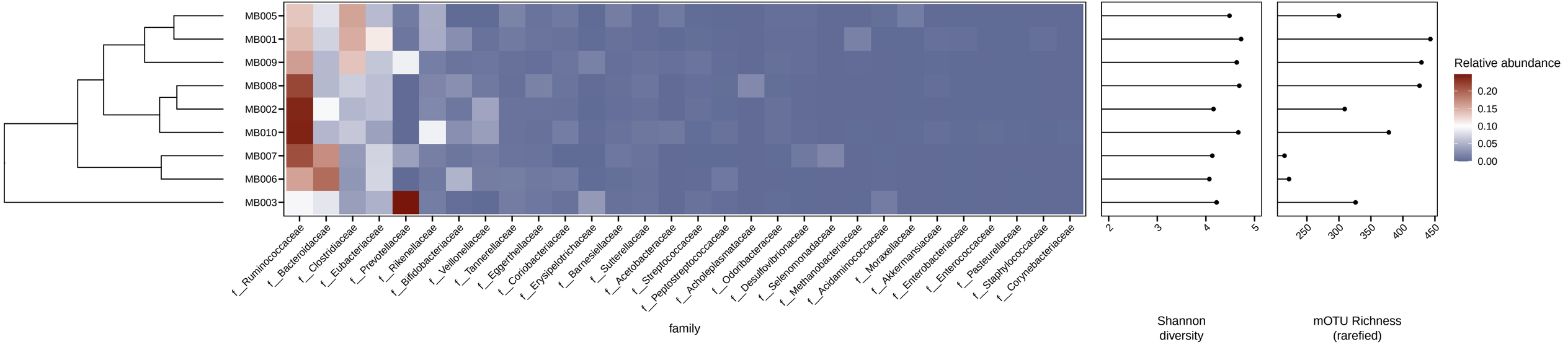
MB005
MB001
MB009
MB008
MB002
MB010
MB007
MB006
MB003
f__Ruminococcaceae
f__Bacteroidaceae
f__Clostridiaceae
f__Eubacteriaceae
f__Prevotellaceae
f__Rikenellaceae
f__Bifidobacteriaceae
f__Veillonellaceae
f__Tannerellaceae
f__Eggerthellaceae
f__Coriobacteriaceae
f__Erysipelotrichaceae
f__Barnesiellaceae
f__Sutterellaceae
f__Acetobacteraceae
f__Streptococcaceae
f__Peptostreptococcaceae
f__Acholeplasmataceae
f__Odoribacteraceae
f__Desulfovibrionaceae
f__Selenomonadaceae
f__Methanobacteriaceae
f__Acidaminococcaceae
f__Moraxellaceae
f__Akkermansiaceae
f__Enterobacteriaceae
f__Enterococcaceae
f__Pasteurellaceae
f__Staphylococcaceae
f__Corynebacteriaceae
family
2
3
4
5
Shannon diversity
250
300
350
400
450
mOTU Richness (rarefied)
Relative abundance
0.20
0.15
0.10
0.05
0.00