

Environmental and Ecological Interactions of Forest Soil Fungi

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The fungal kingdom

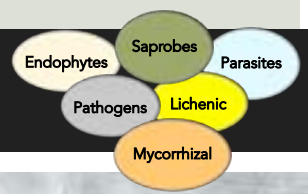


Ascomycota diversity. (A-B) Apothecia, (C-D, F-G) Thallus with apothecia. (E) Bitunicate asci. (H) Perithecia. (I) Earth-tongue apothecia. (J) Cleistothecia (K) Operculate ascus of *Peziza*. (L) *Ascostroma* (M) Unitunicate asci. (N) Prototunicate ascus.



Basidiomycota diversity. (A) *Puccinia iridis*, (B) *Pheogena*, (C) *Coleosporium*, (D) yeast, (E) *Ustilago*, (F) *Exobasidium*, (G) *Monliella*, (H) *Wallemia*, (I) *Phallus*, (J) *Clavaria*, (K) *Amylostereum*, (L) *Clavariadelphis*, (M) *Pycnoporus*, (N) *Russula*, (O) *Boletus*, (P) *Lycoperdon*.

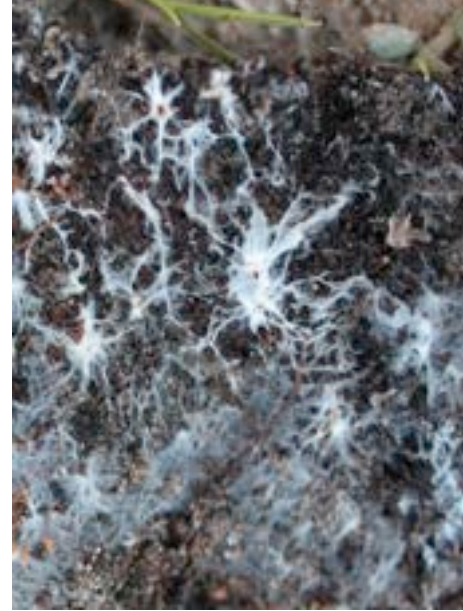
Fungal ecology



White-rot



Brown-rot



Litter-decayers



Mycorrhizal

Fungi play key roles in forest ecosystems



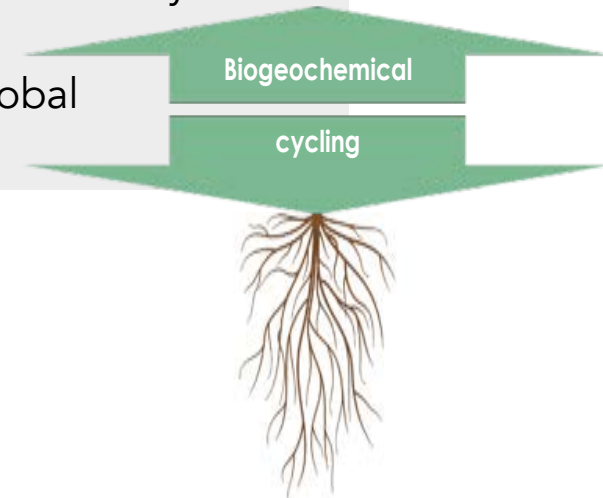
White-rot

Brown-rot

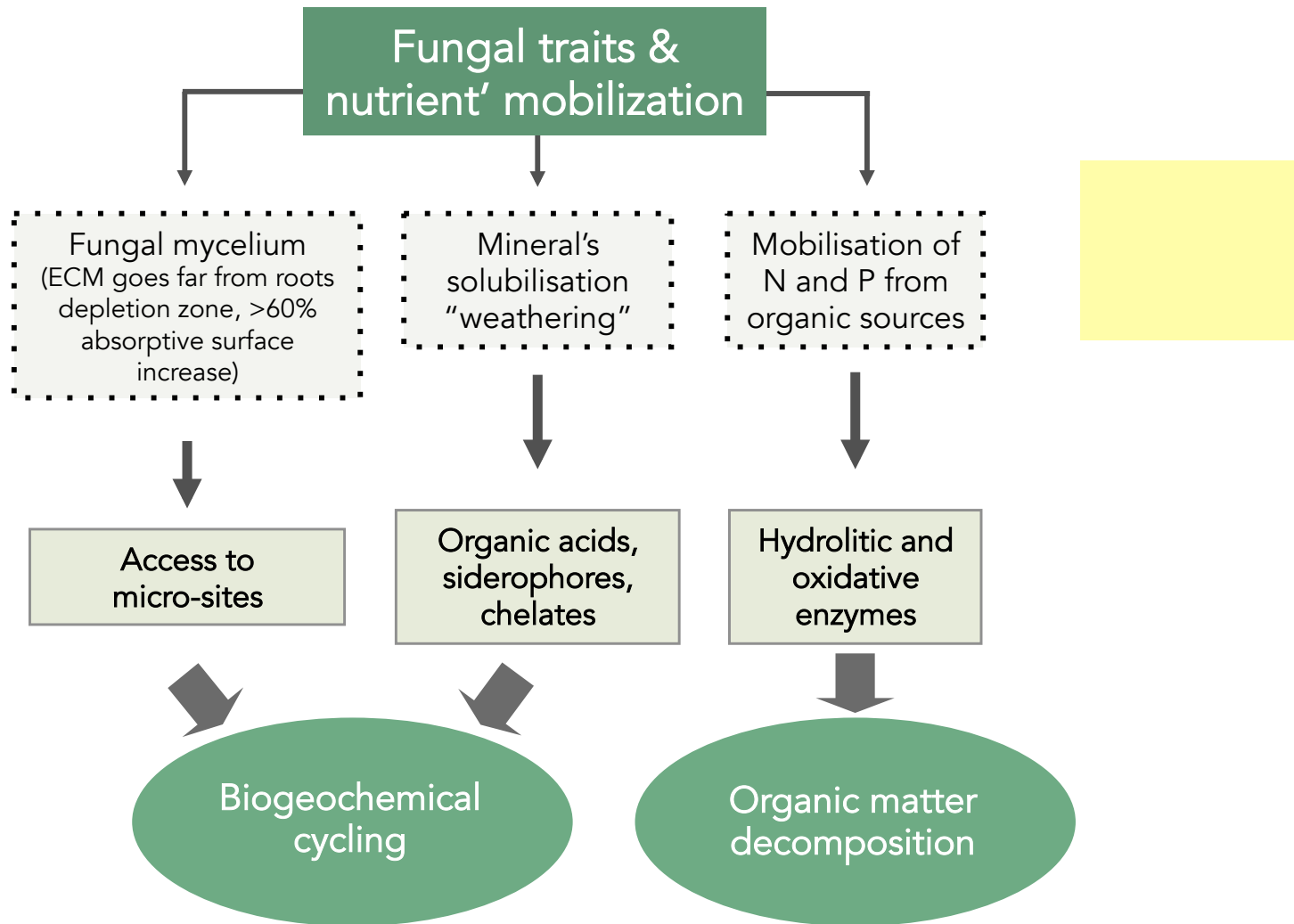
Litter-decayers

Mycorrhizal

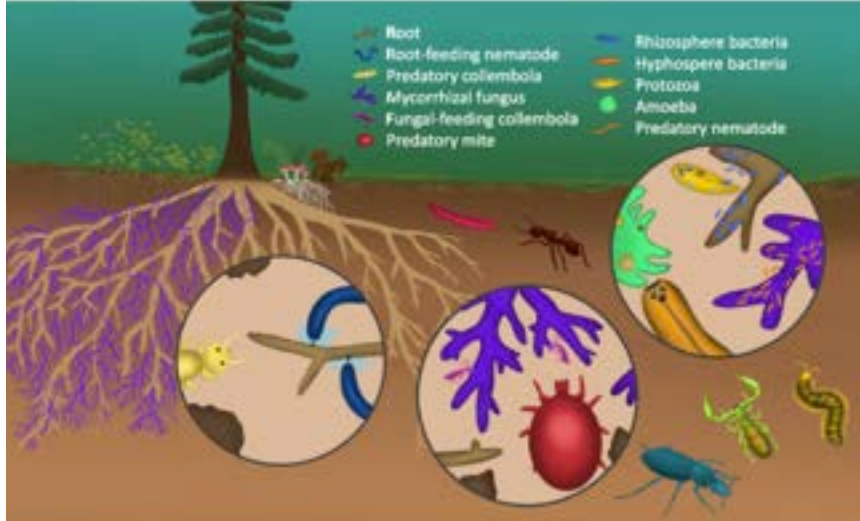
- Soil stabilization → \approx 60-90 % soil microbial biomass in forests
- Trophic base of food webs → relationships with other organisms
- Decomposition and mineralization → extracellular enzymes
- Mutualistic associations → MYC-nutrient acquisition strategies are key functional traits of plants with strong effects on biogeochemical cycles
- Effects from individual plants to ecosystem level, with global consequences → productivity of forest ecosystems



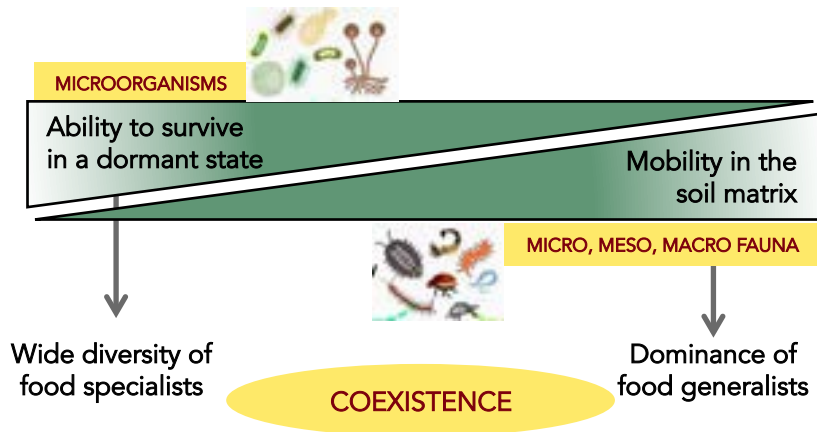
Fungal traits & soil carbon and nutrient cycling



The soil-fungal interface → intra- and inter-kingdom interactions



From: Prescott & Grayston (2023) For Ecol Man 532

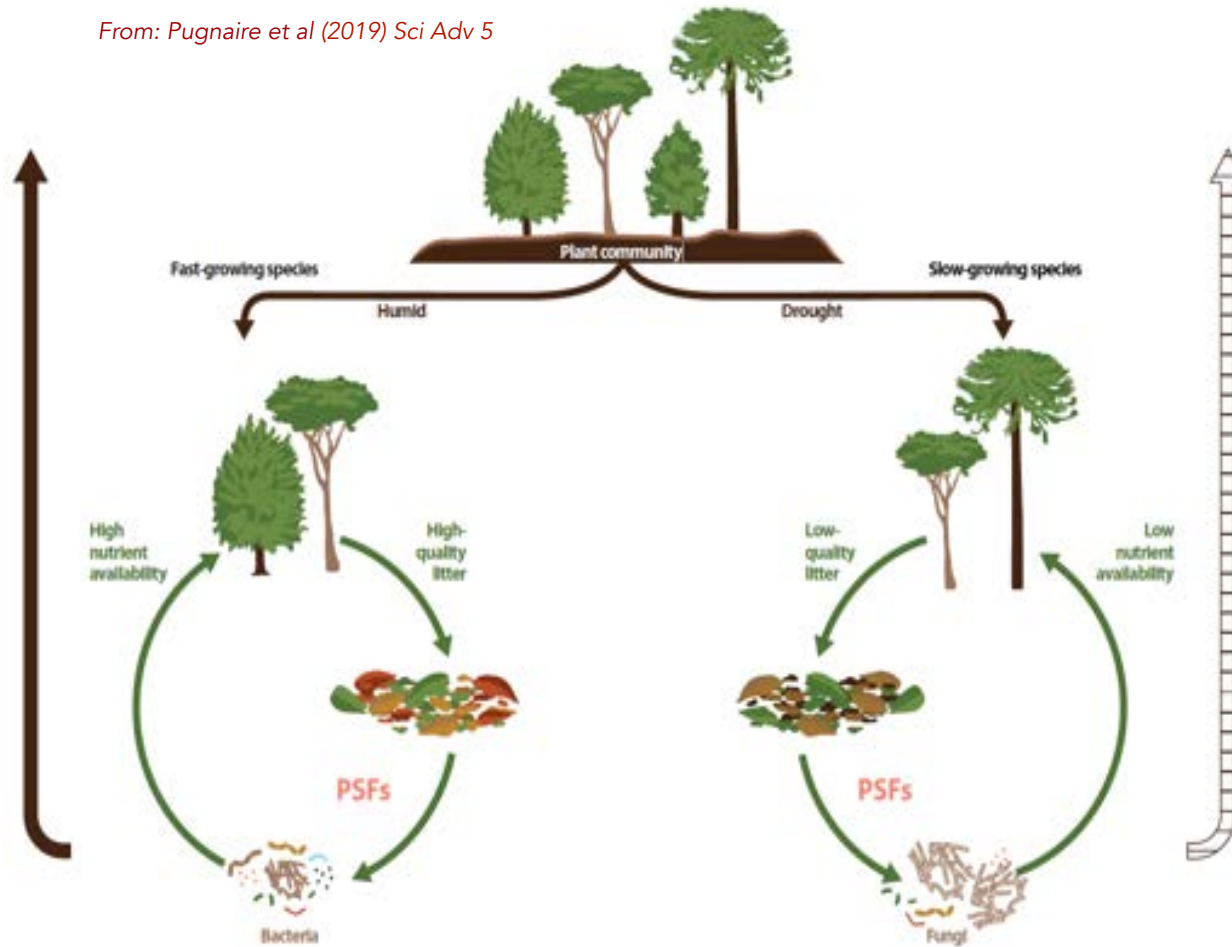


- Soil physical structure (pore size, connectivity, water and nutrient distribution) drives sensing and access to resources, via providing refuge and limiting mobility of organisms in the soil matrix
- Trophic interactions contribute to the formation of soil physical structure via relocating and mixing mineral and organic compounds
- Restrictions of interactions between consumers and preys allows to the coexistence in soil of hyper-diverse microbial communities (fungi, bacteria) and fauna
- Primary producers at the top of forest soil food-webs

Plant-soil feedbacks in forest ecosystems

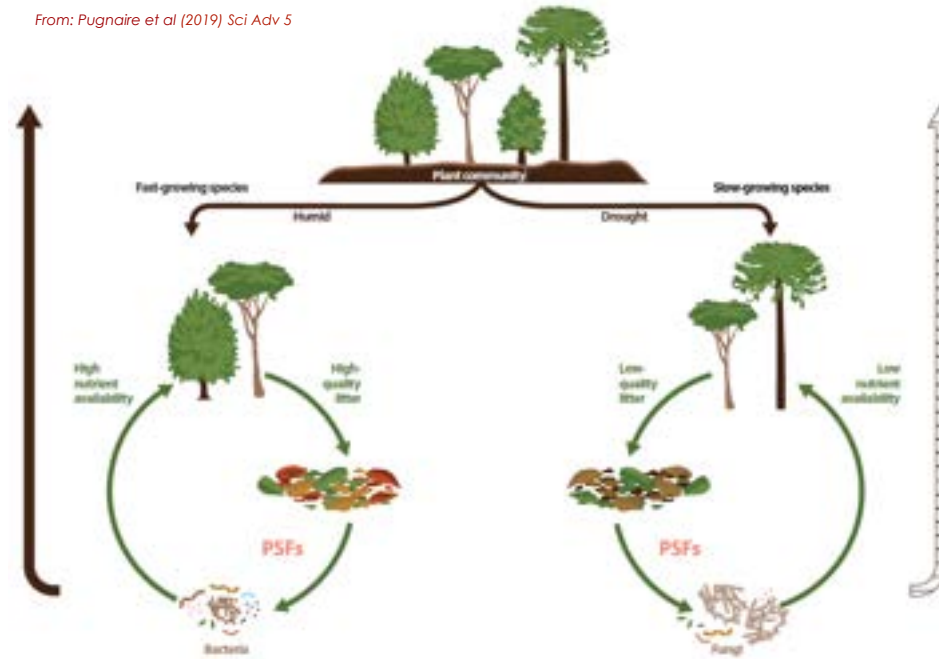
- The quality of plant inputs in soil → a main driver of plant-soil-microbial feedbacks

From: Pugnaire et al (2019) *Sci Adv* 5



Mycorrhizal associated nutrient economy (MANE)

From: Pugnaire et al (2019) Sci Adv 5



AMF
dominated

MANE

ECM
dominated

Philips et al. (2013) New Phytol 199

- P- limiting habitats
- Faster nutrient cycling
- Greater saprotrophs and bacteria activities
- Higher N leaching and nitrification processes
- Biological weathering attributable to bacteria
- Priming effects on bacteria

- N- limiting habitats
- Slower nutrient cycling
- Greater SAP and ECM/ErM fungal activity
- More efficient in fungal mineral weathering (organic acids, quelators)
- Higher priming effects

- ECM dominated habitats can bypass the nutrient mineralization loop for nutrient uptake

Plant Traits and Phylogeny Predict Soil Carbon and Nutrient Cycling in Mediterranean Pine-Quercus Mixed Forests

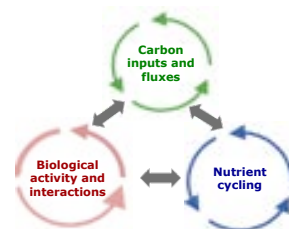


Plant community (32 spp) → AM-type (75%), ECM (25 %)

Plant community phylogenetic diversity

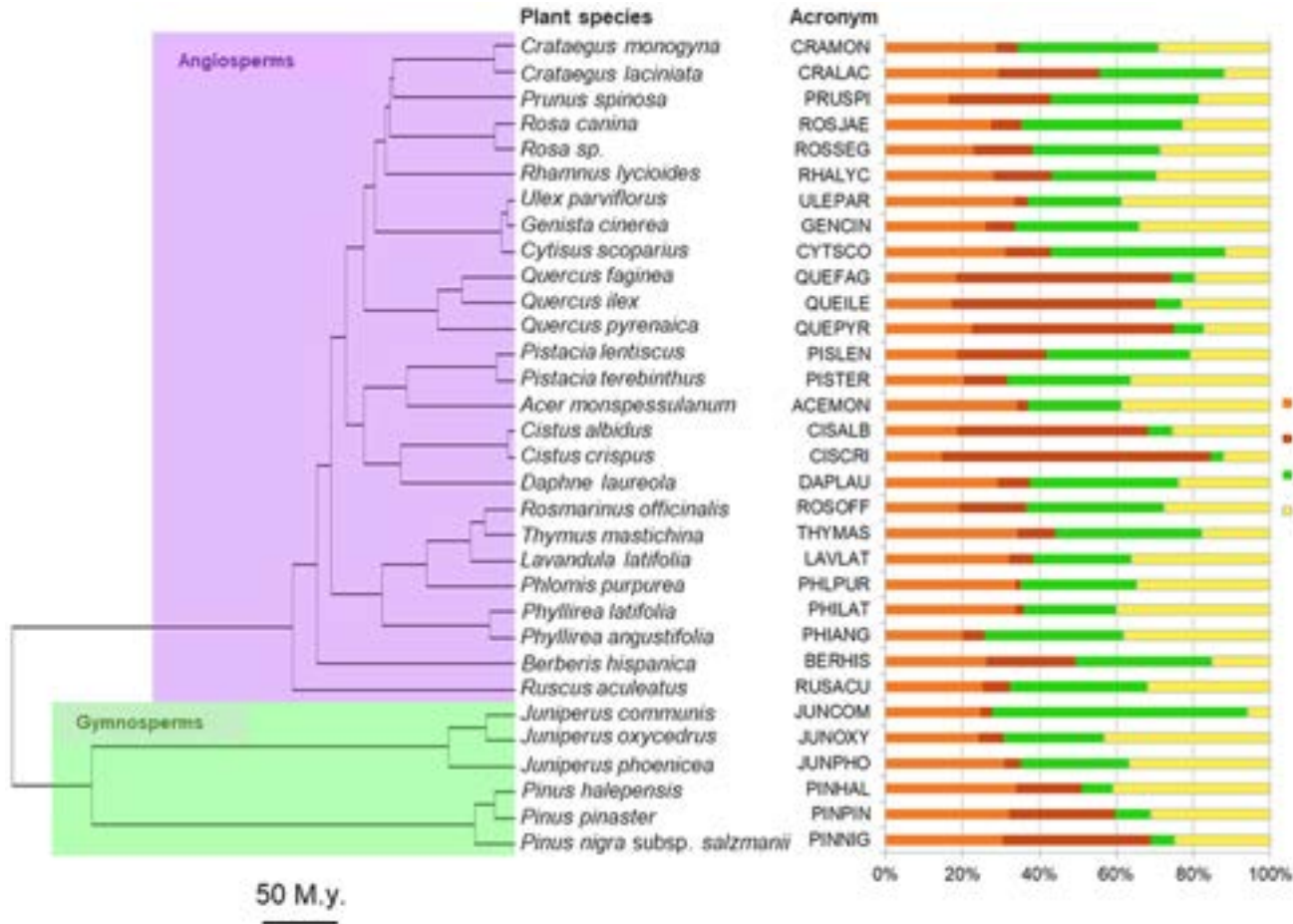
Plant morpho-functional trait database

Enzymatic activities of rhizospheric soils



Prieto-Rubio et al (2023) *Ecosystems* 26

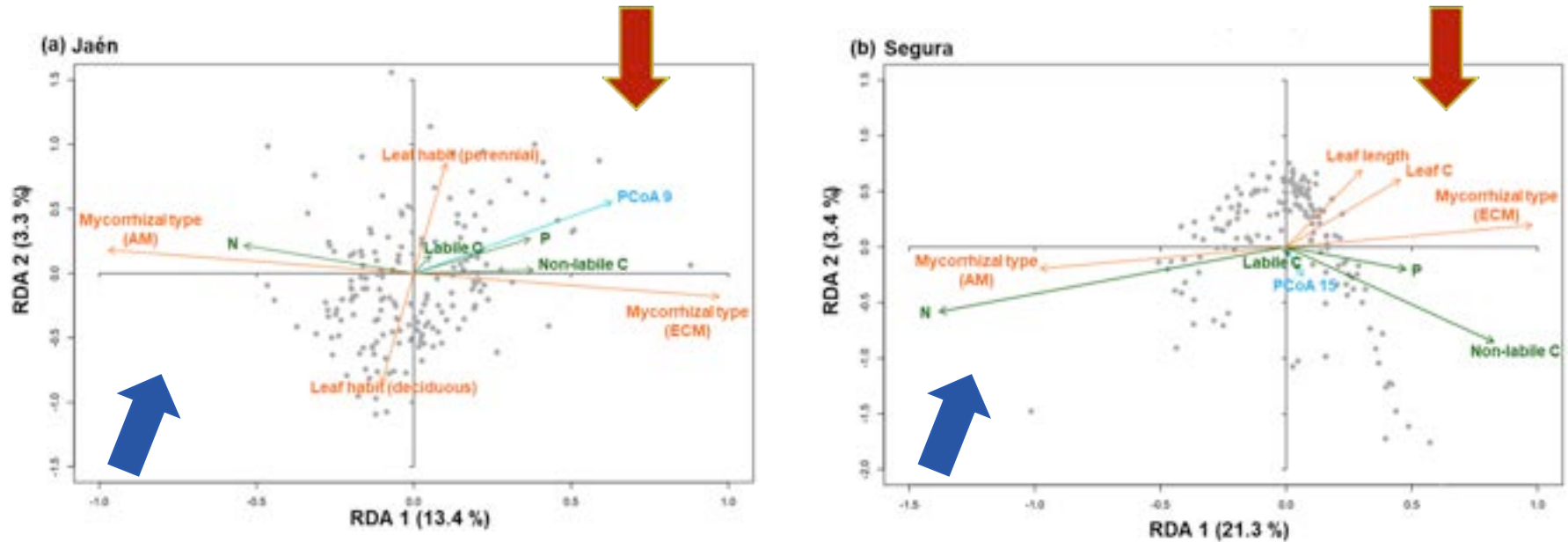
Plant Traits and Phylogeny Predict Soil Carbon and Nutrient Cycling in Mediterranean Pine-*Quercus* Mixed Forests



Significant phylogenetic signal of the plant community on soil functioning (i.e., enzymatic activities related with C and nutrient cycling)



Plant Traits and Phylogeny Predict Soil Carbon and Nutrient Cycling in Mediterranean Mixed Forests

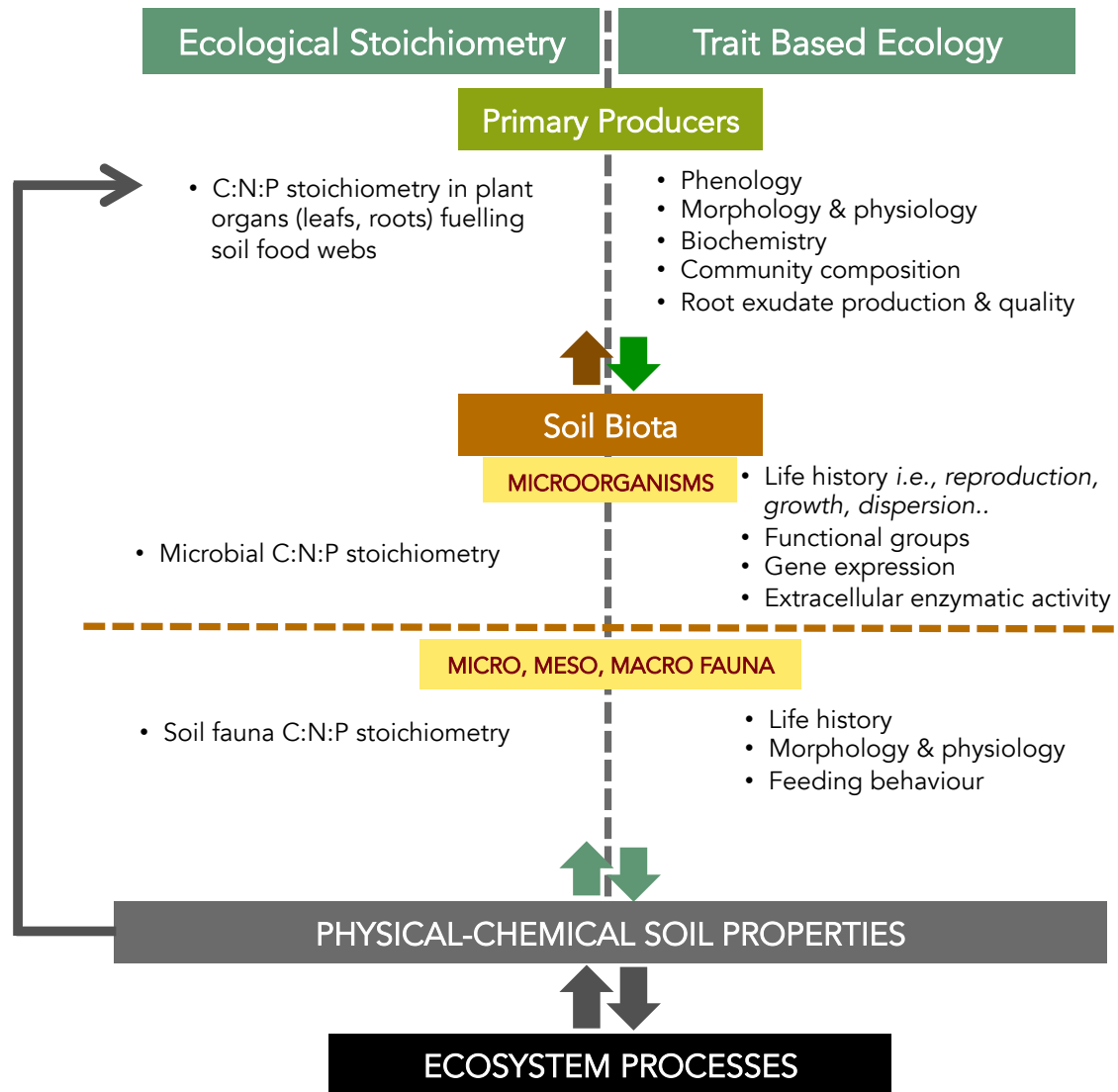


AM-type & N cycling ↔ ECM-type
 Leaf length, leaf C, leaf habit & Non-labile C

Plant community structure, morpho-functional leaf traits and MYC-type are main drivers of soil C and nutrient cycling in Mediterranean mixed forests

Mycorrhizal associated nutrient economy (MANE)

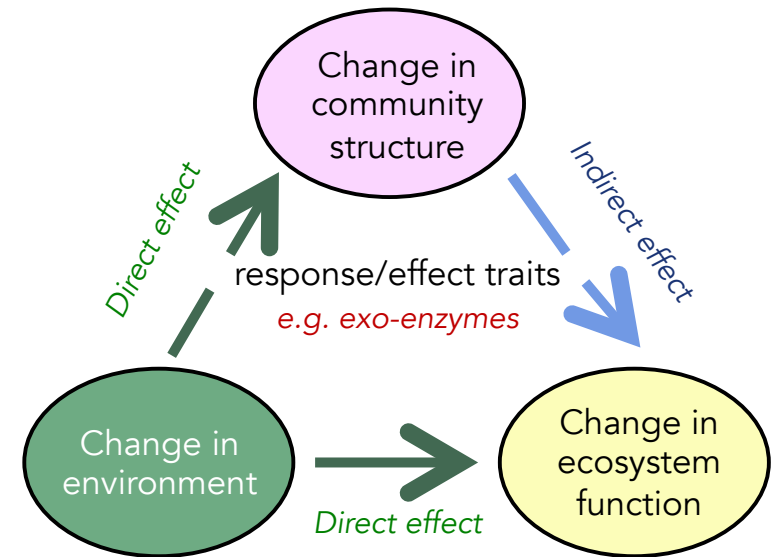
Environmental and ecological interactions in forest ecosystems



From: Maaroufi et al (2020) *Front Glob Chang*

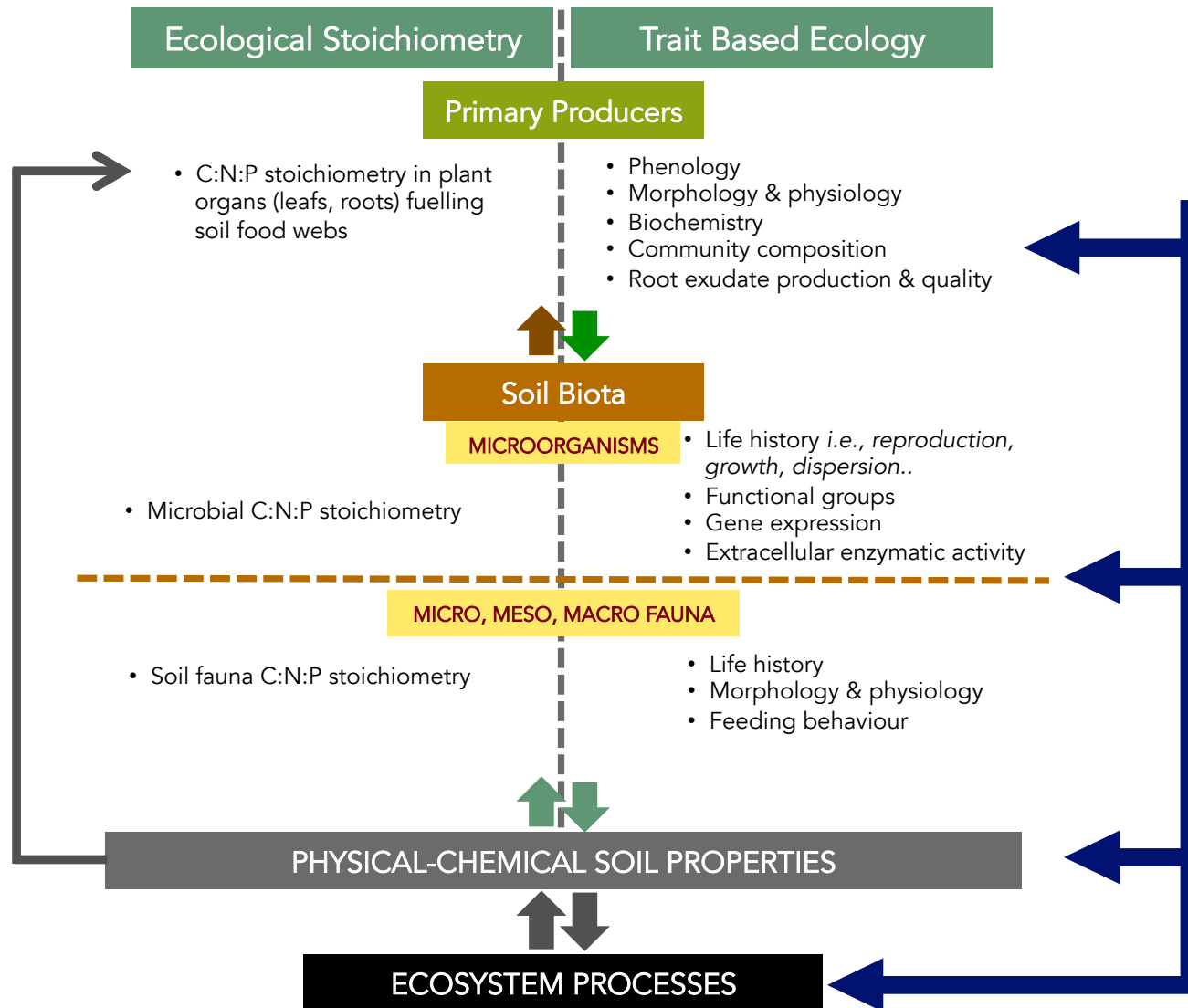
Soil fungal communities → linkages between biodiversity and functioning

- Community ecology goal → predict environmental change effects on ecosystem function
- Hyper-diverse soil fungal communities (taxonomically / phylogenetically / functionally)
- Links between biodiversity and functioning may help to predict ecosystem's response to environmental variations
- Resilience and recovery of forest ecosystems




Koide et al. (2014) *New Phytologist*, 201


Environmental and ecological interactions in forest ecosystems




Forest soil fungi: environmental and ecological interactions



Influence of the plant genotype / phenotype and the plant community on the structure and functioning of soil microbial communities



Relationships between edaphic-climatic conditions and soil fungal communities or individual taxa

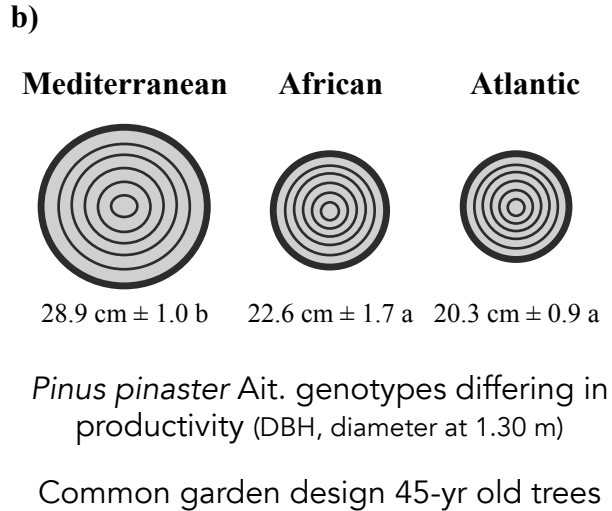


Structure and functional responses of fungal communities to disturbance

Plant intraspecific variation modulates nutrient cycling through its below-ground microbiome

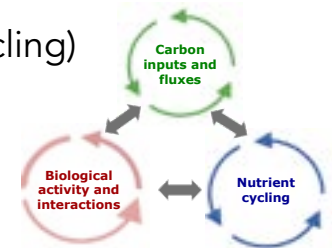


n = 108 trees



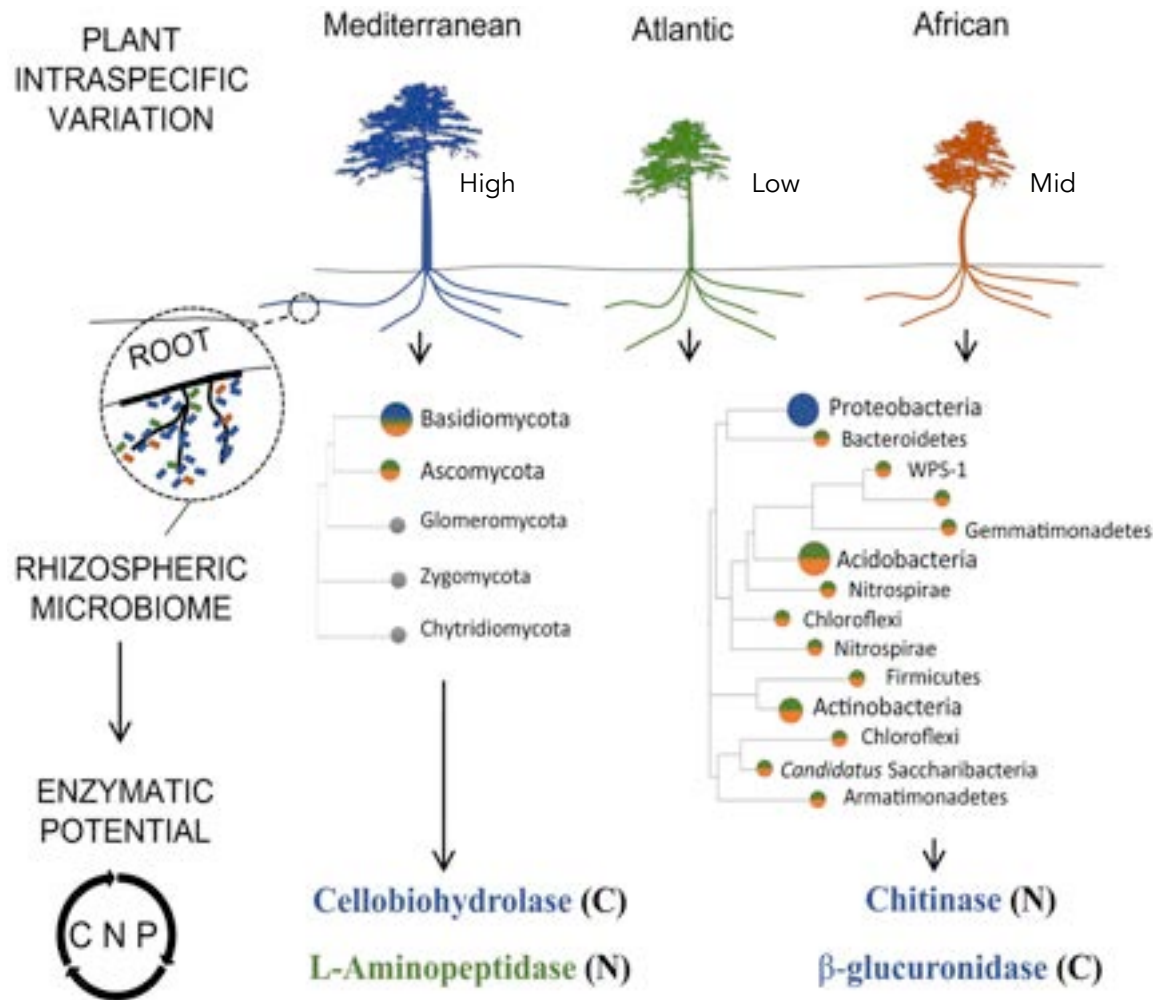
Microbial communities (bulk soil and root-tip) (NGS)

Enzymatic activities (C & nutrient cycling)



Pérez-Izquierdo et al (2017) *Env Microbiol*, 19:1639
Pérez-Izquierdo et al (2019) *J Ecol*, 107:1594

Plant intraspecific variation modulates nutrient cycling through its below-ground rhizospheric microbiome



- Atlantic – low productivity
- Mediterranean – high
- African – mid

Prevalence of Basidiomycetes and Proteobacteria at the rhizosphere of the most productive tree genotype

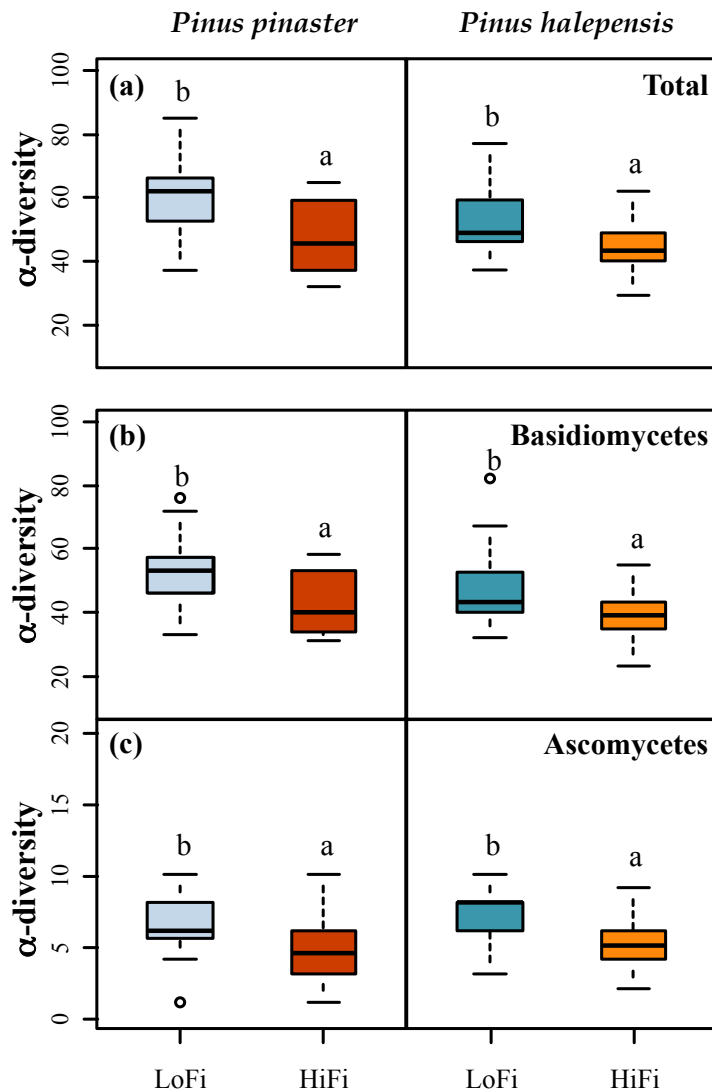
Plant intra-specific variation has a key ecological relevance in modulating the microbial controls of C and nutrient cycling

Ectomycorrhizal fungal diversity decreases in Mediterranean pine forests adapted to recurrent fires



Fig 1. (A) Closed cones of serotinous *Pinus halepensis* Mill. trees. (B) Wildfire in a *Pinus pinaster* Ait. forest in Eastern Spain.

Ectomycorrhizal fungal diversity decreases in Mediterranean pine forests adapted to recurrent fires

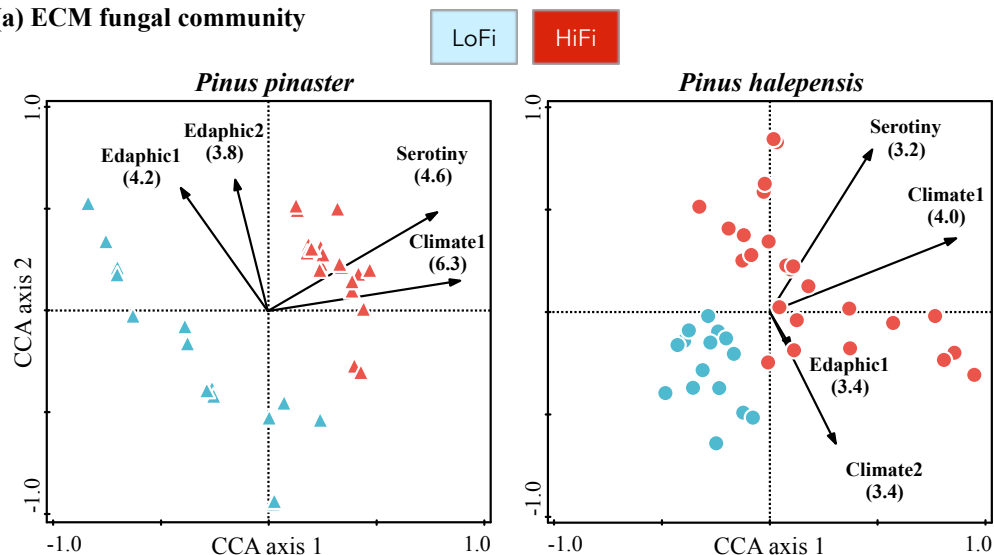


Root-tip ECM diversity declined in pine populations subjected to high fire recurrence, pointing to the community simplification (with prevalence of fire-prone fungi), due to the high selective pressure of fire (i.e., edaphic transformations, drier environment..)

Main drivers of root-tip ectomycorrhizal community structure and function in Mediterranean pine forests adapted to recurrent fires



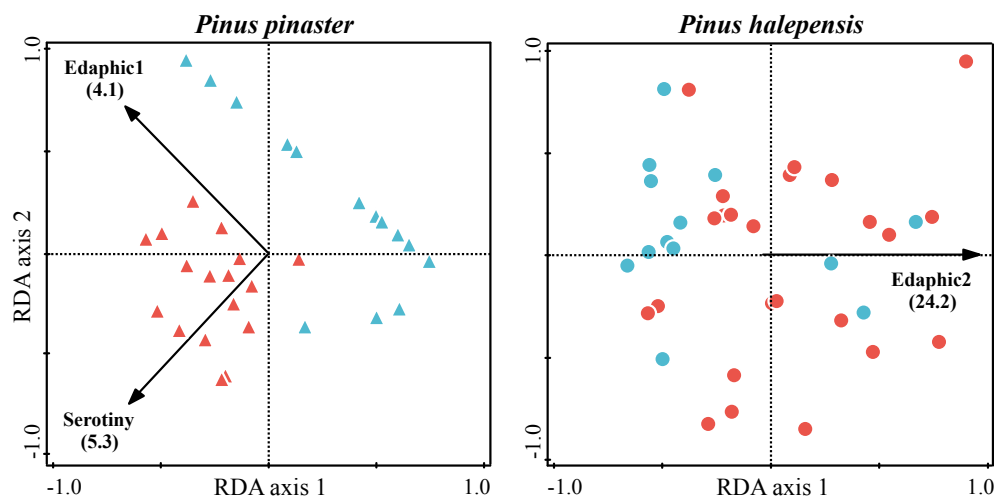
(a) ECM fungal community



Climate and edaphic factors were major drivers of ECM fungal community structure and functioning in the later case

Serotiny (i.e. tree fire-adaptation trait, did significantly explain the structure and, for *P. pinaster* also the functioning, of the root-tip ECM fungal community

(b) Enzymatic activity



Knowing the structure and function of ECM fungal communities of fire-prone pine species may have important implications for the dynamics and resilience of these Mediterranean ecosystems

Forest soil fungi: environmental and ecological interactions

Evaluation of microbial biodiversity-based strategies for soil restoration and remediation assessment

Response to stress (fire, soil degradation)
MIDIVERSOIL



Ecology & applications in truficulture
TUBERLINKS



Deciphering biodiversity and soil-plant feedbacks in the truffle environment for fitting best management practices to optimise the productions of truffle systems



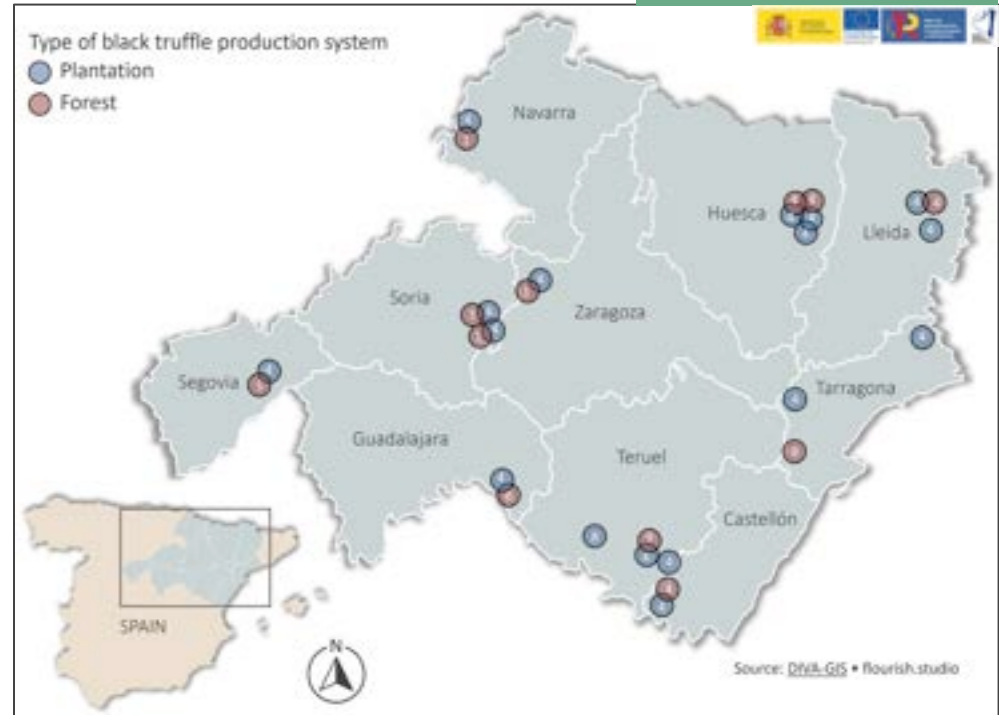
TUBERSYSTEMS and TUBERLINKS Projects



Tuber melanosporum VITTAD.

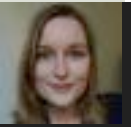


<https://www.tubersystems.com>



- Ascomycete ECM fungus + *Q. ilex*
- Highly appreciated edible fungi
- Mediterranean distribution (Spain is a main producer)
- Types of production: wild and plantations (125 trees)

Modelling environmental drivers of *Tuber melanosporum* mycelium in productive holm oak plantations and forests



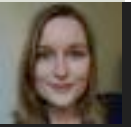
Compared with wild-producing systems, we expected:

- i) higher Tmel mycelium in plantations and less dependency of climate due to the managed-controlled environmental conditions
- ii) Truffle mycelium biomass will respond to different environmental abiotic variables in forests vs plantations

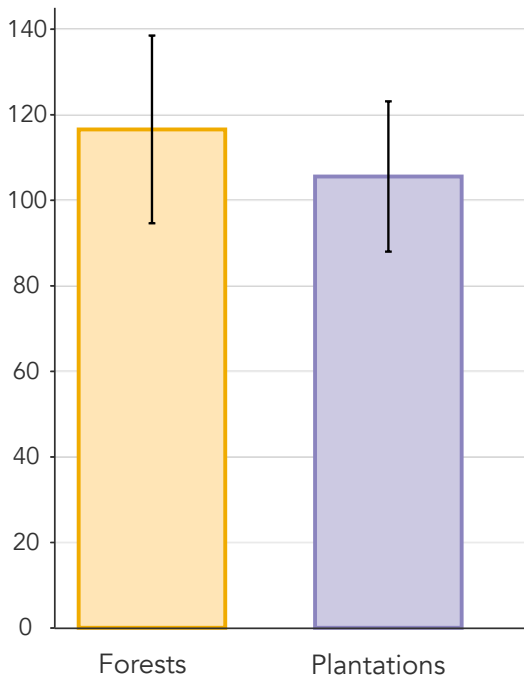
Generalized Additive Modelling (GAM) (≈ 30 abiotic variables)

Stepwise selection of best-fitted GAMs based on AIC

Modelling environmental drivers of *Tuber melanosporum* mycelium in productive holm oak plantations and forests




Mean \pm SE of *Tuber melanosporum* soil mycelium biomass ($\mu\text{g g}^{-1}$)



- No significant differences due to the type of truffle productive system (wild vs plantations)
- Variation in truffle mycelium biomass explained up to the 70 % in plantations and to the 50 % in forests, by the edaphic-climatic variables tested

Some Challenges in Soil Forest Fungal Research

- 
- Functional potential → comparison of genomes of distantly related taxa. Transcriptomics and proteomics approaches are needed to link genes to biological processes (SOM decay, nutrient mobilization, symbiosis establishment) *Lebreton et al. 2021. Annu. Rev. Ecol. Evol. Syst.*
 - Links diversity-function → understand the functional consequences of inter- and intra-specific fungal variability, and the patterns and drivers of fungal (hyper-diverse) communities assembling *Mony et al 2021. Front Ecol Evol*
 - Fungal traits → e.g. the extent of hyphae belowground (alive & necromas) is a challenging but important step for understanding fungal impacts on soil C fluxes (e.g. MYC nets may represent ~50 % of the total soil microbial biomass) *Wurzburger & Clemmensen 2017. J Ecol*

Some Challenges in Soil Forest Fungal Research



- Models → e.g. MYC are a key entry point of C into soil food webs (estimates of 3.58 Gt C or 13.12 Gt CO₂e), but we lack a robust quantitative and mechanistic understanding of the contribution of MYC associations to the global C cycle *Hawkins et al. 2023. Current Biol*
- From basic to applied research → Need of knowledge transference of science-based solutions for forest and agro-forest management and productions optimization (e.g. forest conservation, climate change mitigation, truficulture)

Thank you for
your attention!!

Collaborators

- Concepción Azcón, Álvaro García, José L. Garrido (EEZ-CSIC)
- Marc Buée (INRA-France)
- Marta Goberna (INIA-CSIC)
- Miguel Verdú, Jorge Prieto (CIDE-CSIC)
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