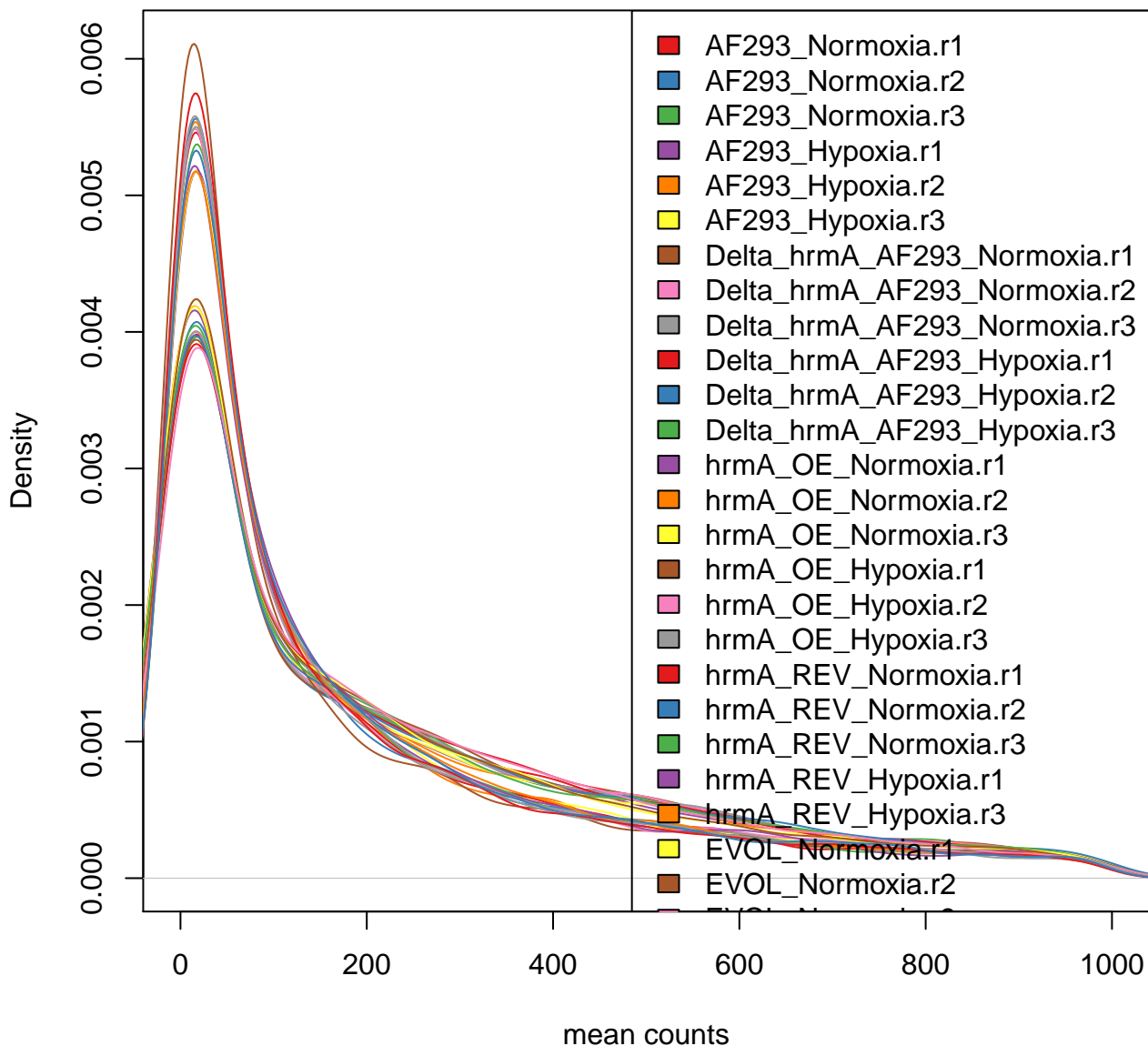
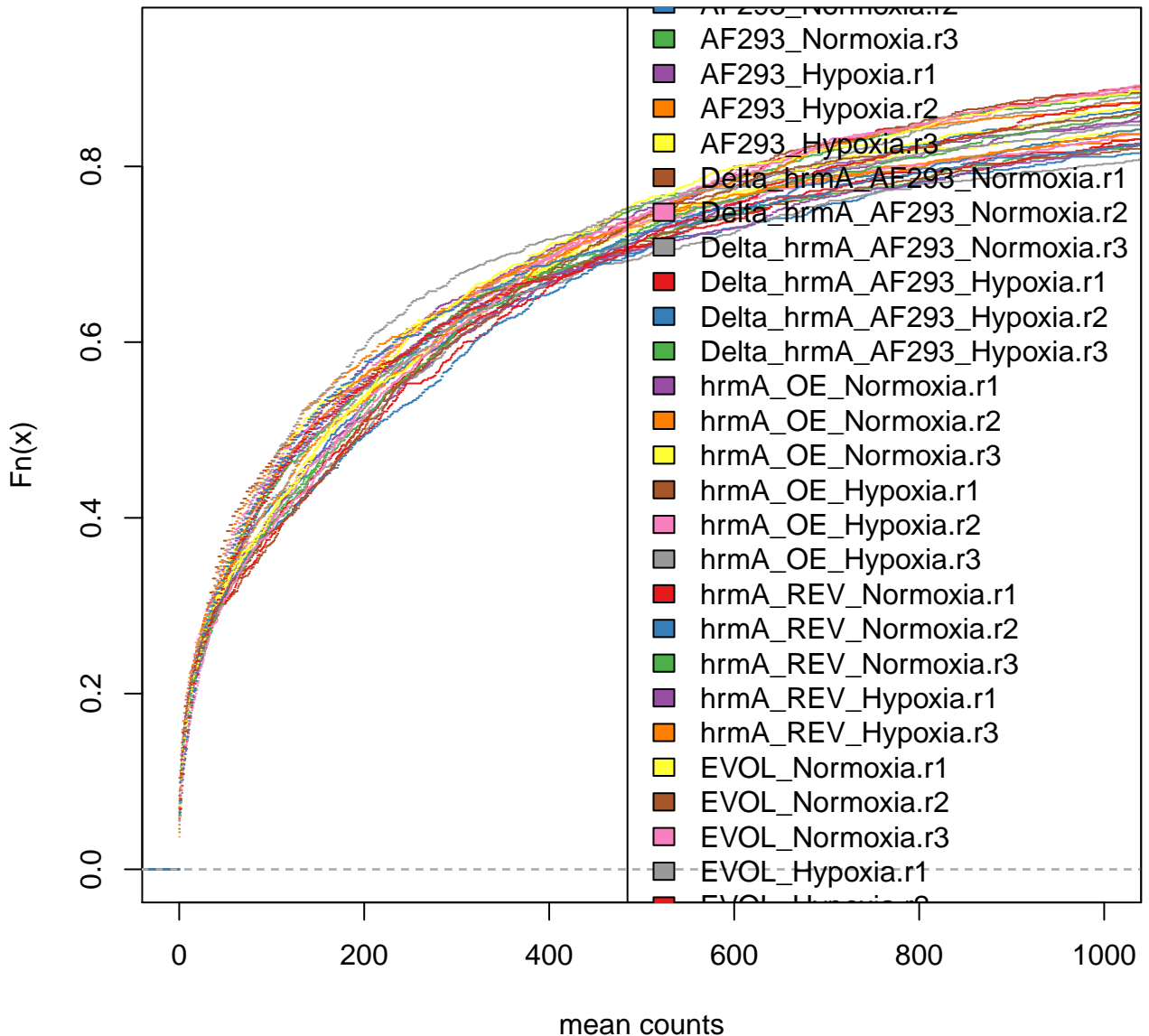


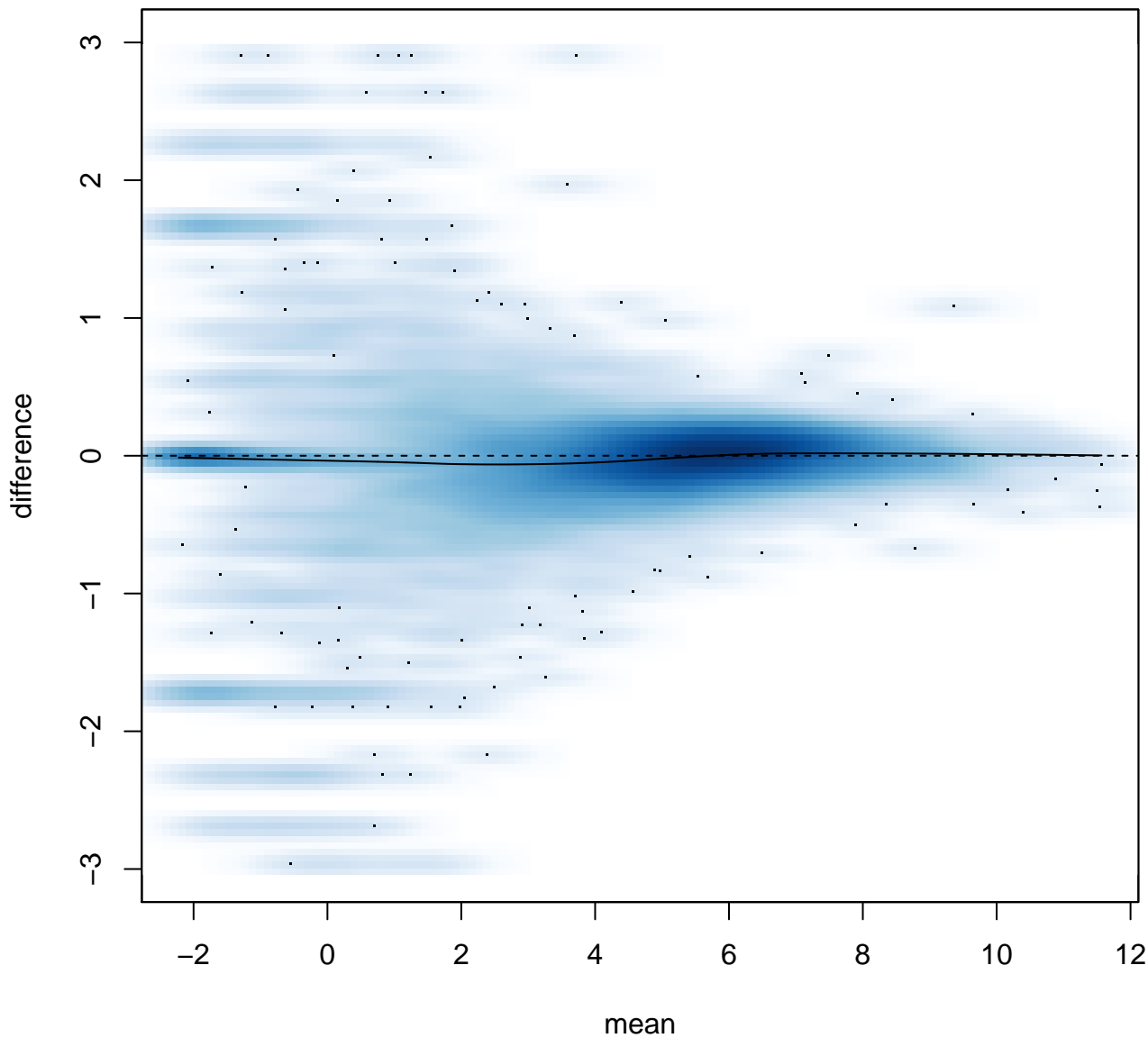
densities



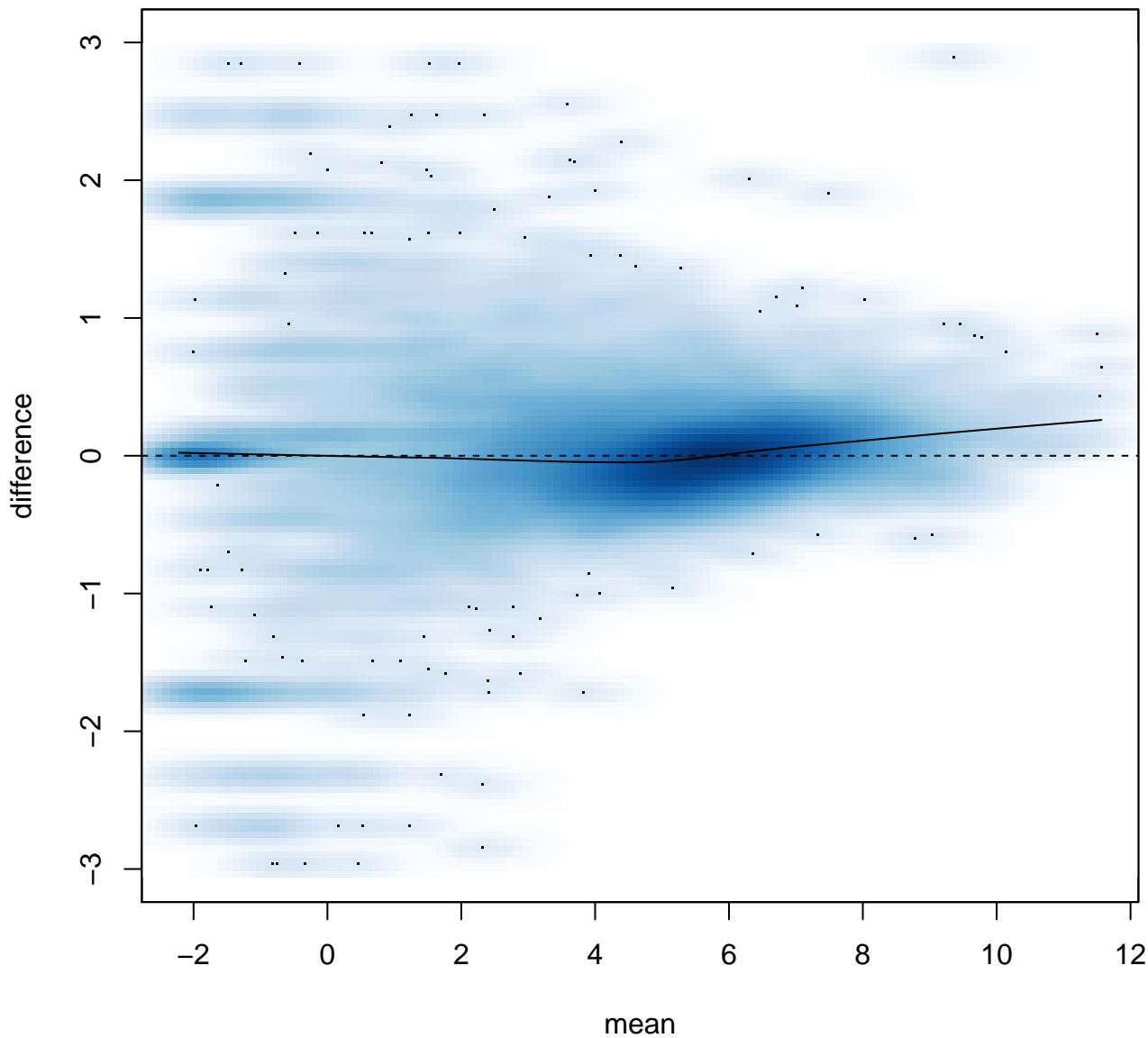
ecdf



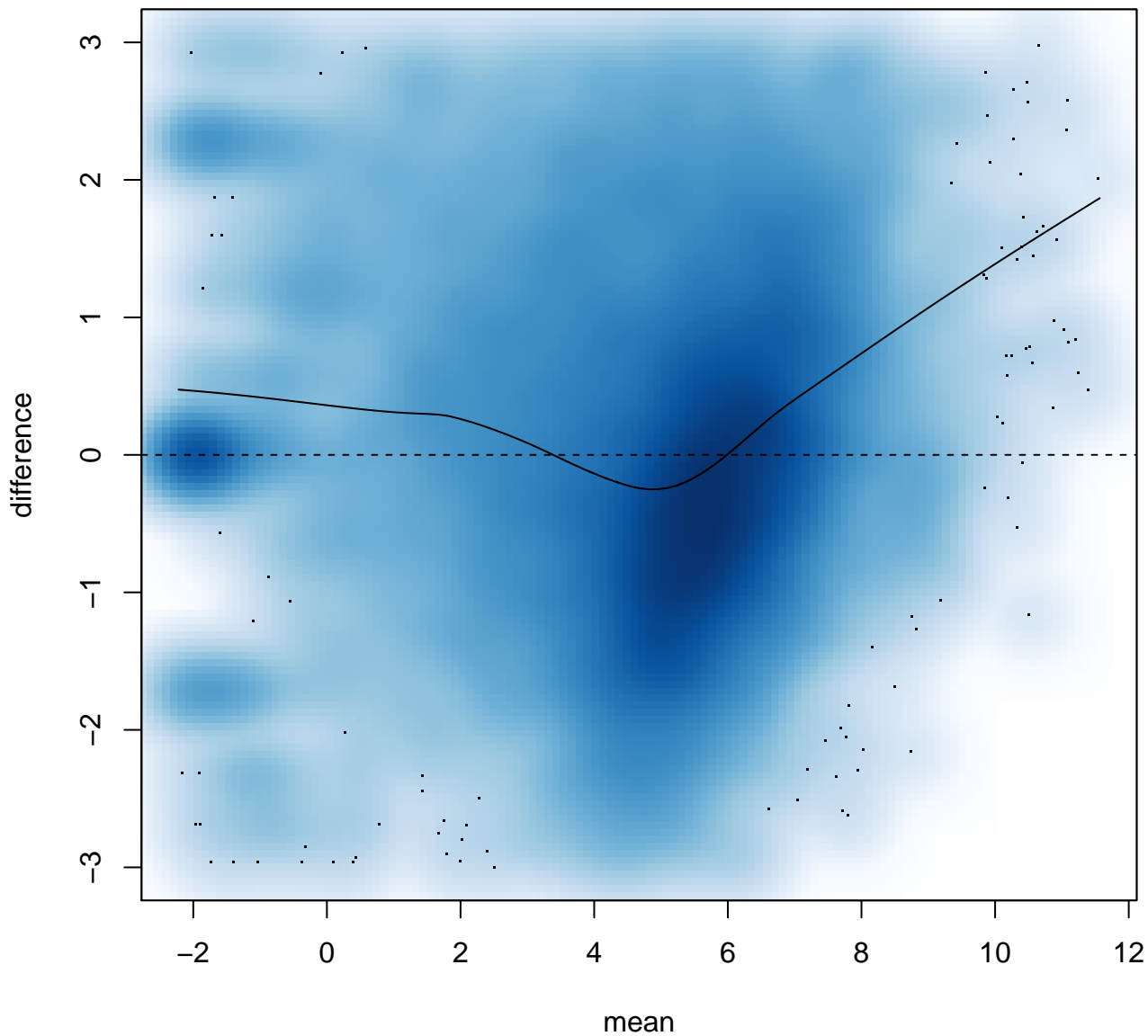
AF293_Normoxia.r1 vs AF293_Normoxia.r2



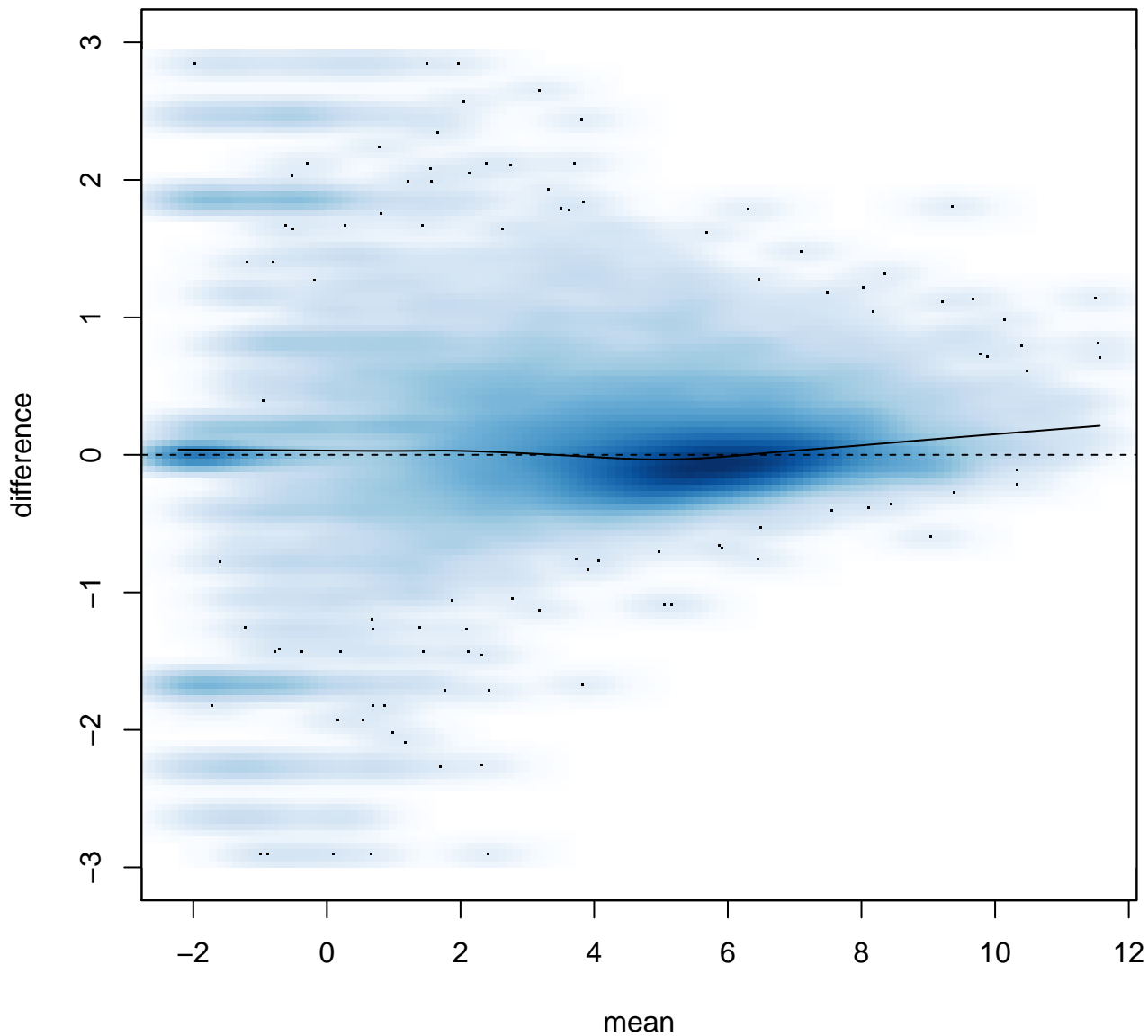
AF293_Normoxia.r1 vs AF293_Normoxia.r3



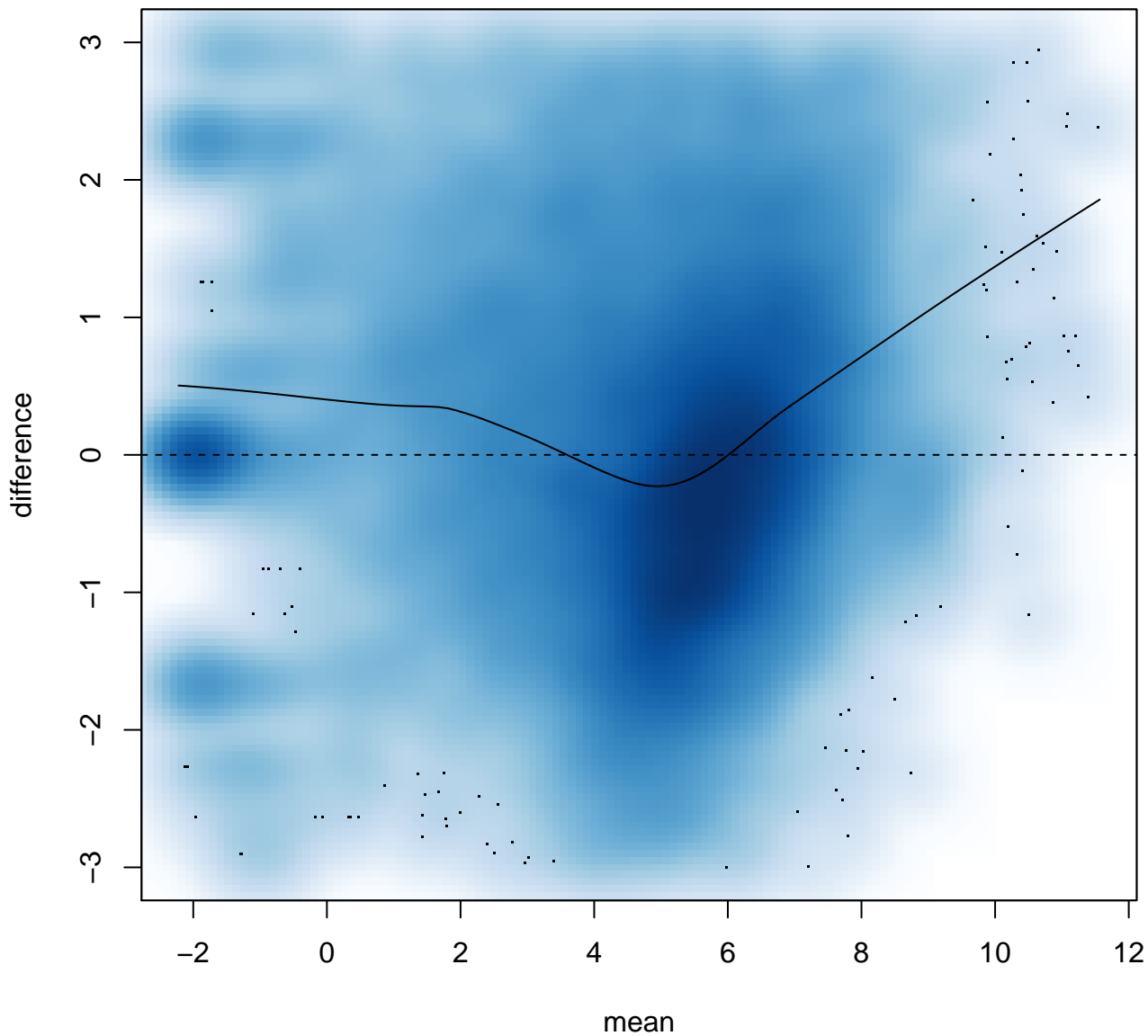
AF293_Normoxia.r1 vs AF293_Hypoxia.r1



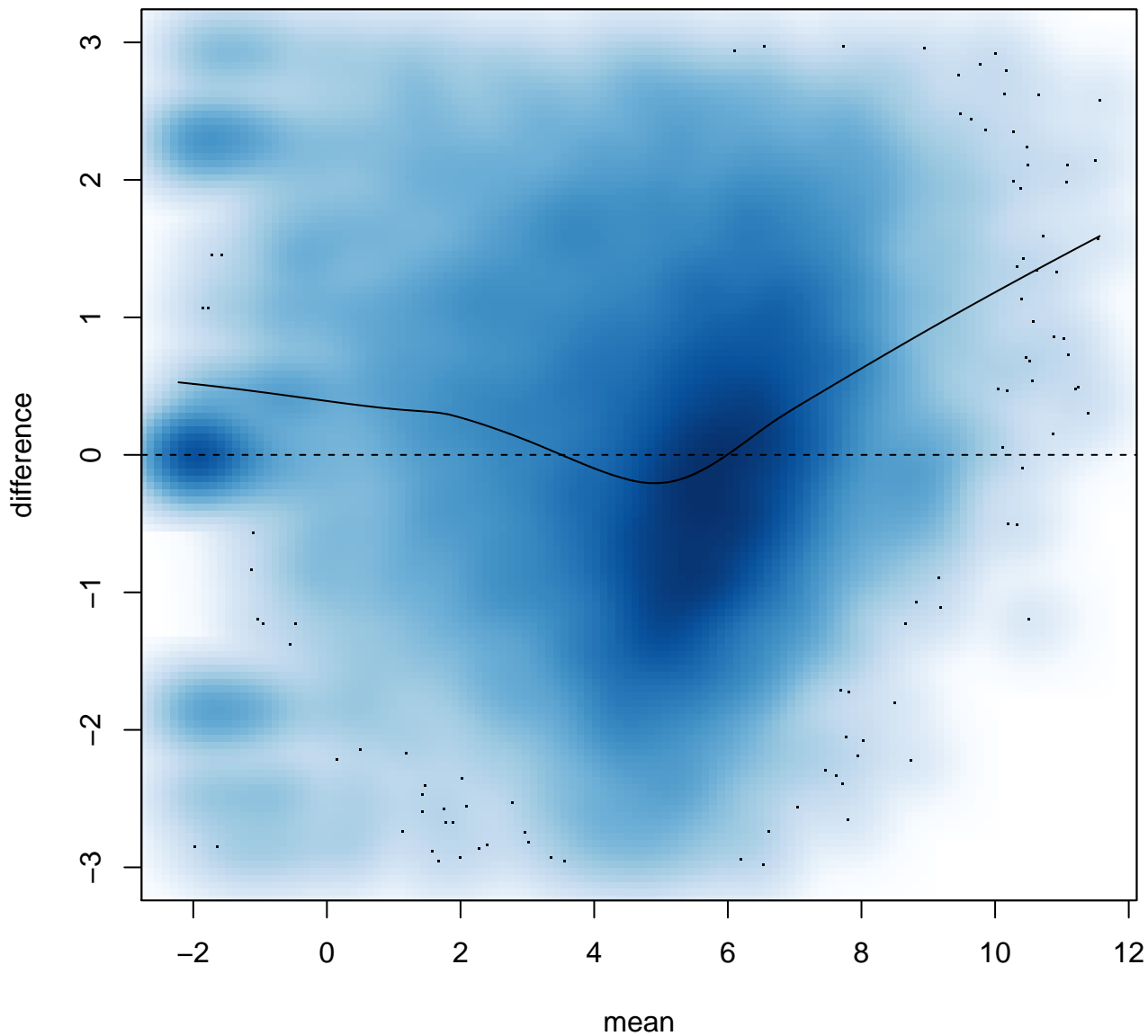
AF293_Normoxia.r2 vs AF293_Normoxia.r3

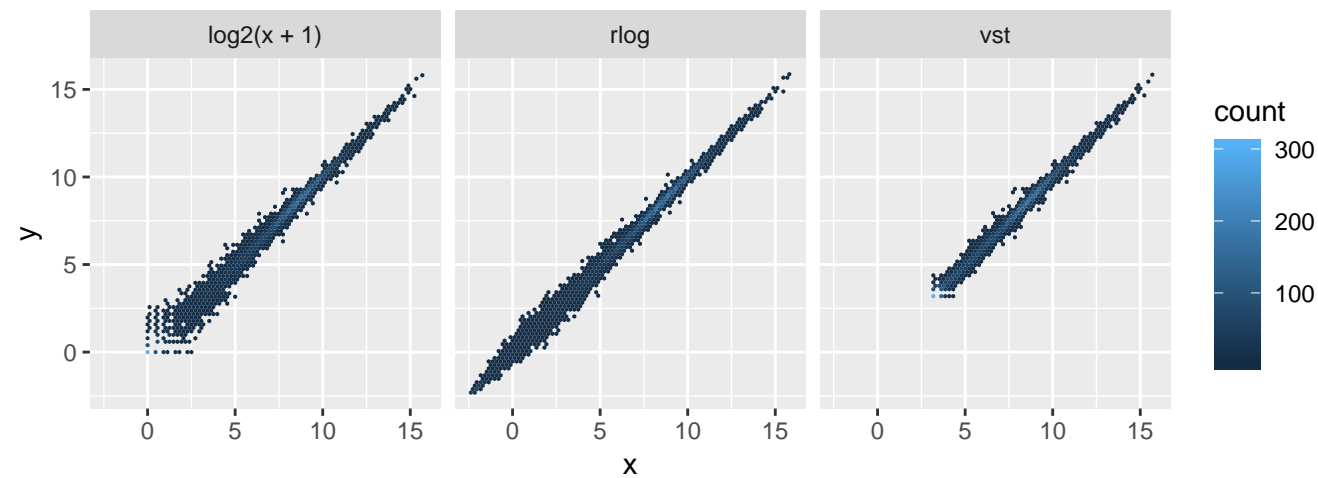


AF293_Normoxia.r2 vs AF293_Hypoxia.r1

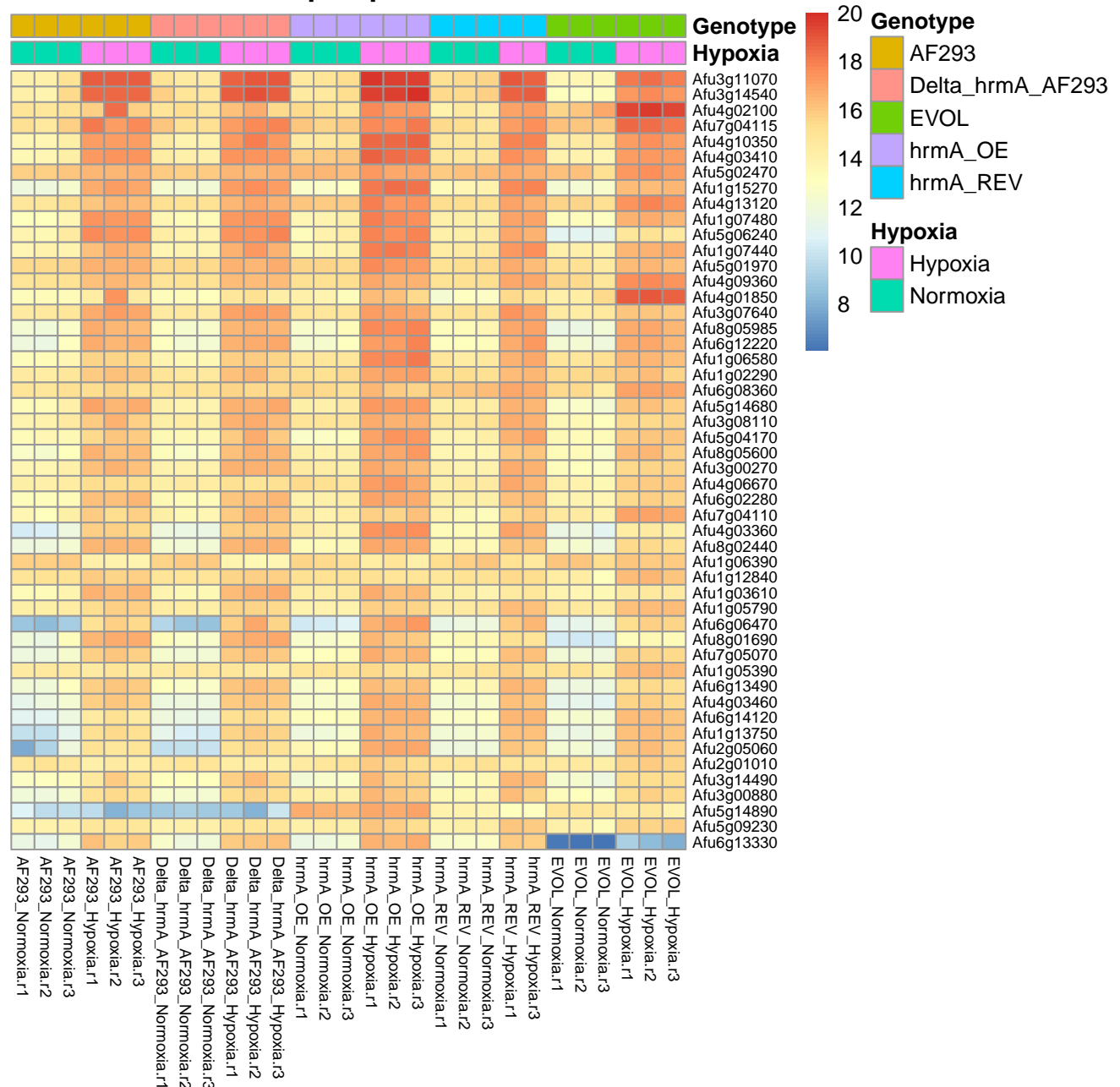


AF293_Normoxia.r3 vs AF293_Hypoxia.r1

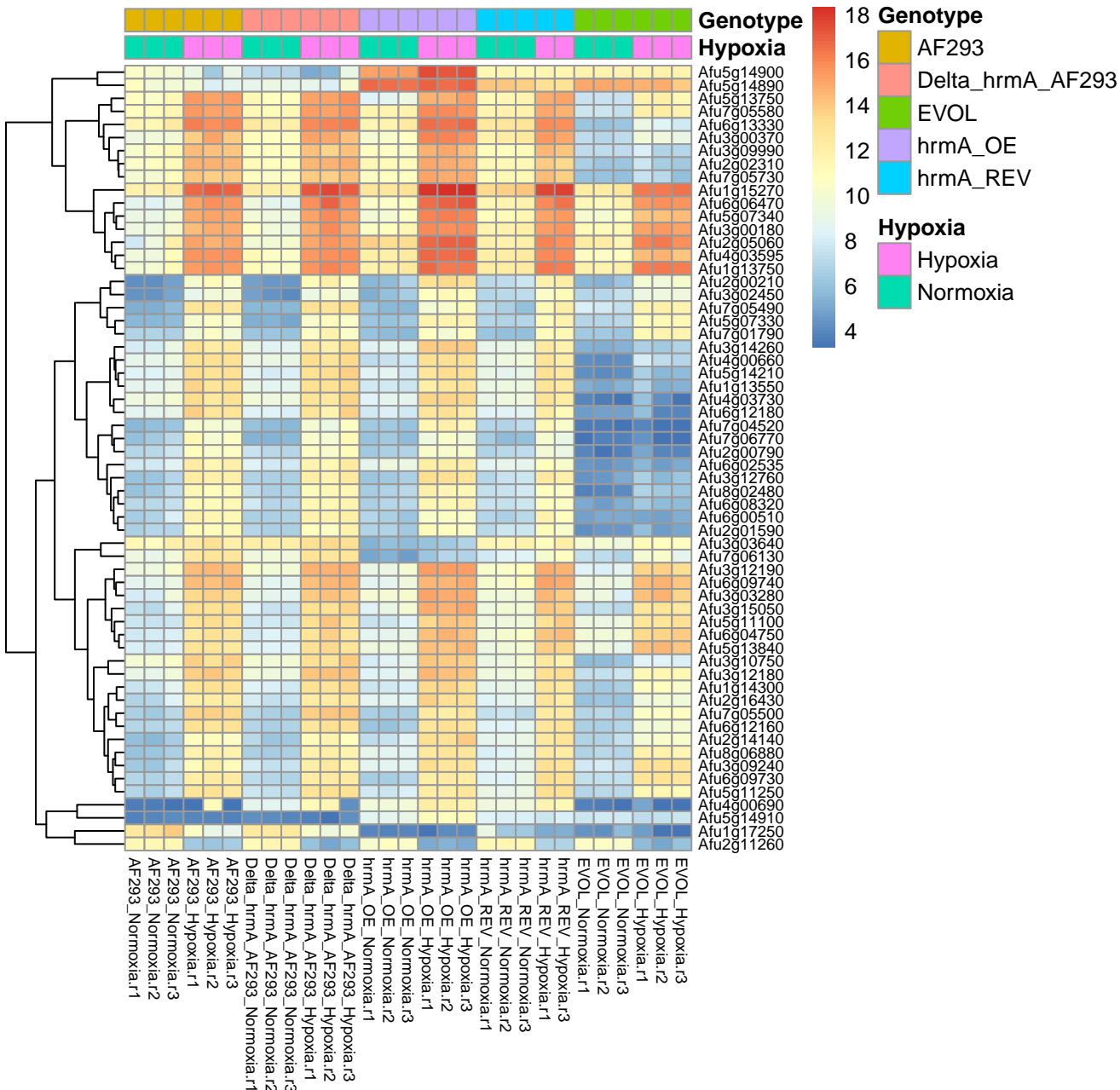




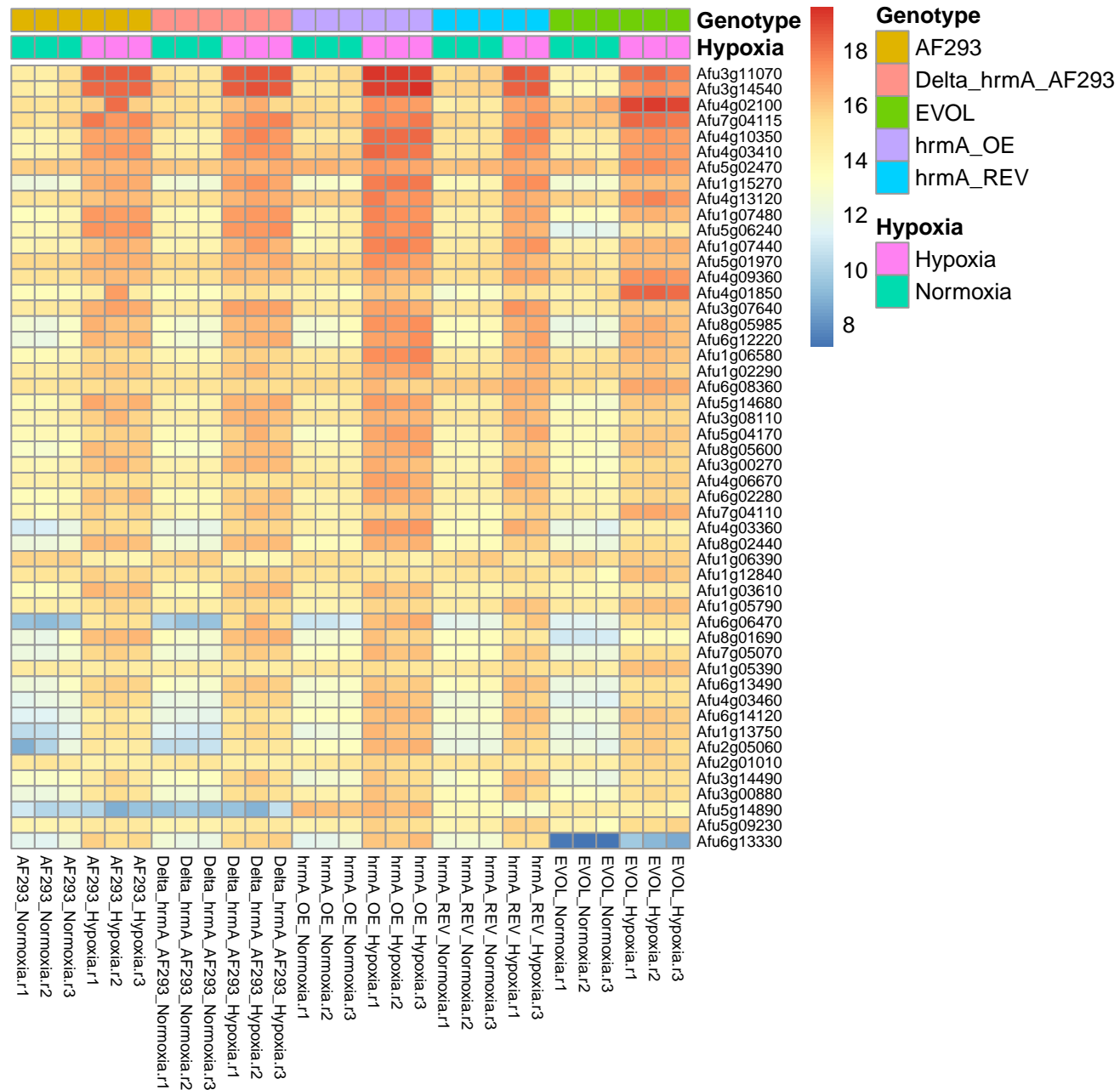
VSD Top Expression



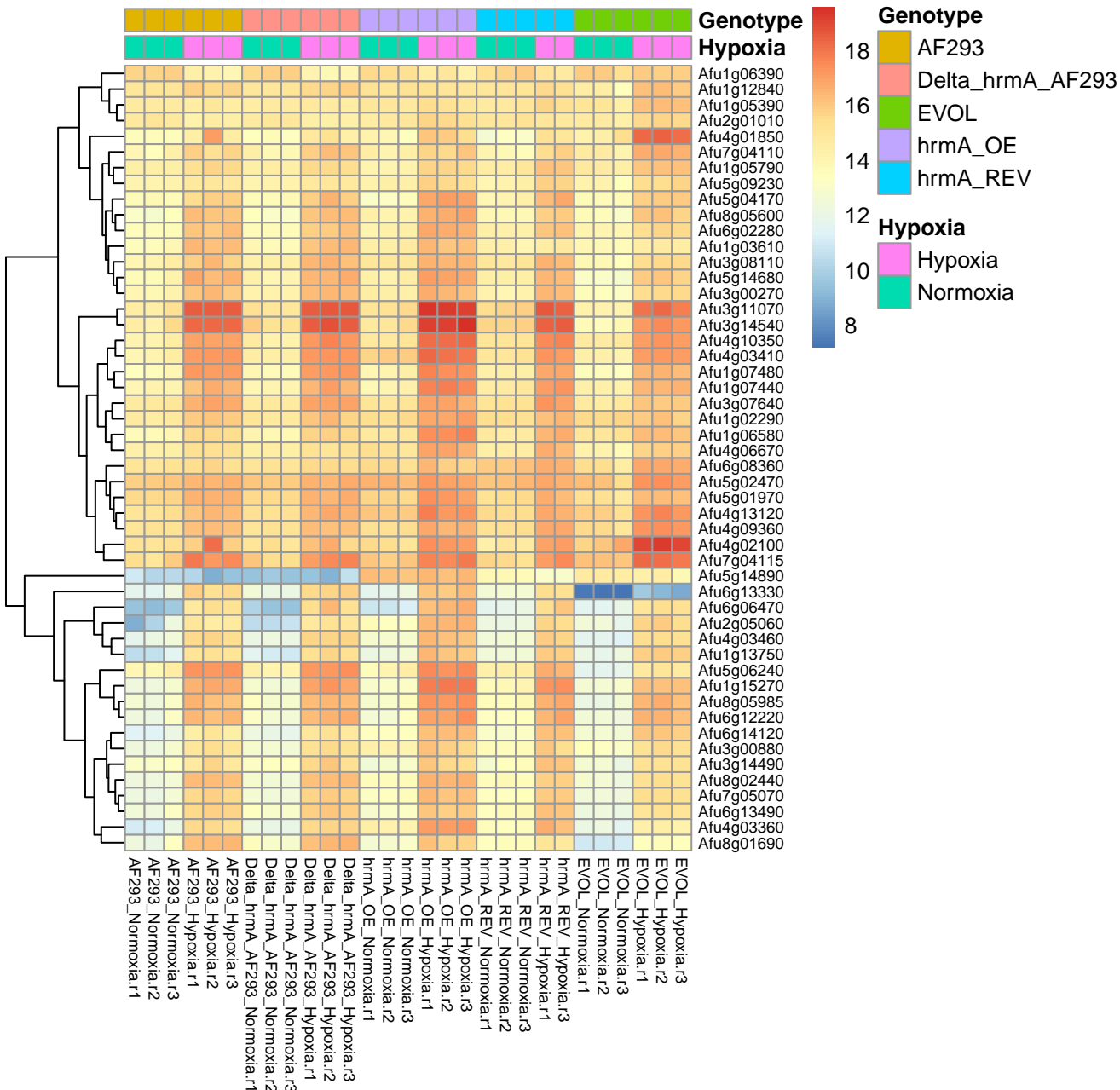
VSD Most different



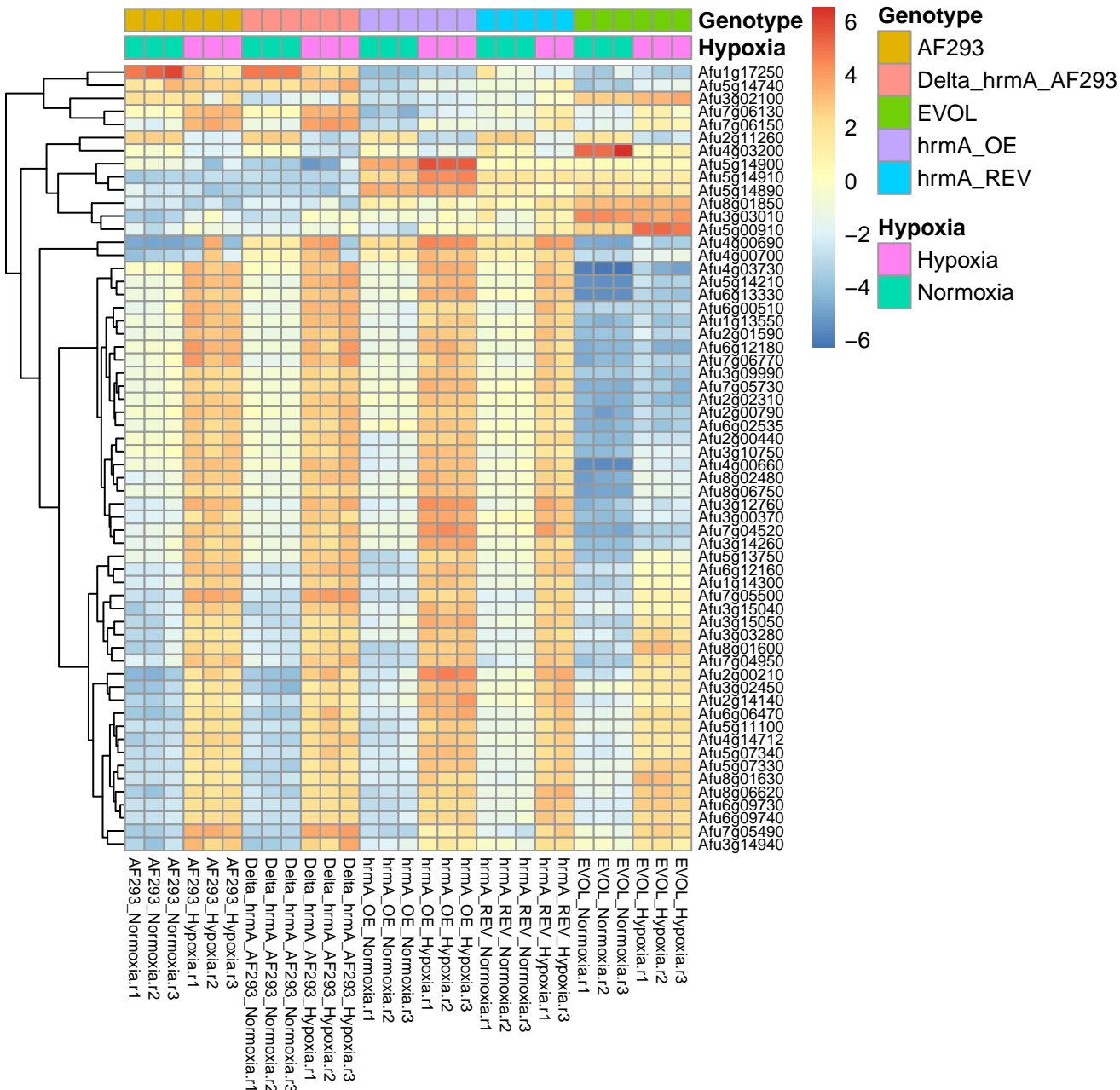
RLD Top Expression

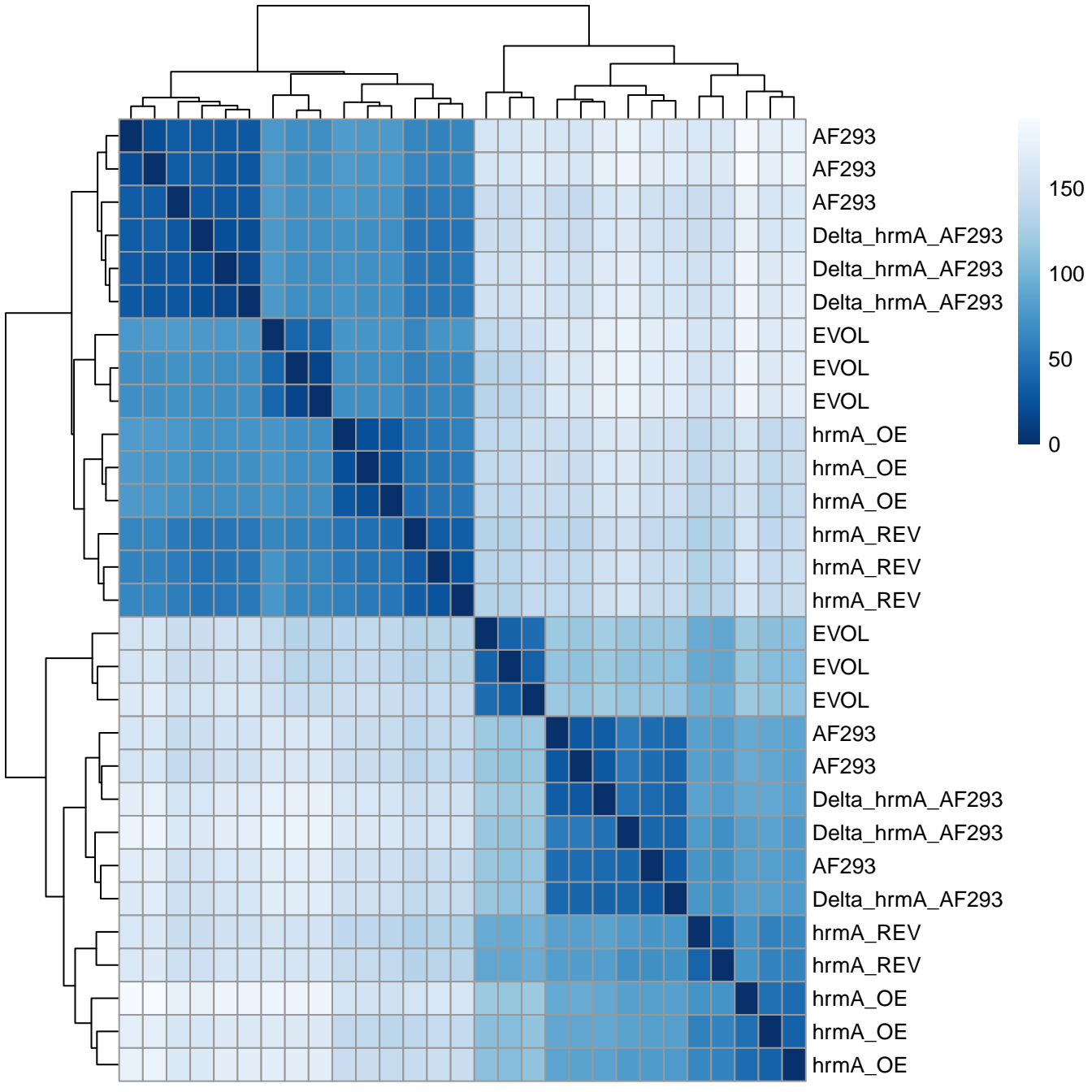


RLD Top Expression

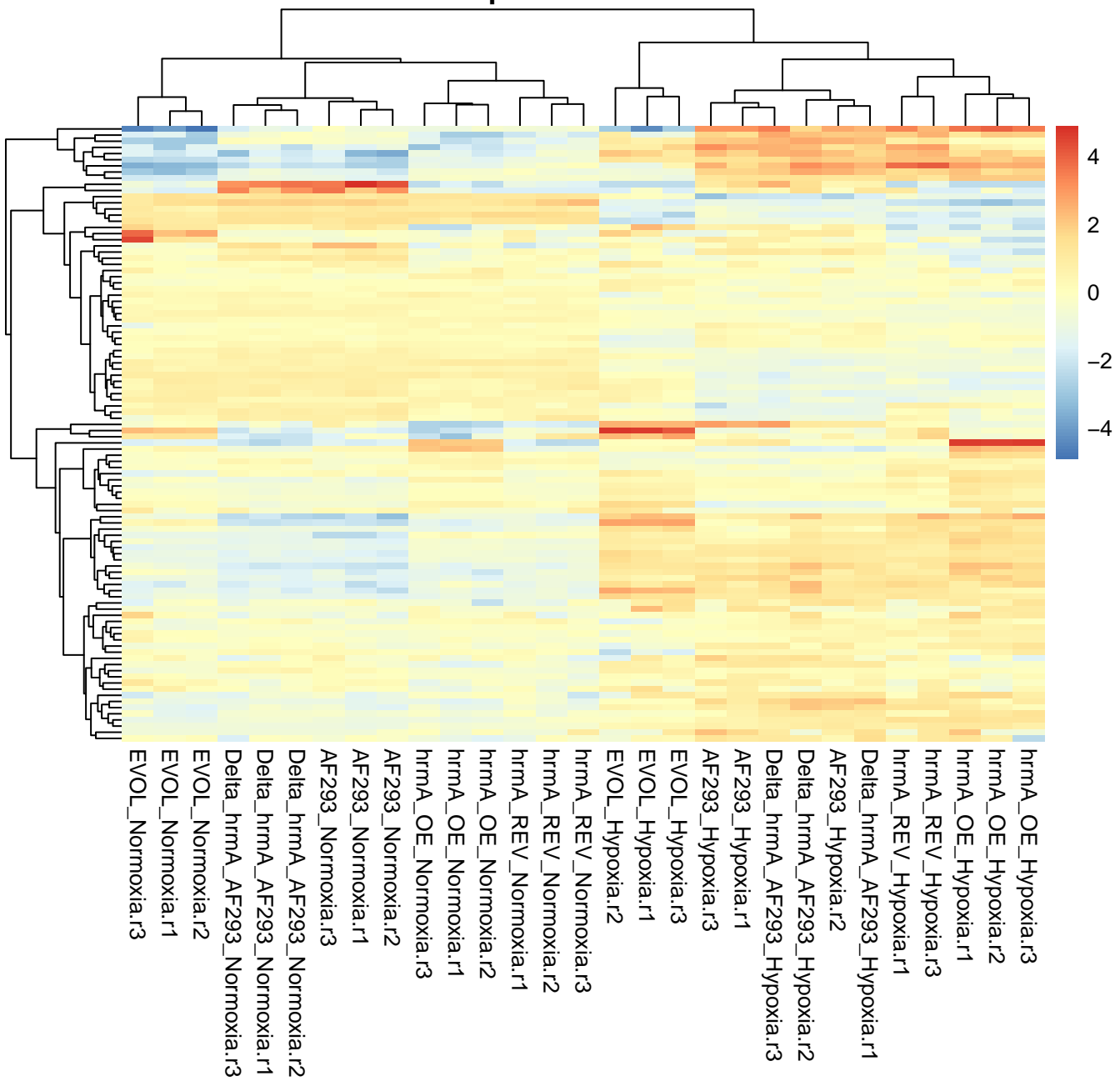


RLD Most different

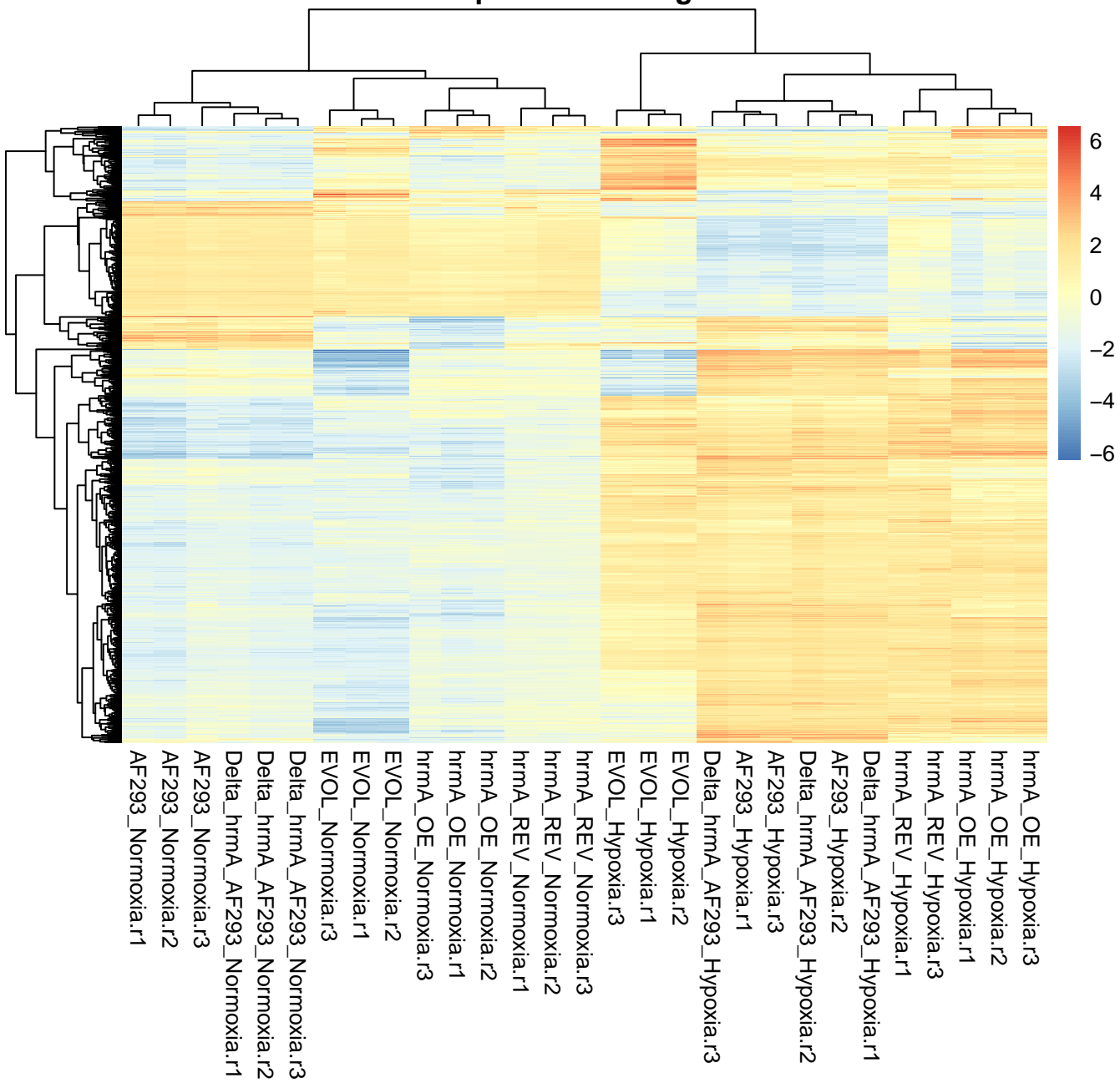


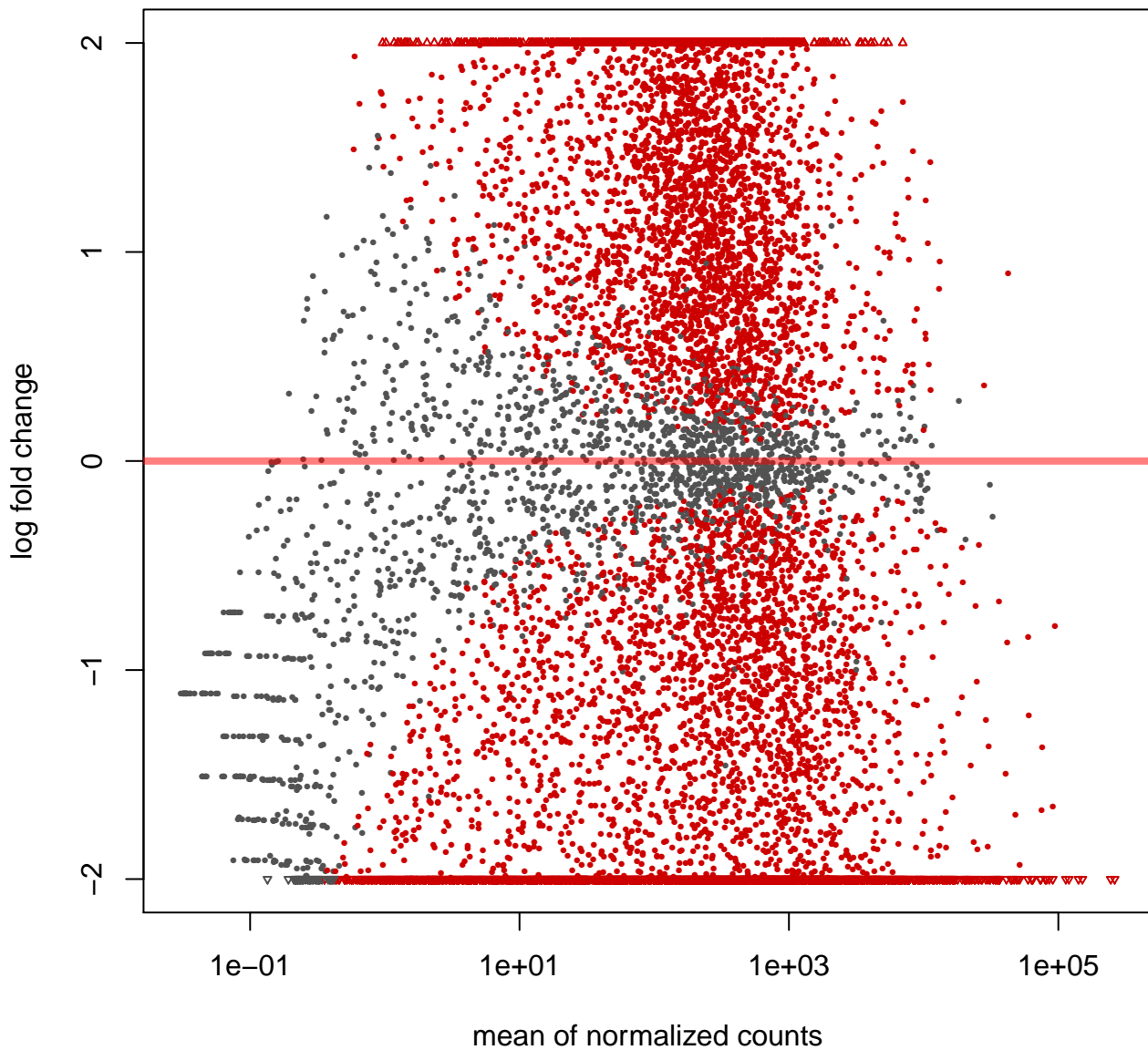


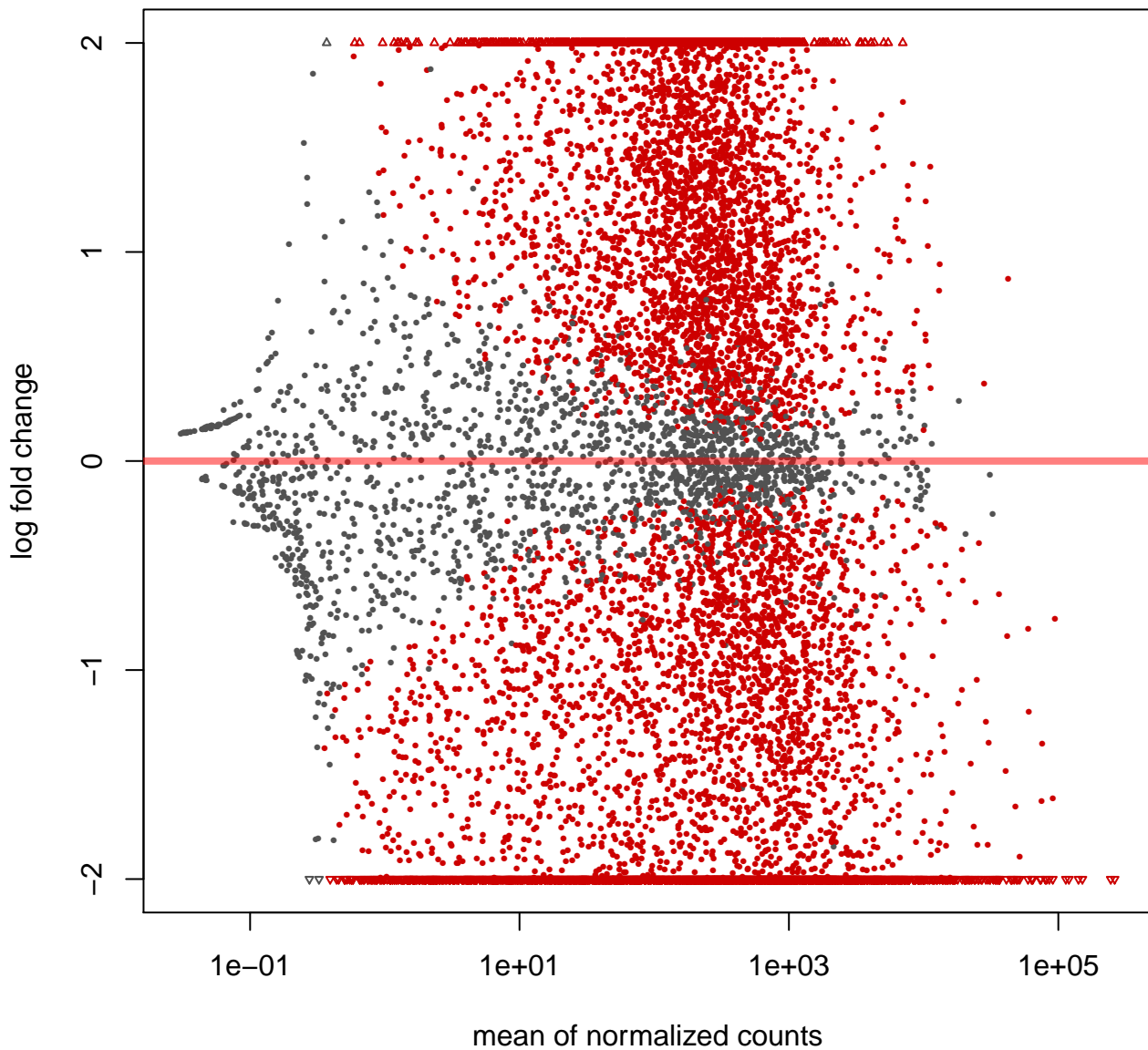
TopVar normalized



Unsupervised 1000 genes







Fold Change Plot Down

