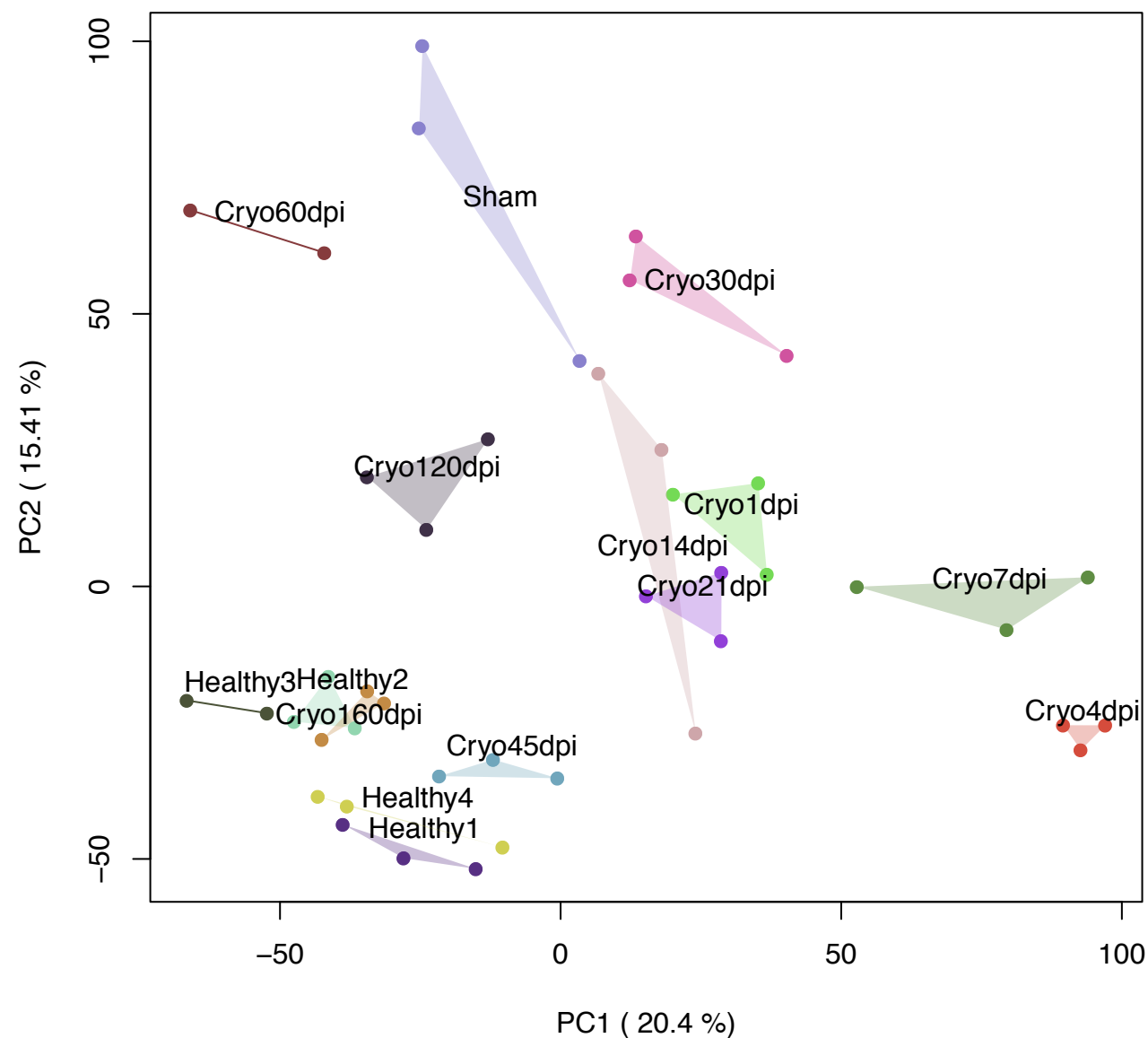


Data before batch correction

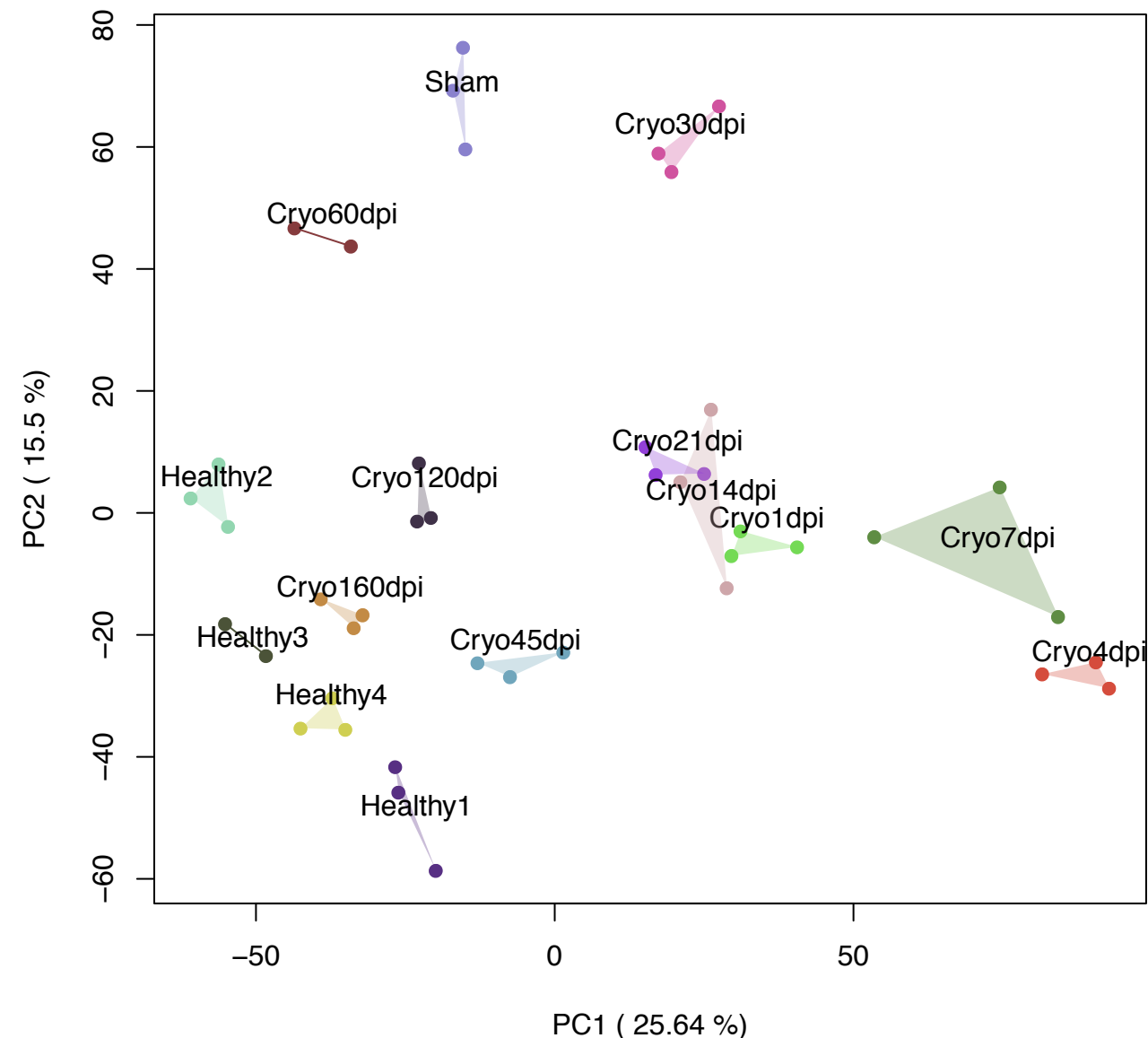
mRNA



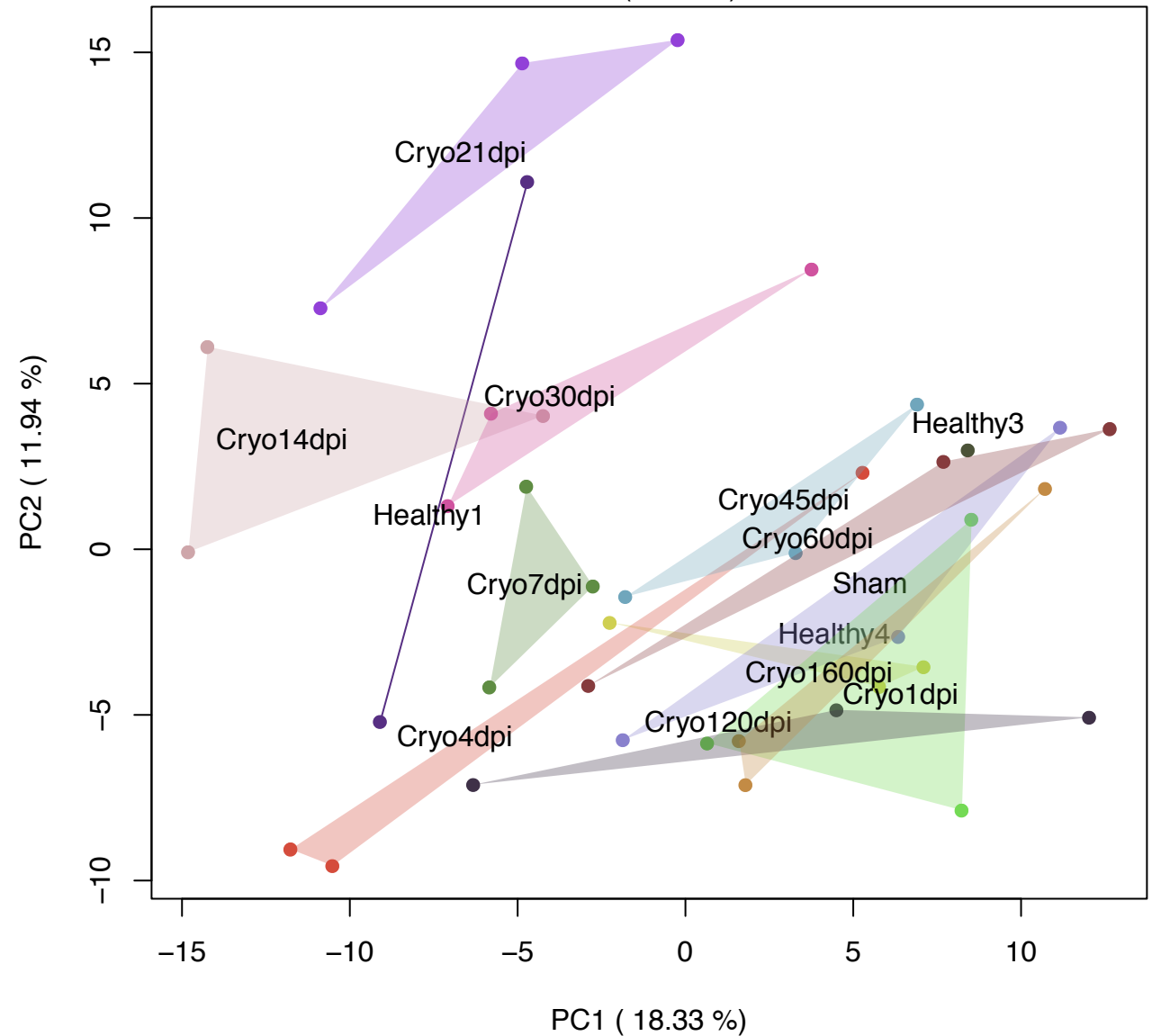
SVA

Data after batch correction

PC2 (15.5 %)



miRNA



SVA

