

# Drivers of spatial variability of soil respiration along altitudinal gradient in Northwest Caucasus Mountains

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## I. MOTIVATION

Mountains cover almost 25% of the world's land surface, so their contribution to the global C cycle can be significant. However, the C fluxes, in particular soil respiration ( $R_s$ ), of these remote areas remain poorly understood. This study focused on estimating the spatial variability of  $R_s$  along the altitudinal gradient as a possible function of changes in vegetation and soil characteristics.

## 2. STUDY AREA

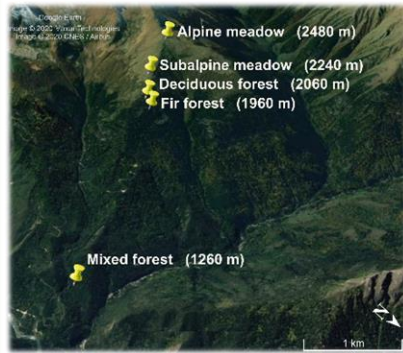
**Location:** Northwest Caucasus Mount.; Russia; 43°40'N / 40°47'E

**Slope:**

- northeastern exposure
- 1.2 km altitudinal gradient
- 5 vegetation belts (ecosystems)

**Climate:** MAT → from 3.5 °C to 5.9 °C  
MAP → from 800 to 1850 mm

**Soils:** Cambisols, Umbrisols, Leptosols

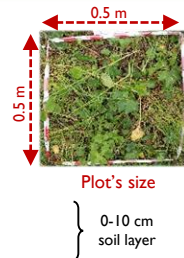


## 3. DESIGN and METHODS

Each ecosystem → 12 randomly distributed plots (totally n=60)

Simultaneously measurement on 11 August 2018:

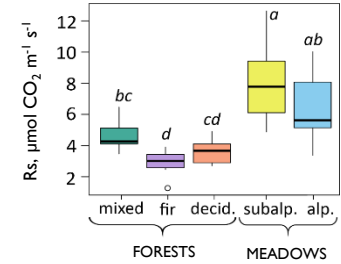
- $R_s$  with closed chamber technique (SBA-5, PP system, USA)
- Vegetation survey (projective cover, richness, Shannon-Wiener index, graminoids, forbs)
- Soil physicochemical (temperature, moisture, C and N contents, C:N ratio, pH)
- Soil microbial (biomass, activities of  $\beta$ -glucosidase, chitinase, leucine aminopeptidase)



## 4. RESULTS

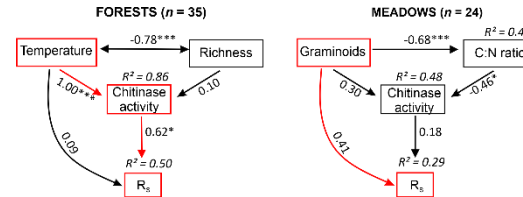
### $R_s$ rate for forests and meadows

- A higher spatial variability of  $R_s$  was found in meadows than in forests.
- On average,  $R_s$  was twice as high in meadows as in forests, reaching 7.3 and 3.7  $\mu\text{mol CO}_2 \text{ m}^{-2} \text{ s}^{-1}$ , respectively.



### Drivers of $R_s$ spatial variability

- (1) The best predictors based on stepwise regression analysis:  
forests → soil temperature, chitinase activity, species richness (29–50% explained variation)  
meadows → chitinase activity and graminoid abundance (19 and 27% explained variation)
- (2) Direct/indirect predictor effects tested by path analysis:

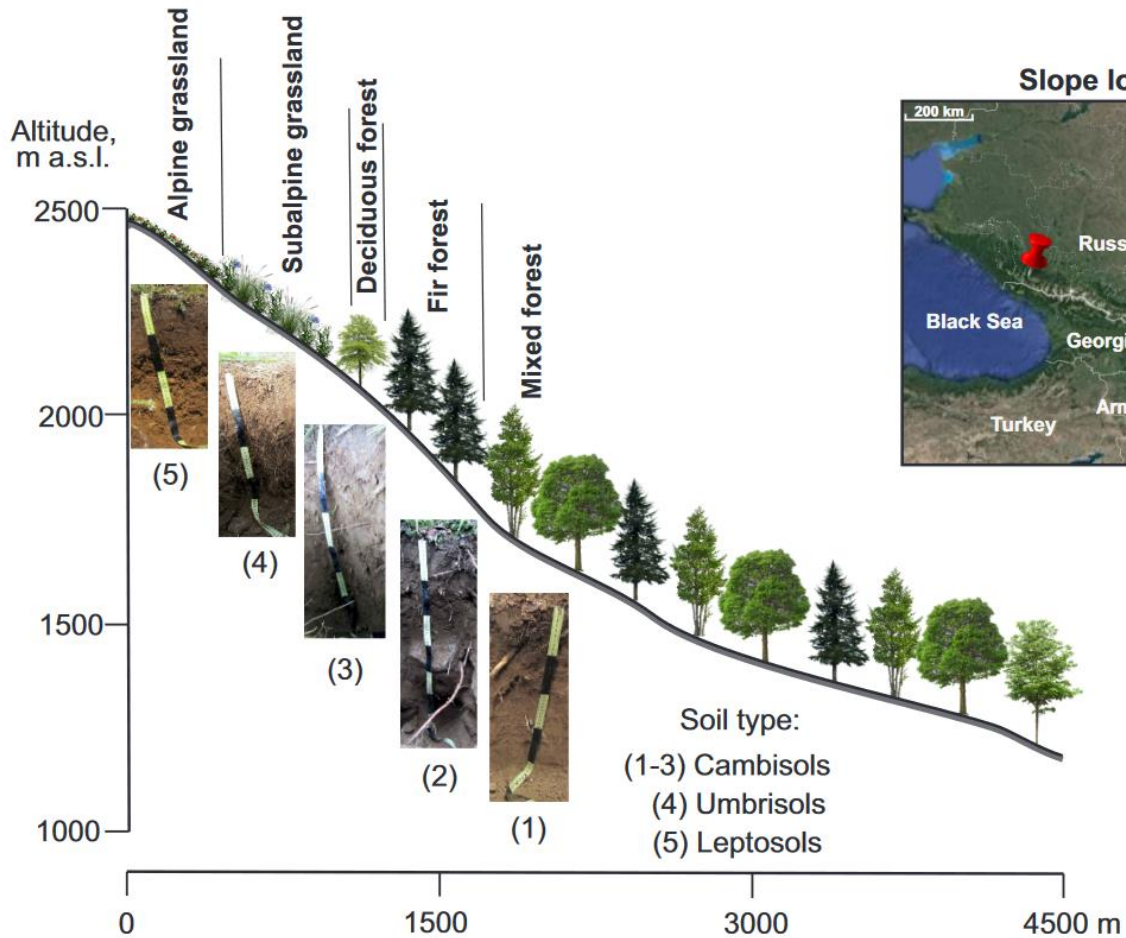


**forests** →  
soil temperature (indirect),  
chitinase (direct)

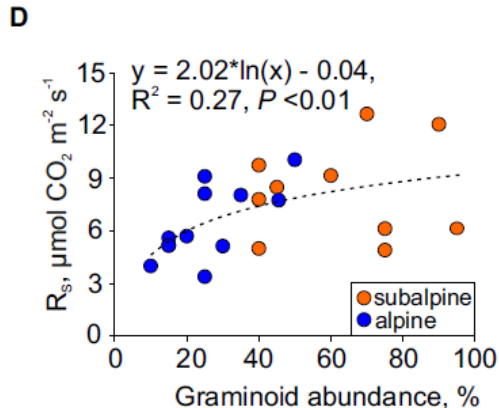
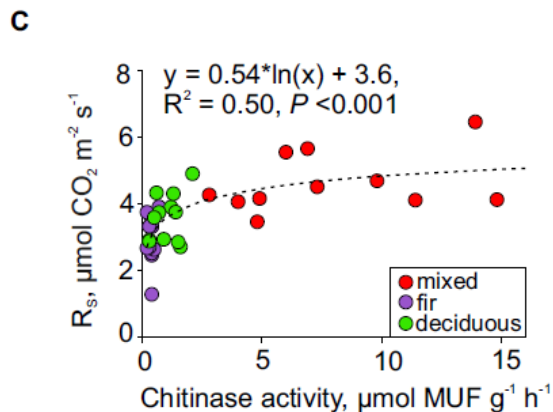
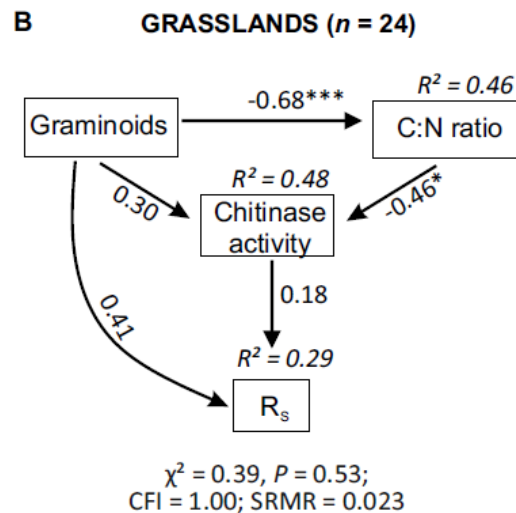
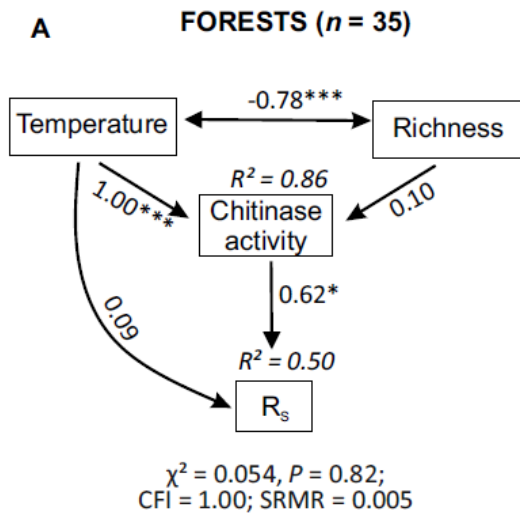
**meadows** →  
graminoid abundance (direct)

## 5. CONCLUSIONS

The results showed principal differences in the main drivers of  $R_s$  spatial variability in forests and meadows. The chitinase dependence of  $R_s$  in forests may be related to soil N limitation. This was confirmed by low N content and high C:N ratio compared to meadow soils. The greater sensitivity of meadow  $R_s$  to vegetation structure may be related to the essential root C allocation for some grasses. Thus, the crucial role of soil microbial activity in forests and vegetation structure in meadows as drivers of  $R_s$  spatial variability has been demonstrated.



**Fig. 1:** Scheme of the northeastern slope of Mt. Tkachiha (Northwest Caucasus, Karachay-Cherkess, Republic of Russia)



**Fig. 2:** Path models and simple regression revealed the effects of soil (temperature, chitinase activity, C:N ratio) and vegetation factors (species richness, graminoid abundance) on  $R_s$  spatial variability in mountain forests (A, C) and grasslands (B, D). In the path model, numbers within double-headed arrows are correlation coefficients between variables, numbers within one-way arrows are standardized path coefficients indicating the size effect of the causal relationship among variables (\* $P \leq 0.05$ ; \*\*0.01; \*\*\*0.001); CFI, comparative fit index; SRMR, standardized root mean square residual.