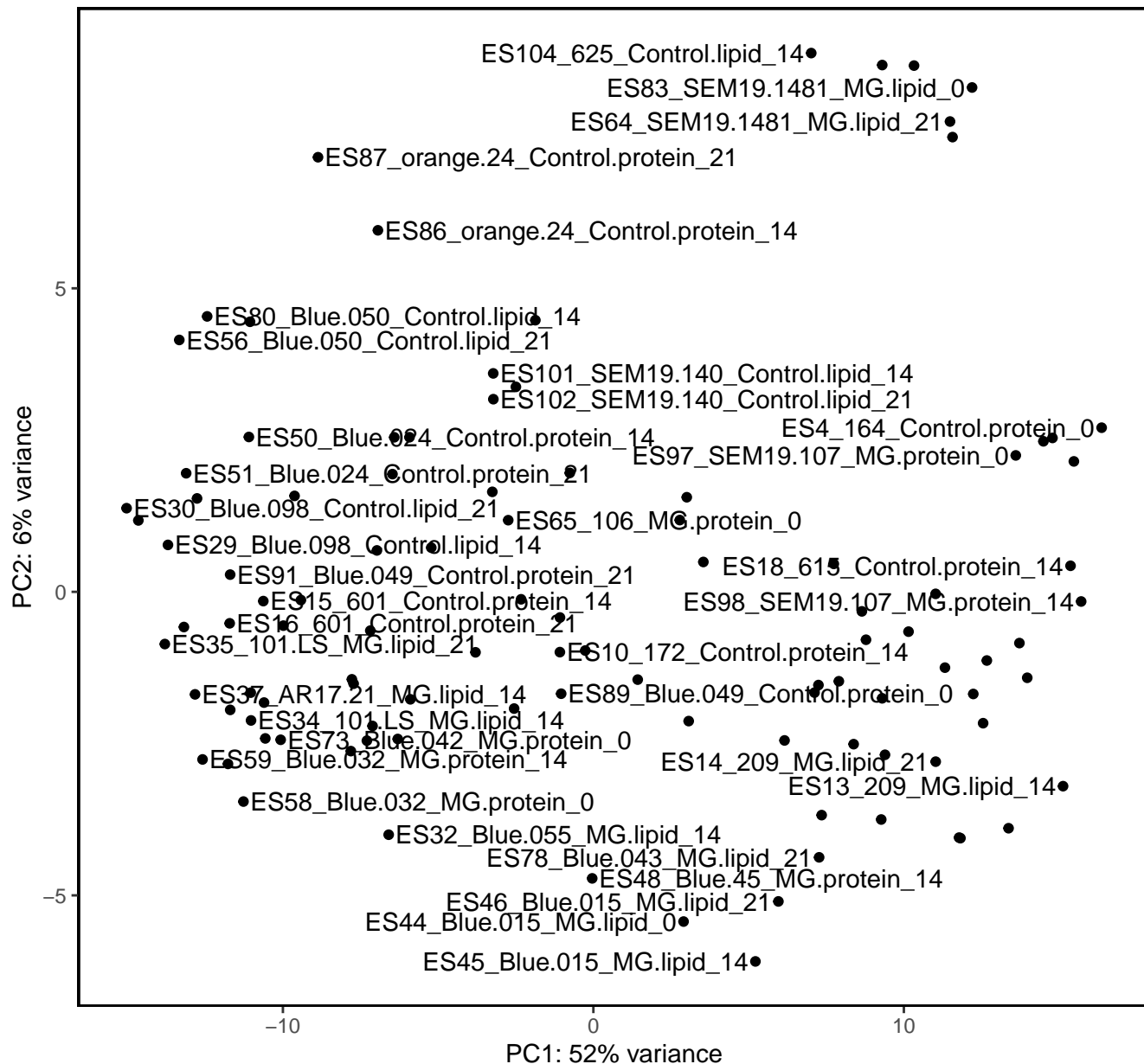
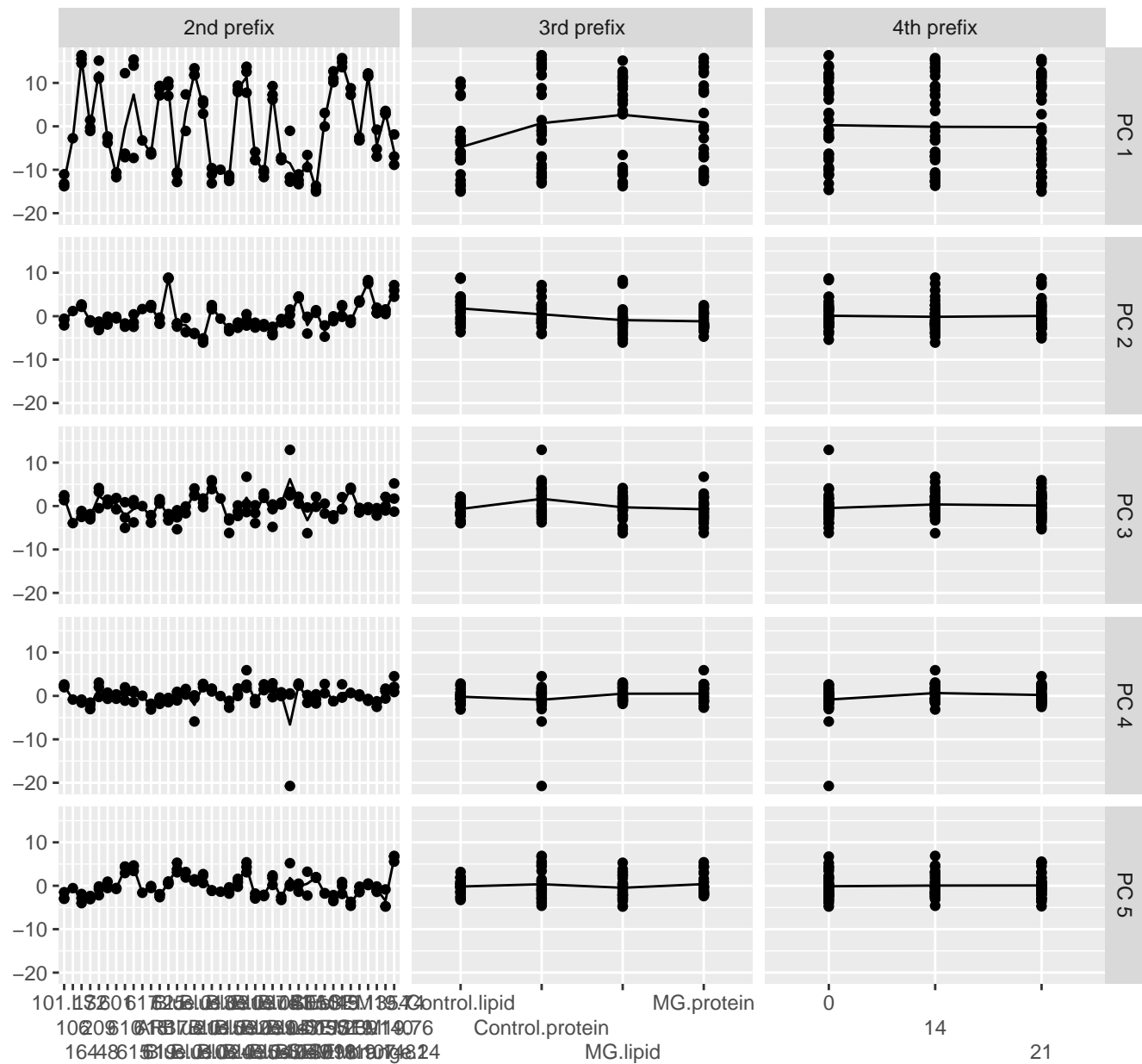


# First PCs on rlog-transformed data

Top 500 genes

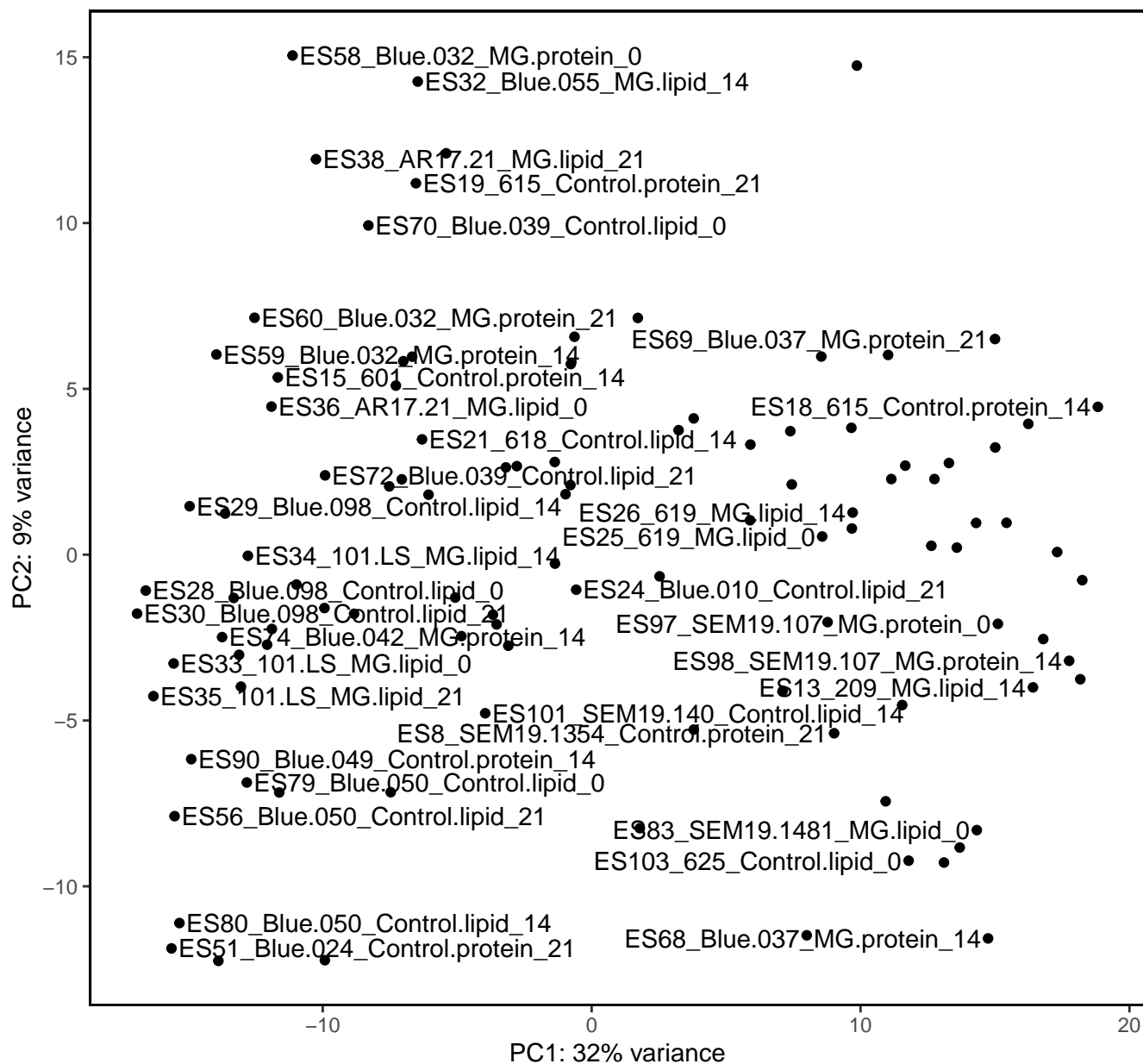


## Top 500 genes



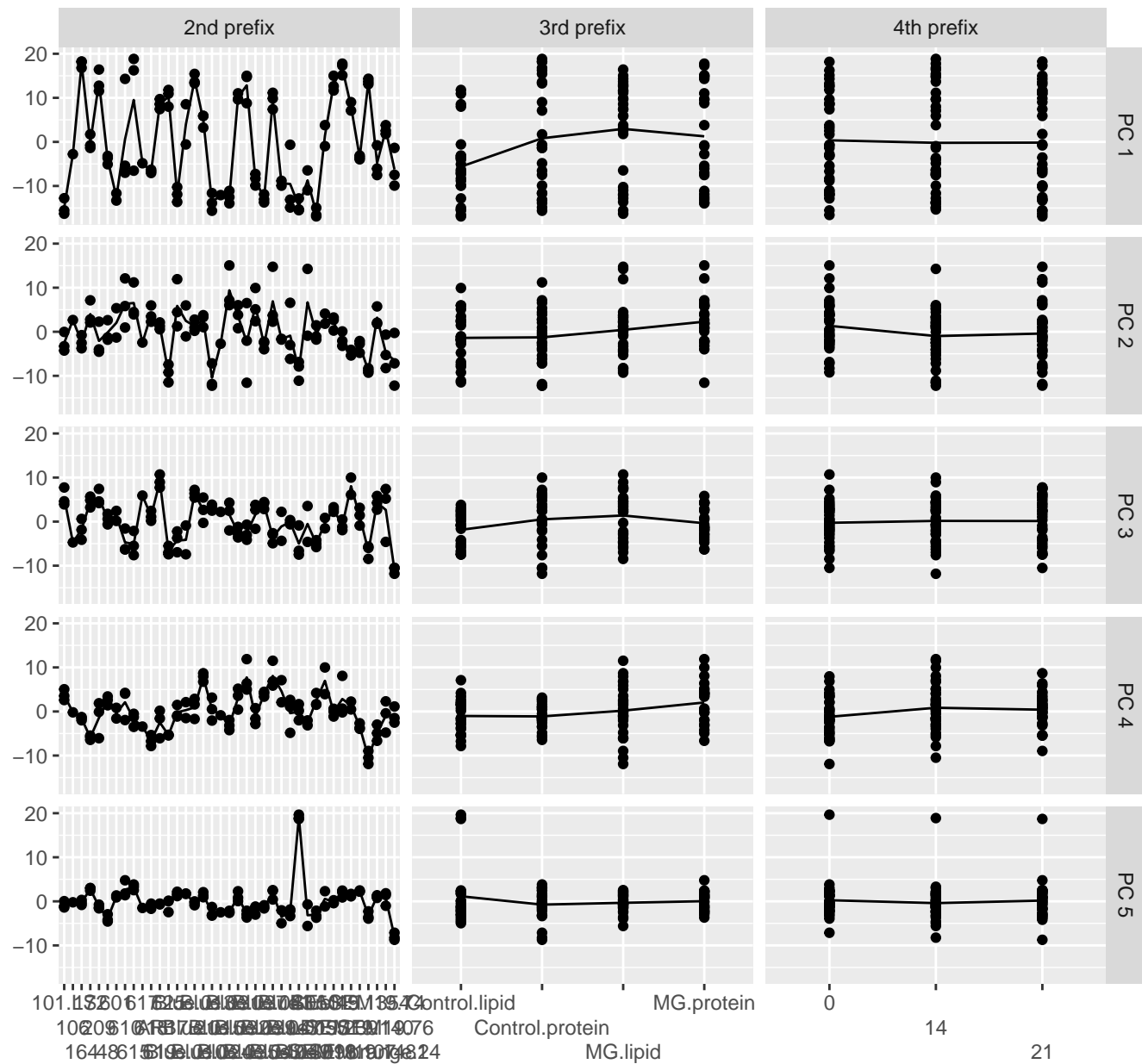
# First PCs on rlog-transformed data

All genes



# PCs split by sample-name prefixes

All genes



## Euclidean distance between rlog of samples

