

# Type II Aerobic Methane Oxidizing Bacteria (AMOB) Drive Methane Oxidation in Pulsed Wetlands as Indicated by <sup>13</sup>C-Phospholipid Fatty Acid Composition



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## INTRODUCTION

> The magnitude of climate change predicted for the United States over the next 100 years will cause significant impacts on temperature and precipitation patterns

> Wetlands are a major source (~25 %) of methane globally<sup>1</sup>

> Aerobic Methane Oxidizing Bacteria (AMOB) have the unique ability to utilize CH<sub>4</sub> as the sole source of carbon and energy by the Methane Mono Oxygenase (MMO) mediated reaction

> A variety of environmental determinants have been implicated in the dominance of AMOB type. Generally, type I AMOB outcompete type II AMOB in low CH<sub>4</sub> (~2ppmv), high oxygen environments, while type II AMOB are favored in high CH<sub>4</sub> (≥40ppmv) environments<sup>2</sup>

> AMOB are active at the oxic sediment - water interface ("pulsing fringe", Fig. 2)

> AMOB consume ~30 Tg-CH<sub>4</sub>-yr<sup>-1</sup>, and potentially can offset CH<sub>4</sub> losses to the atmosphere<sup>1</sup>

> Very little is known about the effects of pulsing hydrology and season on the AMOB ecology

> Therefore, the Objectives of this Study were:

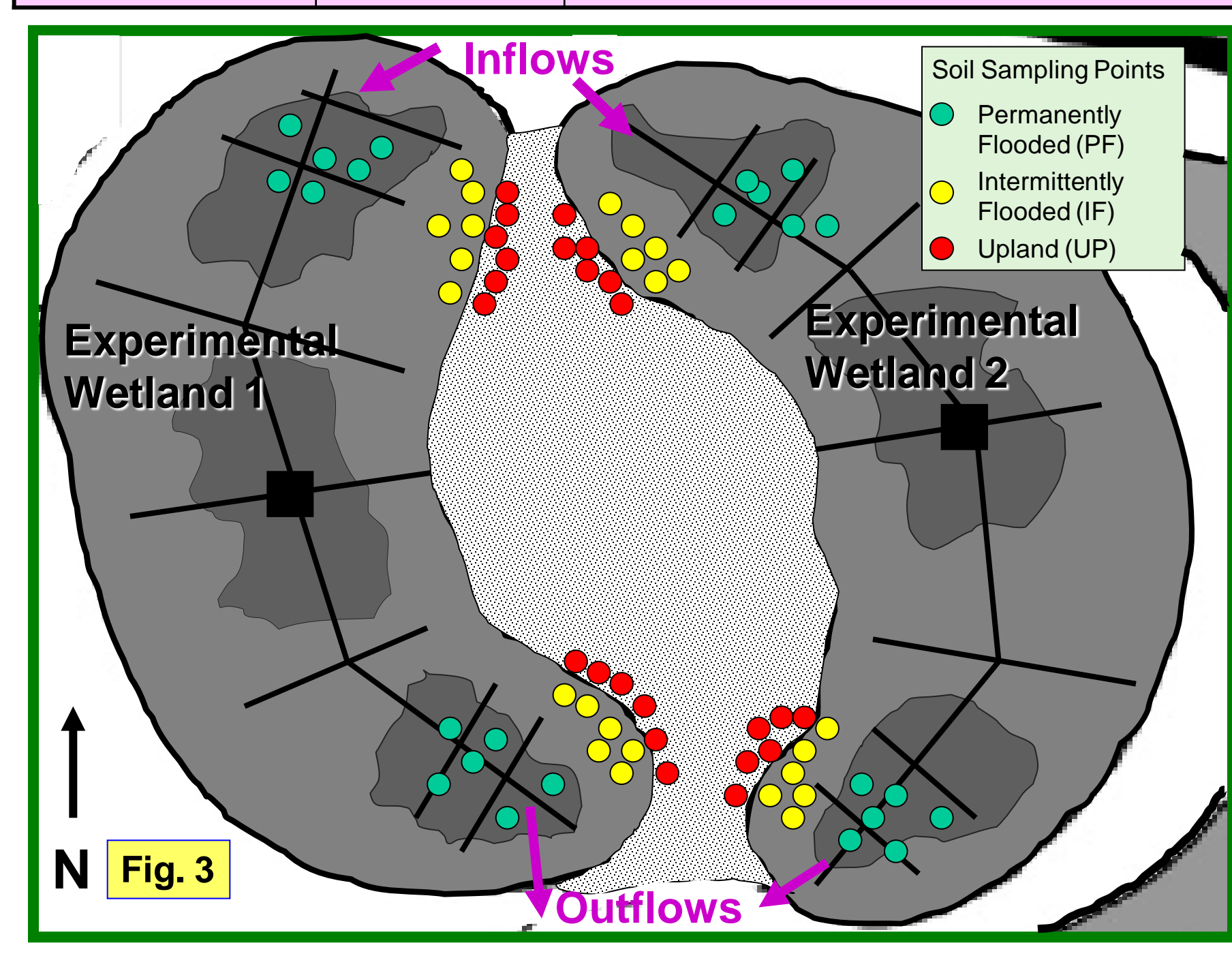
❑ Experiment 1: Determine the effects of Season and Landscape position on Potential Methane Oxidation (PMO) in the two wetlands

❑ Experiment 2: Determine the effects of Season and Landscape position on the Aerobic Methane Oxidizing Bacteria (AMOB) as reflected by their biomarker Phospholipid Fatty Acid (PLFA) Compositions

## EXPERIMENTAL SITE

THE OLENTANGY RIVER "Kidney" WETLANDS, Ohio, U.S.A.  
Soil Sampling Locations (Fig. 3) and Factors:

Factor	Number	Name
Landscape	2	ORW1, ORW2
Hydrology	3	Permanently Flooded (PF) Intermittently Flooded (IF) Upland (UP)
Flow zones	2	Inflow (I), Outflow (O)
Soil Depth	2	0-8 cm 816 cm
Season	4	Fall, Winter, Spring, Summer



## RESULTS: <sup>13</sup>C-Phospholipid Fatty Acids

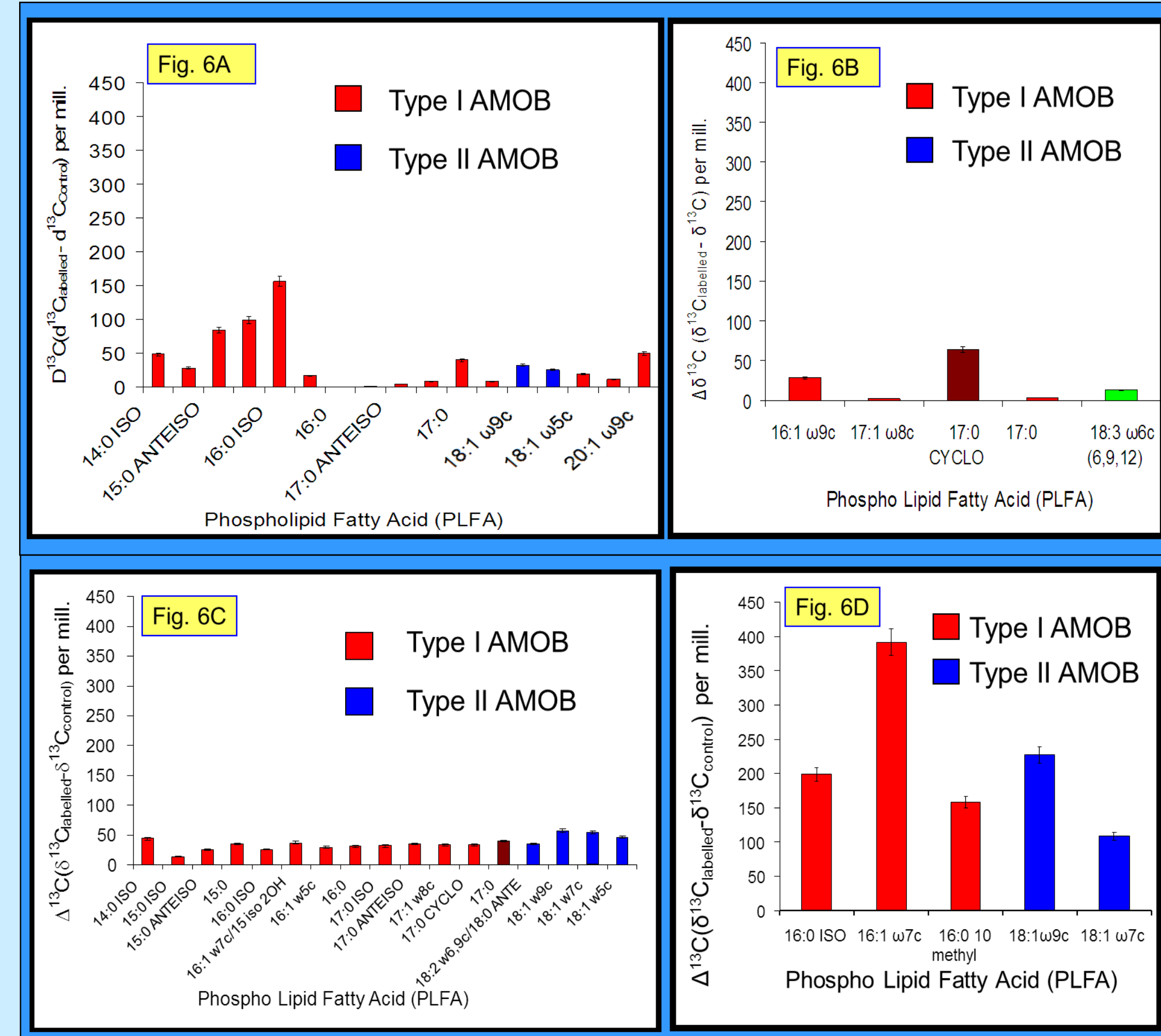
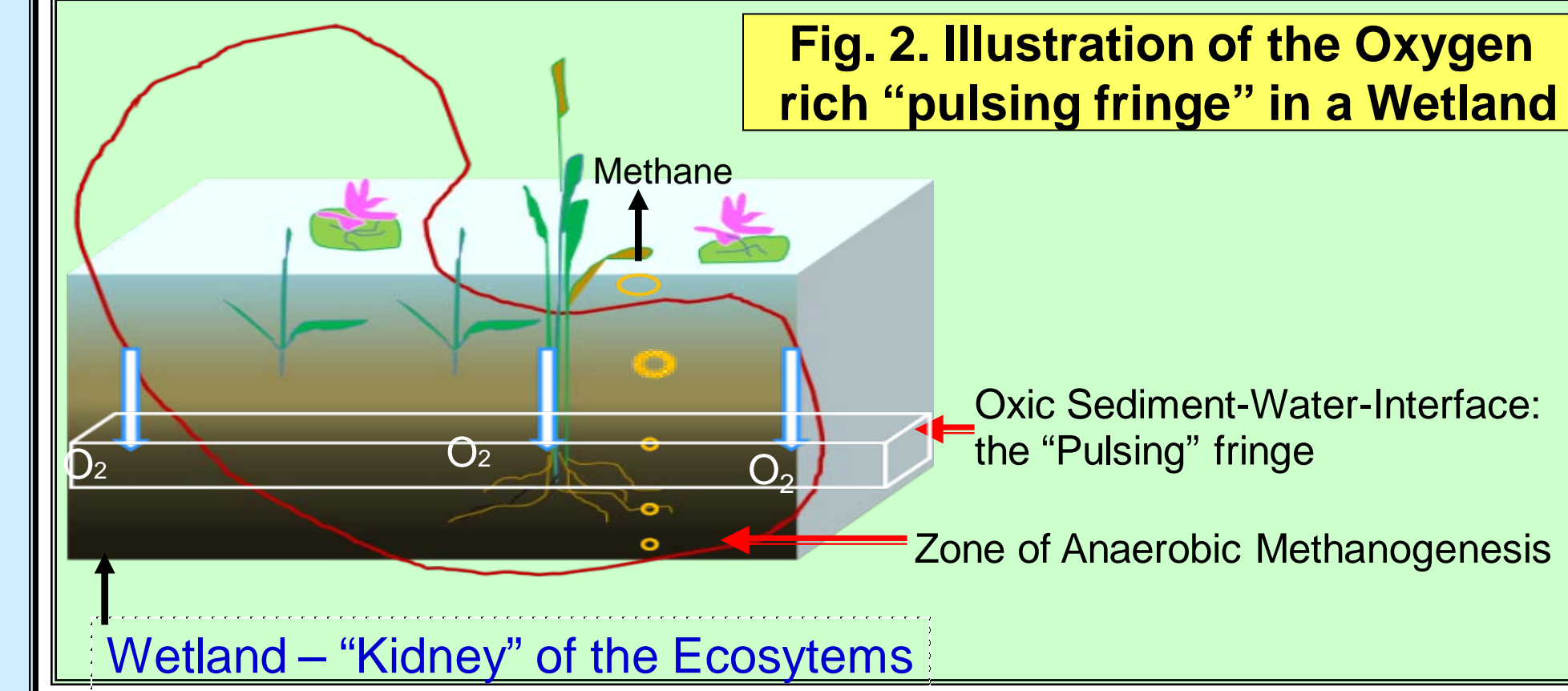
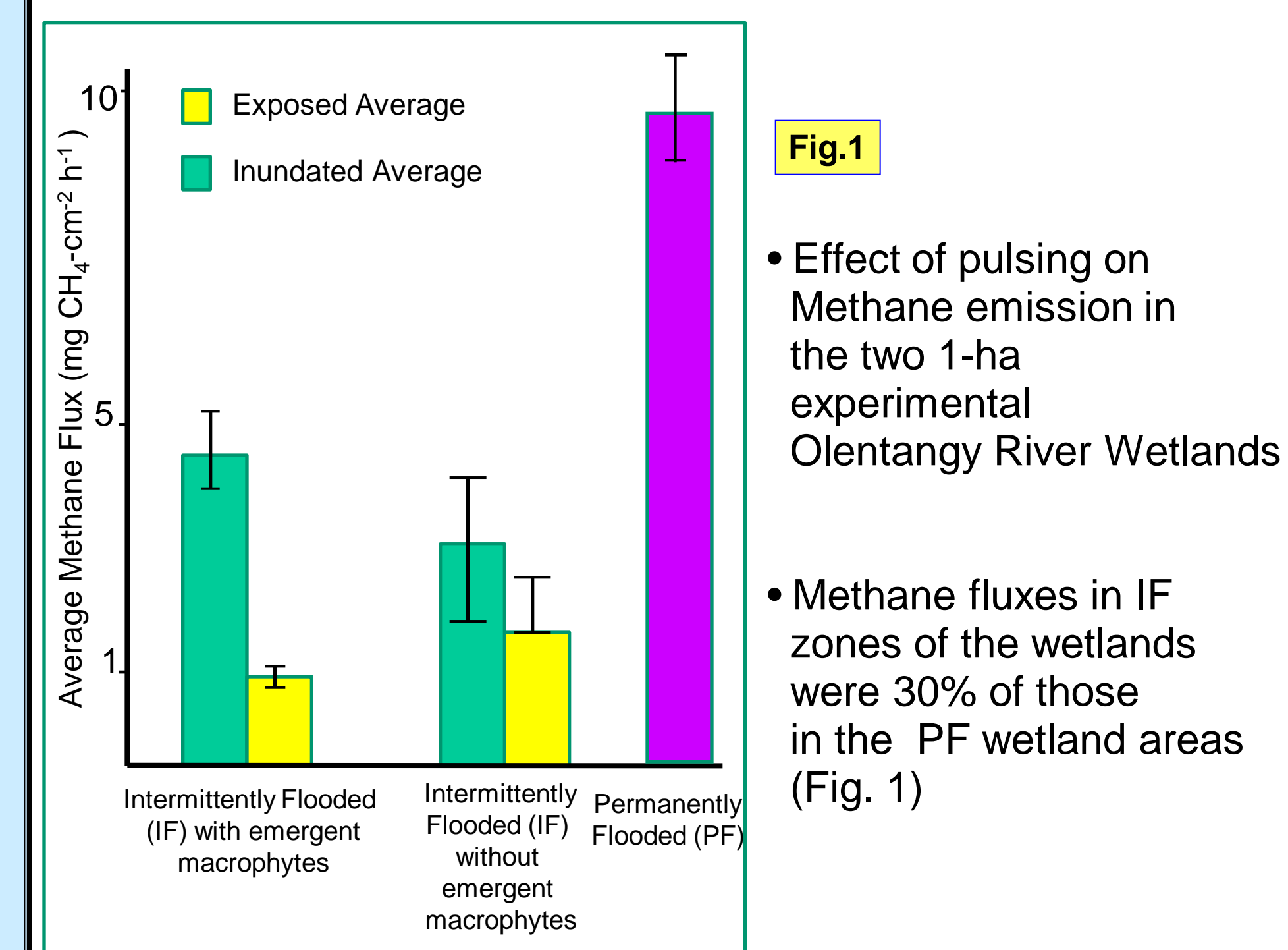


Fig. 6. Compound-specific carbon isotope values of PLFAs incorporating <sup>13</sup>C following incubation of Winter soil samples under Yppmv <sup>13</sup>CH<sub>4</sub> :

A: 0-8cm, PF Site, 2ppmv      B: 0-8cm, IF Site, 2ppmv  
C: 0-8cm, PF Site, 60ppmv      D: 0-8cm, IF Site, 60ppmv

## WETLAND HYDROLOGY EXPERIMENT

Effect of Pulsing on Methane Emission in the Olentangy River Wetlands<sup>4</sup> (Altor and Mitsch, 2006)



## REFERENCES

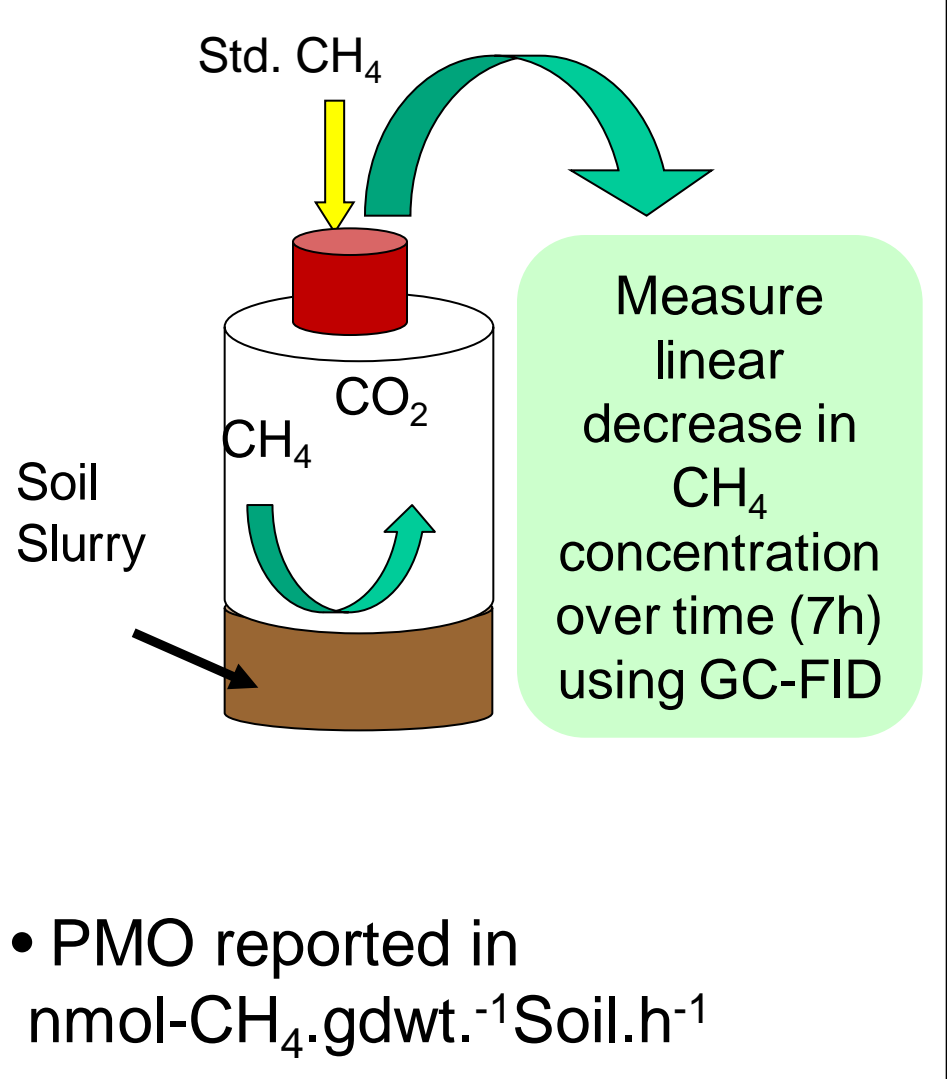
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Figs. 4 & 5: Seasonal average PMO across landscape positions:

- > PMO in Winter significantly higher than in Summer (p < 0.01)
- > PMO statistically higher at 0-8 cm depth of soil in the PF and IF sites (p < 0.05)

## METHODS

• Soil samples injected with known concentration of 99.99% pure CH<sub>4</sub> gas

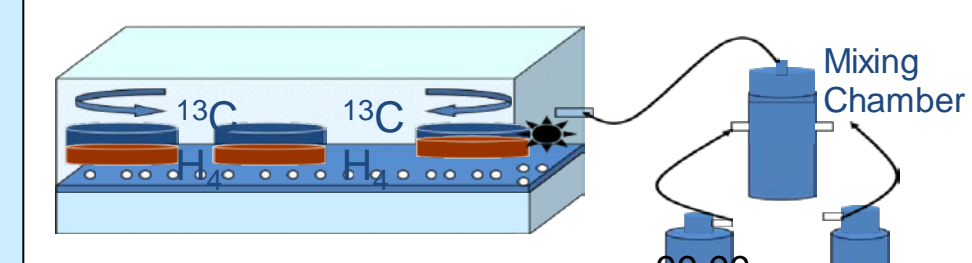


• PMO reported in nmol-CH<sub>4</sub>-gdwt.<sup>-1</sup>Soil.h<sup>-1</sup>

Method 1: Potential Methane Oxidation (PMO), Crossman et al. (2004)

Soils Incubated under two different concentrations of <sup>13</sup>CH<sub>4</sub> : 2ppmv and 60ppmv

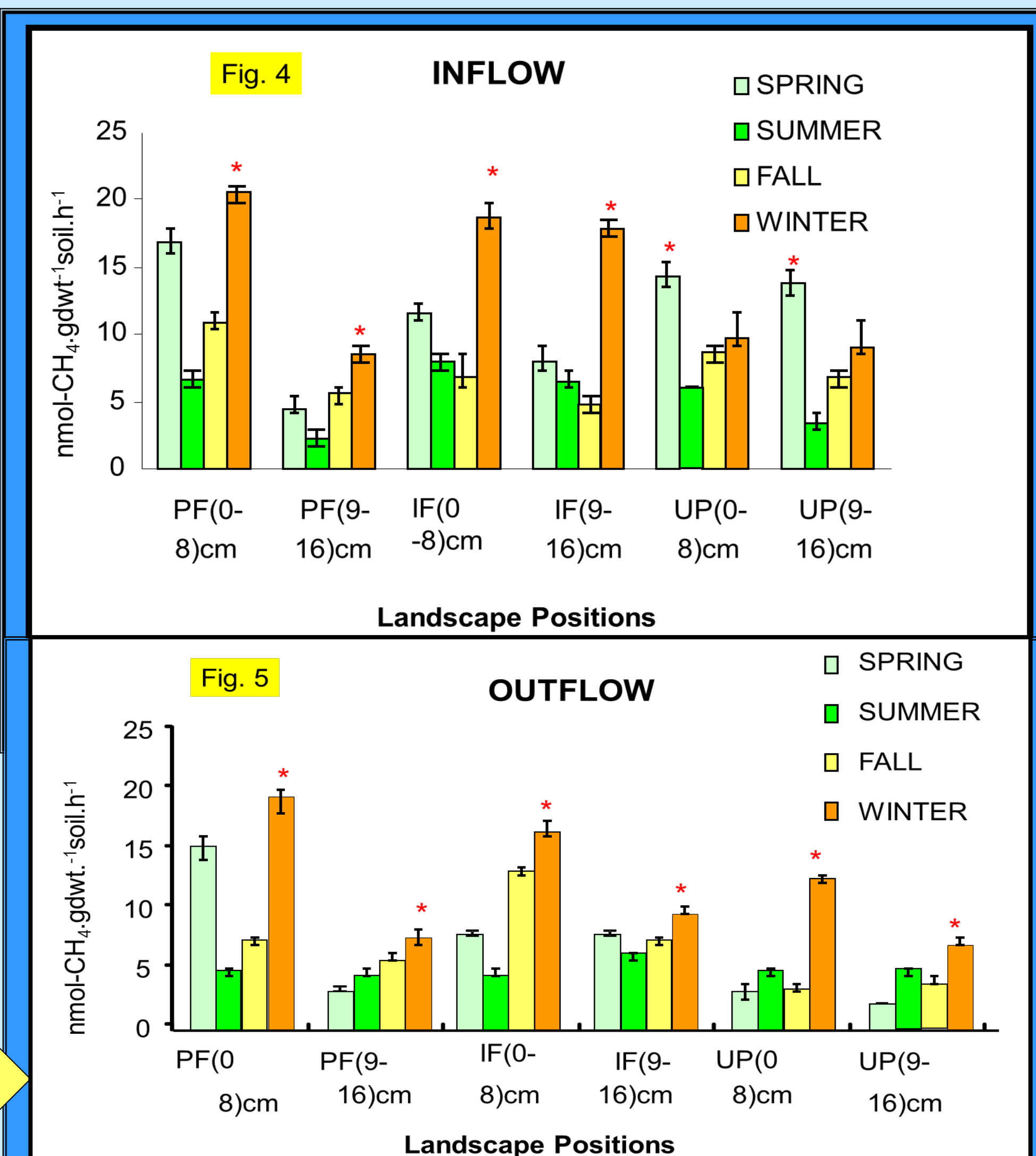
- "Type I AMOB": [CH<sub>4</sub>] < 1.8 ppm
- "Type II AMOB": [CH<sub>4</sub>] > 40 ppm



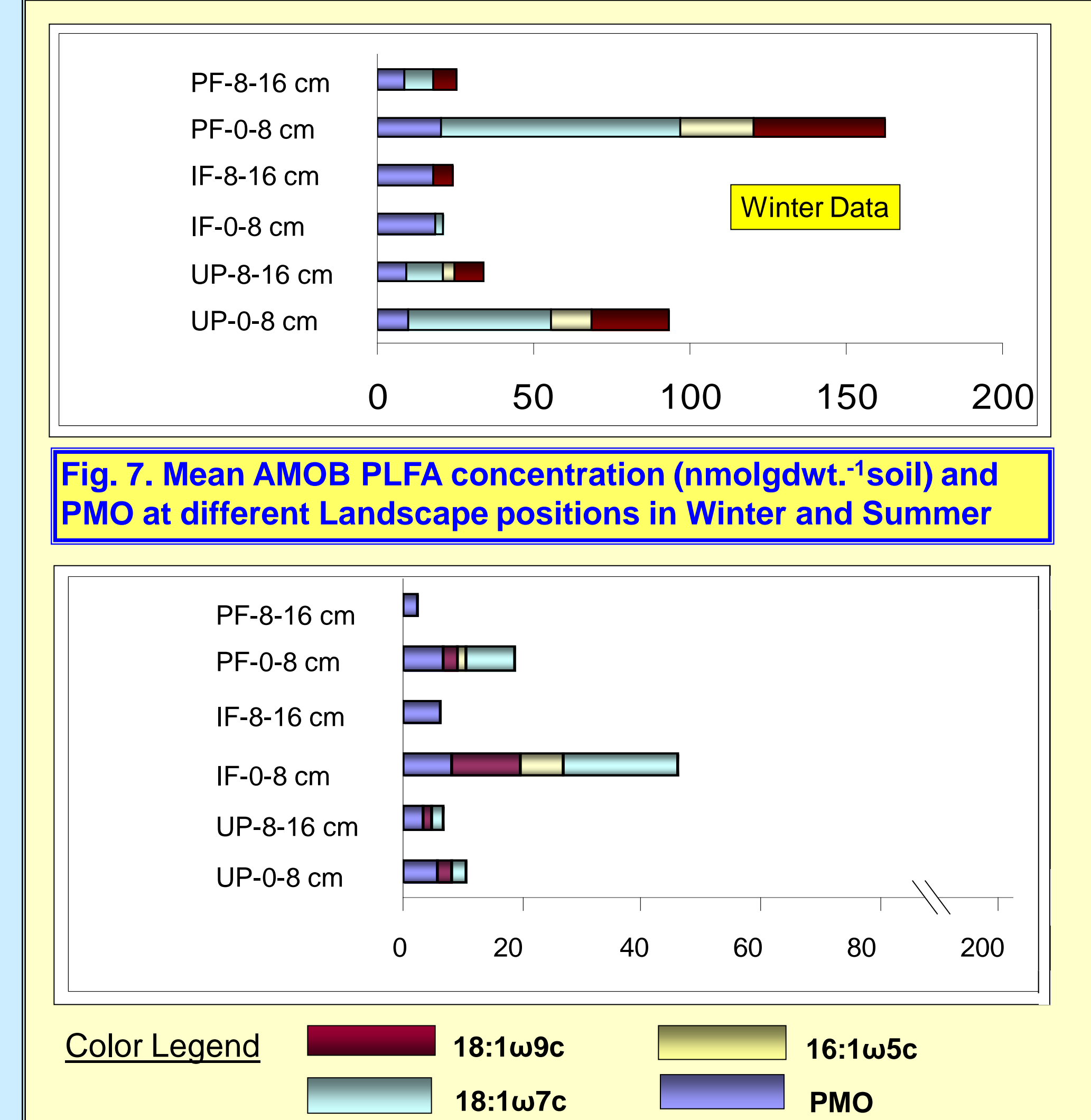
Fingerprint AMOB PLFAs:  
Type I AMOB: 16:1ω5c, 16:1ω7c  
Type II AMOB: 18:1ω7c, 18:1ω9c, 18:1ω8c

Method 2: <sup>13</sup>CH<sub>4</sub> Incubation Pulse-Chase Experimental Setup: Stable Isotope Probing of PLFA using GC-c-IRMS, Maxfield et al., 2006

## RESULTS: Potential Methane Oxidation



## SUMMARY: AMOB PLFA Conc. & PMO



## CONCLUSIONS

- > The concentration profiles of AMOB signature PLFAs, as detected by Stable Isotope Probing (SIP) completely corroborates with the Potential Methane Oxidation (PMO) values at all study sites
- > The highest PMO values in the Permanently Flooded site can be entirely attributed to the AMOB, as reflected by the relative abundance of the signature PLFAs
- > Type I AMOB are dominant under the oxygen rich conditions in contrast to the Type II that are abundant under submerged conditions
- > The three hydrologically distinct landscape positions each present characteristic microbial community structure, as evident in this study

## ACKNOWLEDGEMENT

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