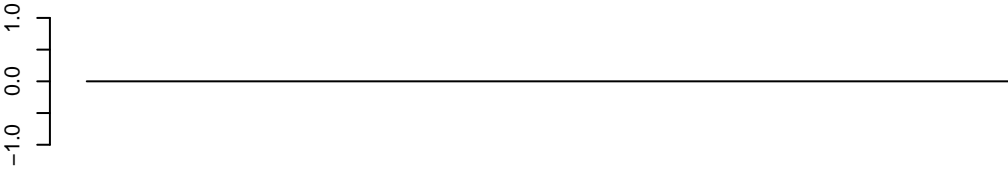
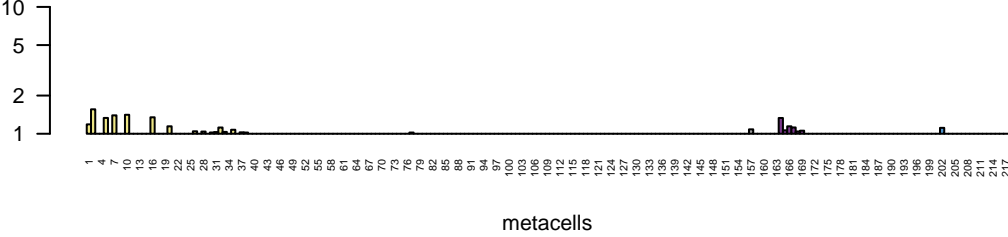


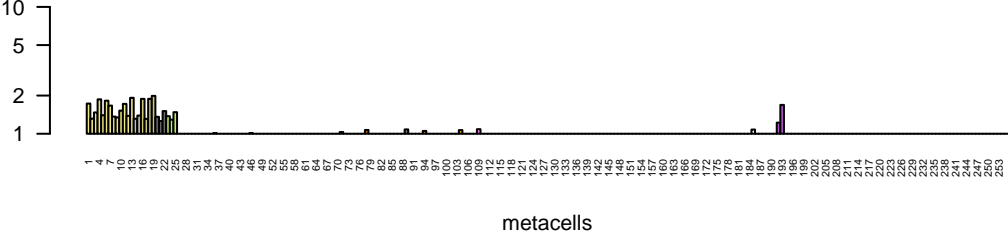
HLH.HG2.53:BHLHA15/BHLHE22/BHLHE23/OLIG1/OLIG2/OLIG3
Tadh | no data



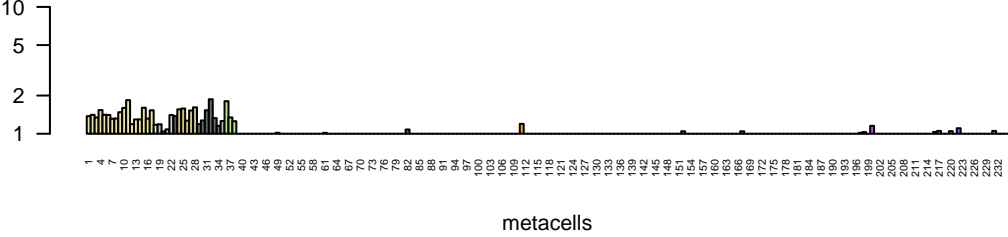
TrH2 OG_3824
TrH2_TrispH2_010486-RA
HLH.HG2.53:BHLHA15/BHLHE22/BHLHE23/OLIG1/OLIG2/OLIG3

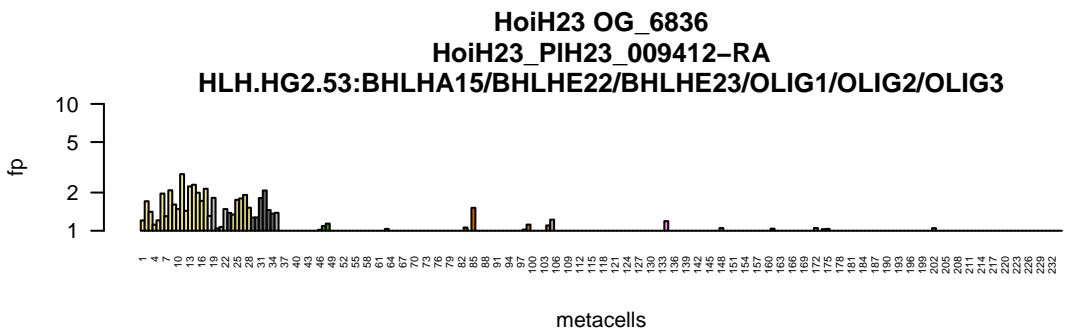
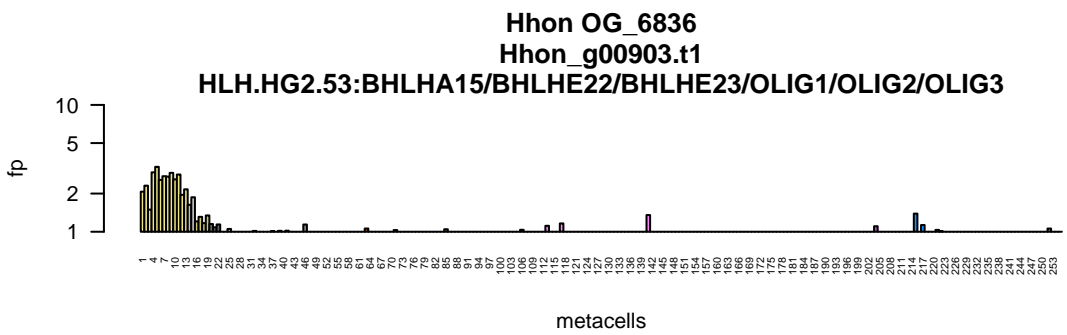
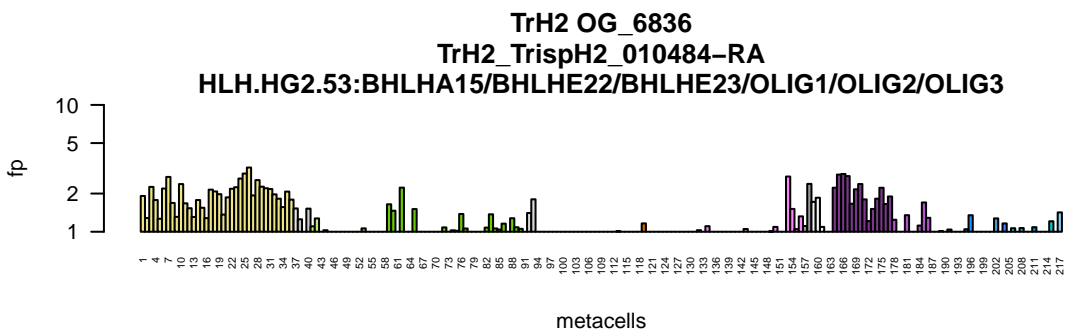
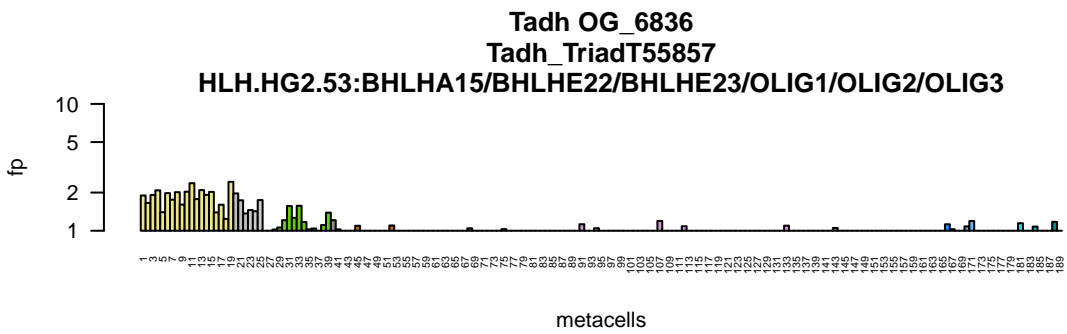


Hhon OG_3824
Hhon_g00902.t1
HLH.HG2.53:BHLHA15/BHLHE22/BHLHE23/OLIG1/OLIG2/OLIG3

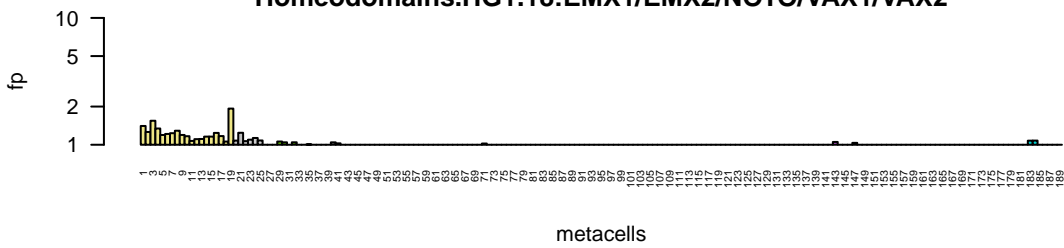


HoiH23 OG_3824
HoiH23_PIH23_009413-RA
HLH.HG2.53:BHLHA15/BHLHE22/BHLHE23/OLIG1/OLIG2/OLIG3

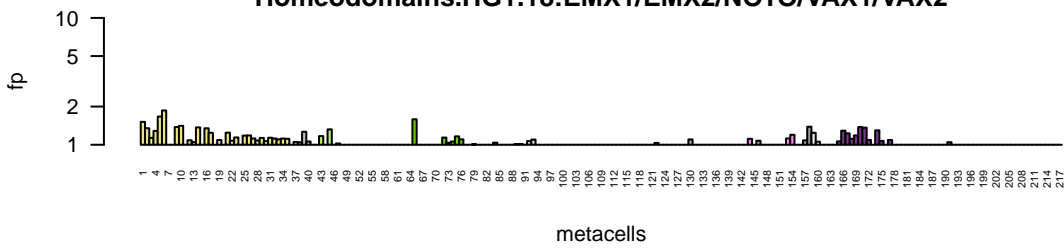




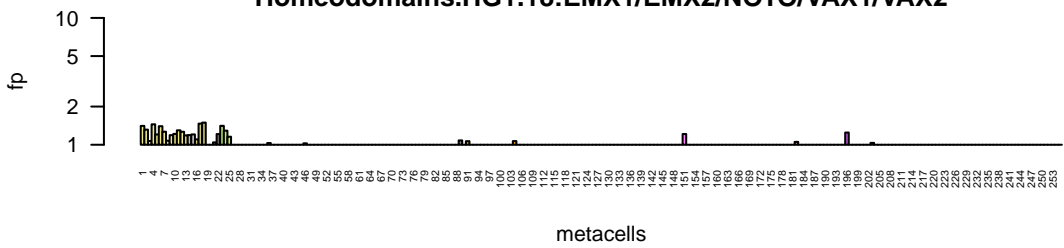
Tadh OG_5112
Tadh_TriadT49524
Homeodomains.HG1.18:EMX1/EMX2/NOTO/VAX1/VAX2



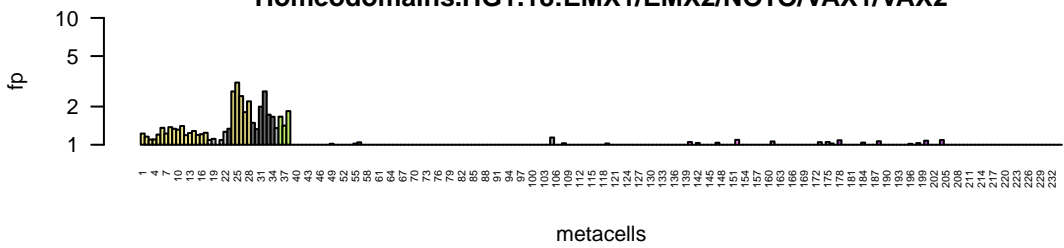
TrH2 OG_5112
TrH2_TrispH2_003519-RA
Homeodomains.HG1.18:EMX1/EMX2/NOTO/VAX1/VAX2



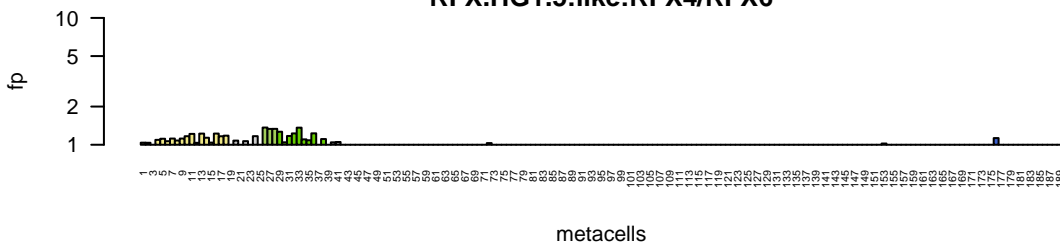
Hhon OG_5112
Hhon_g10555.t1
Homeodomains.HG1.18:EMX1/EMX2/NOTO/VAX1/VAX2



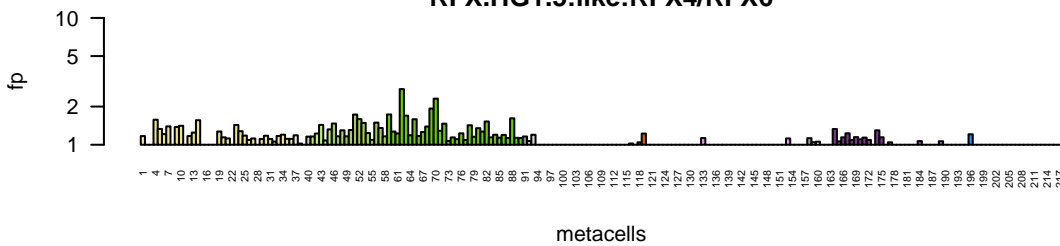
HoiH23 OG_5112
HoiH23_PIH23_007693-RA
Homeodomains.HG1.18:EMX1/EMX2/NOTO/VAX1/VAX2



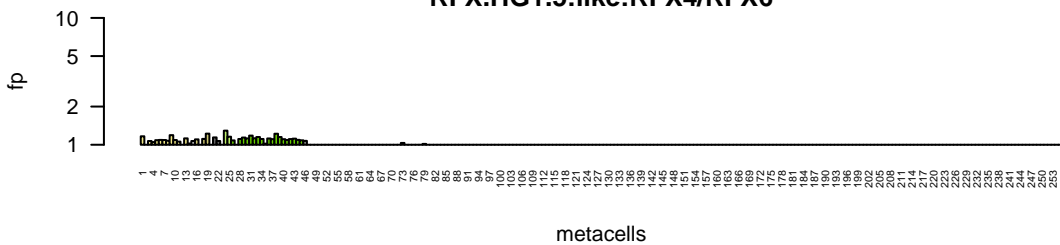
Tadh OG_9882
Tadh_TriadT51266
RFX.HG1.5:like:RFX4/RFX6



TrH2 OG_9882
TrH2_TrispH2_008141-RA
RFX.HG1.5:like:RFX4/RFX6



Hhon OG_9882
Hhon_g00515.t1
RFX.HG1.5:like:RFX4/RFX6



HoiH23 OG_9882
HoiH23_PIH23_008508-RA
RFX.HG1.5:like:RFX4/RFX6

