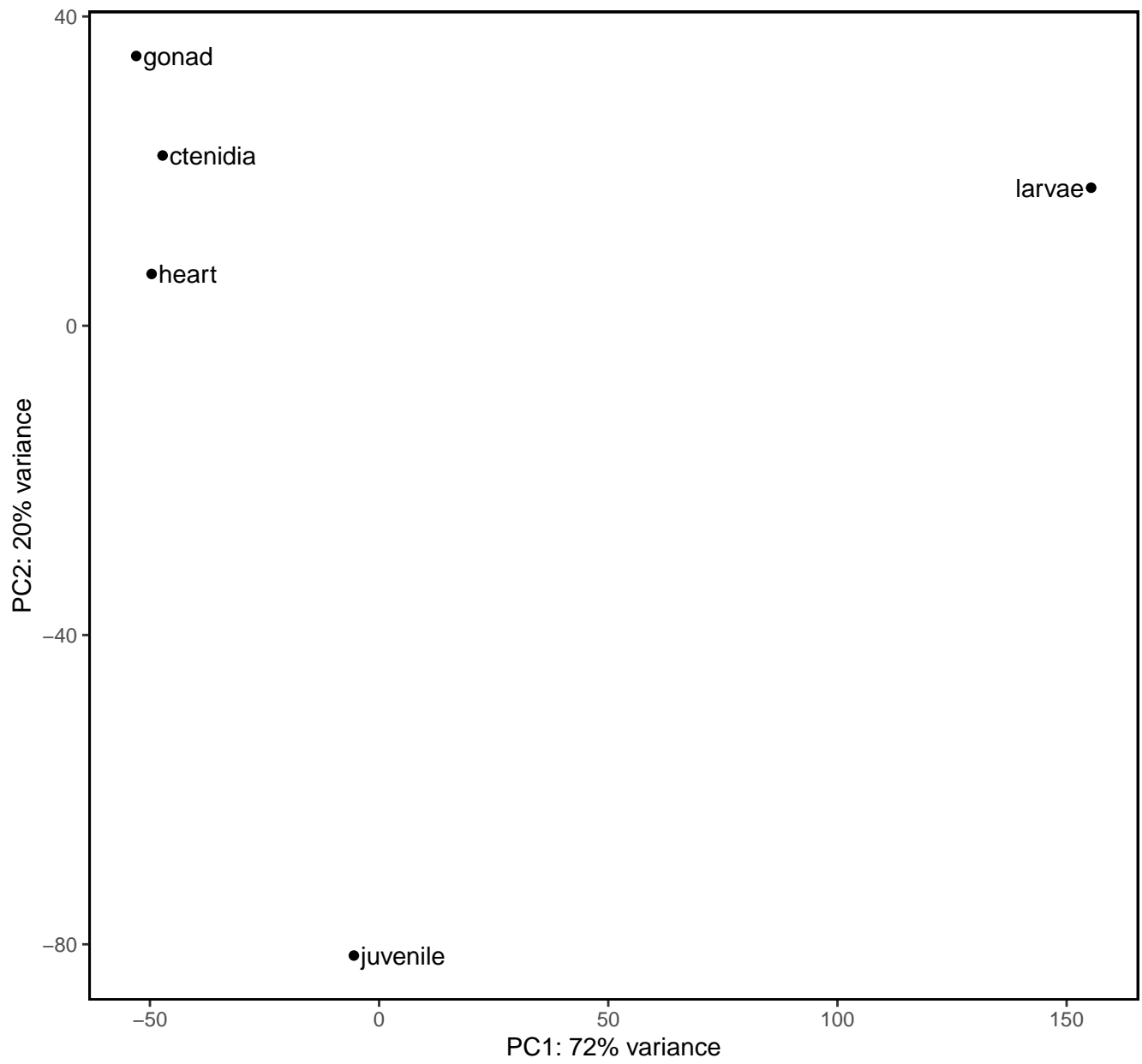


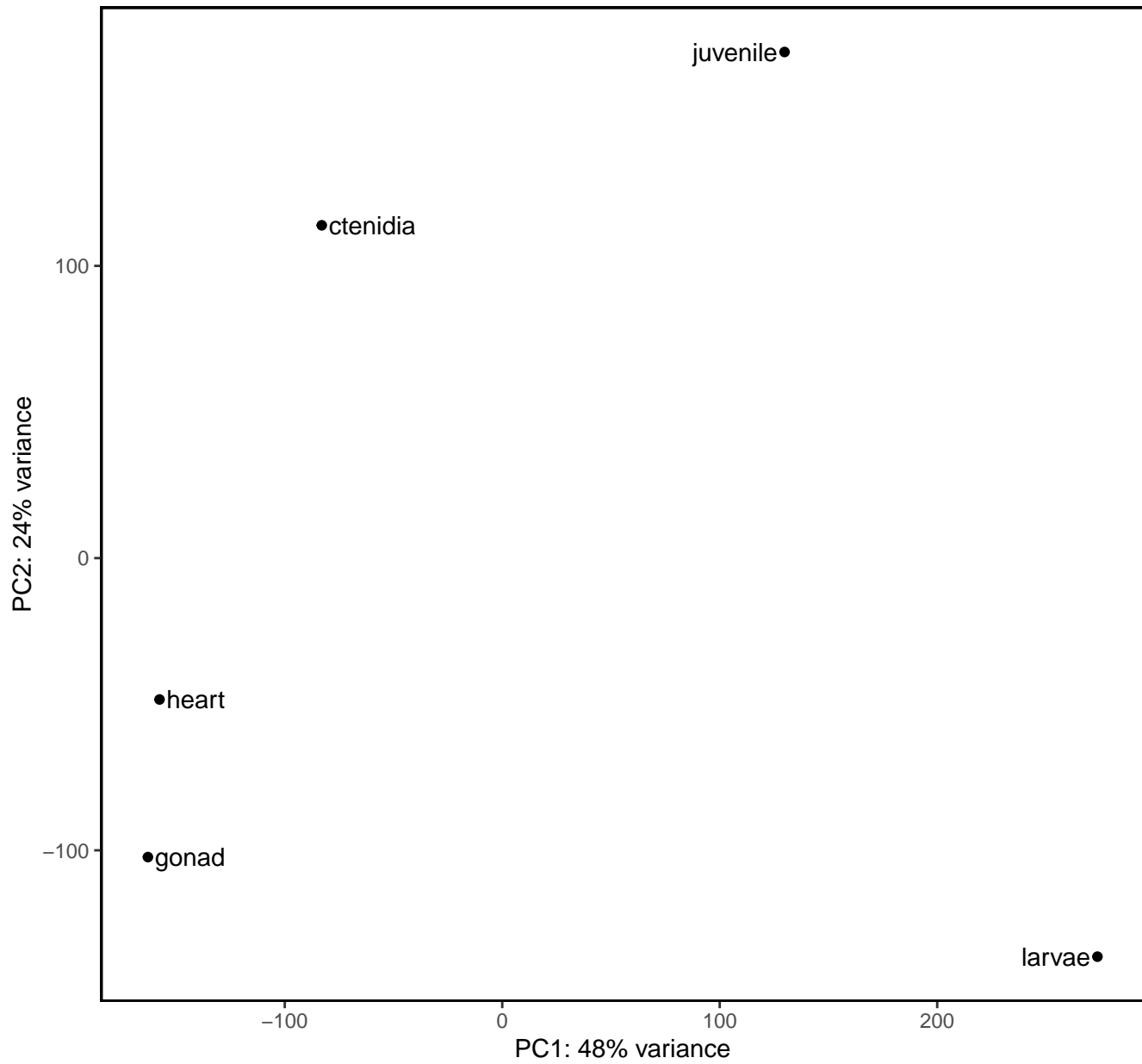
# First PCs on rlog-transformed data

Top 500 genes



# First PCs on rlog-transformed data

All genes



# Euclidean distance between rlog of samples

