

Soil Microbial Diversity and Abundance at Russell Ranch



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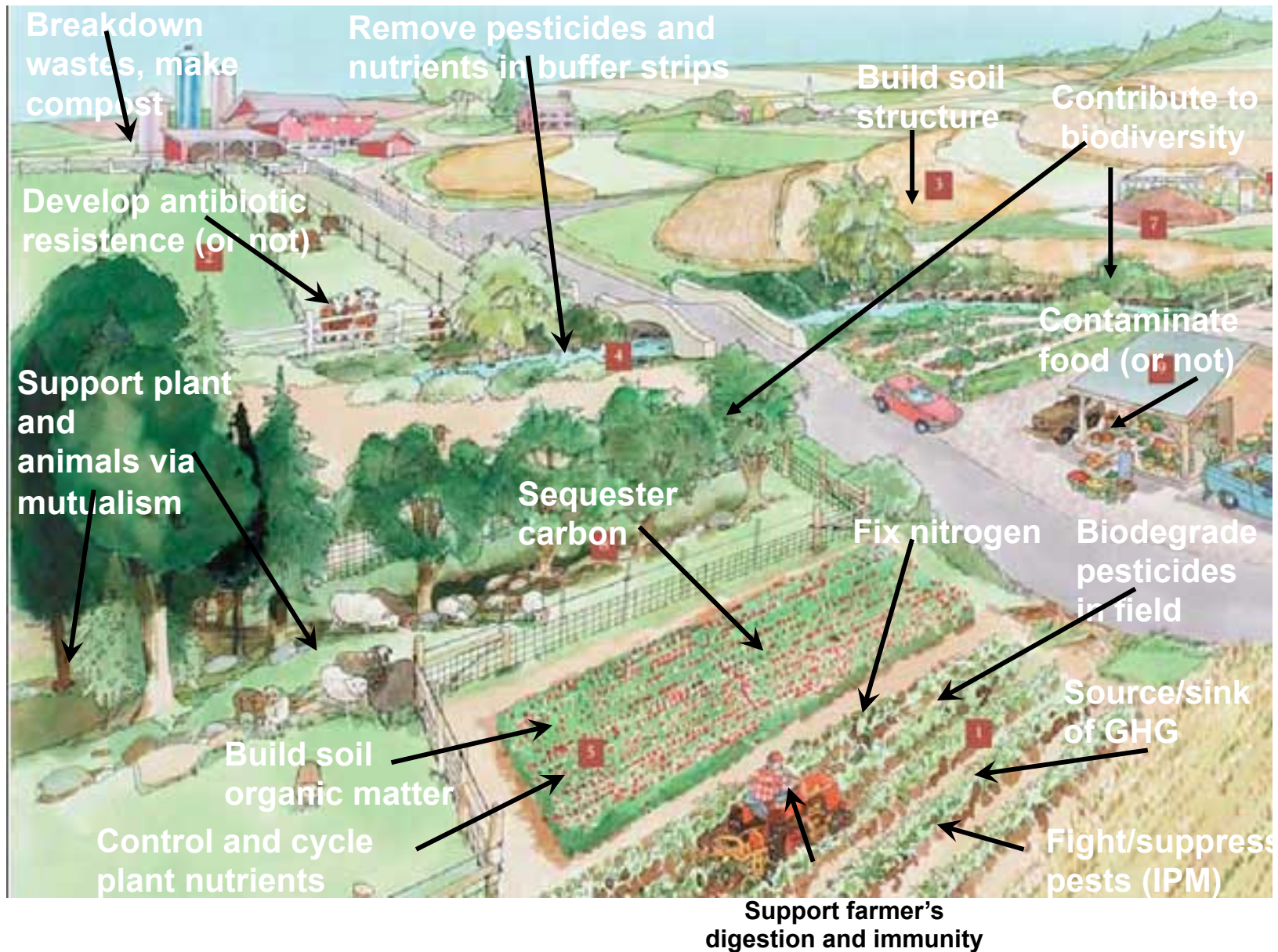
Scow Lab

Soils and Biogeochemistry

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How microorganisms contribute to agriculture

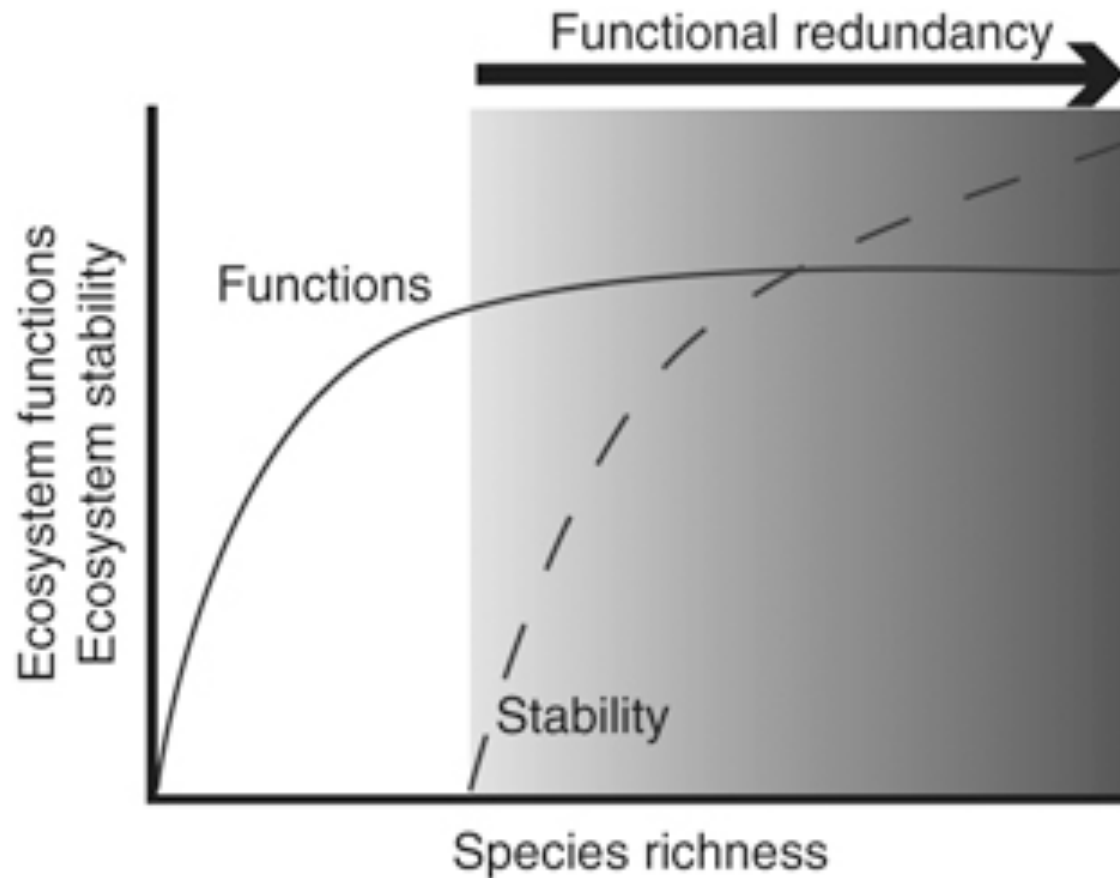
Indicators, metrics for soil biodiversity

Indicator	How measure	Why useful
<u>Identity</u> <u>Community composition</u>	DNA or RNA: clone libraries, pyrosequencing, DNA fingerprinting, metagenomics	Determines WHO is there, what relationships are there, what is “potential” of these organisms
<u>Quantity</u> •Numbers •Biomass	Fumigation-extraction, phospholipids, DNA, microscope counts,	How many or how much? How fast?
<u>Functional traits</u> •Degrades chemicals, produces chemicals •Associates with a symbiont •Kills someone •Competes with someone •Helps build structure	Substrate utilization, formation of products (e.g. CO ₂) Presence/quantity of functional genes (Geochip); RNA expression of functional genes	What are <u>IMPACTS</u> of microbes? What do they eat, excrete, who do they partner with What is activity of genes responsible for specific microbial processes

Background: Microbial biodiversity

- Diversity – degree of variation of life forms in an environment
- **Species diversity** refers to number of species in an environment and their relative abundance.
- Why do we care?
 - To achieve functional redundancy

Functional Redundancy = Stability



Functional redundancy comes from diversity

Research questions

- Are soil microbial communities in unfarmed (fallow) soils more diverse than in soils that are actively farmed?
- Does growing crops without any fertilizer reduce soil microbial diversity?
- Are soil microbial communities more diverse in organic than conventionally managed crops?

Farming Systems Compared

RECEIVING INPUTS (rotation with wheat)

- **Organic** tomato: irrigated, no synthetic pesticides, fertility from compost and cover crops
- **Conventional** tomato: irrigated, pesticides, fertility from ammonium sulfate

NO INPUTS

- Wheat: **no inputs**--not irrigated or fertilized
- Fallow: **uncropped** systems with weeds

Biomass

Total microbial biomass is ~2X higher in organic than other systems.

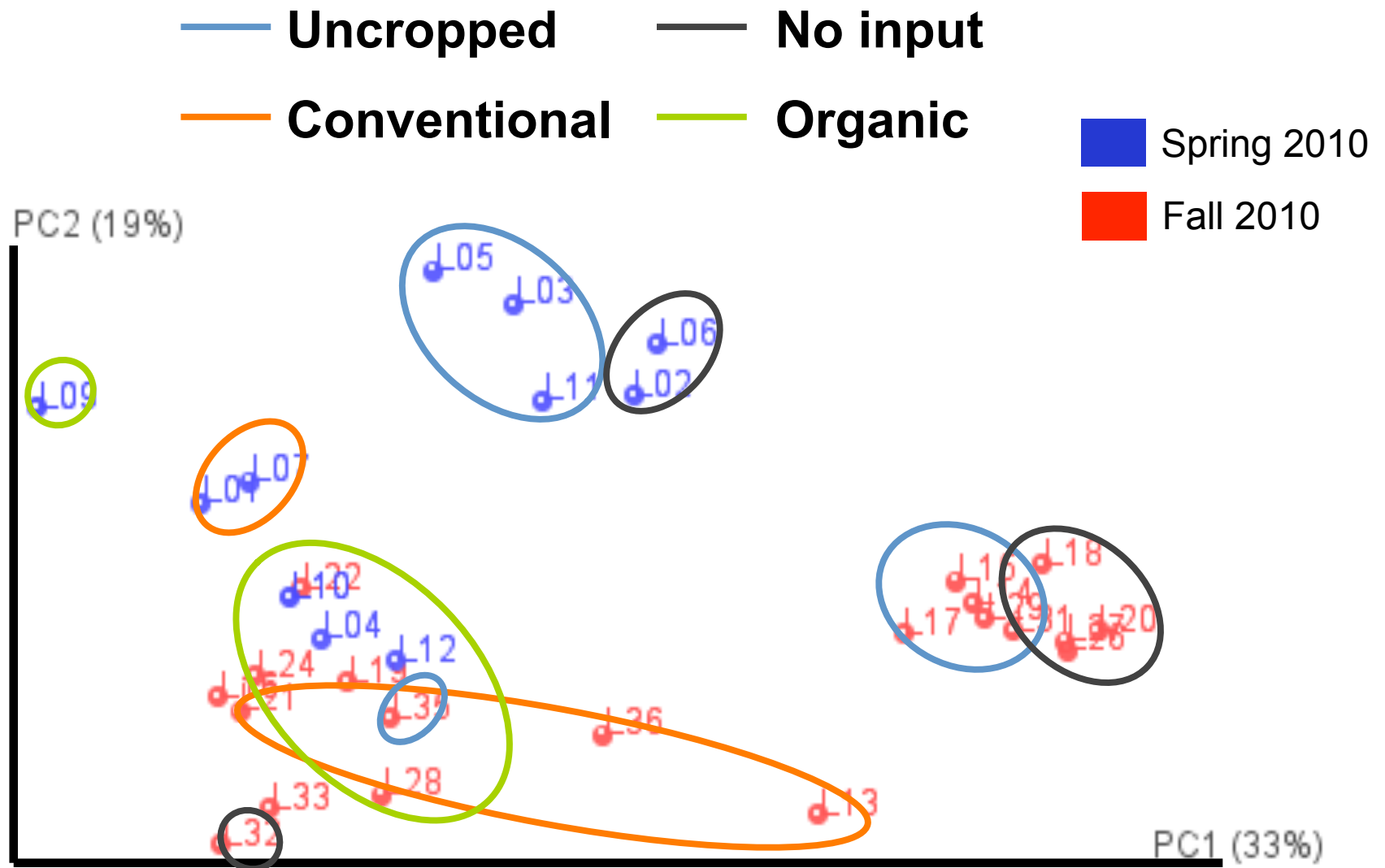
Conv, no input and uncropped not significantly different in biomass.

Species Diversity

We frequently use **DNA sequencing** to measure microbial species diversity. Each “species” has unique DNA sequences that can be used to catalog numbers and relative abundance.

Pyrosequencing is recent technology for characterizing complex microbial communities.

Community Composition by System and Season



RESULTS

- All systems have same **number** of species
- Systems differ in **composition** of their communities.
 - Organic and conventional systems more similar to one another than to no input and uncropped systems
- Most systems showed strong changes in communities in Spring versus Fall.
 - Organic less so?

Conclusions/discussion

- Organic and conventional not very different—why? How about commercial organic farms with greater crop diversity? How about commercial conventional farms with higher fertilizer/pesticide inputs?
- Seasonal variation is high in uncropped, no input systems and conventional but less so in organic. Organic has more resources to maintain and stabilize microbial communities?
- Do differences in community composition translate into
 - Differences in ability of these soils to perform key functions?
 - Differences in resistance and resilience to disturbances?

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