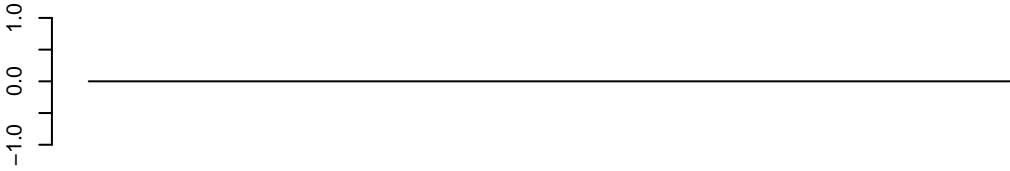
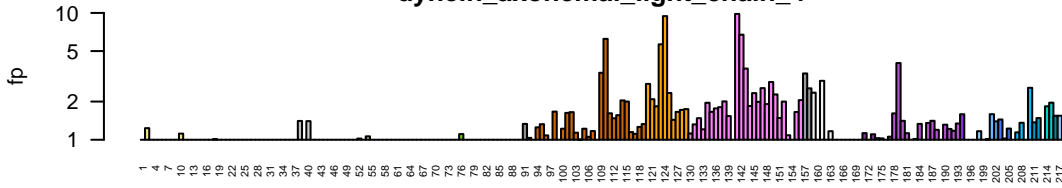


dynein_axonemal_light_chain_4
Tadh | no data

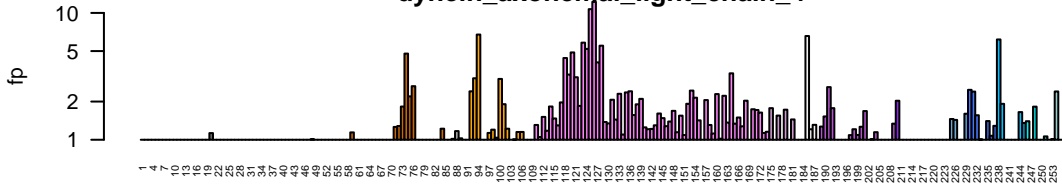


TrH2 OG_8349
TrH2_TrispH2_005999-RA
dynein_axonemal_light_chain_4



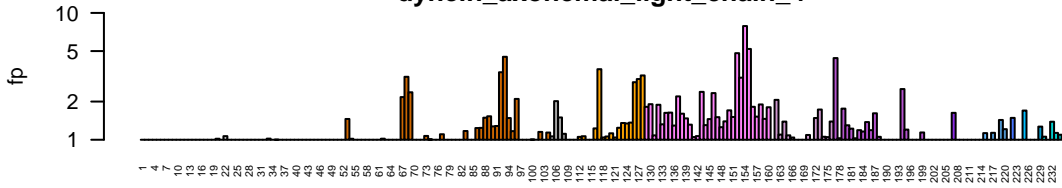
metacells

Hhon OG_8349
Hhon_g01448.t1
dynein_axonemal_light_chain_4

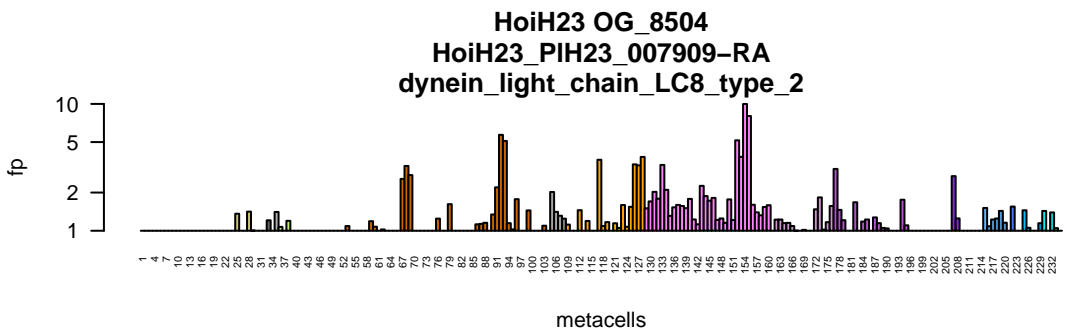
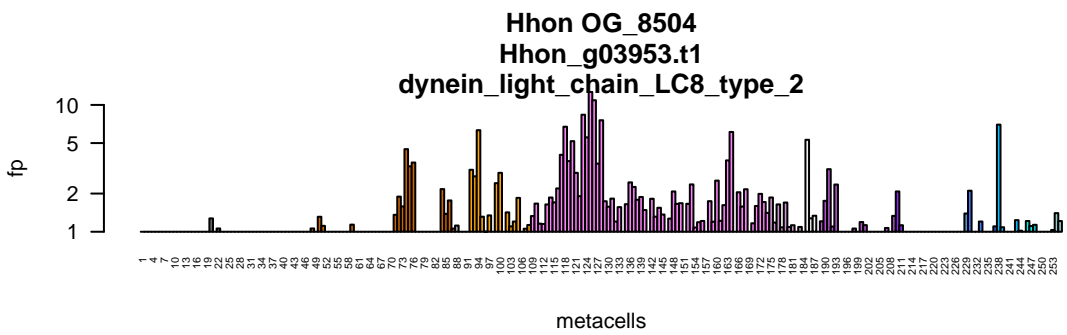
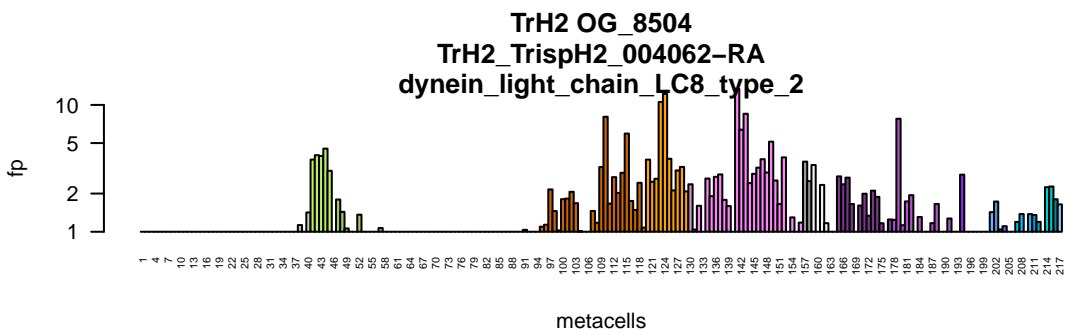
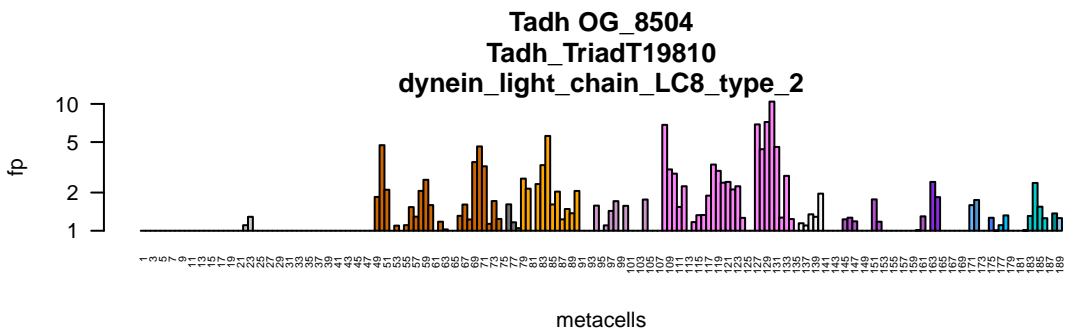


metacells

HoiH23 OG_8349
HoiH23_PIH23_002932-RA
dynein_axonemal_light_chain_4



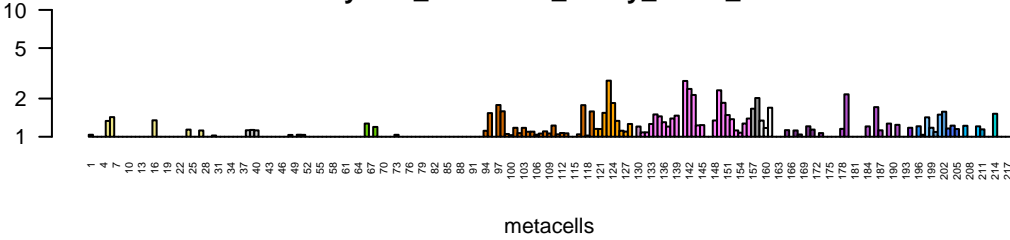
metacells



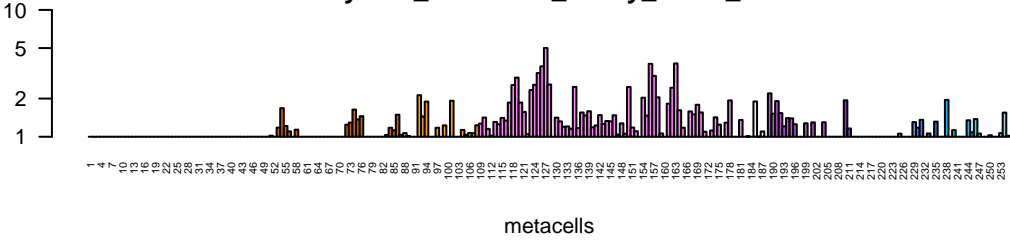
dynein_axonemal_heavy_chain_17
Tadh | no data



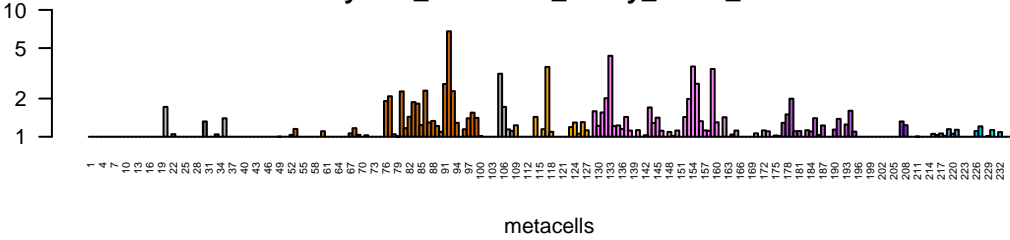
TrH2 OG_8561
TrH2_TrispH2_005969-RA
dynein_axonemal_heavy_chain_17

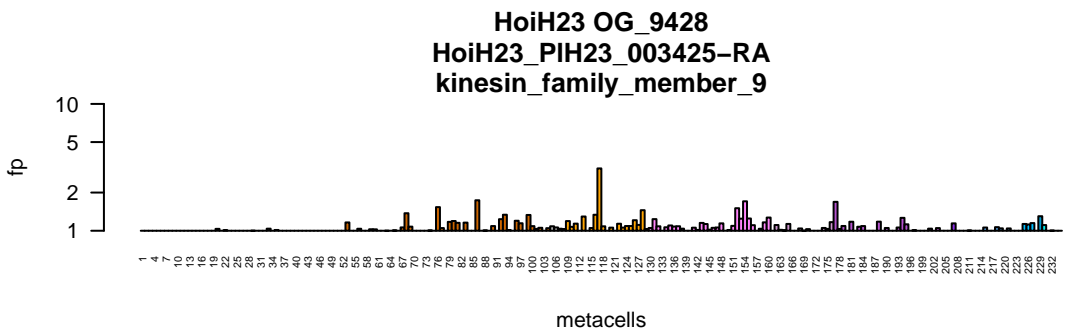
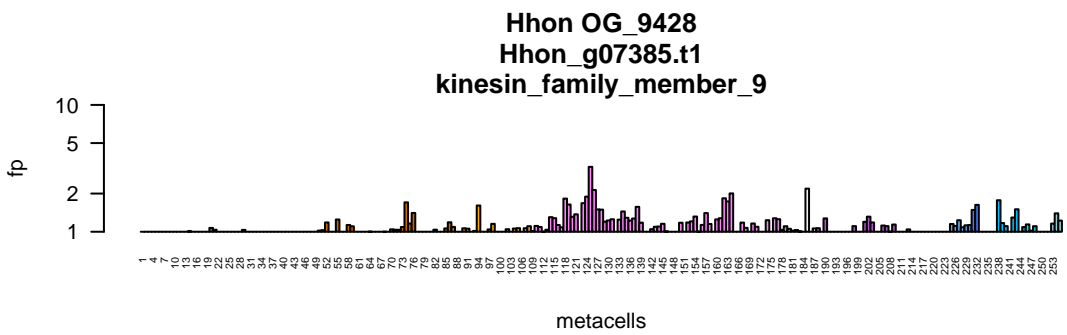
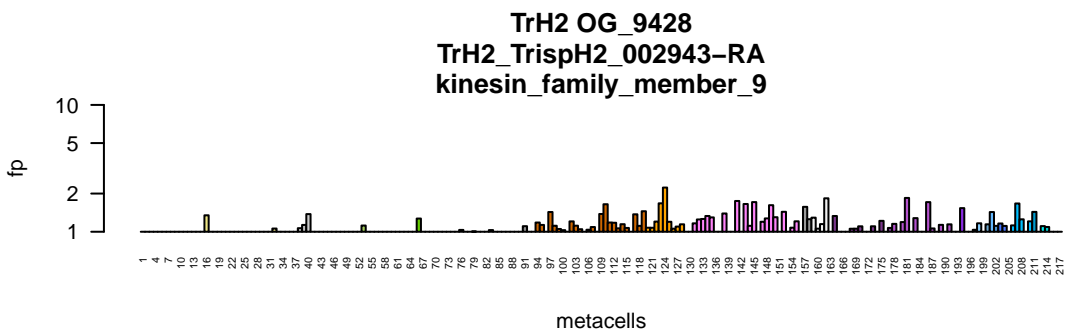
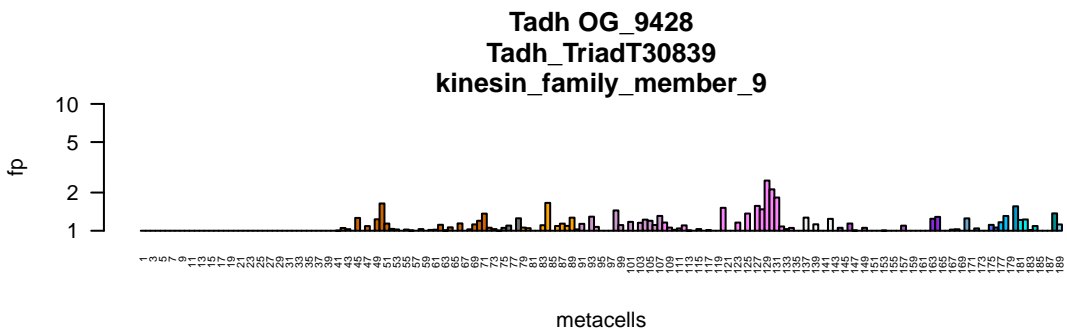


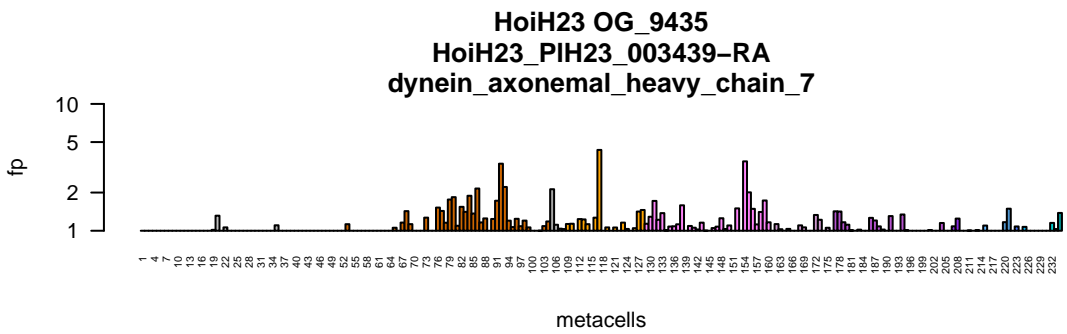
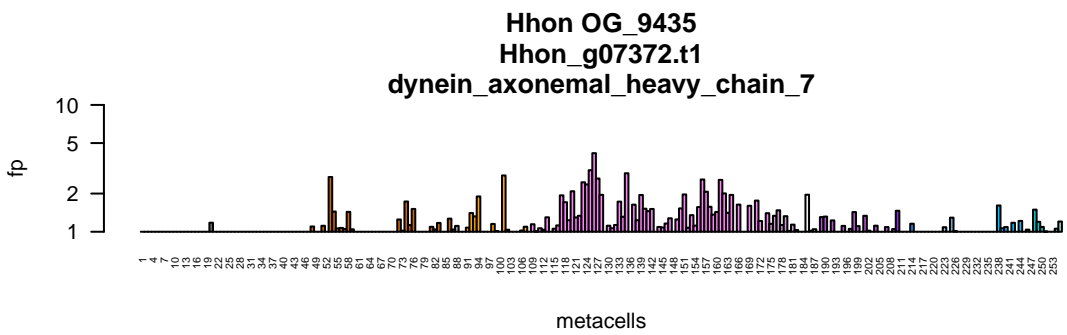
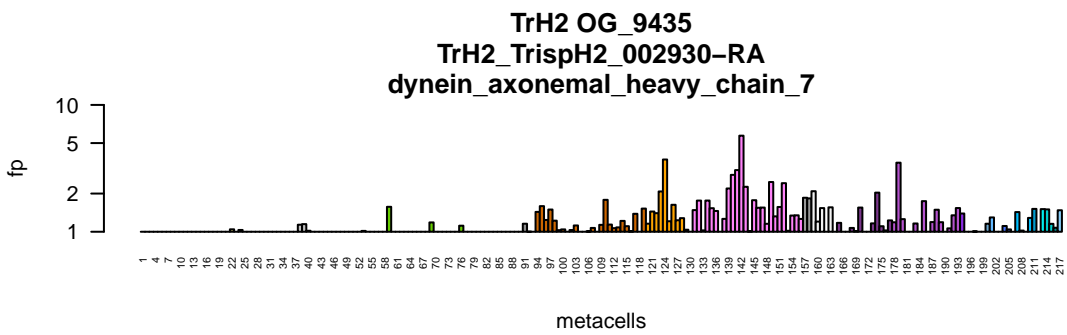
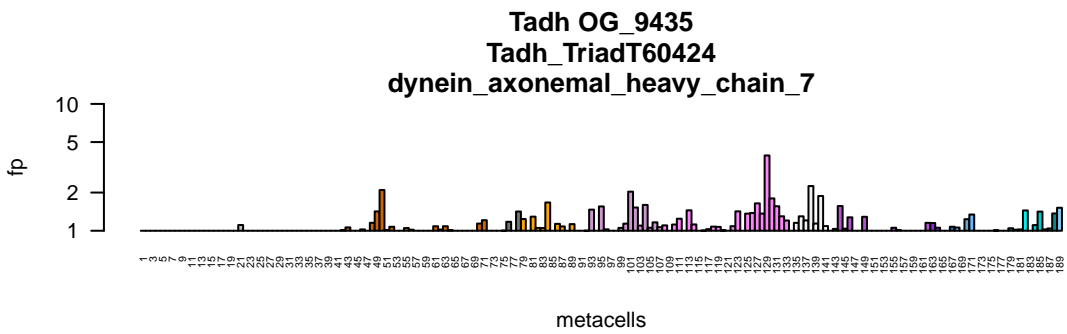
Hhon OG_8561
Hhon_g01475.t1
dynein_axonemal_heavy_chain_17

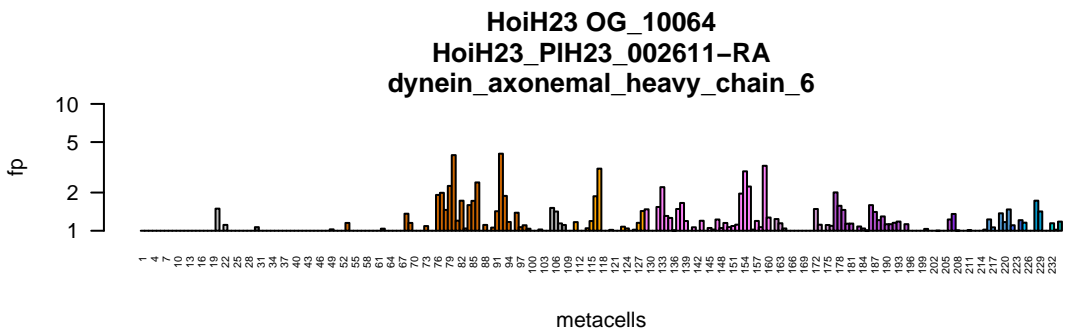
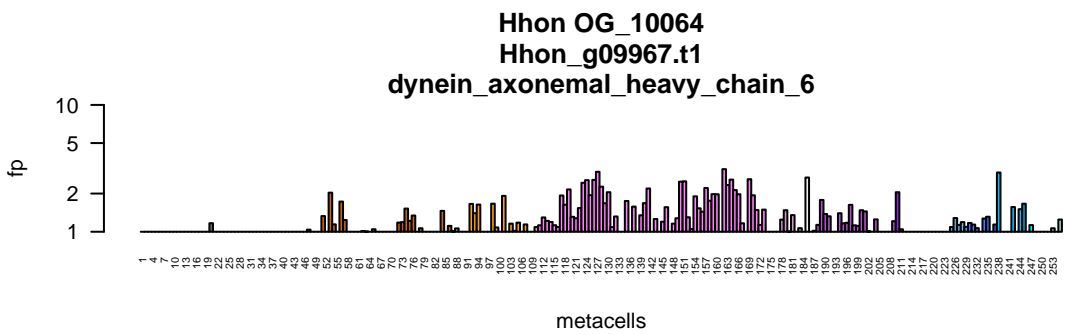
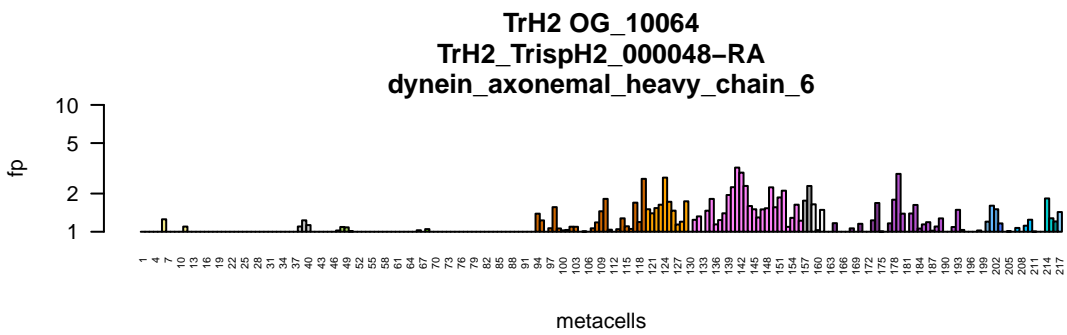
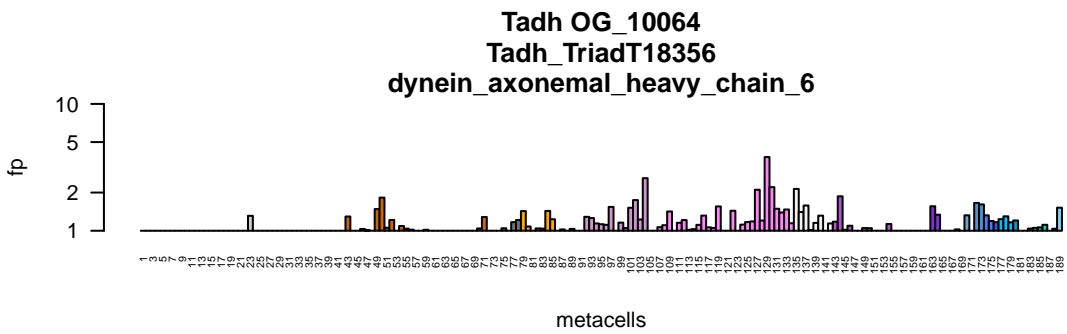


HoiH23 OG_8561
HoiH23_PIH23_001971-RA
dynein_axonemal_heavy_chain_17





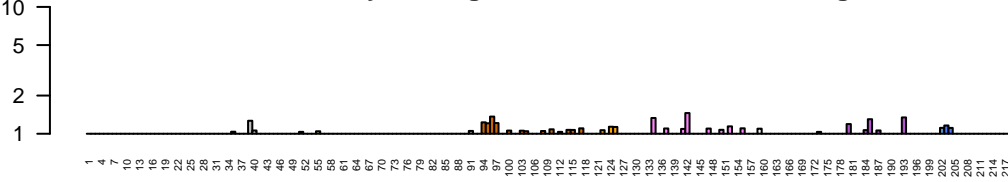




axonemal_dynein_light_chain_domain_containing_1
Tadh | no data

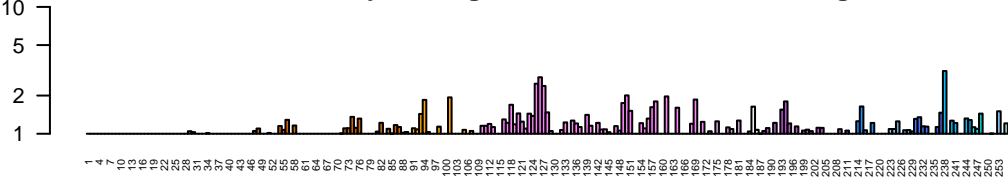


TrH2 OG_441
TrH2_TrispH2_002682-RA
axonemal_dynein_light_chain_domain_containing_1



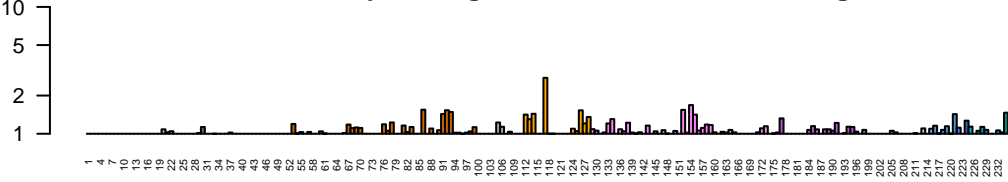
metacells

Hhon OG_441
Hhon_g01591.t1
axonemal_dynein_light_chain_domain_containing_1

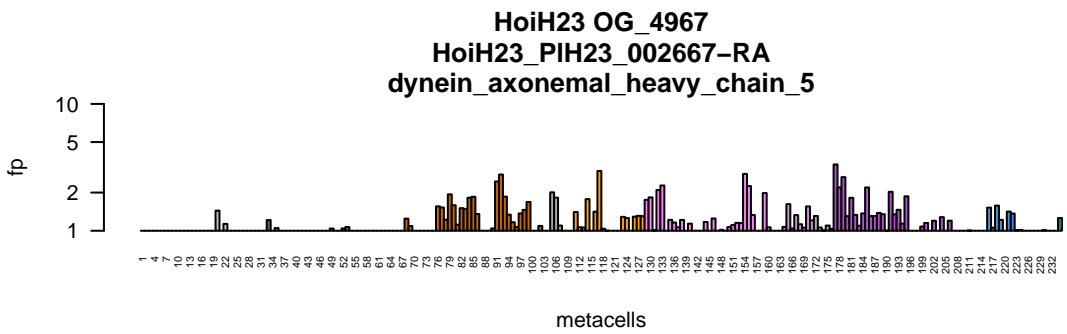
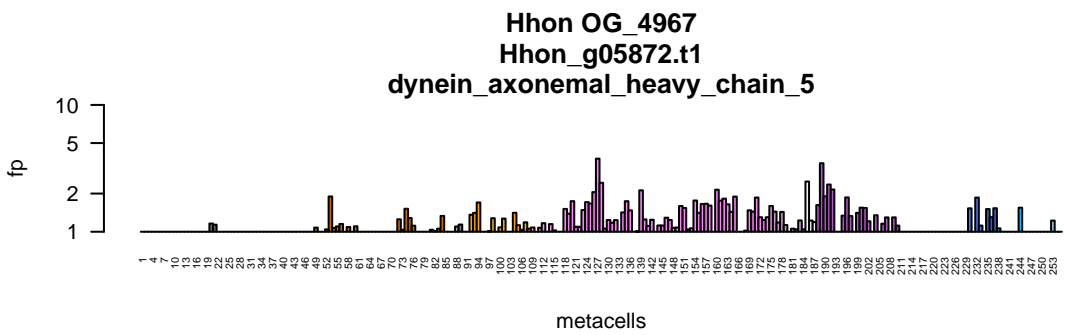
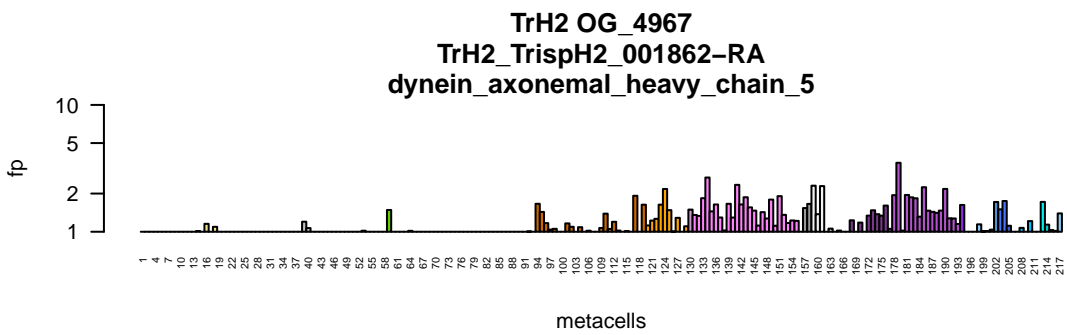
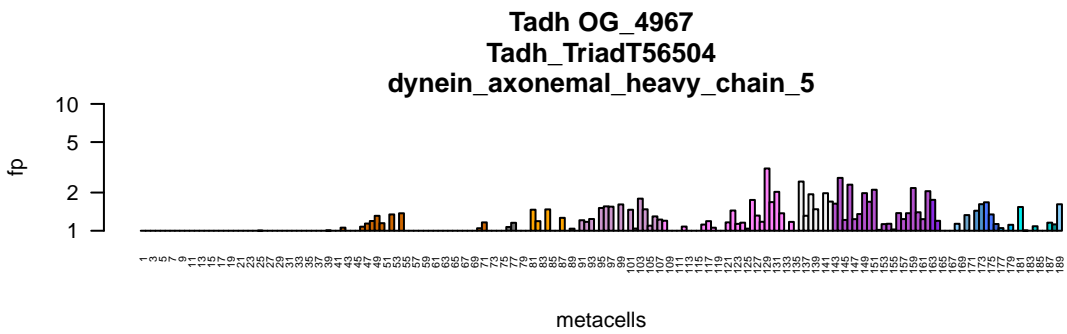


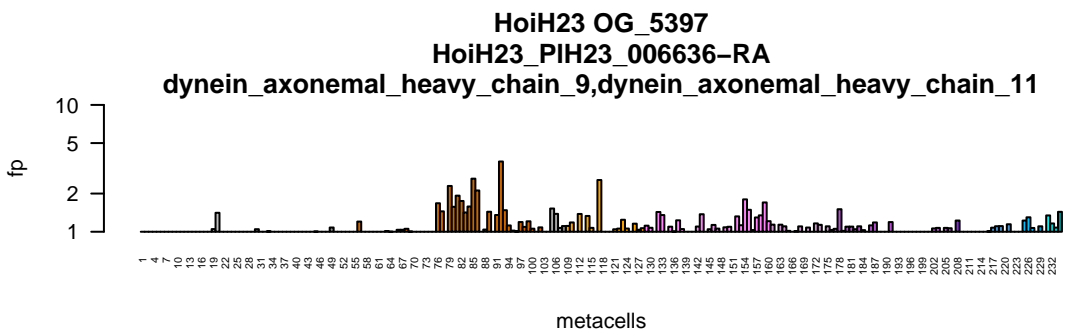
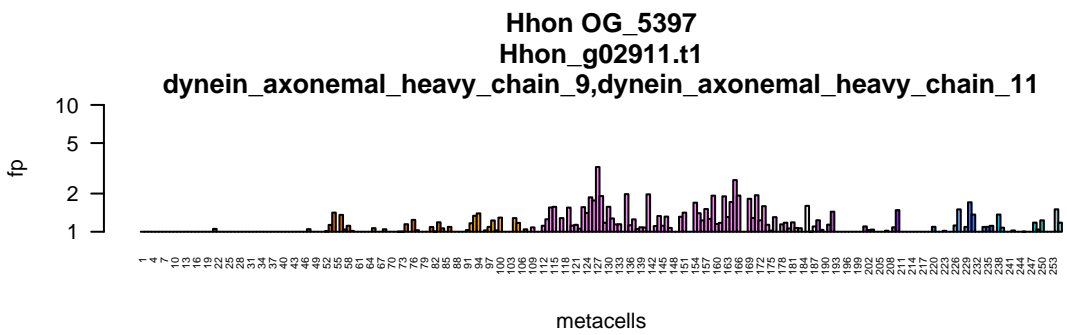
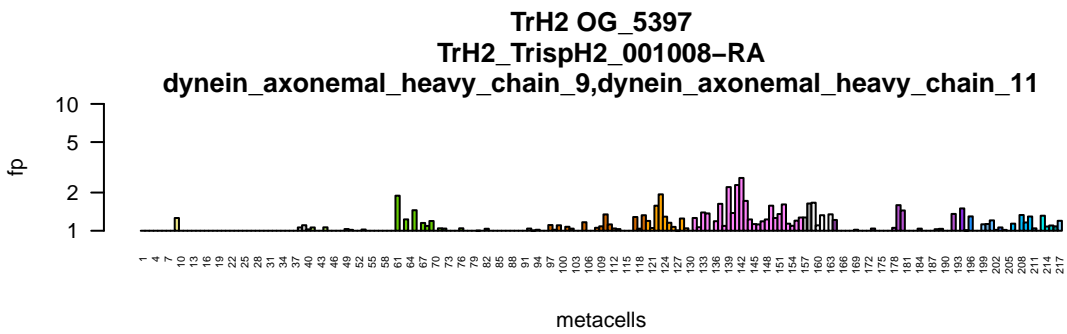
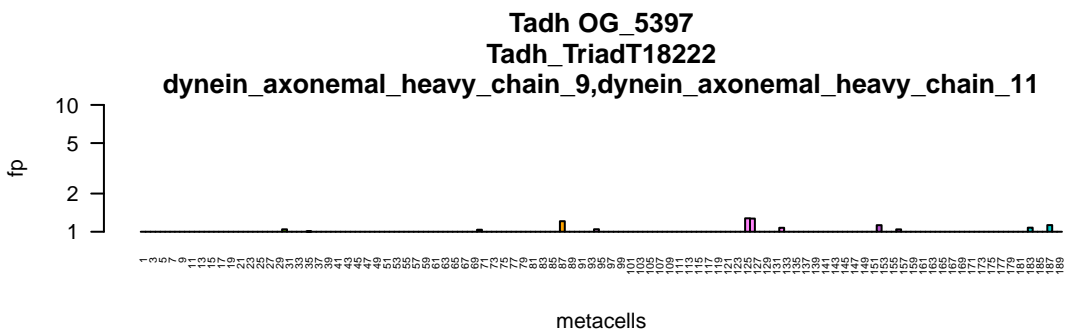
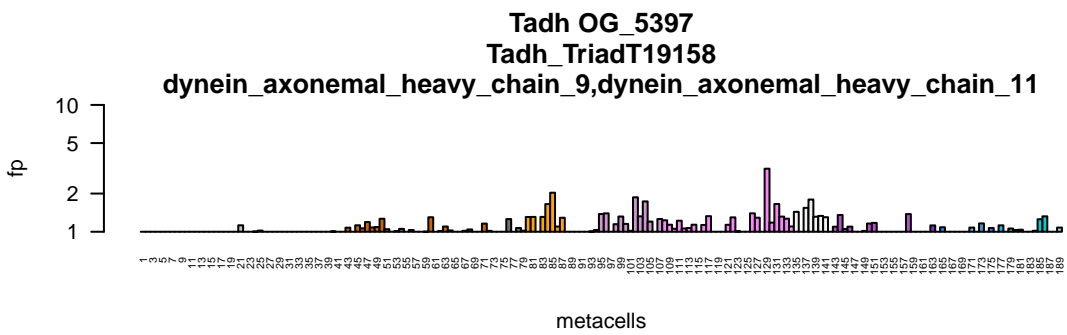
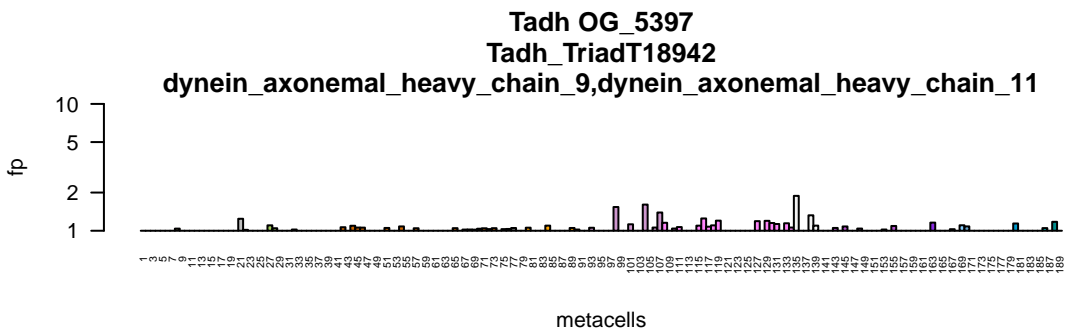
metacells

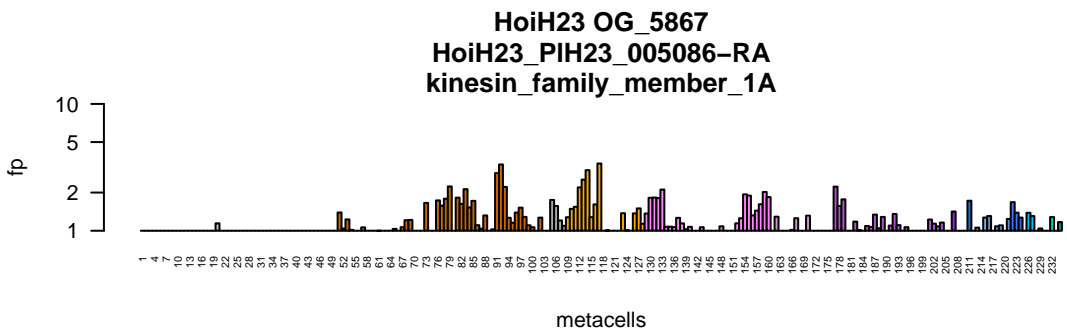
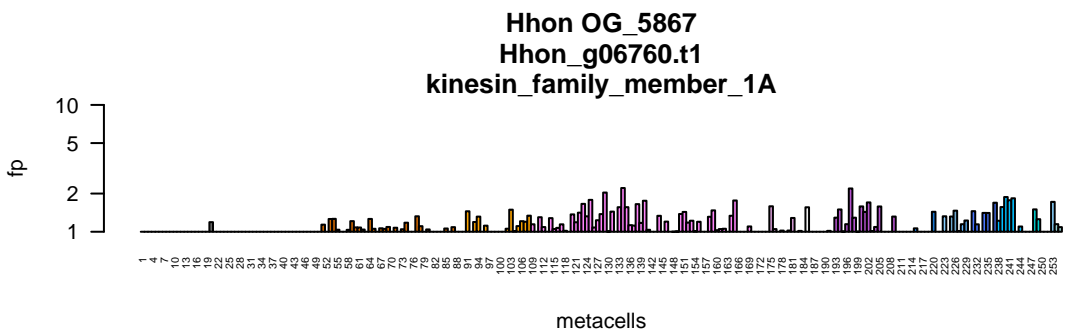
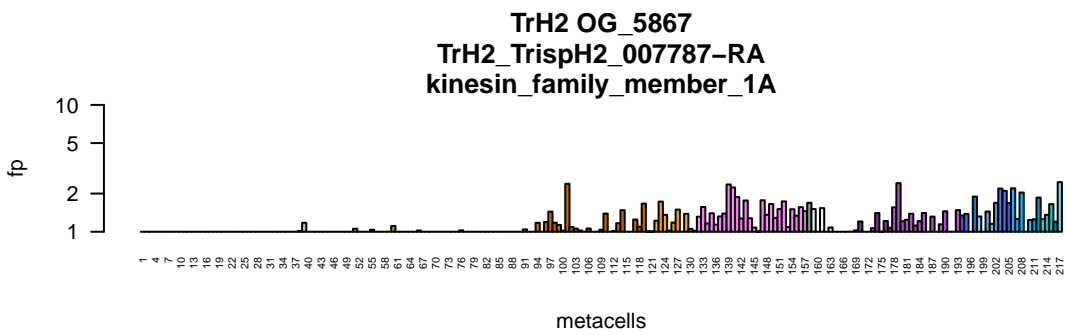
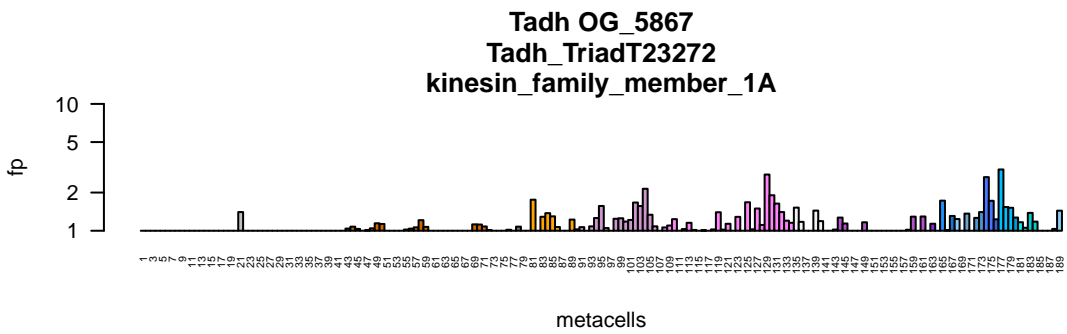
HoiH23 OG_441
HoiH23_PIH23_002503-RA
axonemal_dynein_light_chain_domain_containing_1



metacells



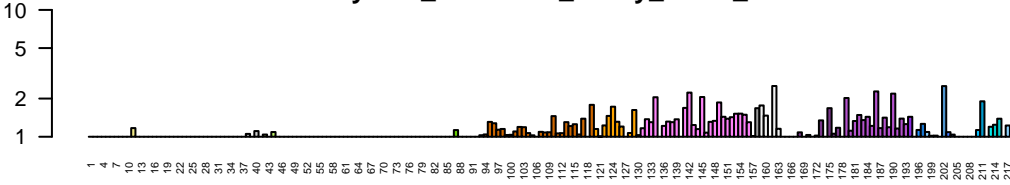




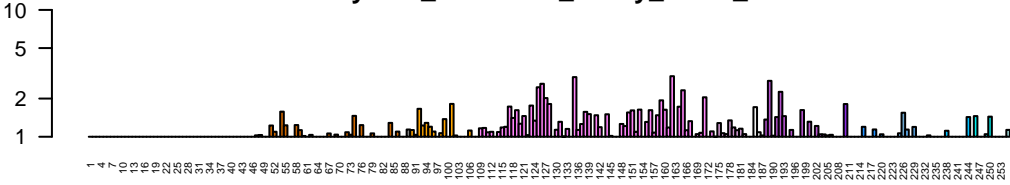
dynein_axonemal_heavy_chain_3
Tadh | no data



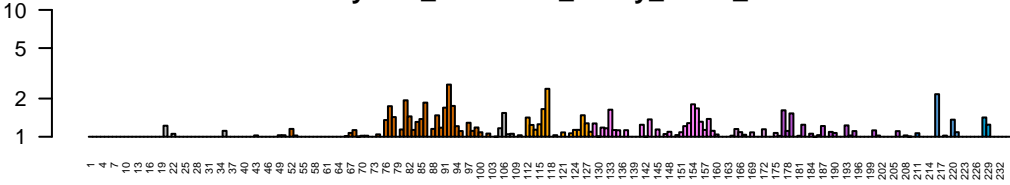
TrH2 OG_6103
TrH2_TrispH2_003413-RA
dynein_axonemal_heavy_chain_3

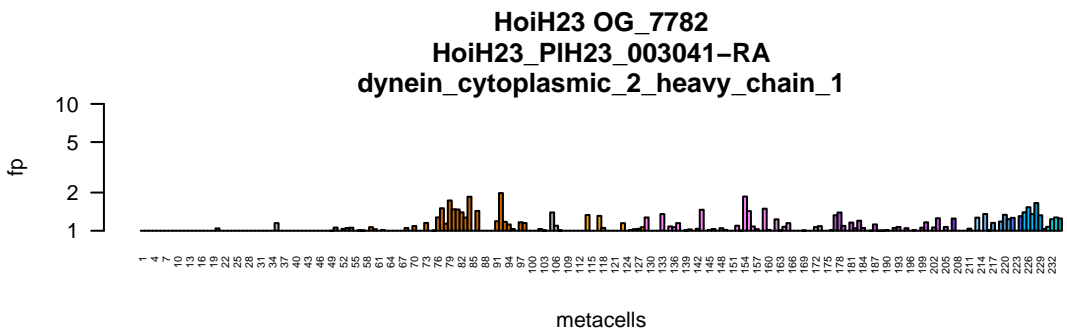
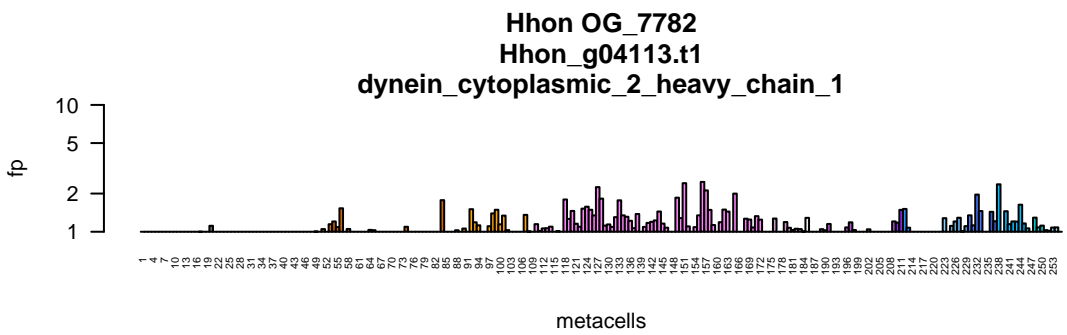
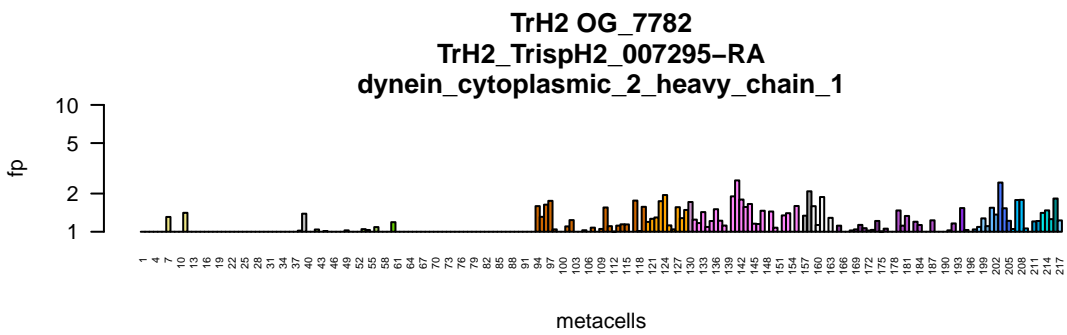
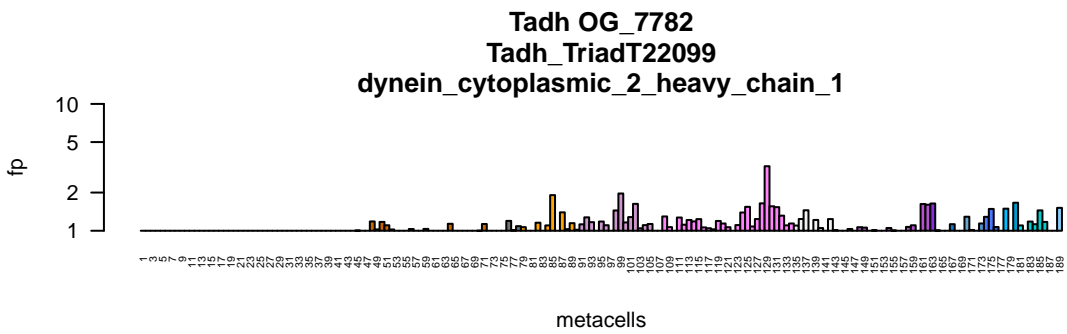


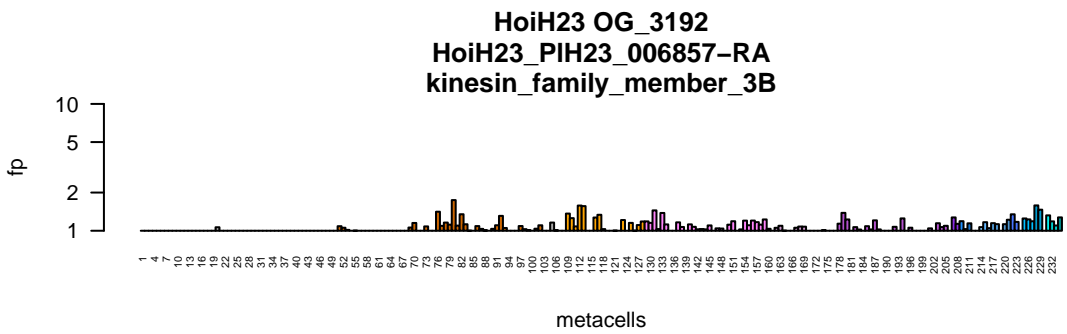
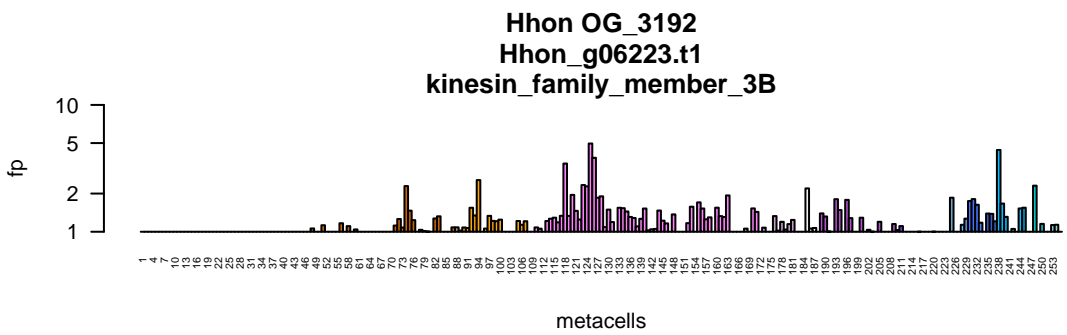
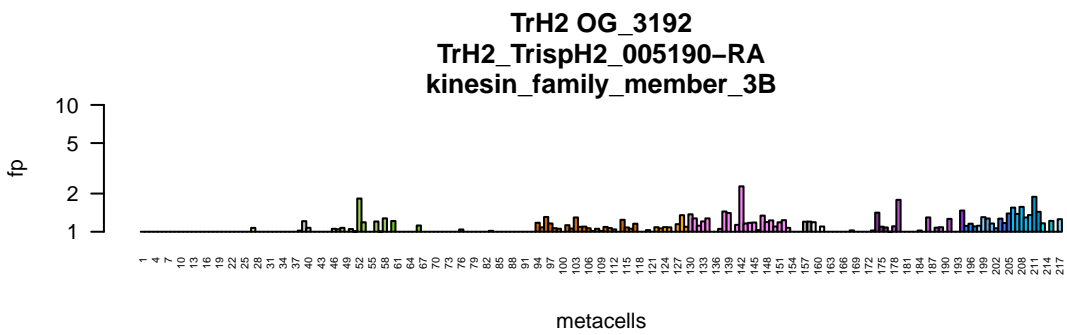
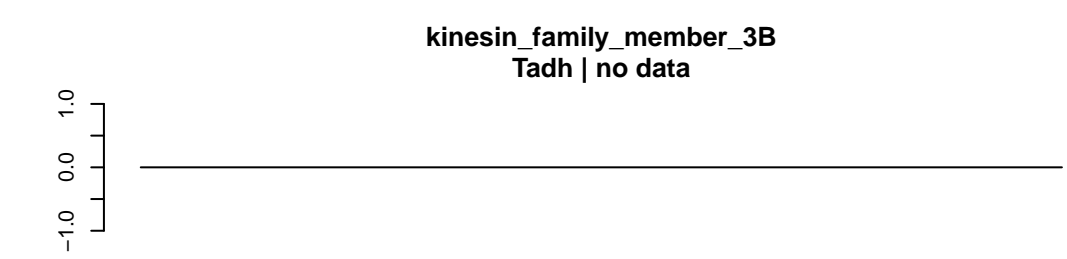
Hhon OG_6103
Hhon_g05504.t1
dynein_axonemal_heavy_chain_3

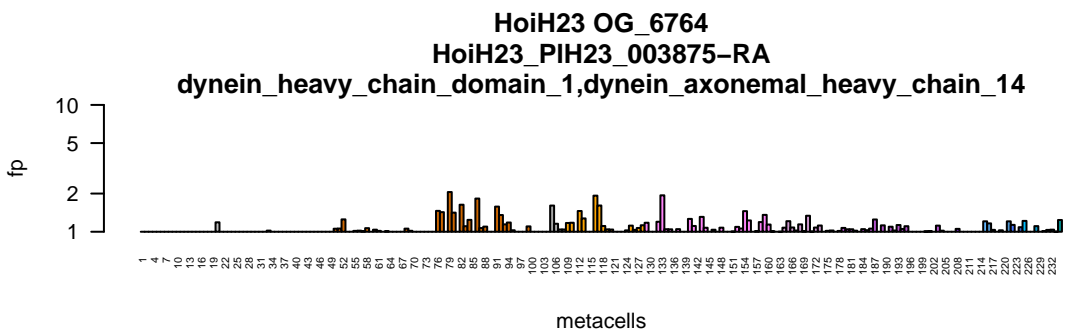
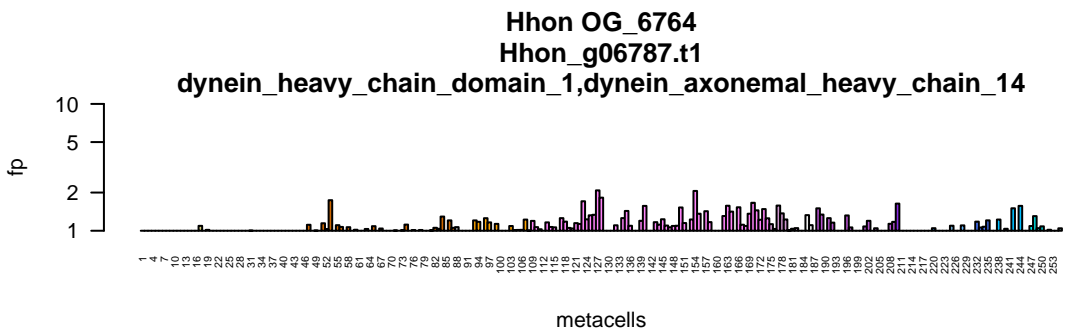
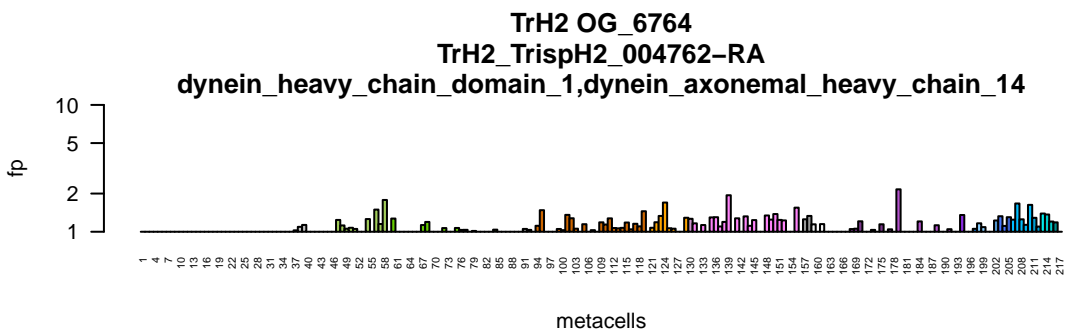
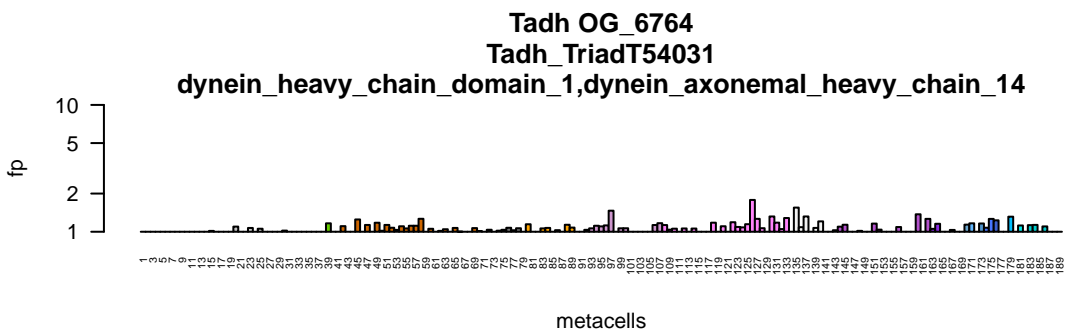
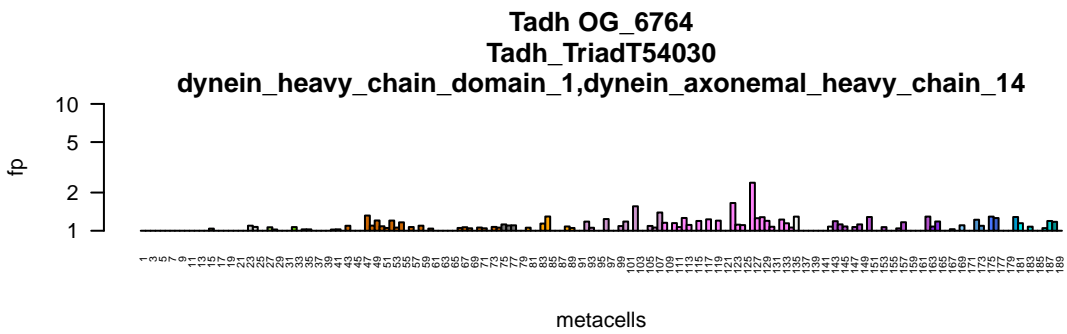


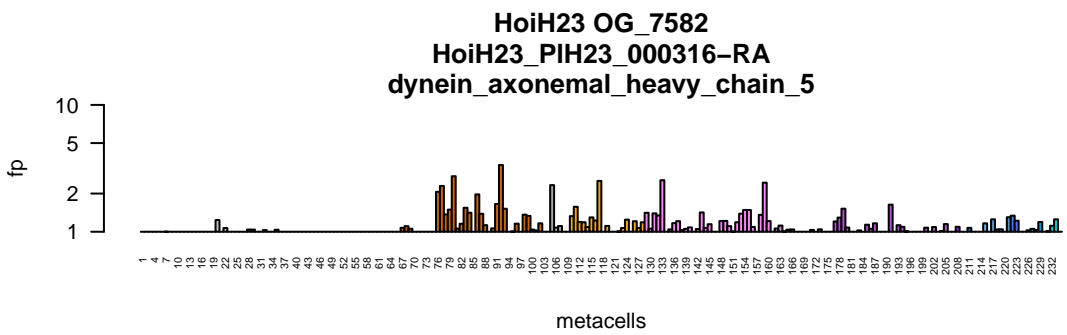
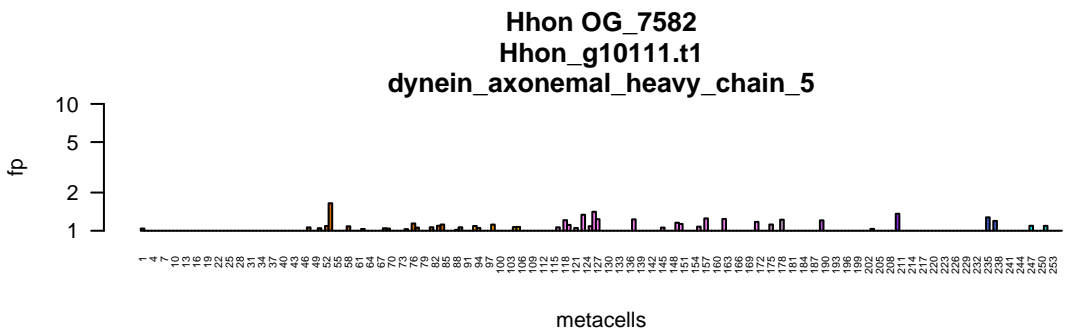
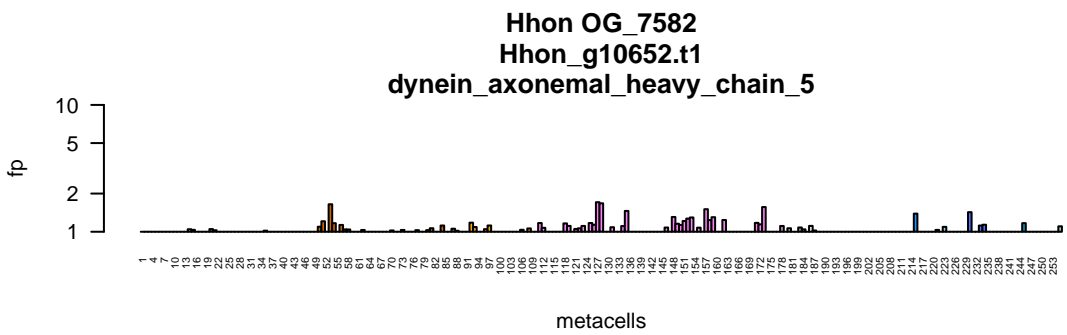
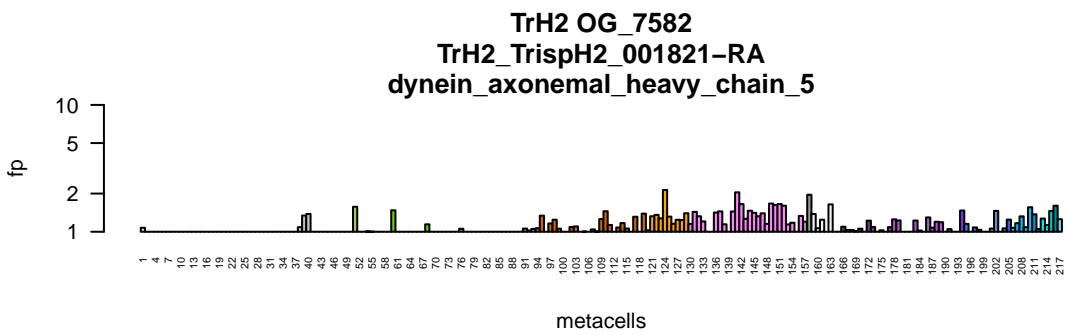
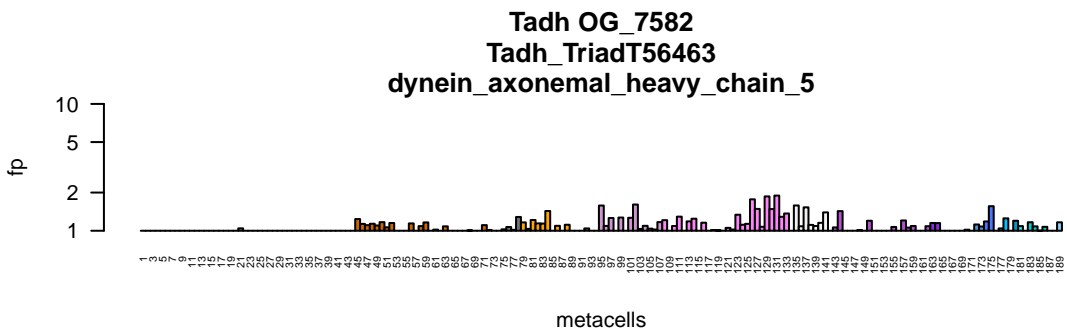
HoiH23 OG_6103
HoiH23_PIH23_000174-RA
dynein_axonemal_heavy_chain_3

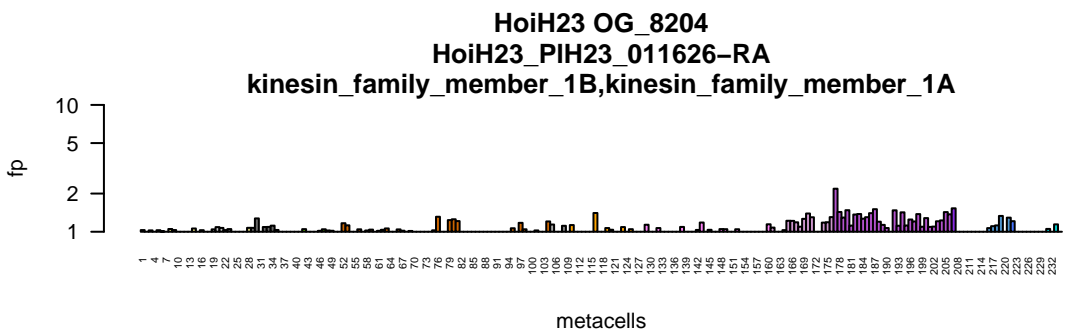
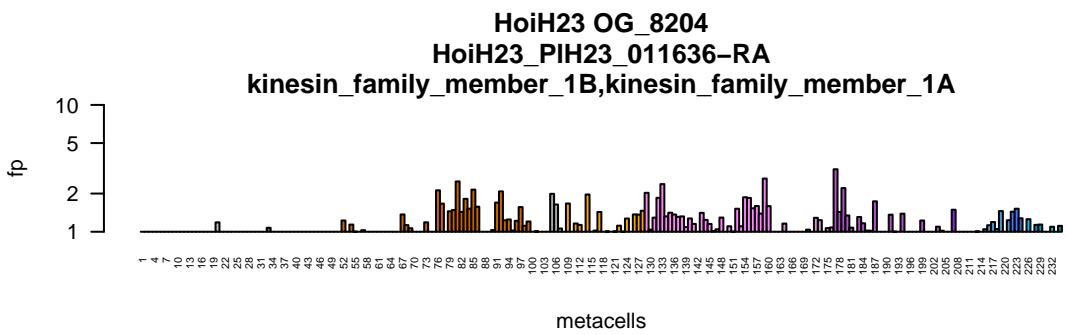
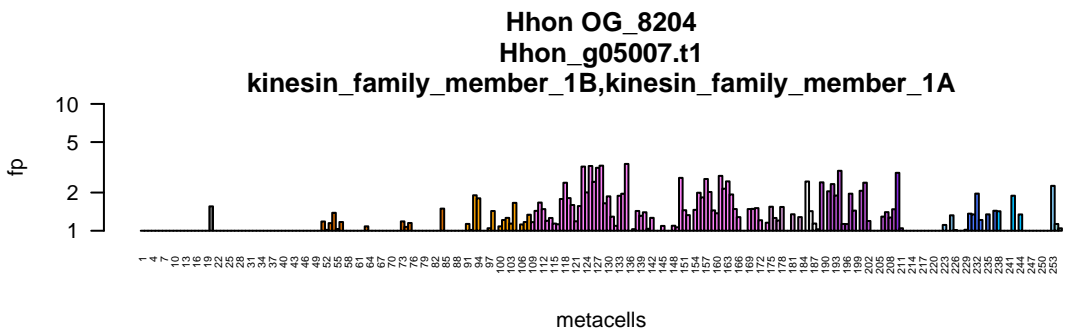
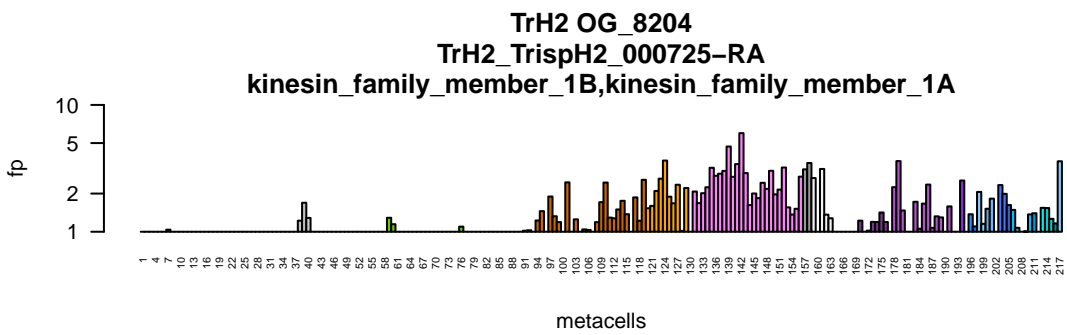
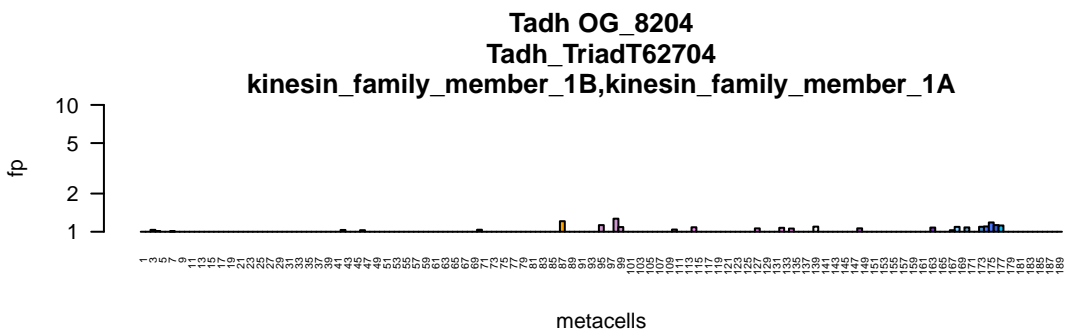
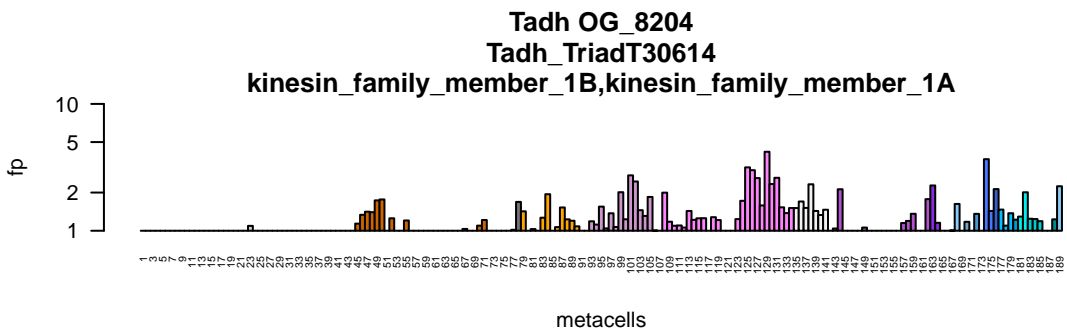


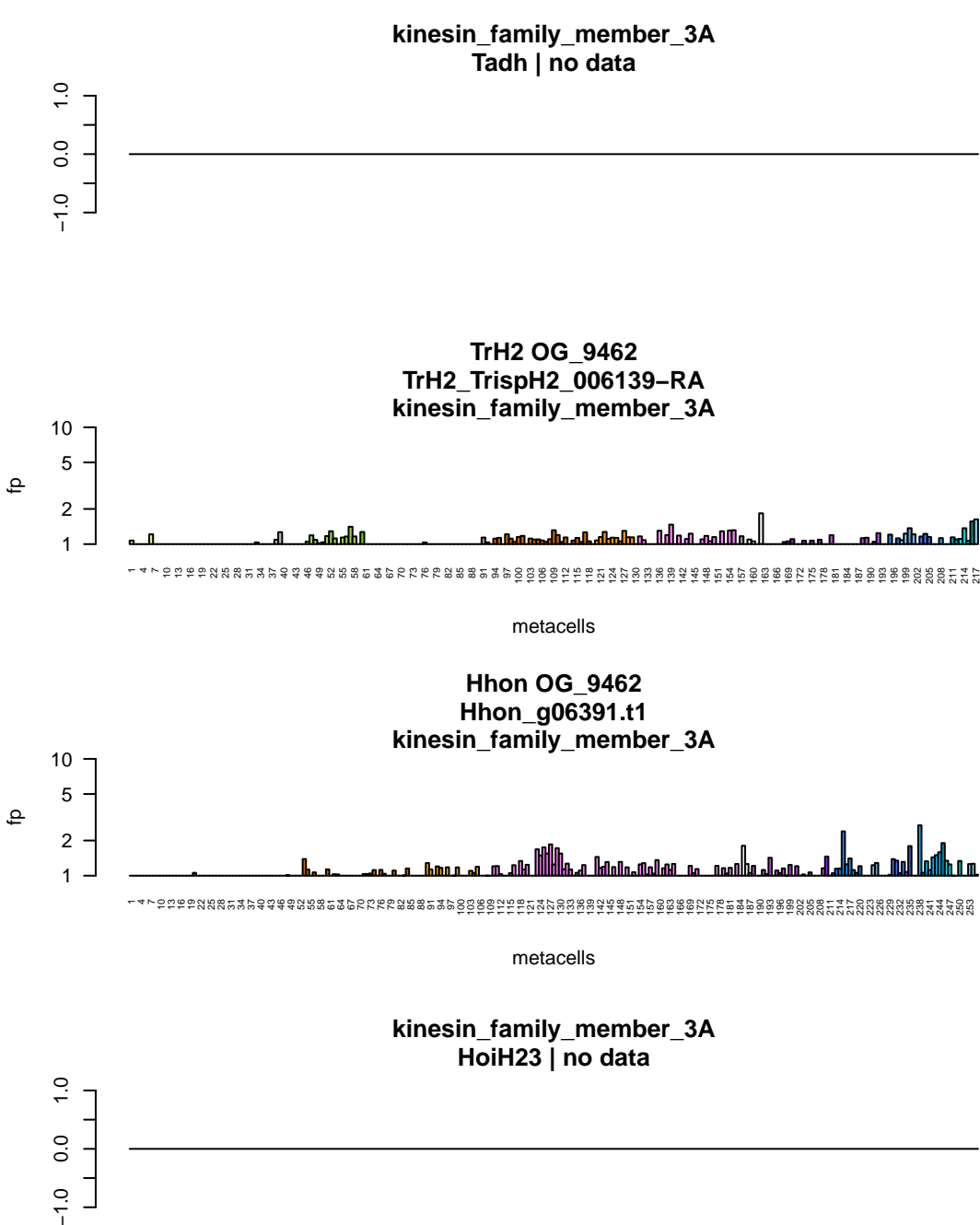








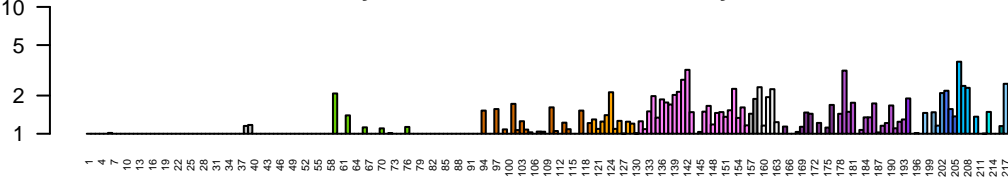




kinesin_family_member_1A,kinesin_family_member_1B
Tadh | no data

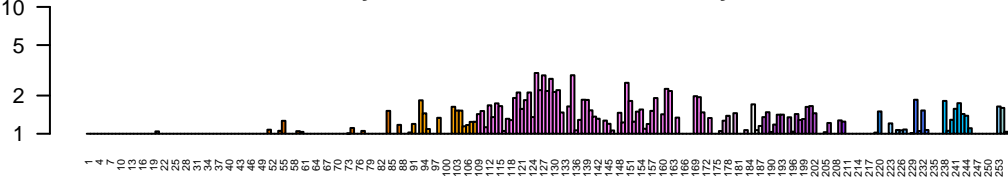


TrH2 OG_9872
TrH2_TrispH2_009952-RA
kinesin_family_member_1A,kinesin_family_member_1B



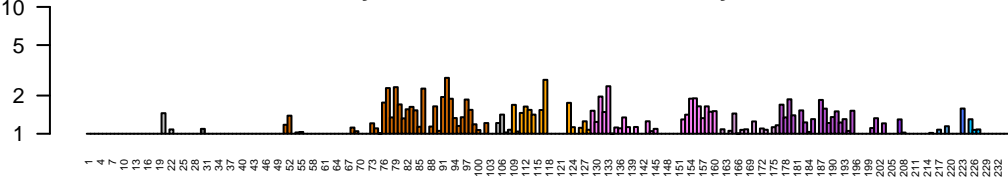
metacells

Hhon OG_9872
Hhon_g04755.t1
kinesin_family_member_1A,kinesin_family_member_1B



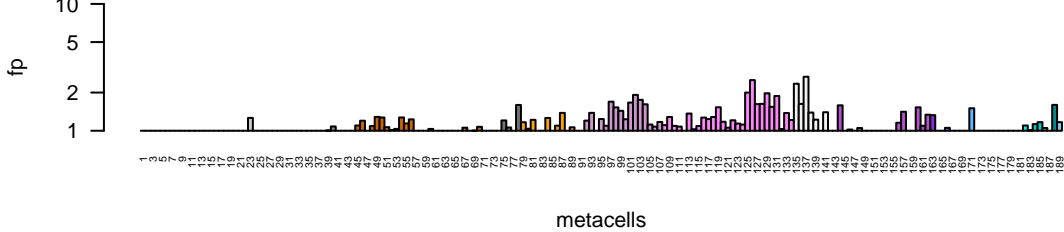
metacells

HoiH23 OG_9872
HoiH23_PIH23_006591-RA
kinesin_family_member_1A,kinesin_family_member_1B

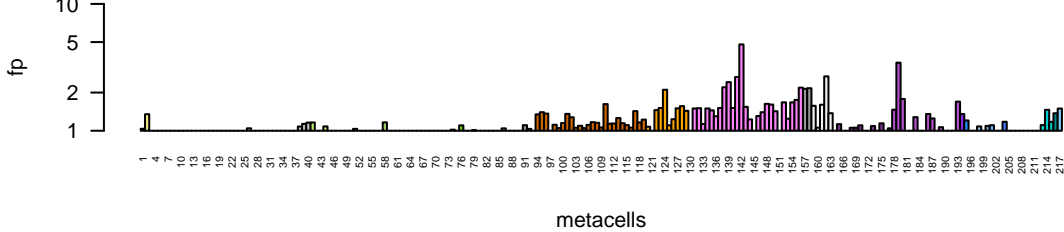


metacells

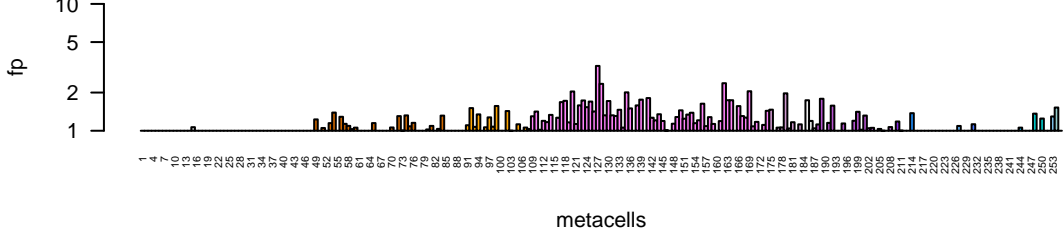
Tadh OG_10098
Tadh_TriadT60039
dynein_axonemal_heavy_chain_3



TrH2 OG_10098
TrH2_TrispH2_010403-RA
dynein_axonemal_heavy_chain_3



Hhon OG_10098
Hhon_g09666.t1
dynein_axonemal_heavy_chain_3



HoiH23 OG_10098
HoiH23_PIH23_006080-RA
dynein_axonemal_heavy_chain_3

