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Supplementary Material

Plant species, nitrogen status and endophytes are drivers of soil microbial communities in grasslands

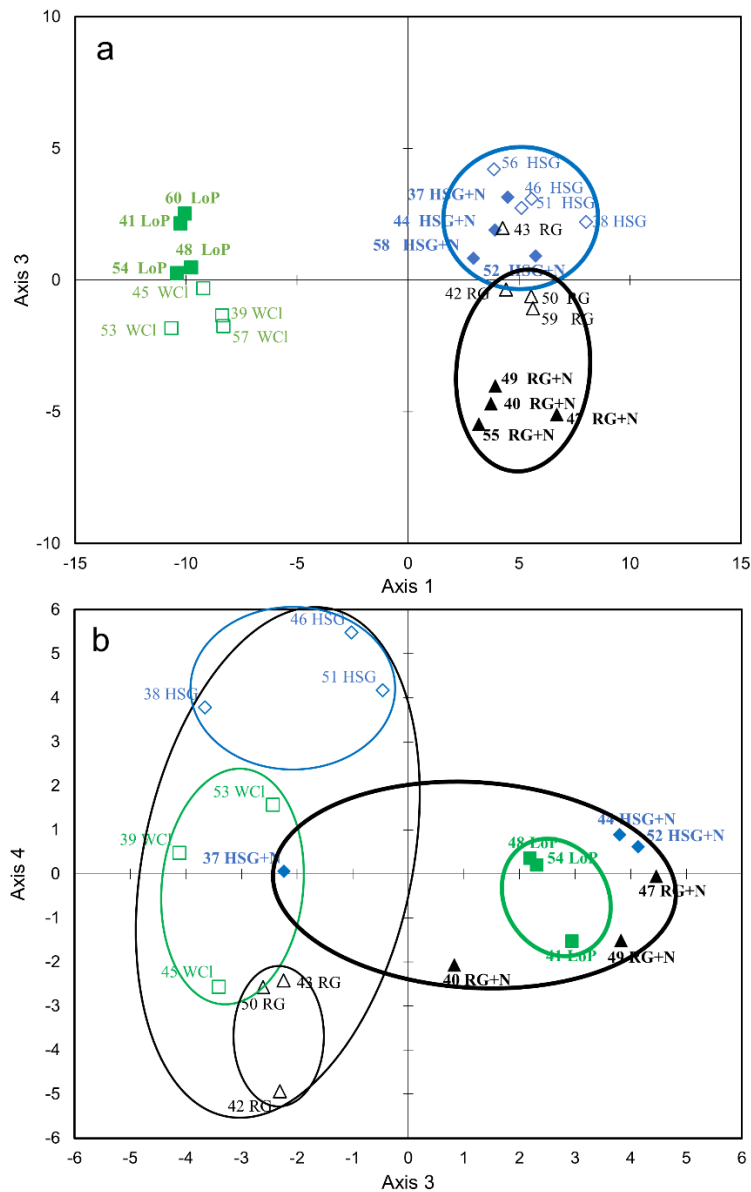
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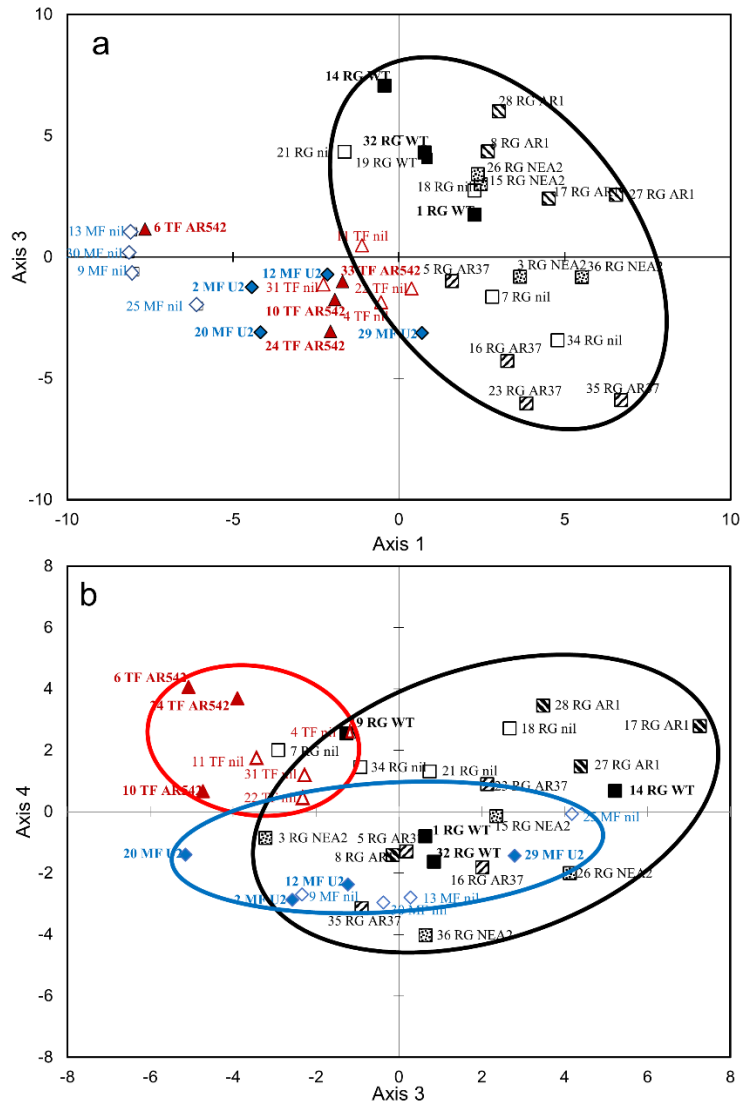
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Supplementary material Fig. S1. Partial Least Square Discrimination Analysis (PLS-DAnalysis) of impacts of plant species/ traits and N fertilization on the abundance of fungal (a) and bacterial (b) genera. Blue open diamonds: HSG (high sugar grass; *Lolium perenne* cv. AberDart), blue closed diamonds: HSG+N, black open triangles: RG (ryegrass; *L. perenne* cv. Impact), black closed triangles: RG+N, green open square: WCI (white clover; *Trifolium repens*), green closed squares: LoP (*Lotus pedunculatus*). Abundancies were box-cox transformed prior to PLS-DAnalysis. Groups showing major spatial separations were confined within ellipses.



Supplementary material Fig. S2. PLS-D analysis of impacts of plant extended phenotype (endophyte) and grass species on the abundance of fungal (a) and bacterial (b) genera. Black open squares: RG_nil (ryegrass without endophyte), black closed squares: RG_WT (ryegrass infected with *Epichloë festucae* var. *lolii* wild type endophyte), black downward striped squares: RG_AR1 (ryegrass infected with *E. festucae* var. *lolii* AR1 endophyte), black upward striped squares: RG_AR37 (ryegrass infected with *Epichloë* sp. LpTG-3 strain AR37 endophyte), black dotted squares: RG_NEA2 (ryegrass infected with *E. festucae* var. *lolii* NEA2 endophyte), blue open diamonds: MF_nil (meadow fescue), blue closed diamonds: MF_U2 (meadow fescue infected with *Epichloë uncinata* U2 endophyte), red open triangles: TF_nil (tall fescue infected

with *Epichloë coenophiala* AR542 endophyte), red closed triangle: TF_AR542. Groups showing major spatial separations were confined within ellipses.