





#### Common mycorrhizal networks of European Beech trees drive belowground allocation and distribution of plant-derived C in soil

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# Common mycorrhizal networks (CMNs)

Mycelial connection between several co-existing plant individuals, even from different species

- Pathogen resistance
- Establishment of seedlings
- Amplification or alleviation of nutrient competition
- Impacts on plant community composition



Van der Heijden M (2016) Underground networking. Science (80- ) 352:288–290. https://doi.org/10.1126/science.aaf3354 (1) Is the total belowground C allocation of plant photosynthates influenced by the size of the mycorrhizal network and its access to resources?

(2) Is the belowground C distribution within a CMN altered if trees have unequal access to C from photosynthesis?

(3) Do CMNs amplify or alleviate competition for nutrients between connected trees?



#### Experimental setup



Without CMN

With CMN

(60 plants -> 4 treatments x 6 replicates + 2 treatments x 3 replicates)

6 cm

3 cm

# Plants relied mostly on their mycorrhizal partners to acquire nitrogen



#### Two-pool mixing model:

63% of plant <sup>15</sup>N derived from hyphalexclusive peat bags while only 37% originated from the soil

No clear effect caused by shading nor extended mycorrhizal network on N uptake

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### CMN increased total assimilation and belowground transfer of C



Differences in pools were analyzed with ANOVA. n = 6 for shaded and n=9 for not shaded treatments. Error bars represent standard error.

(ng excess <sup>13</sup> C)	Donor	Receiver
+CMN	1885	31
-CMN	956	21

(ng excess <sup>13</sup> C/g DW)	Donor	Receiver
Peat bags	15.06	5.08
Soil	8.63	1.25

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



Soil: Higher microbial biomass in soil than other pools, however, receivers were **not** significantly enriched in <sup>13</sup>C

Peat: High <sup>13</sup>C within microbes from **donor and receiver** peat bags, especially **fungal** markers

Sand: Disrupted CMN significantly decreased abundance of fungal markers. CMN affected bacterial and fungal <sup>13</sup>C enrichment

Differences in pools were analyzed with Kruskal-Wallis rank sum test. Significant tests (p < 0.05) were followed by Wilcoxon signed-rank's post-hoc test of multiple comparisons with Benjamini-Hochberg correction (adj. p < 0.05); n = 15. Error bars represent standard error.

# Summary

# (1) Is the total belowground C allocation of plant photosynthates influenced by the size of the mycorrhizal network and its access to resources?

- Yes: presence a larger mycorrhizal network connecting to another plant and an additional N source almost doubled photosynthetic CO<sub>2</sub> assimilation and belowground C allocation by plants
- However, <sup>13</sup>C was similarly transferred in the intact and disrupted CMN treatments to the neighboring pots (despite the fact that disrupted CMN treatments showed a decline in fungal biomass in the sand).

# (2) Is the belowground C distribution within a CMN altered if trees have unequal access to C from photosynthesis?

- No: shading did not affect the belowground distribution of C.
- Plant C was preferentially transferred to mycorrhiza-exclusive N sources from the own and distant partner pot

(3) Do CMNs amplify or alleviate competition for nutrients between connected trees?

- No clear effects of belowground competition for N
- Plants relied mostly on the mycorrhizal fungi to acquire N

#### Conclusions

- Belowground ectomycorrhizal networks represent a significant sink strength for plant photosynthates and may thus be a major driver of C sequestration in beech forest soils.
- The belowground distribution of C via fungal networks is mainly related to the distribution of nutrient-rich patches in the soil and less to differences in the photosynthetic capacity of the host plants.

