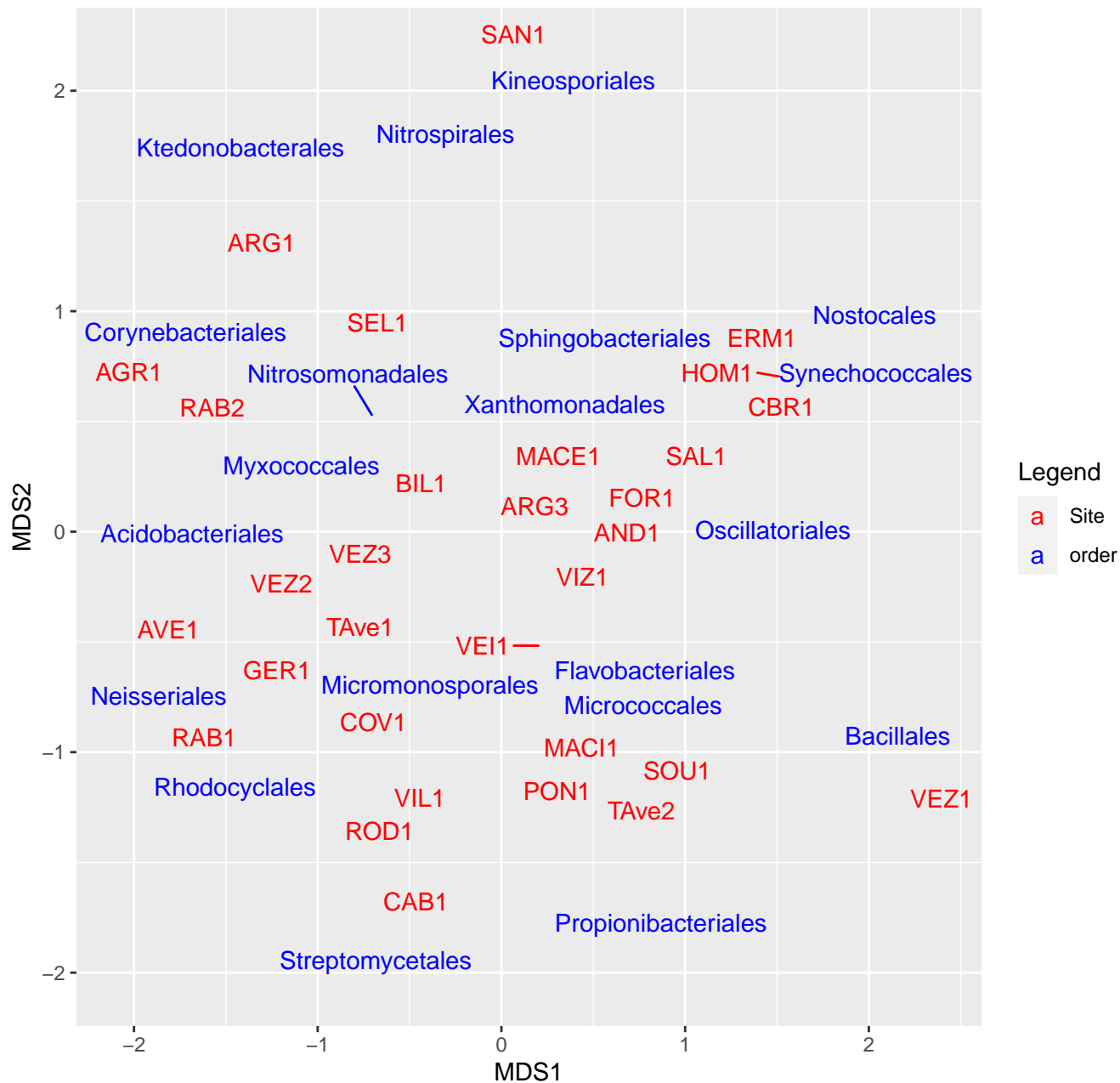


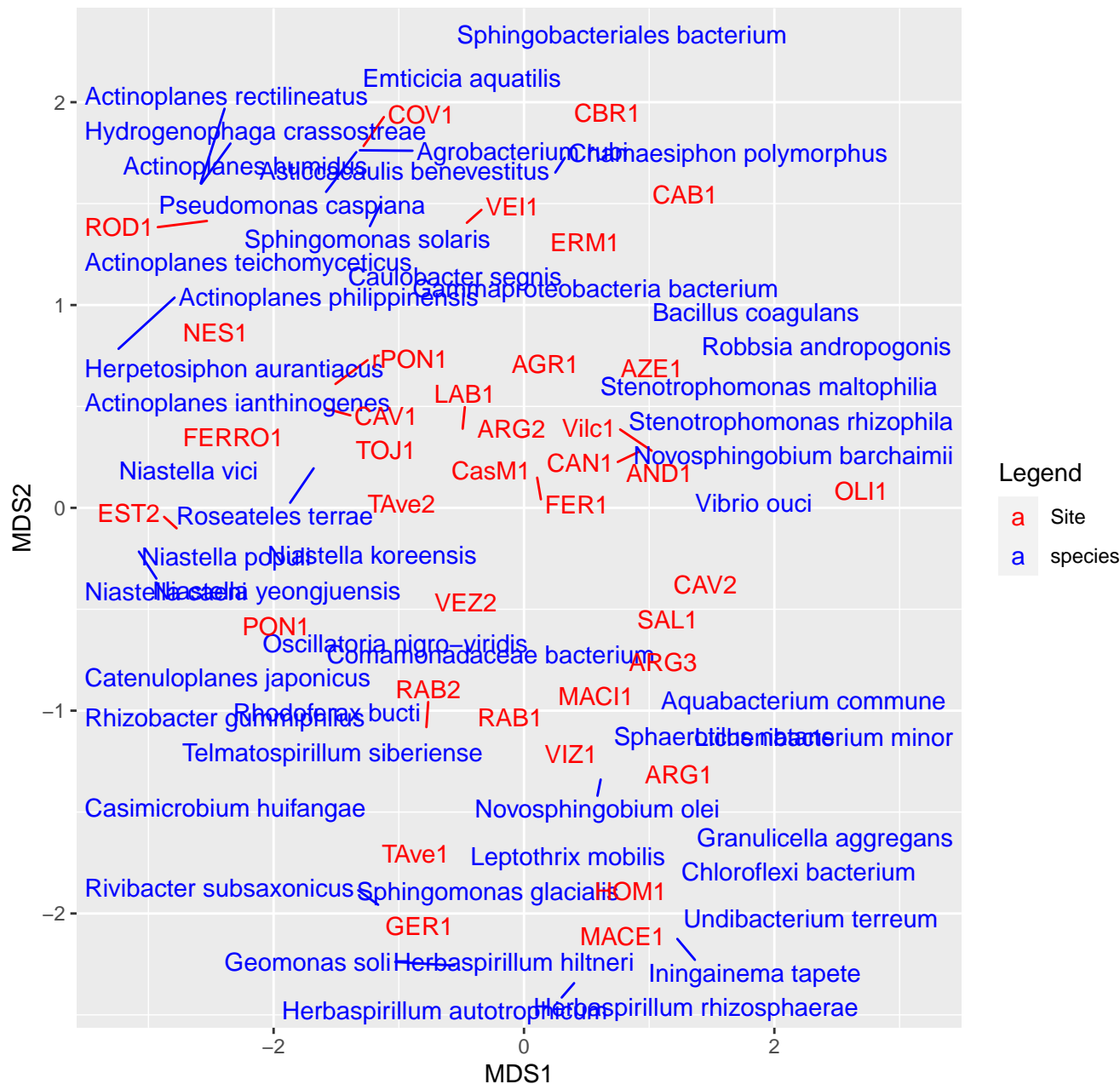
[illegible]

a	Site
a	genus

Bacteria_order



Bacteria_species

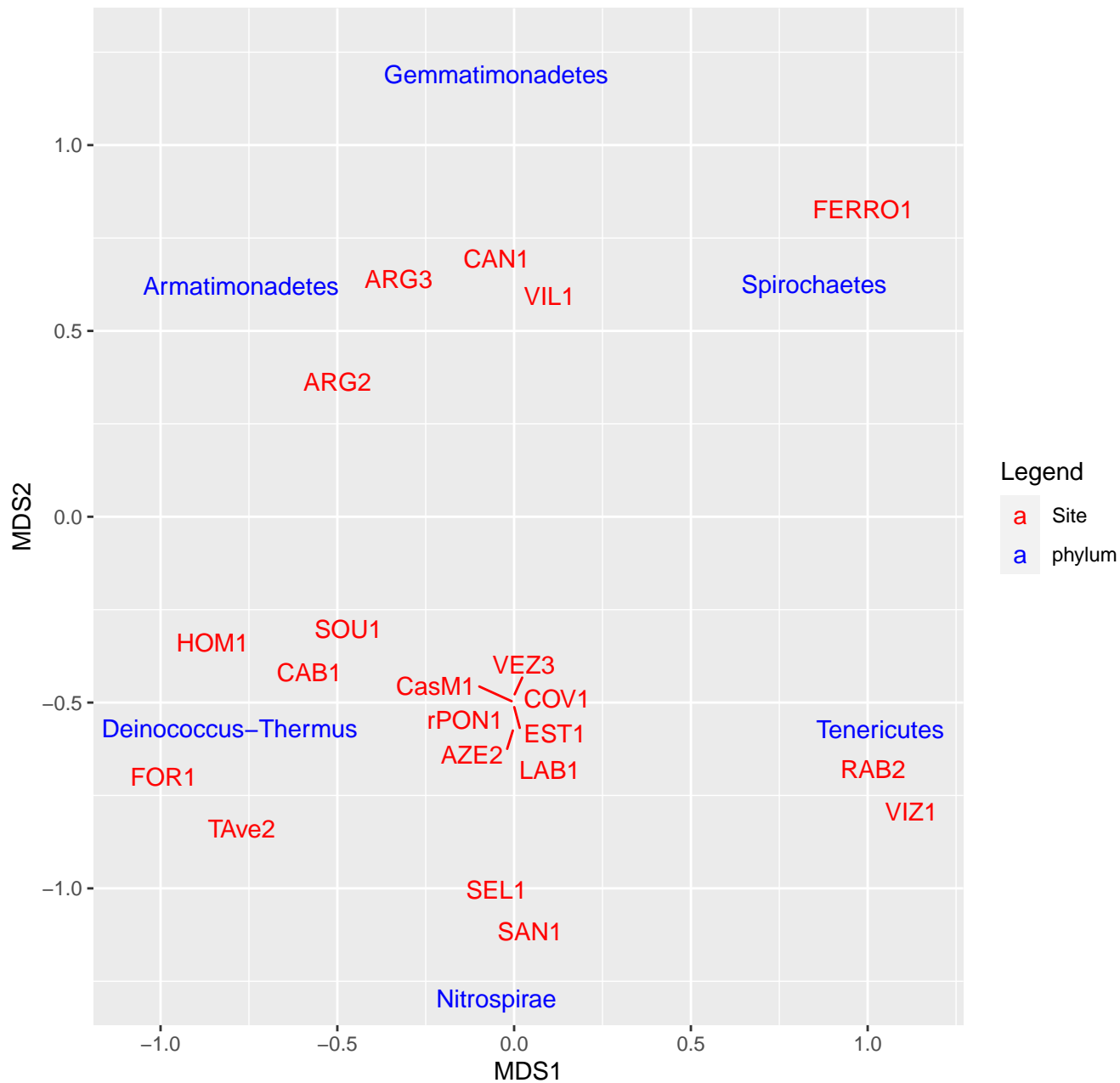


Yersiniaceae
Moraxellaceae
Hymenobacteraceae
Xanthomonadaceae
Nostocaceae
Rhodanobacteraceae
Enterobacteriaceae
Sphingobacteriaceae
Acetobacteraceae
Lichenibacteriaceae
Herpetosiphonaceae
Methylobacteriaceae
Pseudomonadaceae
Rhodospirillaceae
Acidobacteriaceae
Aestuariivirgaceae
Methylocystaceae
Methylophilaceae
Zoogloeaceae
Alcaligenaceae
Micromonosporaceae
Flavobacteriaceae
Streptomycetaceae
Erythrobacteraceae
Chamaesiphonaceae
Nitrospiraceae
Hyphomicrobiaceae
Kineosporiaceae
Microcoleaceae
Microbacteriaceae
Nocardioidaceae
Intrasporangiaceae
SOU1

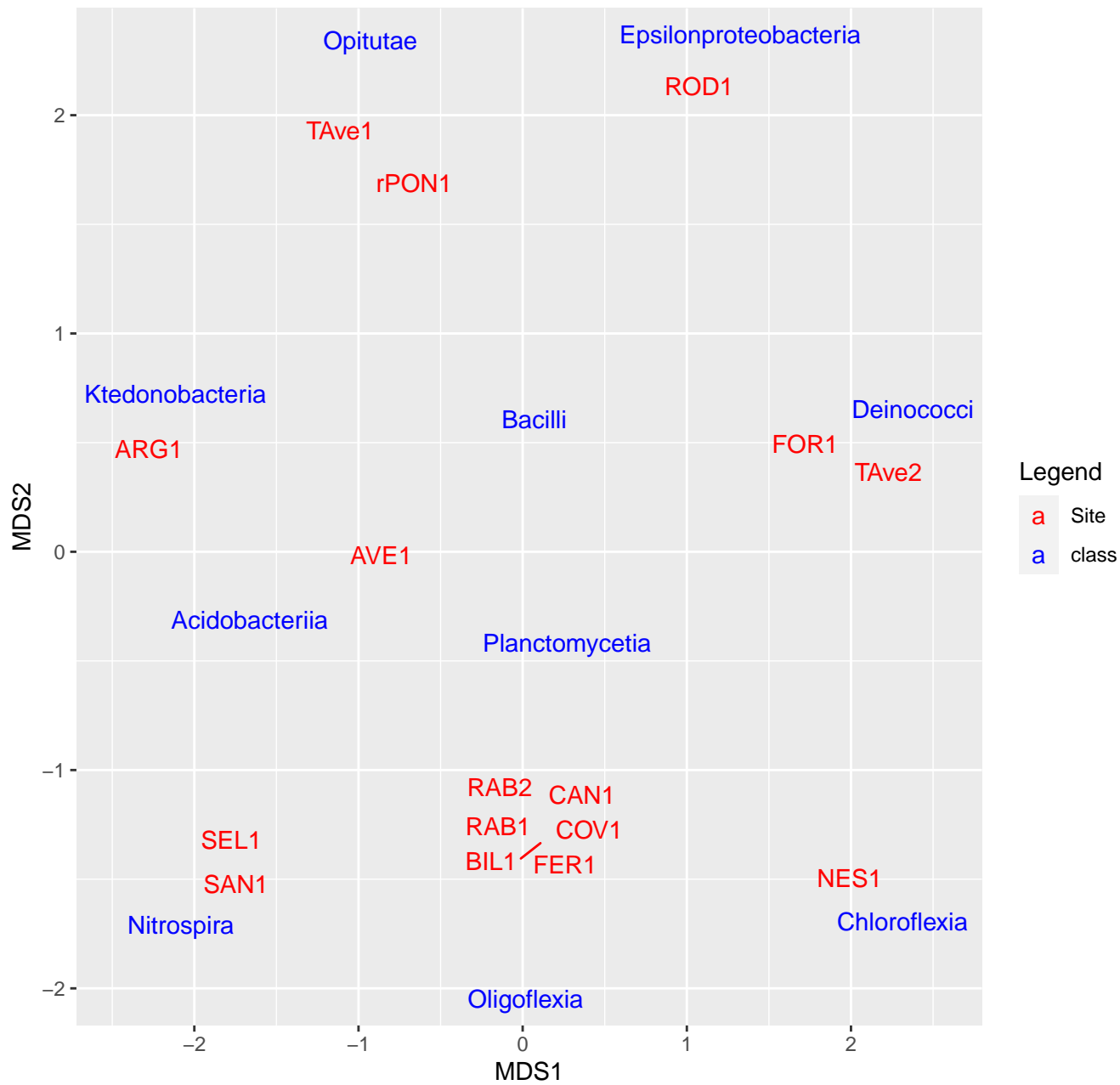
FOR1
SAL1
VIL1
OLI1
HOM1
MACE1
ARG1
AND1
VEZ2
GER1
RAB1
AZE2
AGR1
SEL1
SAN1
CAN1
VEZ1
EST1
EST2
VEZ3
ERM1
CAB1
AZE1
MAC1
PON1
CBR1
TAVe1
TAVe2

a	Site
a	family

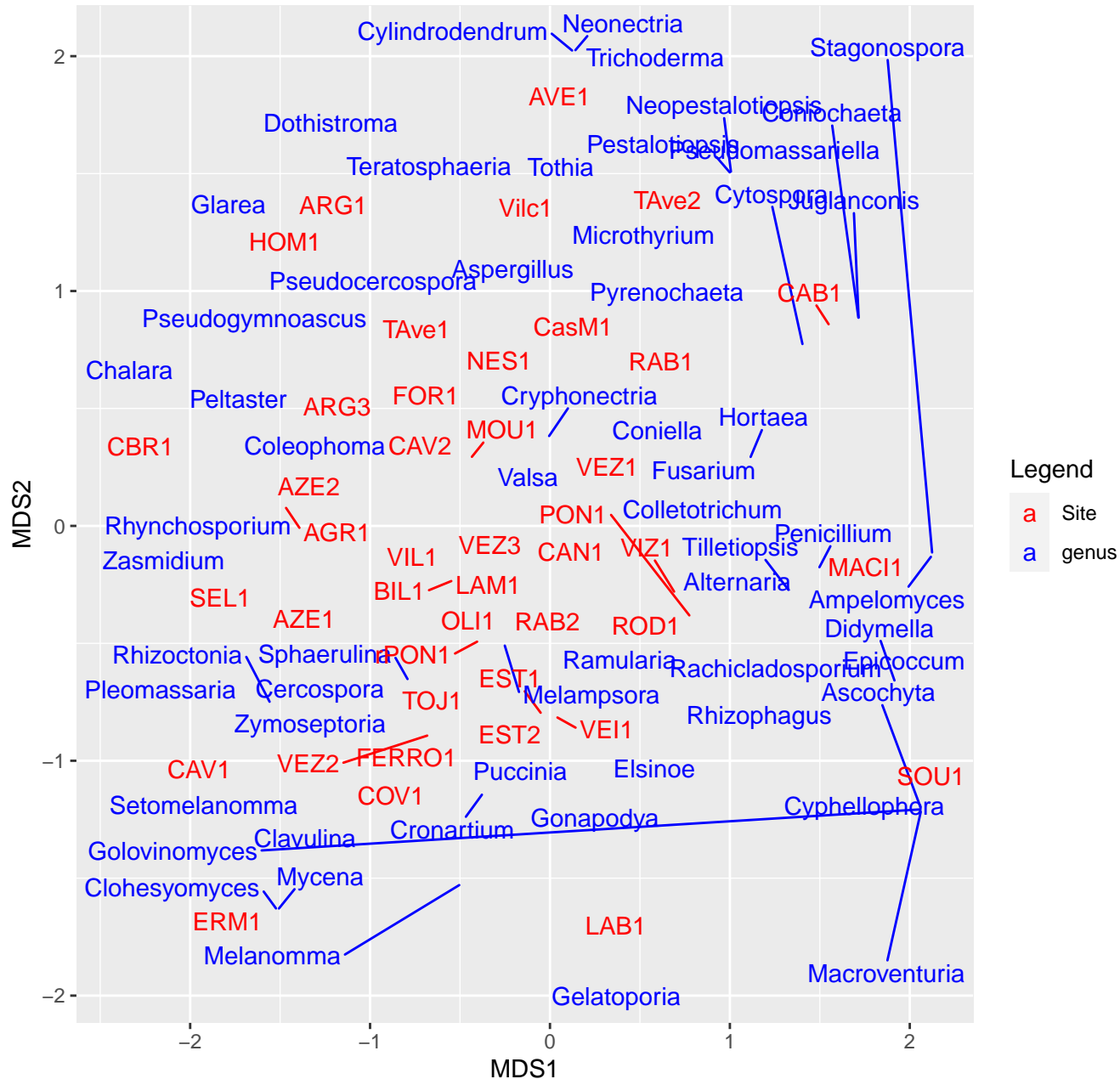
Bacteria_phylum



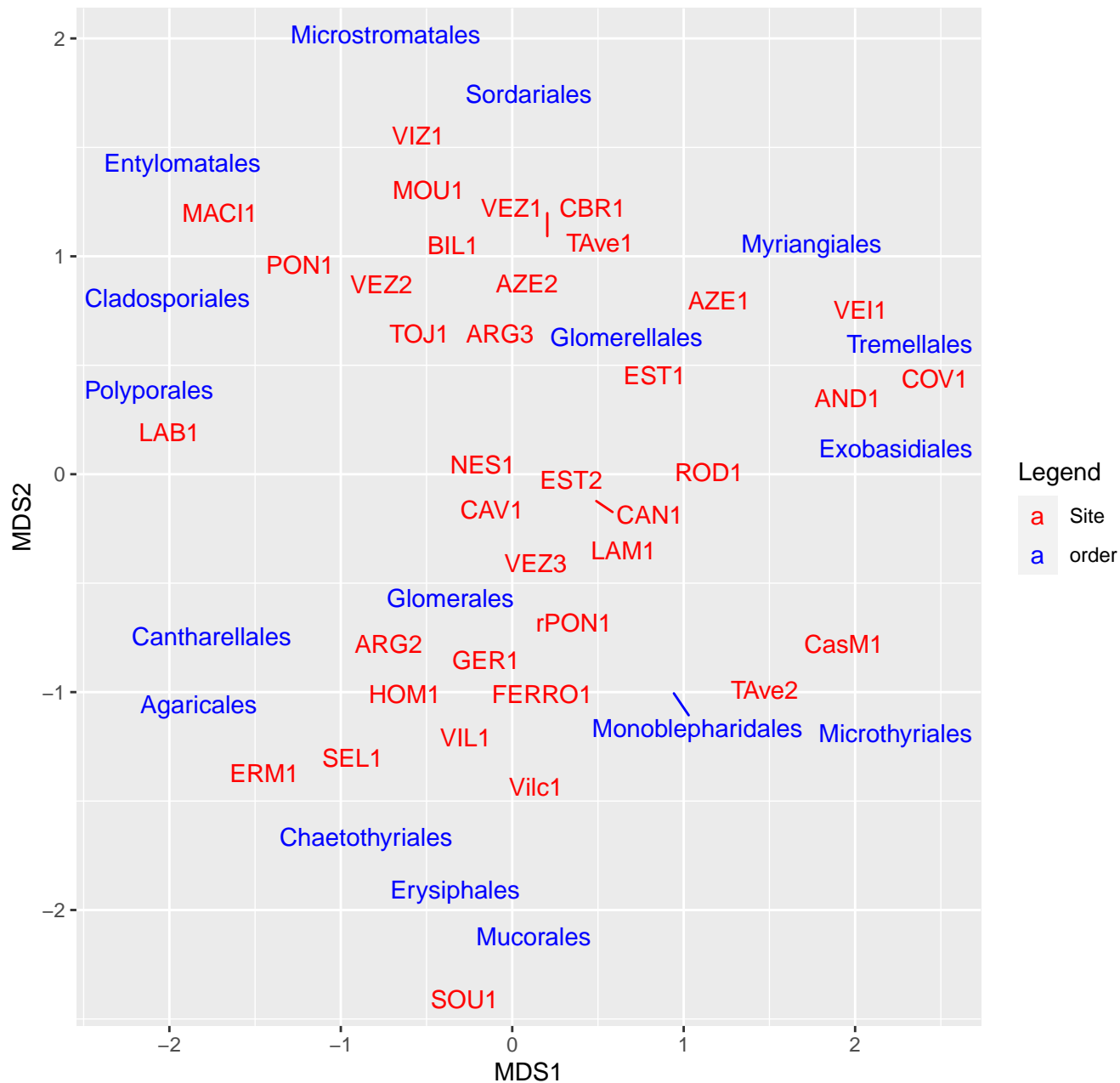
Bacteria_class



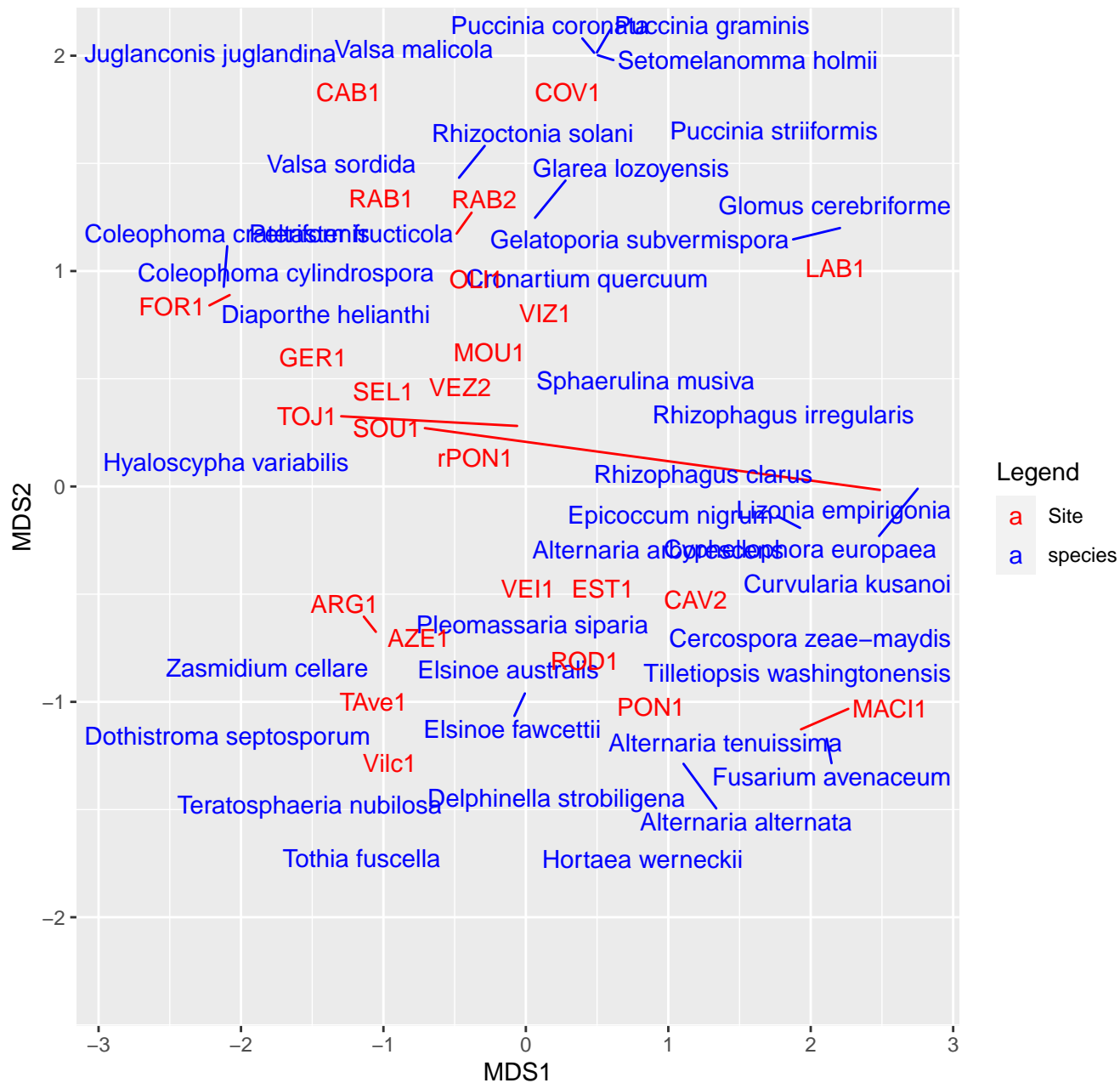
Fungi_genus



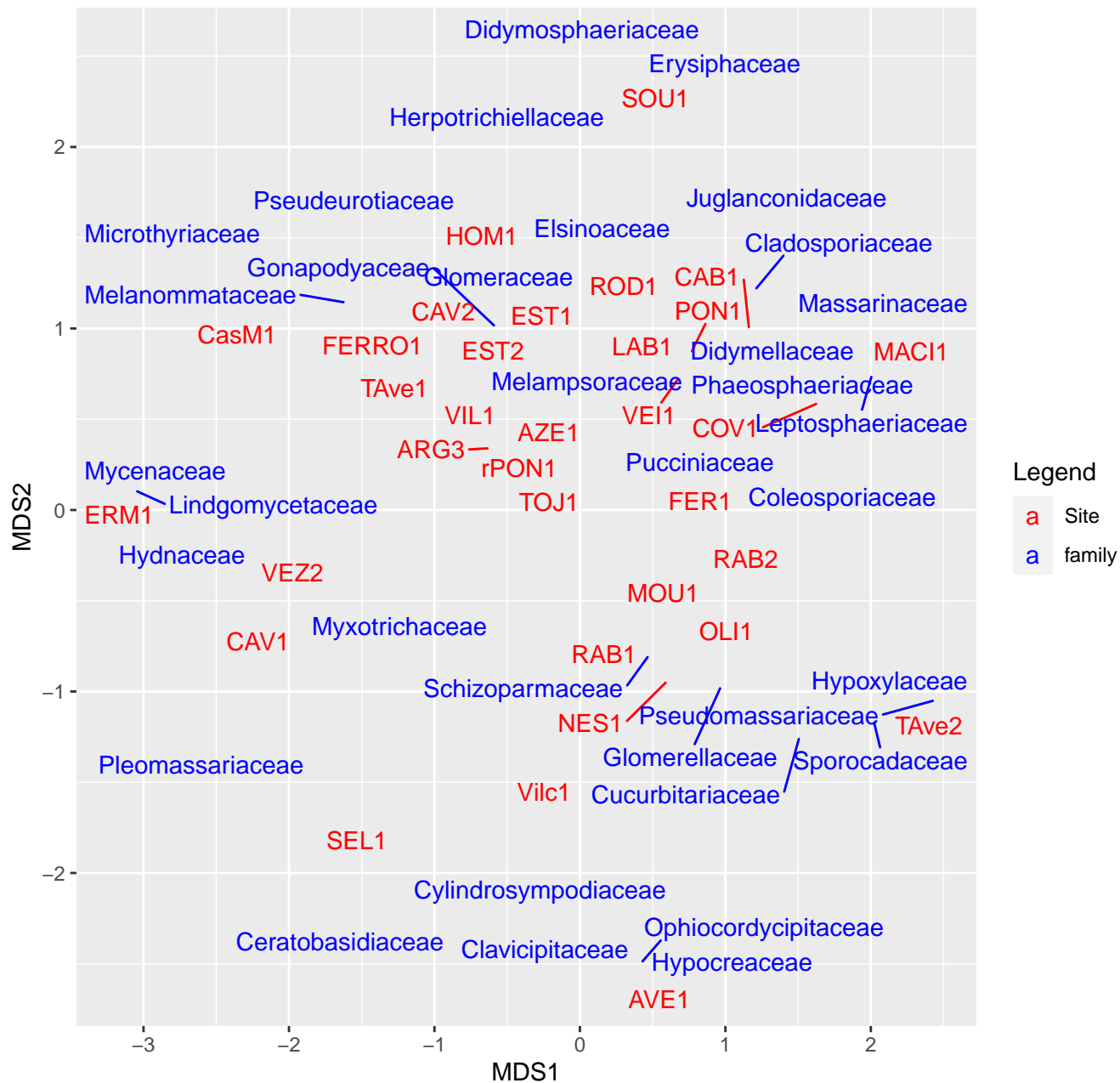
Fungi_order



Fungi_species

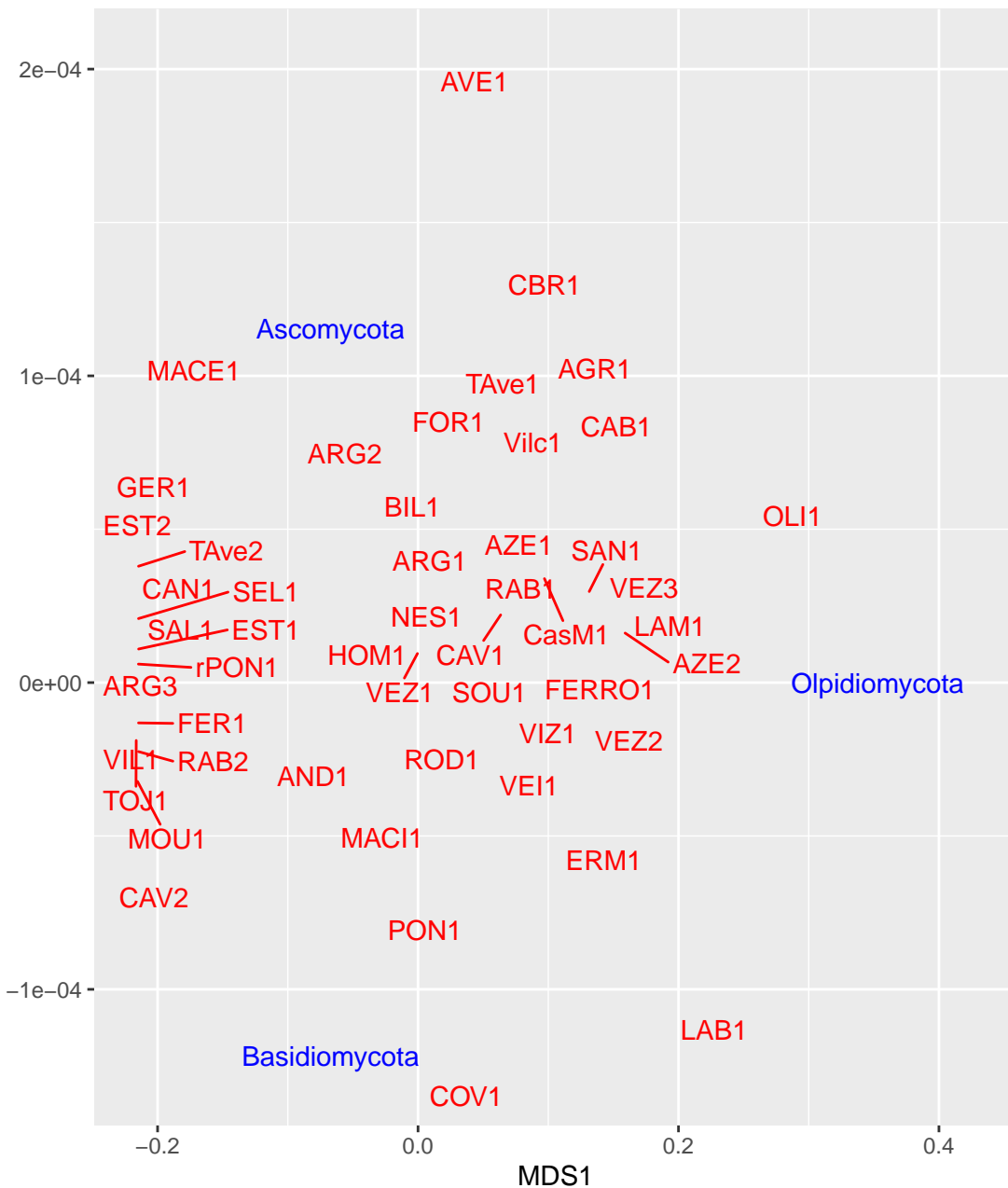


Fungi_family



Fungi_phylum

MDS2



Fungi_class

