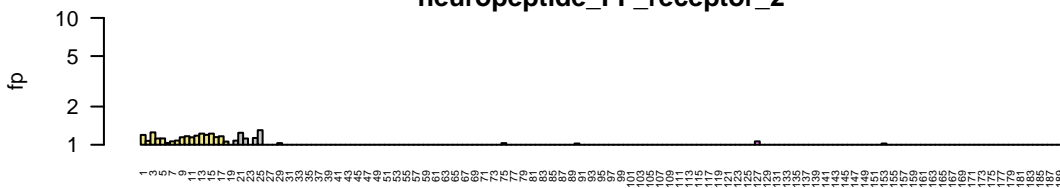
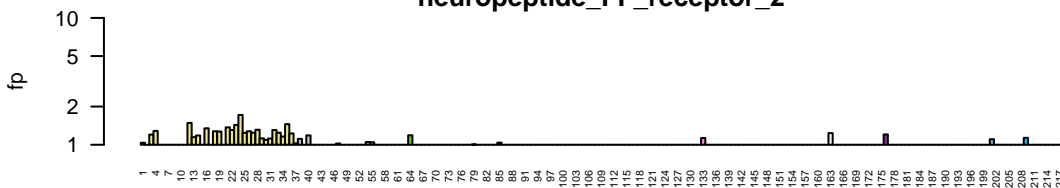


Tadh OG_5409
Tadh_TriadT15472
neuropeptide_FF_receptor_2



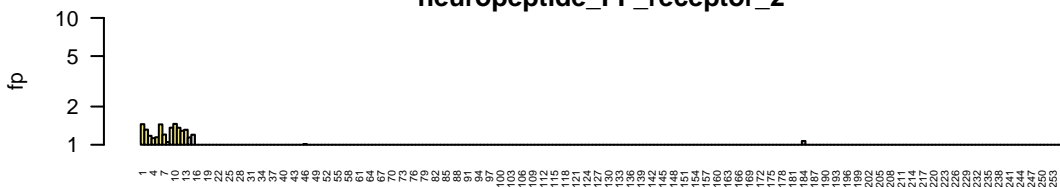
metacells

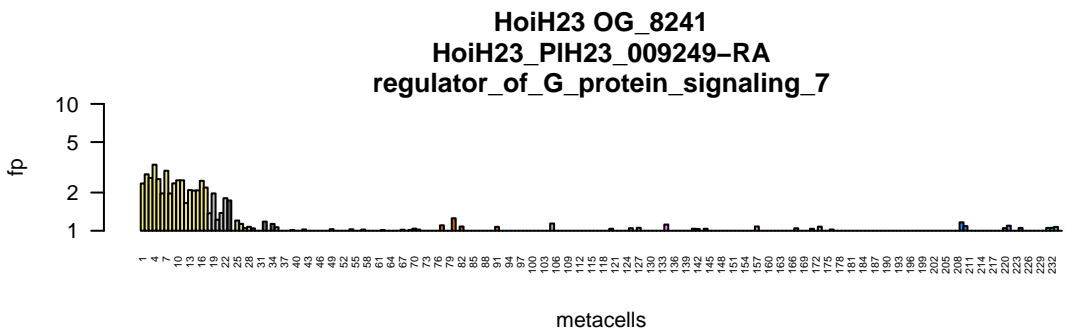
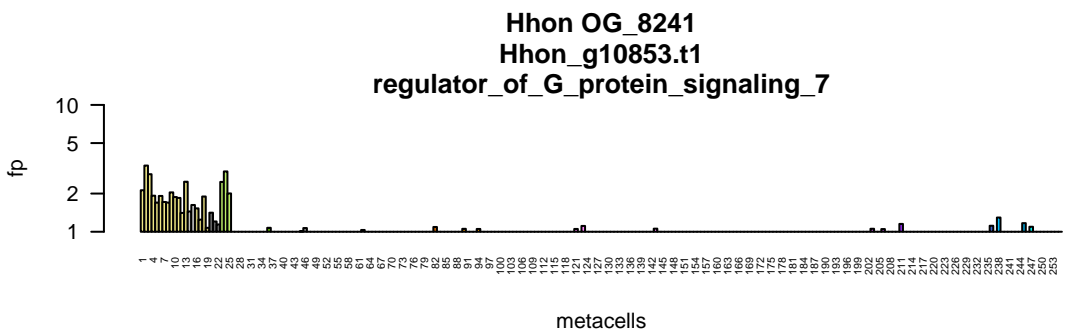
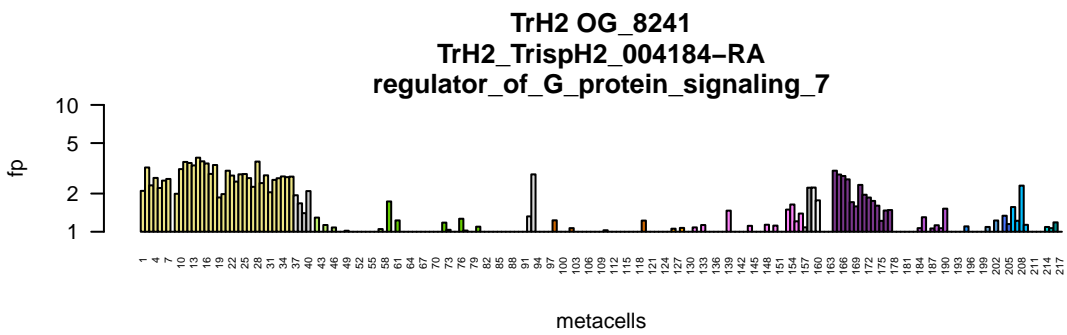
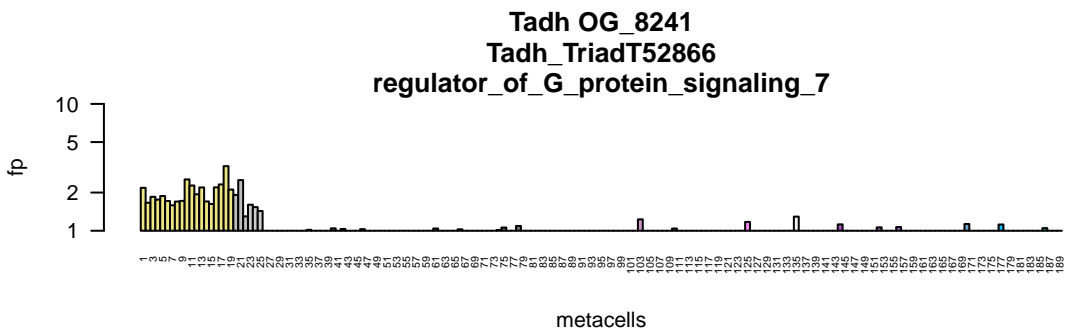
TrH2 OG_5409
TrH2_TrispH2_008515-RA
neuropeptide_FF_receptor_2

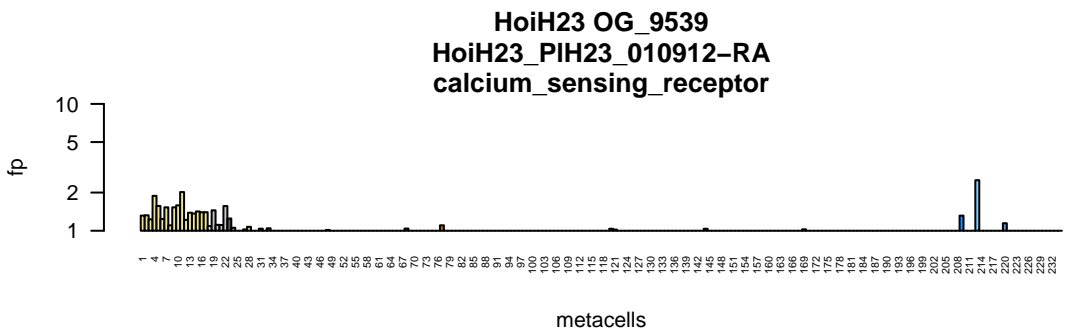
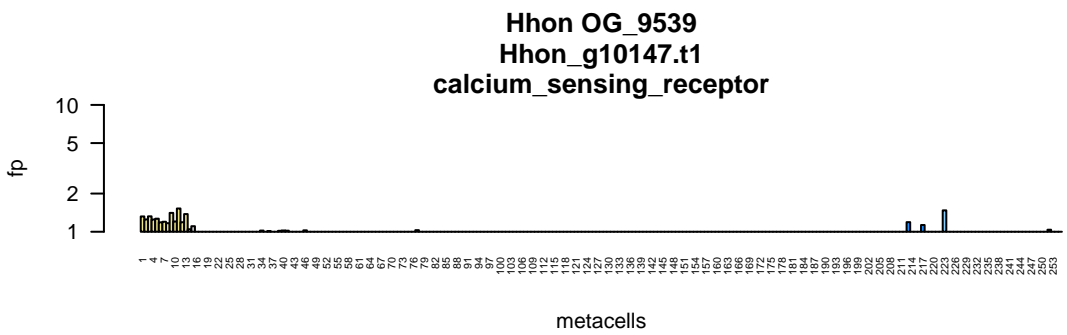
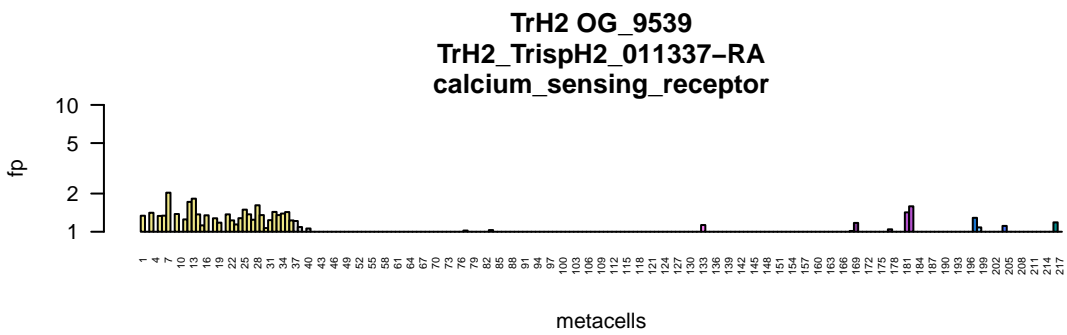
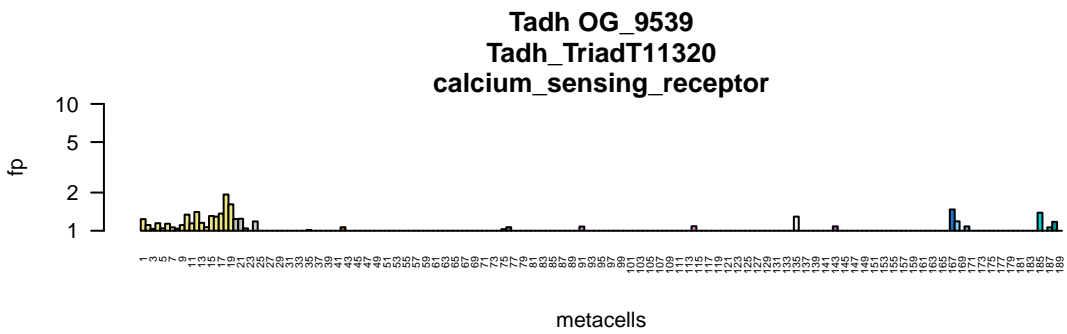


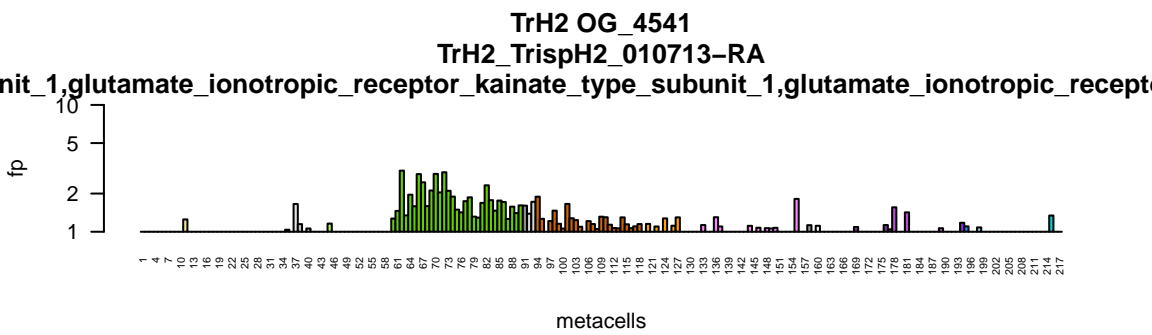
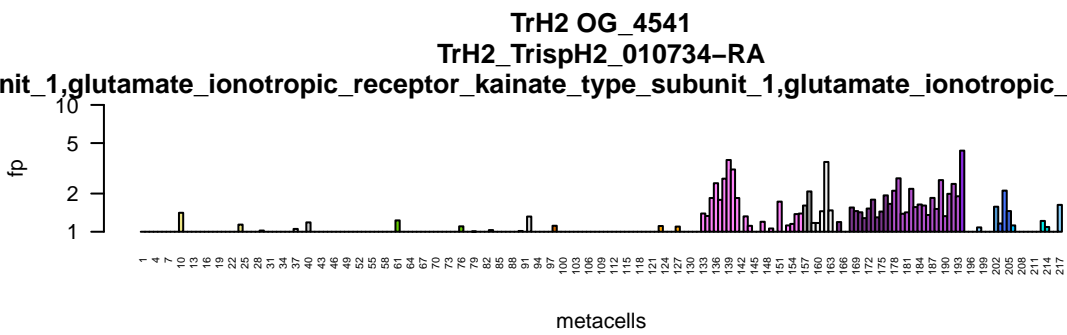
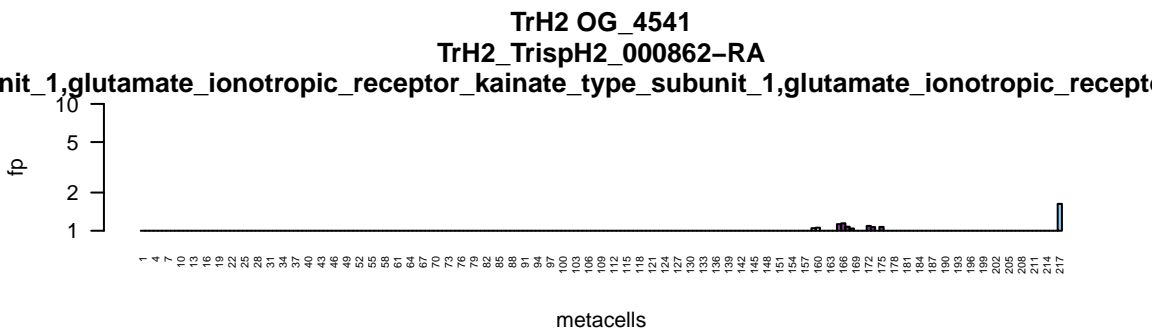
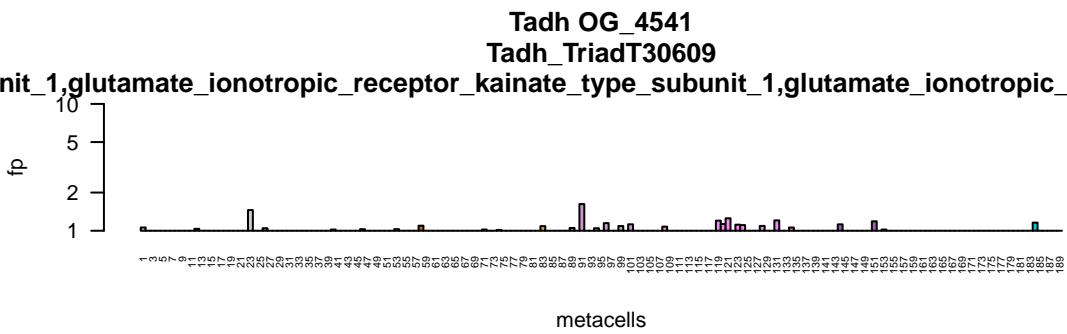
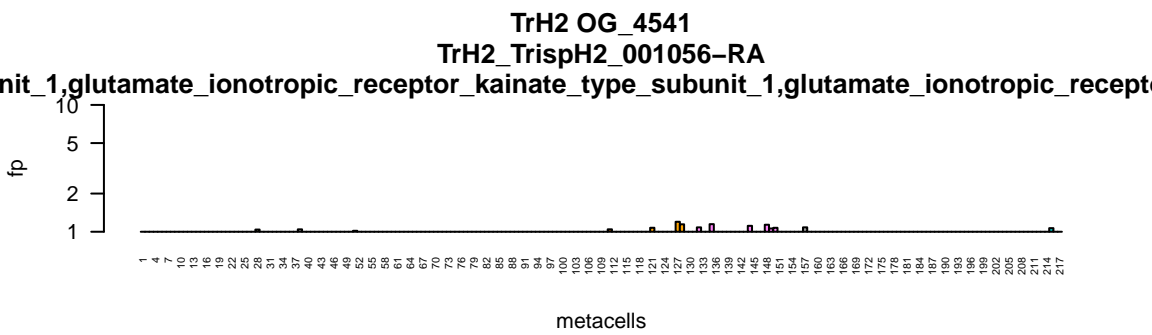
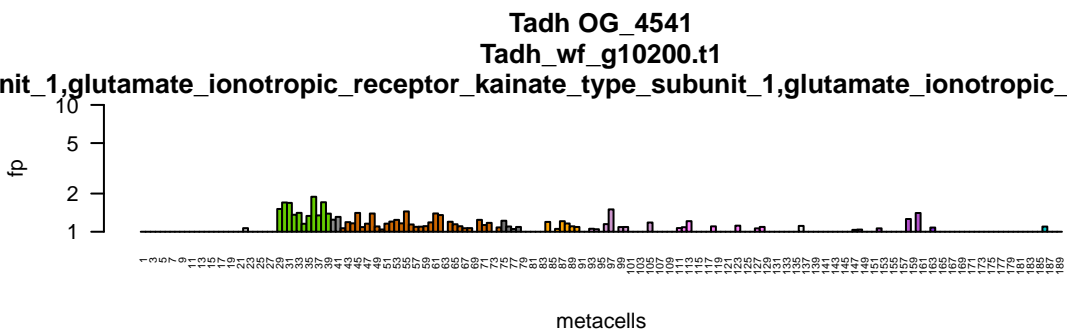
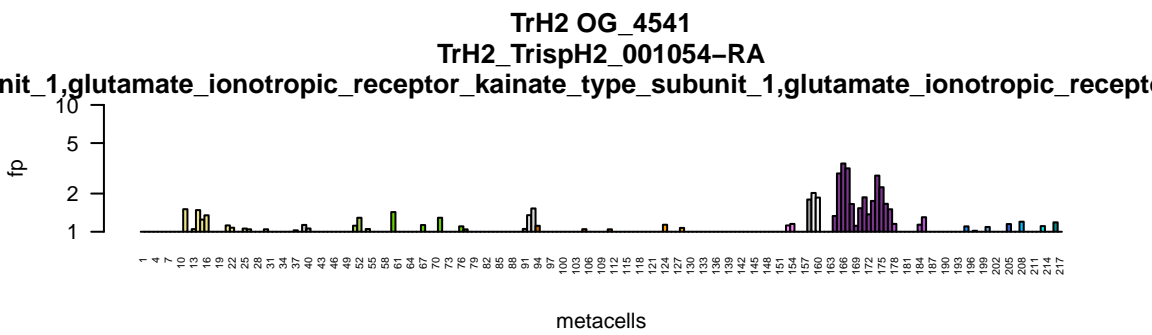
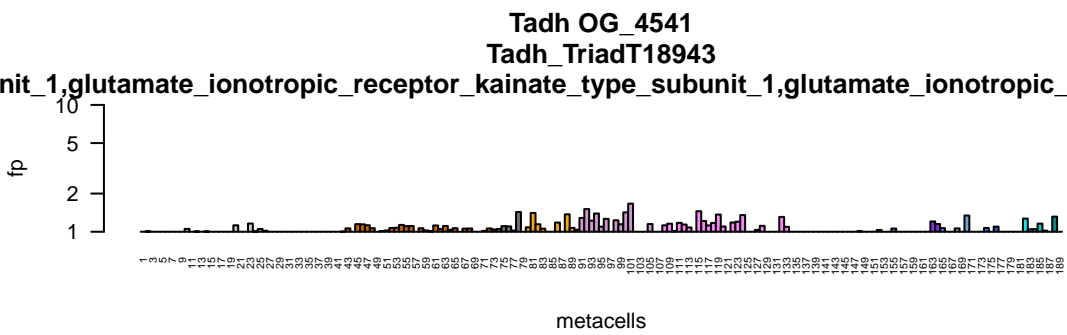
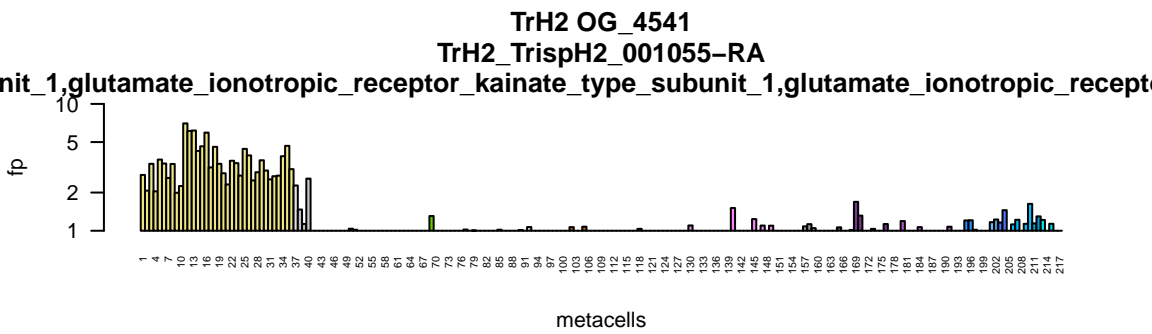
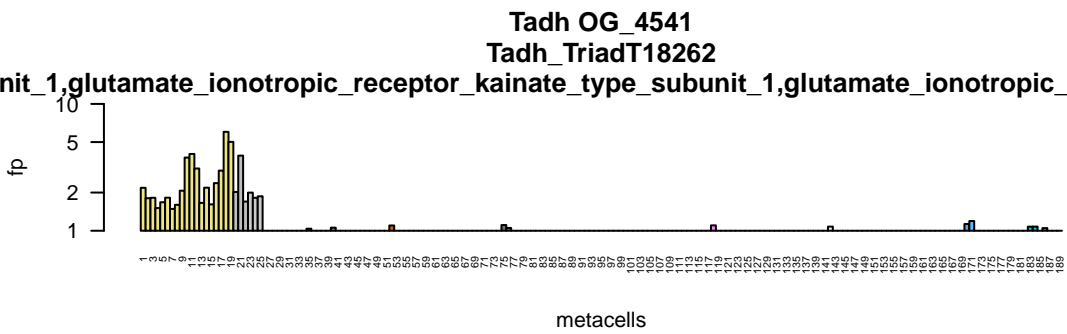
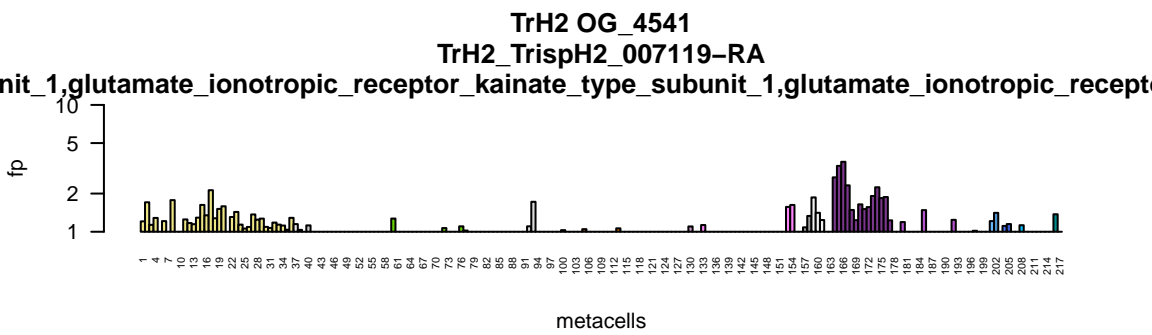
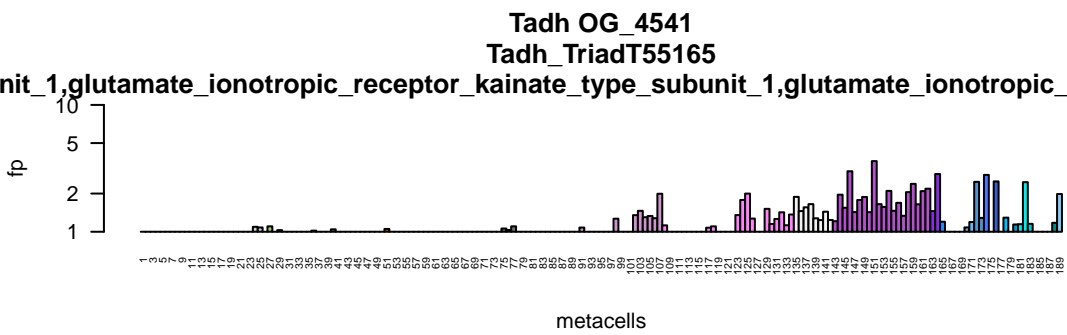
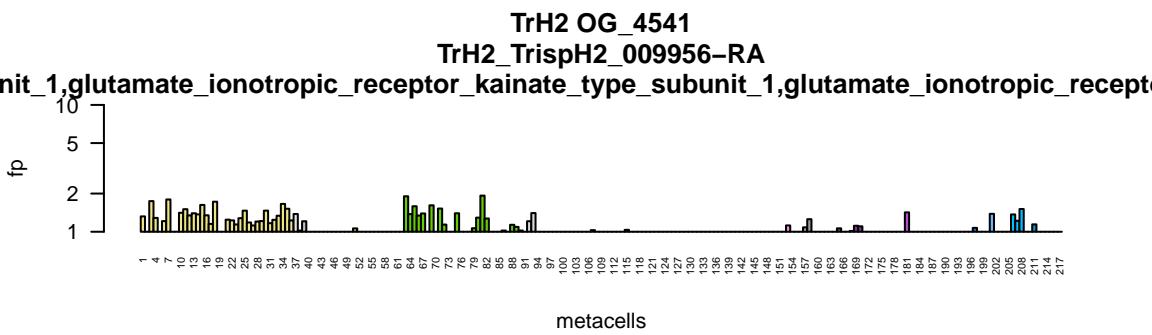
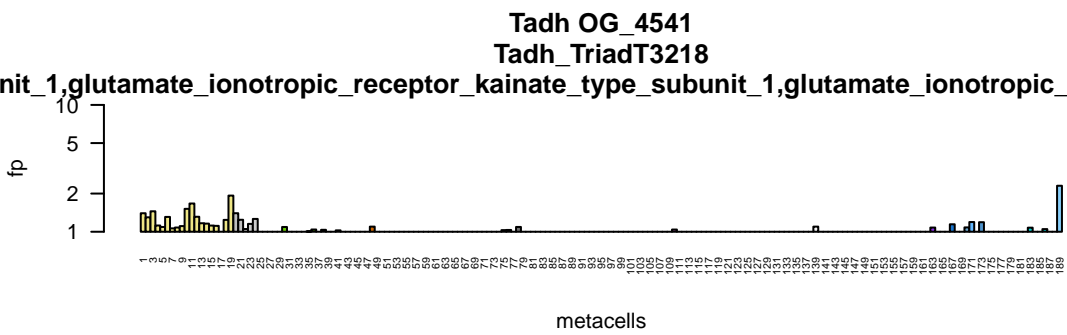
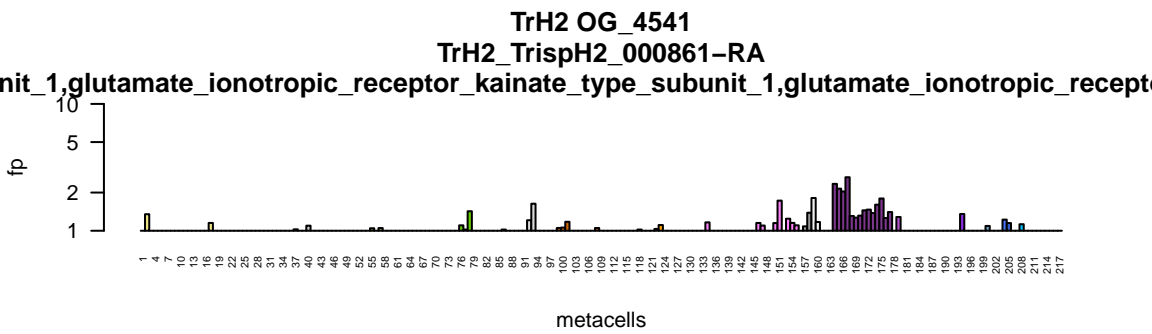
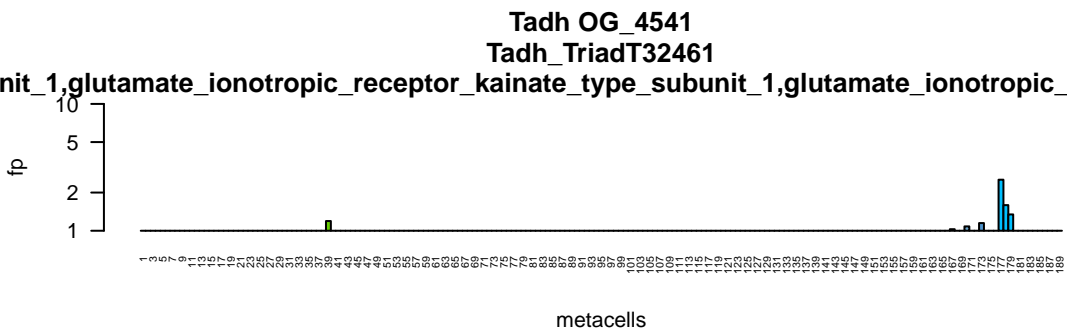
metacells

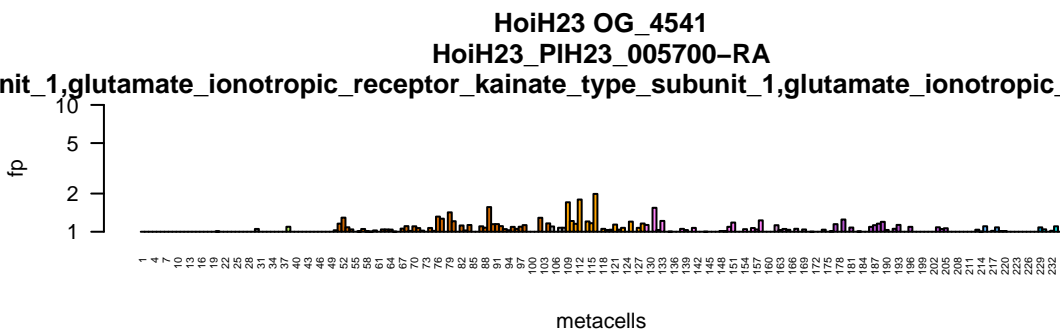
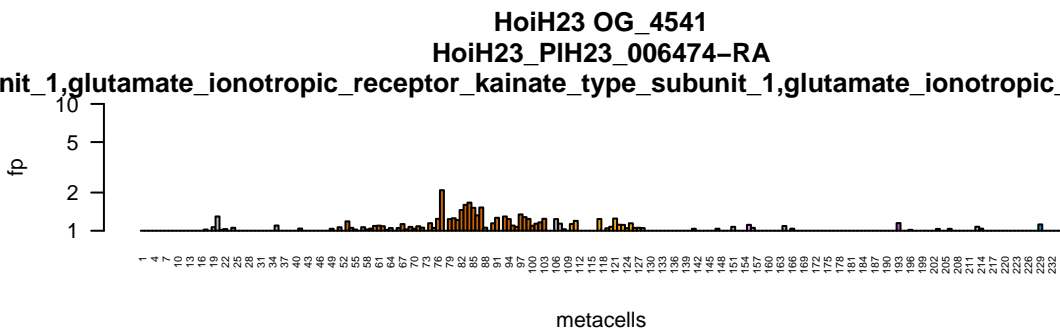
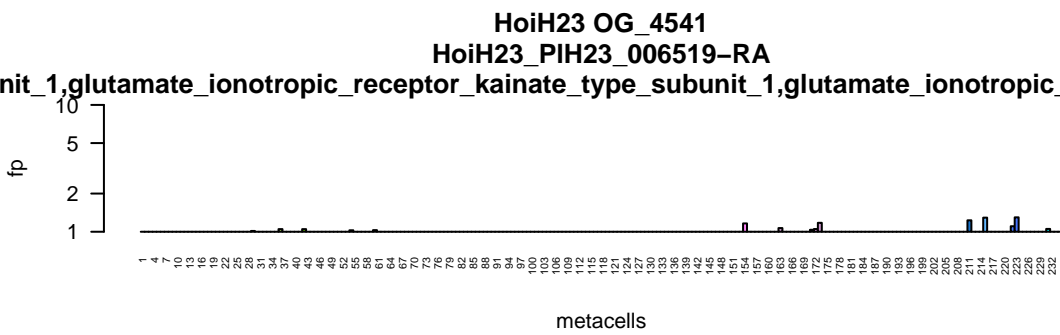
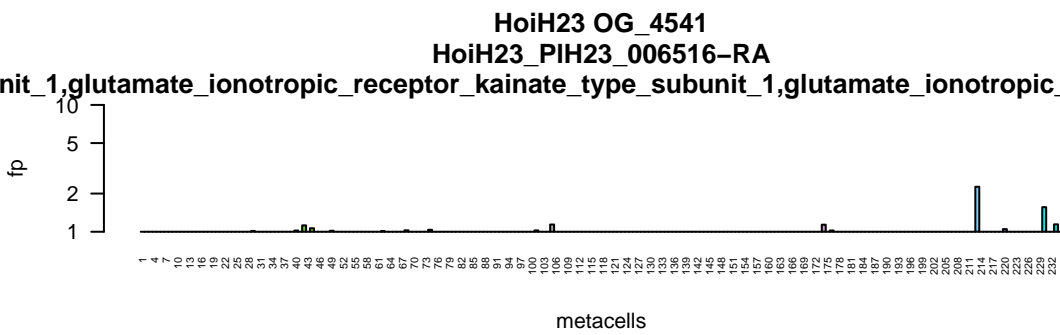
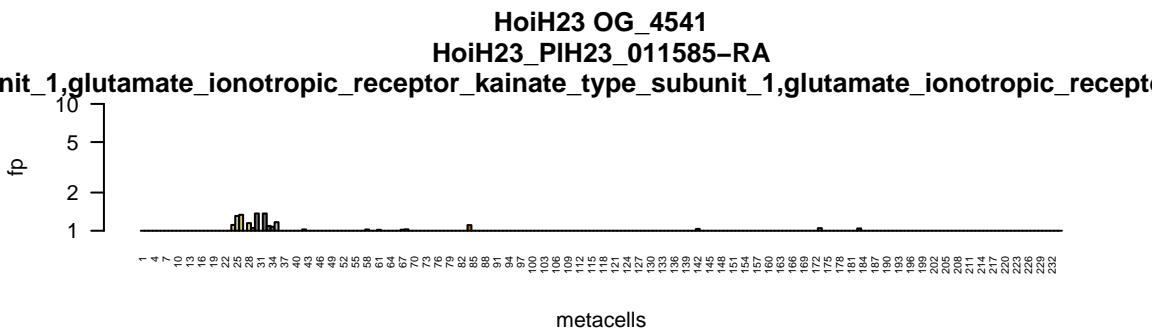
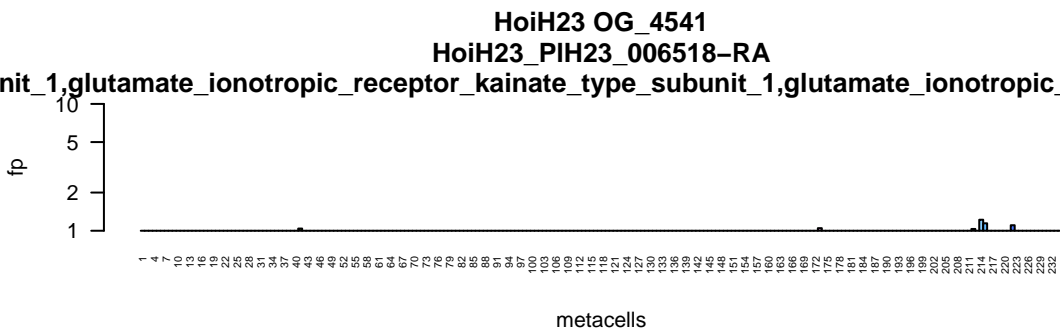
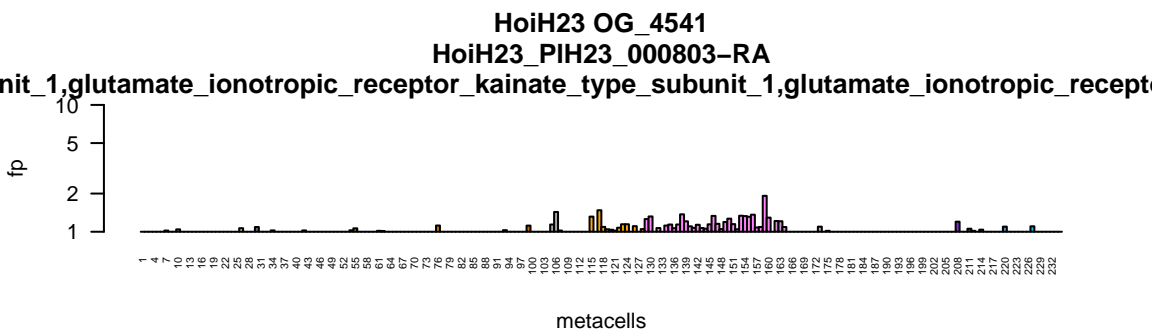
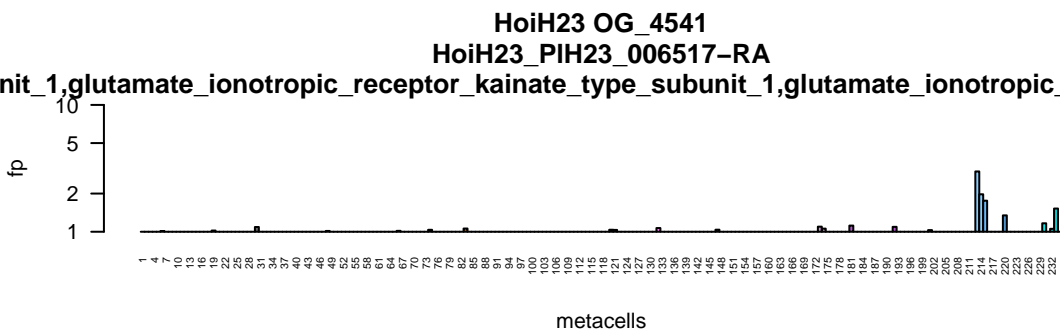
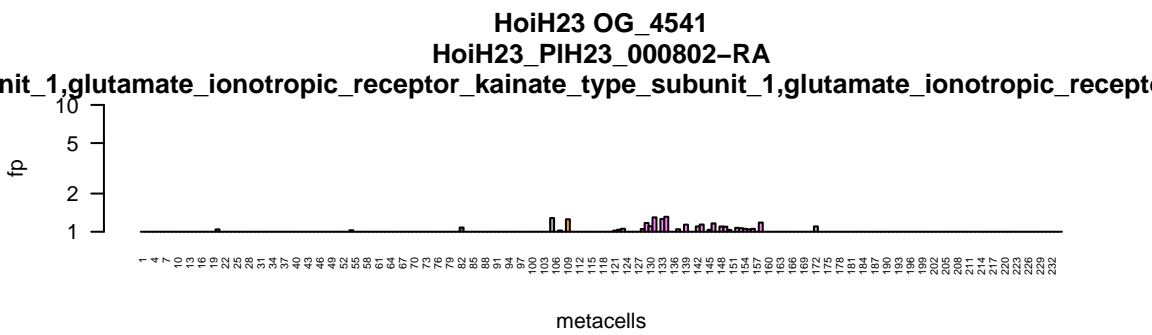
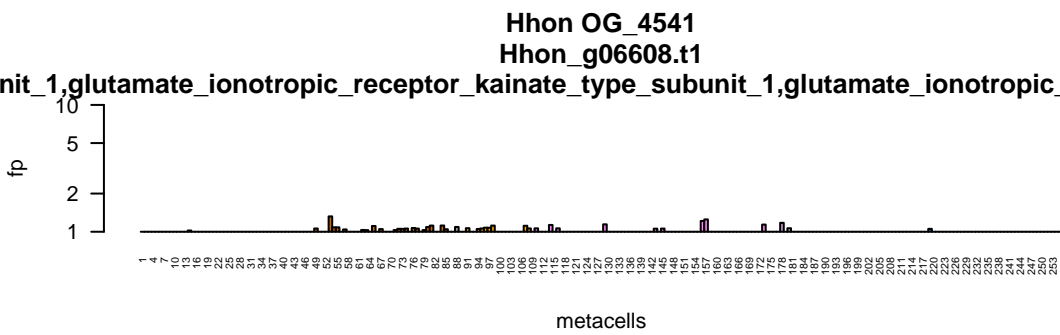
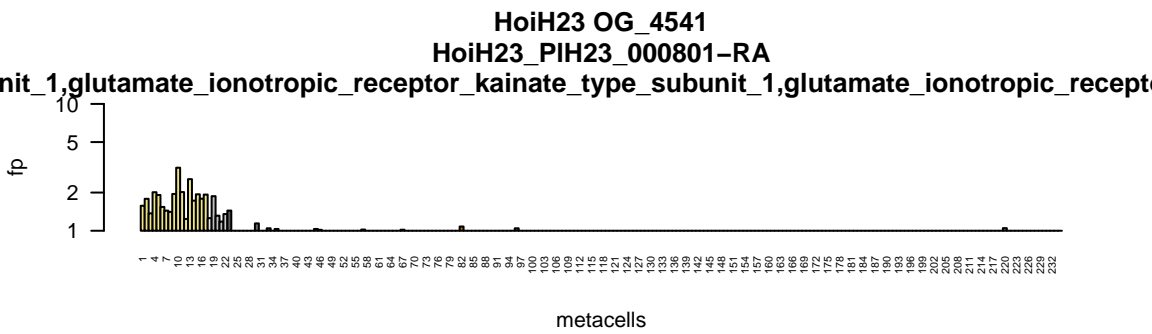
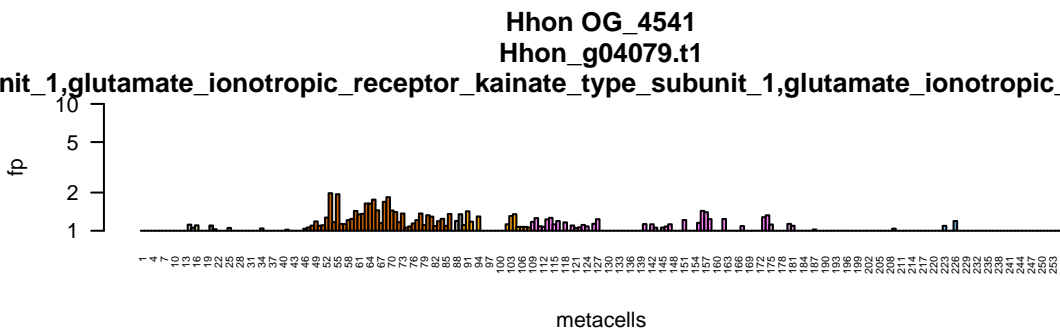
Hhon OG_5409
Hhon_g04180.t1
neuropeptide_FF_receptor_2

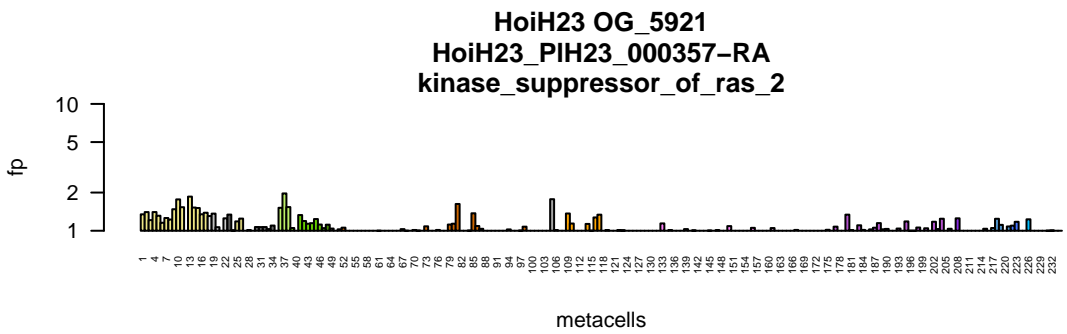
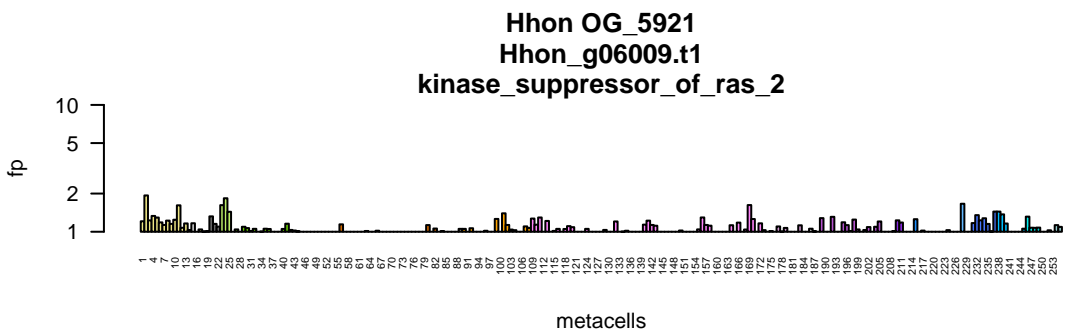
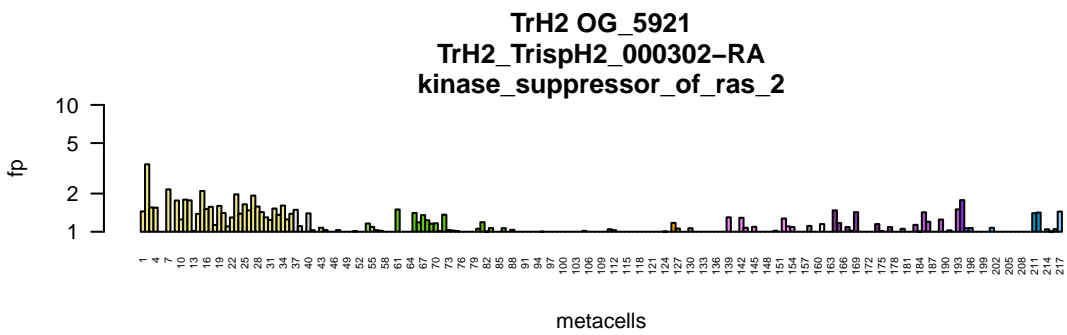
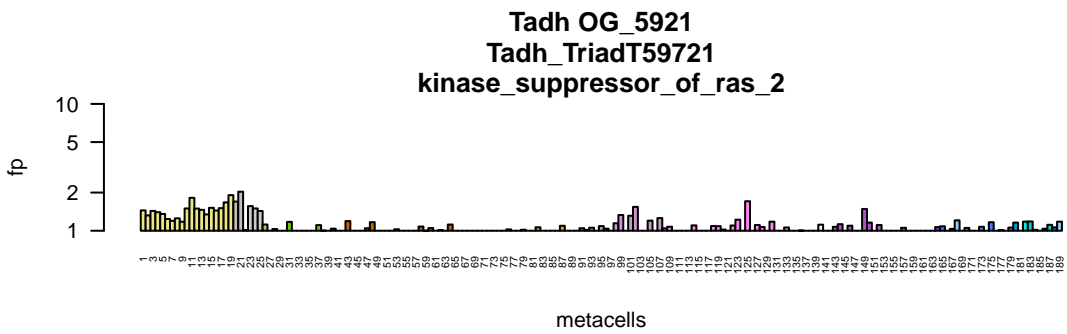


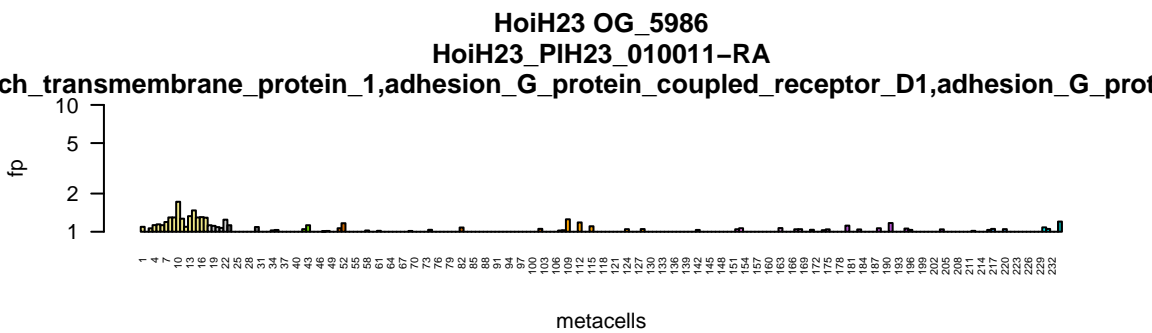
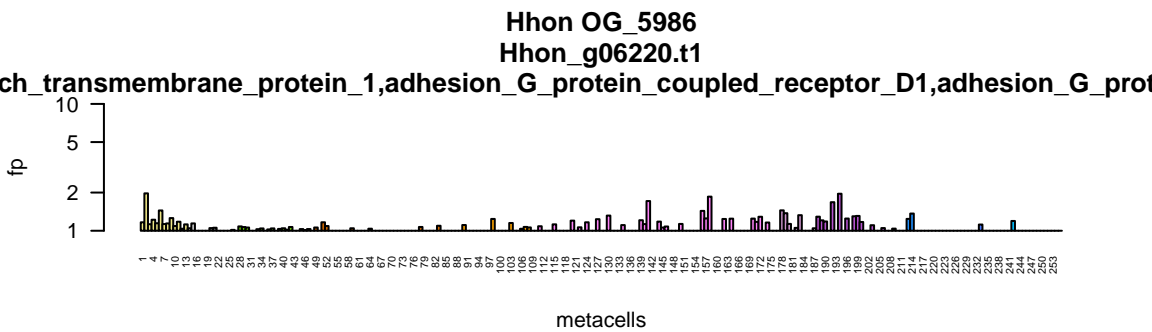
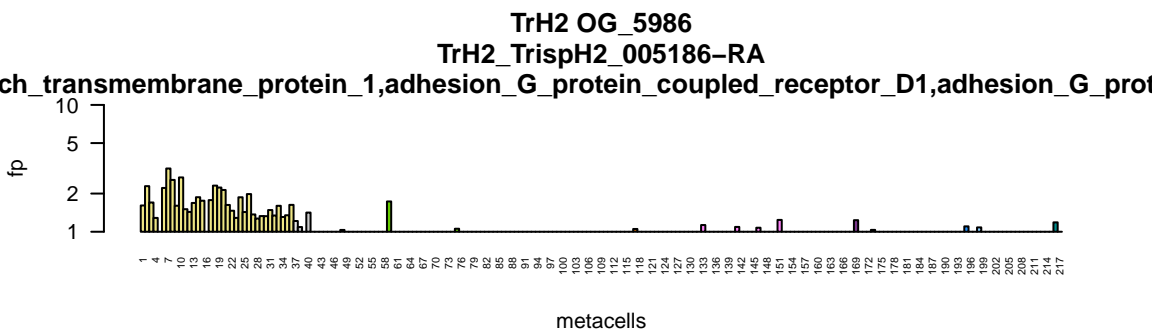
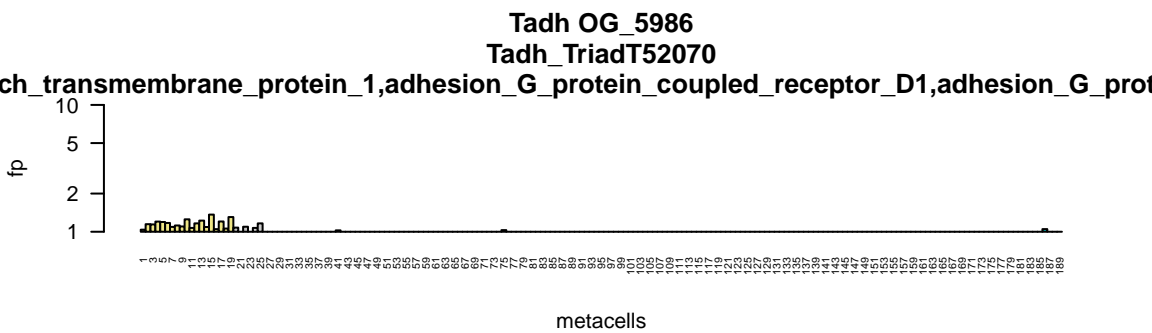
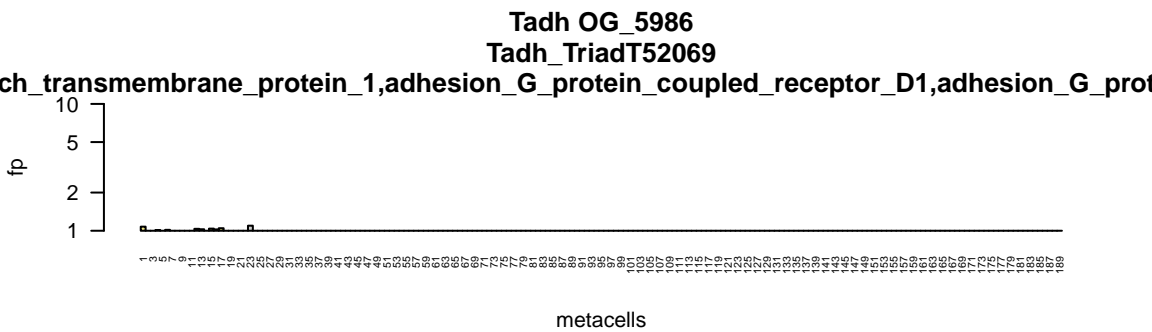


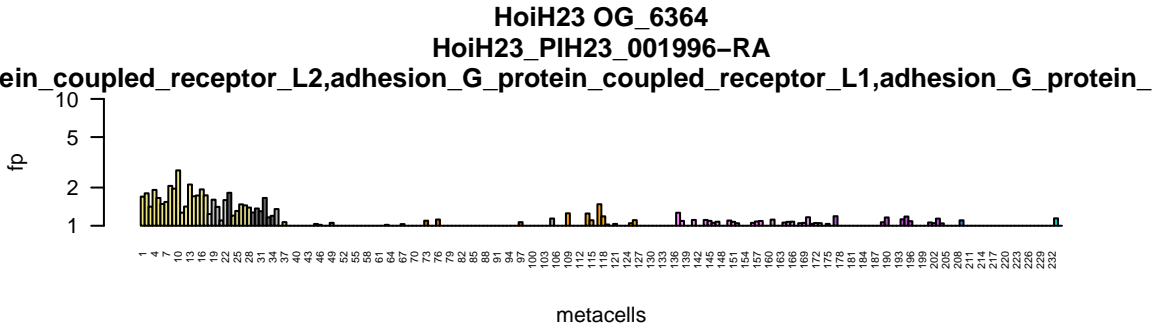
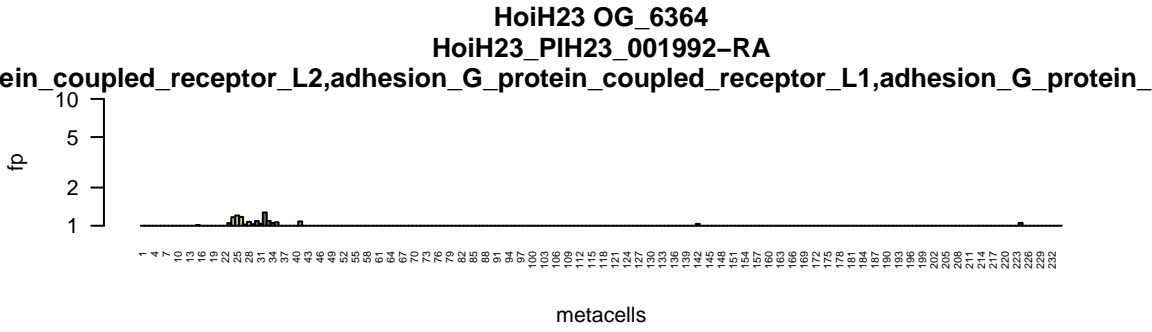
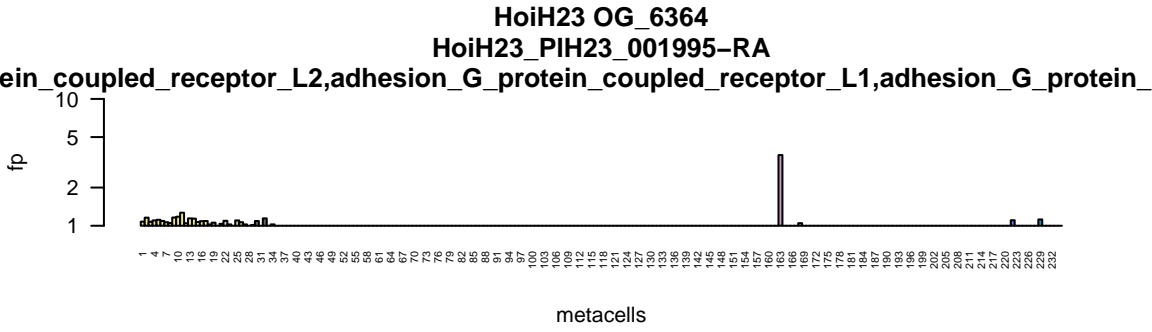
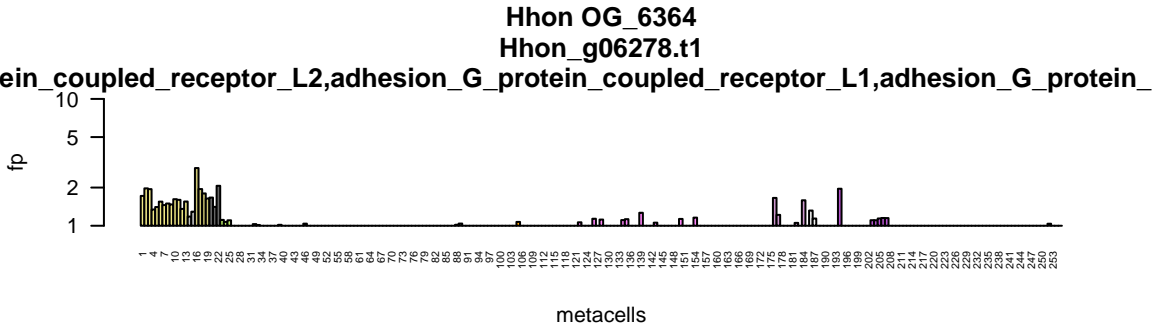
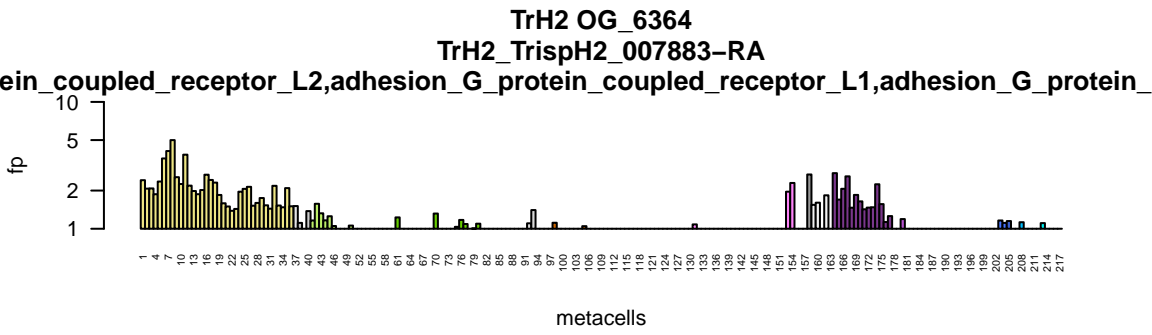
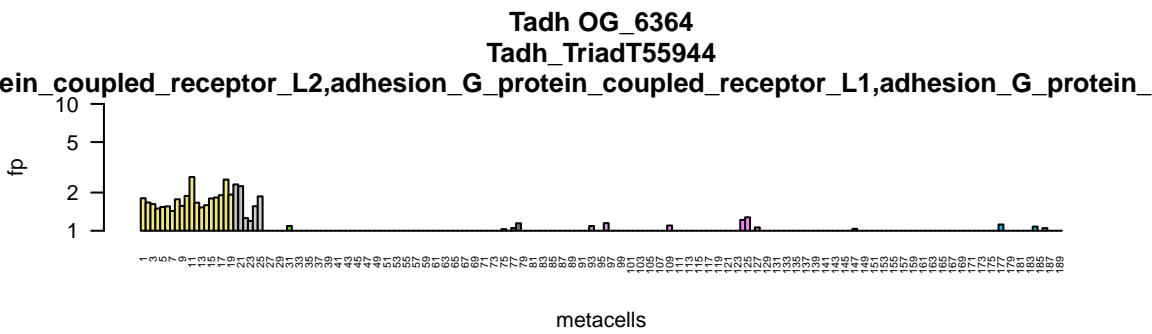


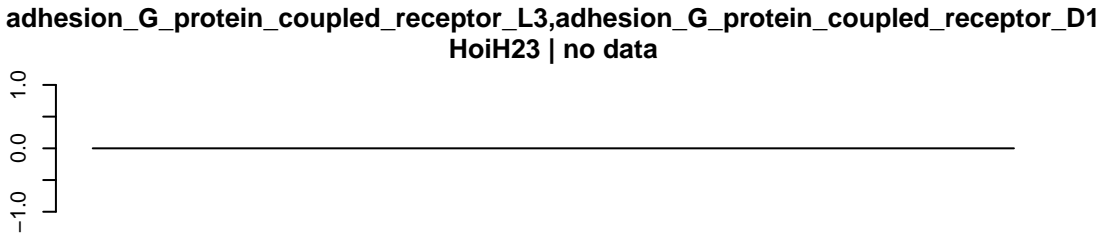
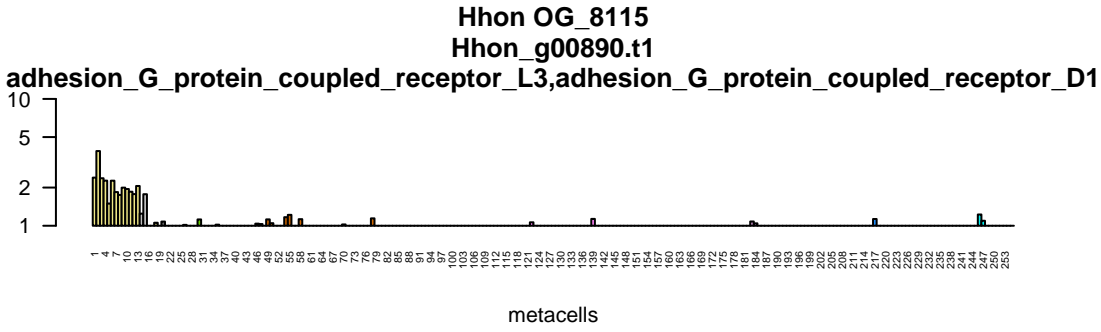
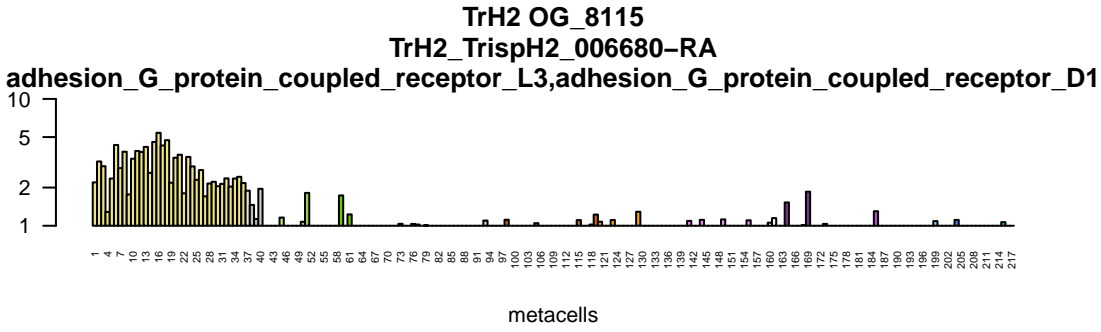
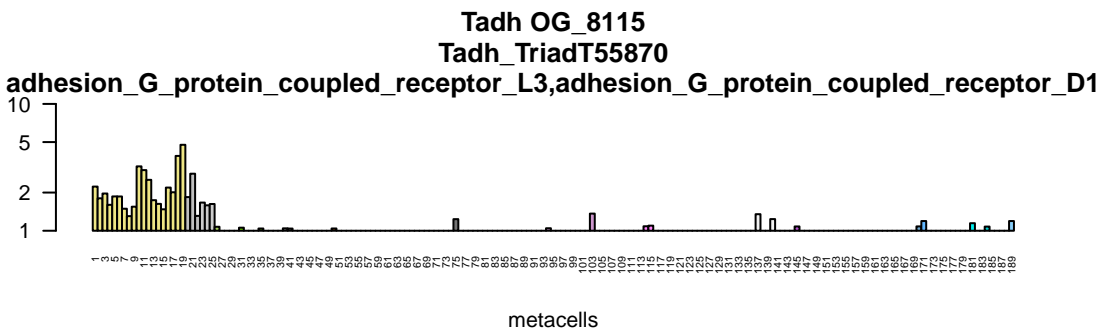


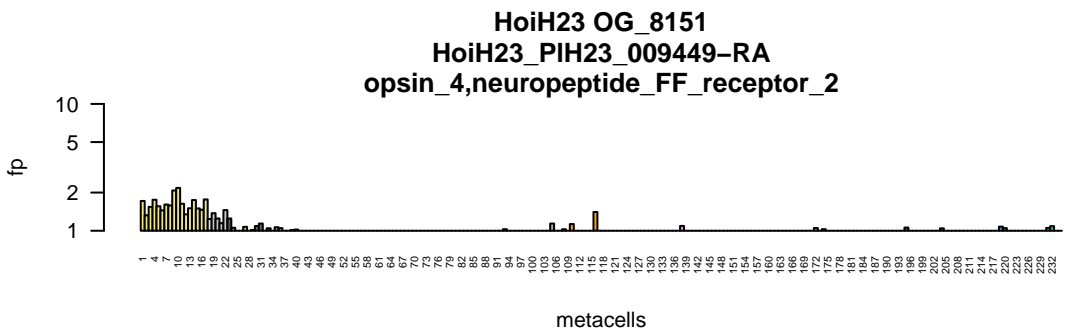
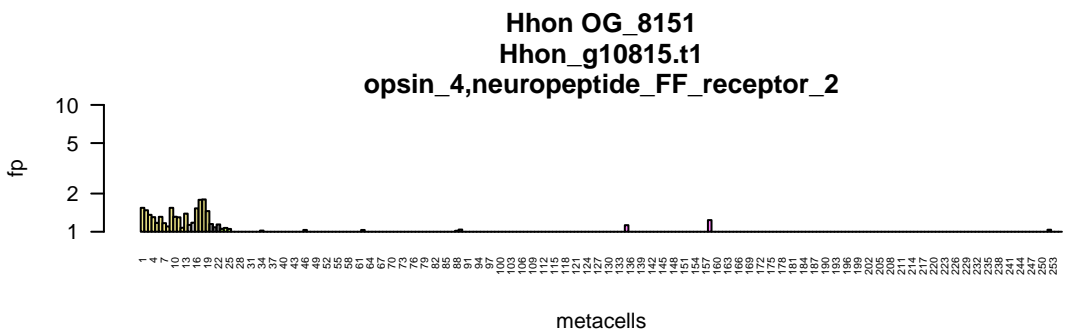
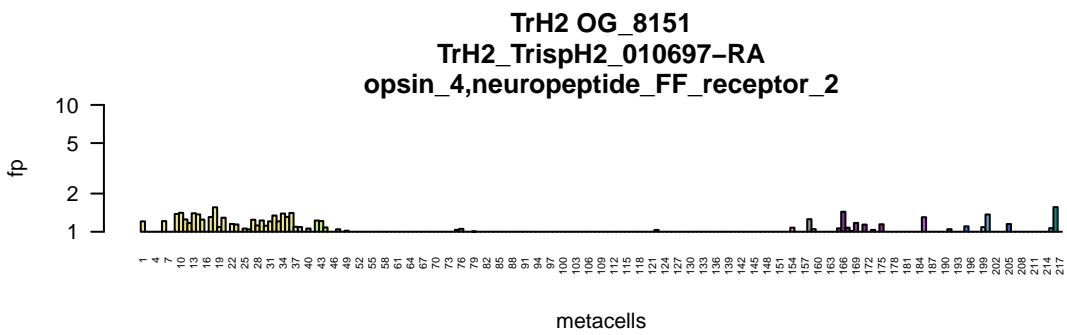
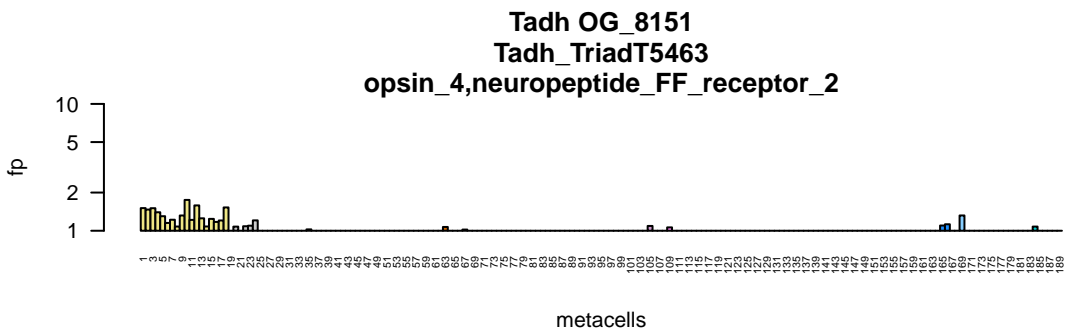








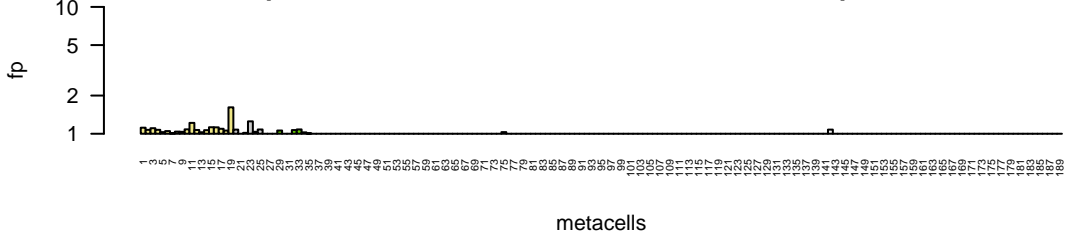




Tadh OG_8575

Tadh_TriadT56776

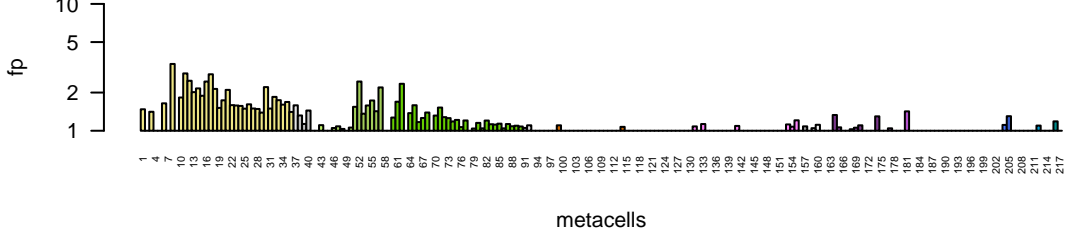
interleukin_1_receptor_associated_kinase_4,interleukin_1_receptor_associated_kinase_4



TrH2 OG_8575

TrH2_TrispH2_008687-RA

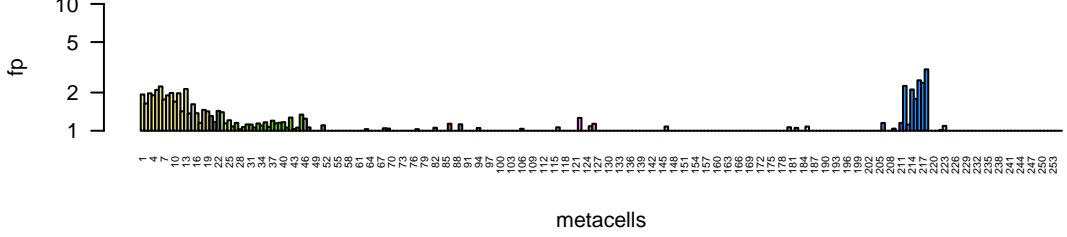
interleukin_1_receptor_associated_kinase_4,interleukin_1_receptor_associated_kinase_4



Hhon OG_8575

Hhon_g09836.t1

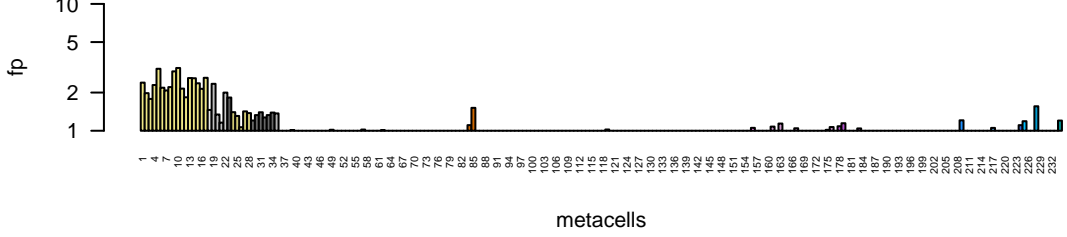
interleukin_1_receptor_associated_kinase_4,interleukin_1_receptor_associated_kinase_4



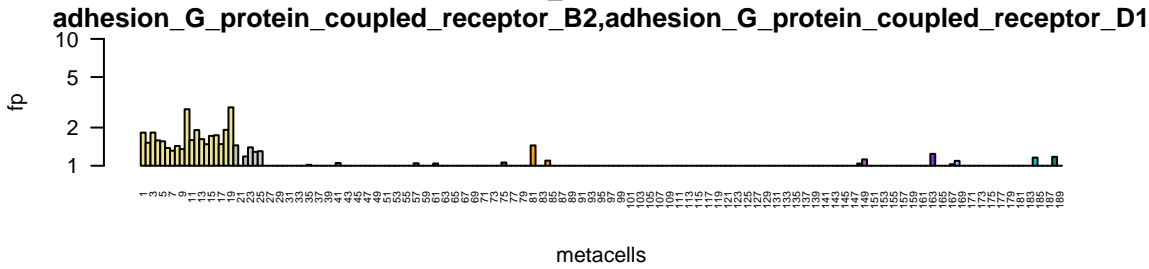
HoiH23 OG_8575

HoiH23_PIH23_007836-RA

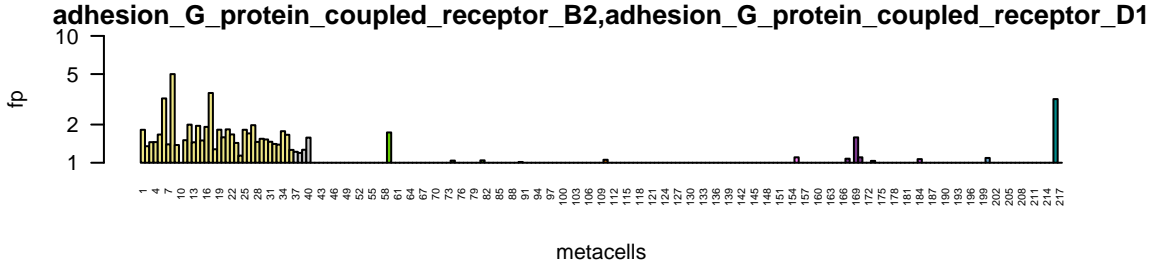
interleukin_1_receptor_associated_kinase_4,interleukin_1_receptor_associated_kinase_4



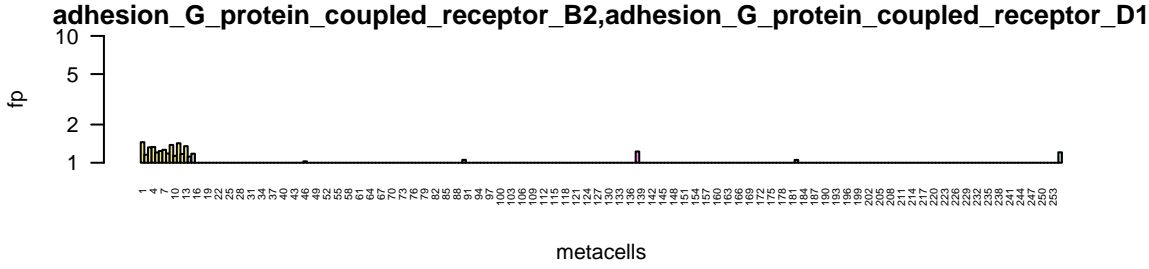
Tadh OG_9913
Tadh_TriadT57964



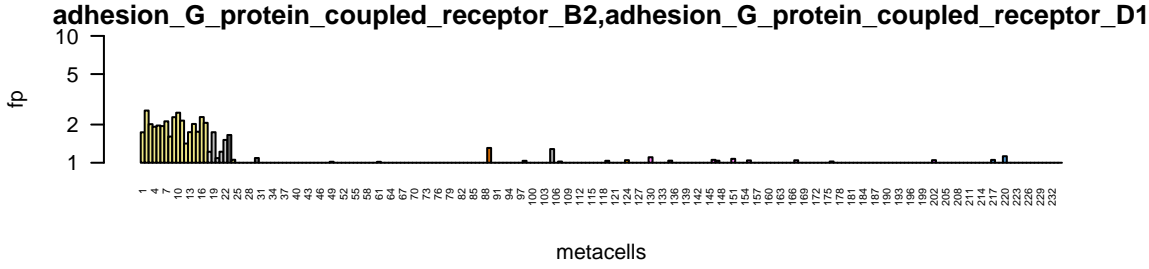
TrH2 OG_9913
TrH2_TrispH2_006500-RA

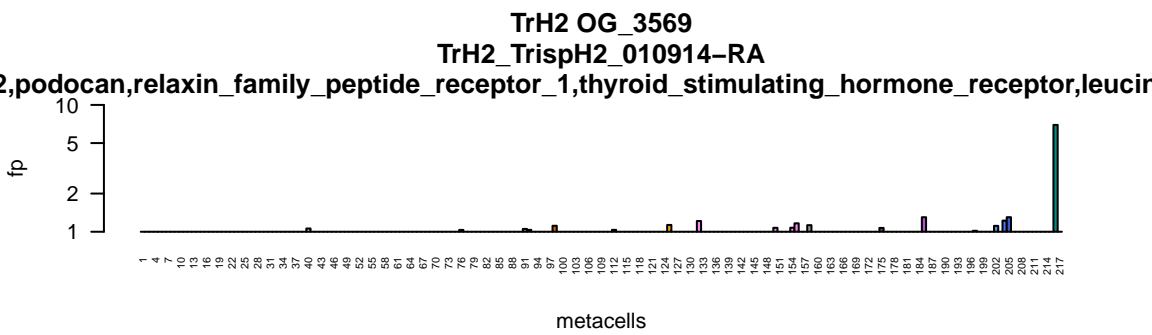
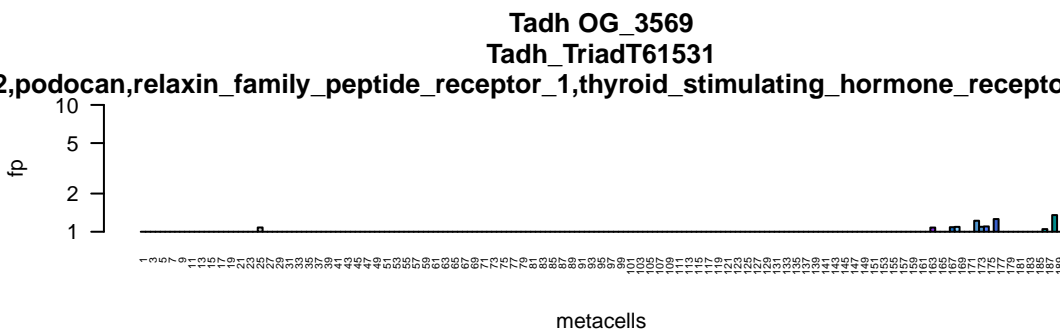
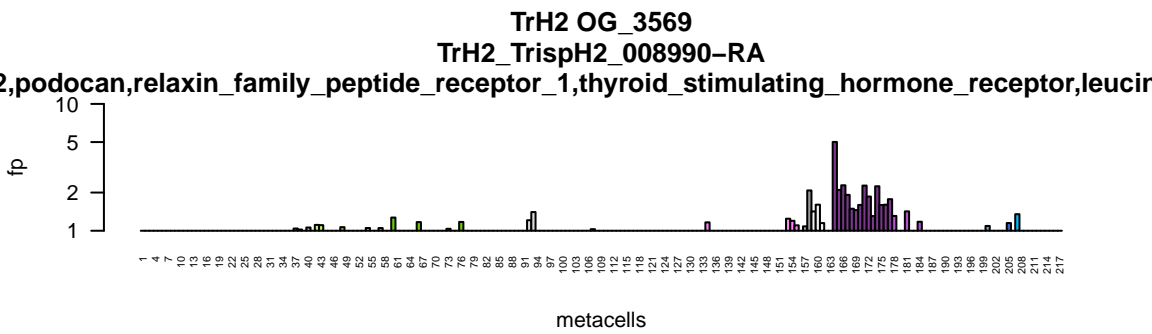
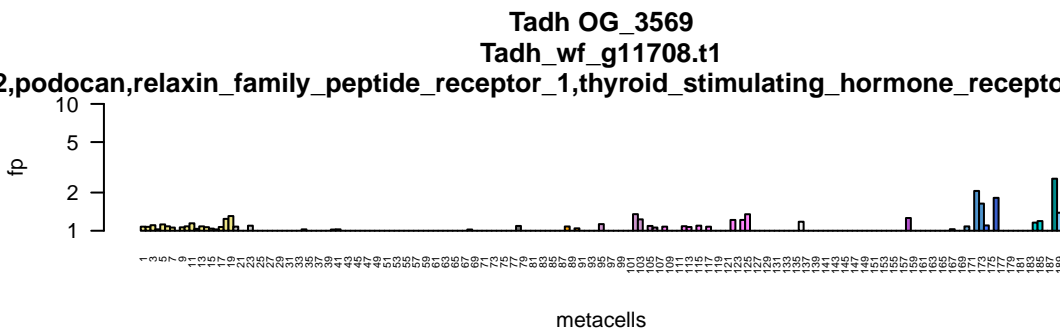
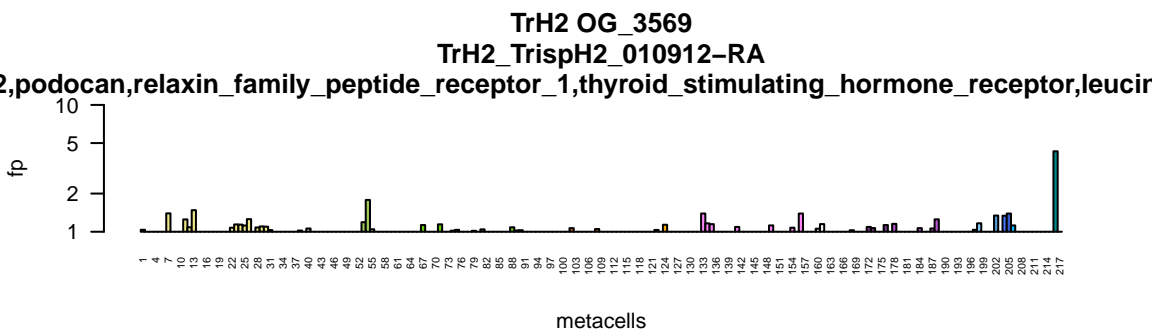
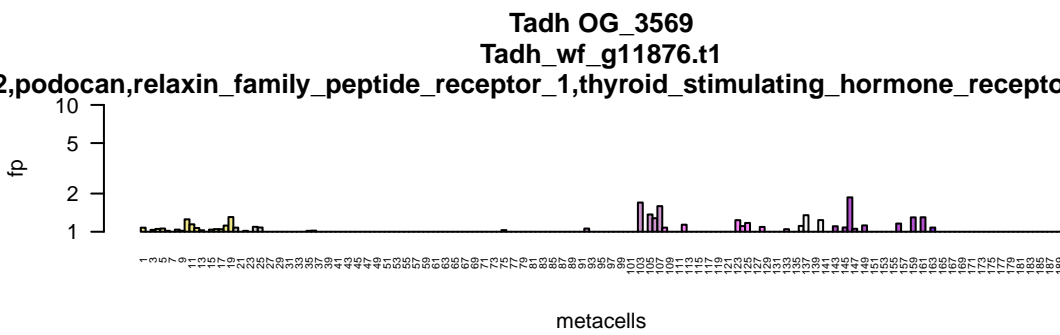
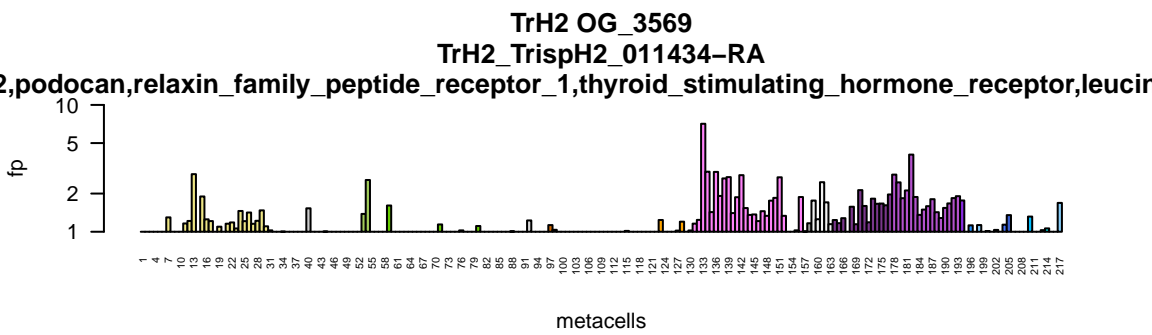
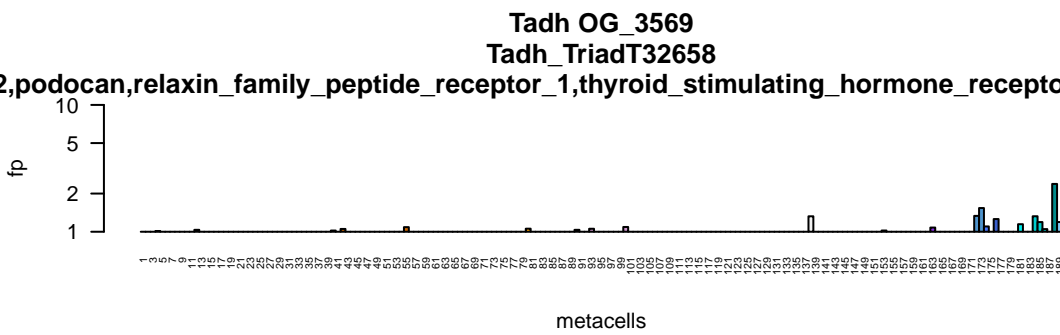
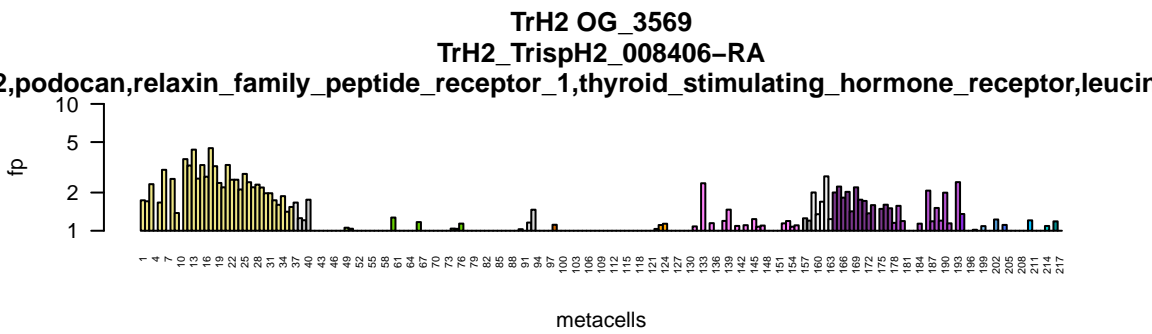
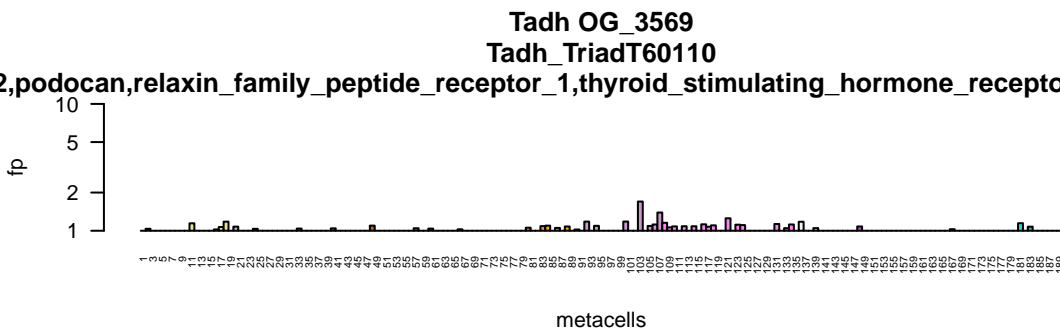
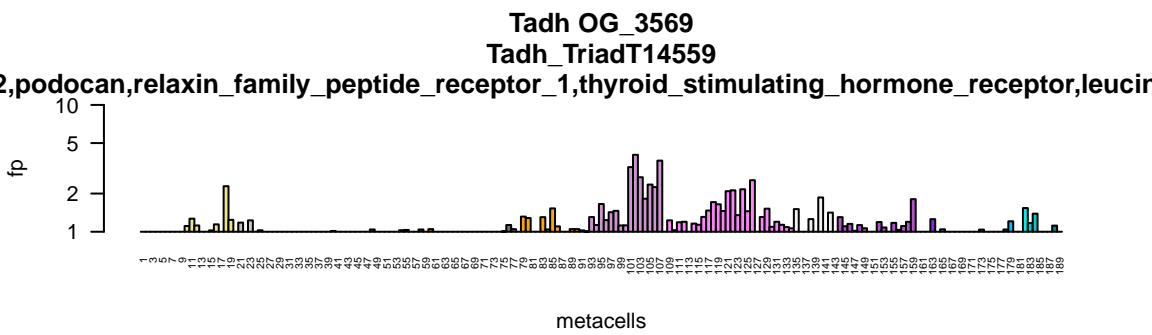
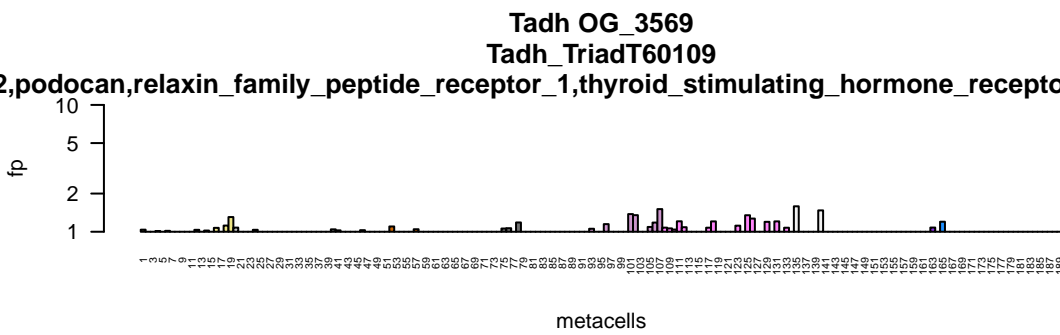
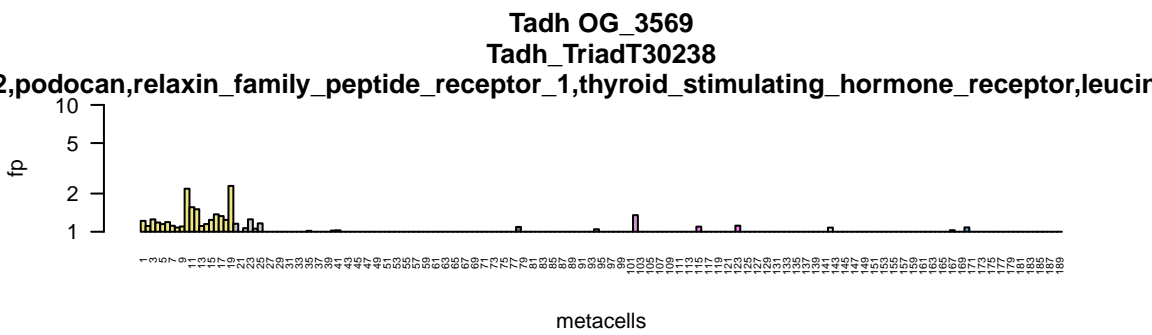
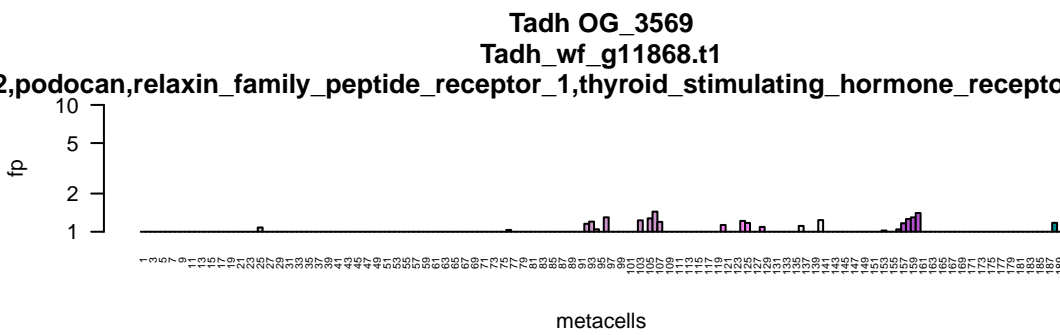
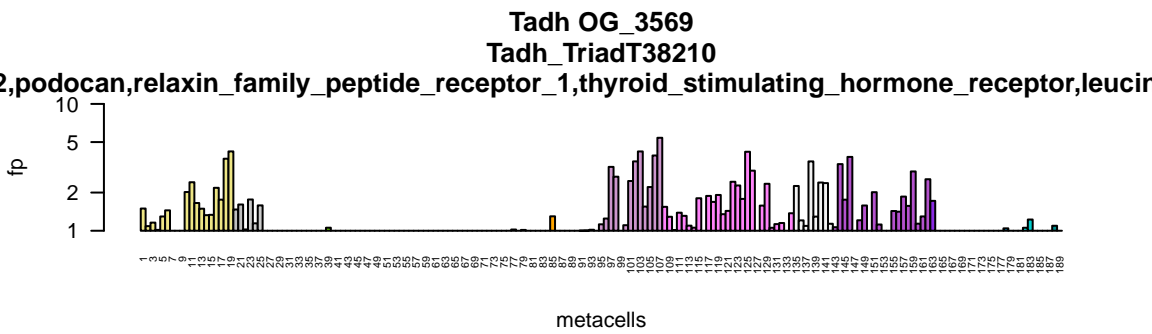
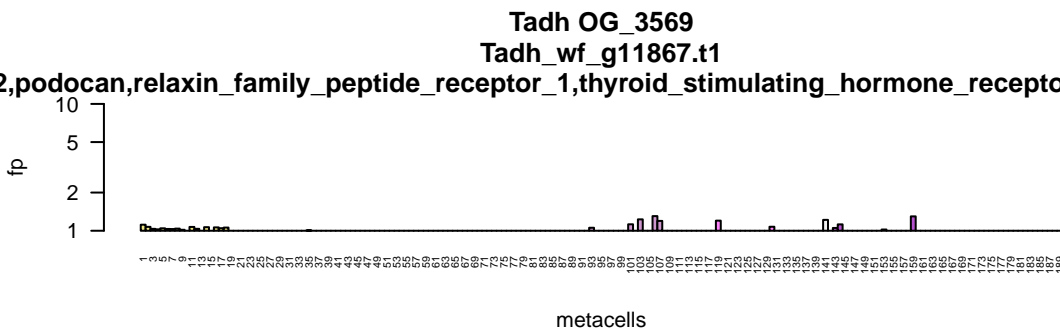


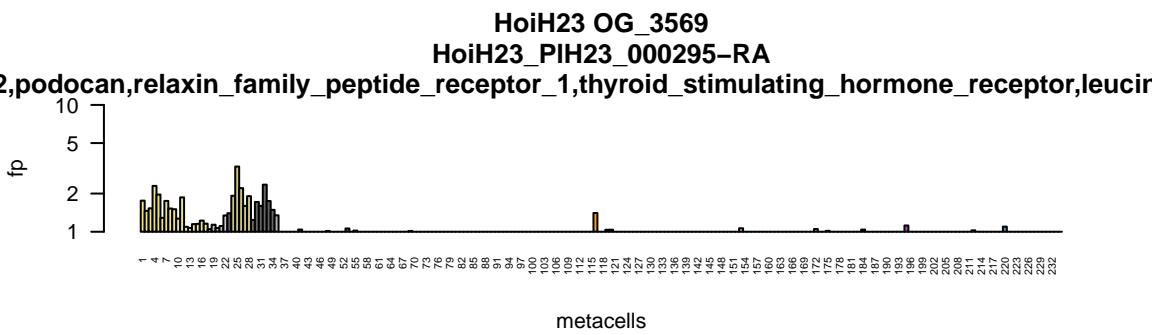
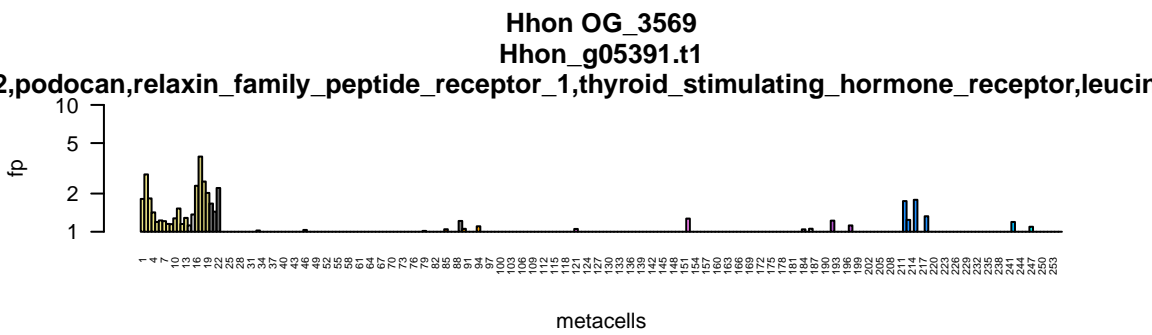
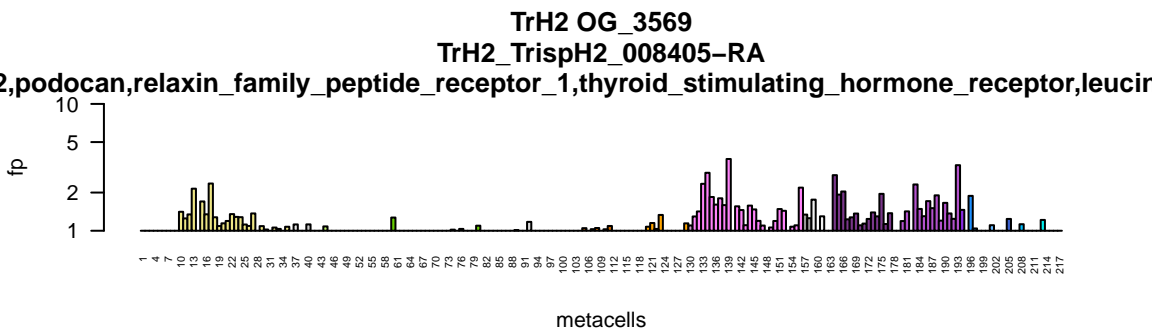
Hhon OG_9913
Hhon_g00250.t1

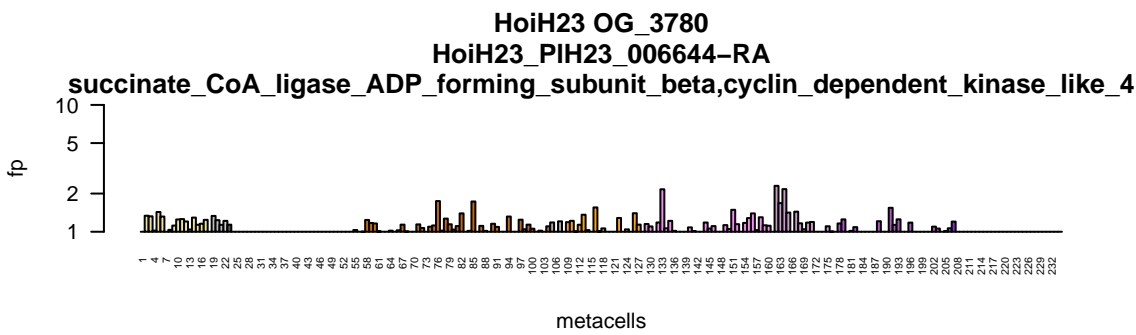
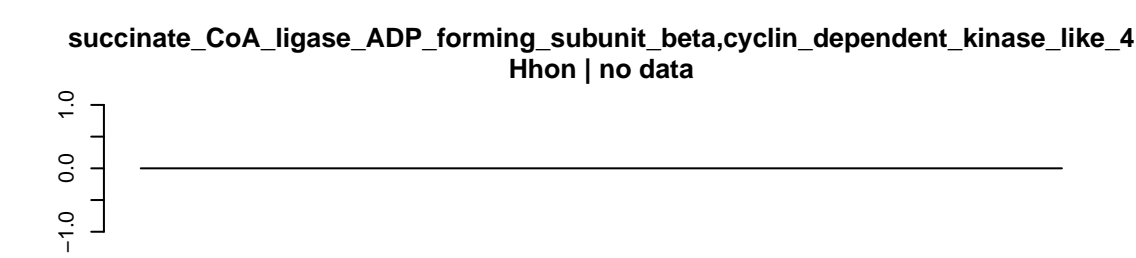
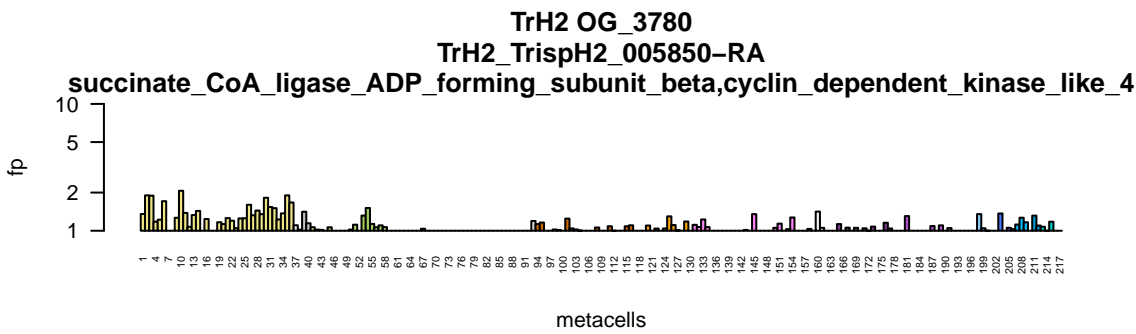
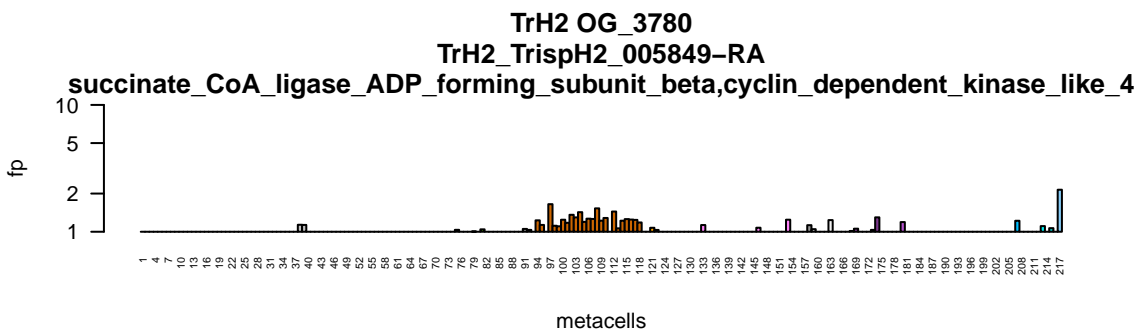
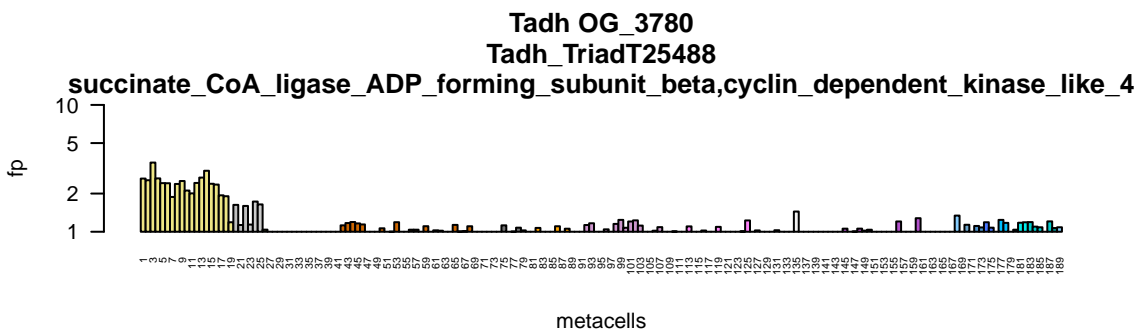
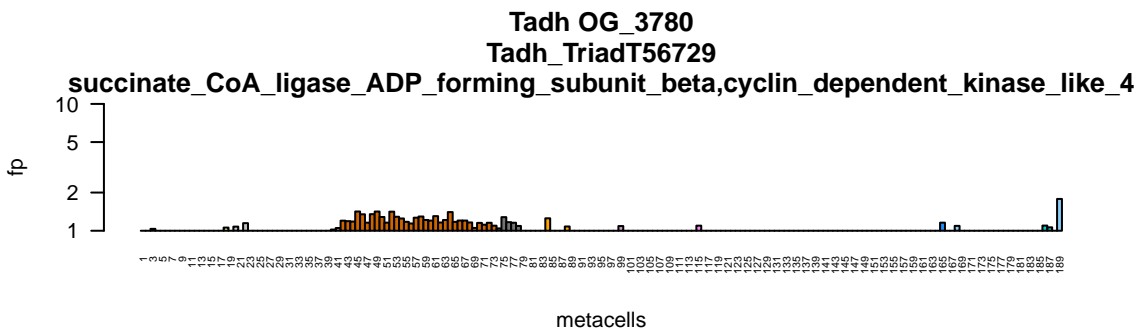


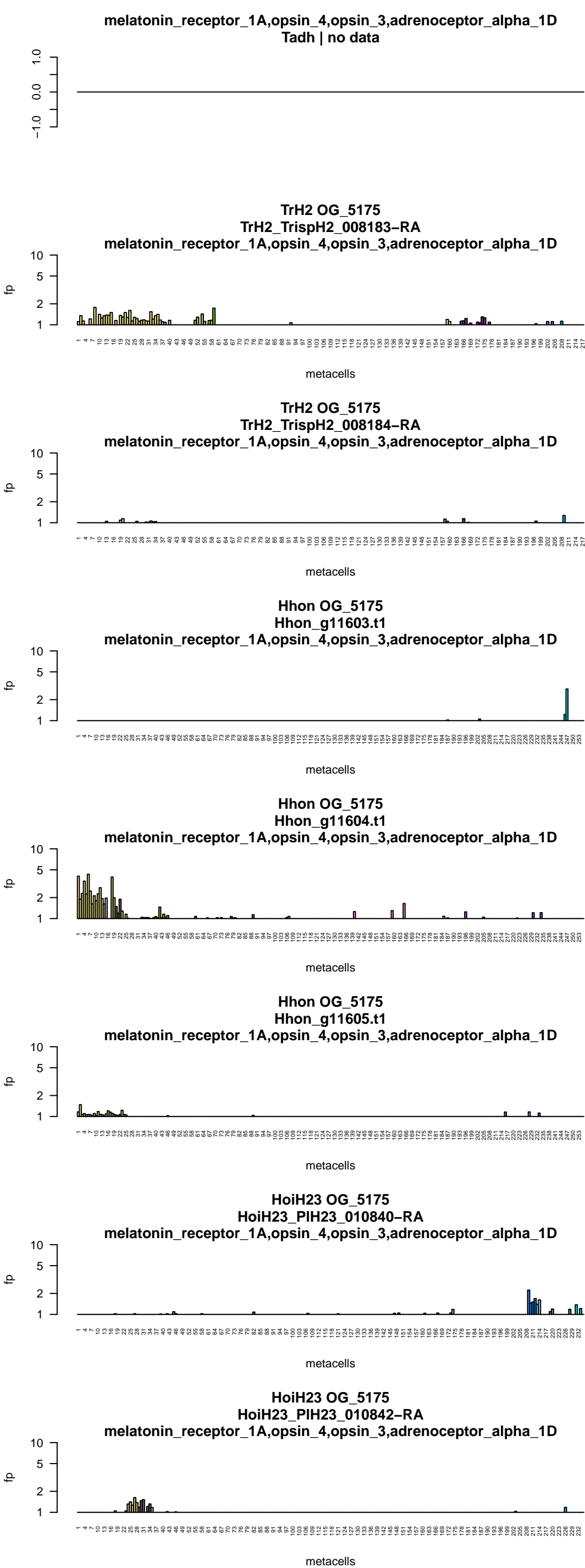
HoiH23 OG_9913
HoiH23_PIH23_006124-RA

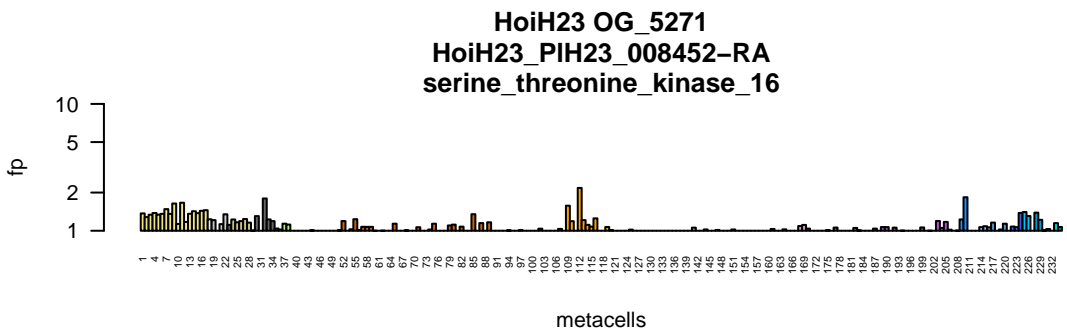
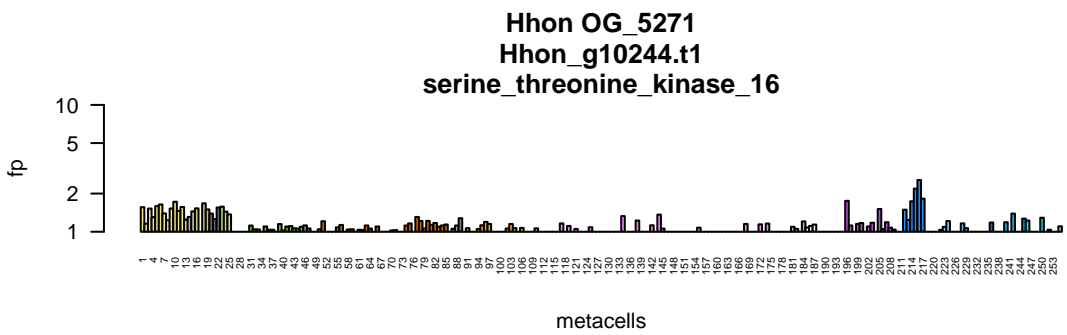
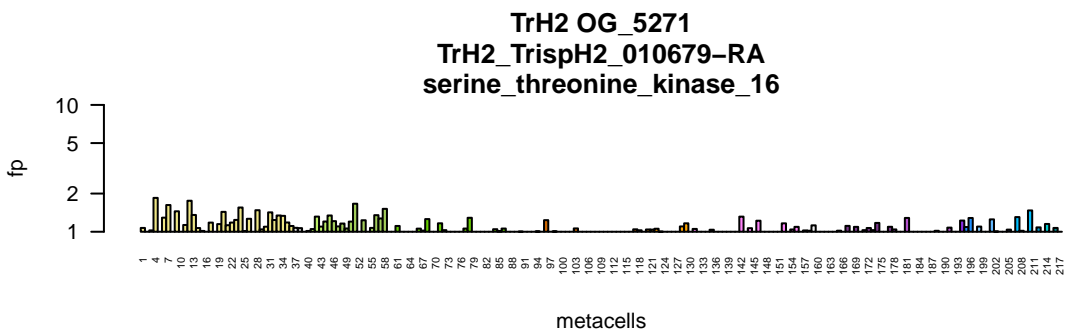
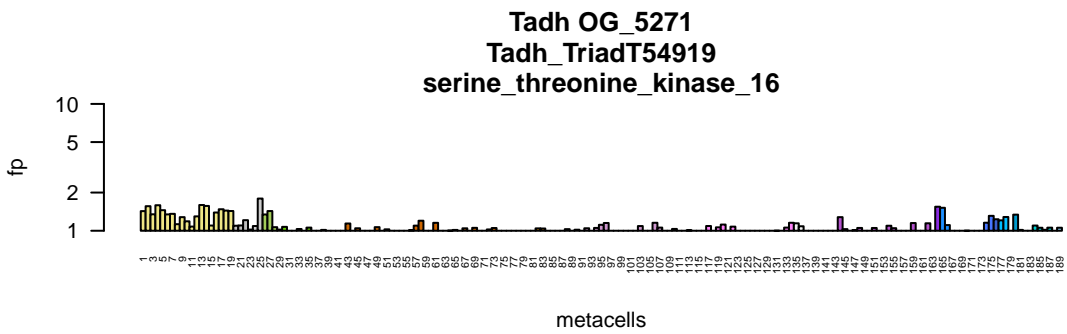


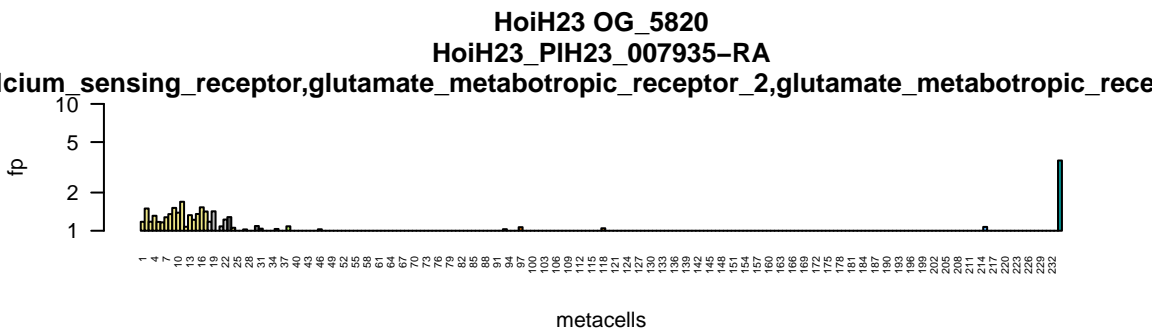
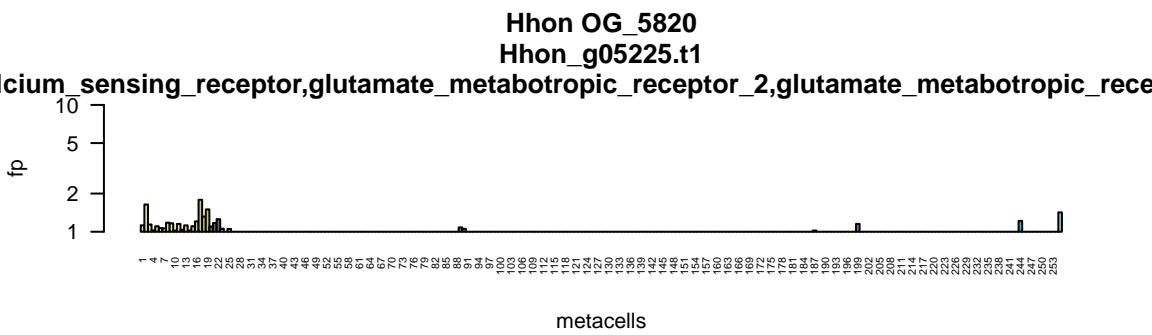
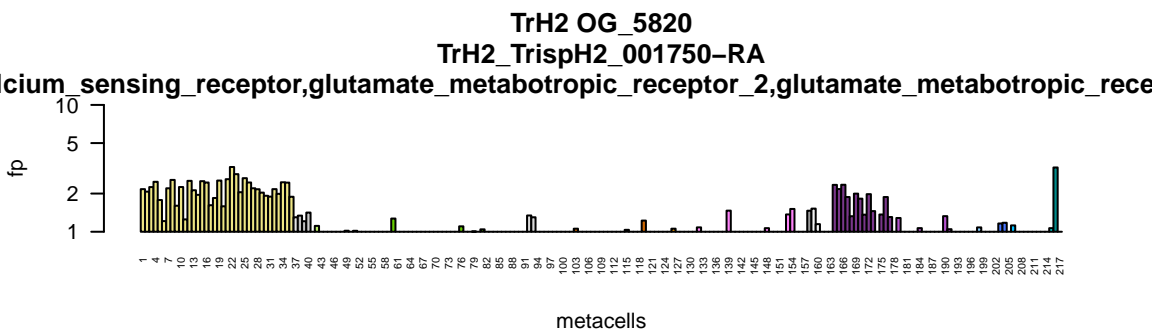
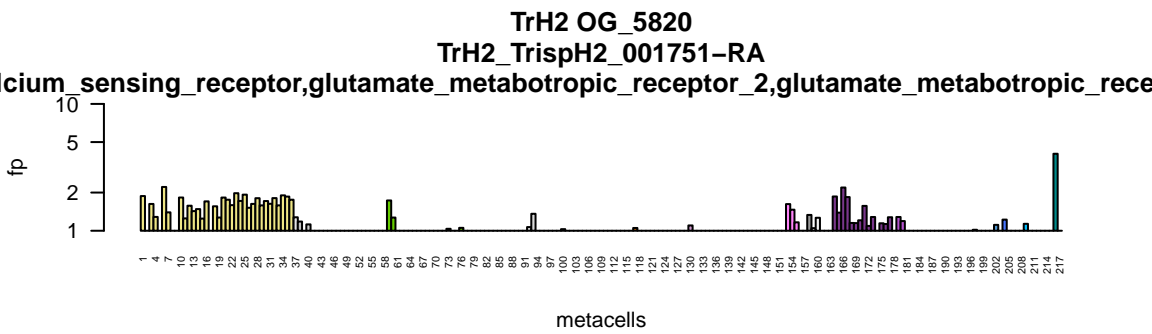
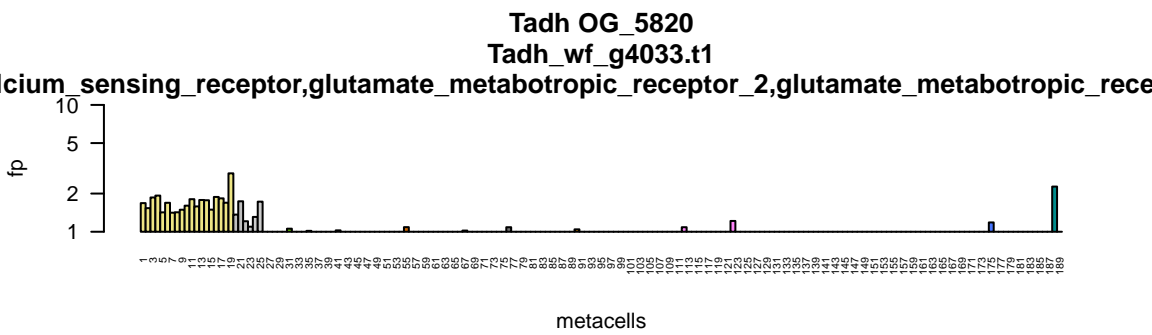
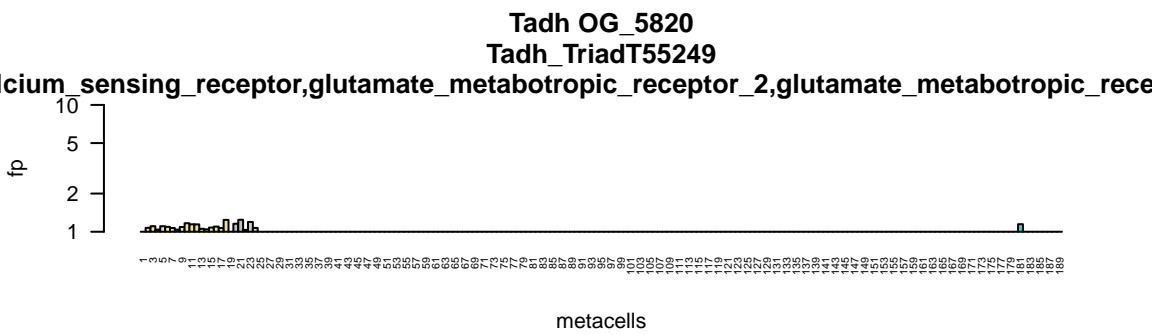
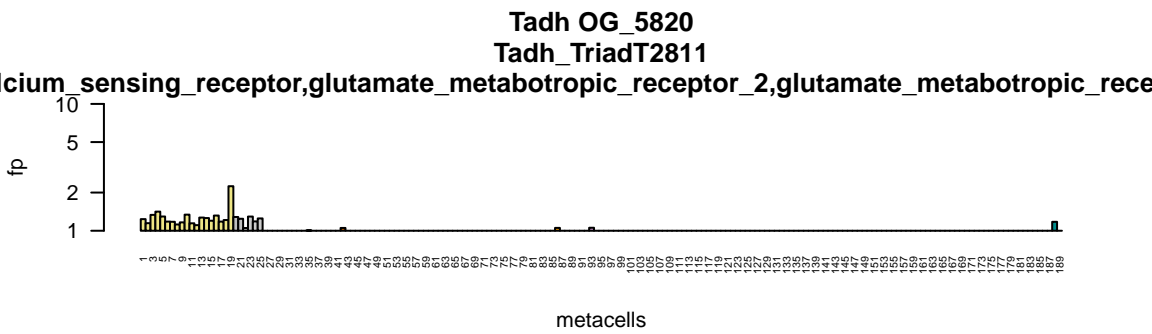


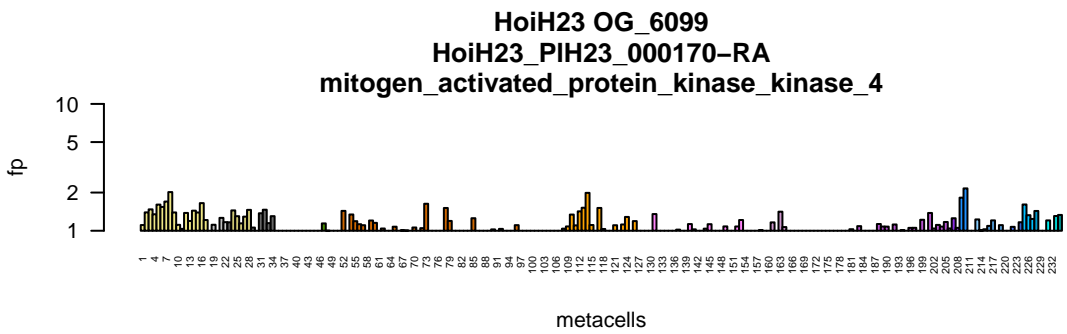
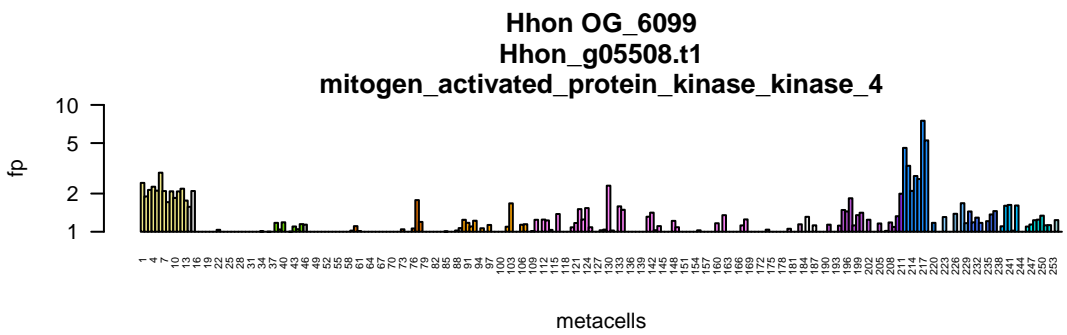
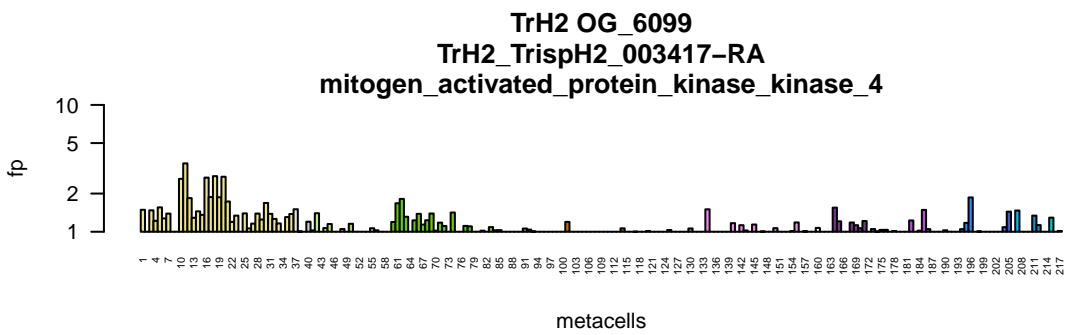
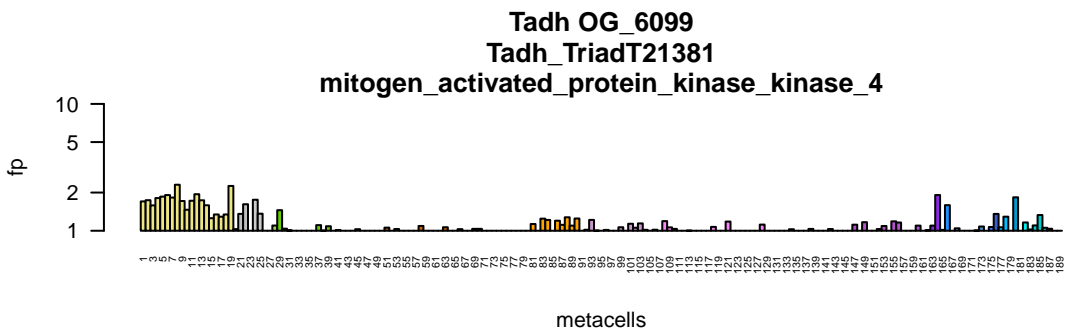


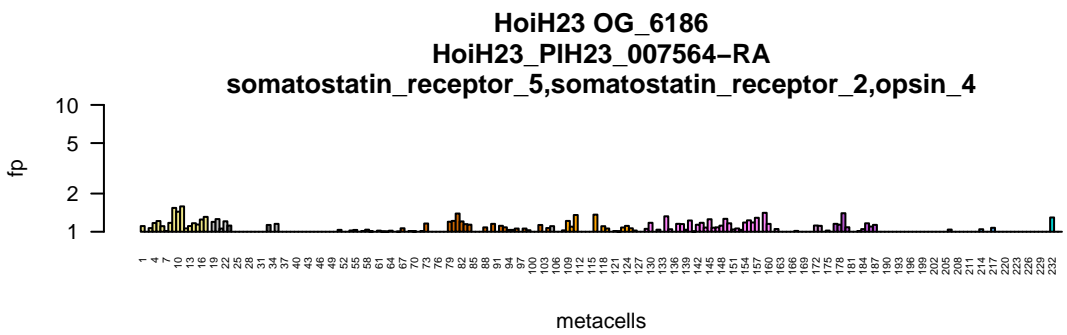
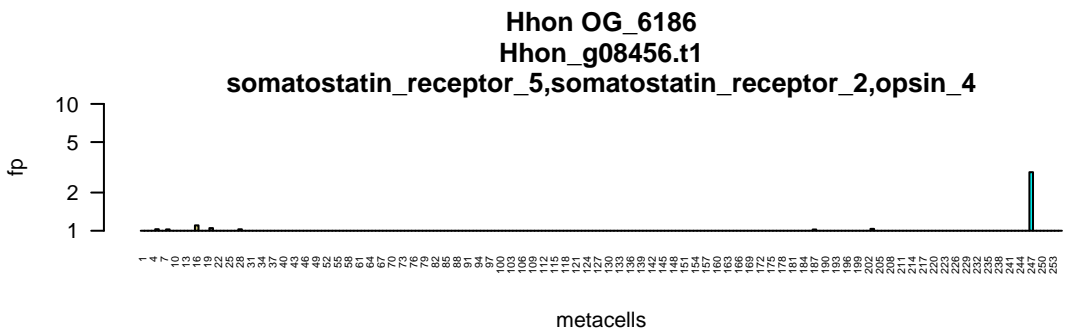
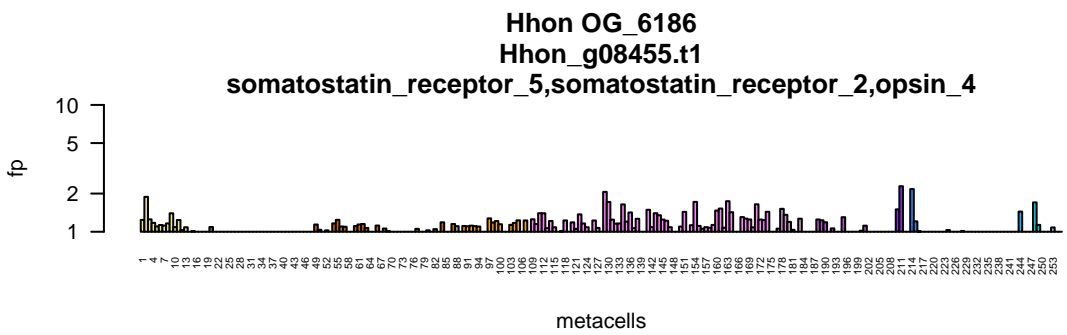
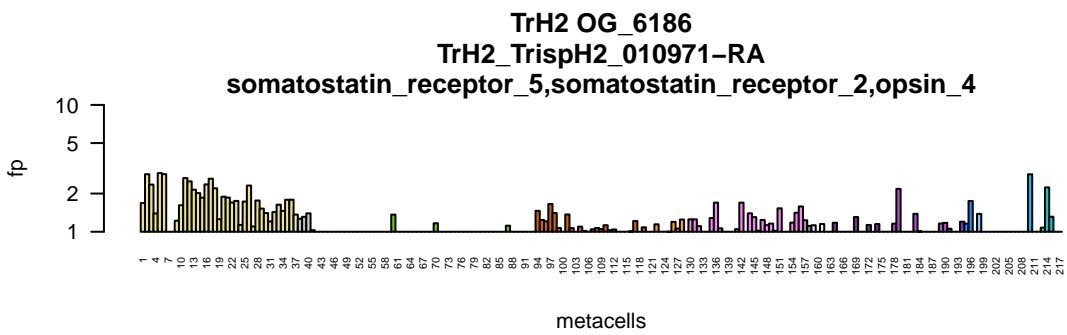
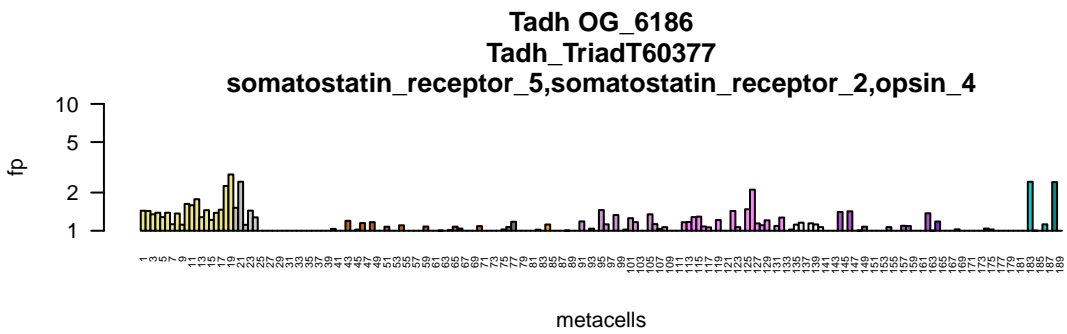


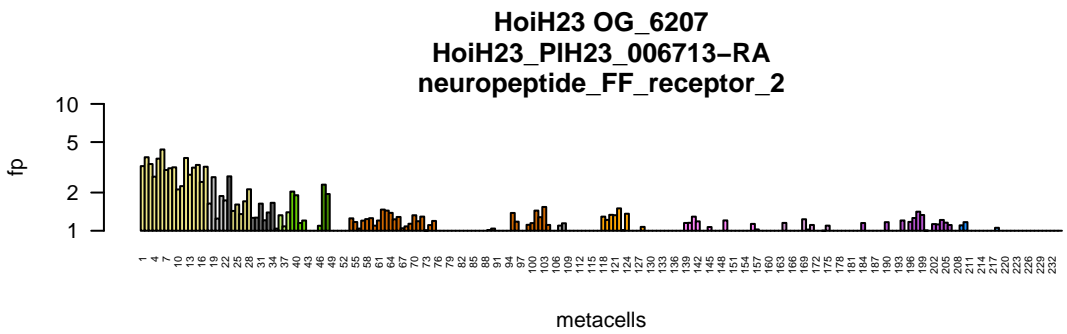
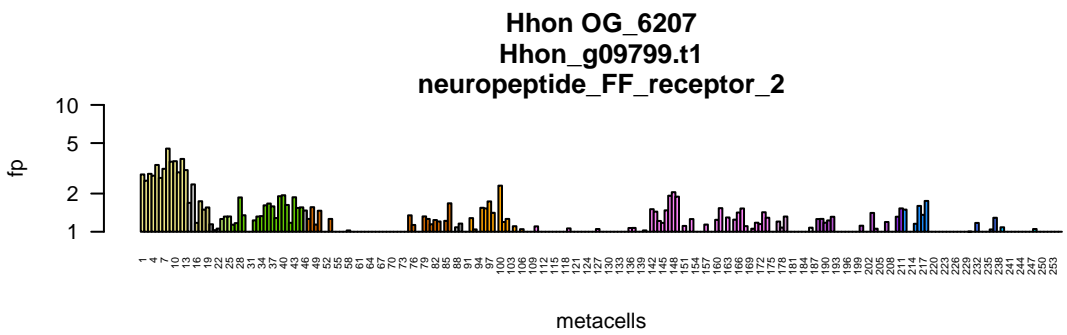
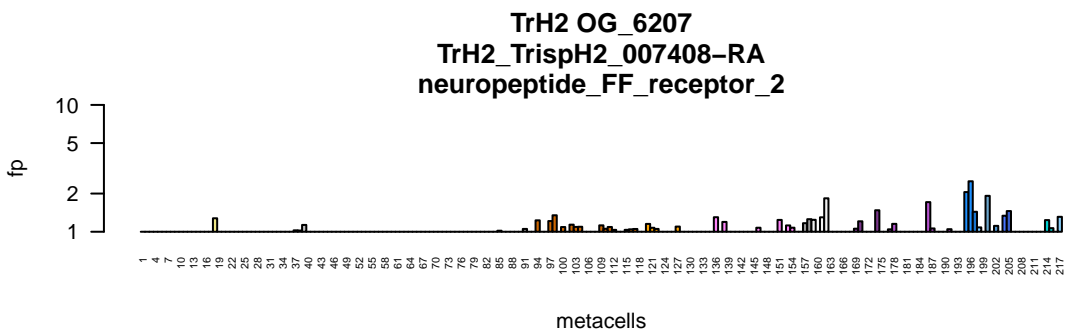
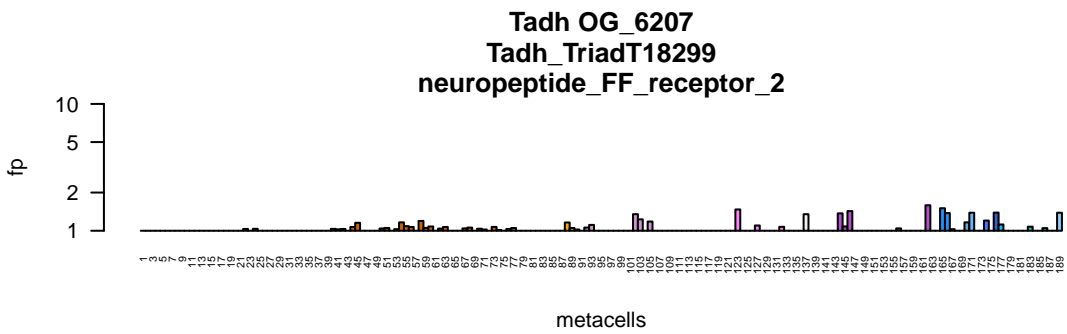


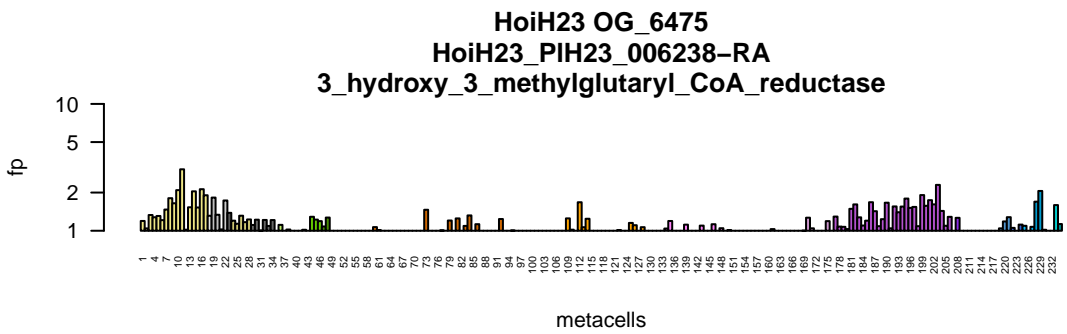
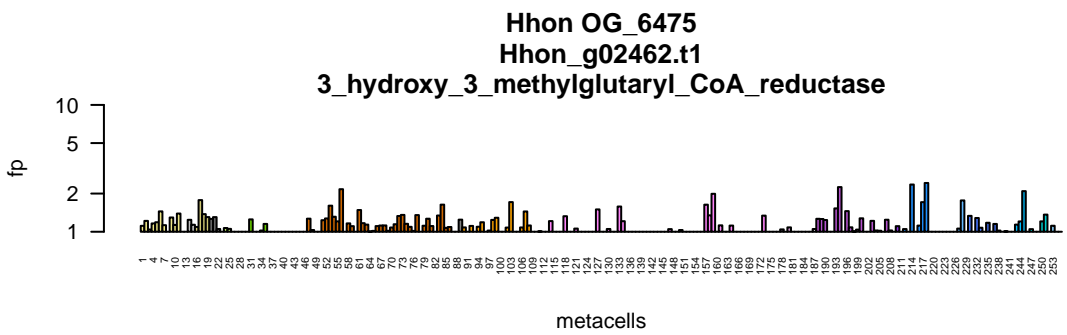
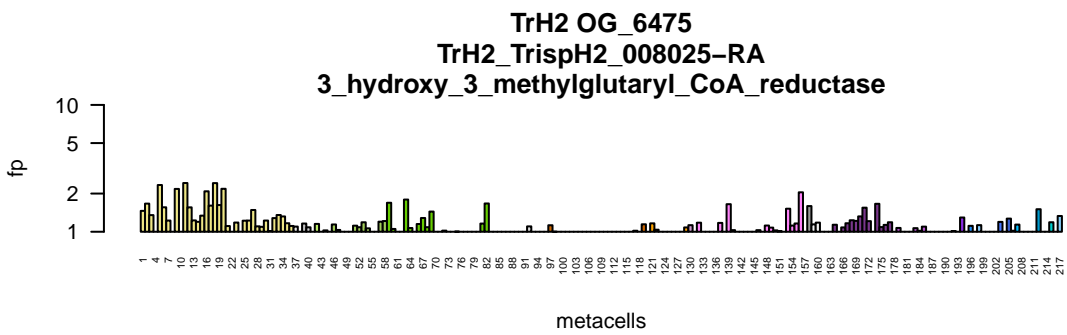
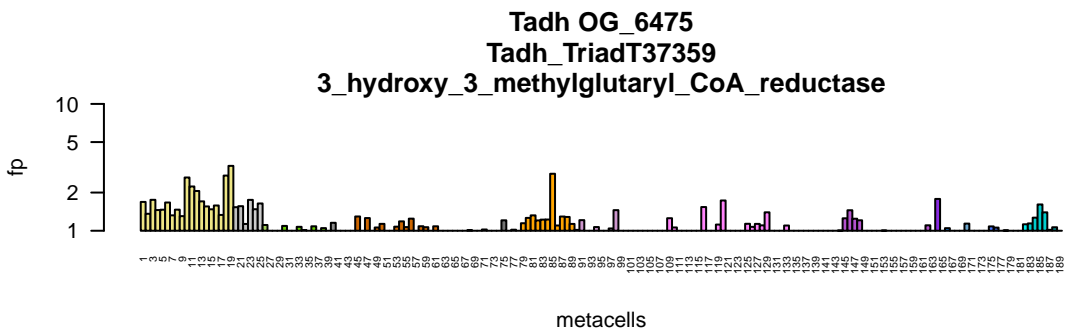


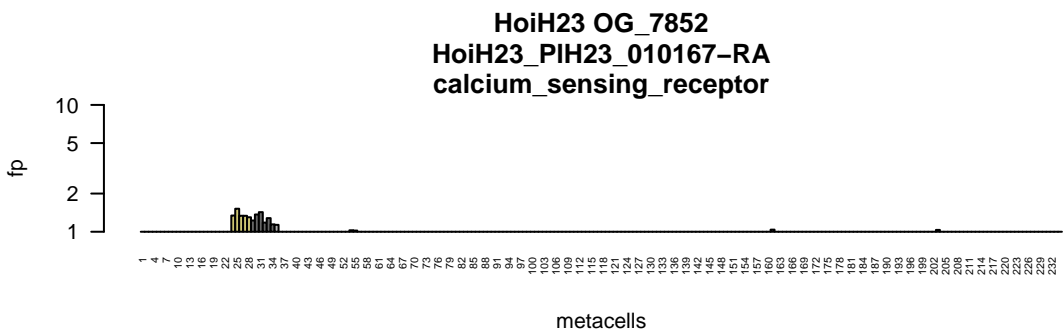
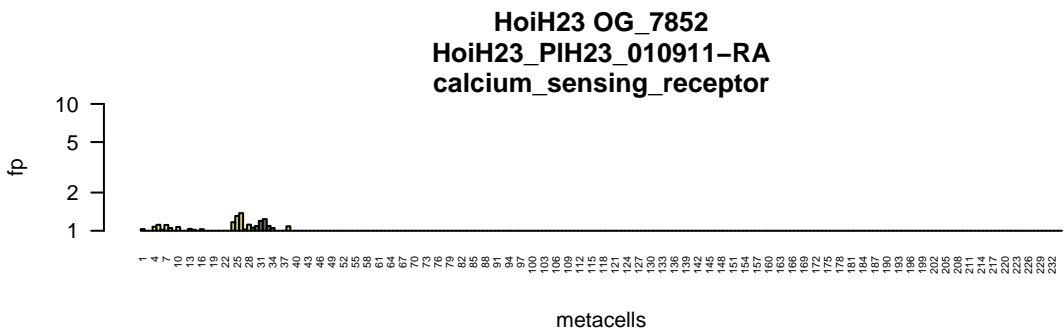
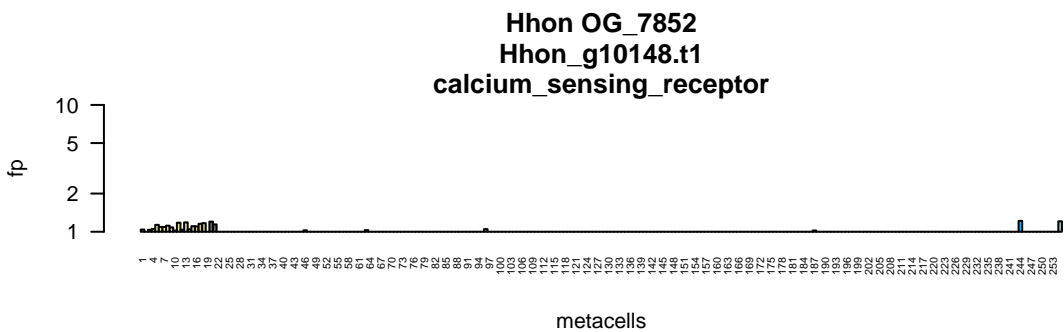
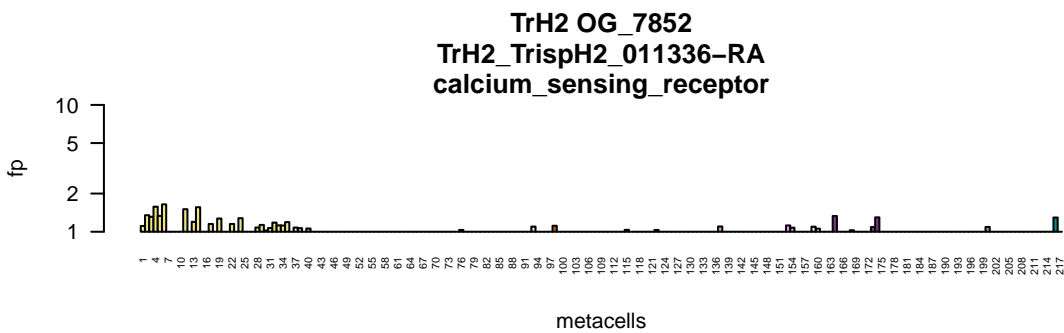
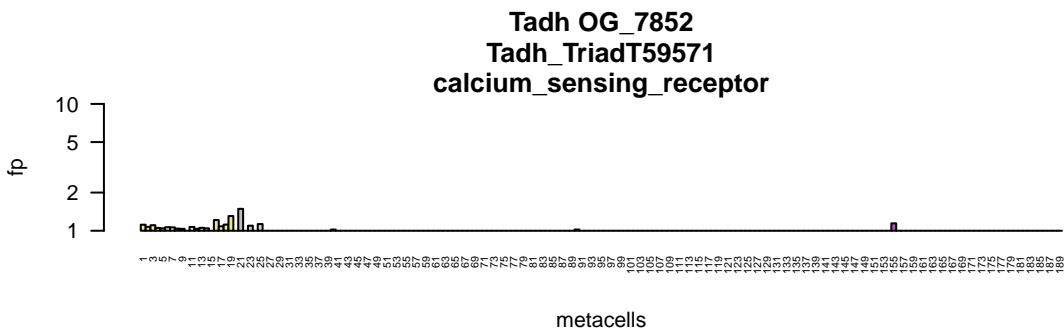


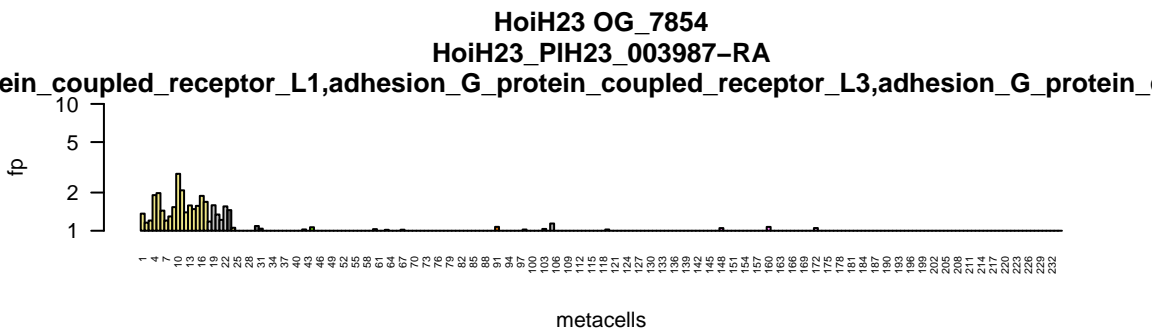
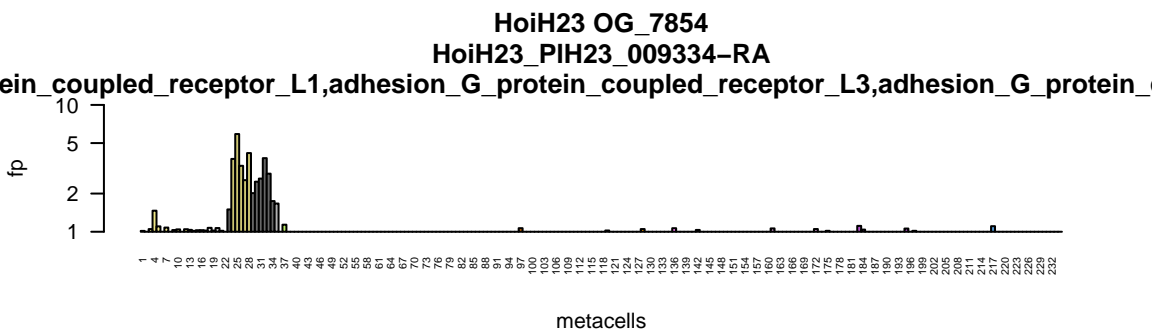
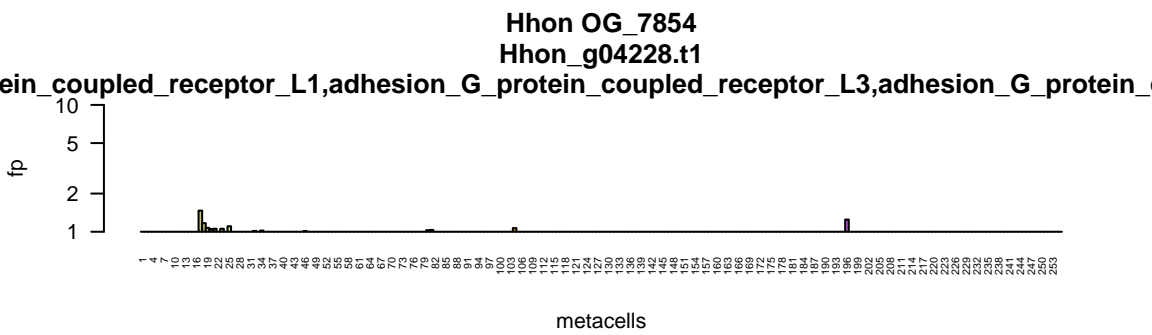
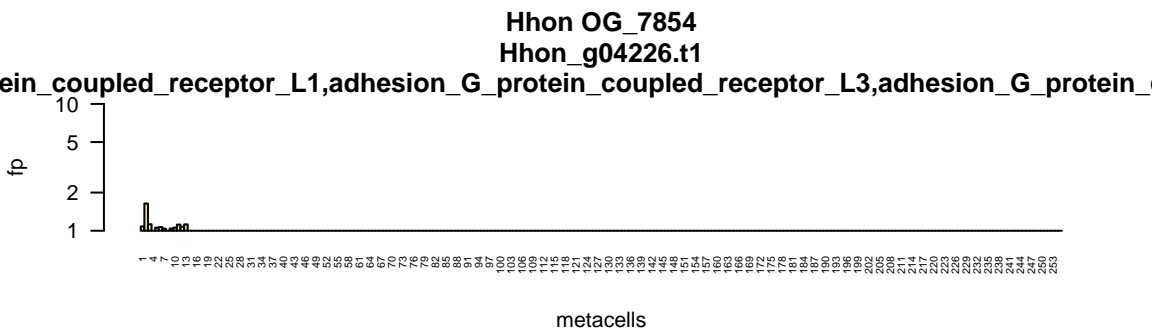
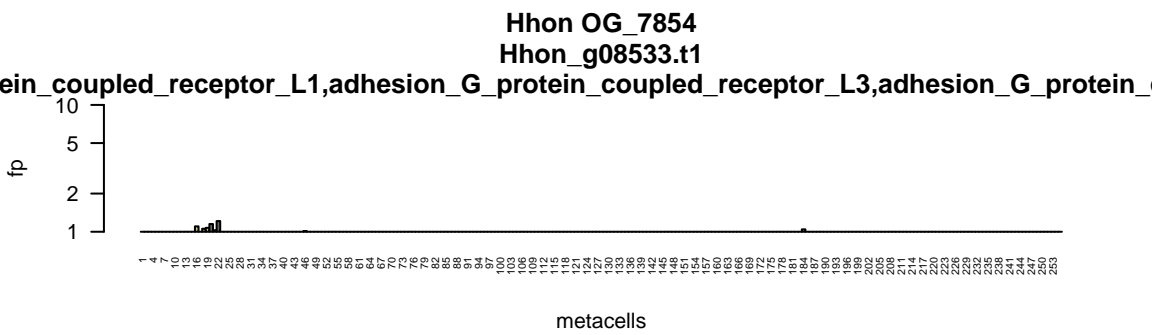
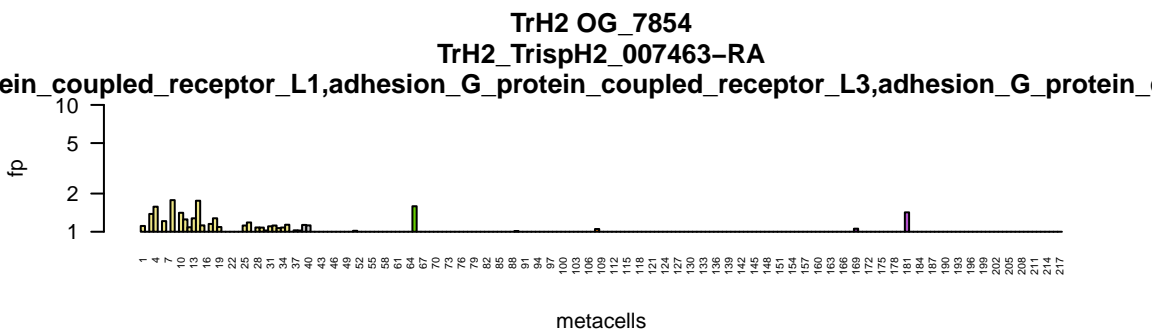
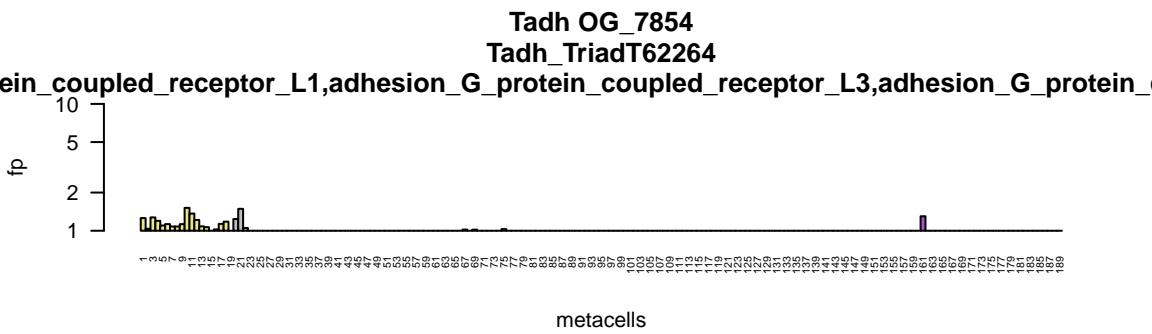








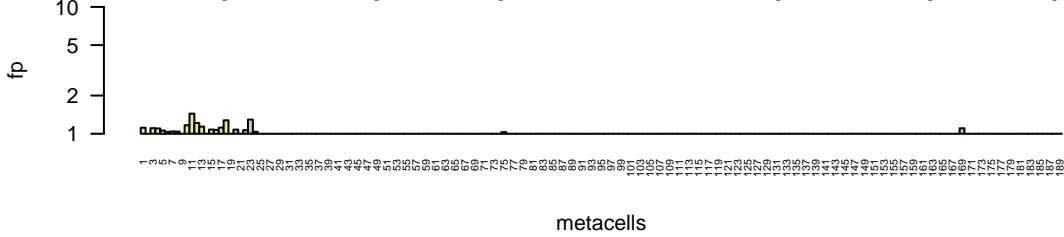




Tadh OG_8671

Tadh_TriadT57692

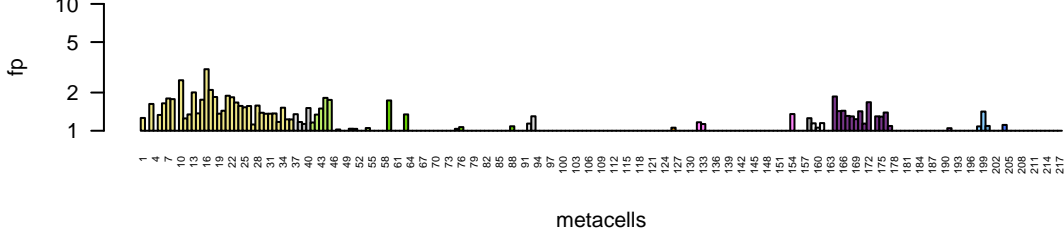
adhesion_G_protein_coupled_receptor_G2,adhesion_G_protein_coupled_receptor_L4



TrH2 OG_8671

TrH2_TrispH2_004680-RA

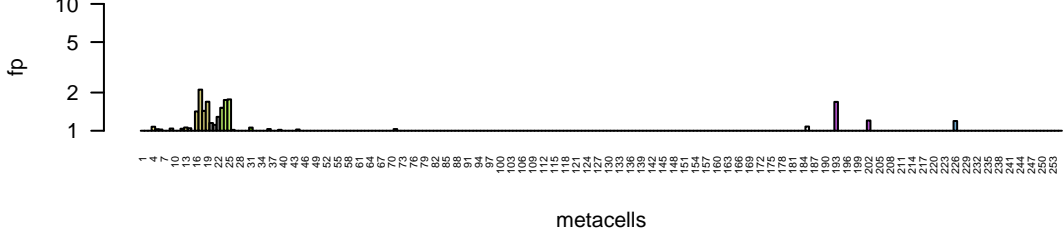
adhesion_G_protein_coupled_receptor_G2,adhesion_G_protein_coupled_receptor_L4



Hhon OG_8671

Hhon_g01778.t1

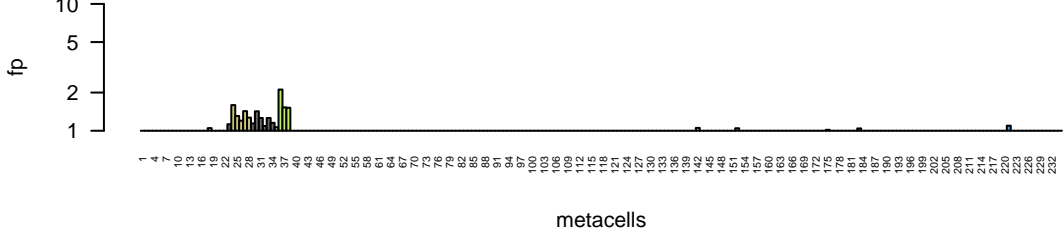
adhesion_G_protein_coupled_receptor_G2,adhesion_G_protein_coupled_receptor_L4



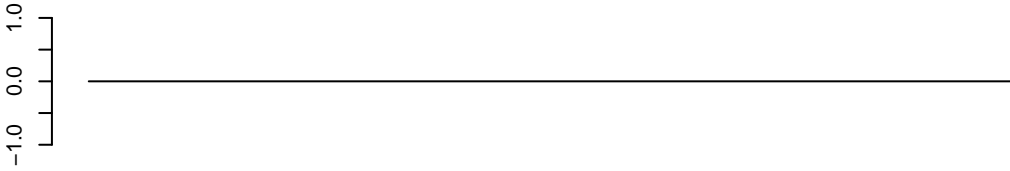
HoiH23 OG_8671

HoiH23_PIH23_004834-RA

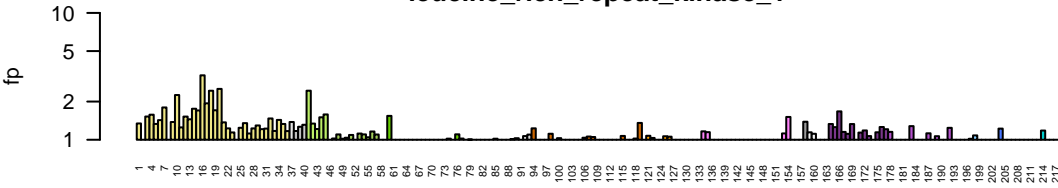
adhesion_G_protein_coupled_receptor_G2,adhesion_G_protein_coupled_receptor_L4



leucine_rich_repeat_kinase_1
Tadh | no data

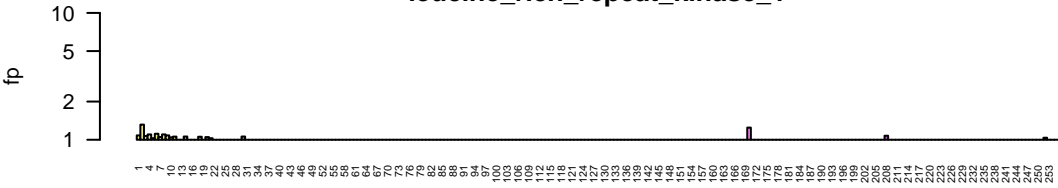


TrH2 OG_10615
TrH2_TrispH2_004289-RA
leucine_rich_repeat_kinase_1



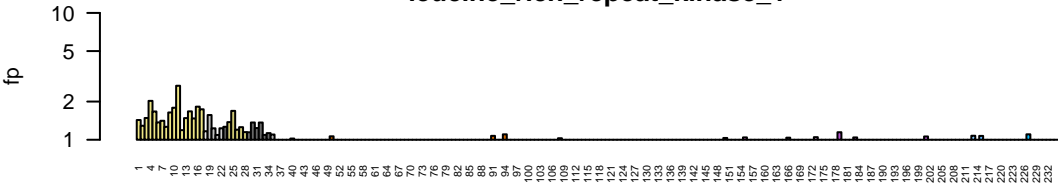
metacells

Hhon OG_10615
Hhon_g06459.t1
leucine_rich_repeat_kinase_1

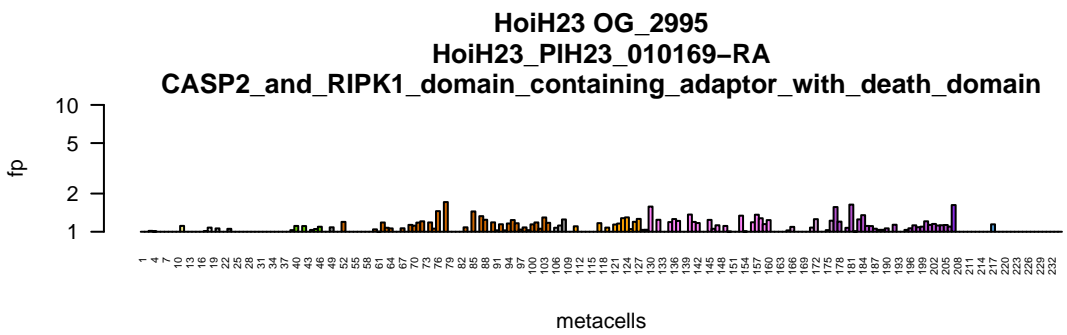
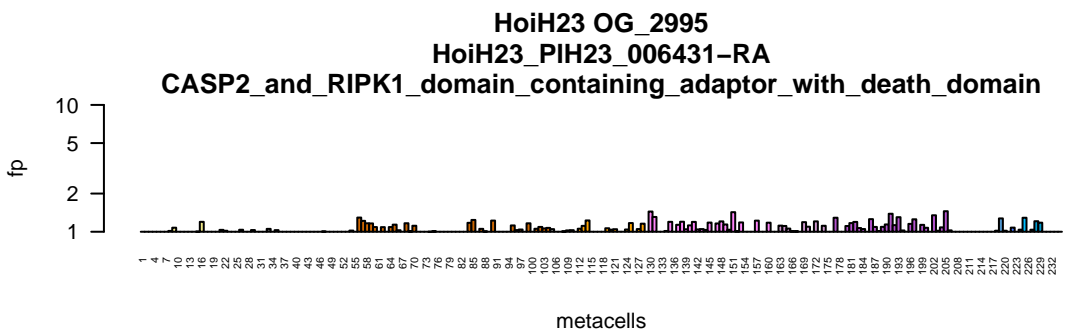
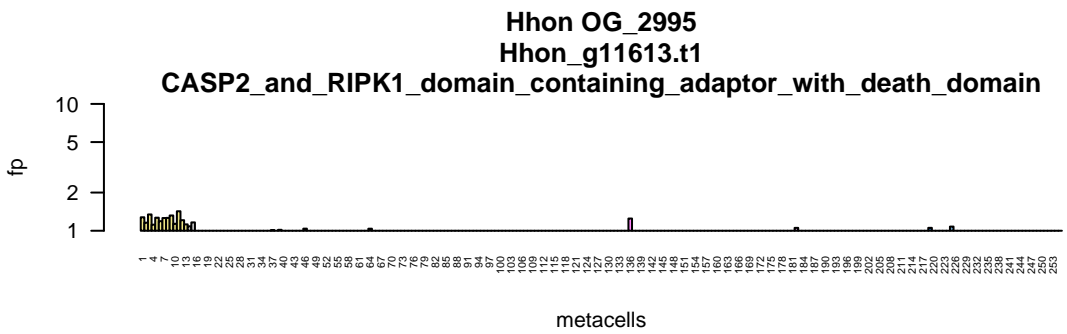
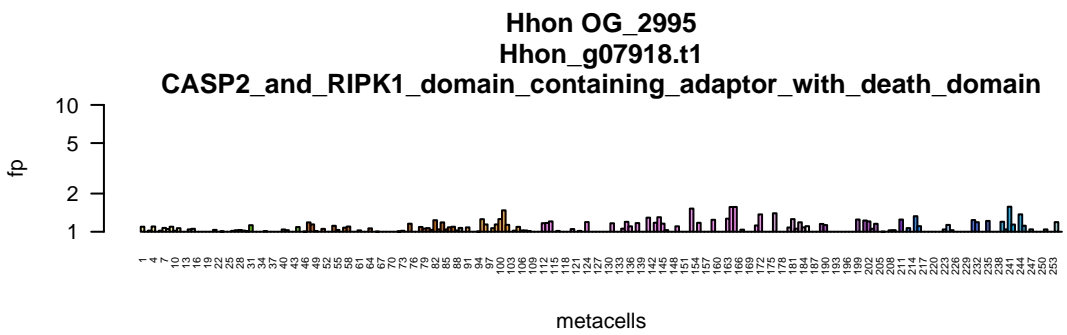
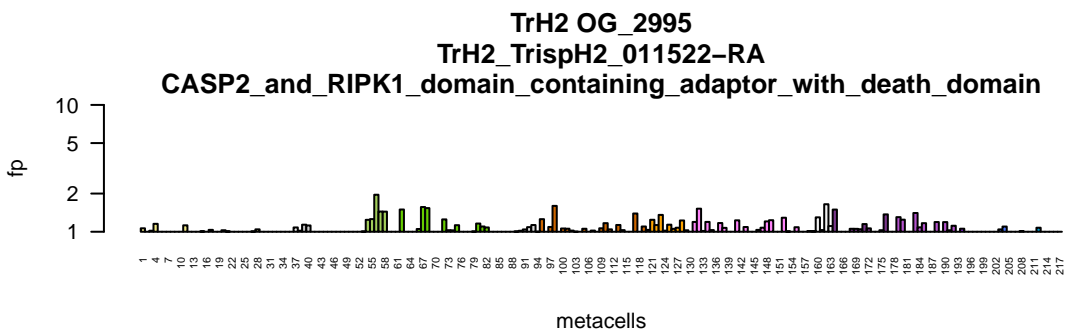
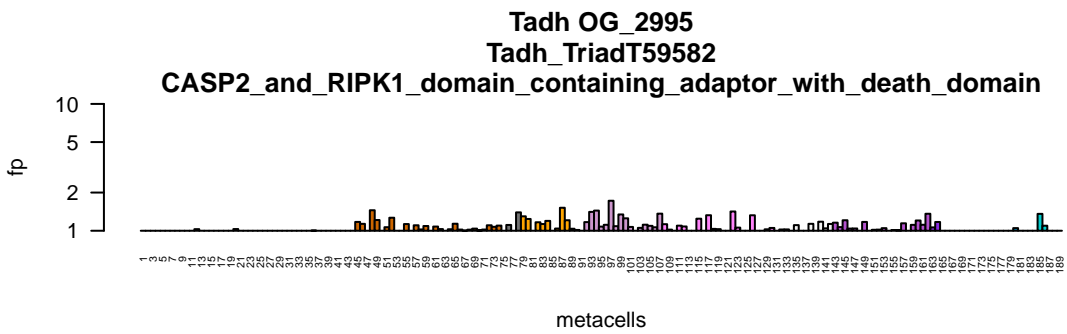


metacells

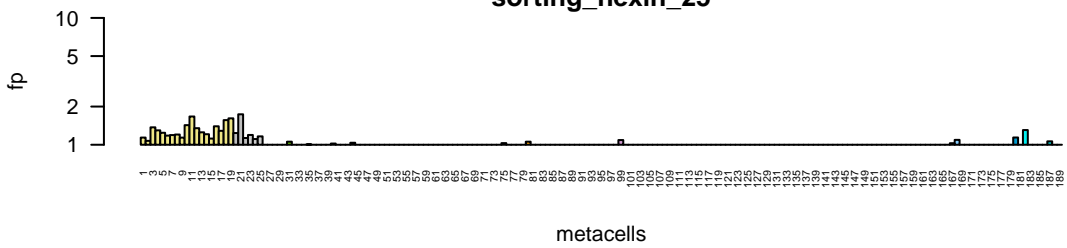
HoiH23 OG_10615
HoiH23_PIH23_003918-RA
leucine_rich_repeat_kinase_1



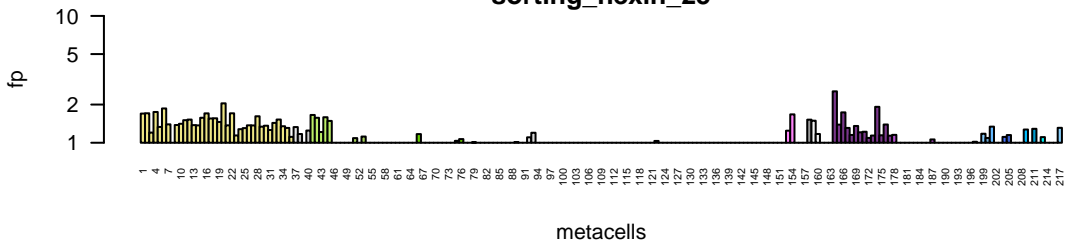
metacells



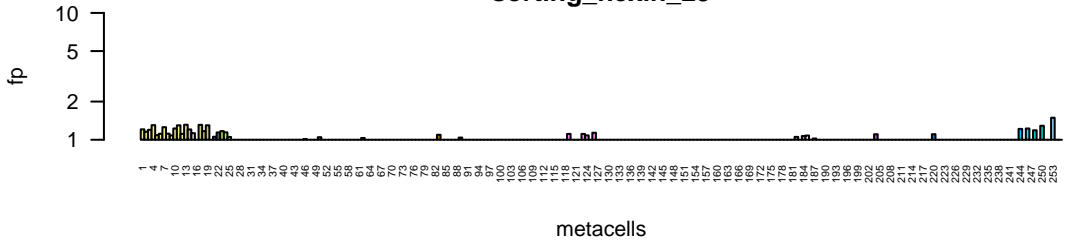
Tadh OG_4176
Tadh_TriadT56593
sorting_nexin_25



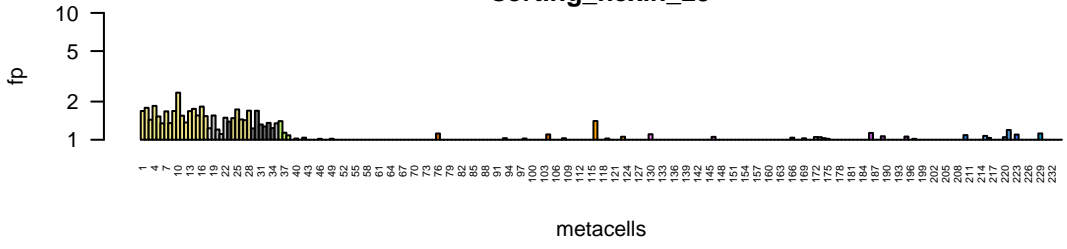
TrH2 OG_4176
TrH2_TrispH2_005862-RA
sorting_nexin_25

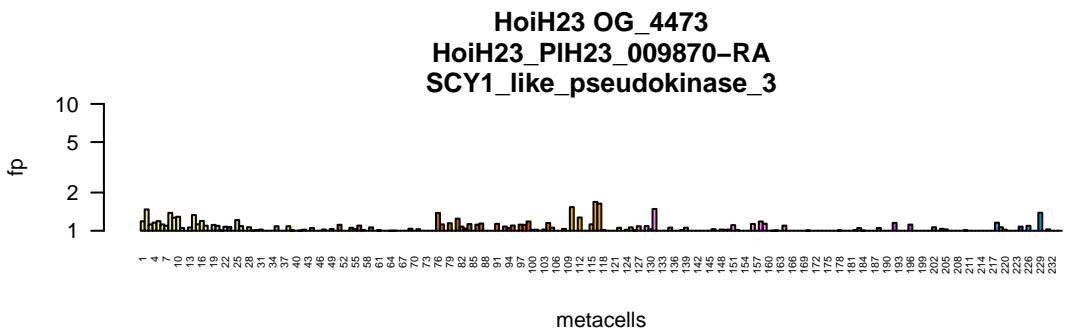
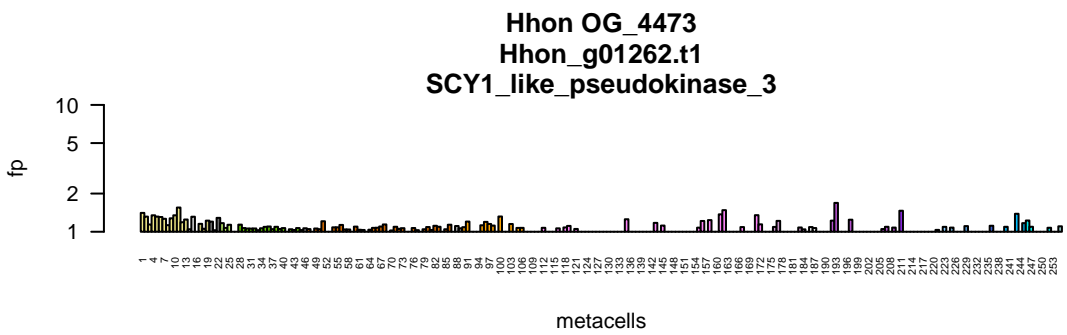
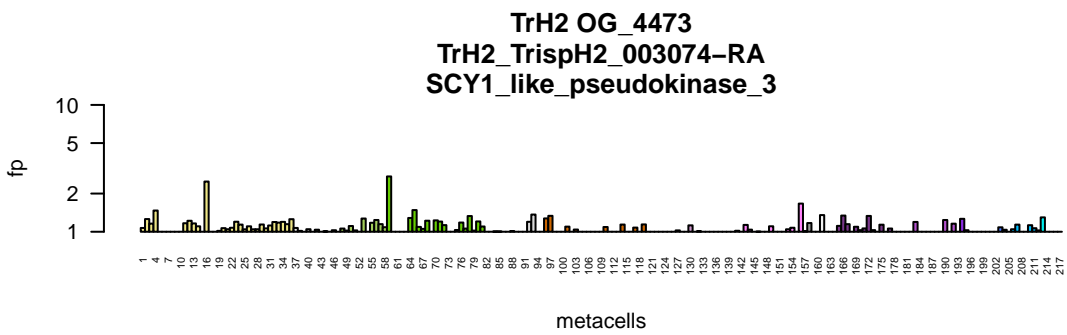
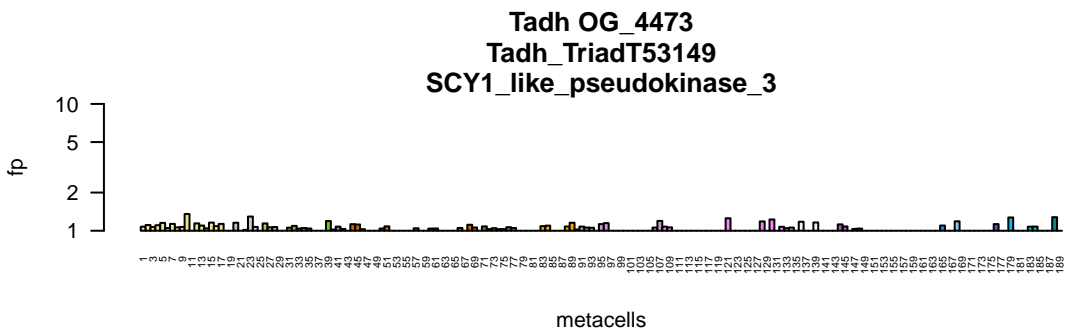


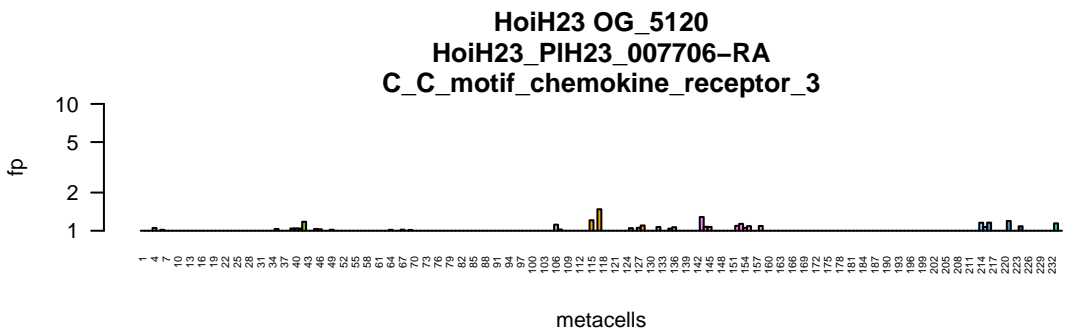
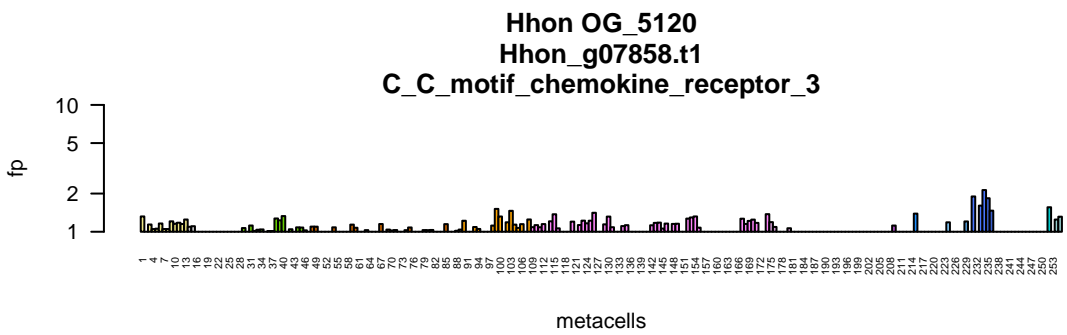
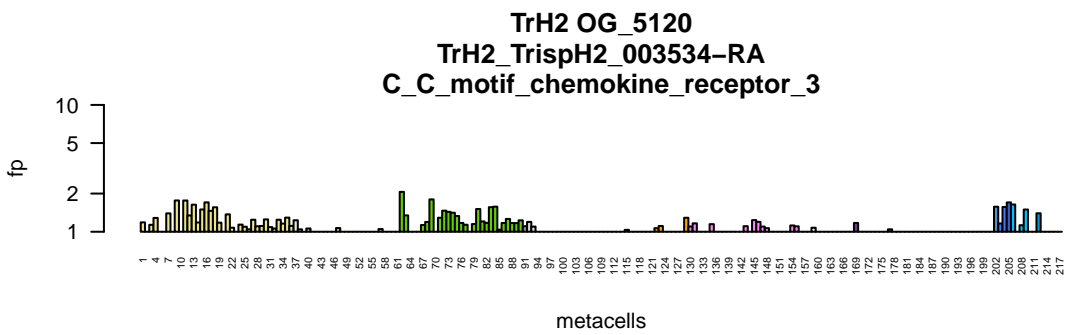
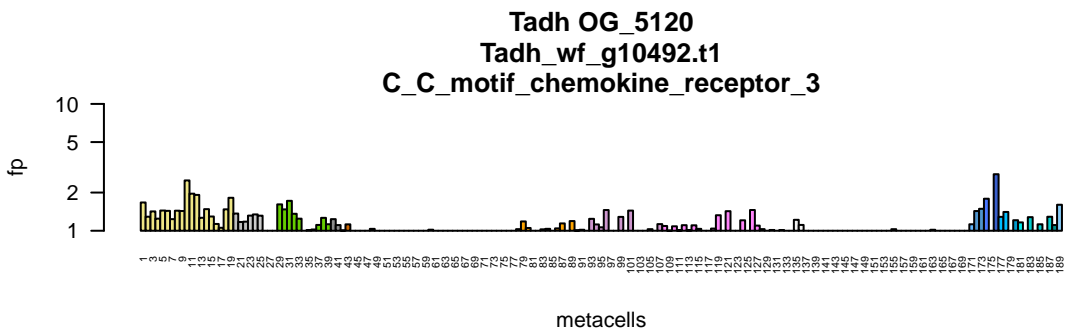
Hhon OG_4176
Hhon_g07573.t1
sorting_nexin_25

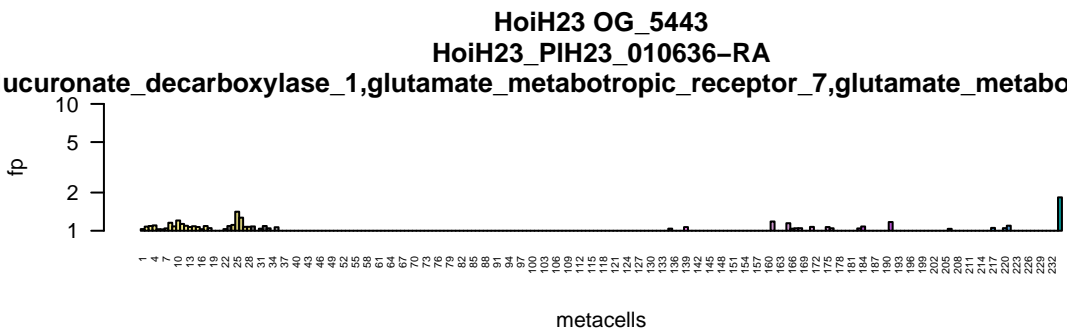
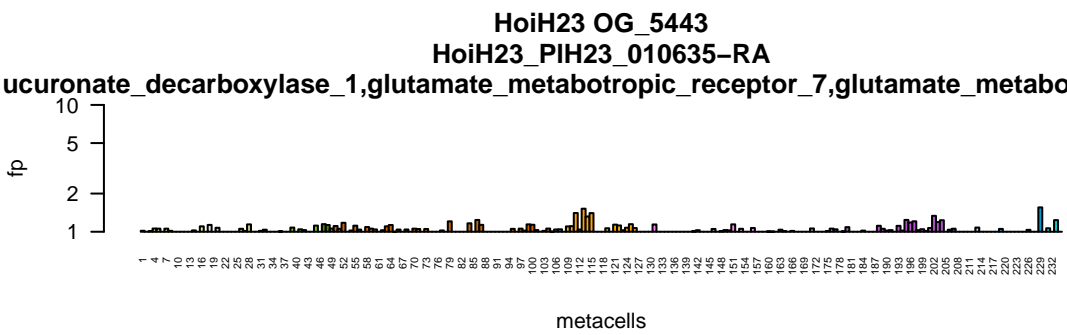
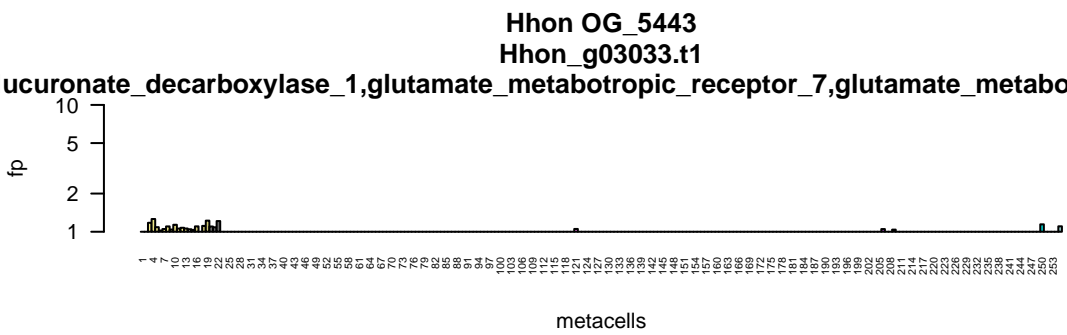
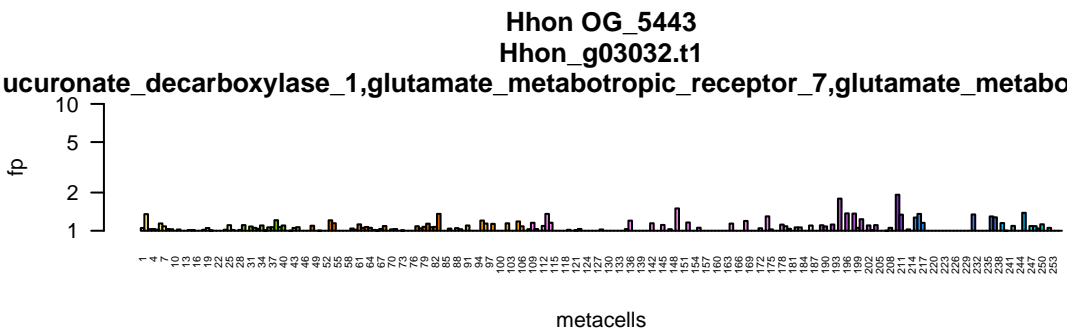
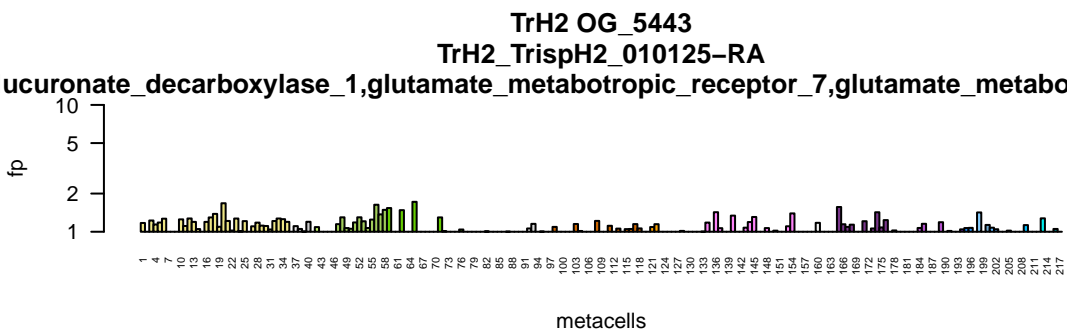
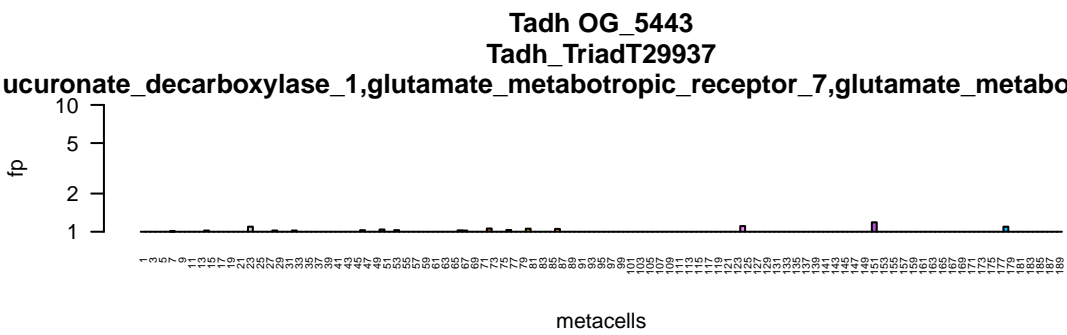
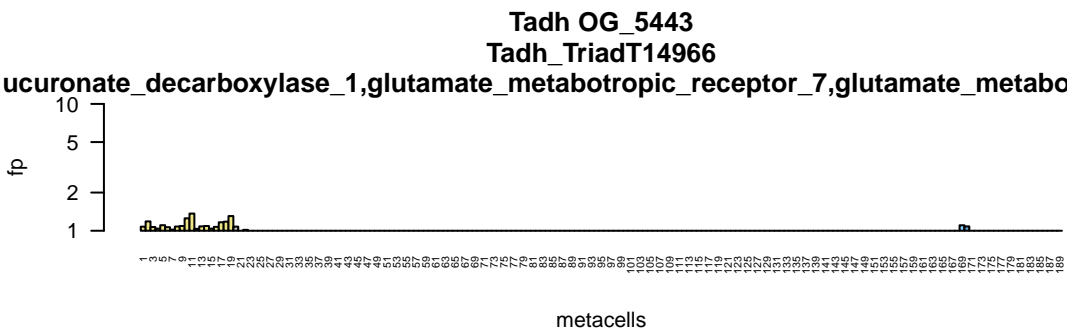


HoiH23 OG_4176
HoiH23_PIH23_009291-RA
sorting_nexin_25









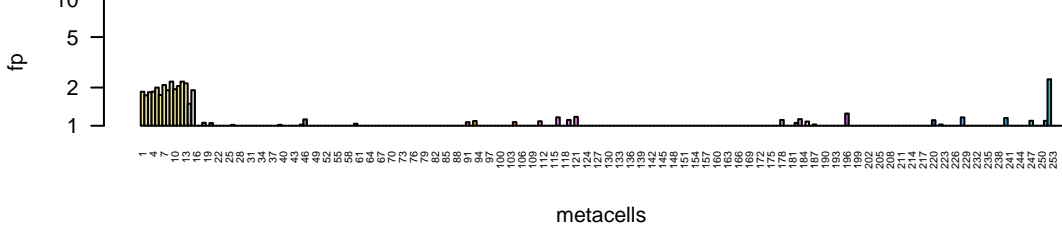
5_hydroxytryptamine_receptor_1D
Tadh | no data



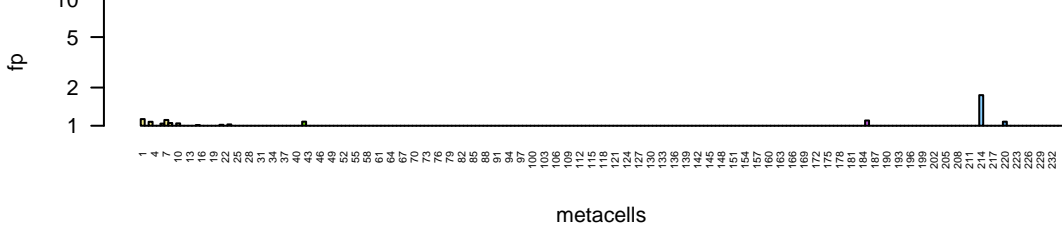
5_hydroxytryptamine_receptor_1D
TrH2 | no data



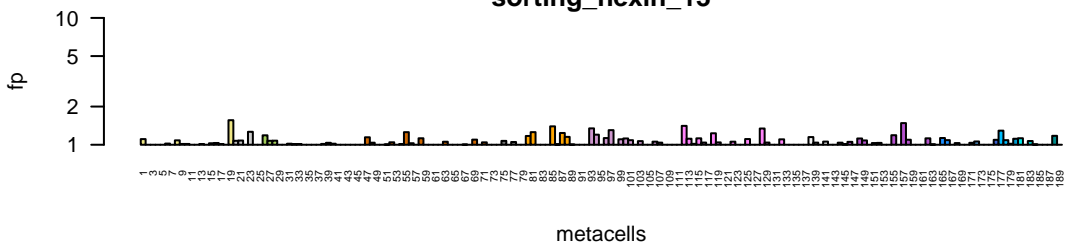
Hhon OG_5843
Hhon_g00398.t1
5_hydroxytryptamine_receptor_1D



HoiH23 OG_5843
HoiH23_PIH23_001773-RA
5_hydroxytryptamine_receptor_1D

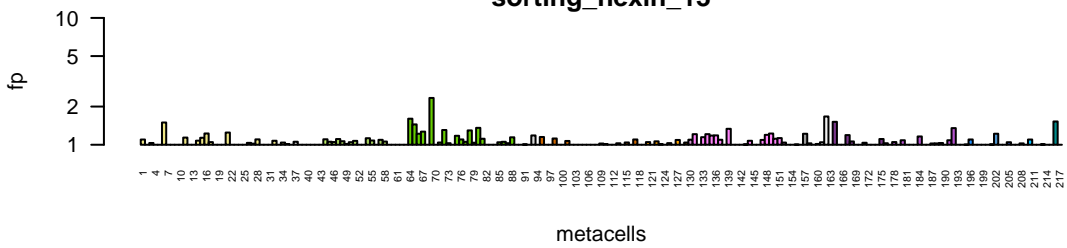


Tadh OG_5872
Tadh_TriadT53998
sorting_nexin_13



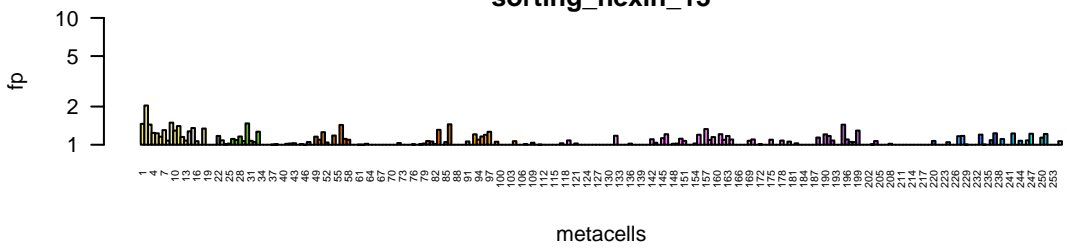
metacells

TrH2 OG_5872
TrH2_TrispH2_007781-RA
sorting_nexin_13



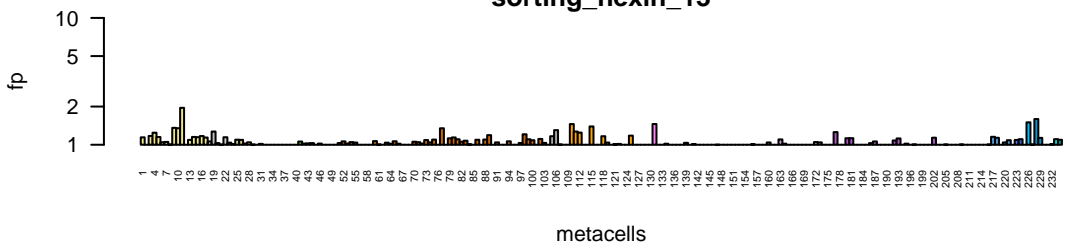
metacells

Hhon OG_5872
Hhon_g10880.t1
sorting_nexin_13

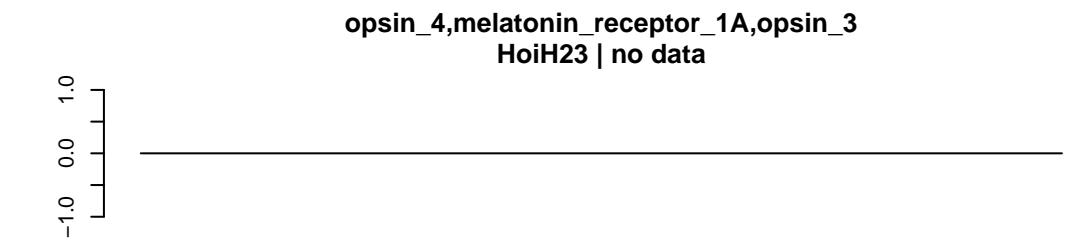
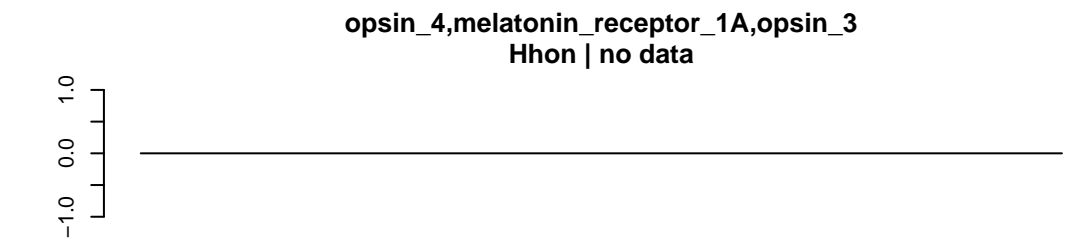
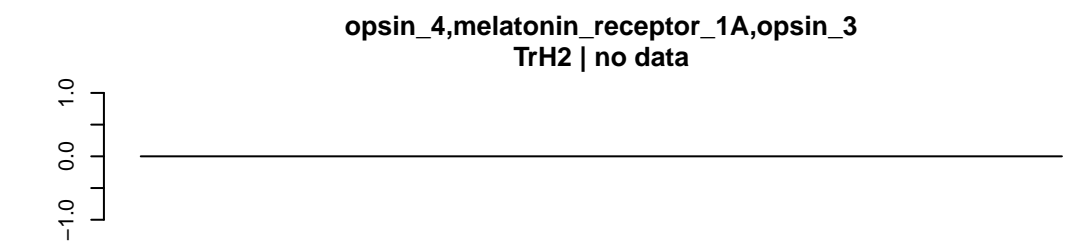


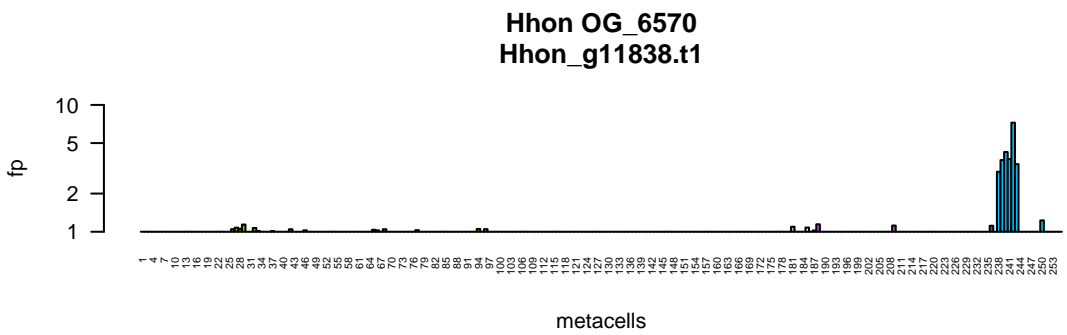
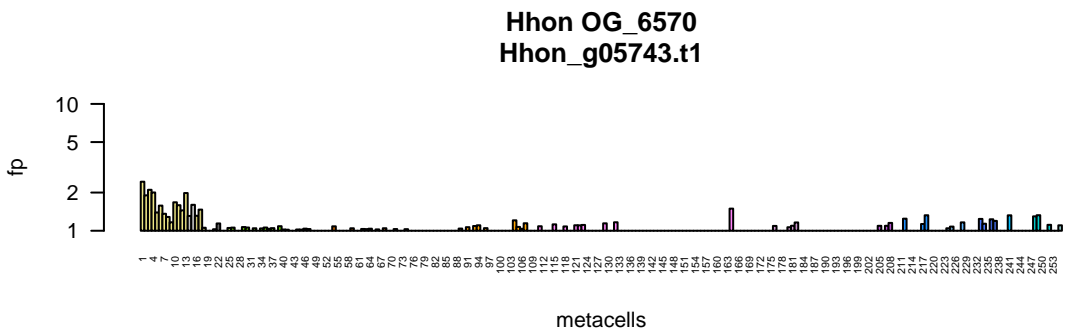
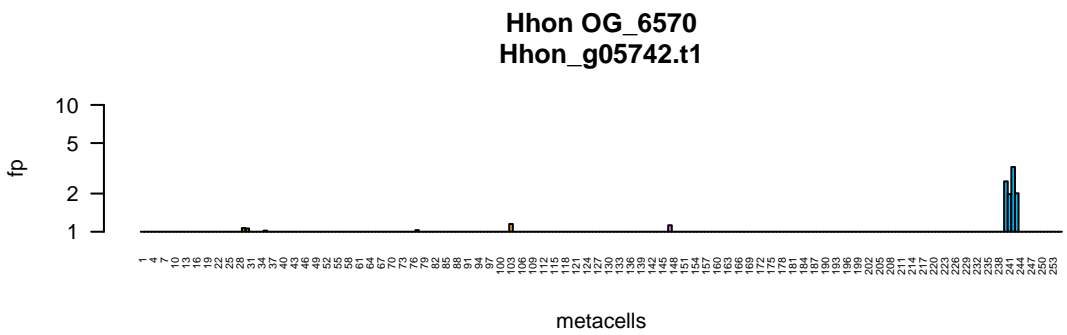
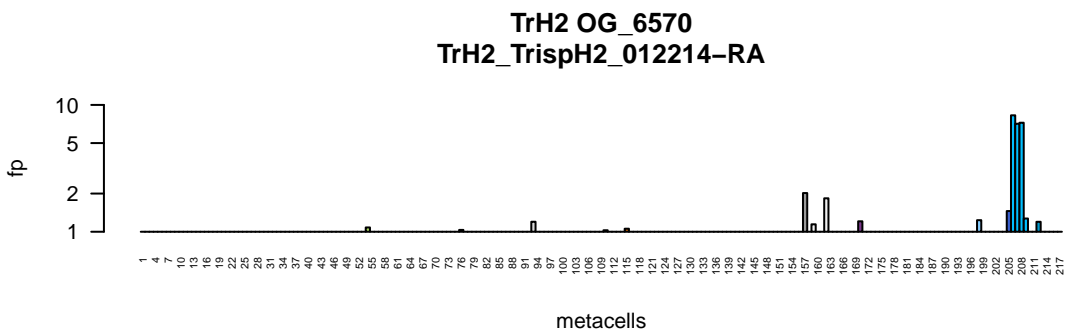
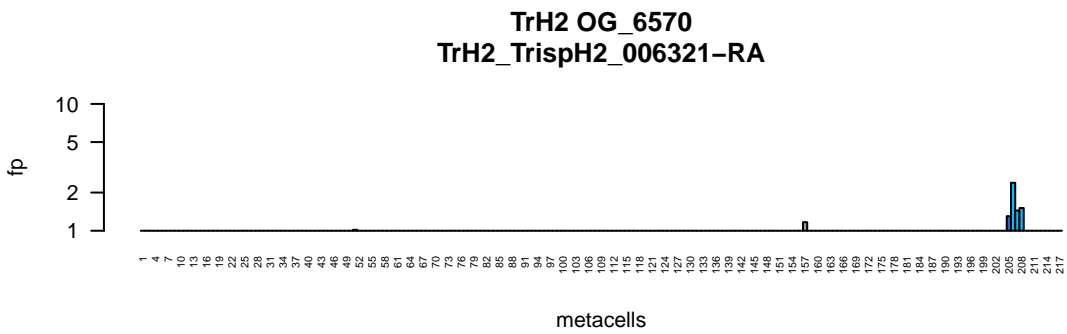
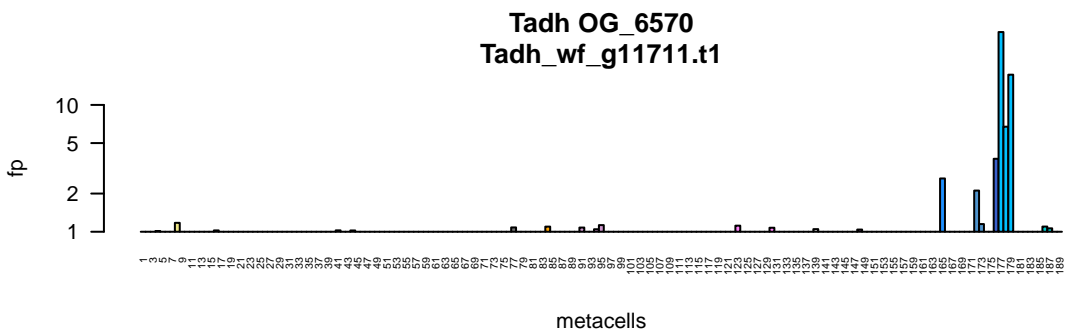
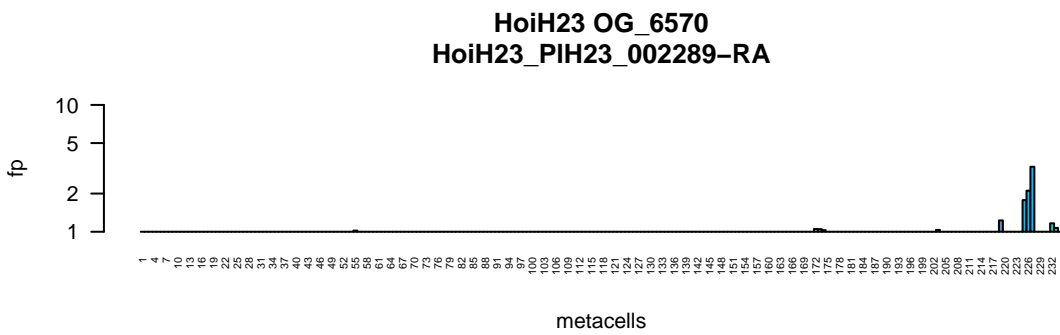
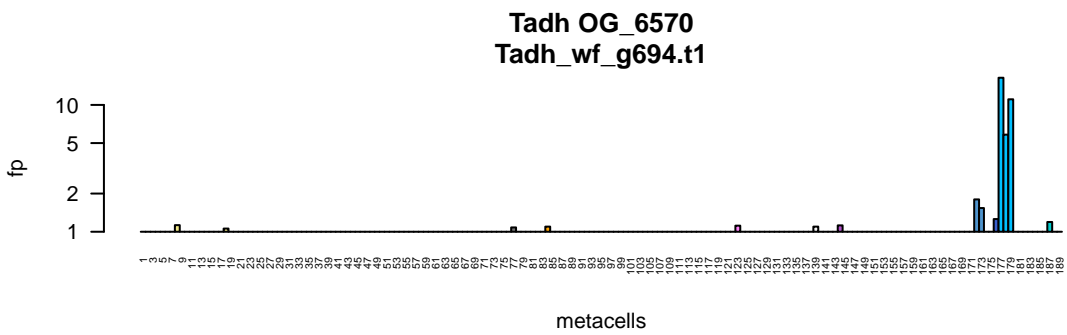
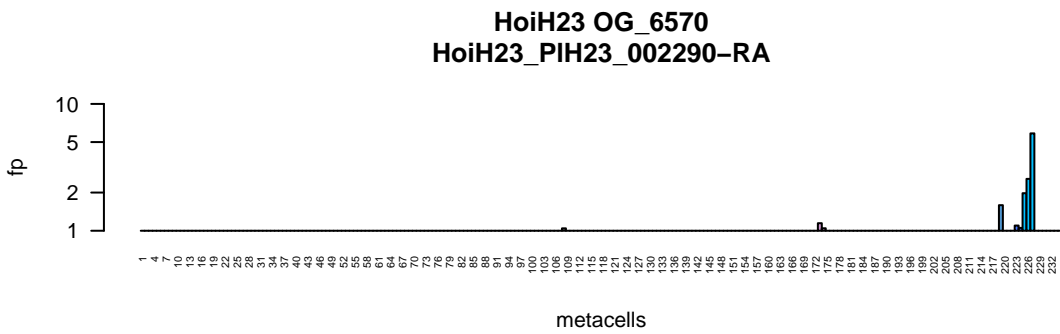
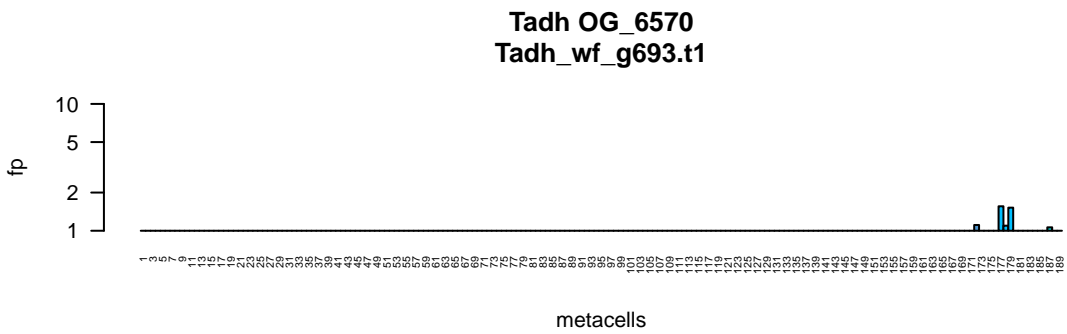
metacells

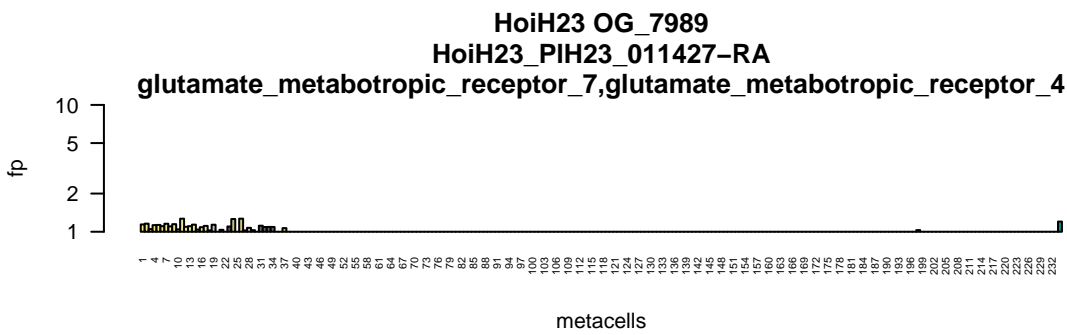
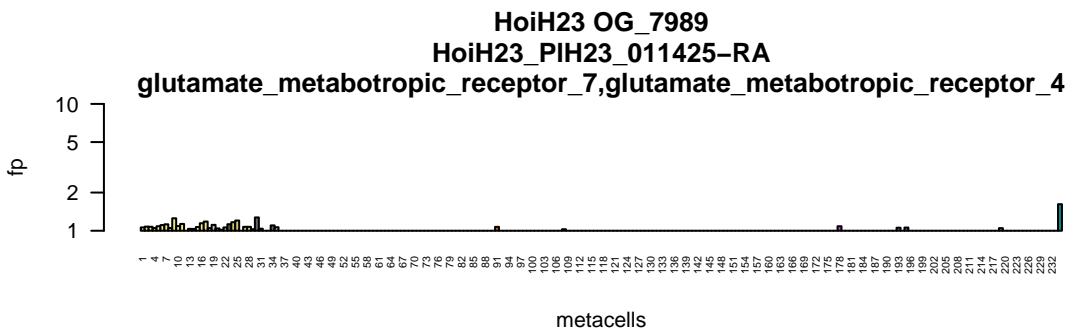
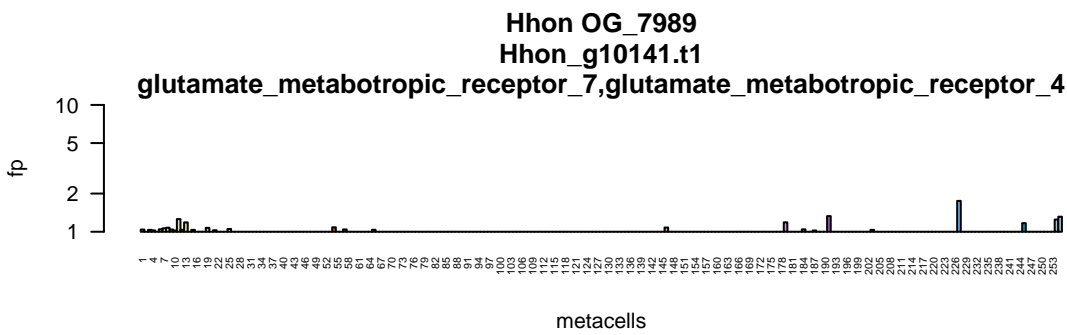
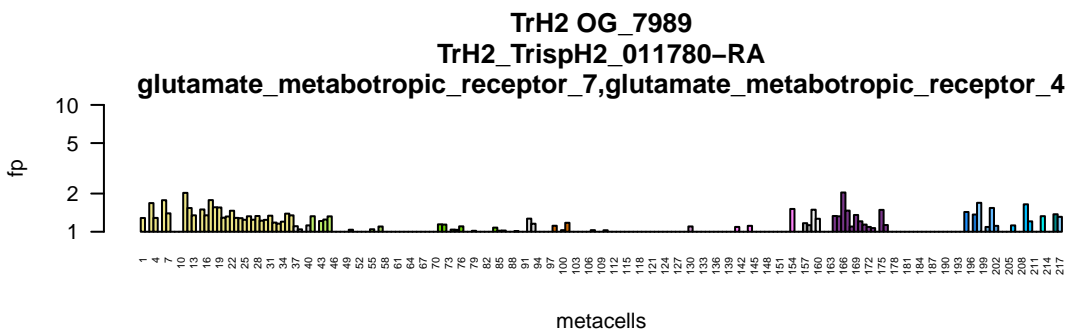
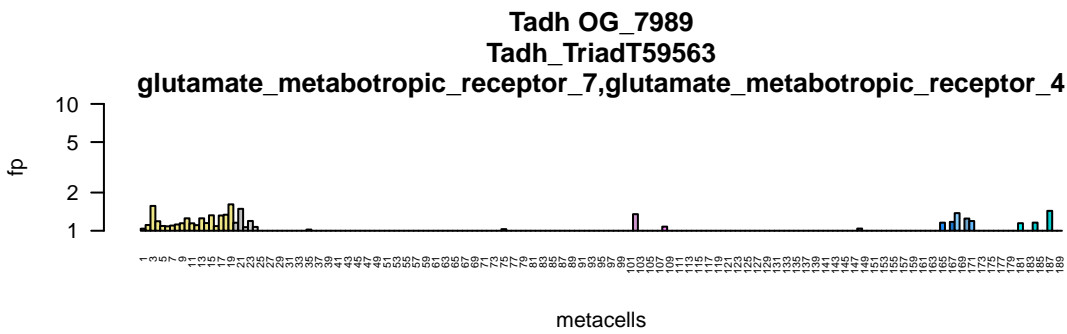
HoiH23 OG_5872
HoiH23_PIH23_005091-RA
sorting_nexin_13

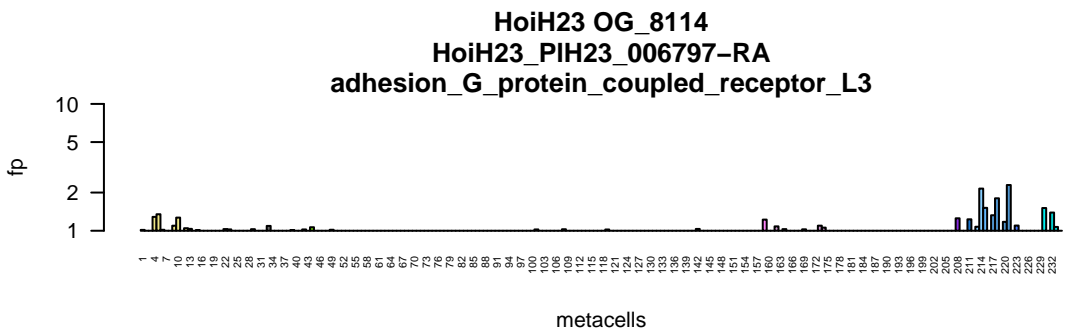
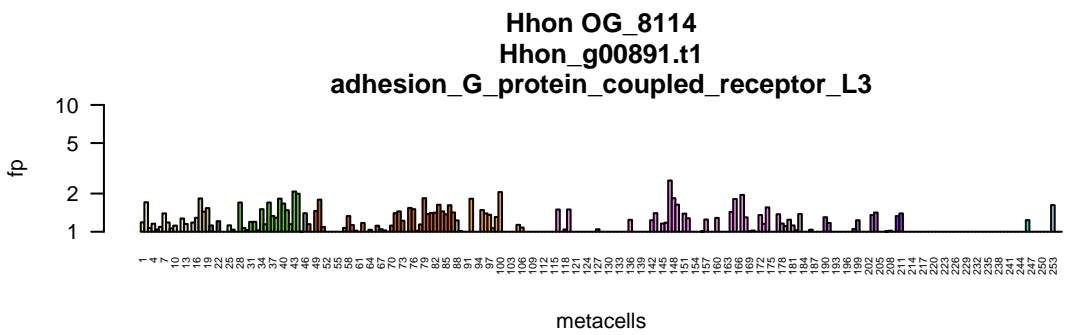
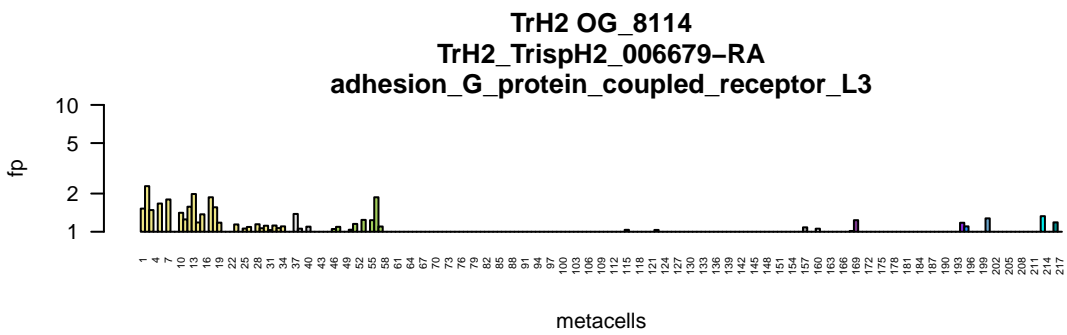
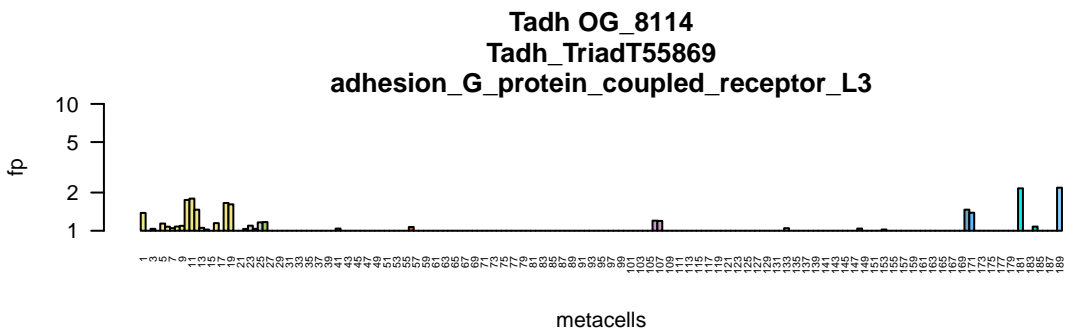


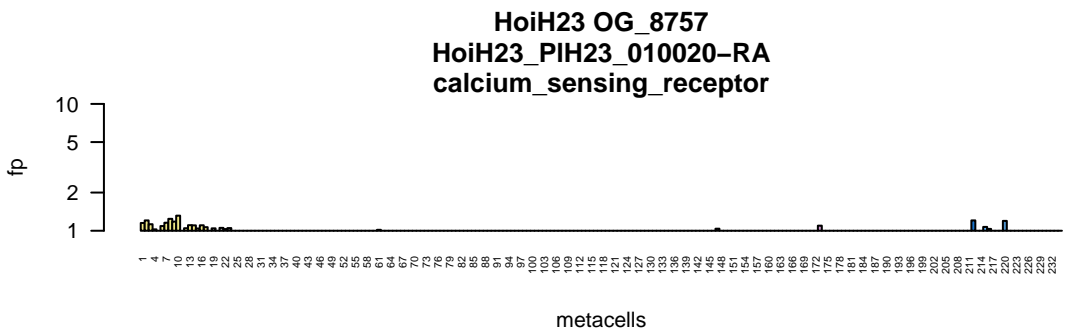
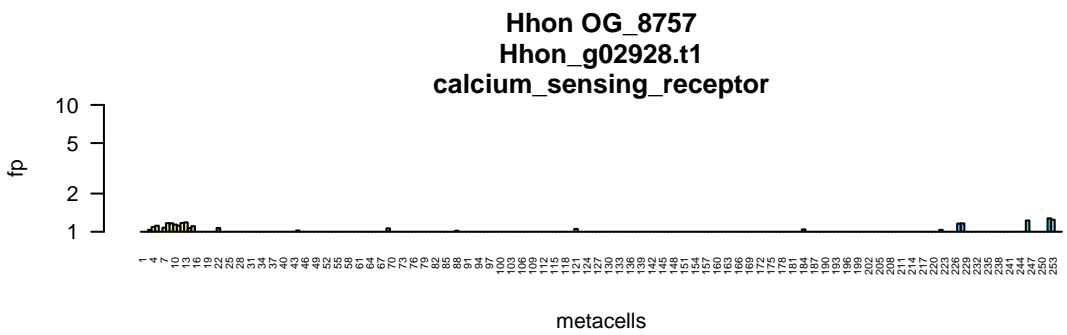
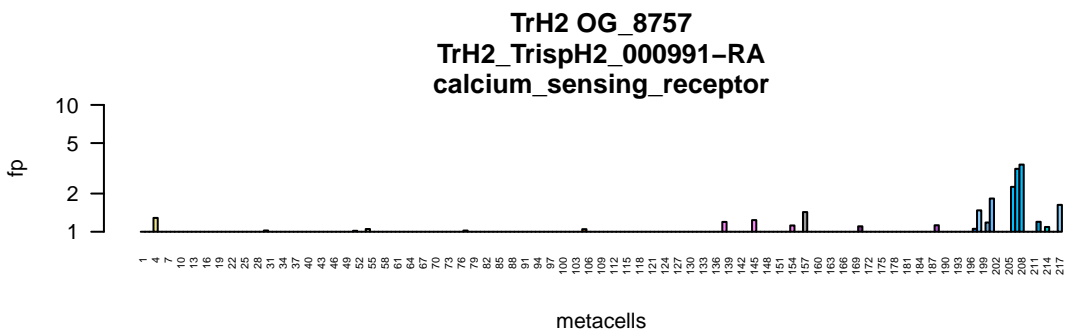
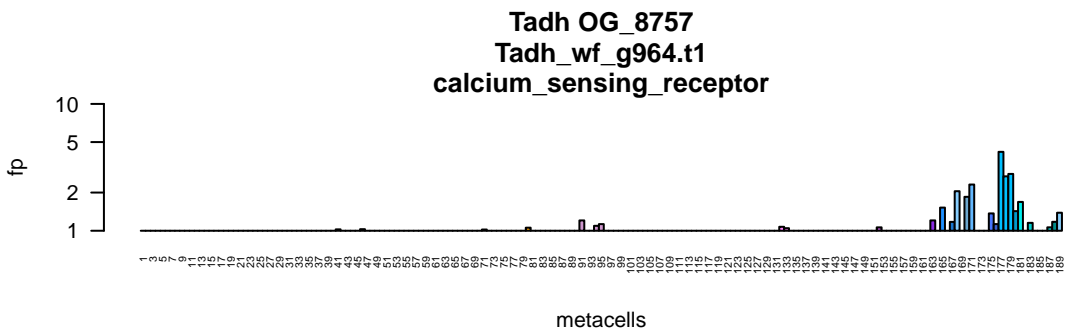
metacells



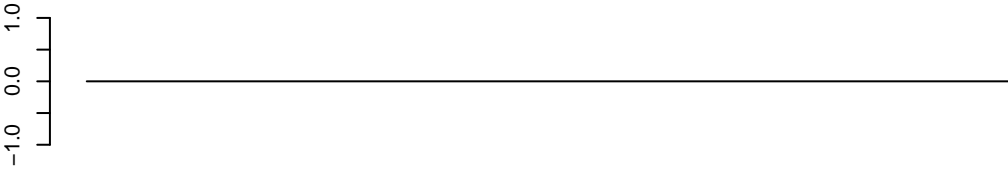




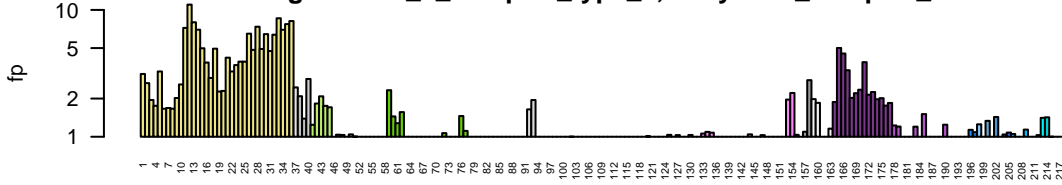




angiotensin_II_receptor_type_2,tachykinin_receptor_1
Tadh | no data

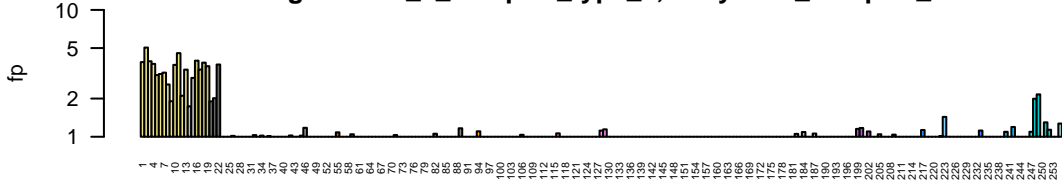


TrH2 OG_9264
TrH2_TrispH2_002603-RA
angiotensin_II_receptor_type_2,tachykinin_receptor_1



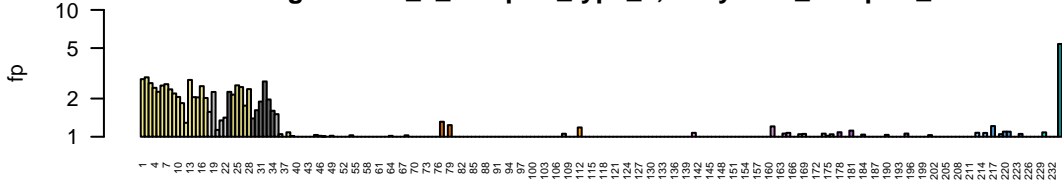
metacells

Hhon OG_9264
Hhon_g05345.t1
angiotensin_II_receptor_type_2,tachykinin_receptor_1



metacells

HoiH23 OG_9264
HoiH23_PIH23_008322-RA
angiotensin_II_receptor_type_2,tachykinin_receptor_1



metacells

