on an Ascarite trap beforehand, made of a 10 cm 1/4 in. stainless steel tube. In this new set-up, N₂O is passed through the system and can be measured in the mass spectrometer.

Air reference injections are realized by switching V1, either mimicking an ice sample by introducing the air into the glass extraction vessel or by bypassing the sample vessel, depending on the position of V2. Following a switch of V3, the air sample is transferred from the charcoal trap to a trap filled with Hayesep D (T3, at −100 °C), where methane is quantitatively trapped while the bulk air (N₂, O₂, Ar) is vented. Residual air components and CH₄ are focussed on T4 (three coils of a GC column (CP-PoraBond Q, 0.32 mm i.d.) at −196 °C) and injected onto a GC column (Carboxene 1010 PLOT column (30 m, i.d. = 0.32 mm)). Valve V5 is switched to route the sample through a new cold trap T5 and towards the pyrolysis furnace only for the time window in which CH₄ is leaving the GC column. T5 replaces Nafion-2 of the old set-up, which was less effective in removing water before the pre- and post-pyrolysis trapping steps, and consists of a
Figure 1. Flow scheme of the new $\delta D(CH_4)$ system including pre- and post-pyrolysis trapping of methane and hydrogen, respectively, and a second short, cooled chromatographic separation column (2nd GC) after the last trap. The coloured boxes highlight the major differences compared to Bock et al. (2010a). Red are the changes made in 2011–2012, blue represents additional changes carried out in 2013 and green in 2014 (compare Table 1). Capillaries are stainless steel (SST) or fused silica (FS), with the latter occurring only in the low-flow part. Inside the GC we use a Carboxene 1010 PLOT column (30 m, i.d. = 0.32 mm). Cold traps are “He purifier” (charcoal, 1/4 in.), T1 (open tube, 1/8 in.), T2 (charcoal, 1/4 in.), T3 (Hayesep D, 1/8 in.), T4 (CP-PoraBond Q, i.d. = 0.32 mm), T5 (open untreated capillary, i.d. = 0.53 mm), T6, T7 and 2nd GC (all GS-CarbonPLOT, i.d. = 0.32 mm). The CO$_2$ and H$_2$O trap after the pyrolysis oven consists of a piece of untreated capillary (i.d. = 0.32 mm) immersed in LN.

U-shaped piece of untreated capillary (i.d. = 0.53 mm) placed in a well-insulated dewar cooled by LN droplets to $-90^\circ$C.

In the following we describe the main new developments. Eluting CH$_4$ from the GC is focused on T6 for 18 s (pre-pyrolysis trapping, prePT) before it is released by passive warming to room temperature. Subsequently, the focussed pulse of CH$_4$ is pyrolysed as described by Bock et al. (2010a), but the produced H$_2$ and the pyrolysis side products are not allowed to enter the mass spectrometer (Isoprime, Elementar Analysensysteme, Hanau, Germany) directly. Instead, the pyrolysis products are trapped on T7 for 40 s (post-pyrolysis trapping, postPT). Both traps – T6 and T7 – are U-shaped, 20 cm long GC columns (GS-CarbonPLOT, ID 0.32 mm, film 1.5 µm, Agilent Technologies, part number 113-3112) retaining CH$_4$ and H$_2$ (and other gases) at LN temperature. After postPT is complete, T7 is lifted out of LN and warmed to room temperature allowing H$_2$ to enter the mass spectrometer via an open split. After T7 the line has been extended by two meters of the same PLOT column (2nd GC in Fig. 1), of which 70 cm are cooled to $-80^\circ$C to ensure baseline separation of H$_2$ from other gases, which produce signals visible in both the $m/z$ 2 and 3 traces (Fig. 2). The column piece of the 2nd GC is cooled on a well insulated brass surface by two thermoelectric coolers/heaters (20 W, 25 g, PKE 36 A 001, Peltron GmbH, Fürth, Germany) with the heat sink (a copper plate) immersed in LN. Temperature is controlled by a thermostat (Jumo) to within 0.5 °C and can be set in a dynamic range of $-70$ to $-100^\circ$C. To prevent CO$_2$ and H$_2$O (eluting from the pyrolysis oven) from accumulating on the cold 2nd GC column, a short piece of untreated fused silica capillary is immersed in LN during the day. Valve V6 is used to bypass the pyrolysis furnace, traps T6 and T7, the CO$_2$ and H$_2$O trap, and the 2nd GC in order to ventilate water eluting from a warm T5 trap.

Note that in an earlier version, our system was only extended by post-pyrolysis trapping, while pre-pyrolysis trapping was implemented later (Table 1 and Fig. 1). The latter is of potential interest as CH$_4$ and CDH$_3$ experience different retention on GC columns (e.g. Bock et al., 2010a), leading to a “time shift” (Ricci et al., 1994) or “time displacement” (Meier-Augenstein, 1999). As both CH$_4$ and CDH$_3$ are held on T6, this pre-pyrolysis trapping step resets the chromatographic separation introduced by the GC, allowing for pyrolysis of a non-fractionated methane peak. Furthermore, prePT allows for a shorter post-pyrolysis trapping time, as H$_2$ from CH$_4$ pyrolysis elutes during a shorter time interval. This is advantageous because H$_2$ cannot be held on T7 for an extended time under the described conditions. Instead, strong chromatographic separation between H$_2$ and HD for a system using a long post-pyrolysis trapping time leads to strong intra-peak fractionation visible in a large time shift.

Pyrolysis of CH$_4$ is achieved using a custom-made high-temperature furnace (Bock et al., 2010a). Using a brand new thermocouple indicated that the optimal pyrolysis temperature in our case is 1400 °C. However, due to ageing of the thermocouple, the read-out of the temperature is considerably reduced over a time period of several months. To determine the optimal pyrolysis temperature, we introduce
CH\textsubscript{4} peaks via V7 (e.g. 3 times 10\,$\mu$L loop with roughly 500 ppb CH\textsubscript{4} in He) at different temperatures. We observe two plateaus at different temperatures, one for $\delta^{13}$CH\textsubscript{4} values and one for peak areas. The plateau of $\delta$D(CH\textsubscript{4}) at higher temperature is favoured for high-precision isotope measurements because the small but inevitable temperature fluctuations in the reactor then lead to smaller scatter in isotope values. Too high temperatures lead to shortened lifetimes of pyrolysis reactors which becomes noticeable through higher backgrounds of nitrogen and argon caused by ambient air. Typically, a reactor (stone-ware GmbH, Switzerland, DEGUSSIT\textsuperscript{®} Al\textsubscript{2}O\textsubscript{3}, length = 420 mm, i.d. = 0.5 mm, o.d. = 1.5 mm) facilitates reproducible results for about half a year. When a new reactor has to be installed, it is heated up using a ramp of 5 h and pre-conditioned over the course of a day by injecting 10\,$\mu$L loops of the previously mentioned mixture of 500 ppb CH\textsubscript{4} in He every 40 s (without using any trap).

As a second major improvement to the system following a development by Schmitt et al. (2014), we can now measure N\textsubscript{2}O concentration, $\delta^{15}$N and $\delta^{18}$O of N\textsubscript{2}O on the same sample. Therefore, after the H\textsubscript{2} acquisition for methane is completed, the pyrolysis reactor is bypassed using valve V6 and a peak jump is performed in order to focus the mass spectrometer to the N\textsubscript{2}O configuration measuring m/z 44, 45, and 46. After a second peak jump, we measure xenon (as $^{132}$Xe\textsuperscript{2+} and $^{136}$Xe\textsuperscript{2+}) using beams m/z 66 and 68. Xenon is considered a proxy for total air content and is used to calculate CH\textsubscript{4} and N\textsubscript{2}O concentrations. For detailed descriptions of N\textsubscript{2}O and Xe analytics, we refer the reader to a companion publication by Schmitt et al. (2014) reporting on a new system to simultaneously measure $\delta^{13}$CH\textsubscript{4}, isotopes of N\textsubscript{2}O, Xe and more trace gas concentrations.

2.2 Data processing

We use custom-made Python (http://www.python.org/) scripts to process the raw beam data, to organise peak data of references, standards and samples in specific libraries and to perform the calibration to the international VSMOW (Vienna standard mean ocean water) scale. The peak integration method is similar to that described by Bock et al. (2010a). Integration limits are found based on the major beam time series and also applied to the minor beam. In commercially available mass spectrometer software, the integration limits are determined based on the derivative of the beam time series according to thresholds of the slope (e.g. Ricci et al., 1994). We chose a different approach here: we determine the peak maximum and set the integration limits to fixed numbers of data points before and after the peak maximum, that is, we use a fixed peak width. This choice is not critical as an alternative peak evaluation using the commonly used start- and end-slope criteria (Ricci et al., 1994) led to the same results within the given error limits (re-evaluated data not shown). In contrast to our previous procedure, pre\&postPT removes the isotopic fractionation induced by the chromatographic separation resulting in nearly unfractionated H\textsubscript{2} peaks in the current set-up. Hence, we do not perform a time shift correction of the m/z 3 beam. Generally, the background is determined as the median of data points 6 s before the peak start (see Fig. 2).
Table 1. Time line of our measurement system. Shown are the key properties of the system that changed over the course of the past four years. Colours refer to successive changes indicated in Fig. 1 and are in line with Fig. 4.

<table>
<thead>
<tr>
<th>Name</th>
<th>Colour</th>
<th>2010</th>
<th>2011–2012</th>
<th>2013</th>
<th>2014</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reference</td>
<td>black &amp; white</td>
<td>Bock et al. (2010)</td>
<td>Post-pyrolysis trapping, separation afterwards</td>
<td>Pre- &amp; post-pyrolysis trapping, enhanced 2nd GC separation afterwards</td>
<td>As 2013 plus H₂ peak free of known interferences (except unavoidable H₂O formed in ion source)</td>
</tr>
<tr>
<td>Short description</td>
<td>GC-pyrolysis-IRMS</td>
<td>Post-pyrolysis trapping, basic 2nd GC separation afterwards</td>
<td>Pre- &amp; post-pyrolysis trapping, basic 2nd GC separation afterwards</td>
<td>Pre- &amp; post-pyrolysis trapping, basic 2nd GC separation afterwards</td>
<td>Pre- &amp; post-pyrolysis trapping, basic 2nd GC separation afterwards</td>
</tr>
<tr>
<td>Key property</td>
<td>All mentioned substances</td>
<td>Separation of unpyrolysed CH₄ and Kr from H₂</td>
<td>Separation of unpyrolysed CH₄ and Kr from H₂</td>
<td>Separation of unpyrolysed CH₄ and Kr from H₂</td>
<td>Separation of unpyrolysed CH₄ and Kr from H₂</td>
</tr>
</tbody>
</table>

In order to calibrate samples, it is essential to compare samples to reference measurements that are sufficiently stable over time and match the sample size. If this cannot be achieved, one has to correct for any drift and signal dependency (e.g. Schmitt et al., 2003; Potter and Siemann, 2004; Bock et al., 2010a; Brass and Röckmann, 2010). In our case this is essential, because we observe a clear signal dependency of the δD(CH₄) values (Fig. 3a). We note that the H₂ factor did not change compared to Bock et al. (2010a) and is accounted for during evaluation of chromatograms. Hence, residual signal dependency is due to processes upstream of the mass spectrometer (e.g. pyrolysis conditions). The observed signal dependency is stable and reproducible over long time intervals and can therefore be precisely corrected for without compromising the overall precision of the measurement (see Sect. 3.2). When a new pyrolysis reactor is installed, the signal dependency may change, and a new interval of our data analysis has to be started to account for this change. We developed a new software routine to correct for any system (time) drift and signal dependency simultaneously; this is presented in detail in the Appendix of this article and Fig. 3a and b. It takes standard measurements of known isotopic signature and iteratively fits parameters for (temporal) drift and signal dependency at the same time in order to minimize the standard deviation of δD(CH₄) of this reference air. The latter assumes constant signal dependency within a certain time period (typically a few weeks). The same assumption holds for laboratories determining signal dependency on a periodic schedule, but we see two advantages of our approach: (1) no extra day is needed to examine signal dependency and (2) if signal dependency changes slightly during the chosen time interval, this change is already accounted for by our reference measurements covering this interval. The fit parameters and daily mean values of our reference “Air Controlé” are used to calibrate the samples.

3 System performance

3.1 Accuracy

Our reference used to calibrate all samples is Air Controlé, a recent clean air tank (CH₄ concentration = [CH₄] = 1971 ± 7 ppb) for medical purposes (bottle 541659, filled February 2007 in Basel, Switzerland, Carbagas). Air Controlé was cross-referenced to −93.6 ‰ with respect to Vienna standard mean ocean water (VSMOW) using bottled air from Alert station “Alert 2002/11” (Bock et al., 2010a; Poss, 2003; Marik, 1998), previously measured at the Institute of Environmental Physics in Heidelberg (IUP, University of Heidelberg, Germany). At the IUP two scales for δD(CH₄) co-exist; one is based on mass spectrometric measurements of water-derived H₂ (named MAT), and the other is based on methane-in-air gases measured using a tunable diode laser system (named TDLAS). The anchors
are IAEA (International Atomic Energy Agency) water standards VSMOW and VSLAP (Vienna standard light Antarctic precipitation) for MAT and CH$_4$ in air standards, the latter originally calibrated by the Bundesanstalt fuer Geowissenschaften und Rohstoffe (BGR) Hannover, Germany for TDLAS (Bergamaschi et al., 1994, 2000). The two scales agree within 1.0‰ (MAT > TDLAS), which is within their precisions of 2.4‰ and 1.0‰ for MAT and TDLAS, respectively. $\delta D$(CH$_4$) for Alert was measured in Heidelberg twice using each method and was calibrated with respect to the mean of both scales ($-82.2\pm1.0‰$, C. Veidt, personal communication, 2014). Error propagation of the measurements performed in Heidelberg and Bern to get from primary standards to the value of Air Controlé, which we use to calibrate samples, leads to an uncertainty of 3.5‰.

In our previous report (Bock et al., 2010a), we presented 4 air samples (Dome 6, Dome 13, Groningen Air, NAT-332 air) in the $\delta D$(CH$_4$) interval $[-70, -110‰]$ in good agreement...
with Bräunlich et al. (2001) and younger measurements performed by the Institute for Marine and Atmospheric research Utrecht (IMAU) (Sapart et al., 2011), who so far use the same TDLAS scale as IUP. Two of these air samples (at the margins of the δD(CH₄) interval) have been re-measured with our improved set-up (Table 1, Fig. 1), again with good agreement (Table 3). Note that a new independent scale for δD(CH₄) is currently being established at MPI for Biogeochemistry, Jena. Accordingly, stringent Round Robin tests for methane isotopes will make it possible to check the agreement of different δD(CH₄) scales in the future.

In summary, we are confident that our measurements are close to the VSMOW scale (to about 3.5‰); however, we note and will show later in this section that effects due to differences in matrix and/or concentration of samples and references can hamper highly accurate results, while deviations are difficult to pinpoint for individual laboratories and inter laboratory comparison exercises.

As an update from Bock et al. (2010a), we introduce a new standard gas here: “Saphir 4” (bottle 4405, Carbagas, artificial clean air mixture with 761 ppb CH₄ and no krypton). Saphir injections through the melt water of a previously extracted ice core sample are slightly depleted in deuterium (ca. 2‰, see Table 3) compared to Saphir injections bypassing the sample container, but the mean values are within the combined error. Taking the small difference at face value, this fits to our expectation of preferential dissolution of the heavy isotope in the denser medium. If the extraction efficiency is equal for CH₄ and CDH₄ (or less for the heavier isotopologue), we expect lighter values for air standard injections flowing through melt water. Unfortunately, we cannot quantify the process, as there is no ice sample with known isotopic composition of the occluded air. Note, that any bias would only be relevant for intercomparison exercises with other labs, but would not influence the interpretation of time series or interpolar difference studies consistently carried out with our set-up. To conclude, it is not clear whether the aforementioned offset prevails for ice samples, or if the effect only occurs after ice sample extractions. We therefore chose not to correct for this (potential but) insignificant offset.

Note that our results for WAIS (West Antarctic ice sheet, core WDC05A, tube 184, depth range: 172.74–173.03 m, age approximately 410 a BP) are 15‰ more enriched in deuterium compared to data presented in Mischler et al. (2009). This offset is similar to the one observed for Boulder air (Bock et al., 2010a) compared to measurements performed at the Stable Isotope Lab of the Institute of Arctic and Alpine Research (INSTAAR, University of Colorado, Boulder, CO, USA) and reflects the fact that the laboratories in the US and Europe are tied to different primary standard air bottles. Note that no internationally accepted isotope reference material for CH₄ from air samples is yet available. At the time of writing, the aforementioned lab offsets are being addressed in a Round Robin organized by T. Sowers and E. Brook using WAIS ice and bottled air samples with varying methane concentrations.

Post-pyrolysis trapping and subsequent gas chromatographic separation enables the measurement of a pure H₂ peak in the mass spectrometer. Recently Schmitt et al. (2013) demonstrated that krypton (Kr) interference is possible during carbon isotopic analyses of CH₄. Conventional stable isotope analysis of CH₄ using GC-IRMS without postconversion separation leads to insufficient separation of CH₄ and Kr. In the case of δ¹³CH₄, the influence of Kr on δ¹³C analysis leads to a significant alteration of the results. Similarly, Meier-Augenstein et al. (2009) reported interference of N₂ for H₂ analyses. In the discussion version of this contribution, we thoroughly demonstrate the influence

<table>
<thead>
<tr>
<th>Substance</th>
<th>RT (s)</th>
<th>Origin</th>
<th>Influence on target beams</th>
<th>Visible on beam(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>H₂</td>
<td>0</td>
<td>pyrolysis-derived sample-CH₄</td>
<td>target (positive peak on both m/z 2 and 3)</td>
<td>m/z 2 and 3</td>
</tr>
<tr>
<td>Ar</td>
<td>9</td>
<td>collected background</td>
<td>negative signal on m/z 2 (none on m/z 3)</td>
<td>m/z 16, 32</td>
</tr>
<tr>
<td>O₂</td>
<td>20.4</td>
<td>CH₄ pyrolysis side product</td>
<td>positive signal on m/z 2 (none on m/z 3)</td>
<td>m/z 12, 16, 28 (44)</td>
</tr>
<tr>
<td>N₂</td>
<td>0.5</td>
<td>in situ production</td>
<td>positive signal on m/z 2 (none on m/z 3)</td>
<td>m/z 16, 17, 18</td>
</tr>
<tr>
<td>CO</td>
<td>9.3</td>
<td>in mass spectrometer</td>
<td>positive signal on m/z 2 (small on m/z 2)</td>
<td>m/z 15, 16</td>
</tr>
<tr>
<td>CH₄</td>
<td>185</td>
<td>only from Kr containing air samples</td>
<td>positive signal on m/z 2, negative on m/z 3</td>
<td>m/z 43</td>
</tr>
<tr>
<td>Kr</td>
<td>210</td>
<td>only from Kr containing air samples</td>
<td>positive signal on m/z 2, negative on m/z 3</td>
<td>m/z 43</td>
</tr>
</tbody>
</table>

Table 2. Sequence of detected species using the 2014 set-up (Fig. 1 and Table 1). Retention time (RT) is given relative to the H₂ peak maximum. Ar, O₂ and N₂ are not well-separated from each other under the described conditions. For CH₄ and Kr, the start of peaks is given, as the two substances elute very broadly from the cooled PLOT column. H₂O does not enter the mass spectrometer but is produced in situ (see text).
unpyrolysed CH$_4$ and Kr on the target beams m/z 2 and 3 when only a basic chromatographic separation is used after the conversion step (Bock et al., 2013). Based on that, we assess in the following the origin and influence of peaks showing up after the CH$_4$-derived H$_2$ peak when a cooled PLOT column is used for separation, subsequently referred to as post-peaks (Fig. 2). Implementation of the cooled 2nd GC revealed additional substances causing signals in the mass spectrometer either due to a direct effect or via the production of a third substance from the eluting substance plus background. The latter effect, occurring in the ion source, is named in situ production.

For pure CH$_4$ in He injections, signals on m/z 12, 14, 15, 16, 17, 18, 20, 28, 32, 40 and 44 are found for the respective focussing of the mass spectrometer (data not shown). In Table 2 we list all species measured with the 2014 set-up (Fig. 1 and Table 1). From the measurements of the different species at different focus settings, we conclude that the major components are N$_2$ and CO measured ca. 9 and 20 s after the H$_2$ peak maximum, respectively (Fig. 2). Furthermore, Ar, O$_2$, unpyrolysed CH$_4$ and Kr elute from the 2nd GC and are baseline-separated from our target H$_2$ (Fig. 2 and Table 2). H$_2$O, N$_2$O and CO$_2$ cannot pass the cold trap between the pyrolysis oven and T7 but are visible when the mass spectrometer is focussed to their typical m/z settings. Therefore, these gases must be produced from eluting peaks plus background gases present in the ion source. It is evident that H$_2$ peaks produce a signal on m/z 18 (and 17), with the latter peaking shortly (ca. 0.5 s) after the H$_2$ peak maximum. We propose that water is produced in situ within the ion source from oxygen-containing species already present in the ion source (CO, CO$_2$, O$_2$, H$_2$O). Tests indicate that the changes in the history of H$_2$O-generating background levels in the mass spectrometer can significantly alter the isotopic signature of pure (rectangular) H$_2$ peaks. Specifically, we find decreased $\delta^4$D(H$_2$) values for higher water levels (generated in the ion source). Performing the same measurements at lower electron voltages (from 90 to 70 eV) reduces the effect and is generally recommended for $\delta^4$D(H$_2$) analysis due to the formation of $^4$He$^{2+}$ at voltages $> 79$ eV (Denifl et al., 2002) (and comments by W. Brand in the Isogeochem archive, 2002 and 2005, http://list.uvm.edu/cgi-bin/wa?A0=ISOGEOCHEM).

In any case, it is possible to create and maintain stable conditions for our complete system enabling robust $\delta^4$D(CH$_4$) measurements, that is, strictly following the “identical treatment” (IT) principle of samples and references (Werner and Brand, 2001). To provide each CH$_4$-derived H$_2$ peak with identical background conditions, we stick to the previously described regular injections of either pure CH$_4$ in He or sample/reference air-derived CH$_4$ every 20 min (Bock et al., 2010a) and leave the open split inserted over the course of the day. Obviously, also pure H$_2$ rectangular monitoring peaks are injected on a regular schedule.

Table 3. Results obtained with the new $\delta^4$D(CH$_4$) system (green boxes in Fig. 1) in comparison to our previous set-ups (Table 1). Mean values are given in column 3; columns 4 and 5 show standard deviations (1σ) of samples and Air Controlé reference air measurements, respectively. Air Controlé measurements are used to calibrate the samples to the international VSMOW scale. Air Controlé has been cross-referenced with respect to Alert (see text). The value given for Alert in the column for 2010 is given by C. Veidt (personal communication, 2014). “N” represents the number of measurements used: subscripts “SA” and “REF” in columns 2 and 6 denote sample and reference (i.e. Air Controlé), respectively. Columns 7–11 are arranged in the same pattern for the data with only basic GC separation after pre/postPT (Bock et al., 2013) (red and blue boxes in Fig. 1). Columns 12–14 show values obtained with the previous set-up presented in Bock et al. (2010a). Ice sample results are not corrected for any firm diffusion process. Gas ages of the ice samples are estimated as follows: B30 – 670 a BP, B34 – 1400 to 1530 a BP, and WAIS – 410 a BP. The WAIS samples are from core WDC05A, tube 184, depth range: 172.74–173.03 m. NGRIP gas samples date from between 870 and 9000 a BP.

<table>
<thead>
<tr>
<th>Sample description (sample size, origin CH$_4$ concentration)</th>
<th>This study (green)</th>
<th>Bock et al. (2013) (blue)</th>
<th>Bock et al. (2010a)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>N$_{SA}$</td>
<td>$\delta^4$D(CH$_4$)</td>
<td>1σ sample (%)</td>
</tr>
<tr>
<td>Air reference and samples</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Alert (<strong>2002/11</strong>, [CH$_4$]= 1833 ppb)</td>
<td>6</td>
<td>–81.4</td>
<td>0.5</td>
</tr>
<tr>
<td>Air Controlé (all injections, 4–40 mL)</td>
<td>47</td>
<td>–93.6</td>
<td>1.1</td>
</tr>
<tr>
<td>Air Controlé (only larger loops (18–40 mL)</td>
<td>21</td>
<td>–93.6</td>
<td>0.6</td>
</tr>
<tr>
<td>Saphir 4 ([CH$_4$]= 761 ppb)</td>
<td>5</td>
<td>–169.6</td>
<td>1.5</td>
</tr>
<tr>
<td>Saphir 4 (loop after sample)</td>
<td>3</td>
<td>–171.6</td>
<td>1.1</td>
</tr>
<tr>
<td>Saphir 3 ([CH$_4$]= 1004 ppb)</td>
<td>2</td>
<td>–173.4</td>
<td>0.4</td>
</tr>
<tr>
<td>Boulder (CA08289 [CH$_4$]= 1500 ppb)</td>
<td>4</td>
<td>–81.2</td>
<td>0.5</td>
</tr>
<tr>
<td>NAT-332 ([CH$_4$]= 2141 ppb)</td>
<td>2</td>
<td>–107.7</td>
<td>1.1</td>
</tr>
<tr>
<td>Dome 6 (thin air [CH$_4$]=1718 ppb)</td>
<td>2</td>
<td>–71.8</td>
<td>0.1</td>
</tr>
<tr>
<td>Ice core replicates</td>
<td></td>
<td>N$_{SA}$</td>
<td>depth intervals</td>
</tr>
<tr>
<td>B30 (Greenland, pre-industrial, depth range 2 m)</td>
<td>2</td>
<td>–91.5</td>
<td>0.8</td>
</tr>
<tr>
<td>WAIS (Antarctica, pre-industrial, parallel replicates)</td>
<td>4</td>
<td>–73.0</td>
<td>0.5</td>
</tr>
<tr>
<td>B34 (Antarctica, late Holocene, depth range 9 m)</td>
<td>4</td>
<td>–71.4</td>
<td>1.5</td>
</tr>
<tr>
<td>Ice core replicates</td>
<td></td>
<td>N$_{SA}$</td>
<td>depth intervals</td>
</tr>
<tr>
<td>B34 ice (parallel replicates, late Holocene)</td>
<td>37</td>
<td>17</td>
<td>2.2</td>
</tr>
<tr>
<td>NGRIP (bag replicates of gas cut, Holocene)</td>
<td>27</td>
<td>13</td>
<td>2.3</td>
</tr>
</tbody>
</table>
In the following we will discuss the accuracy of our new system and its consistency with previous versions of our δD(CH₄) analysis (Table 1). As presented in Table 2, the first post-peak (mostly N₂) is due to trapped background nitrogen in the He carrier gas stream. Assuming constant background (leak rate) conditions over the course of a day, any effect will cancel out by adhering to the IT principle of samples and references (Werner and Brand, 2001). The second post-peak (CO) is produced as a side product of CH₄ pyrolysis. In our previous report (Bock et al., 2010a), we showed that within the precision of that time, neither changes in CH₄ concentration nor sample volume influence δD(CH₄) for our extraction/conversion line. Hence, the IT principle is also valid here, as we match the peak sizes (i.e. methane amount) for samples and references. On the contrary, differences may occur in systems with or without post-conversion GC separation caused by unpyrolysed CH₄ and Kr. Results for Saphir 3 (Table 3), an artificial air sample containing no krypton, which was re-measured with a −5.8‰ offset and with a combined error of 3.5‰ (the square root of the sum of the squared standard deviations of samples and reference measurements) using the system with only basic 2nd GC separation (blue in Fig. 1 and Table 1), may indicate such an effect. Although this offset is still within 2σ of the error, we speculate that this is related to the Kr effect for the older measurements without post pyrolysis GC separation. In the discussion version of this contribution, we demonstrate that Kr produces a positive signal on m/z 2 and a negative signal on m/z 3 (Bock et al., 2013). The effect for Saphir is maximal as we reference this gas mixture to a standard containing recent Kr and CH₄ concentrations (Air Control®) (see also Schmitt et al., 2013). Using a 2nd GC separation, separating Kr from the CH₄-derived H₂ peak as in the latest version of our method, a Kr effect is completely avoided.

Generally, we have found our system to be remarkably stable over the past several years, despite the implemented changes (Table 1 and Fig. 1). Table 3 summarizes our results and shows no significant differences for Alert, Boulder, NAT-332, Dome6 or ice samples when comparing the different stages of our system.

Using the 2014 set-up, Alert, our primary standard and anchor with respect to VSMOW, has been re-measured with a difference of +0.8‰ compared to Bock et al. (2010a), which is not significant with respect to the measurement errors. We decided not to shift our scale based on these six measurements.

### 3.2 Precision and sample size

In this section we describe the improvements concerning precision and sample size due to pre- and post-pyrolysis trapping of methane and hydrogen, respectively. In our old system (without pre&postPT) a typical sample (up to 500 g of polar ice with CH₄ concentrations between 350 and 700 ppb) showed peak heights of the major beam between 0.6 and 1.3 nA (for ice core samples presented in Bock et al., 2010b). While peak areas are still in the same range for identical amounts of CH₄, major peak heights are increased roughly fourfold due to postPT.

In Table 4 and Fig. 4, we present 51 ice core samples (see Sect. 3.3) of a core dry drilled next to the EPICA...
Figure 4. δD(CH₄) of B34 ice core samples measured during the years 2011 to 2014 (using the different set-ups shown in Table 1 and Fig. 1) on a depth scale. Error bars represent the pooled standard deviation of B34 replicates (2.3 ‰) for δD(CH₄) and the total depth range of each individual sample. The 10 m depth interval presented here corresponds to an age range of approximately 150 a; accordingly, a depth change of 1 m represents a nominal age increase of about 15 a, which is significantly smaller than the width of the age distribution of the bubbles in the ice.

3.3 First results

B34 ice core samples described in the last section have been analysed thoroughly using the different set-ups. No B34-specific gas age scale has been established; however, due to its vicinity to EDML, we make use of the Antarctic Ice Core Chronology 2012 (AICC2012) (Veres et al., 2013) to derive gas age estimates: on the EDML scale the depth range 181–191 mbs corresponds to an age of the occluded air of 1401–1532 a BP (Table 4). Note that in ice cores, the extracted air is integrated over a large number of individual air bubbles in the ice, which have somewhat different gas ages due to the bubble close-off process in the firn column. Accordingly, ice cores provide only a low-pass filtered signal of the atmospheric concentration. Using the firn model described by Spahni et al. (2003), we calculated the age distribution for EDML in the Holocene, representing the distribution of the gas age in each bubble: the peak in the age distribution is at 33 a and the width at half-maximum is 67 a. Figure 4 shows δD(CH₄) measured on B34 ice samples on a depth scale. The same data are also presented in Table 4, which additionally shows gas age, the measurement date and the weight of samples. Overall, we are confident that the described system was stable in terms of accuracy over the past few years, and the pooled standard deviation of all replicates (N = 41, identical depths = 17) is 2.3 ‰.
Along with our measurements on standard air tanks, the ice sample data for all set-ups agree within the overall scatter of all data. Despite the fact that the samples have been measured with three different measurement systems and over a time span of several years, all samples agree within $2\sigma$ for any given depth interval and most samples agree within $1\sigma$, showing that they are statistically identical. Despite this statistical agreement there seems to be the tendency of samples taken at exactly the same depth to agree better than samples that have been cut from adjacent ice, which, due to the slow bubble enclosure process, should be regarded as replicates as well. Moreover, there seems to be some variability in the average $\delta D(\text{CH}_4)$ value on the metre depth scale, which is unexpected given the wide age distribution of the air bubbles, which quite effectively smoothes out multi-annual atmospheric variability. Whether this is due to an incomplete understanding of the bubble close-off process in this core, reflects limited potential in situ CH$_4$ formation (Rhodes et al., 2013) or remains just statistical coincidence requires further dedicated studies in the future using an identical measurement system.

Mean values for WAIS (Antarctica) and B30 (Greenland) from similar (pre-industrial) time periods (around 410 and 670 a BP, respectively) are $-73.0$ and $-91.5$ ‰. This difference of $18.5$ ‰ with a combined error of $1.9$ ‰ (determined as above) can be largely explained by the expected inter-polar difference in $\delta D(\text{CH}_4)$, which can be explained by the geographical distribution of CH$_4$ sources with different $\delta D(\text{CH}_4)$ signature in combination with the inter-hemispheric air mass exchange and the lifetime of CH$_4$ of the order of 8–10 years.

It is beyond the scope of this article to discuss the biogeochemical implications of this finding, but we note that the inter-polar difference derived from our measurements is in line with earlier work (Sowers, 2010; Quay et al., 1999).

4 Conclusions

We presented pre- and post-pyrolysis trapping of methane and hydrogen, respectively, combined with post-trapping GC separation on a cooled PLOT column to improve accuracy and precision and reduce sample amount in $\delta D(\text{CH}_4)$ analysis of atmospheric and ice core samples. We showed that the precision for 350 g of ice (or roughly 30 mL of air) with 350 ppb of methane is approximately 2.3 ‰. This corresponds to recent tropospheric air samples (roughly 1900 ppb CH$_4$) of about 6 mL (STP) or about 500 pmol of pure CH$_4$. In contrast, 30 mL (STP) samples with recent tropospheric CH$_4$ concentration can be determined with a precision of better than 1 ‰. Compared to our old set-up (Bock et al., 2010a), this translates into improvement factors for sample size (350 g)/(500 g) and precision (2.3 ‰)/(3.4 ‰) of 0.7.

We note, however, that the high standard in accuracy and precision for such small samples is achieved at the cost of sample throughput; typically we can measure eight standards in addition to at most two ice core samples or four atmospheric samples a day.

We showed that the accuracy of systems without pre&postPT and subsequent chromatographic separation can be potentially biased depending on pyrolysis efficiency and varying methane/krypton ratios in samples and the reference. However, for atmospheric samples (ice and tropospheric air samples), the updated method did not measurably change in terms of accuracy of $\delta D(\text{CH}_4)$ values compared to our initial set-up described in Bock et al. (2010a).

We propose that water produced in situ in the ion source of the mass spectrometer from hydrogen plus oxygen-containing background species may be an important factor affecting precision and accuracy of $\delta D$ measurements. In our case we take advantage of simple chromatograms and regular injections to ensure identical background levels for each sample or reference peak.
Appendix A: Correction for system drifts and signal dependency (linearity)

In order to calibrate samples measured on any isotope system, it is essential to compare samples to standard measurements that are sufficiently stable in time and match the sample size, or to correct for any drift and signal dependency. As amount errors alter isotopic results simultaneously with (time) drift effects, both errors should be corrected at the same time and not consecutively. A decoupling of the corrections is only possible when standards of constant peak size are measured to monitor the time trends only. Effects of signal dependency can be assessed by performing standard runs of different peak sizes; however, this is quite time-consuming. Hence, we present an approach which allows for simultaneous corrections of system drifts and signal dependency effects. For optimum conditions we choose size matching and bracketing standards for individual samples and pool standards measured over several days (assuming constant signal dependency over this time period) to cover the samples’ size range. To correct for both signal dependency and drift effects, we use the following approach.

Any measured isotope value $\delta X_{\text{meas}}$ is composed of the true value $\delta X_{\text{true}}$, any signal dependency, which is a function of peak area, $A$, and a drift correction, which is a function of time, $t$:

$$\delta X_{\text{true}} = \delta X_{\text{meas}} - f_{\text{lin}}(A) - f_{\text{drift}}(t).$$  \hspace{1cm} (A1)

In the following, signal dependency is characterized by a polynomial of order $N$:

$$f_{\text{lin}}(A) = \sum_{n=1}^{N} x_n A^n.$$  \hspace{1cm} (A2)

System drift is decomposed into two additive terms:

$$f_{\text{drift}}(t) = f_{\text{drift}}^1(t) + f_{\text{drift}}^2(t).$$  \hspace{1cm} (A3)

The first term is a drift over the course of a day, which is fitted to a polynomial of order $M$:

$$f_{\text{drift}}^1(t) = \sum_{m=1}^{M} y_m t^m \Theta(t - [\bar{t}_i - \Delta t]) \Theta([\bar{t}_i + \Delta t] - t),$$  \hspace{1cm} (A4)

which is a function of time $t$. Here, $\bar{t}_i$ represents the time during day $i$ at which the current sample was measured. Thus, $\bar{t}_i$ describes the mean measurement time of all samples measured during one day. Since temporal system drifts typically occur on timescales of weeks to months, the size of the drift within a day is usually small. Accordingly, our software allows for the calculation of $\delta D(\text{CH}_4)$ values with or without a diurnal drift correction (the latter is usually our preferred setting). The Theta function, $\Theta$, is zero if its argument is $< 0$ and one if its argument is $> 0$. This efficiently allows for the determination of the drift for each single measurement day in the program code. To discriminate between two consecutive laboratory days, $\Delta t$ is defined as 0.4 days. The number of standard data points for each day should be greater than or equal to $M$.

The second term represents the drift of the reference values between days. The mean isotopic reference signatures of all days are assumed to change in a stepwise linear fashion:

$$f_{\text{drift}}^2(t) = \sum_{i=1}^{L} (m_i t + n_i) \Theta(\bar{t}_i - t) \Theta(t - \bar{t}_{i-1}),$$  \hspace{1cm} (A5)

where $m_i$ quantifies the slope and $n_i$ the intersection with the ordinate on measuring day $i$, and $L$ is the number of all measurement days.

Slope and intersect for each day $i$ are calculated with respect to the previous day $i - 1$.

$$m_i = \frac{\bar{t}_i - \bar{t}_{i-1}}{\delta X_{\text{meas}}(i) - f_{\text{lin}}(A_i) - \left[ \delta X_{\text{meas}}(i-1) - f_{\text{lin}}(A_{i-1}) \right]}$$  \hspace{1cm} (A6)

$$n_i = \left[ \delta X_{\text{meas}}(i) - f_{\text{lin}}(A_i) \right] - m_i \bar{t}_i.$$  \hspace{1cm} (A7)

Influences of signal dependency have to be corrected for before calculating the mean standard isotopic signal of each day.

We can express all quantities given in Eq. (A1) as functions of peak area, $A$, and isotopic signature $\delta X_{\text{meas}}$ at each point measured at time, $t$. The true value of the standard $\delta X_{\text{true}}$ is known. Thus, Eq. (A1) can be used to fit all measured data points. The fit parameters determine both the signal dependency and the drifts during and between days and are found by minimizing the standard deviation of all drift and signal dependency corrected standard values. The fitted parameters are then used to ultimately calibrate the samples.

Our routine is written in Python (www.python.org). The actual optimization uses the function scipy.optimize.fmin(). Figure 3a and b are produced by our routine and show uncalibrated and calibrated data, respectively. Each subfigure shows signal dependency in the left panel and time drift in the right panel (using the same data in the two panels).
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