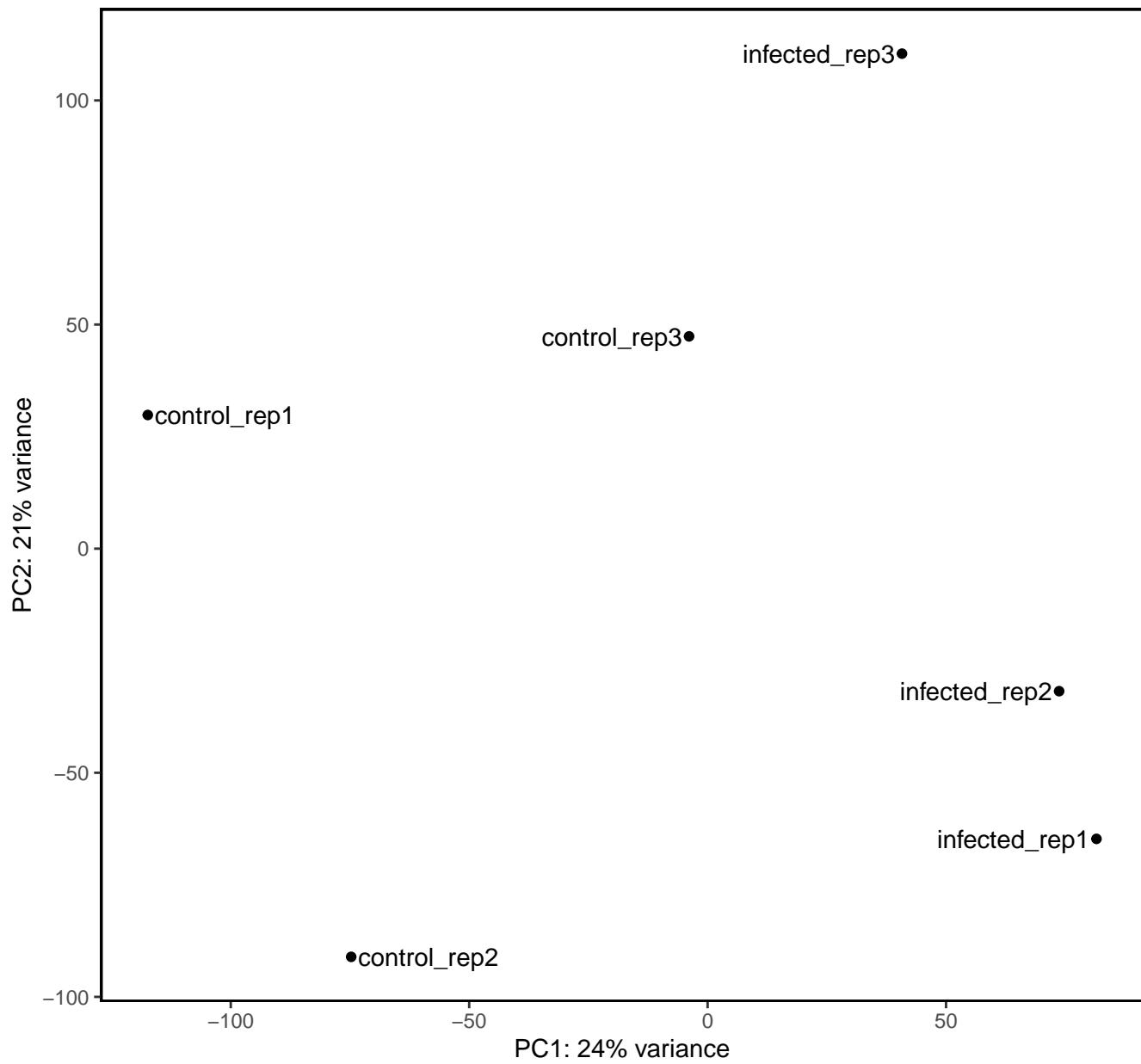


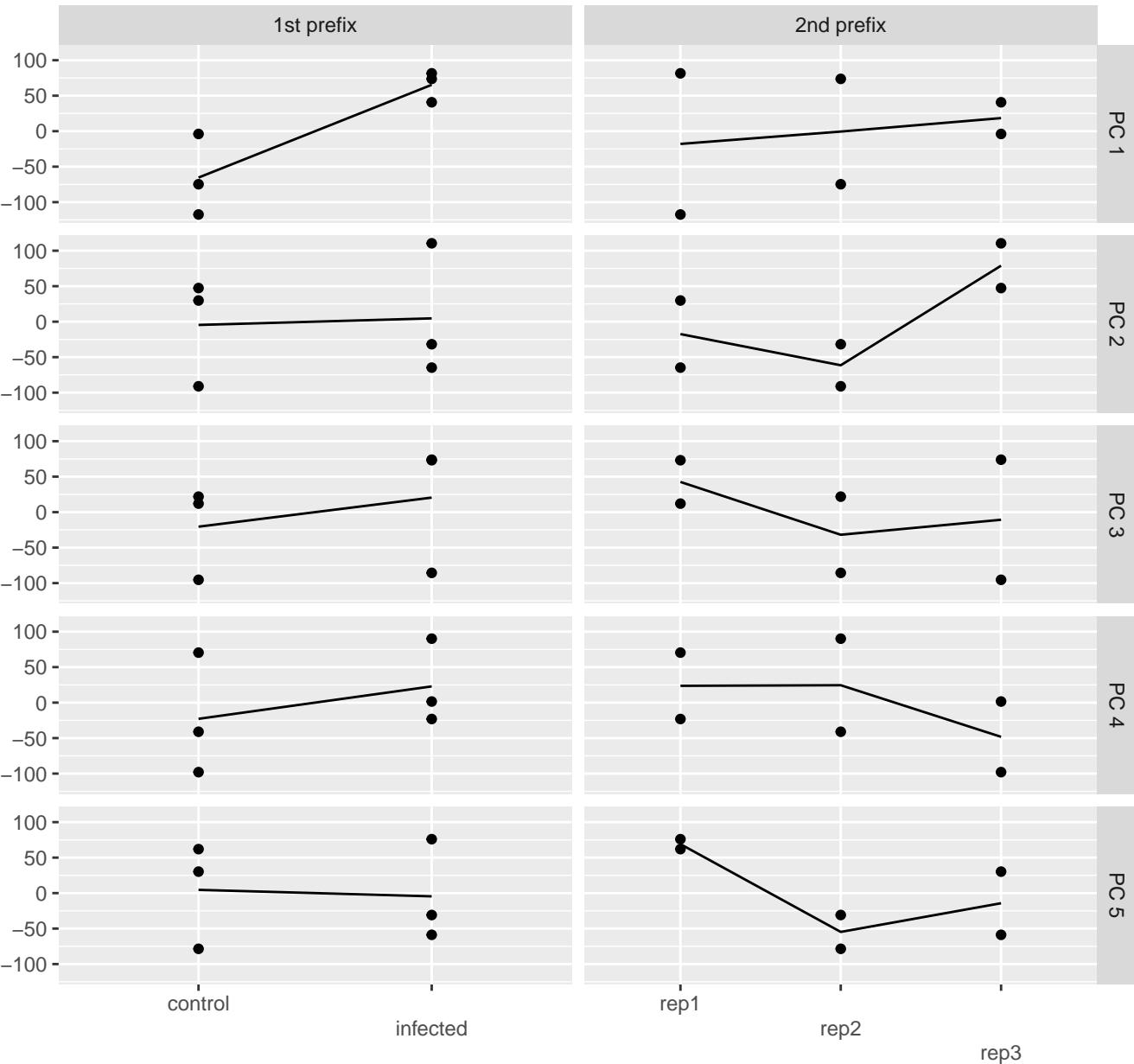
# First PCs on vst-transformed data

Top 500 genes



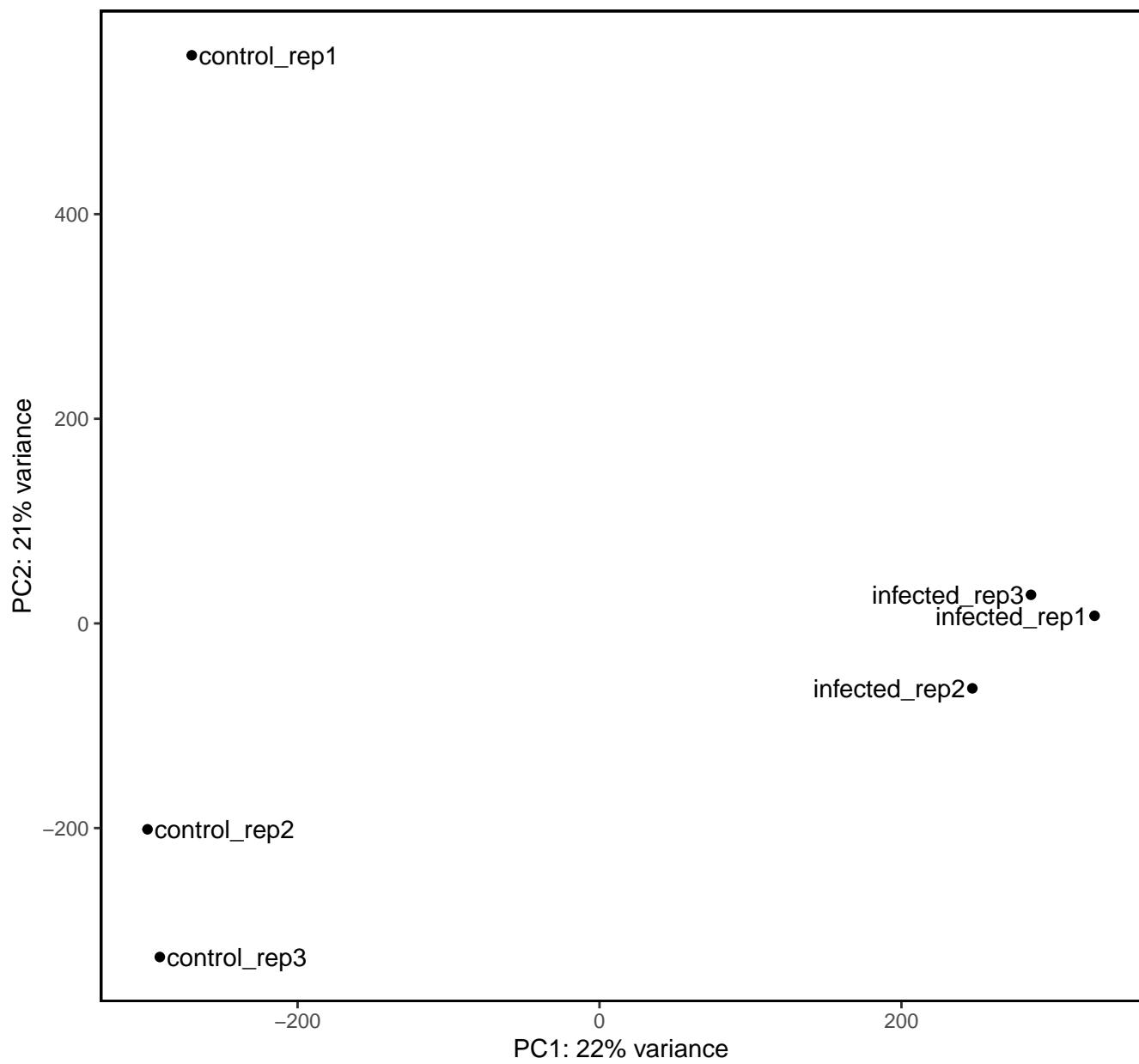
# PCs split by sample-name prefixes

Top 500 genes



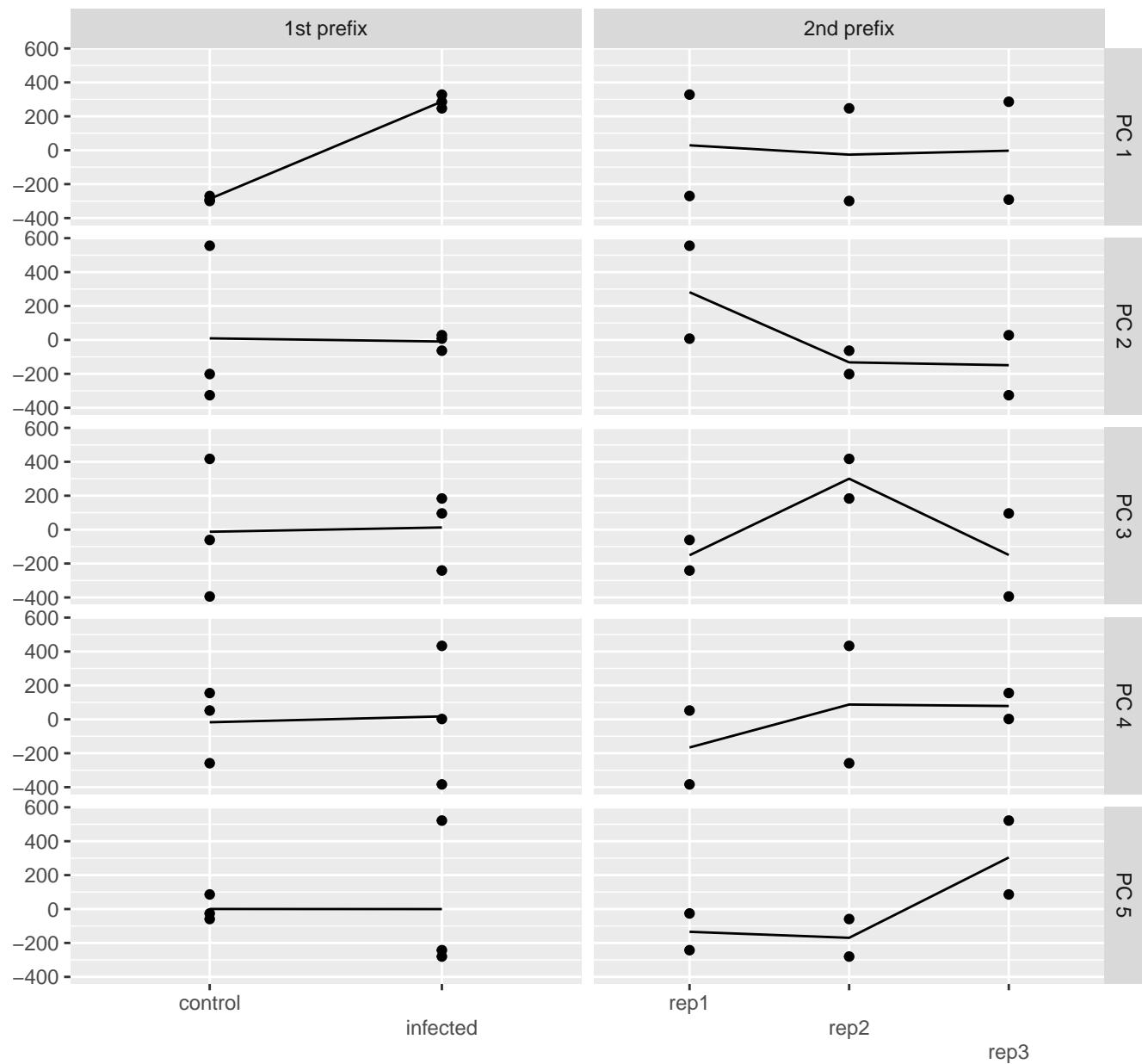
# First PCs on vst-transformed data

All genes



# PCs split by sample-name prefixes

All genes



# Euclidean distance between vst of samples

