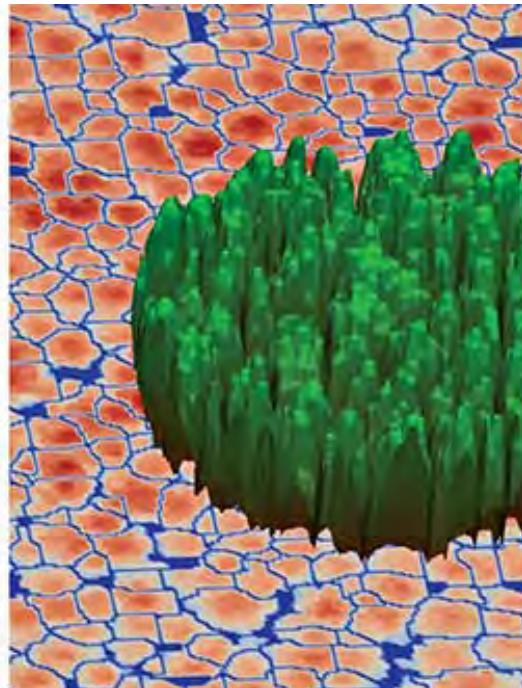


Ecosystem genomics – lessons from other sectors

Authors:
Steve A Wakelin



Humans are not alone



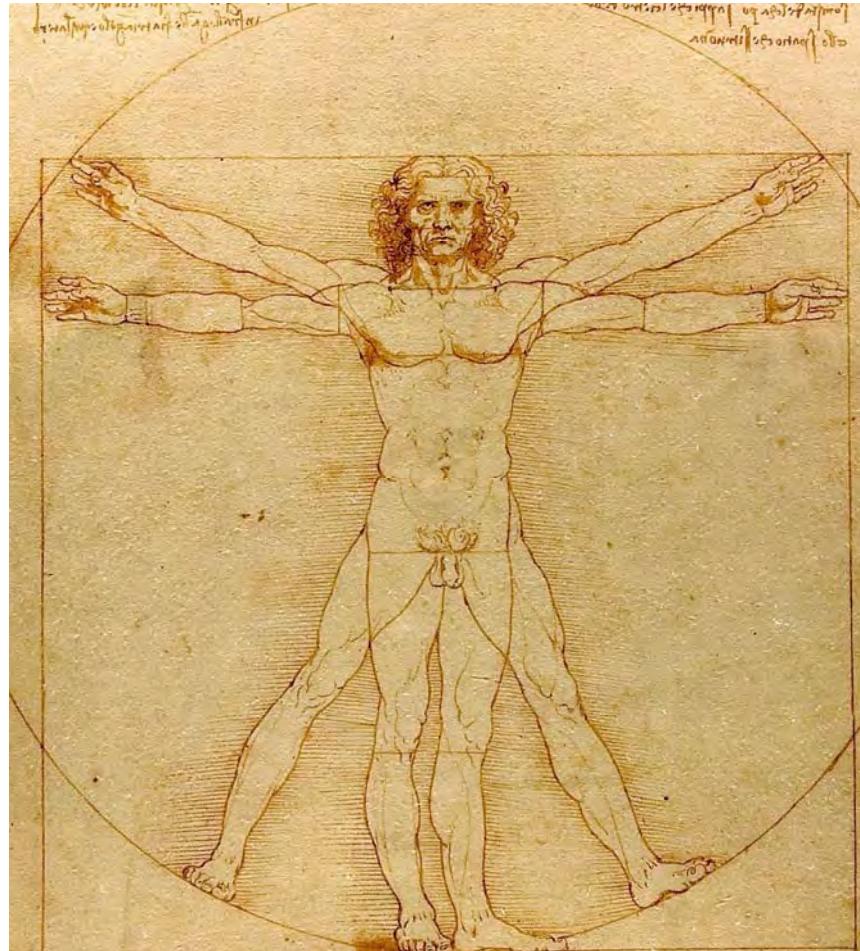
- We share our bodies with microorganisms.
- Spent decades and trillions of \$ trying to eradicate.
- Small fraction of microbes on & in us are pathogenic.



What other microbes live on and in us,
and *what do they do?*

Human microbiome project

- 2008, 5 year, \$115 Million 'feasibility' study.
- Define 'what's there', metabolism, links with human health.
- 242 volunteers.
- 5000 samples: mouth, nose, skin, gut, lower intestine (stool), vagina.

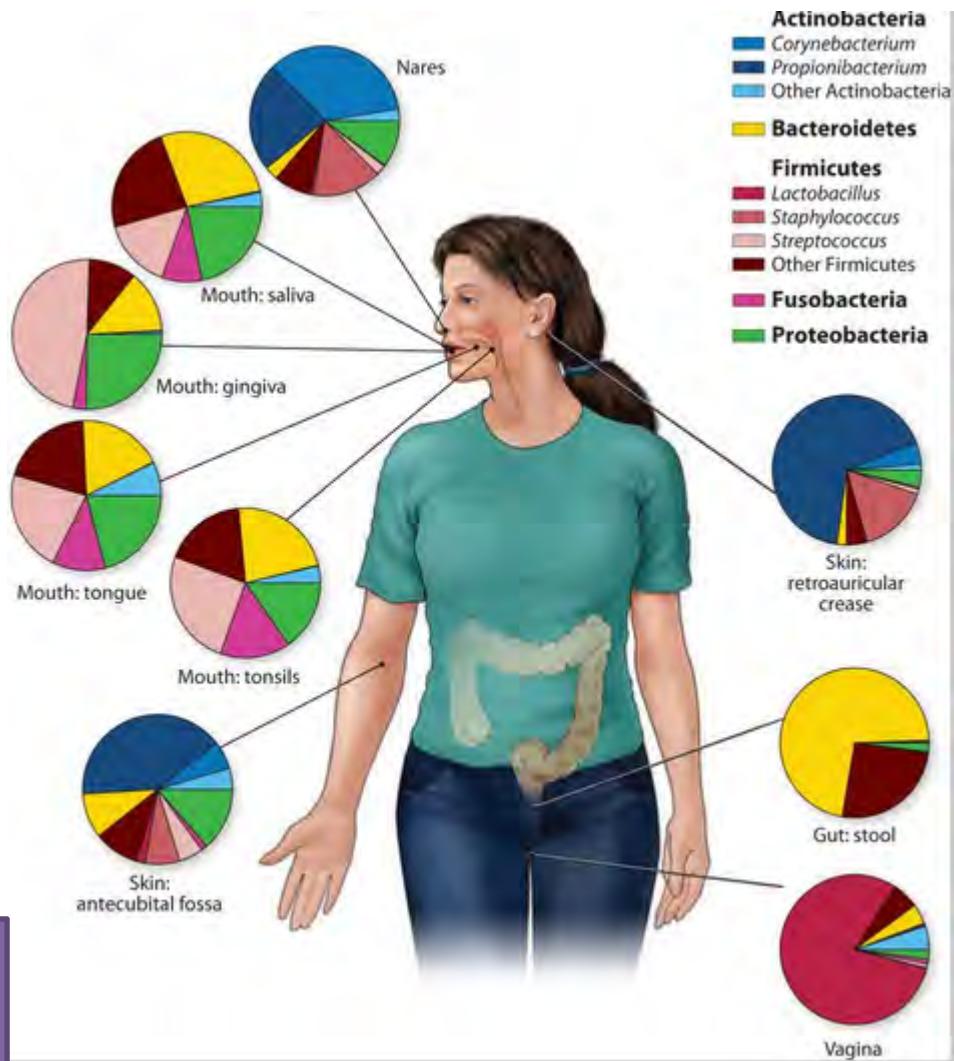


Results are so profound and unexpected, its still going.
Revolutionising human health and medicine.

Human microbiome project

- > 10,000 bacterial species;
100 million million cells
- 1:1 microbial:human cells
- Each body part is a distinct
'ecosystem'.
- Vary by individual, health,
age, gender, etc.

All ecosystems x human genome
= entire 'holobiome'



Functional significance?

Gut ecosystem

- 1000 species of and **3.3 million** genes
- Dwarfs the entire human genome's **23,000** genes
- Microbes contribute more genes for human survival than humans own genes.
- Vitamin production, complex decomposition, steroid & hormone.
- Rather than evolving new functions, we have *acquired microbial symbionts* that conduct specialised functions for us.

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Published online 3 March 2010 | Nature | doi:10.1038/news.2010.104

News

Gut bacteria gene complement dwarfs human genome

Sequencing project finds that Europeans share a surprising number of bacteria.

Article

Cell

Gut Microbiota Regulate Motor Deficits and Neuroinflammation in a Model of Parkinson's Disease

Is chronic fatigue linked to gut bacteria?

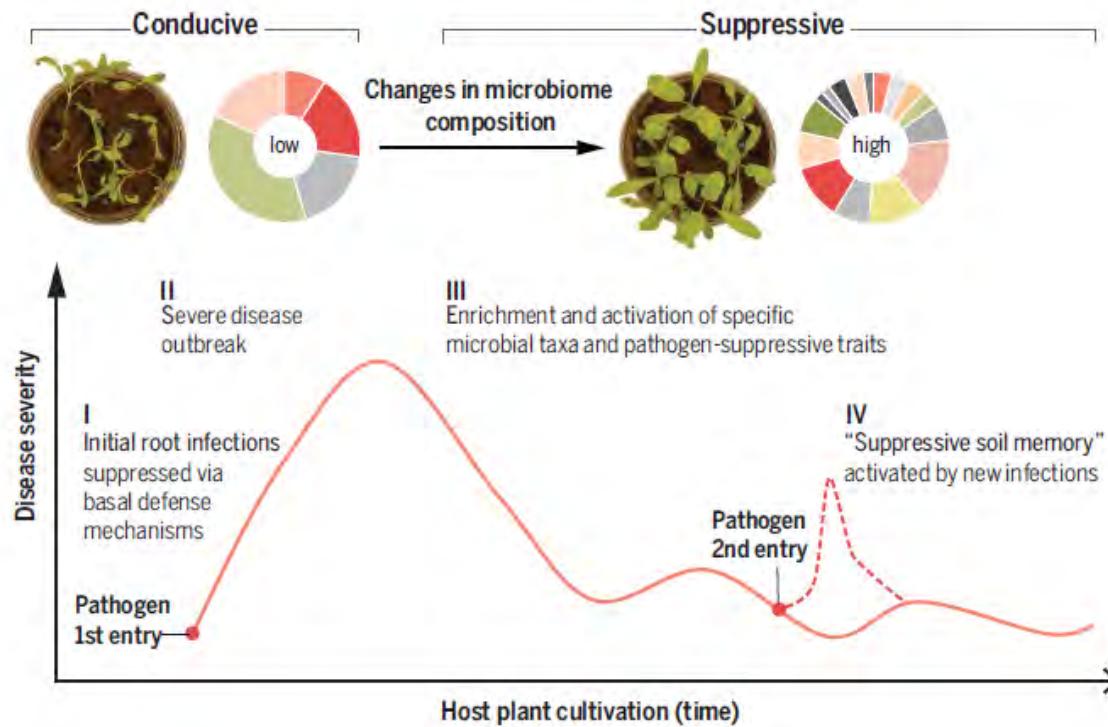
By Ariana Eunjung Cha

6:00 PM Tuesday Jul 5, 2016

Terrible twos blamed on bacteria (especially boys)



Where the gut goes, soils and plants can follow...



Science, 352 (2016)

ECOLOGY

Soil immune responses

Soil microbiomes may be harnessed for plant health

frontiers in
MICROBIOLOGY

PERSPECTIVE ARTICLE
published: 03 December 2013
doi: 10.3389/fmicb.2013.00355



Bespoke microbiome therapy to manage plant diseases

Murali Gopal*, Alka Gupta and George V. Thomas

Microbiology Section, Central Plantation Crops Research Institute, Kudlu, Kasaragod, India

Vineyard microbiome → grape quality

The Soil Microbiome Influences Grapevine-Associated Microbiota

Iratxe Zarraonaindia,^{a,b} Sarah M. Owens,^{a,c} Pamela Weisenhorn,^c Kristin West,^d Jarrad Hampton-Marcell,^{a,e} Simon Lax,^e Nicholas A. Bokulich,^f David A. Mills,^f Gilles Martin,^g Safiyh Taghavi,^d Daniel van der Lelie,^d Jack A. Gilbert^{a,e,h,i,j}

Argonne National Laboratory, Institute for Genomic and Systems Biology, Argonne, Illinois, USA^a; IKERBASQUE, Basque Foundation for Science, Bilbao, Spain^b; Computation Institute, University of Chicago, Chicago, Illinois, USA^c; Center of Excellence for Agricultural Biosolutions, FMC Corporation, Research Triangle Park, North Carolina, USA^d; Department of Ecology and Evolution, University of Chicago, Chicago, Illinois, USA^e; Departments of Viticulture and Enology; Food Science and Technology; Foods for Health Institute, University of California, Davis, California, USA^f; Sparkling Pointe, Southold, New York, USA^g; Department of Surgery, University of Chicago, Chicago, Illinois, USA^h; Marine Biological Laboratory, Woods Hole, Massachusetts, USAⁱ; College of Environmental and Resource Sciences, Zhejiang University, Hangzhou, China^j

Proceedings of the National Academy of Sciences

PNAS | January 7, 2014 | vol. 111 | no. 1 | 5–6

Microbial terroir for wine grapes

Jack A. Gilbert^{a,b,1}, Daniel van der Lelie^c, and Iratxe Zarraonaindia^{a,d}

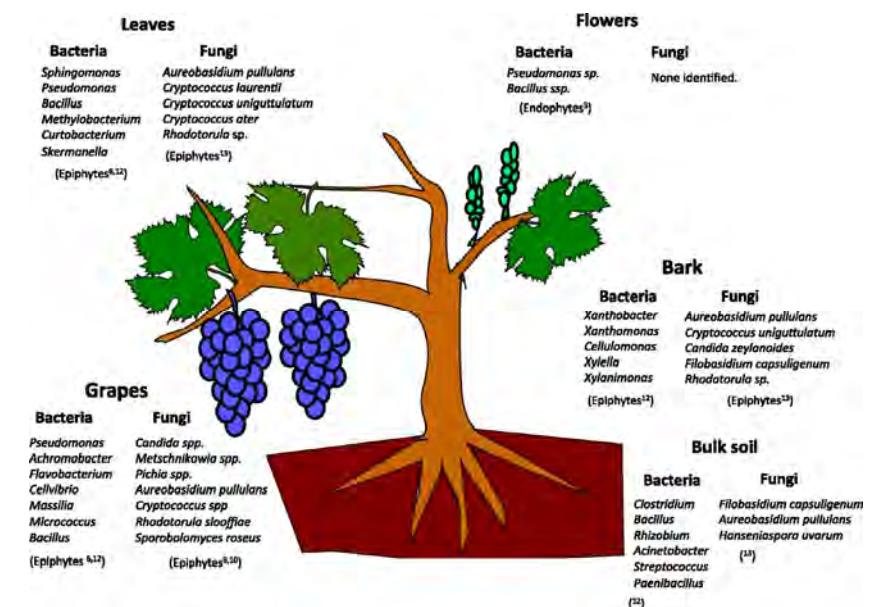
^aInstitute for Genomic and Systems Biology, Argonne National Laboratory, Argonne, IL 60439;

^bDepartment of Ecology and Evolution, University of Chicago, Chicago, IL 60637; ^cCenter of

Excellence for Agricultural Biosolutions, FMC Corporation, Research Triangle Park, NC 27703;

and ^dIkerbasque, Basque Foundation for Science, 48011 Bilbao, Spain

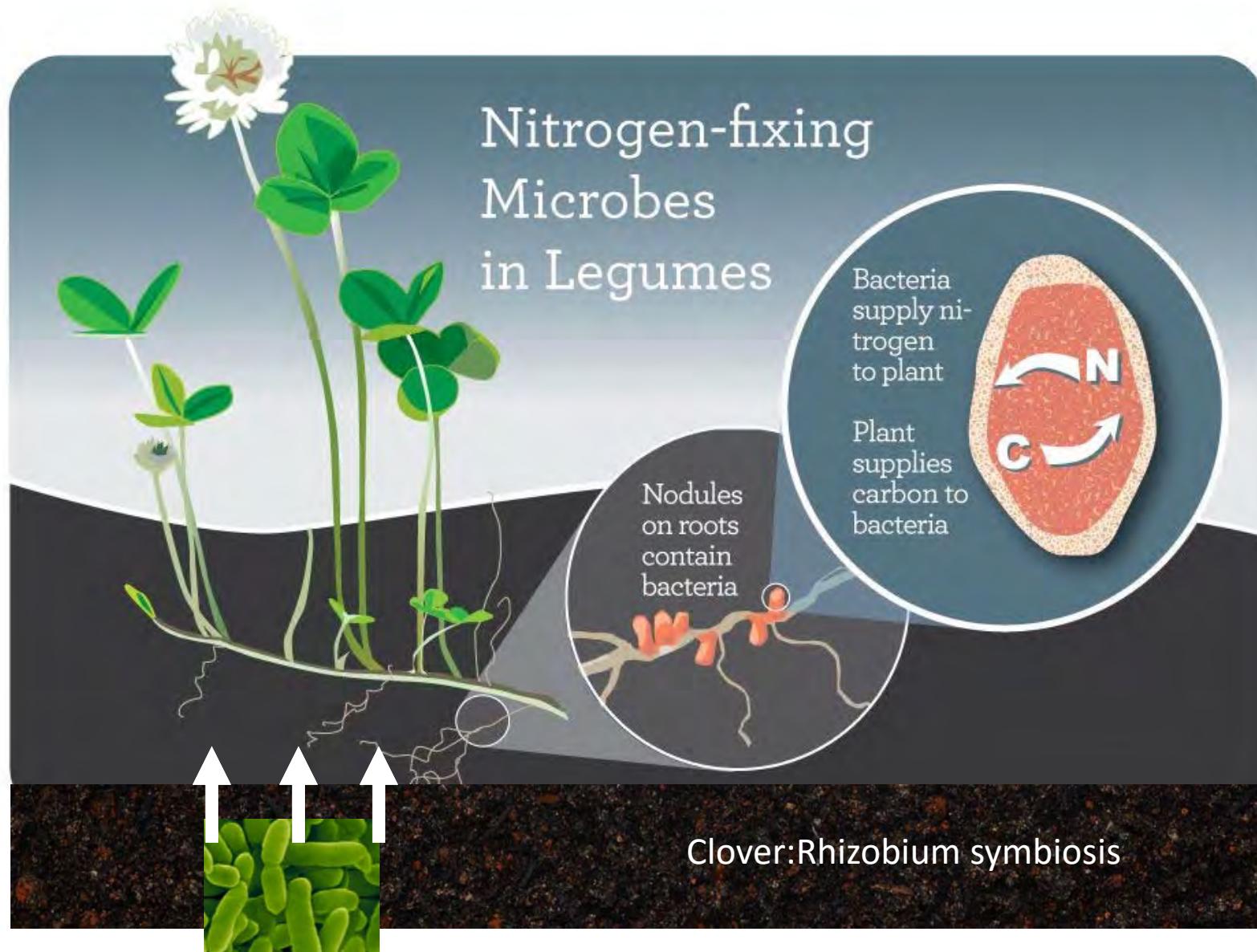
presence of ph
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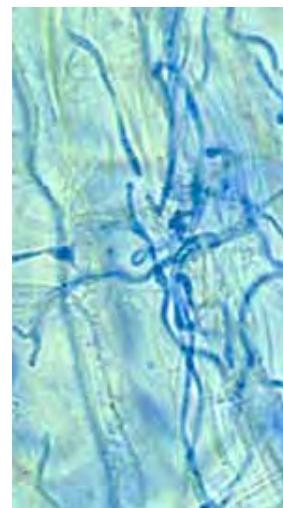
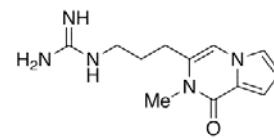
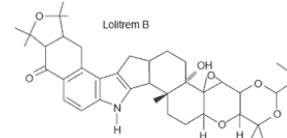
Dairy sector based on a pasture 'holobiome'



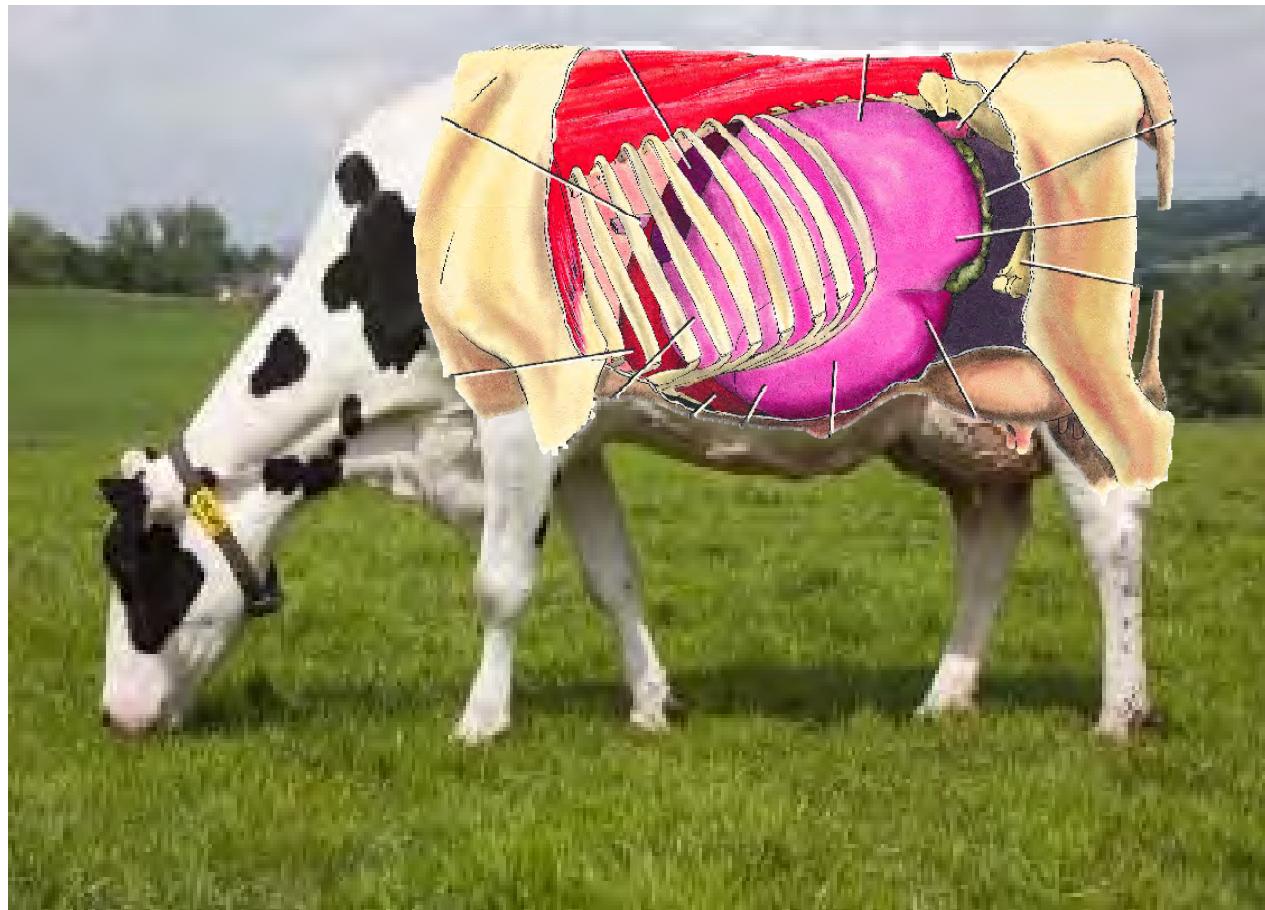
Clover ecosystem: supplies nitrogen to ryegrass



Ryegrass ecosystem: biomass for livestock



Livestock ecosystem: fiber and food



Rumen

2×10^{24} organisms

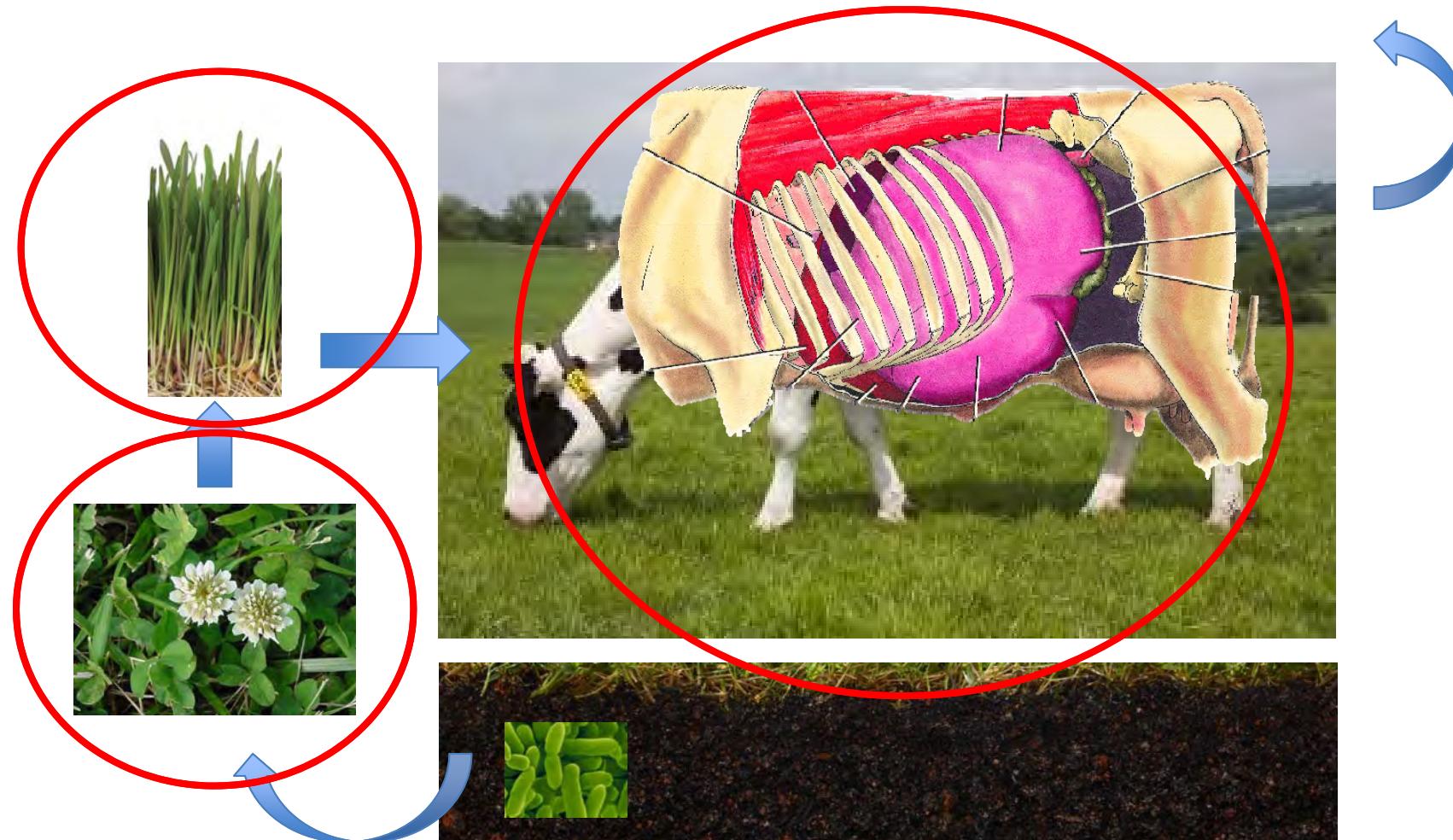
Convert plants to
'metabolisable energy'.

Unwanted byproducts

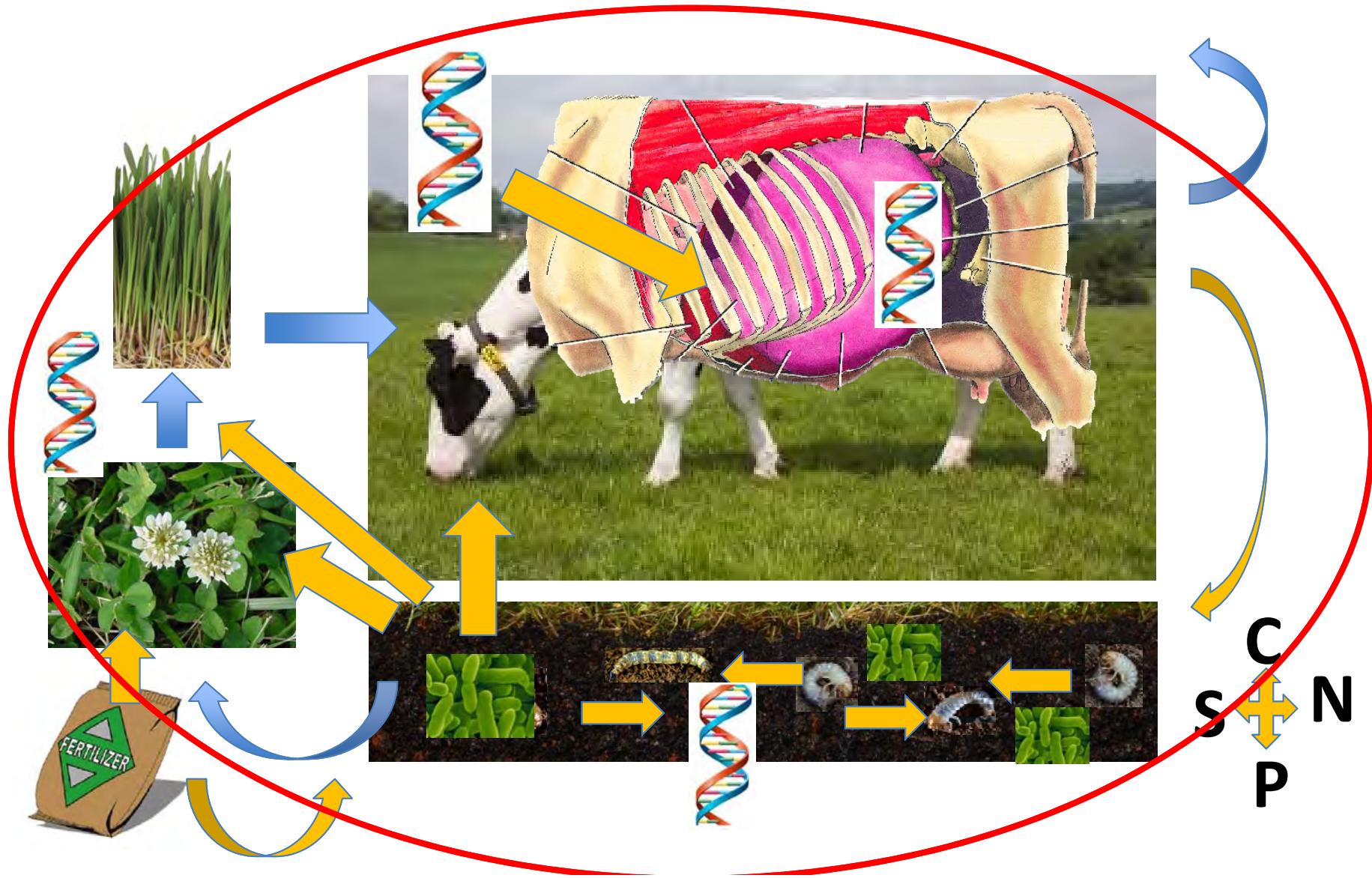
Huge \$:

1. increase feed efficiency and alter milk characteristics
2. reduce CH_4

Meta-ecosystem: integration of several ecosystems



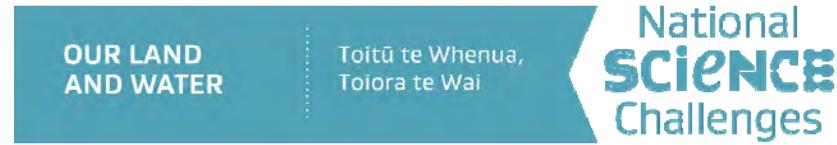
Sub-systems highly coupled and interdependent = holobiome



Dairy Sector: Identifying opportunities for multiple ecosystem interactions

- Experimentally testing interactions among bovine genomes x plant genomes x soil genomes x env (G³xE)

“Understanding the connections and interactions between soil, water, plant and animal microbiomes to form innovative microbiome-informed technologies to drive greater agricultural productivity with reduced environmental impacts”.



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Te Ahuwhenua, Te Kai me te Whai Ora. **Tuatahi**



Landcare Research
Manaaki Whenua

Plant & Food
RESEARCH
RANGAHAU AHUMARA KAI



Microbial ecosystems underpin forest success



Social and cultural “licence to operate” and market access



Tree Pests and diseases

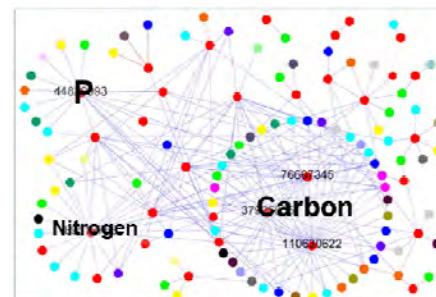
Climate change, land use change, soil ecosystem services, & NRM



Invasion of weeds and pests



Plant cultivar
X
Microbiome
X
Environment

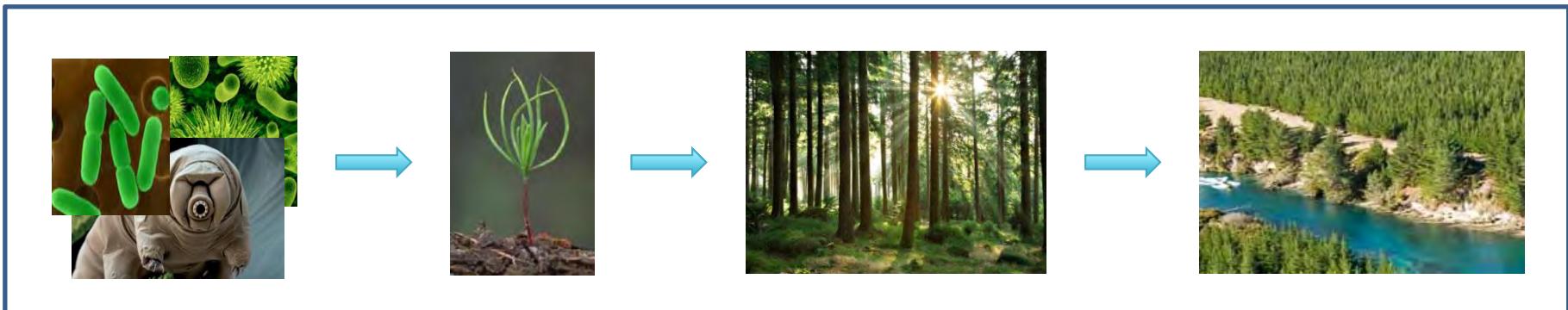


Biogeochemistry C, N, P, S,

Massive opportunity to manage the forest holobiome

Assess ecology within many sub-ecosystems and expand to entire forest holobiome

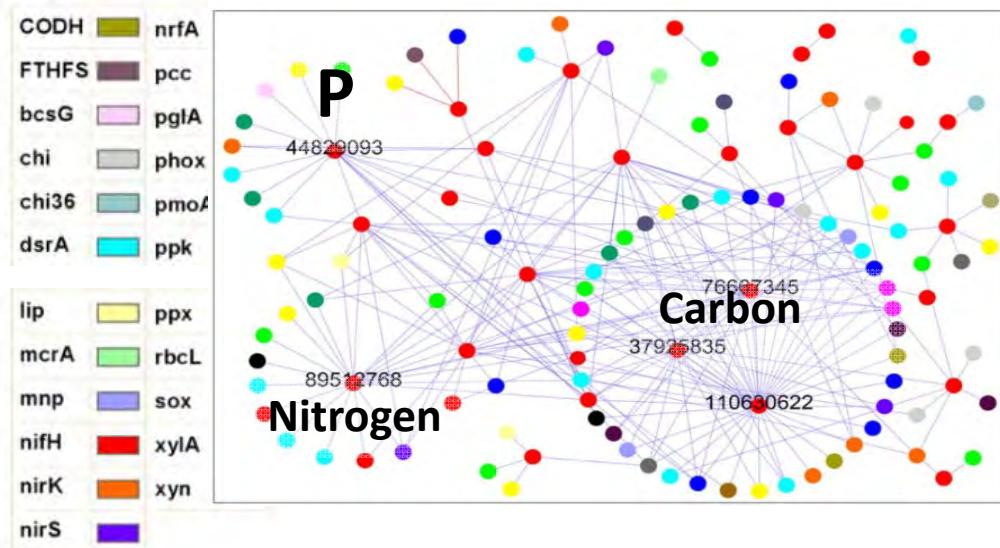
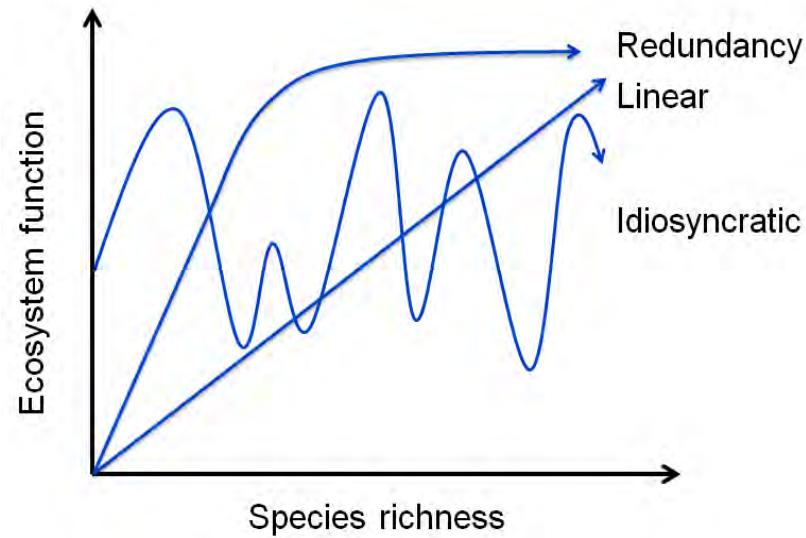
- Huge potential for genetic gain across a wide range of functions (nutrients, disease, NRM).



If it was easy, it would have been done. Its not easy....

- Massive richness of species and highly complex ecosystems.
- Interactions often not predictable.
- Multiple process with emergent properties

Functions do not exist in isolation



Species-function relationships:

- Not immediately apparent
- Difficult to assess
- Don't follow linear responses

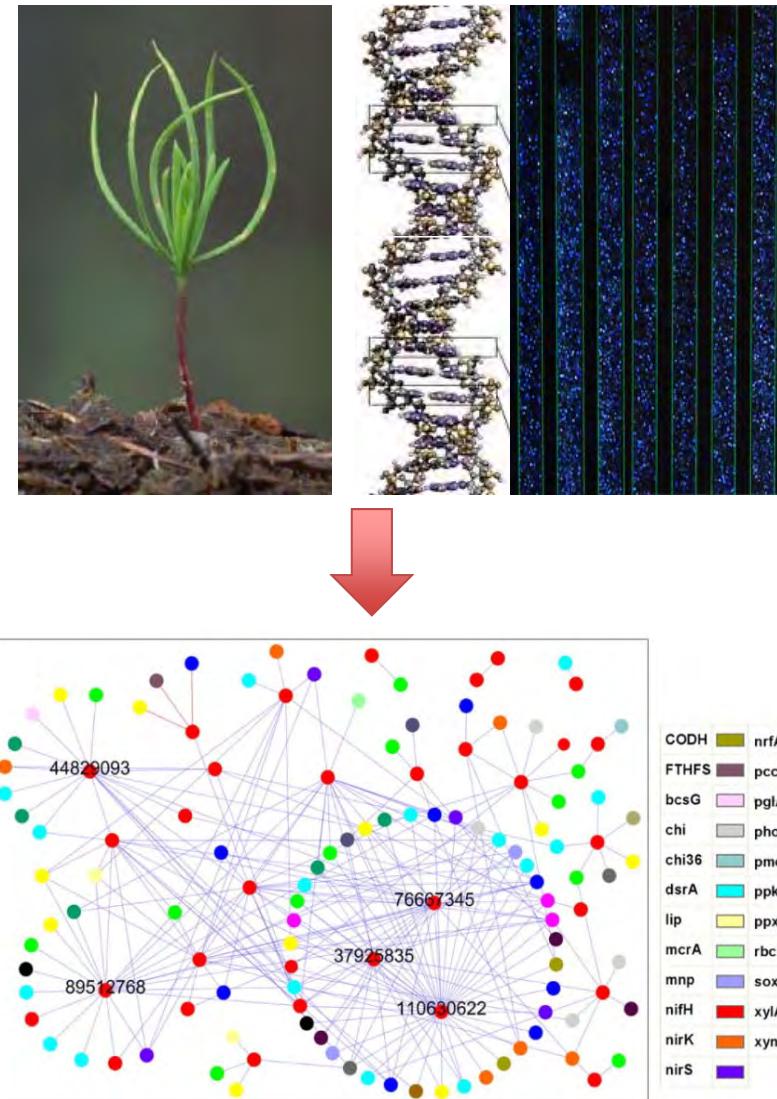
Ecosystem phenotype / functional state

- Emergent outcome of all linked processes and interactions
- High functional integration at an ecosystem level

How do we get to this level of understanding?

Approach: Environmental Genomics

- Reconstruct each sub-ecosystems from the gene up.
- Approaches based on function, species, or combination.
- Complex systems analysis and network theory to account for process coupling and interdependencies.



Building 'genetic gain' in pasture ecosystems

- Recently described a pipeline / approach to build 'genetic gain' in pastoral soils

Wakelin et al., 2016

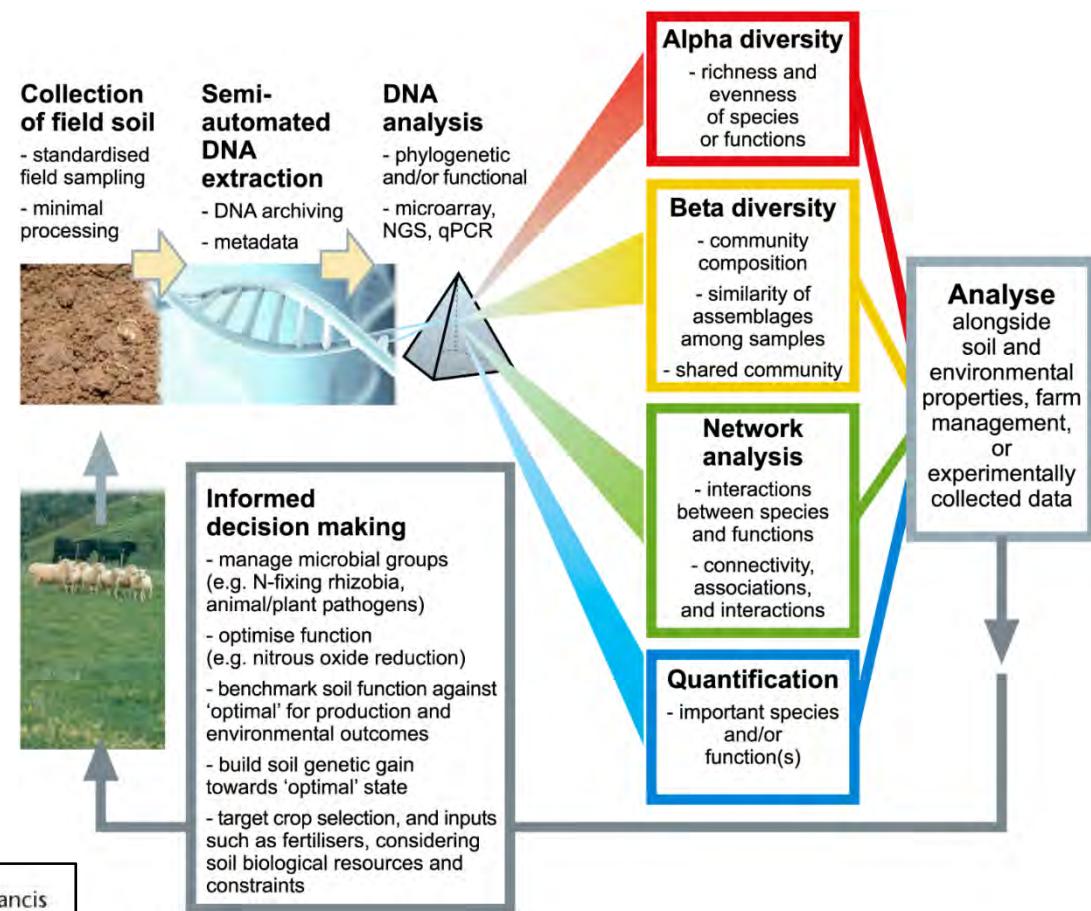
NEW ZEALAND JOURNAL OF AGRICULTURAL RESEARCH, 2016
VOL. 59, NO. 4, 333–350
<http://dx.doi.org/10.1080/00288233.2016.1209529>



RESEARCH ARTICLE

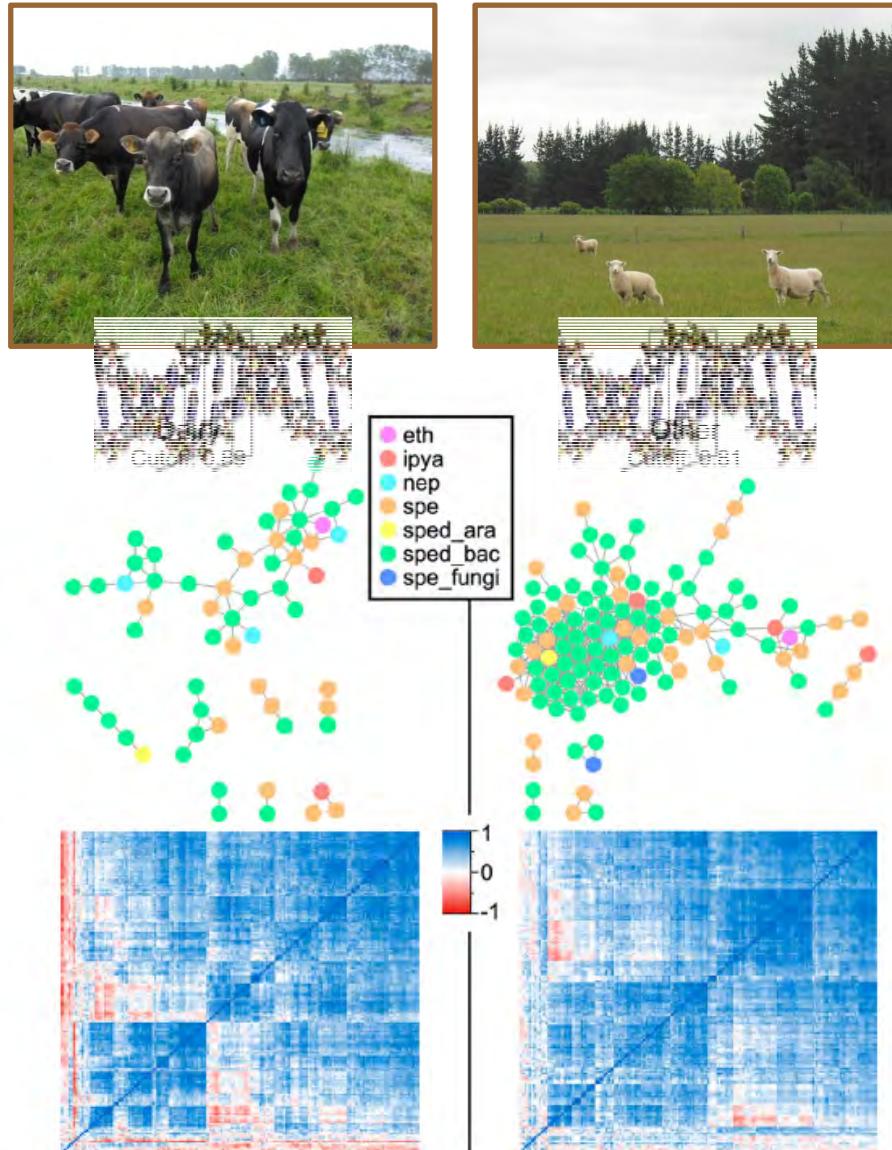
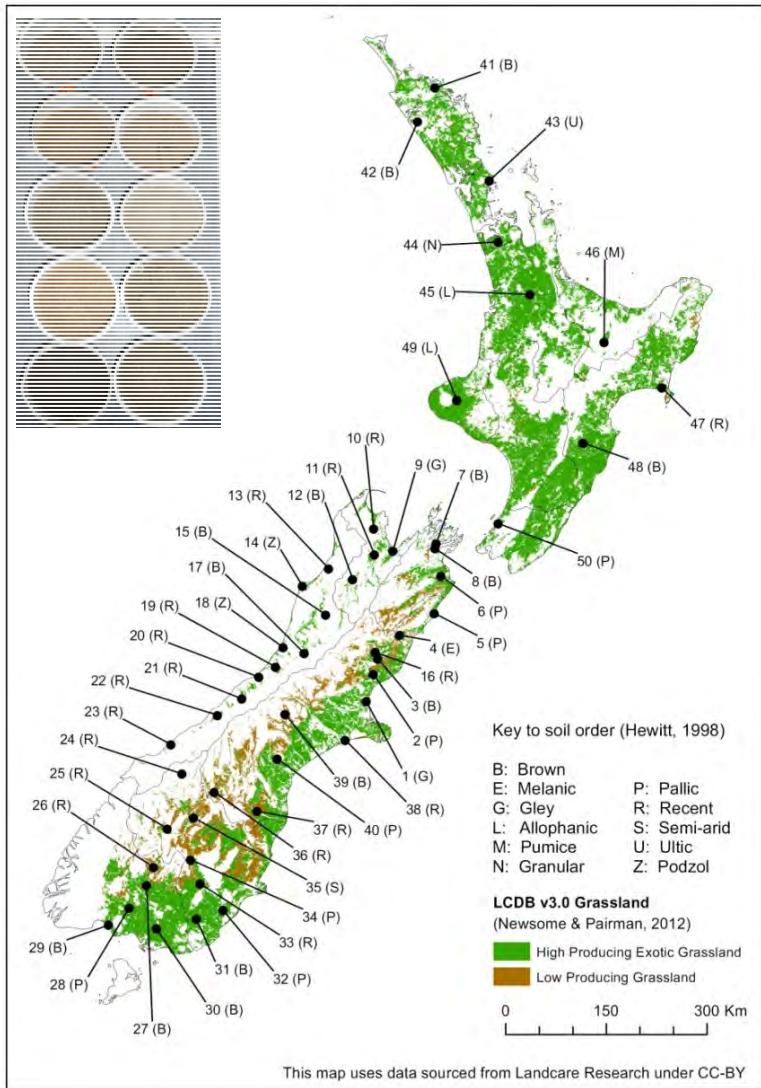
Analysis of soil eDNA functional genes: potential to increase profitability and sustainability of pastoral agriculture

SA Wakelin^a , VM Cave^a, BE Dignam^{a,b}, C D'Ath^a, M Tourna^{a,c}, LM Condron^{b,d}, J Zhou^{e,f,g}, JD Van Nostrand^e and M O'Callaghan^a



Translatable to other land-uses

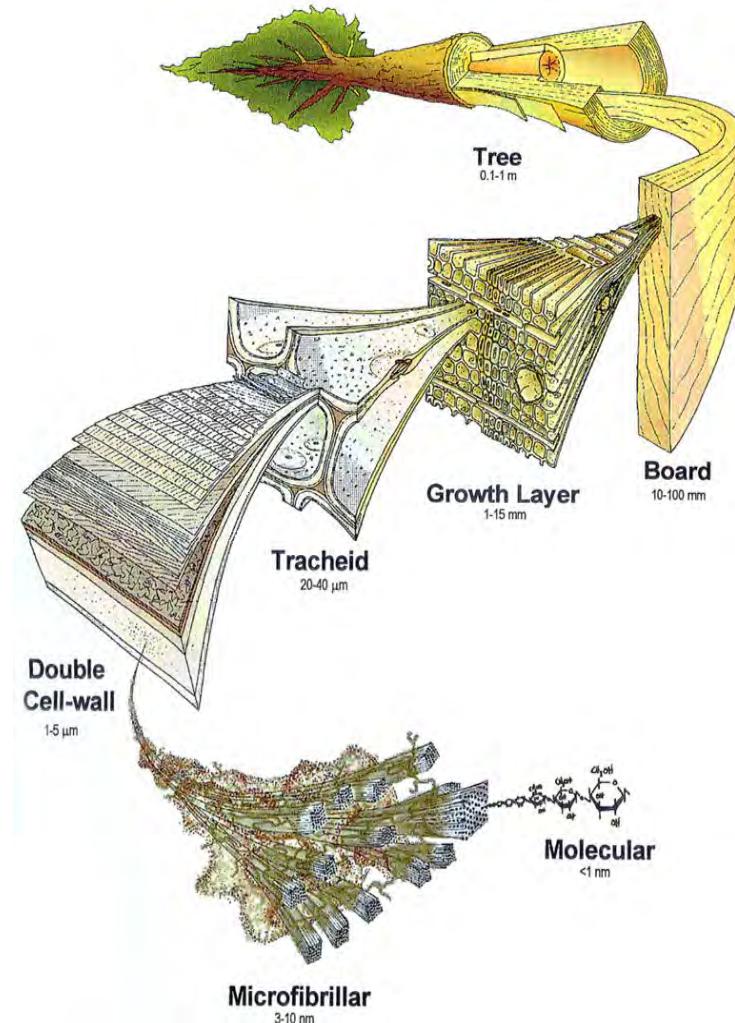
Example: Environmental genomics in NZ pastures



Current opportunities

Determining the role of the ‘pine microbiome’ – from the plant to the soil

- Direct plant productivity – beneficial v deleterious biota
- Stress tolerance, environmentally resilient forests
- Wood formation and quality (cambial, root, soil microbiomes)
- Root-microbe-soil interactions for water and fertiliser use efficiency
- Disease resistant forest soils (e.g. *Phytophthora* resistant)



Future opportunities

Introduction of novel plant microorganism associations. New traits via the 2nd genome

Utilise knowledge of the soil microbiome x environment to inform planting (genotype) and silvicultural processes.

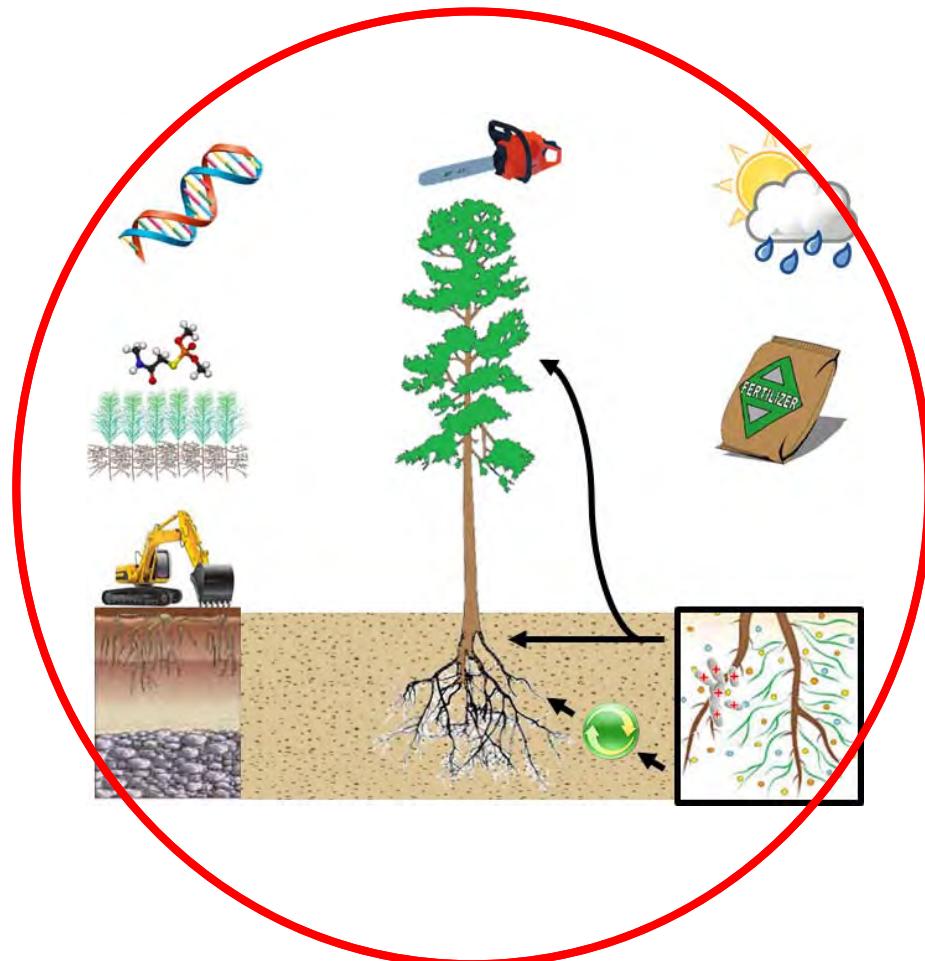
Provenance of wood/products from NZ forests – understanding ecosystem health and protection of native microbiota.

More profitable, resilient, sustainable, and socially acceptable forestry.



Ecological genomics for planted forests

- Like the human genome, there will be a lot of things to discover, that we don't even know to look for yet.
- Many things we know that we don't know.
- But it's the unknown unknowns that will have the most disruptive influence on opportunities for future forest.
- Start now, to build foundation-base for research in 5-10 years time....





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Date: March 2017

