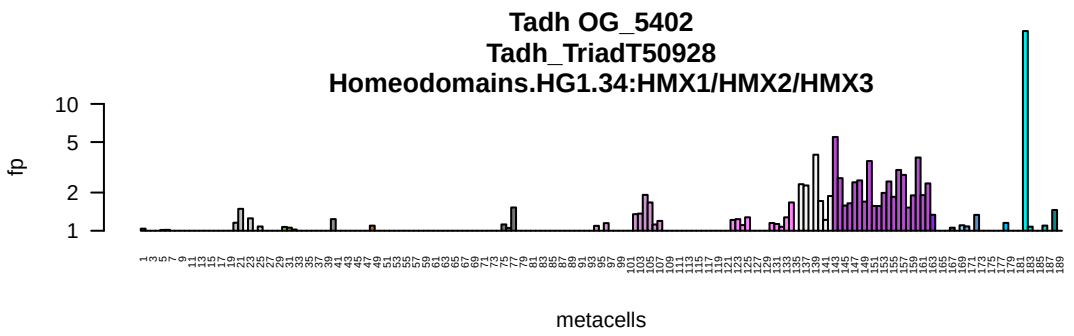
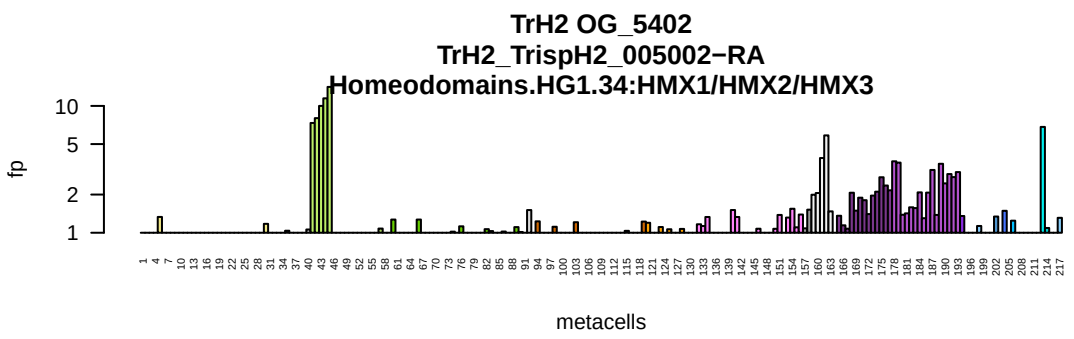
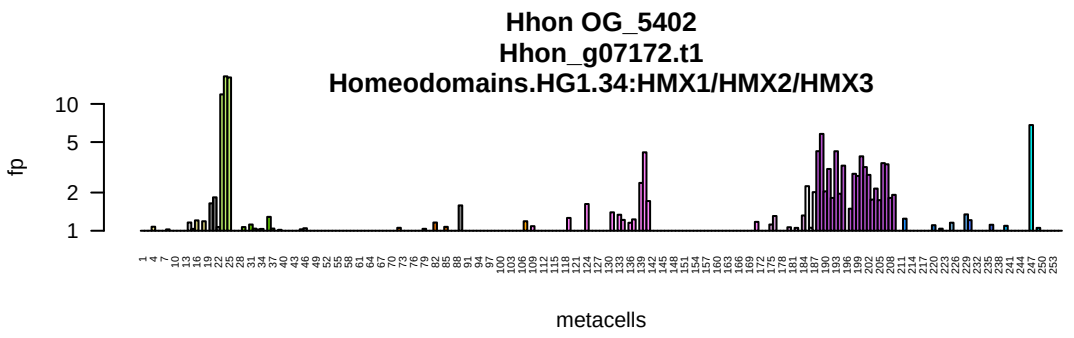
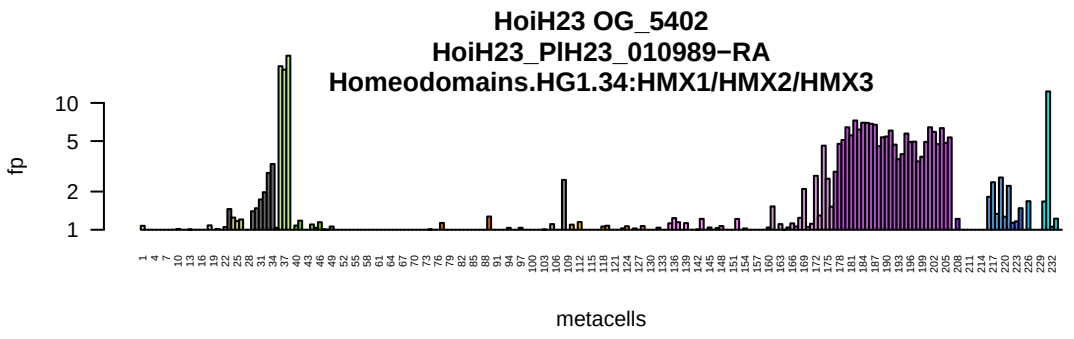
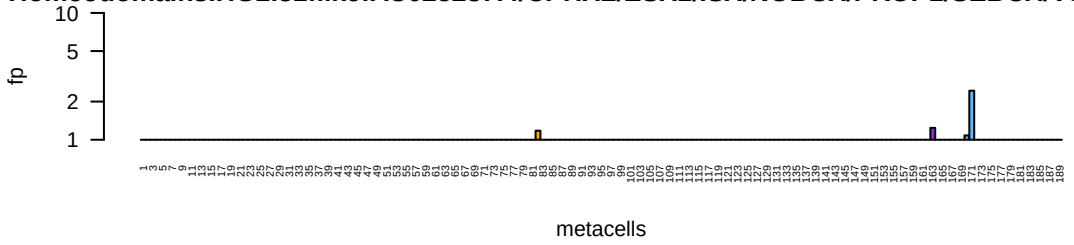

Tadh OG_5402
Tadh_TriadT50928
Homeodomains.HG1.34:HMX1/HMX2/HMX3
fp
metacells


TrH2 OG_5402
TrH2_TrispH2_005002−RA
Homeodomains.HG1.34:HMX1/HMX2/HMX3
fp
metacells

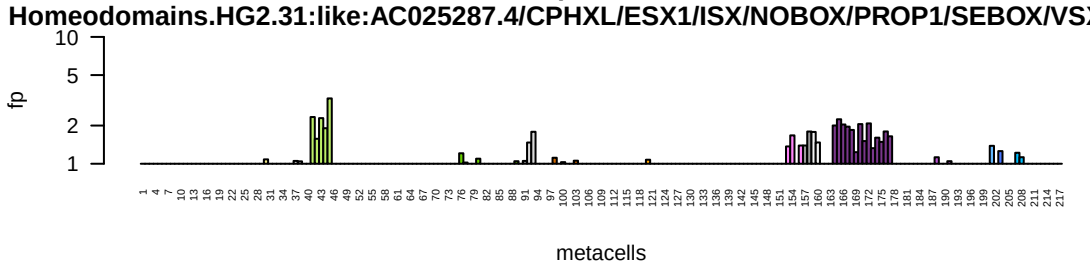

Hhon OG_5402
Hhon_g07172.t1
Homeodomains.HG1.34:HMX1/HMX2/HMX3
fp
metacells


HoiH23 OG_5402
HoiH23_PlH23_010989−RA
Homeodomains.HG1.34:HMX1/HMX2/HMX3
fp
metacells

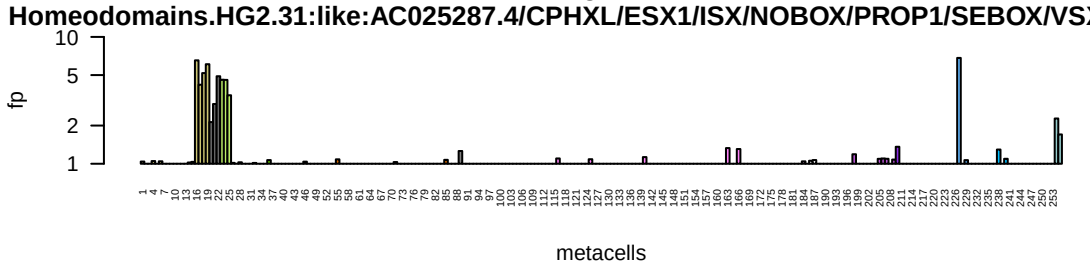
Tadh OG_5876
Tadh_TriadT57021
Homeodomains.HG2.31:like:AC025287.4/CPHXL/ESX1/ISX/NOBOX/PROP1/SEBOX/VSX1/VS



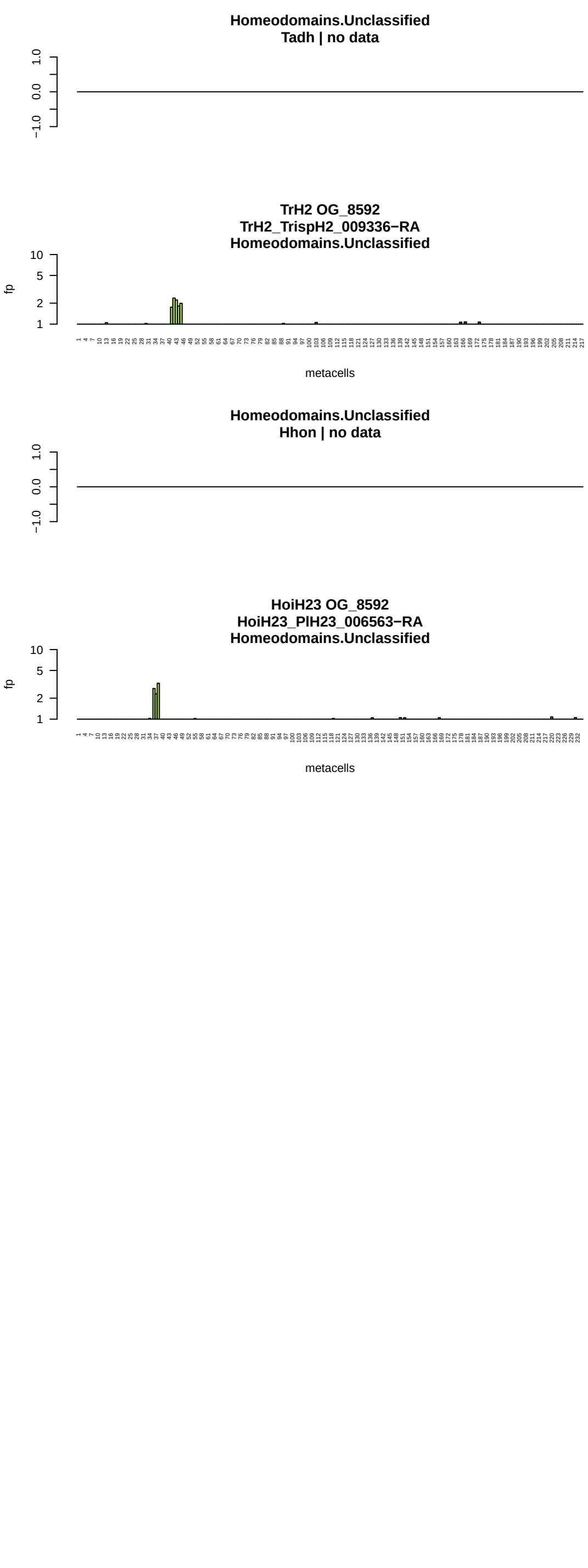
TrH2 OG_5876
TrH2_TrispH2_002361−RA
Homeodomains.HG2.31:like:AC025287.4/CPHXL/ESX1/ISX/NOBOX/PROP1/SEBOX/VSX1/VS



Hhon OG_5876
Hhon_g07457.t1
Homeodomains.HG2.31:like:AC025287.4/CPHXL/ESX1/ISX/NOBOX/PROP1/SEBOX/VSX1/VS



HoiH23 OG_5876
HoiH23_PlH23_002938−RA
Homeodomains.HG2.31:like:AC025287.4/CPHXL/ESX1/ISX/NOBOX/PROP1/SEBOX/VSX1/VS


Homeodomains.Unclassified
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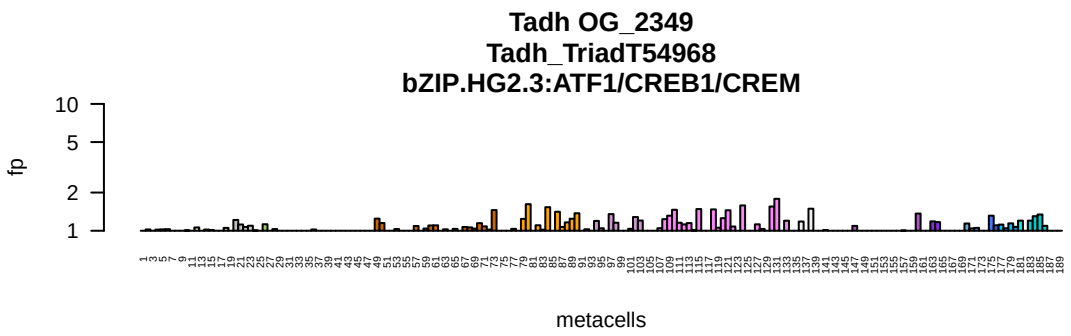
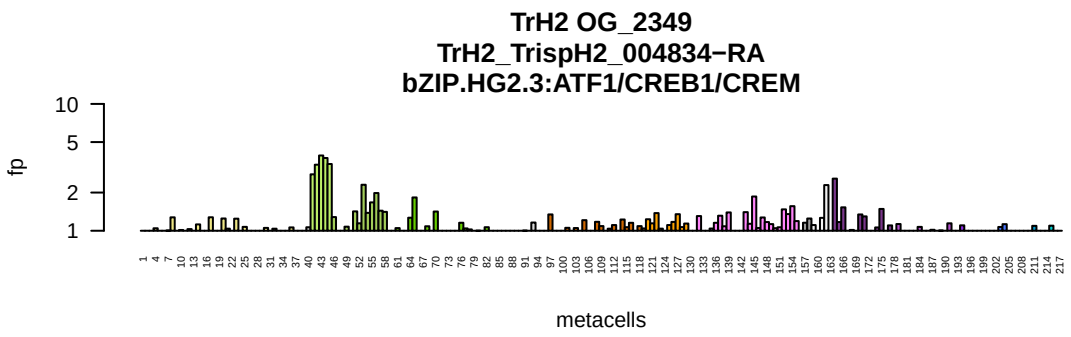
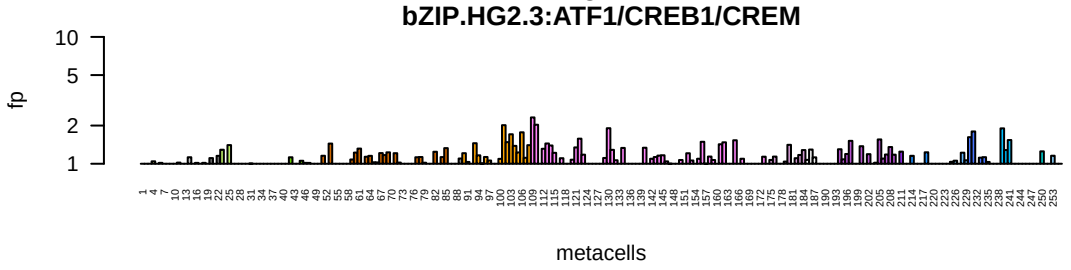
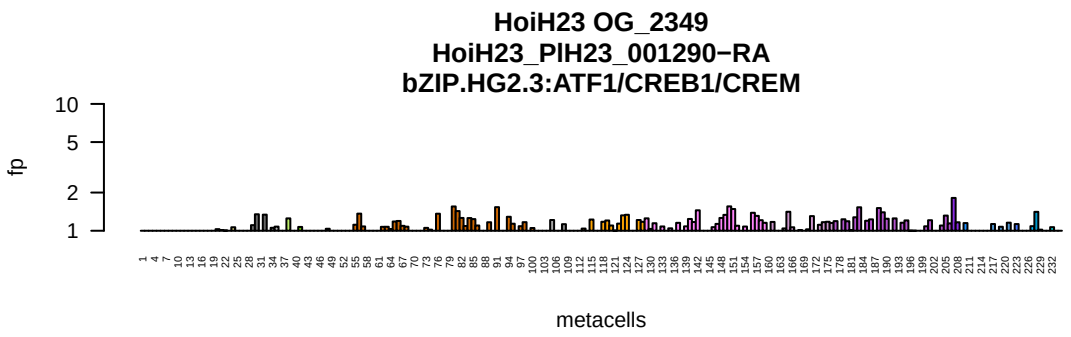
−1.0
TrH2 OG_8592
TrH2_TrispH2_009336−RA
Homeodomains.Unclassified
fp
10
5
2
1
metacells
Homeodomains.Unclassified
Hhon | no data
1.0
0.0
−1.0
HoiH23 OG_8592
HoiH23_PlH23_006563−RA
Homeodomains.Unclassified
fp
10
5
2
1
metacells

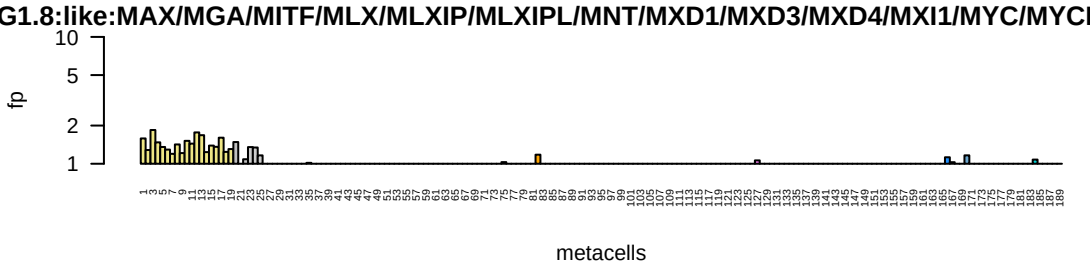

Tadh OG_2349
Tadh_TriadT54968
bZIP.HG2.3:ATF1/CREB1/CREM
fp
metacells


TrH2 OG_2349
TrH2_TrispH2_004834−RA
bZIP.HG2.3:ATF1/CREB1/CREM
fp
metacells

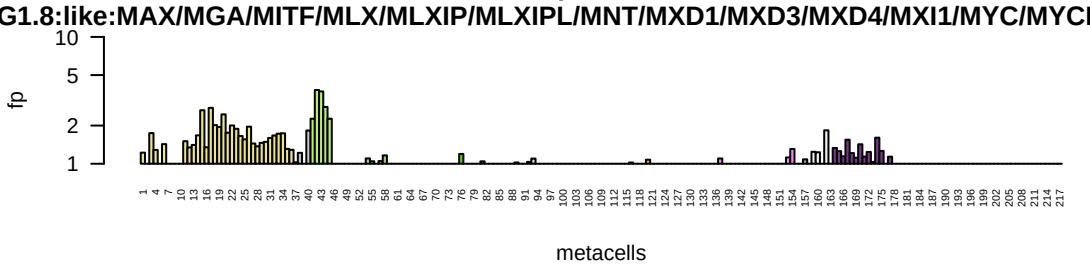

Hhon OG_2349
Hhon_g08642.t1
bZIP.HG2.3:ATF1/CREB1/CREM
fp
metacells


HoiH23 OG_2349
HoiH23_PlH23_001290−RA
bZIP.HG2.3:ATF1/CREB1/CREM
fp
metacells

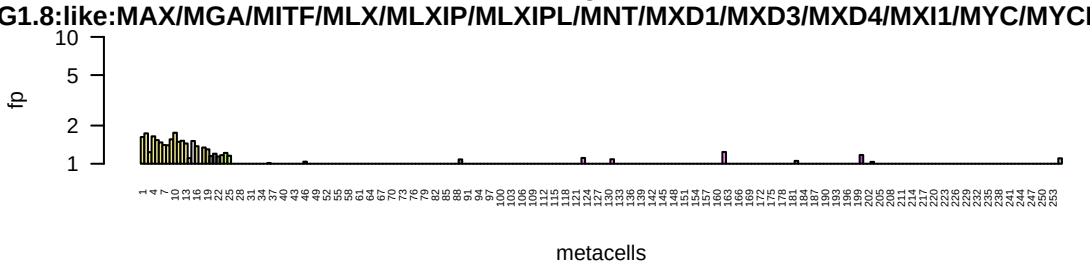
Tadh OG_3756
Tadh_TriadT56276
G1.8:like:MAX/MGA/MITF/MLX/MLXIP/MLXIPL/MNT/MXD1/MXD3/MXD4/MXI1/MYC/MYCL/MY



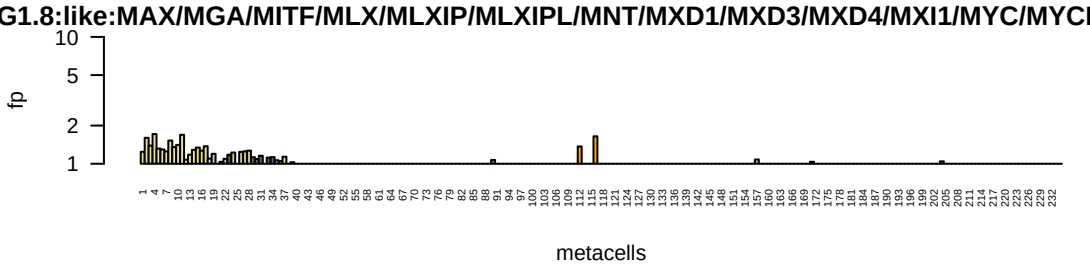
TrH2 OG_3756
TrH2_TrispH2_007744−RA
G1.8:like:MAX/MGA/MITF/MLX/MLXIP/MLXIPL/MNT/MXD1/MXD3/MXD4/MXI1/MYC/MYCL/MY

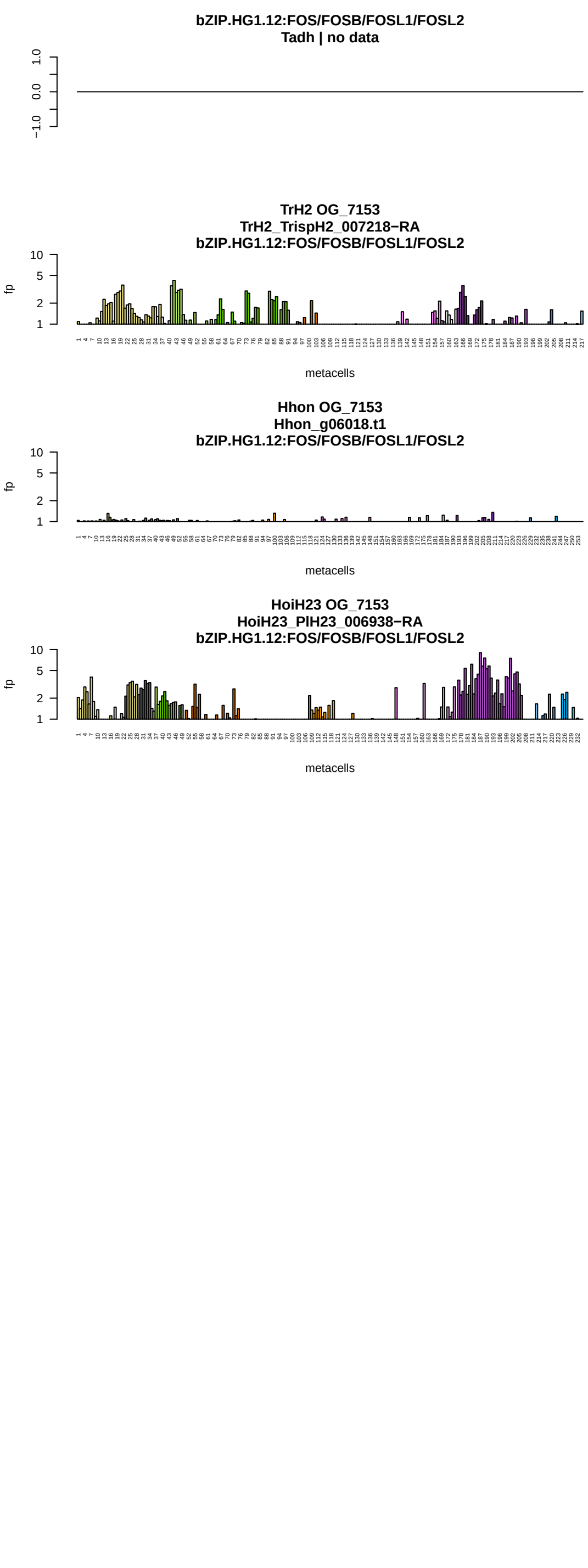


Hhon OG_3756
Hhon_g02951.t1
G1.8:like:MAX/MGA/MITF/MLX/MLXIP/MLXIPL/MNT/MXD1/MXD3/MXD4/MXI1/MYC/MYCL/MY



HoiH23 OG_3756
HoiH23_PlH23_010554−RA
G1.8:like:MAX/MGA/MITF/MLX/MLXIP/MLXIPL/MNT/MXD1/MXD3/MXD4/MXI1/MYC/MYCL/MY




bZIP.HG1.12:FOS/FOSB/FOSL1/FOSL2
Tadh | no data
1.0
0.0
−1.0
TrH2 OG_7153
TrH2_TrispH2_007218−RA
bZIP.HG1.12:FOS/FOSB/FOSL1/FOSL2
fp
metacells
Hhon OG_7153
Hhon_g06018.t1
bZIP.HG1.12:FOS/FOSB/FOSL1/FOSL2
fp
metacells
HoiH23 OG_7153
HoiH23_PlH23_006938−RA
bZIP.HG1.12:FOS/FOSB/FOSL1/FOSL2
fp
metacells