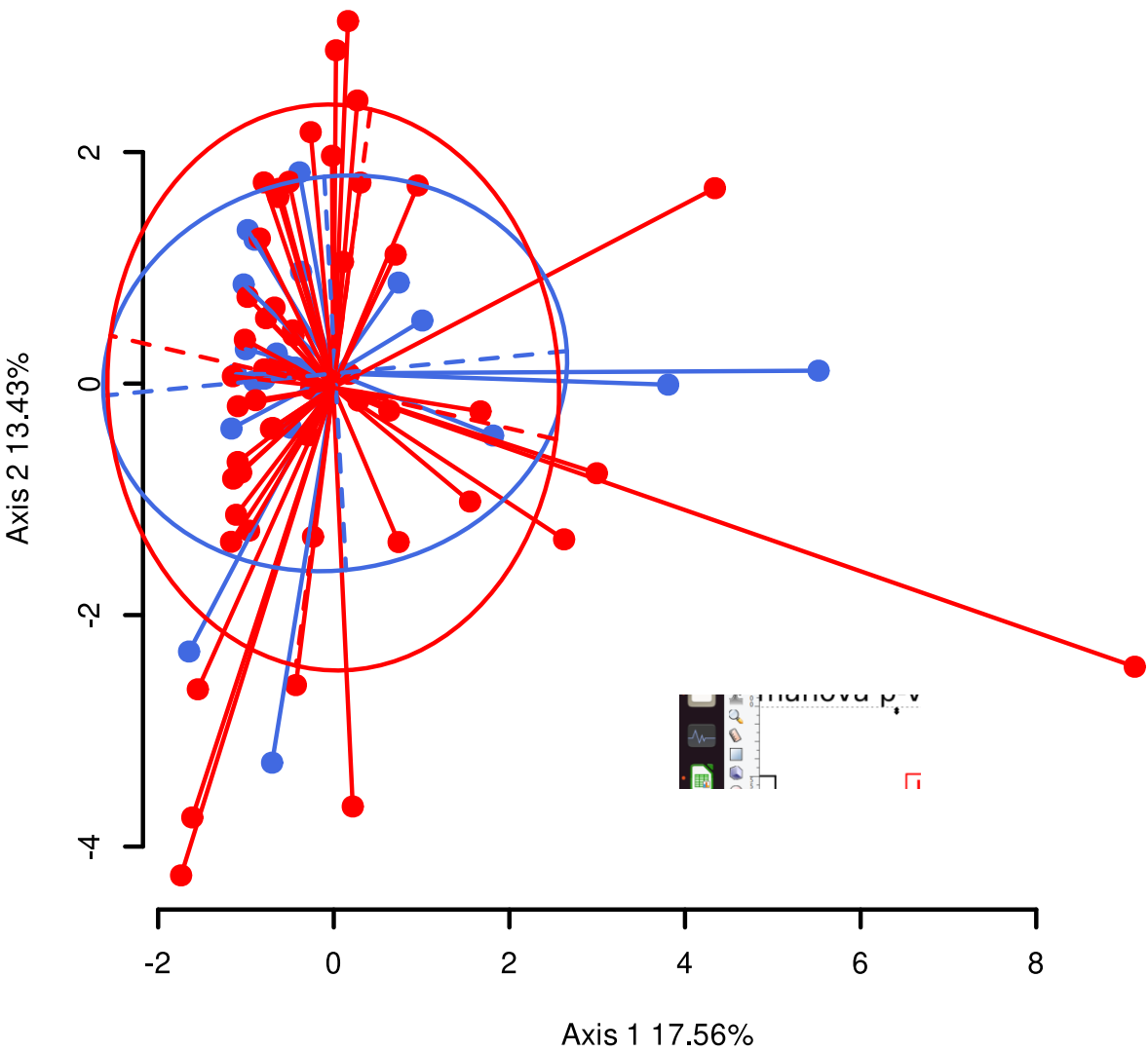
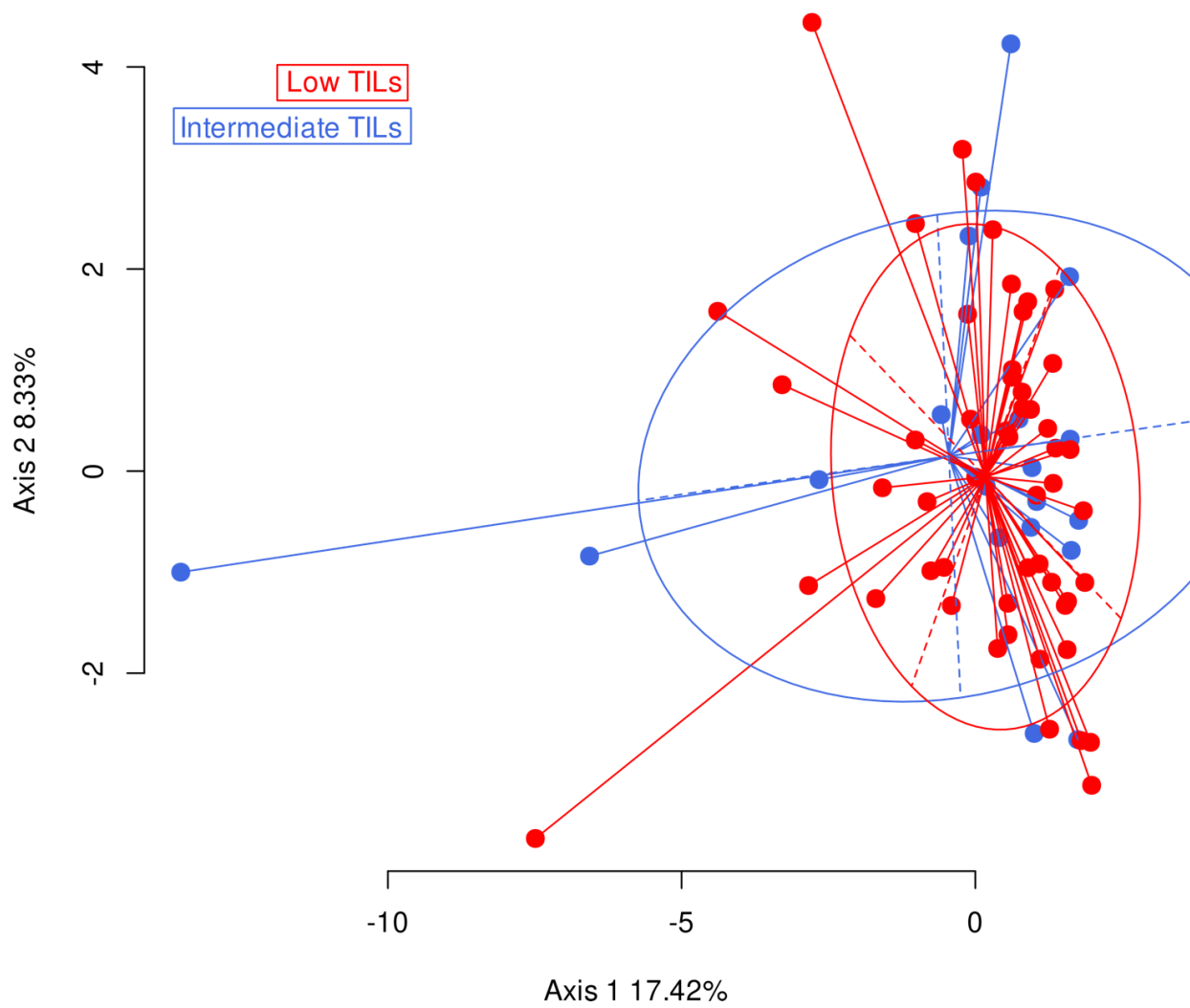


Phylum
PCoA- axes 1 and 2
Permanova p-value: 0.965

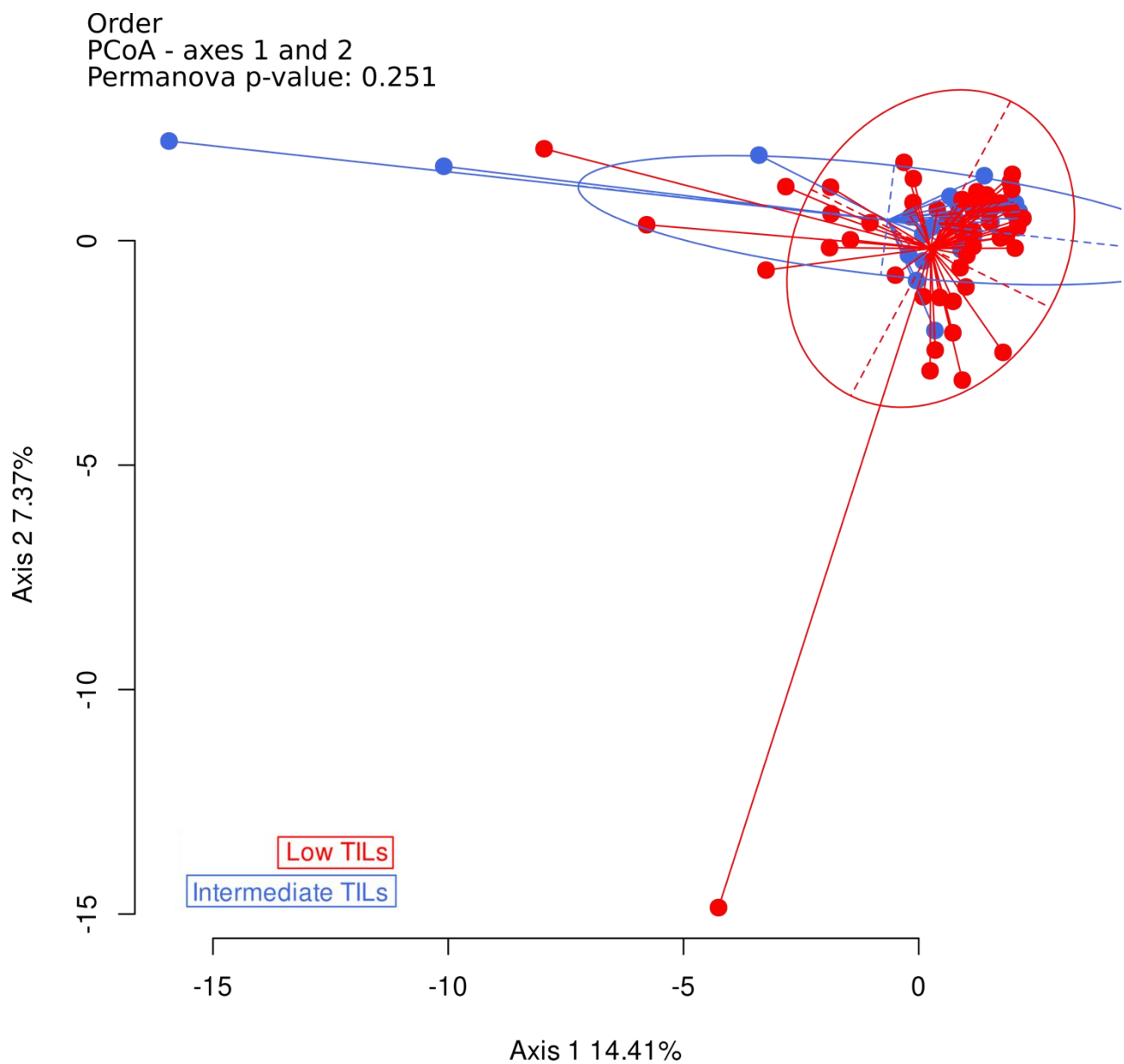


(a)

Class
PCoA - axes 1 and 2
Permanova p-value: 0.503



(b)



(c)

Supplementary Figure S2. Beta-diversity according to TILs levels. The microbiota distances were evaluated through the Bray–Curtis dissimilarity matrix at the taxonomic level of phylum (a), class (b) and order (c) and visualized through Principal Coordinates Analysis (PCoA). Each point represents the microbiota composition of one sample.