



atmospheric CH_4 (top) and the trend in the ratio of the isotope carbon-13 (bottom) from samples collected by NOAA's Global Greenhouse Gas Reference Network. (Credit: Xin Lan and Sylvia Michel)

Methods

2015

• Measurements of the relative abundance of ¹³C to ¹²C (denoted as $\delta^{13}C-CH_{A}$) is a useful tracer for disentangling the contribution of the sources and sinks of CH₄ (Fig. 2, Lan et al., 2021)



Fig. 2. (left) Bar chart of global mean CH₄ emissions from microbial, fossil, and pyrogenic source sectors. (right) Global mean flux-weighted CH₄ isotopic signature (δ^{13} C-CH₄) of microbial, fossil, and pyrogenic sources.

• We updated our CarbonTracker-CH₄ inversion system by jointly assimilating measurements of CH, and $\delta^{13}C-CH_{a}$, optimizing fluxes at a grid scale, and incorporating δ^{13} C-CH₄ signatures of sources. The system has been extended to estimate fluxes through 2021 (Bruhwiler et al., 2014; Basu et al., 2022)

Results 2. Total emissions reach to 650 Tg CH, yr⁻¹ in 2021, ~18% increase 1. Joint inversion that assimilates both $CH_{A} + \delta^{13}C-CH_{A}$ since 2000. The averaged contribution of fossil sources to total measurements matches with both observations, where as emissions is larger (~15 Tg CH₄ yr⁻¹) when assimilating δ^{13} C-CH₄ measurements (joint scenario). joint inversion yields more reasonable partitioning of sources. Fig. 3. Model-data South Pole Barrow, Alaska Fig. 4. Optimized annual global Global Optimized CH, emissions comparison of (top) CH₄ emissions simulated by joint CH_₄ and (bottom) (blue) and CH₄-only (red) $\delta^{13}\dot{C}$ -CH, between scenarios of CarbonTracker-CH observation (black) and for total (o), microbial (x), and simulations from joint fossil (Δ) sources in 2000-2021 (blue) and CH₄-only (red) scenarios for (left) Barrow, Alaska, and (right) South Pole baseline observatories. ---- \rightarrow CH₄ + δ^{13} C-CH₄: Microbial \rightarrow CH₄ + δ^{13} C-CH₄: Foss CH₄+δ¹³CH. CH₄ only Observed 4. Spatial changes in source-specific CH₄ emissions in 2020-2021. • In 2020, the inversion attributes more than half of the microbial increases to tropical regions and the decrease in fossil emissions to



CH₄-only inversion does not match with $\delta^{13}C$ -CH₄. Therefore, the 3. Microbial emissions caused increase in atmospheric methane

- The joint $CH_{A} + \delta^{13}CH_{4}$ inversion simulates the large increase in microbial sources and slight decrease in fossil sources (Red in Fig. 5).
- [OH] decrease can contribute to 30-50% of atmospheric CH increase (Yellow in Fig. 5).



scenarios.

What drove record increases in global atmospheric methane in 2020 and 2021? Youmi Oh^{1,2}, Lori Bruhwiler², Xin Lan^{1,2}, Sourish Basu³, Sylvia Michel⁴, John Miller², Ed Dlugokencky², Lei Hu², and Arlyn Andrews²

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in 2020-2021.

Fig. 5. Changes in global CH_4 emissions (Tg CH4 yr⁻¹) between 2020-2021 and 2018-2019 for total (left), microbial (middle), and fossil (right) emissions. Different color bars represent CH_{a} -only (blue), joint (CH_{a} and $\delta^{13}C$ - CH_{a}) (red), and joint + OH decrease (yellow) inversion

industrialized regions (Fig. 6). In 2021, microbial sources increase in both temperate and tropical regions, while fossil emissions rebounce to 2019 levels.







Fig. 6. The difference of global gridded optimized emissions between 2019 and 2020 for (top-right) total, (top-left) microbial

(bottom-left) fossil. and (bottom-right) pyrogenic emissions simulated by the joint scenario of CarbonTracker-CH

Conclusions

- CarbonTracker-CH_{$_{A}$} is developed to jointly assimilate CH₄ and δ^{13} C-CH₄ measurements and estimate source-specific emissions.
- Source-specific emissions from CH_{4} and $\delta^{13}C-CH_{4}$ should be more realistic because CH₄-only inversion yield δ^{13} C-CH₄ field is inconsistent with observations.
- The total global CH, emission increased more than 100 Tg yr⁻¹ after 2000 and reached \sim 650 Tg yr⁻¹ in 2021.
- A joint $CH_4 + \delta^{13}C CH_4$ inversion suggests that the post-2006 and 2020-2021 atmospheric CH increases are due to microbial sources, although we cannot rule out a role of inter-annual variability in [OH].

References

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