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## **Microbial utilization of residue carbon depends on placement and differs in humid and semi-arid climates**

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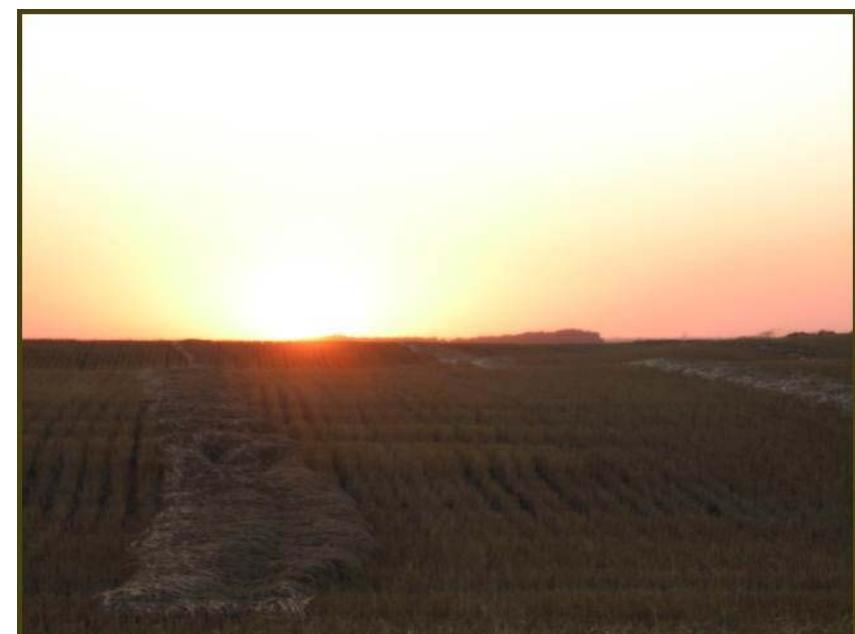
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# Introduction

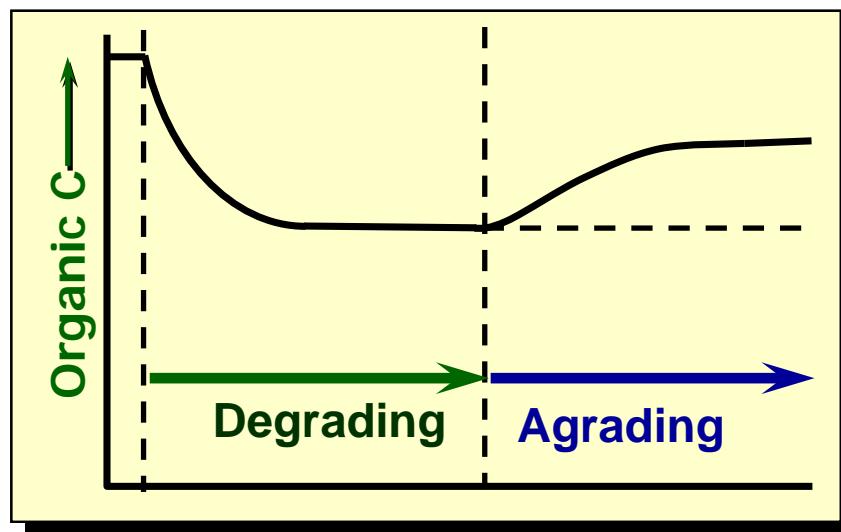
- Producing harvestable C is the main goal of agriculture
- Crop residues are the major source of new C in agricultural soils

Sustain long-term soil productivity

- rates and mechanisms of residue C decomposition
- microbial regulation



# Introduction



$C \text{ storage} = C \text{ input} - \text{decomposition}$

H. Janzen

- By manipulating residue placement we can alter the rate and perhaps pathway of C decomposition



# Introduction

Microbes are primary regulators of litter decay, but pathways of decomposition are only vaguely understood and poorly quantified



Breton: triticale



Swift Current: lentil

Use of  $^{13}\text{C}$  enriched plant material allows us to track fate of C in various physical, chemical and biological pools

“stable isotope probing”

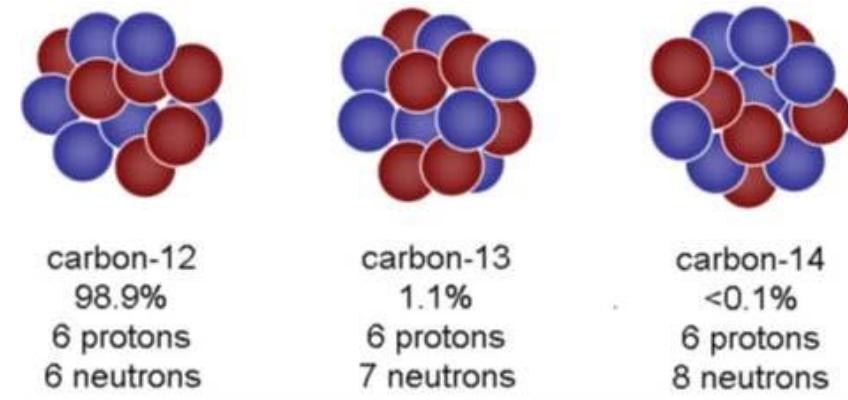


# Background: Stable Isotope Probing

- What is a stable isotope?

Common carbon isotopes:  $^{12}\text{C}$ ,  $^{13}\text{C}$ ,  $^{14}\text{C}$   
6 protons (6, 7 and 8 neutrons)

- $^{12}\text{C}$  and  $^{13}\text{C}$  are stable (non-radioactive) and occur naturally at a ratio of ca. 99:1

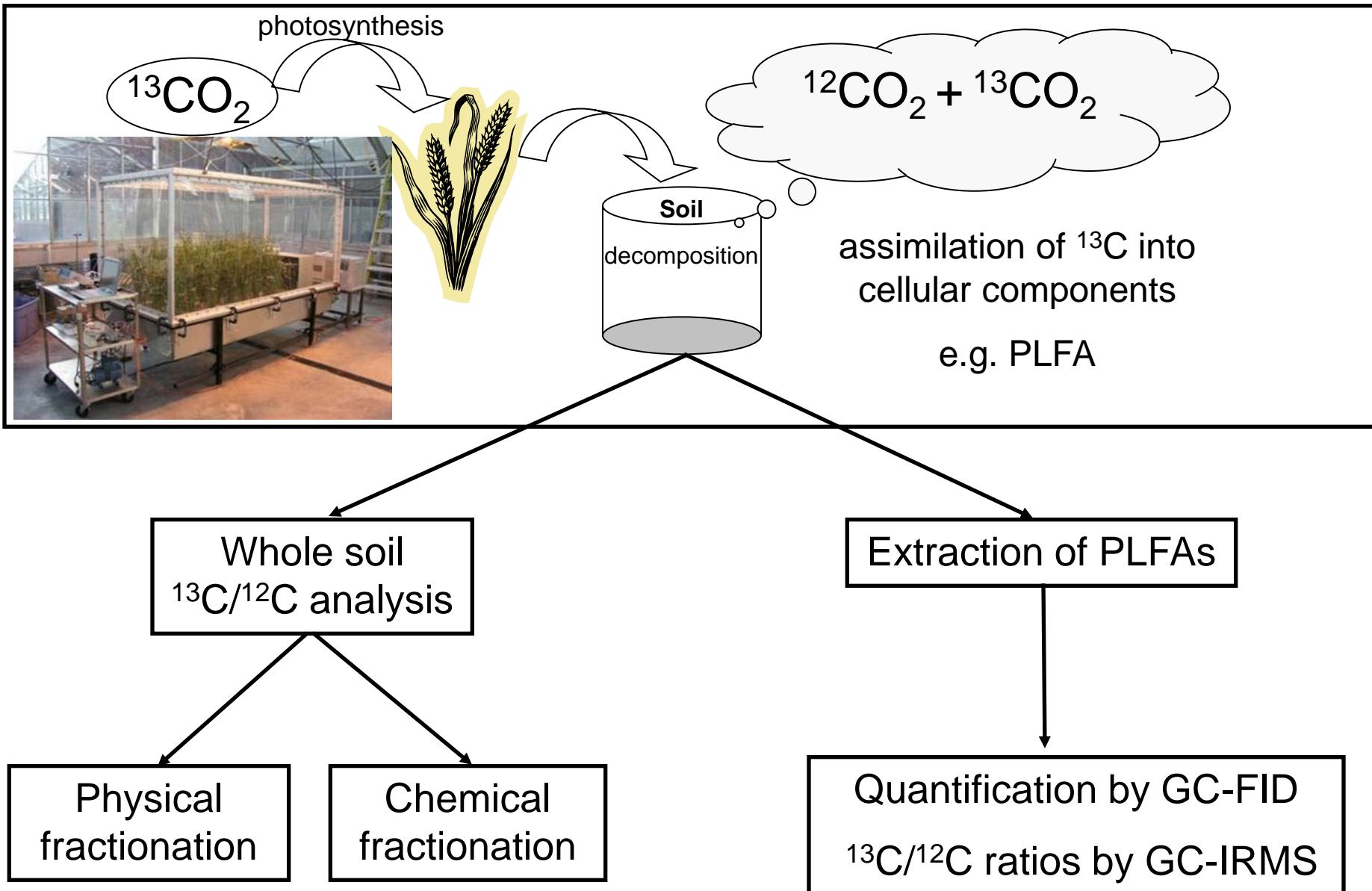


<http://wordpress.mrreid.org/2011/03/23/>

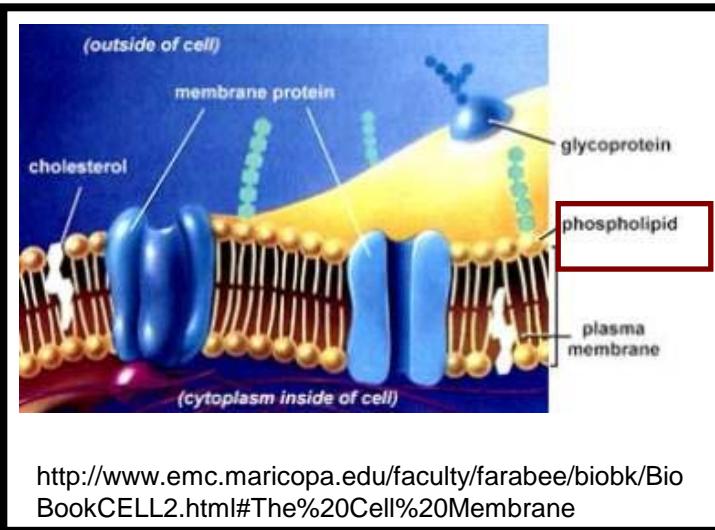
- For our purposes  $^{13}\text{C}$  acts as a *tracer*.



# Background: Stable Isotope Probing



# SIP/Phospholipid fatty acid analysis



Active populations:

1. abundance
2. microbial community structure

Microorganisms metabolize C obtained from their environment ( $^{12}\text{C} + ^{13}\text{C}$ )

Coupled with  $^{13}\text{C}$ -SIP, PLFA indicates *functionally active* components of the soil microbial community

# **Microbial utilization of residue carbon depends on placement and differs in humid and semi-arid climates**

## **Objective:**

**to determine the effect of residue placement on  
retention of  $^{13}\text{C}$  in the microbial biomass in humid  
and semi-arid climates**

# Experimental Design

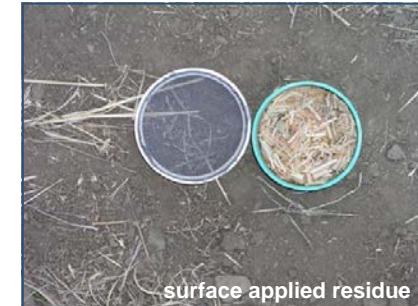
Fall 2007, microcosms set up in transects



Each transect ( $n=4$ ) has 10 microcosms

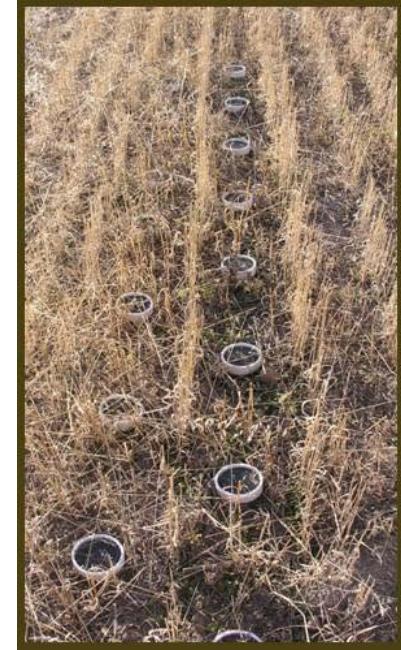
Barley residue ( $^{13}\text{C}$  at ca. 10 atom %) added at a rate of  $2 \text{ Mg C ha}^{-1}$ , either:

- a. incorporated (0-10cm)
- b. surface applied



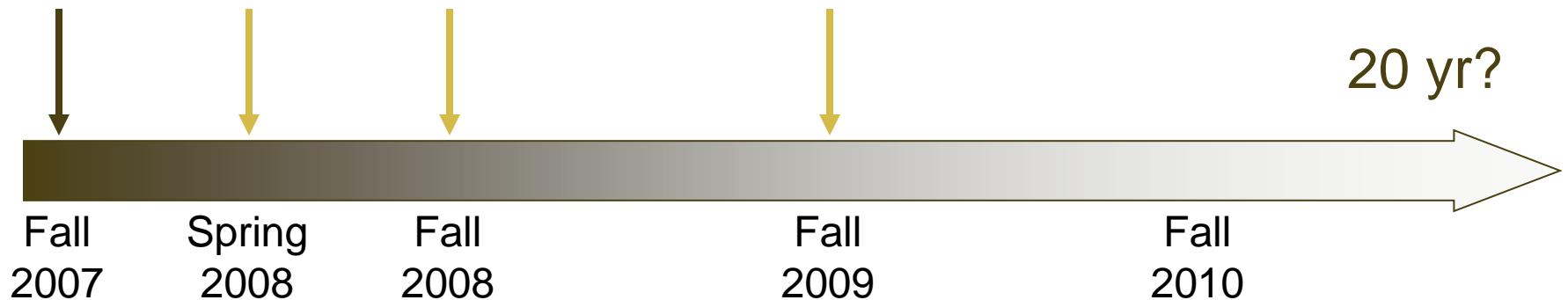
# Experimental design

\*\*Microcosms are amended annually in fall with *non-labeled* barley residue



$^{13}\text{C}$  residue added

Microcosm destructive sampling



# Microcosm sampling

- Microcosms (n=4) destructively sampled:
  - 6 mo (spring 2008)
  - 12 mo (fall 2008)
  - 24 mo (fall 2009)



- Cores divided into 2 depths:
  - 0-5cm
  - 5-10cm
- PLFA analysis:
  - GC-FID, MIDI (quantitative)
  - GC-IRMS ( $^{13}\text{C}/^{12}\text{C}$  of 26 individual PLFAs)

# Experimental design:

Basic Site characteristics:

Ottawa:      humid

                fine sandy loam

MAP: 882mm; MAT: 5.8°C

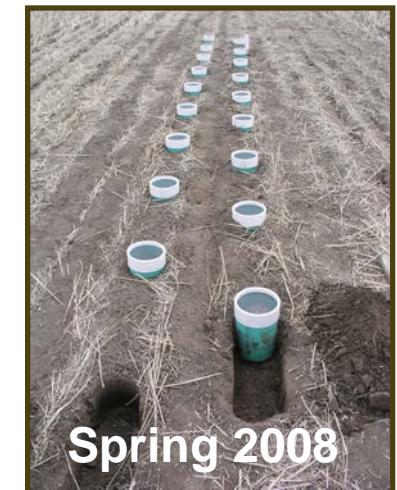


Spring 2008

Lethbridge: semi-arid

                sandy clay loam

MAP: 386mm; MAT: 5.7°C

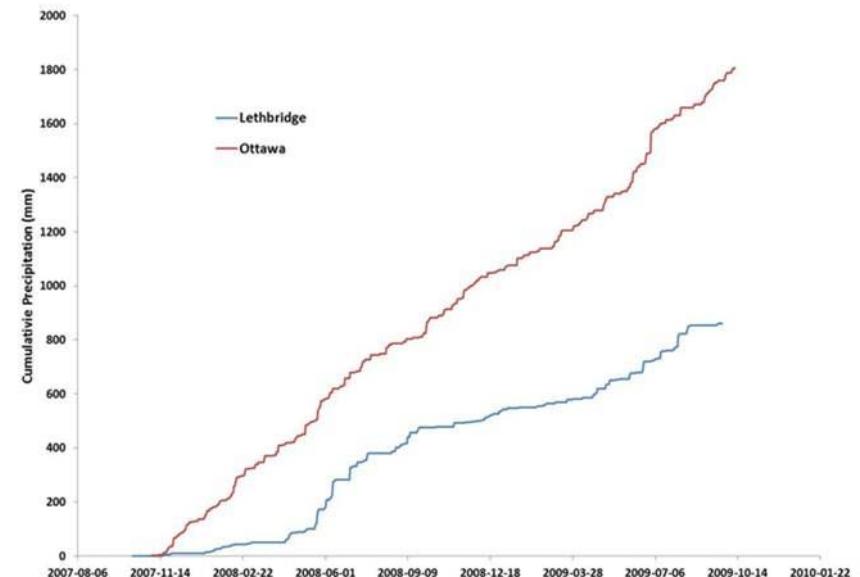
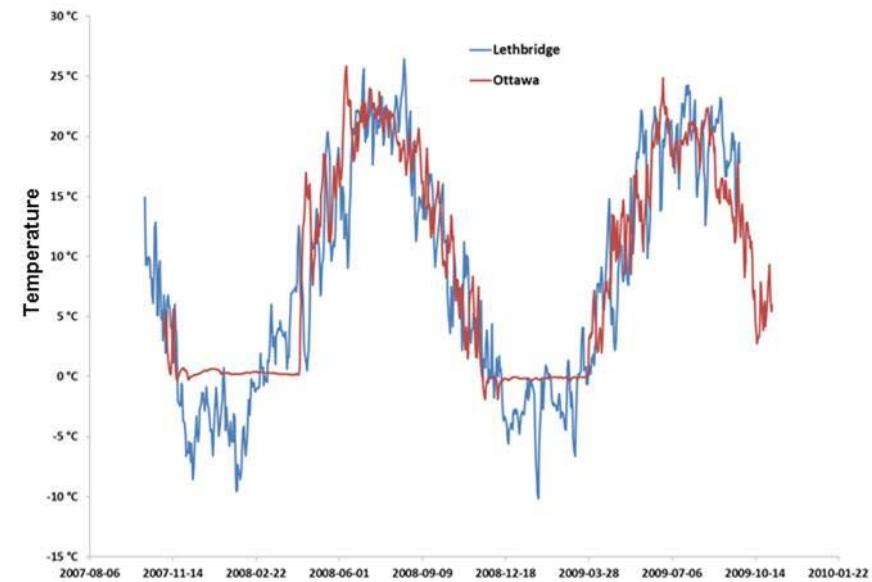


Spring 2008

# Experimental design

## Basic Site characteristics:

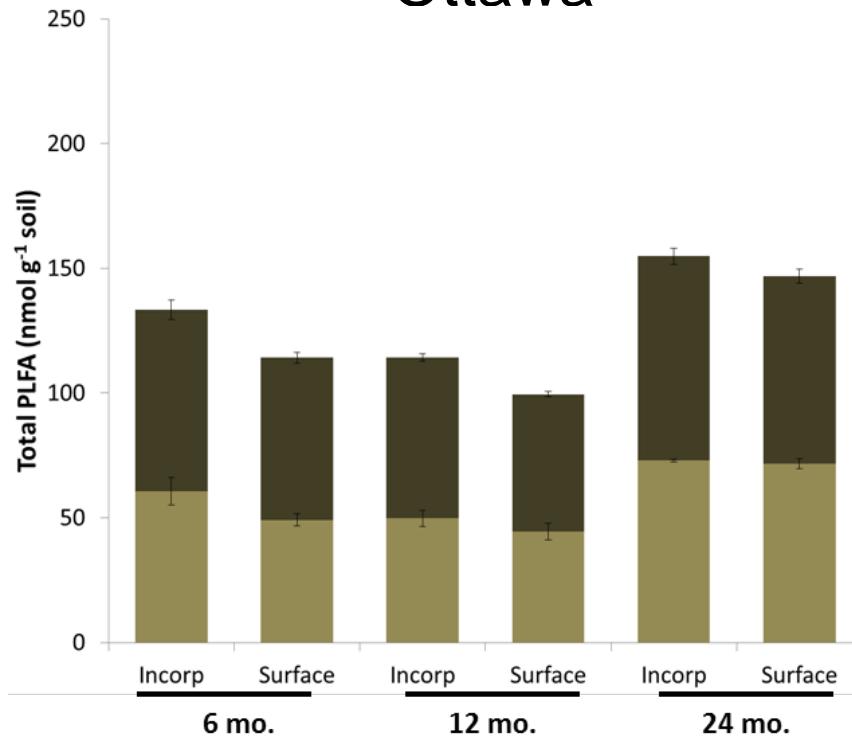
- similar mean annual temperature
- very different temporal patterns of soil temperature
- very different precipitation patterns



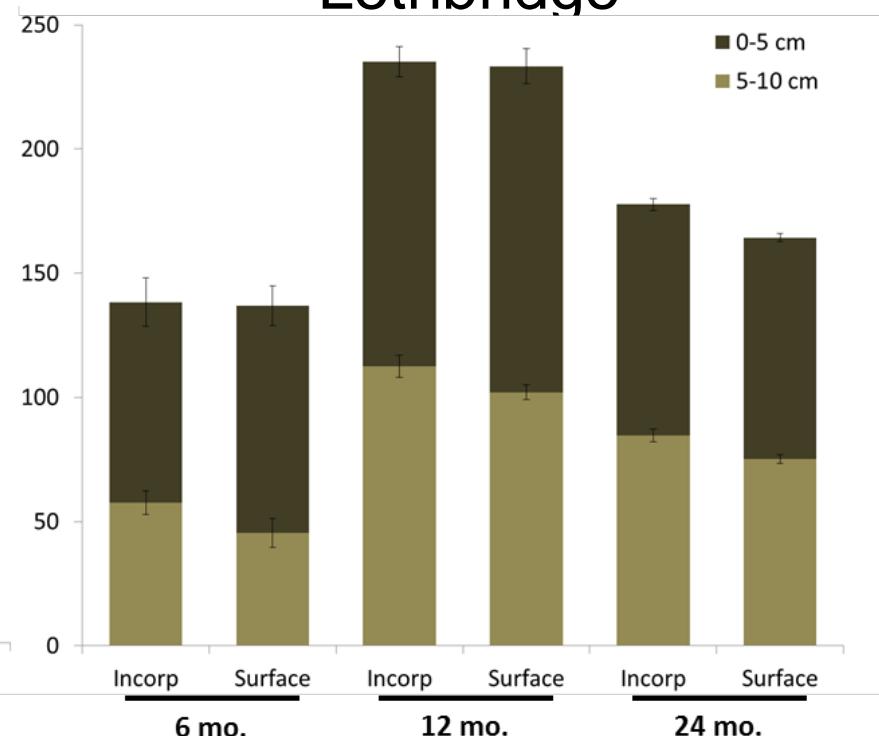


# Microbial abundance – $^{12}\text{C}$ + $^{13}\text{C}$ PLFA

Ottawa



Lethbridge

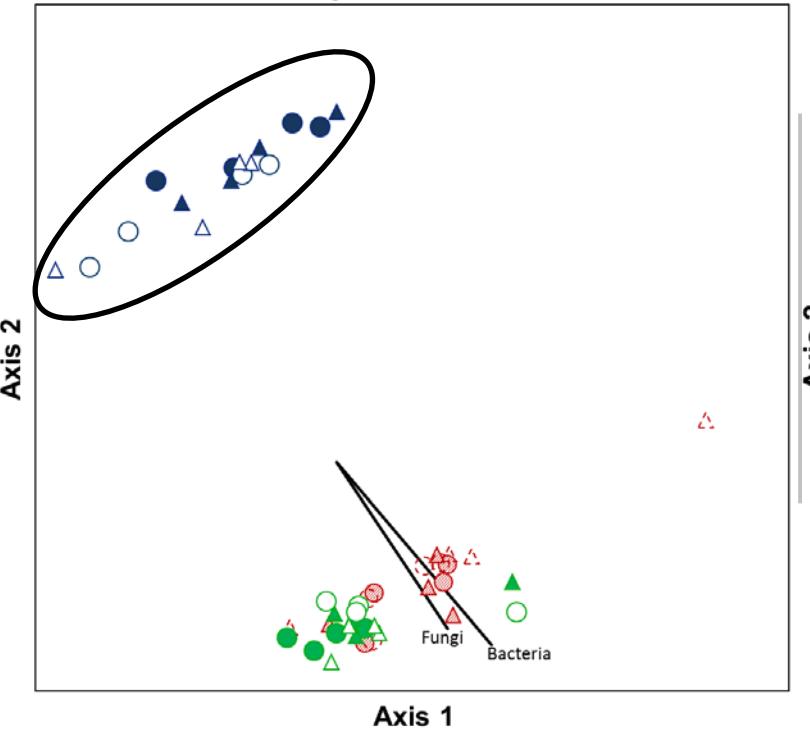


- Higher fall biomass at Lethbridge than Ottawa
- Significant biomass at 0-5 and 5-10 cm depths

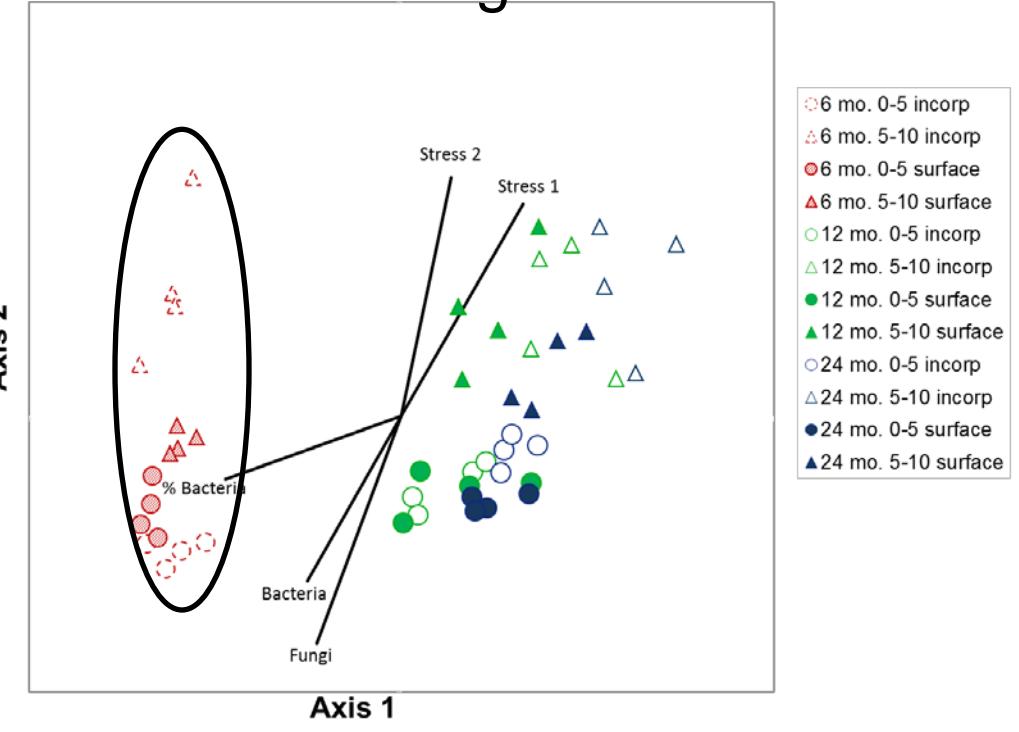


# Community structure – $^{12}\text{C}$ + $^{13}\text{C}$ PLFA

Ottawa



Lethbridge

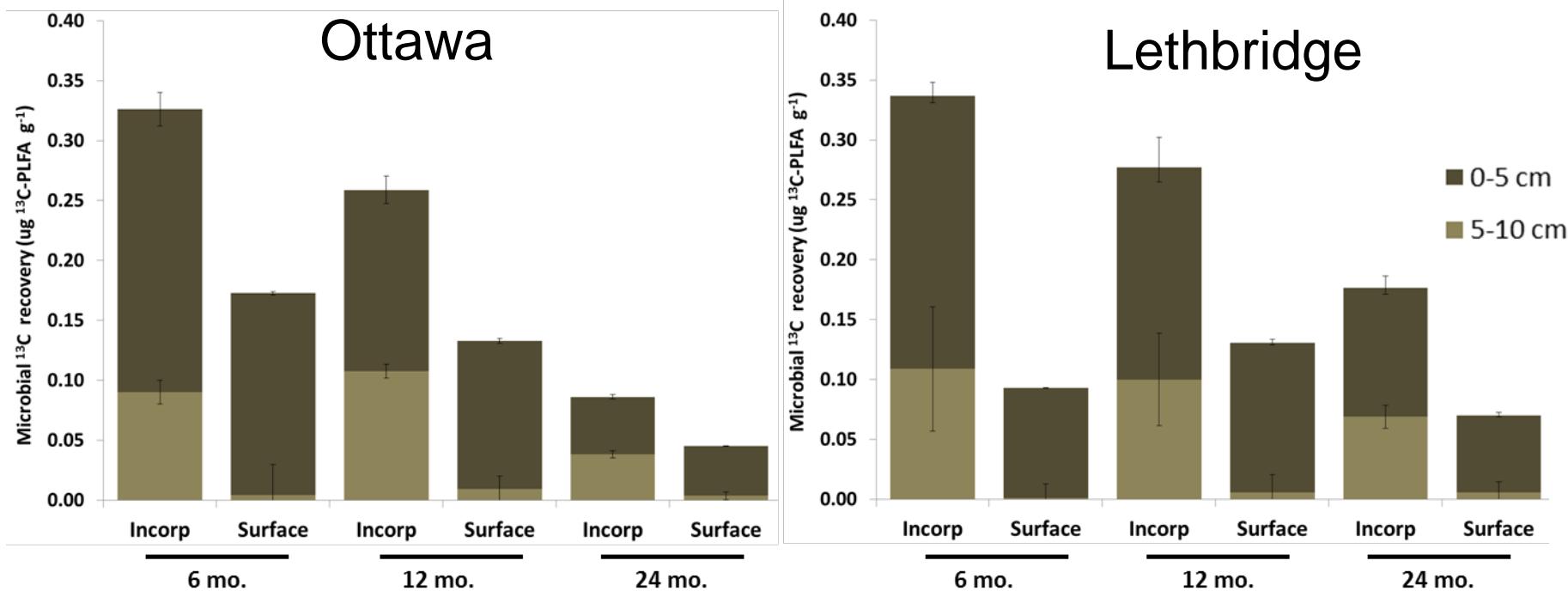


**Ottawa:** communities similar in season 1, major shift by 24 mo.

**Lethbridge:** strongest effects of season and depth



# $^{13}\text{C}$ recovery in the microbial biomass

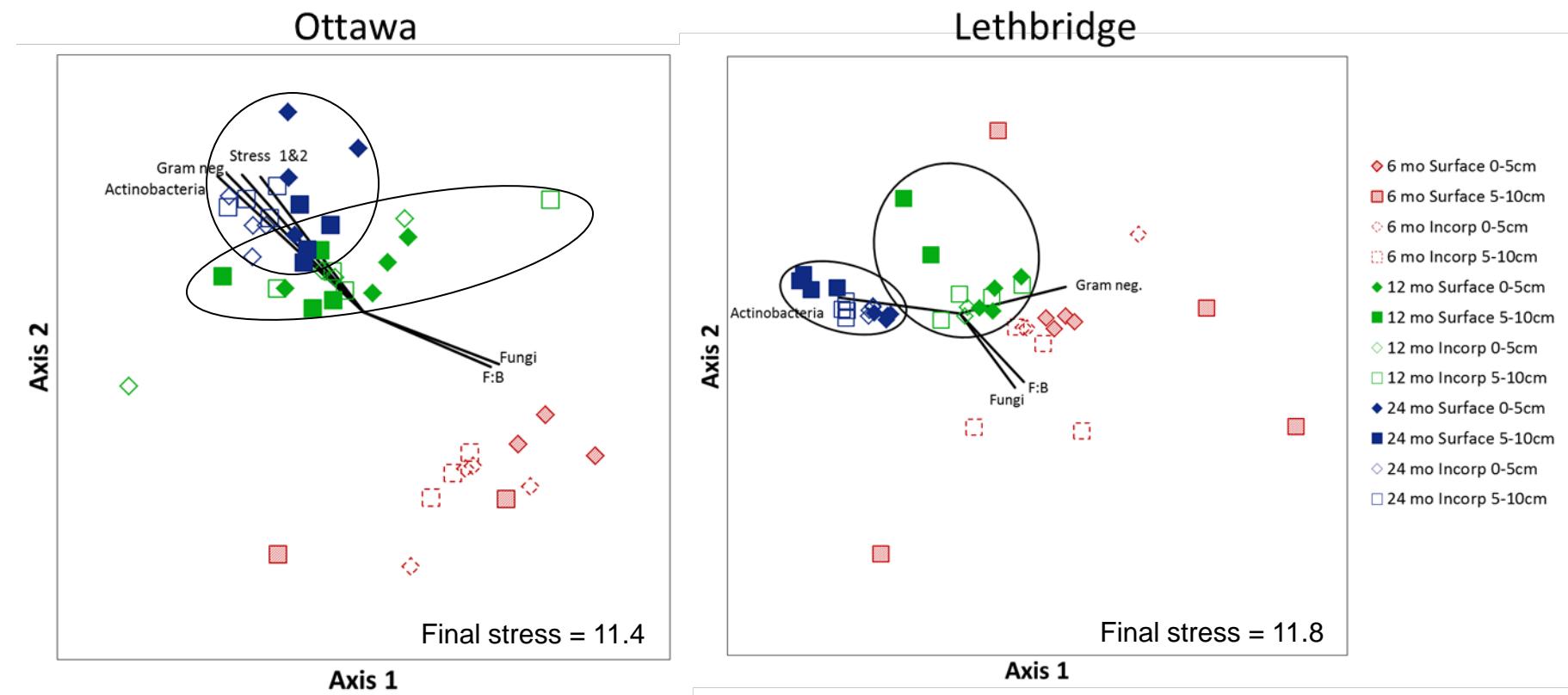


**Incorporated:** Highest recovery at 6 mo.  
Little difference between sites

**Surface applied:** Only 50% as much  $^{13}\text{C}$  recovered  
Negligible  $^{13}\text{C}$  recovery below 5 cm after 2 yr

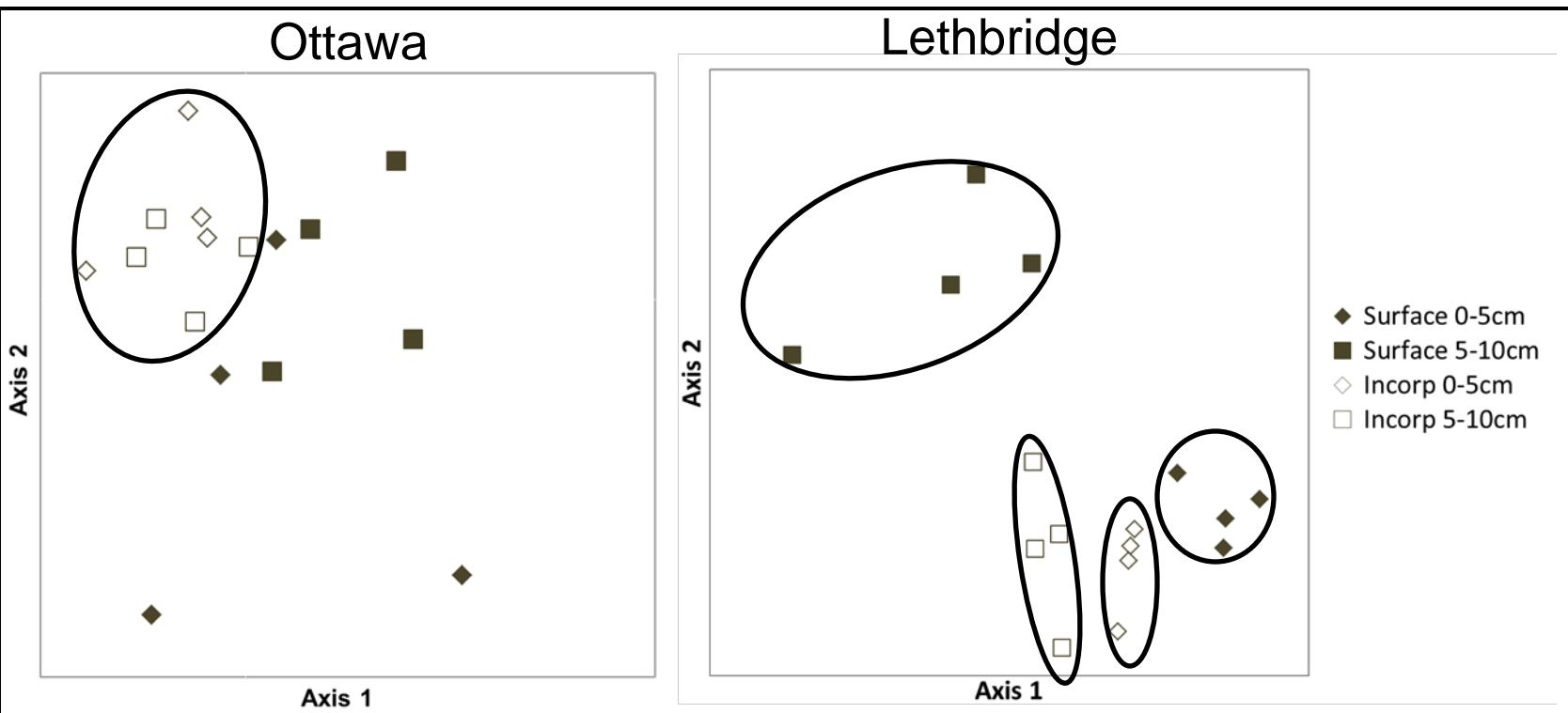


# <sup>13</sup>C recovery in the microbial biomass



- Initially, <sup>13</sup>C labelled community structure highly variable
- Stronger effect of depth at Lethbridge than at Ottawa

# $^{13}\text{C}$ PLFA Community structure: 24 mo.



2 years after the application of the  $^{13}\text{C}$  label:

- distinct community structure reflected earlier residue placement
- there was a strong effect of depth at Lethbridge

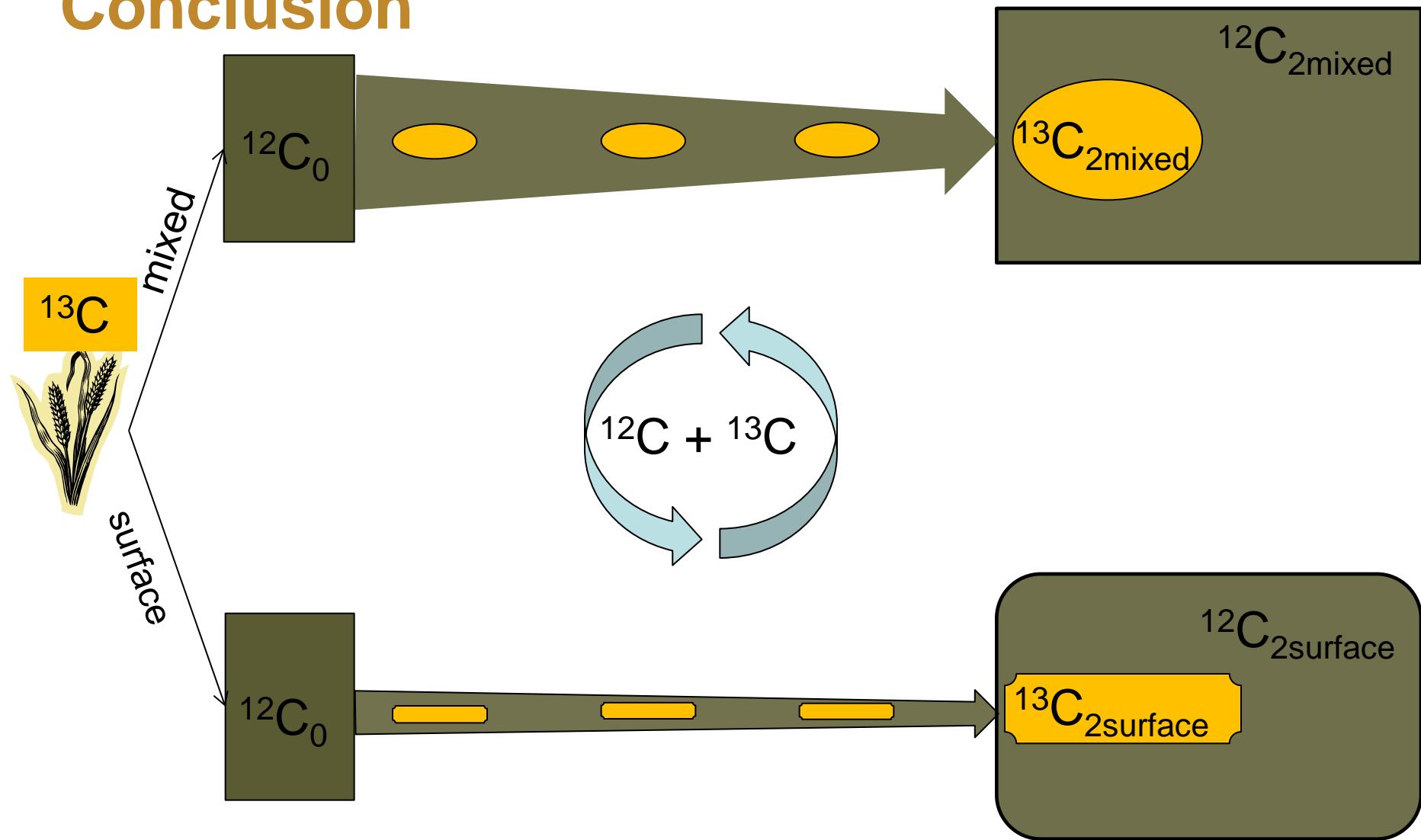
# Conclusion

The effect of residue placement was:

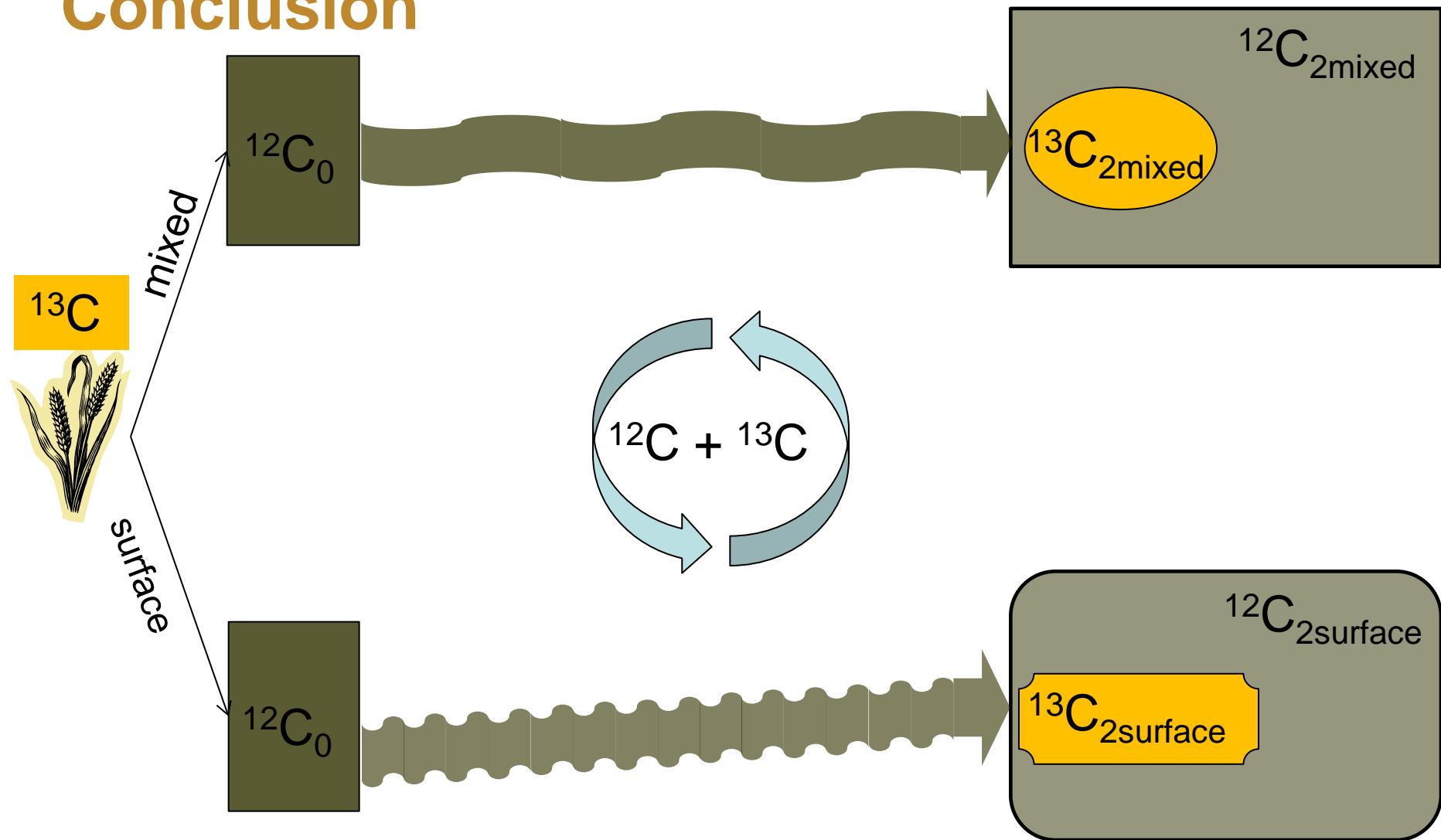
- *site-specific* with,
- a gradual shift toward a *defined community structure* of the labeled microbial biomass that reflected whether residues had been incorporated into the soil or applied to the surface *two years earlier*.



# Conclusion



# Conclusion



# Implication

Climate has a greater influence on microbial utilization of residues when they are placed on the surface than when they are mixed into the soil.

- Consequently, residue C dynamics may show much higher-site-to-site variability in no-tillage systems than in conventionally tilled systems
- This may explain why NT increases soil C in some cases, but not in others

# Acknowledgements

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