

Supplementary material

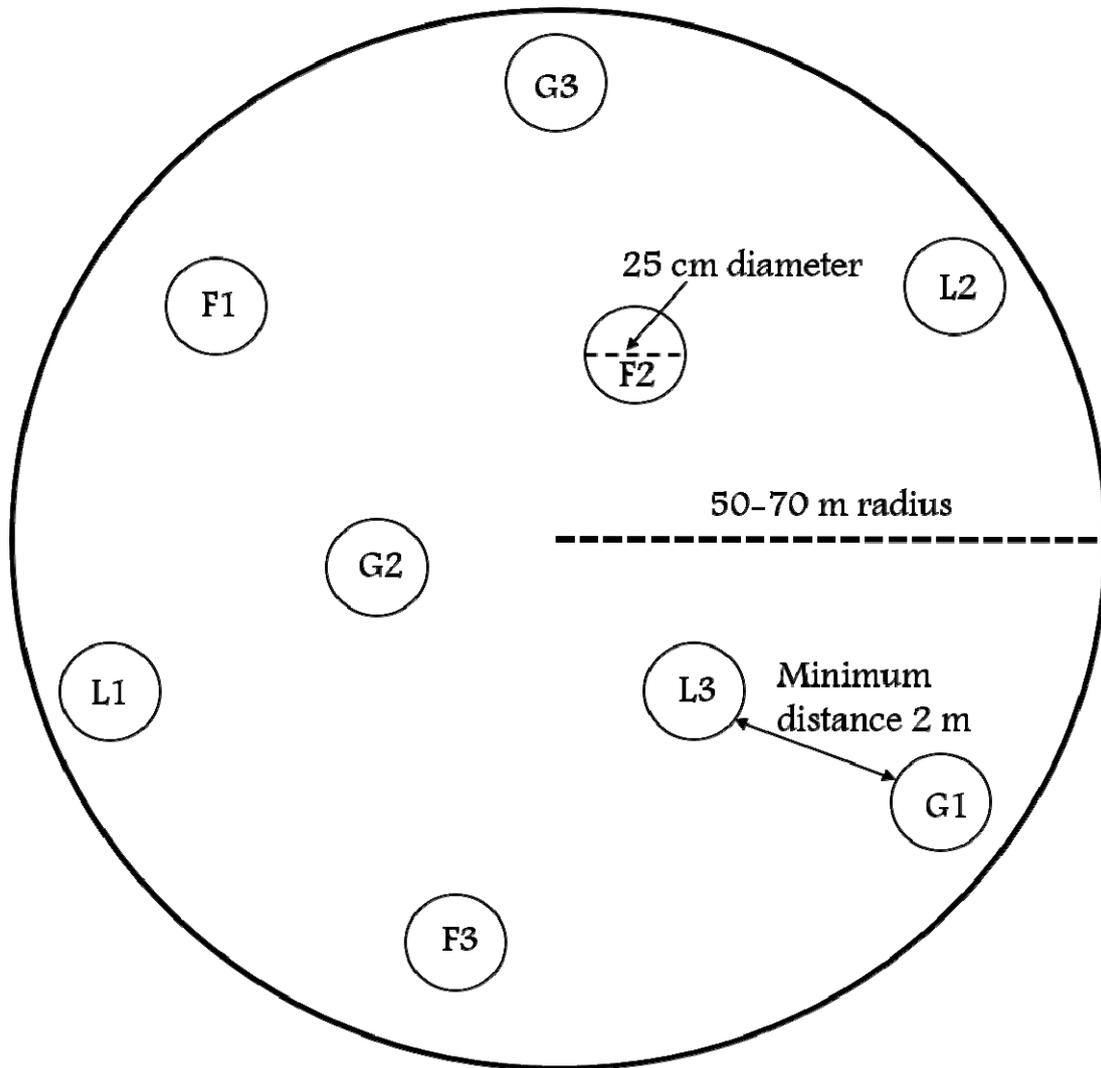


Figure S1. Schematic diagram depicting the sampling design and distribution of samples in each study site. Circles refer to the points (cylinders placed 3 cm deep into the ground: height = 8cm; diameter = 25 cm) from which vegetation and soil samples were collected. F, G and L refer to sampling points dominated by forbs, grasses and legumes, respectively. The numbers following the letters F, G and L refer to the number of replicate, with 3 replicates per PFT (total of 3 PFTs per site: F, G, L) and thus nine sampling points per study site.

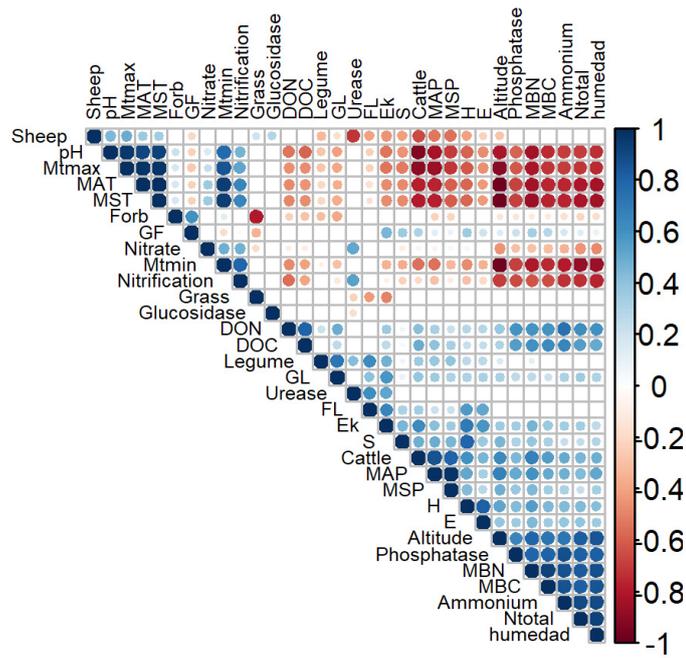


Figure S2. Correlation matrix of the variables used in the models. Positive correlations are displayed in blue and negative correlations in red color. Color intensity and the size of the circle are proportional to the correlation coefficients. In the right side of the correlogram, the legend color shows the correlation coefficients and the corresponding colors.

Table S1. Generalized diversity-interaction models for soil activity and fertility parameters (response variables). The explanatory variables include climatic variables: MTmax = mean annual maximum air temperature; MAP = mean annual precipitation; MTmin = mean annual minimum air temperature; MST = mean summer air temperature; management variables: Cattle and sheep grazers; and plant functional type diversity variables: G = Grass, L = Legume, F = Forb; FG = forb-grass, GL = grass-legume and FL = forb-legume pairwise interactions; Ek = evenness.

| Soil parameter | Model | Adjusted r ² | p-value |
|-------------------------|---|-------------------------|---------|
| | <i>Model 1. Only environmental variables included</i> | | |
| Nitrate | -2.2 + 1.4MTmax ^{***} + 3.0Cattle ^{**} | 0.38 | <0.001 |
| Ammonium | -1.18 - 1.0MTmax ^{***} - 0.2 MAP ⁺ + 1.5Sheep ^{***} + 0.7Cattle | 0.84 | <0.001 |
| DOC | -7.7 + 6.8MTmax ^{**} - 3.4 MTmin ^{***} + 0.9MAP ^{**} + 1Sheep ^{***} + 9.7Cattle ^{***} | 0.79 | <0.001 |
| MBC | -0.6 - 1.9MTmax ^{***} + 0.5MTmin [*] - 0.5MAP [*] + 1.4Sheep ^{***} | 0.75 | <0.001 |
| Glucosidase | -4.6 + 6.0MTmax [*] - 3.2MTmin + 0.8MAP ⁺ + 6.2Cattle ⁺ 0.6 + 0.7MTmax ^{***} + 0.1MAP - 0.8Cattle ^{***} | 0.17 | <0.05 |
| MBN | <i>Model 3. Environmental variables + PFT identity effects + PFT pairwise interactions</i> | | |
| DON | -0.6 + 0.5MTmin ^{***} - 1.9MTmax ^{***} - 0.4MAP ^{**} + 1.4Sheep ^{***} - 16.8 G - 16.5 F - 10.6 L + 0.2FG [*] - 0.1 GL + 0.0FL | 0.60 | <0.001 |
| Phosphatase | -74.1MTmin:G [*] - 73.1 MTmin:F ⁺ - 47.7MTmin:L -0.6 + 0.8MTmin ^{***} - 1.3 MST ^{**} + 1.4 Sheep ^{***} - 82.6 G - 81.3 F ⁺ - 53.1 L + 0.2FG + 0.4GL ^{**} + 0.2FL | 0.63 | <0.001 |
| | <i>Model 4. Environmental variables + PFT identity effects + evenness</i> | | |
| Total N | -0.4 - 0.6MTmin ^{***} + 0.9Sheep ^{**} - 90.7 G - 89.5 F ⁺ - 857.8 L + 0.5 FG ^{***} - 0.1 GL + 0.0FL | 0.84 | <0.001 |
| Urease | -0.3 - 0.4MTmin ^{***} - 0.4MST ^{**} + 0.78Sheep ^{***} - 11.6 G - 11.5 F - 7.5 L + 0.2Ek [*] | 0.79 | <0.001 |
| Nitrification Potential | -56.7 + 29.6MTmin ^{**} + 208.9MTmax ^{***} - 194.4MST ⁺ - 30.3Sheep ^{***} + 93.3Cattle ^{***} - 41.4G - 40.9F ^{**} + 26.4L + 0.33Ek ^{**} | 0.78 | <0.001 |