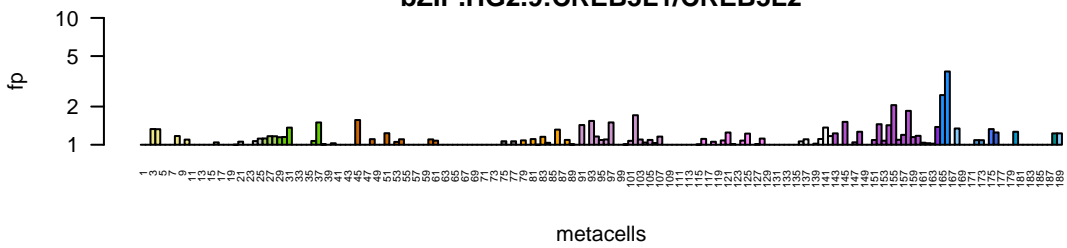
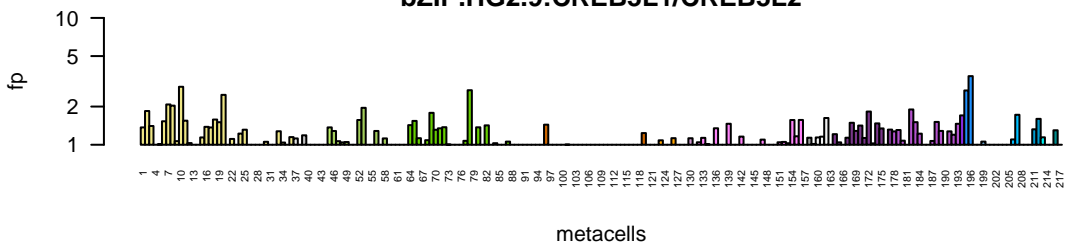


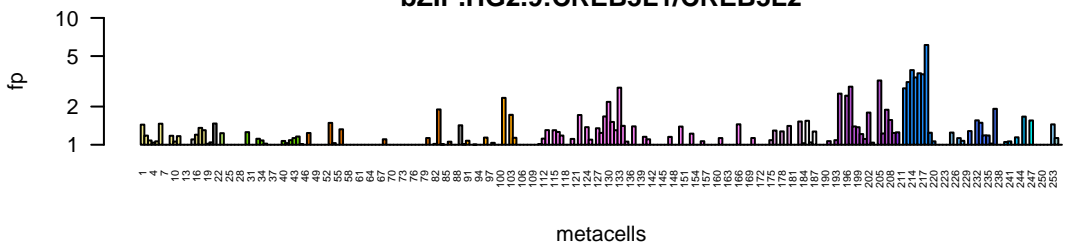
Tadh OG_1314
Tadh_TriadT54788
bZIP.HG2.9:CREB3L1/CREB3L2



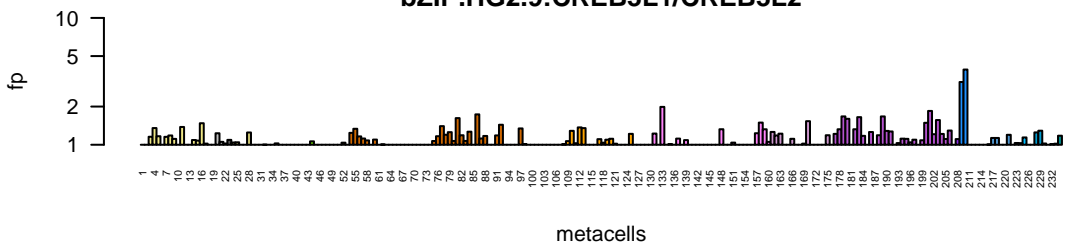
TrH2 OG_1314
TrH2_TrispH2_002162-RA
bZIP.HG2.9:CREB3L1/CREB3L2



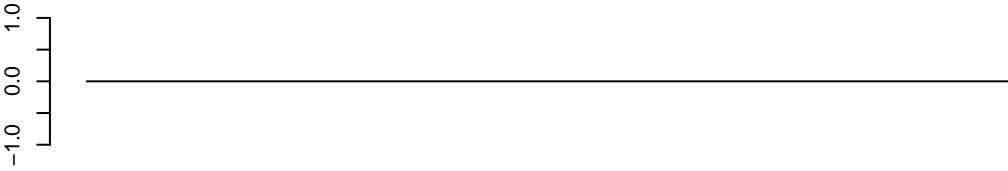
Hhon OG_1314
Hhon_g04701.t1
bZIP.HG2.9:CREB3L1/CREB3L2



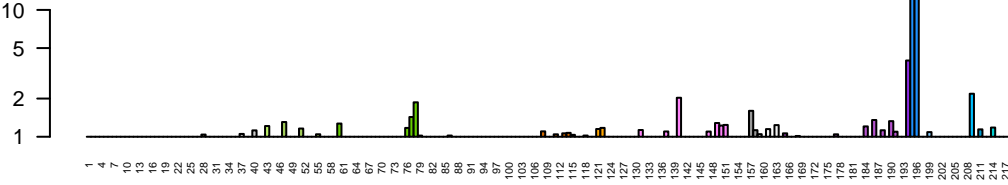
HoiH23 OG_1314
HoiH23_PIH23_002144-RA
bZIP.HG2.9:CREB3L1/CREB3L2



Homeodomains.HG1.4:DBX1/DBX2/HLX
Tadh | no data

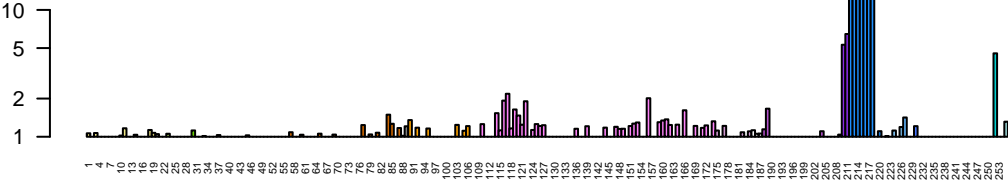


TrH2 OG_6496
TrH2_TrispH2_005279-RA
Homeodomains.HG1.4:DBX1/DBX2/HLX



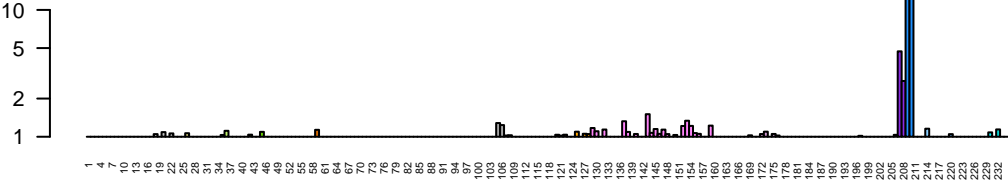
metacells

Hhon OG_6496
Hhon_g08608.t1
Homeodomains.HG1.4:DBX1/DBX2/HLX

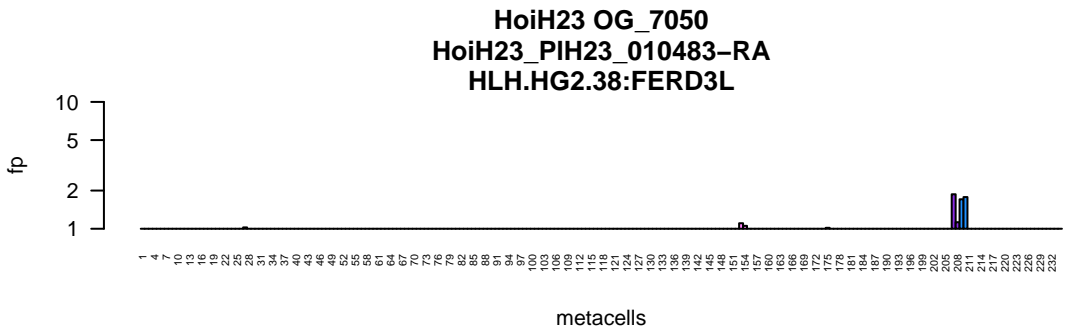
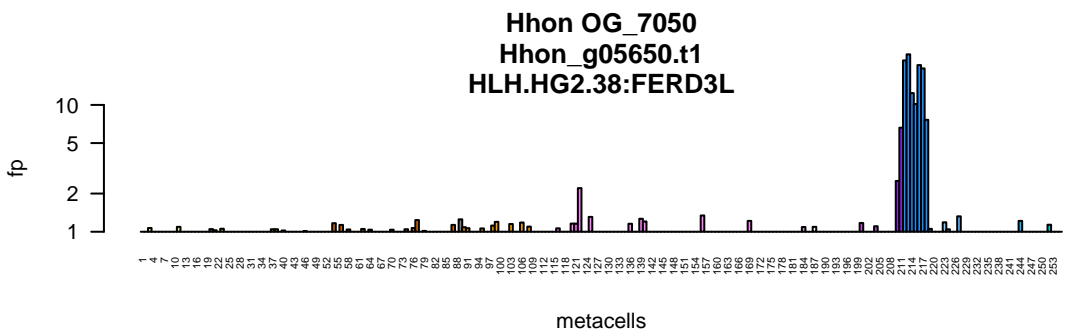
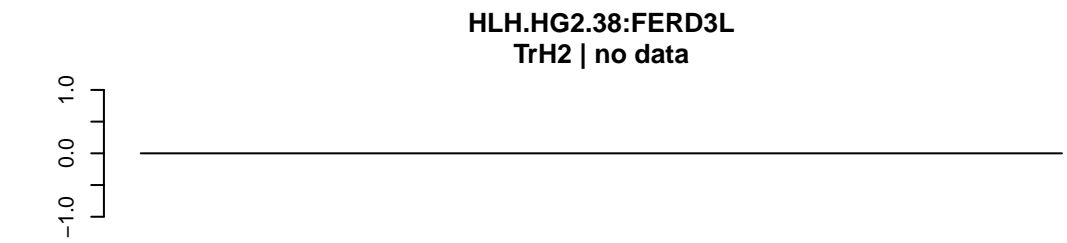
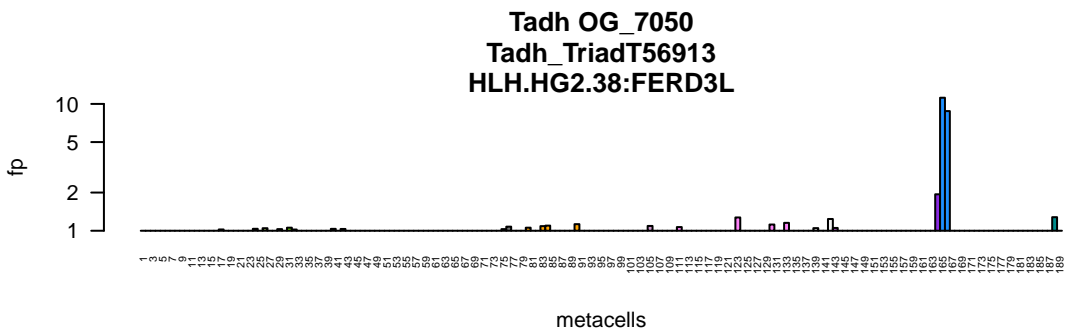


metacells

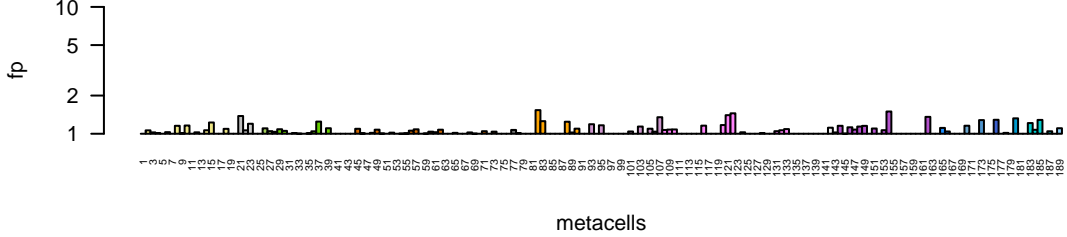
HoiH23 OG_6496
HoiH23_PIH23_004726-RA
Homeodomains.HG1.4:DBX1/DBX2/HLX



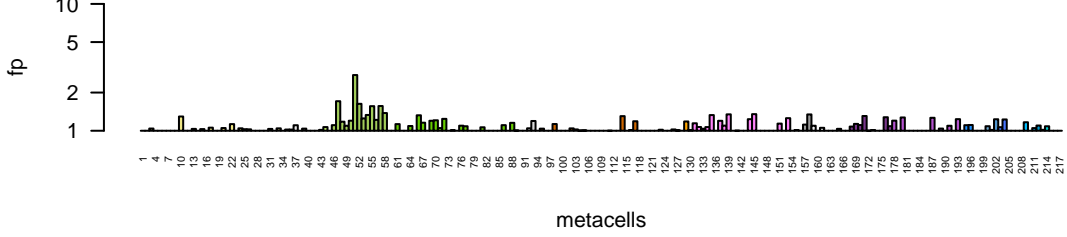
metacells



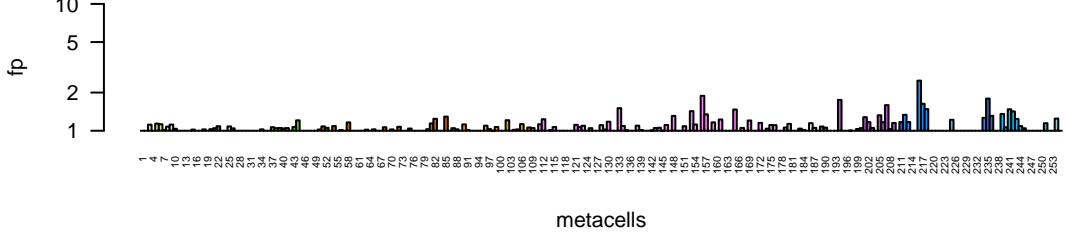
Tadh OG_2014
Tadh_hyp_clust2100.t1
NFYB_NFYC.HG3.2:DRAP1



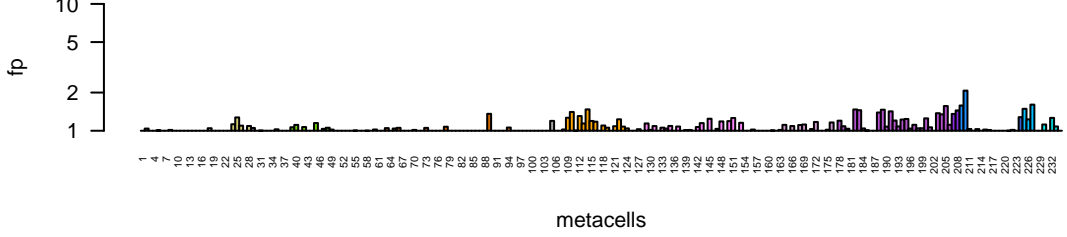
TrH2 OG_2014
TrH2_TrispH2_009775-RA
NFYB_NFYC.HG3.2:DRAP1

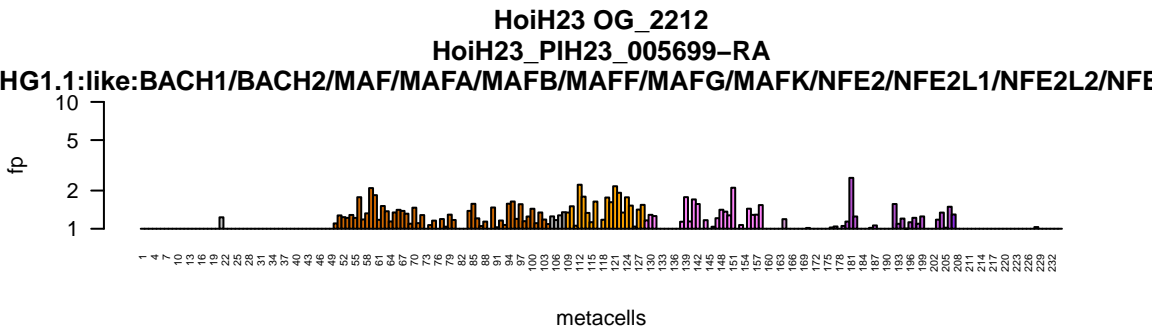
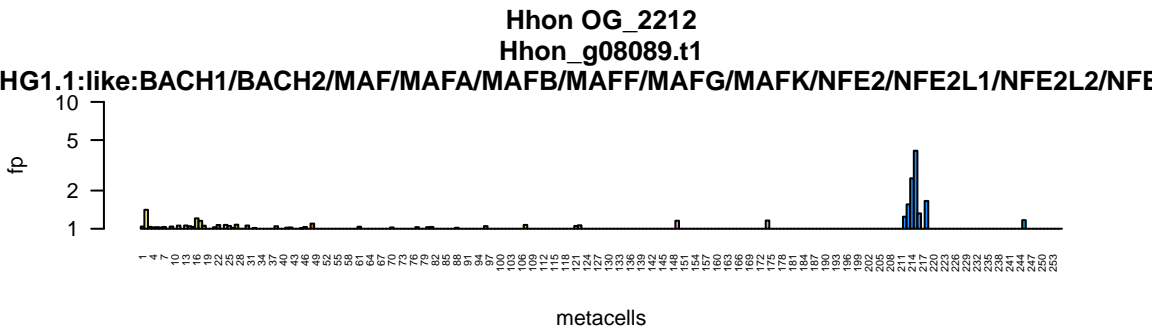
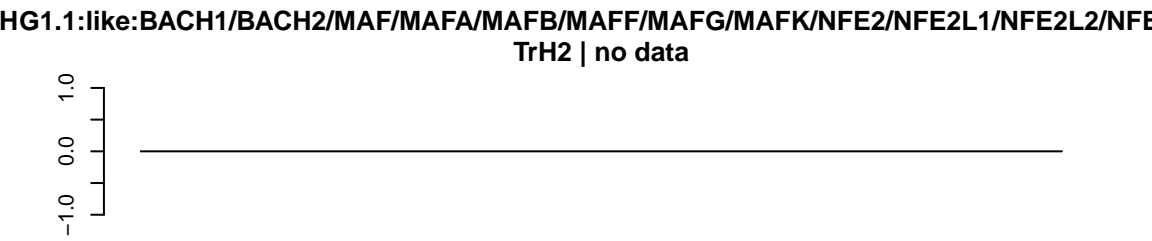
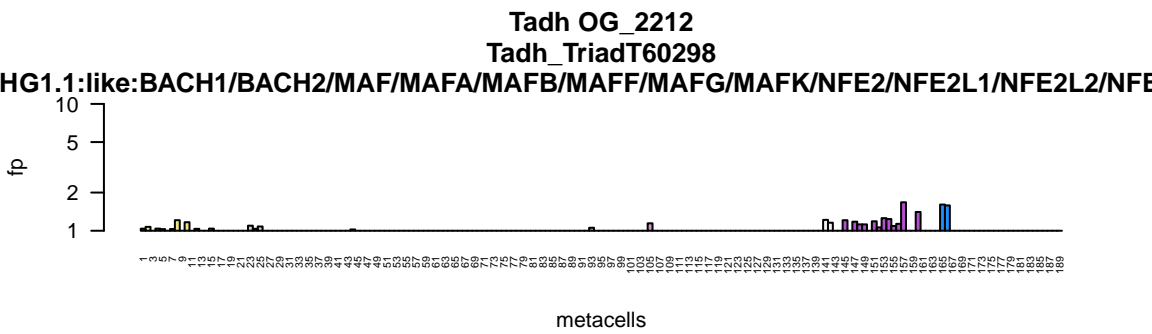


Hhon OG_2014
Hhon_g07246.t1
NFYB_NFYC.HG3.2:DRAP1



HoiH23 OG_2014
HoiH23_PIH23_010470-RA
NFYB_NFYC.HG3.2:DRAP1

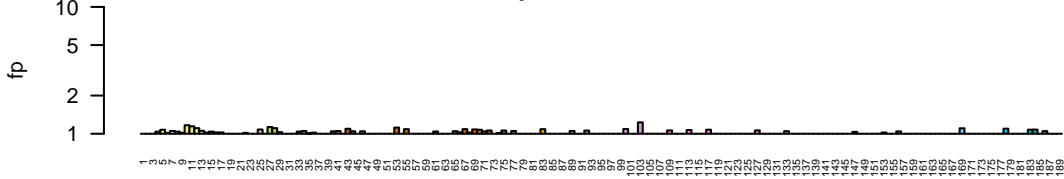




Tadh OG_3669

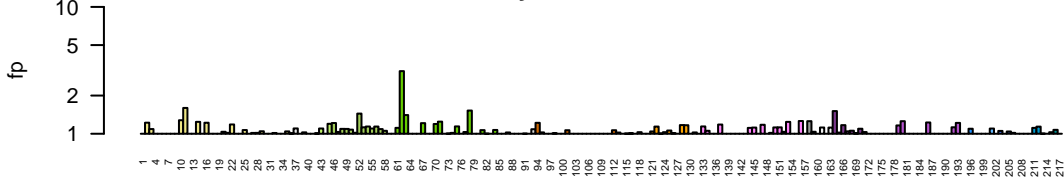
Tadh_TriadT53135

Myb.HG1.0:ZZZ3



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TrH2 OG_3669
TrH2_TrispH2_003060-RA
Myb.HG1.0:ZZZ3

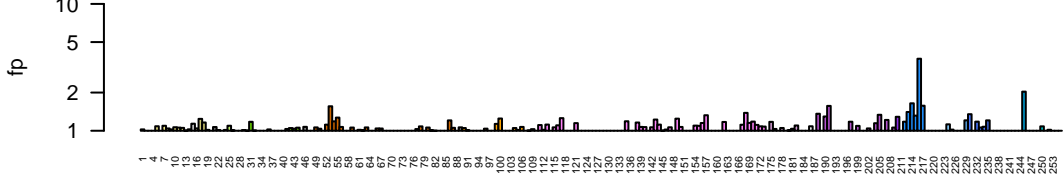


metacells

Hhon OG_3669

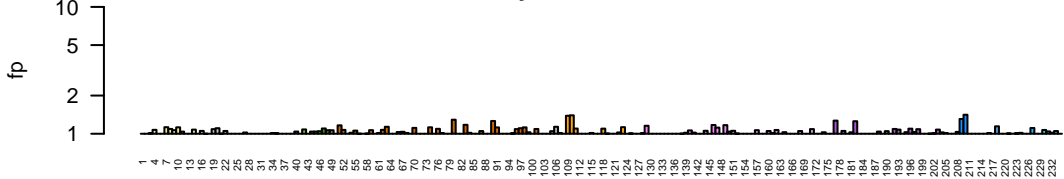
Hhon_g09423.t1

Myb.HG1.0:ZZZ3

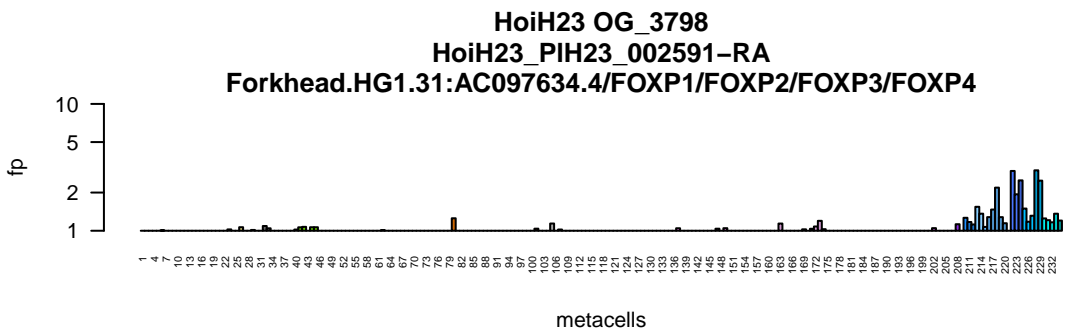
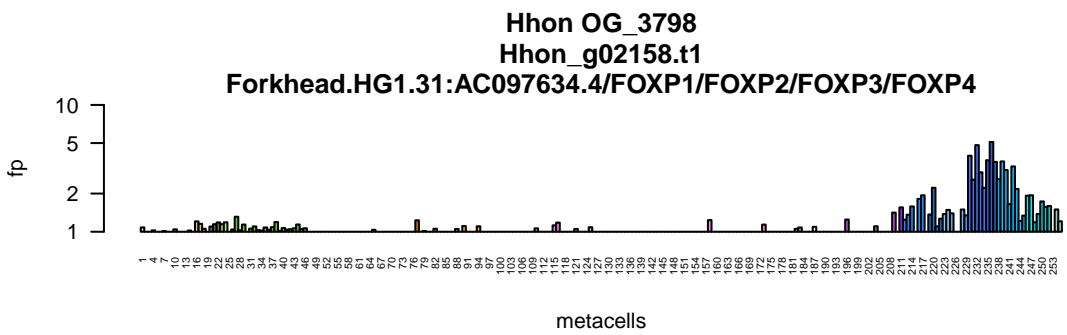
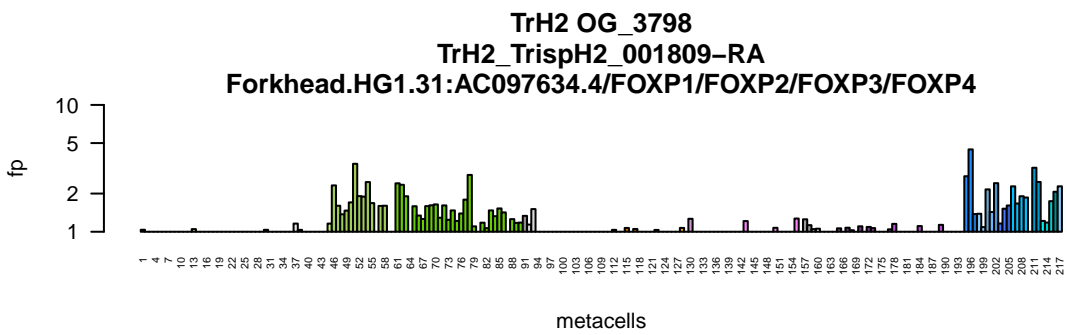
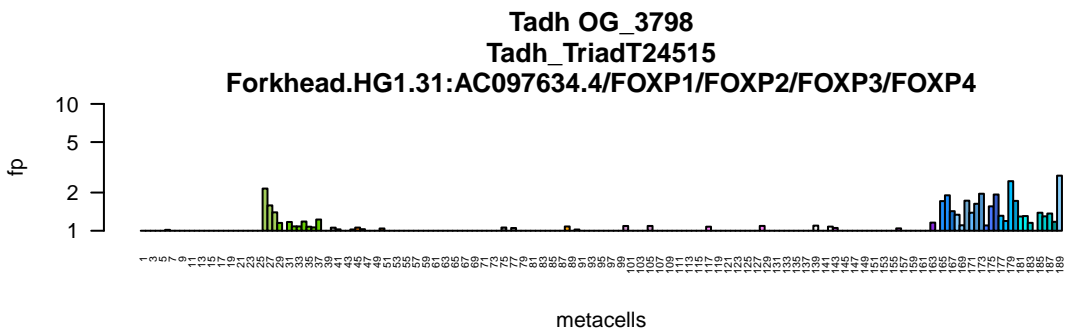


metacells

HoiH23 OG_3669
HoiH23_PIH23_000017-RA
Myb.HG1.0:ZZZ3

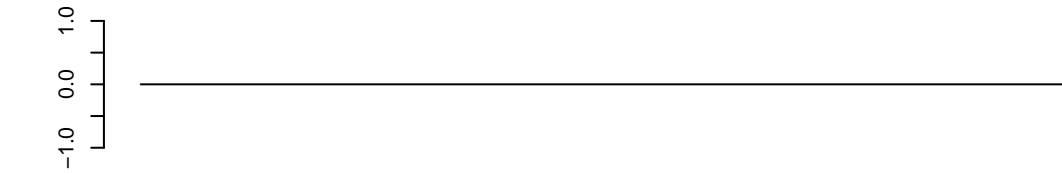


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HMGbox_Sox.HG1.23:AC105001.2/SOX4/SOX7/SOX8/SOX9/SOX10/SOX11/SOX12/SOX17/SO

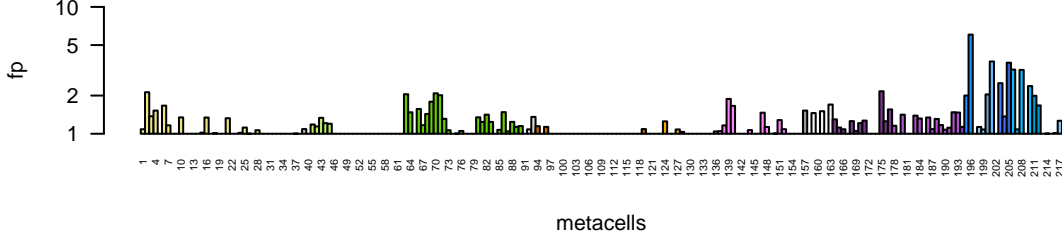
Tadh | no data



TrH2 OG_5274

TrH2_TrispH2_011376-RA

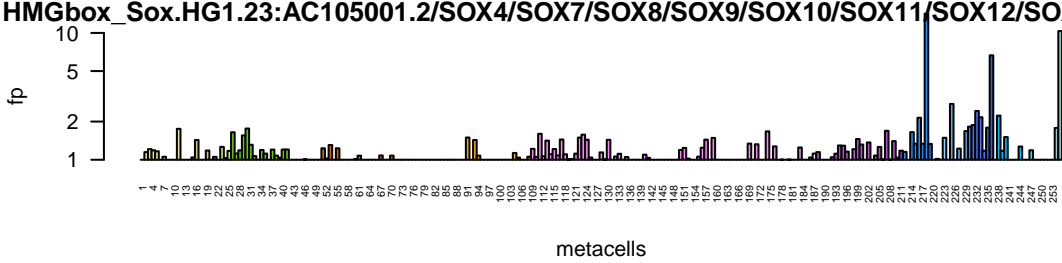
HMGbox_Sox.HG1.23:AC105001.2/SOX4/SOX7/SOX8/SOX9/SOX10/SOX11/SOX12/SOX17/SO



Hhon OG_5274

Hhon_g07787.t1

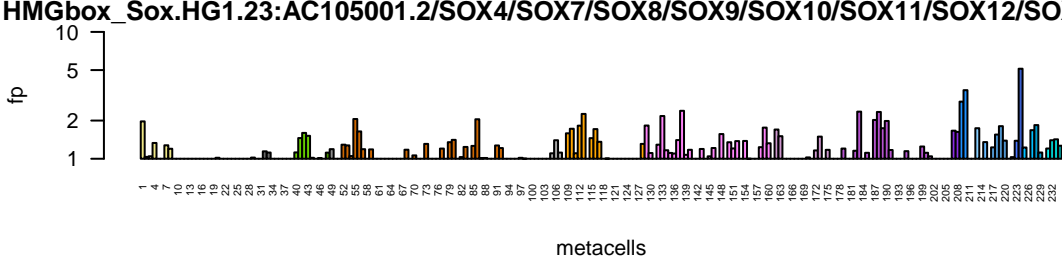
HMGbox_Sox.HG1.23:AC105001.2/SOX4/SOX7/SOX8/SOX9/SOX10/SOX11/SOX12/SOX17/SO

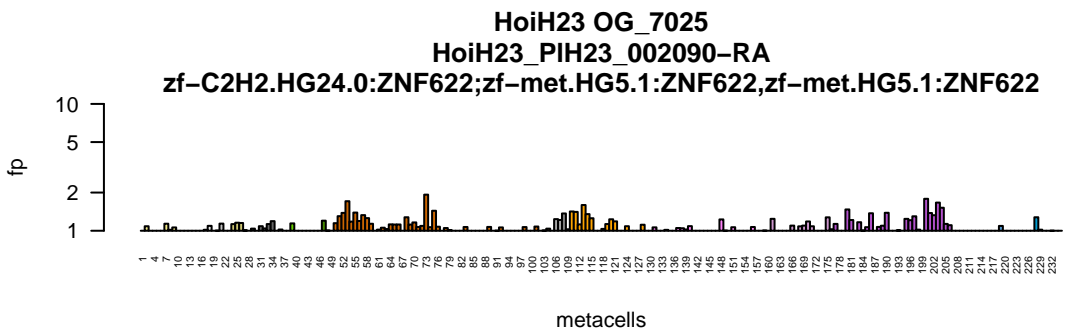
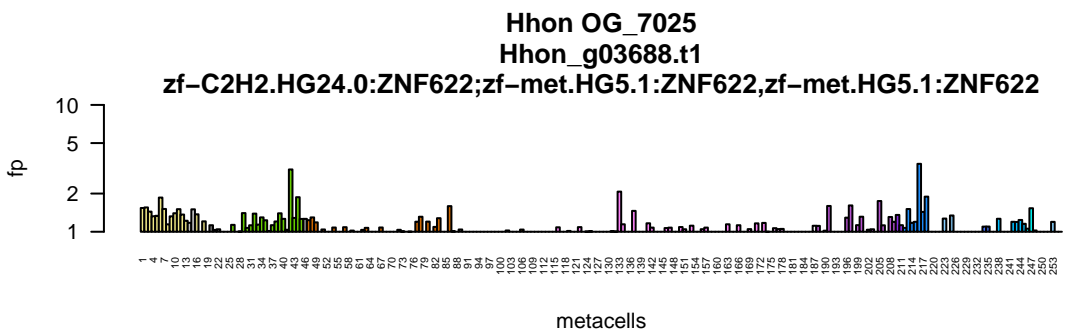
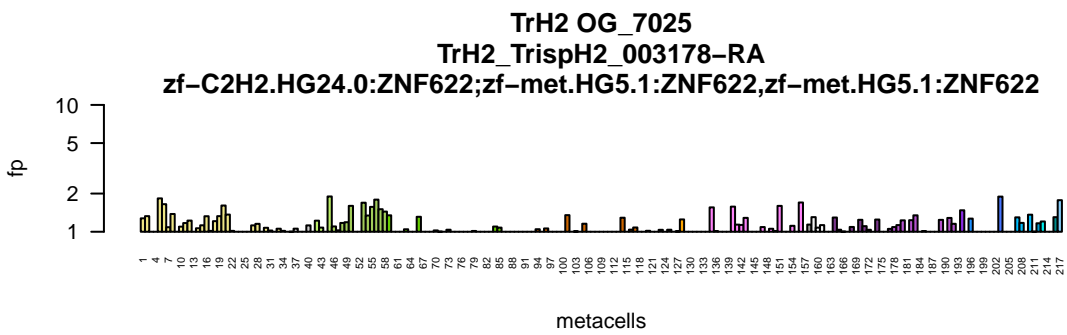
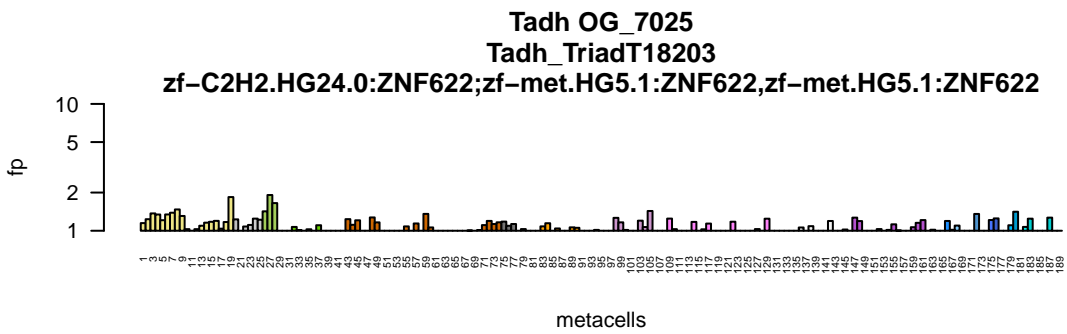


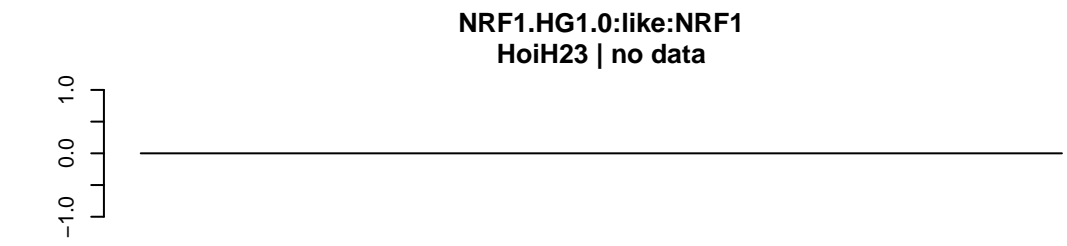
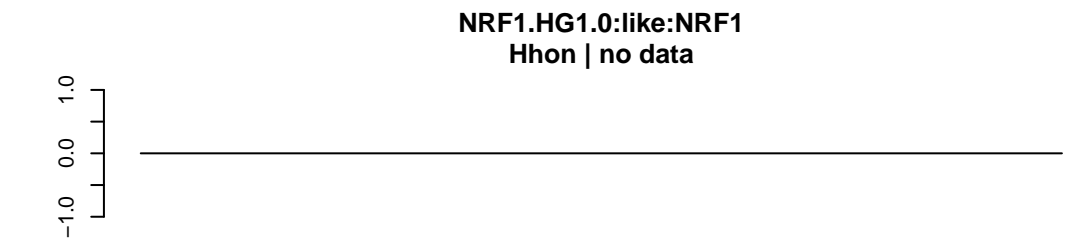
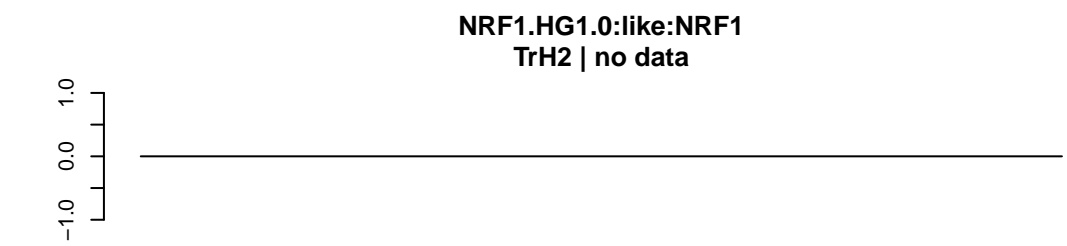
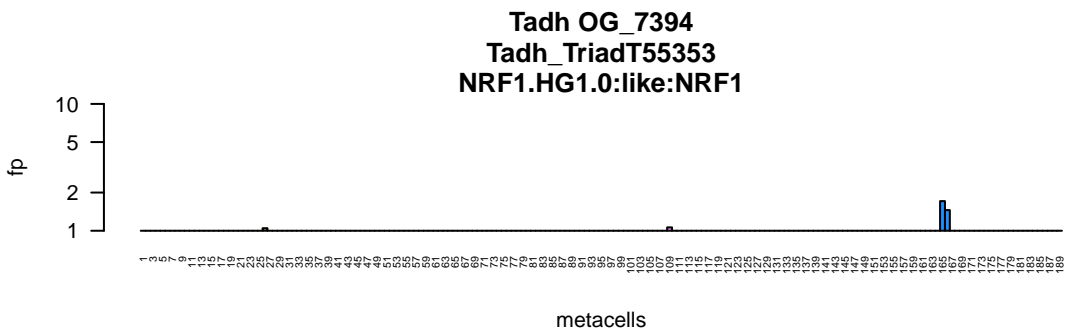
HoiH23 OG_5274

HoiH23_PIH23_010696-RA

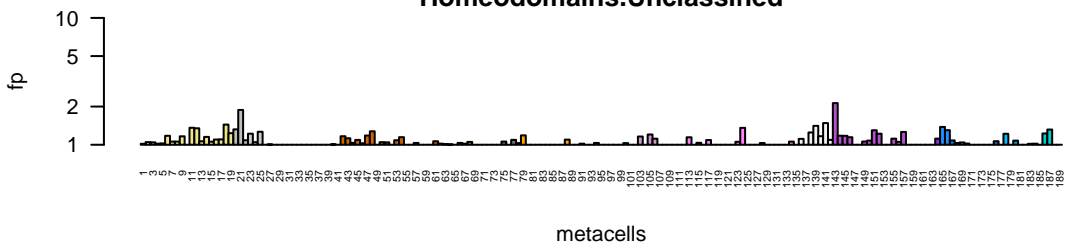
HMGbox_Sox.HG1.23:AC105001.2/SOX4/SOX7/SOX8/SOX9/SOX10/SOX11/SOX12/SOX17/SO



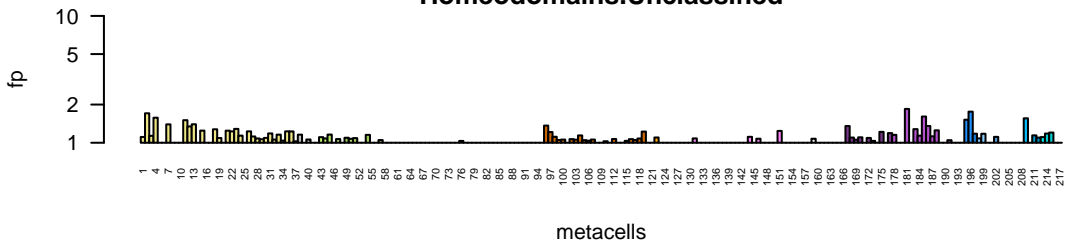




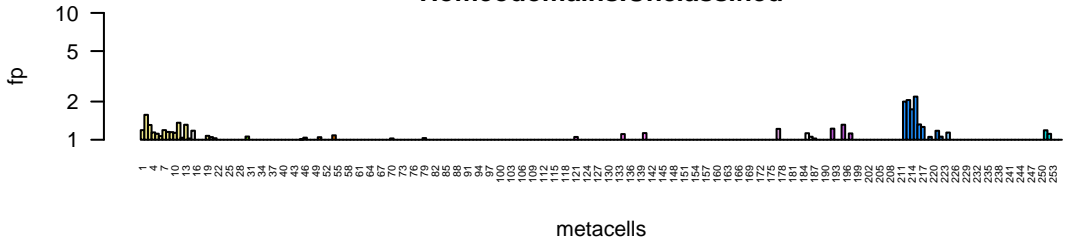
Tadh OG_8656
Tadh_TriadT30944
Homeodomains.Unclassified



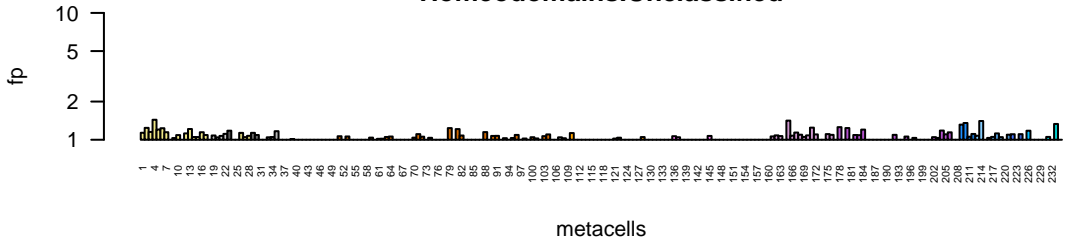
TrH2 OG_8656
TrH2_TrispH2_002999-RA
Homeodomains.Unclassified

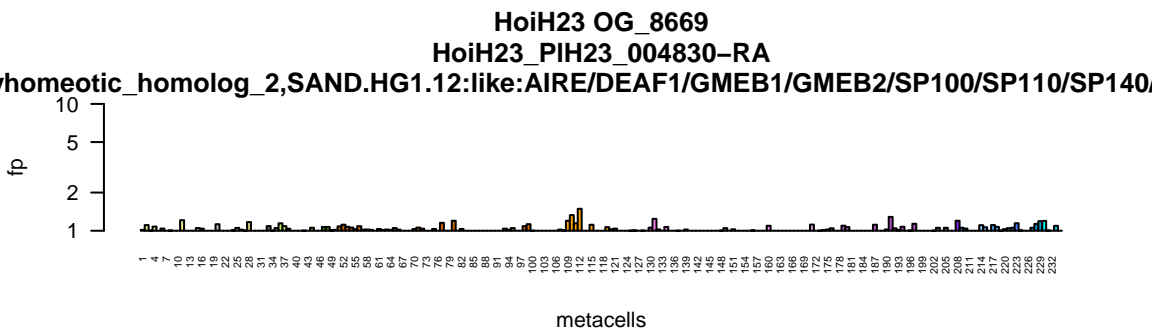
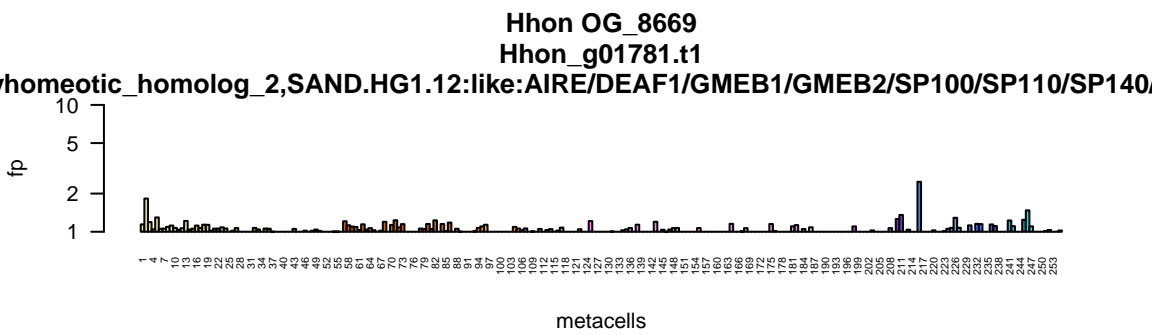
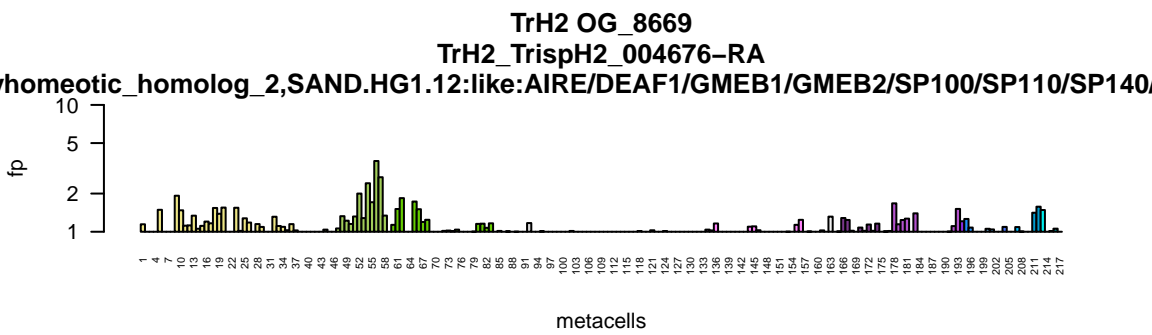
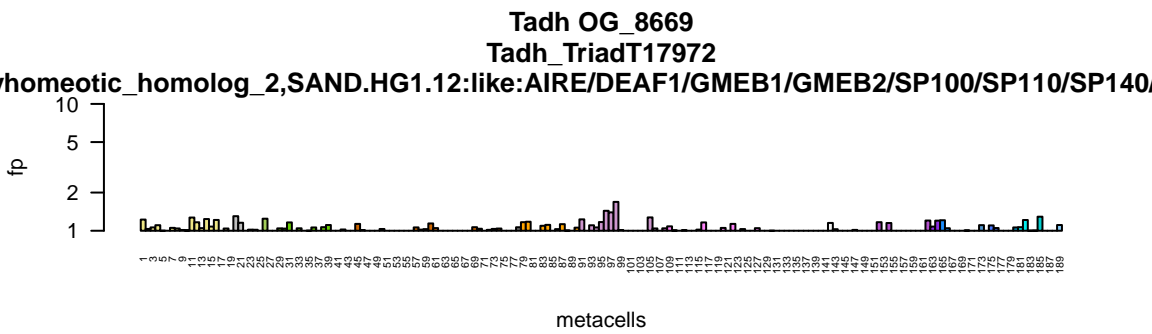


Hhon OG_8656
Hhon_g03907.t1
Homeodomains.Unclassified

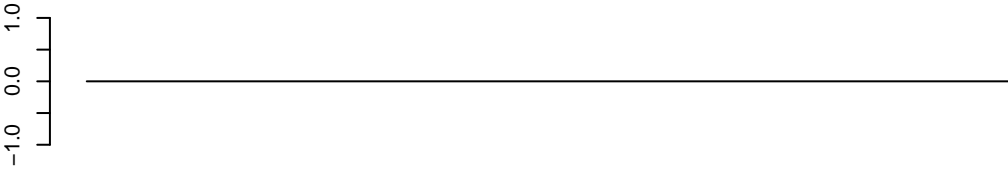


HoiH23 OG_8656
HoiH23_PIH23_004895-RA
Homeodomains.Unclassified

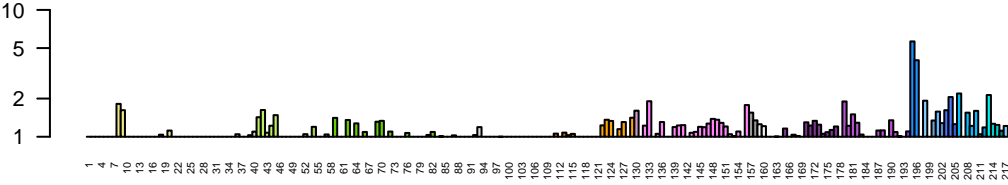




Forkhead.HG1.30:FOXN2/FOXN3
Tadh | no data

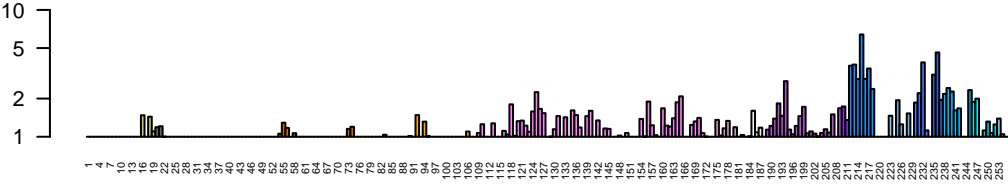


TrH2 OG_8934
TrH2_TrispH2_007536-RA
Forkhead.HG1.30:FOXN2/FOXN3



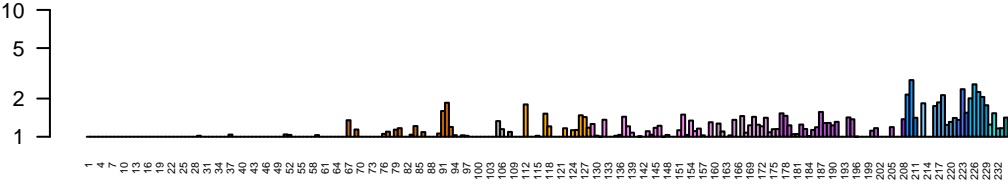
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Hhon OG_8934
Hhon_g07630.t1
Forkhead.HG1.30:FOXN2/FOXN3



metacells

HoiH23 OG_8934
HoiH23_PIH23_002310-RA
Forkhead.HG1.30:FOXN2/FOXN3



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