

Tadh_TriadT62237



Tadh_TriadT62236



Tadh_wf_g11342.t1



Tadh_TriadT62238



Tadh_wf_g11350.t1



Tadh_TriadT3775



Tadh_wf_g11493.t1



2_TrispH2_008282-RA



TrH2_TrispH2_010704-RA



TrH2_TrispH2_010702-RA



TrH2_TrispH2_010703-RA



TrH2_TrispH2_010706-RA



TrH2_TrispH2_011713-RA



TrH2_TrispH2_007495-RA

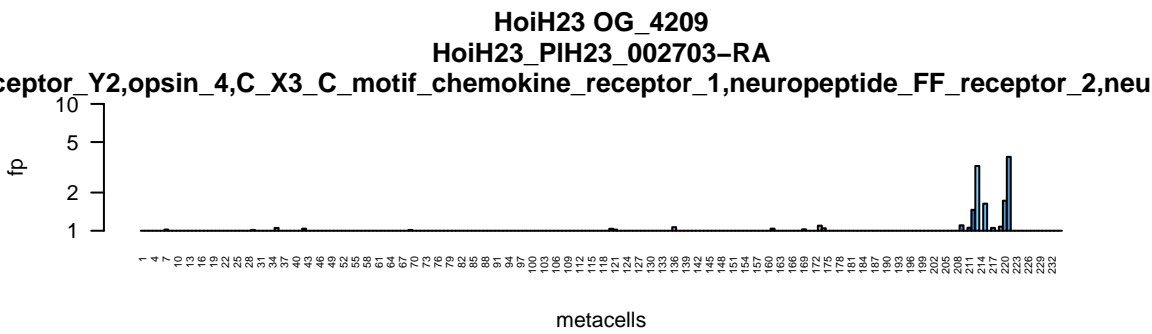
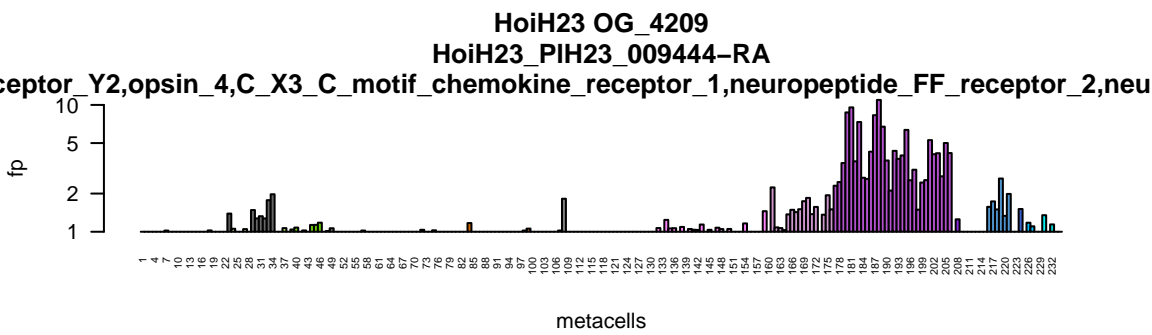
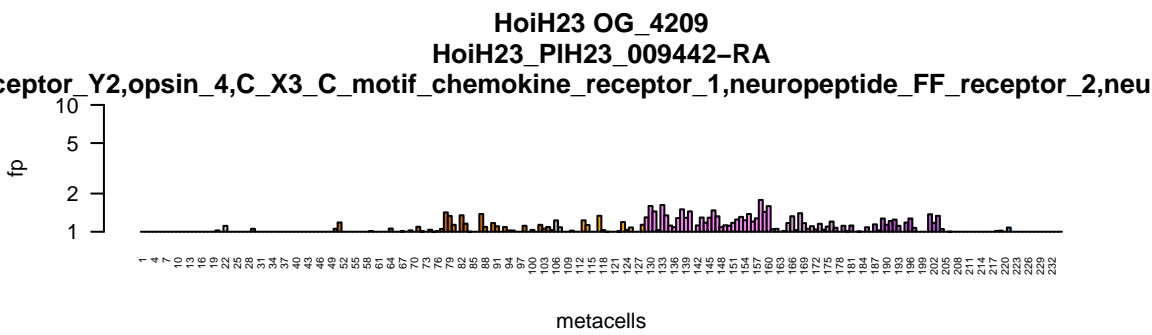
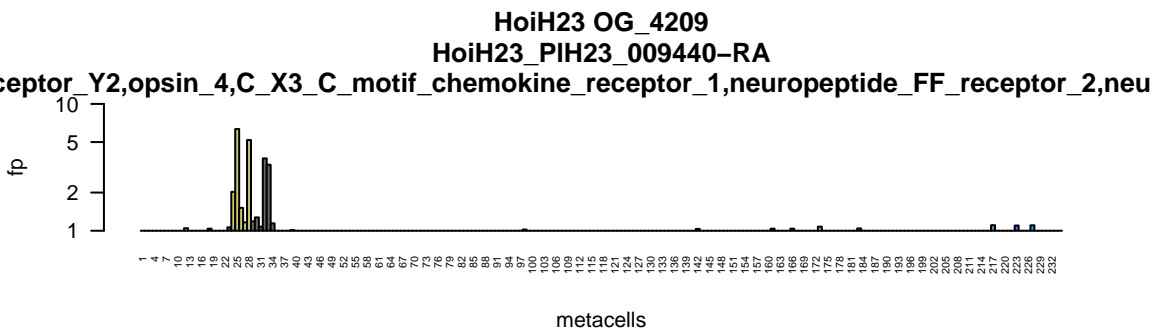
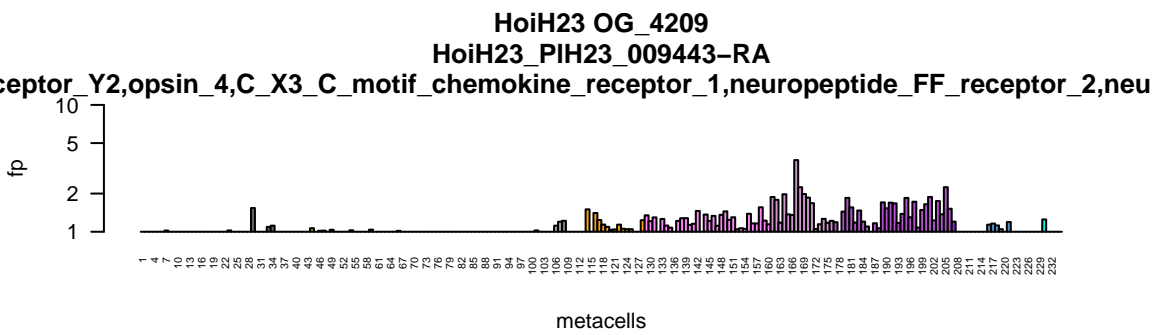
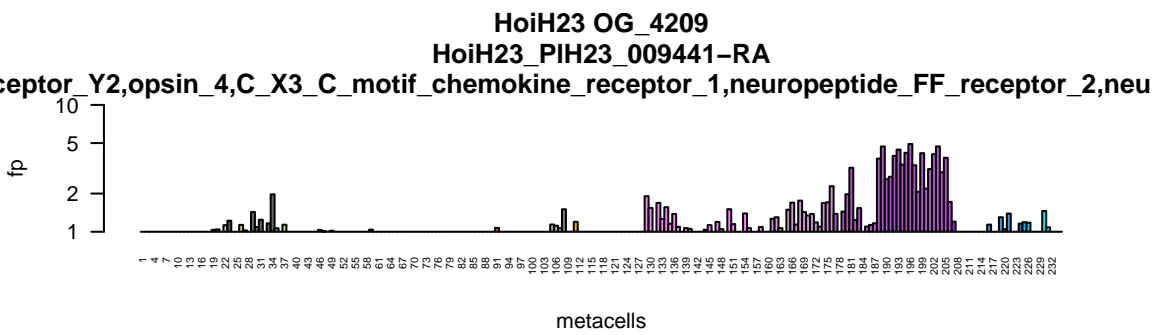
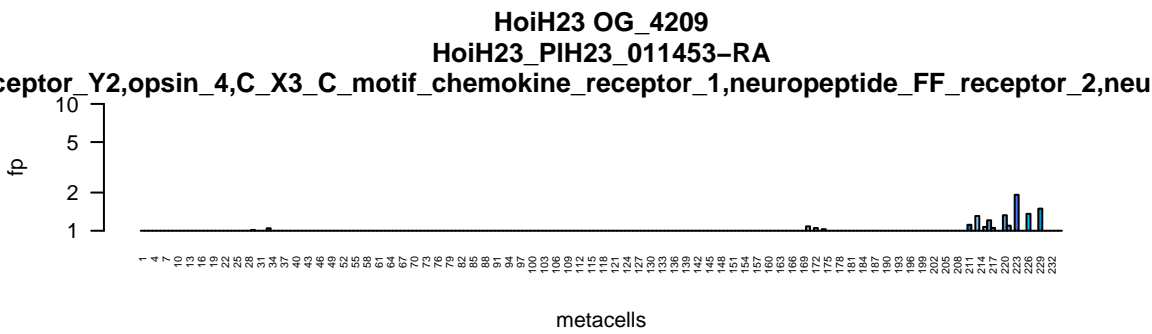


Phon_g11321.t1



Phon_g07103.t1





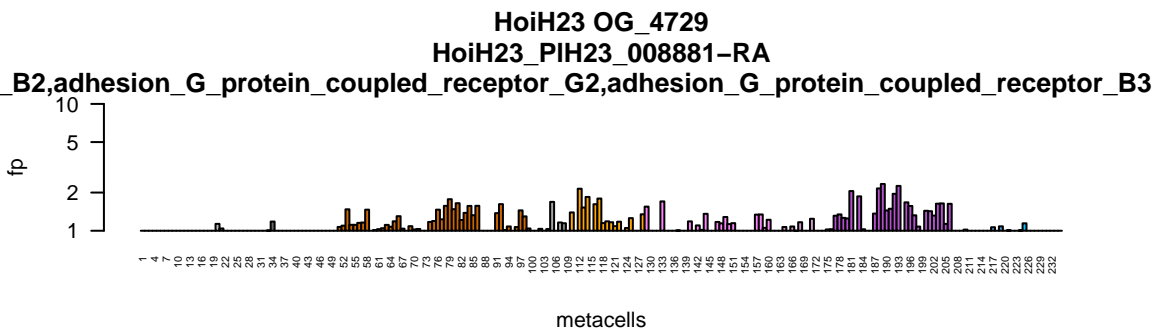
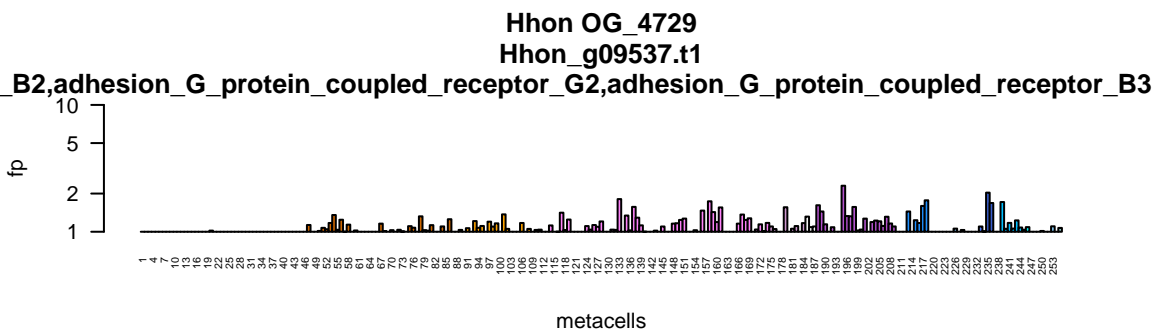
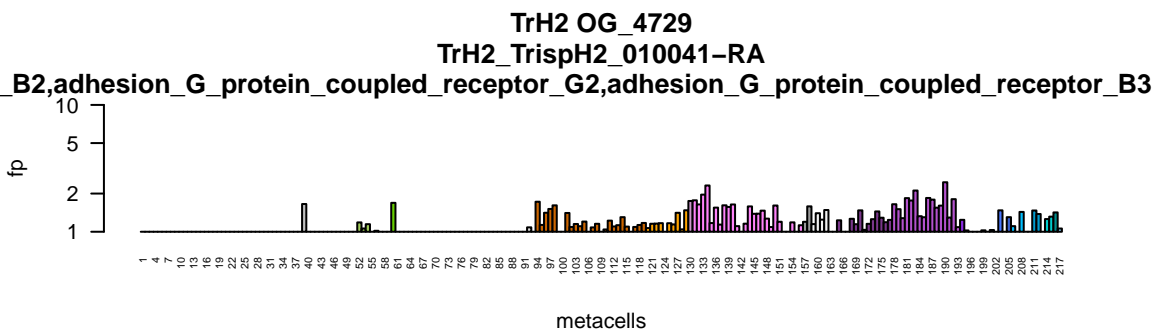
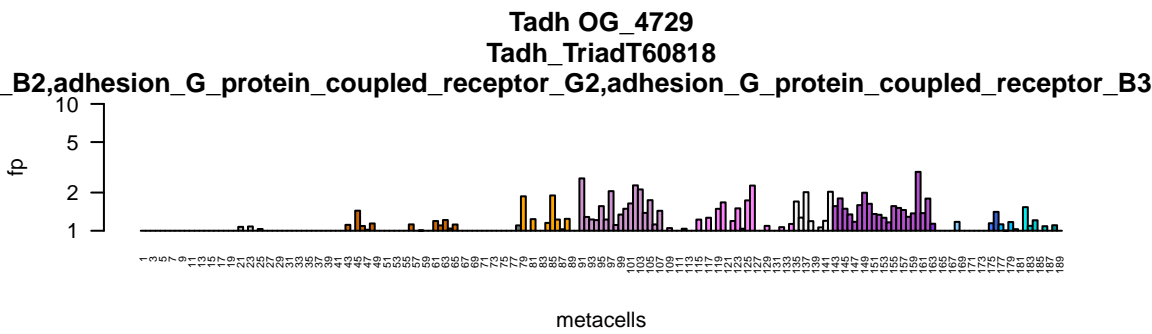
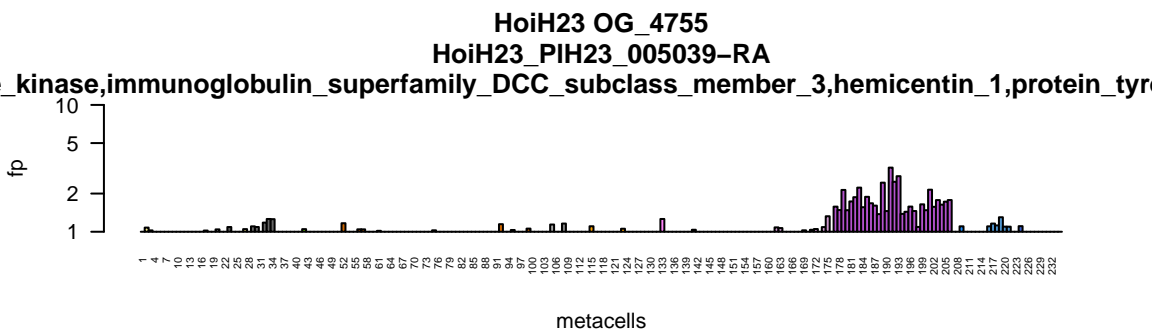
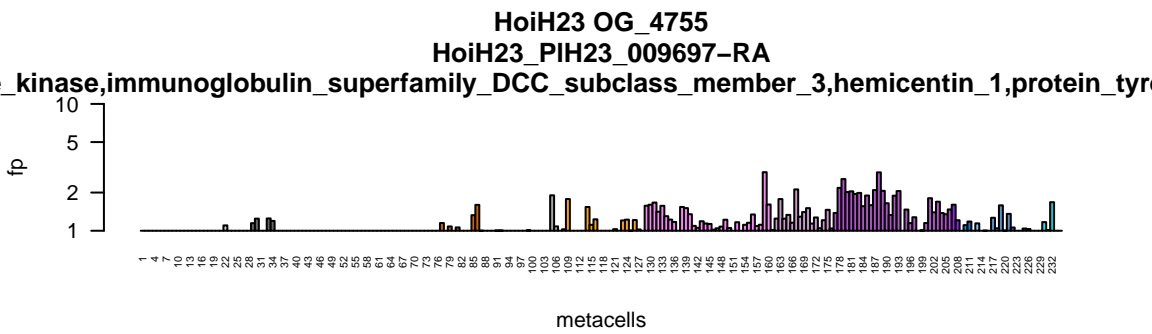
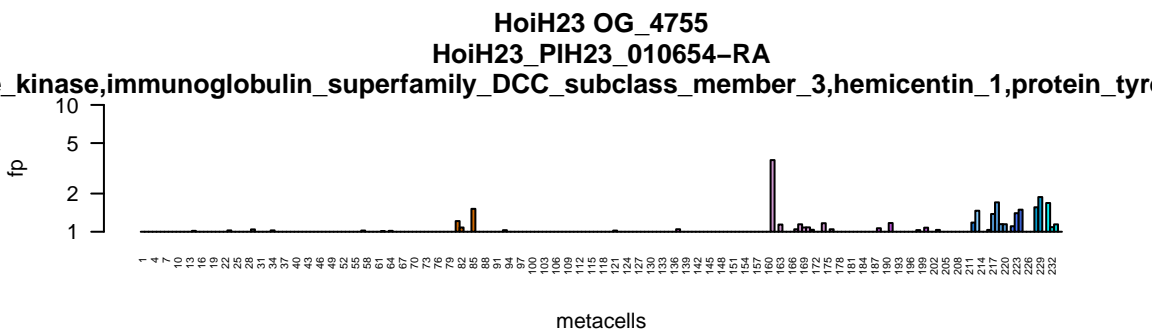
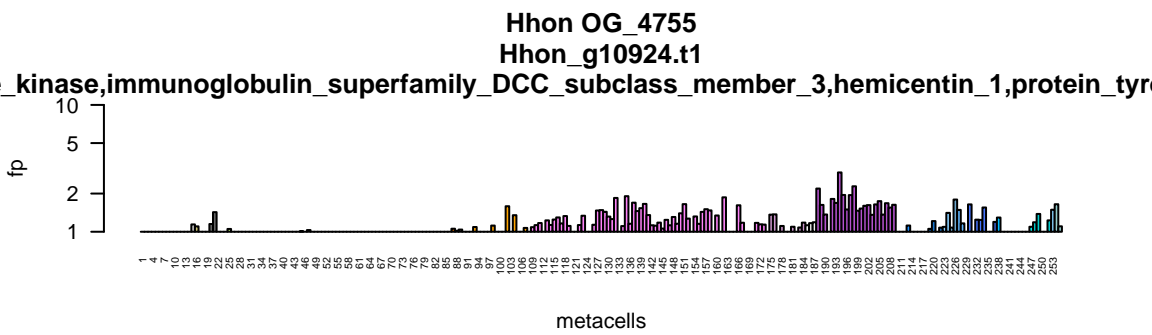
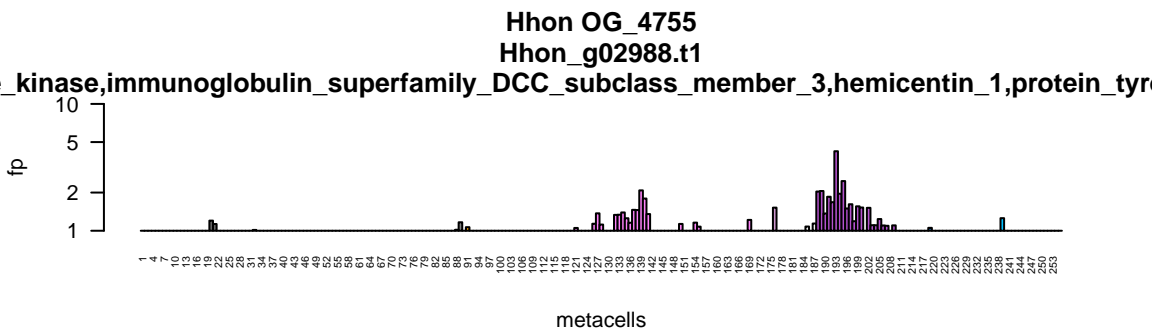


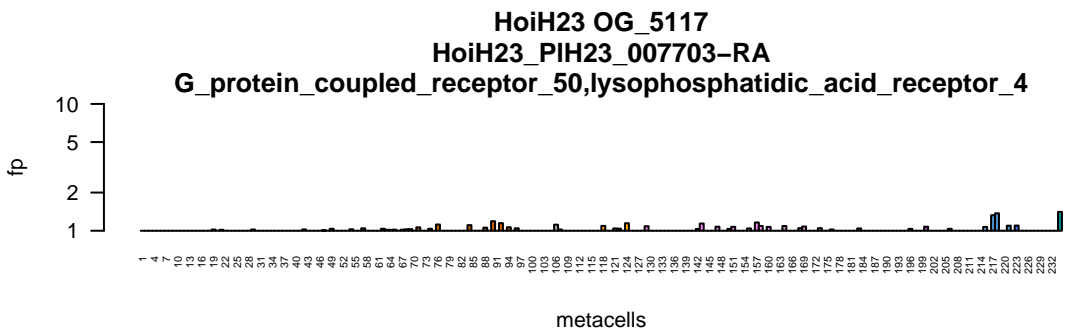
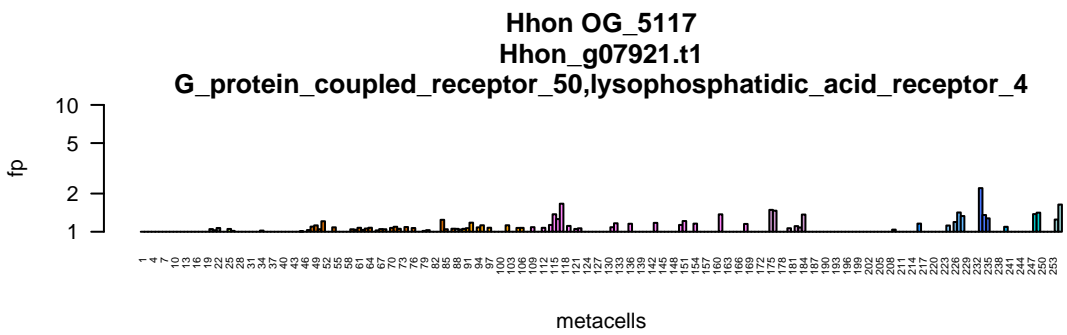
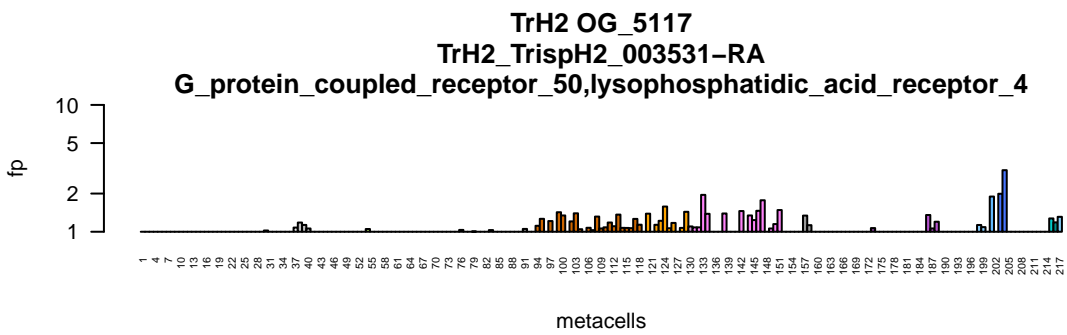
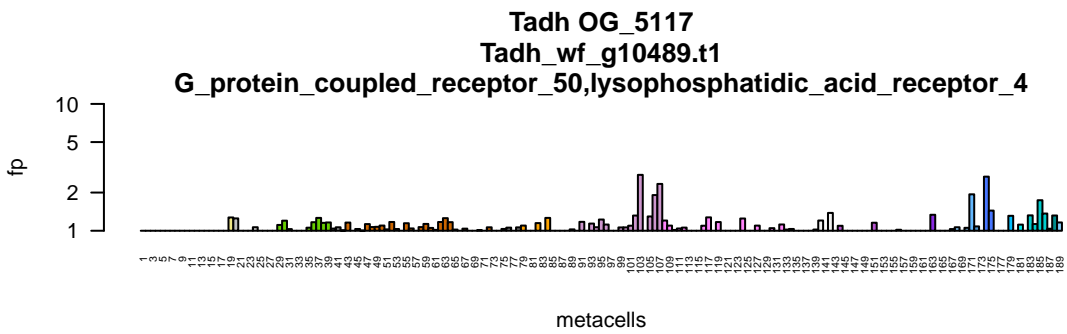
Figure 10: Frequency of metacells. The figure consists of two bar charts. The left chart shows the frequency of metacells for cell types 1 to 189, and the right chart shows the frequency of metacells for cell types 1 to 217. Both charts have a y-axis labeled 'fp' ranging from 1 to 10. The x-axis is labeled 'metacells'. The left chart shows a peak at cell type 189 with a frequency of 1. The right chart shows a peak at cell type 189 with a frequency of 1. Both charts show a general trend of increasing frequency with increasing cell type index.

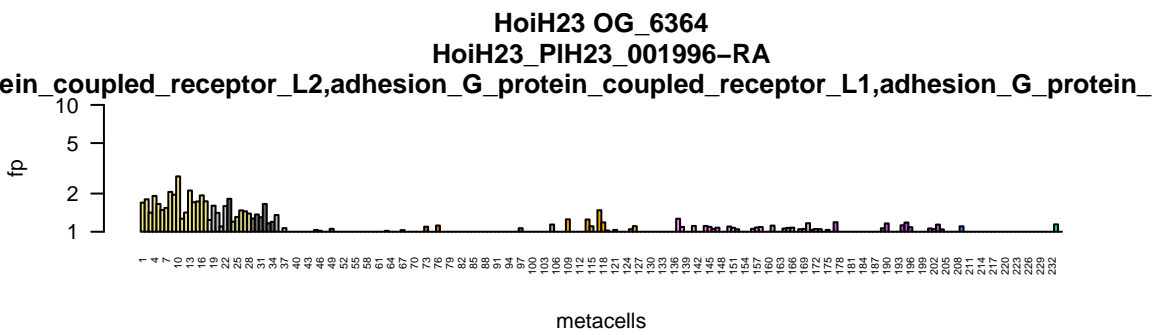
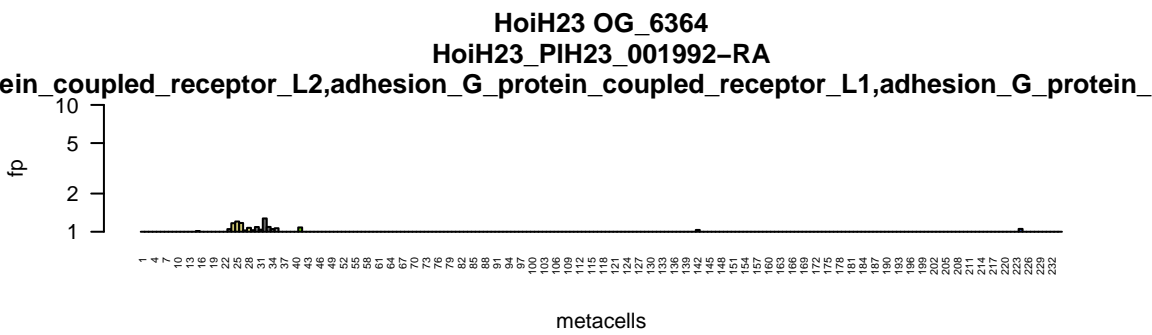
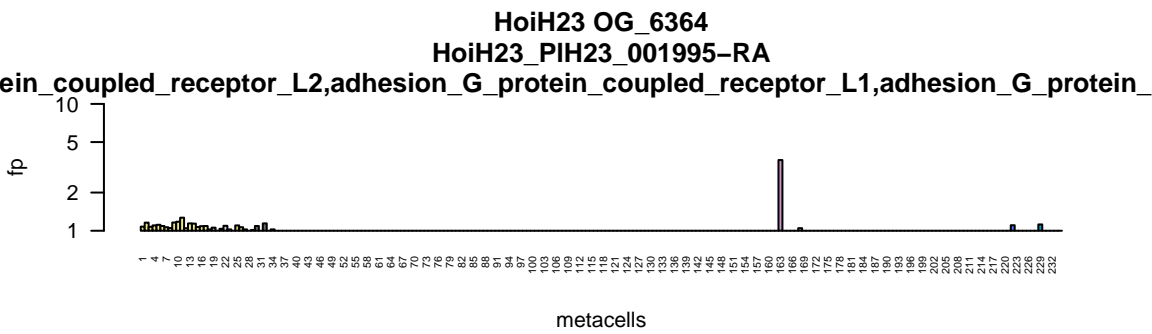
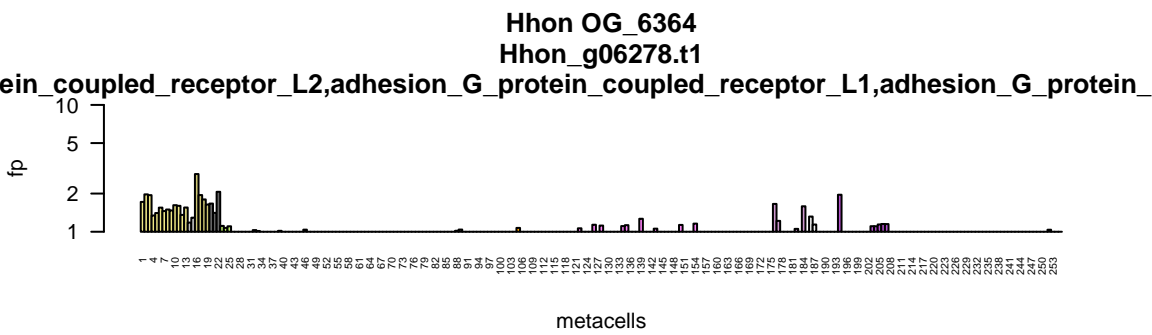
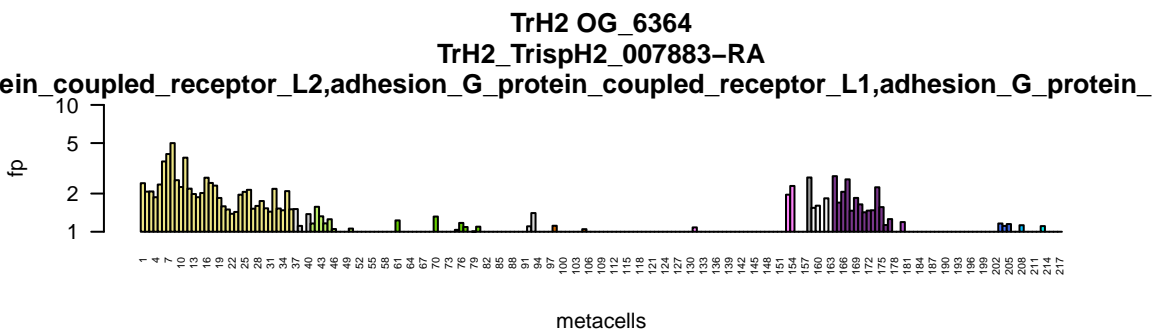
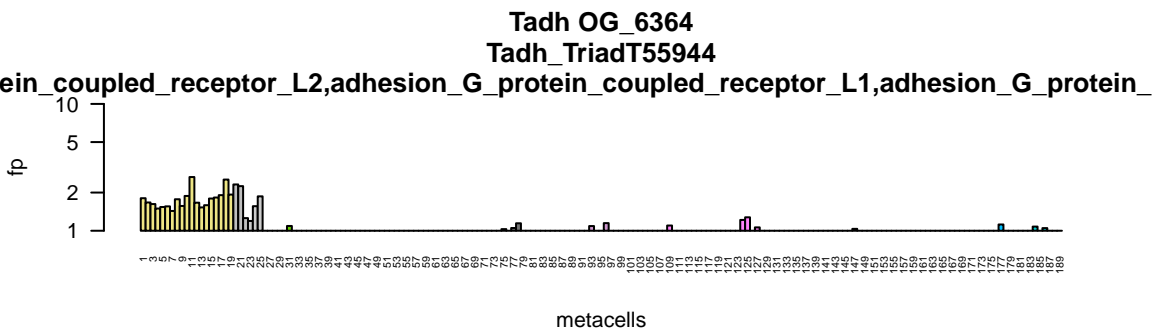
Two bar charts showing the frequency of metacells for different cell types. The left chart shows the frequency of metacells for the 'metacells' category, with a peak at metacell 135. The right chart shows the frequency of metacells for the 'metacells' category, with a peak at metacell 163. Both charts have a y-axis labeled 'fp' ranging from 1 to 10.

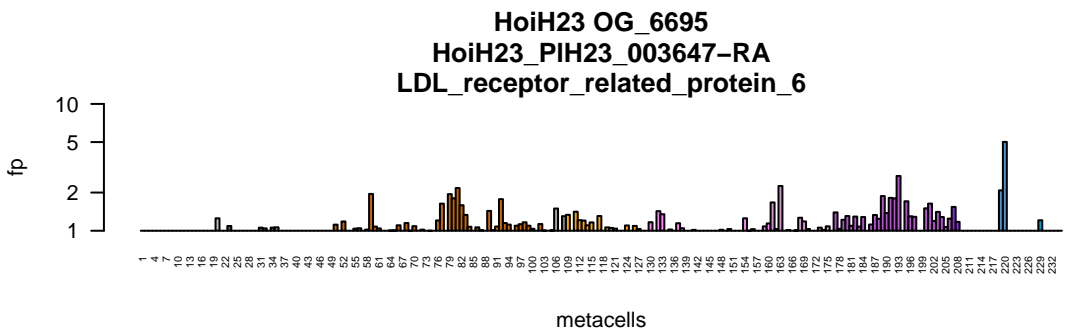
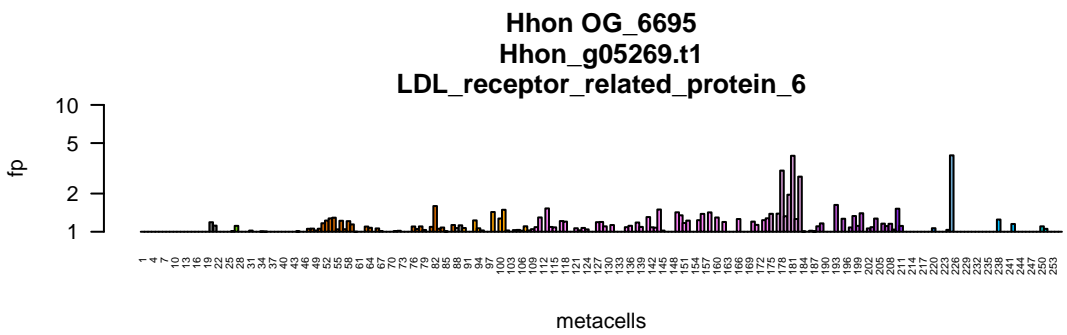
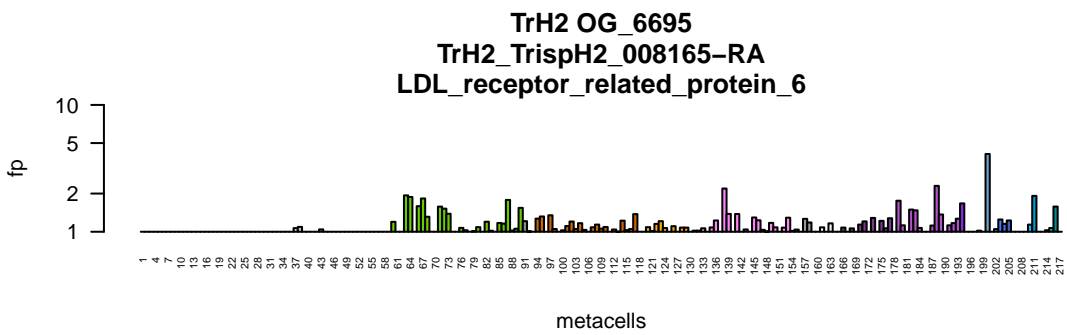
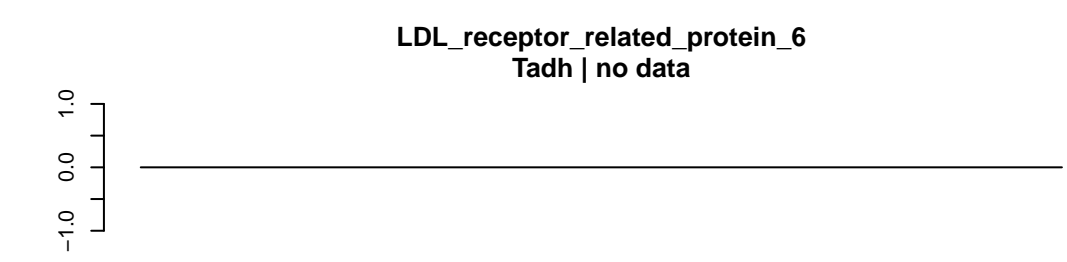
The figure consists of two side-by-side bar plots. Both plots have 'metacells' on the x-axis and 'fp' (false positive rate) on the y-axis. The y-axis has labels at 1, 2, 5, and 10.

- Left Plot:** Shows very low fp values across all metacells. Most bars are at or below 1. There are a few small peaks above 1, notably around metacell 133 and 189.
- Right Plot:** Shows higher fp values, particularly for metacells 160-217. The fp values range from approximately 1 to 3. There are several distinct peaks, with the highest being around metacell 160 (fp ≈ 3).

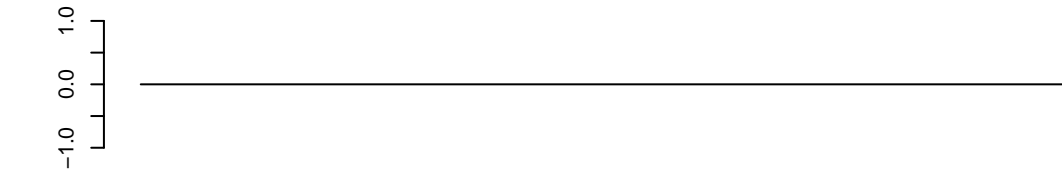




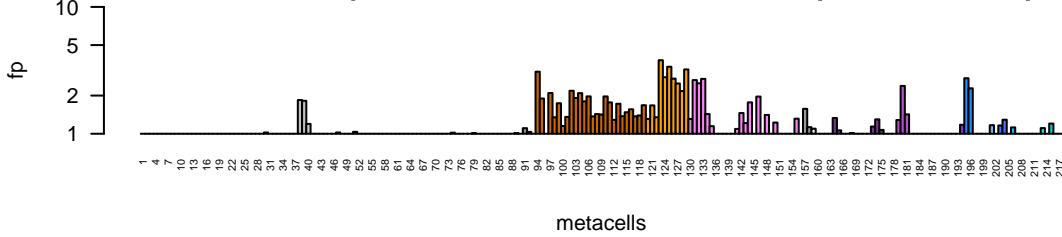




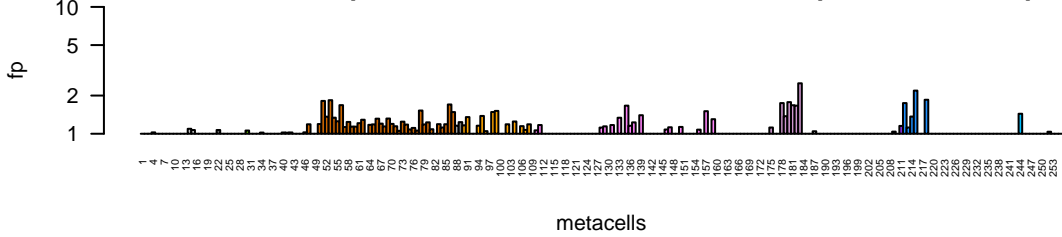
C_C_motif_chemokine_receptor_1,trace_amine_associated_receptor_8,adrenoceptor_alpha2
Tadh | no data



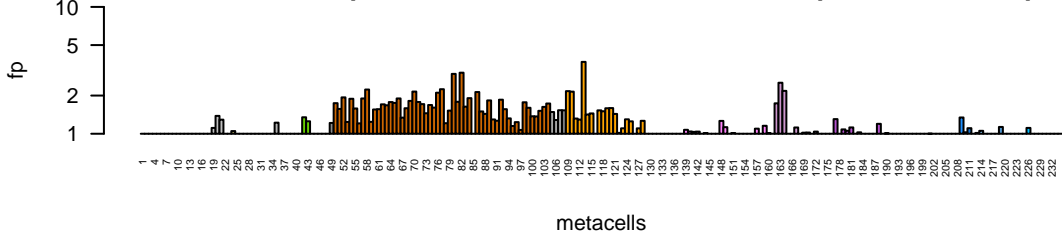
TrH2 OG_7015
TrH2_TrispH2_003193-RA
C_C_motif_chemokine_receptor_1,trace_amine_associated_receptor_8,adrenoceptor_alpha2

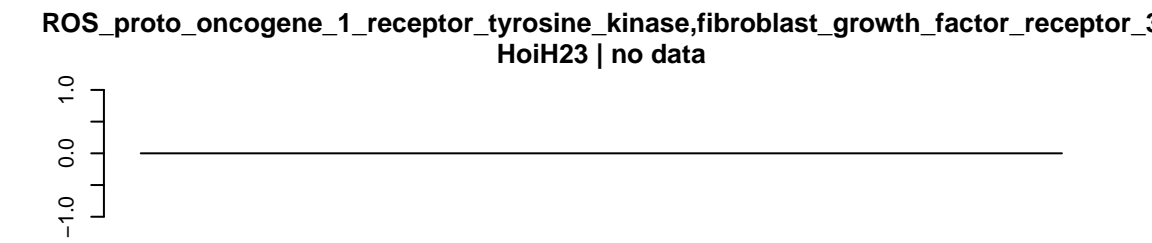
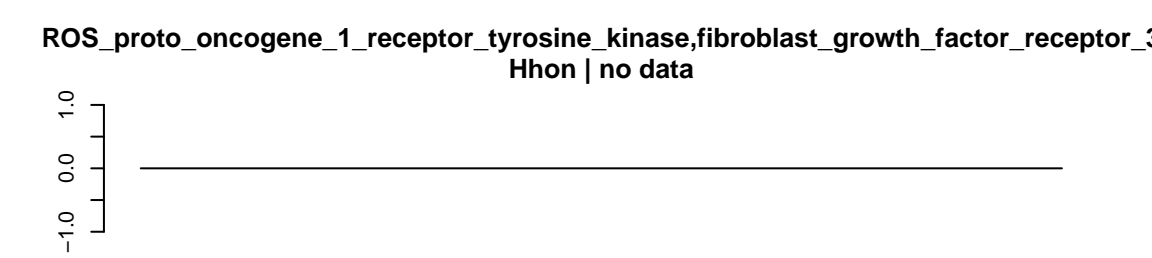
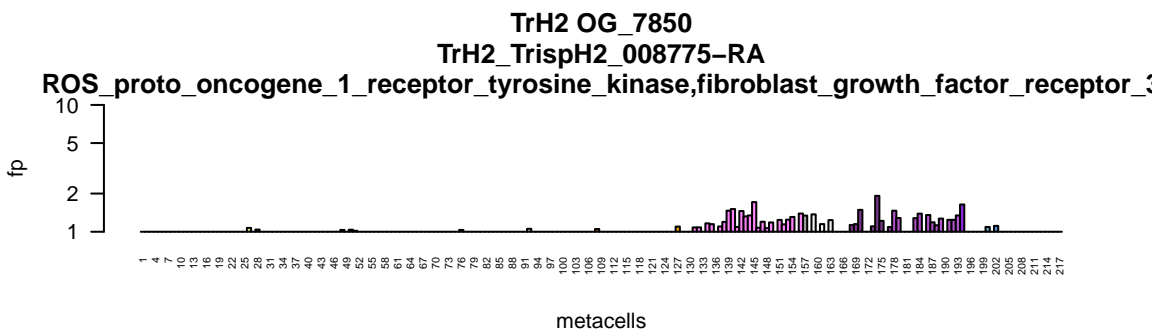
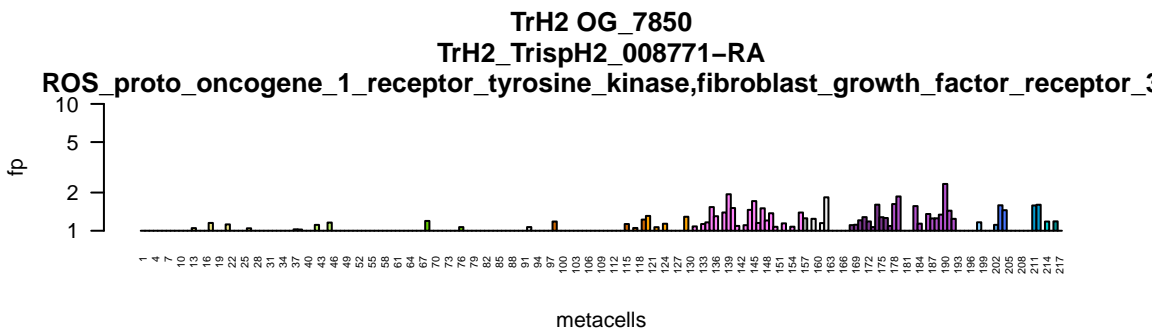
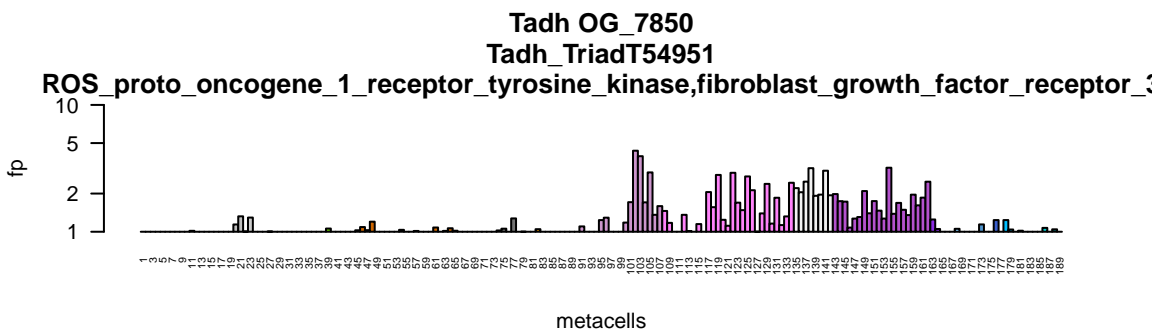
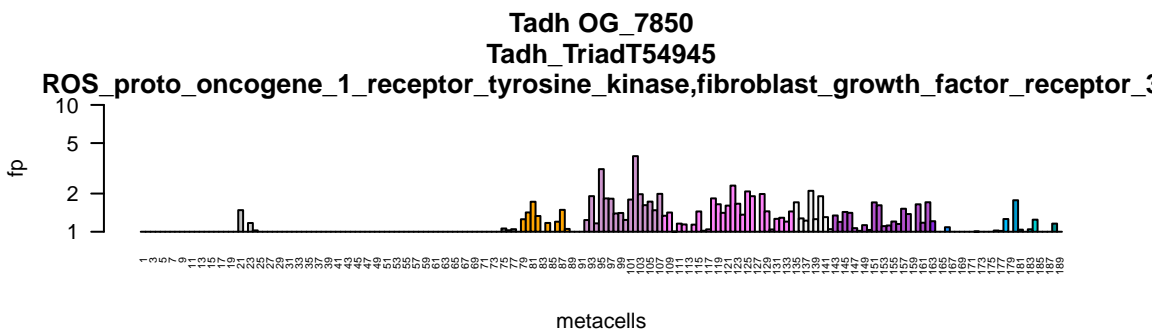
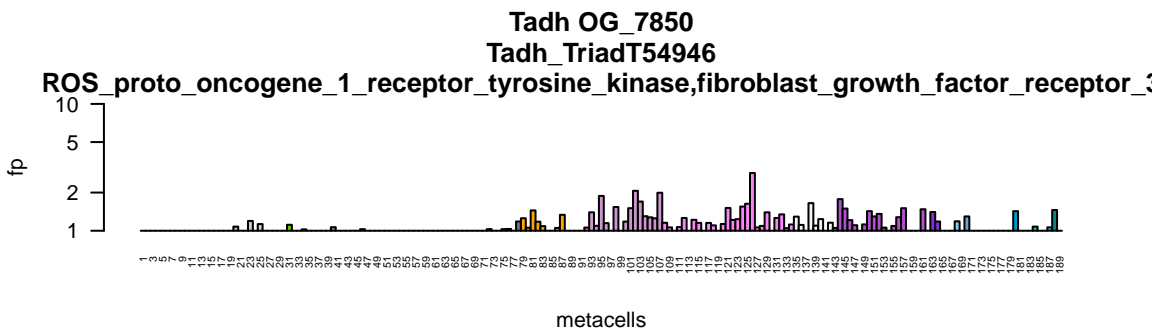


Hhon OG_7015
Hhon_g03672.t1
C_C_motif_chemokine_receptor_1,trace_amine_associated_receptor_8,adrenoceptor_alpha2

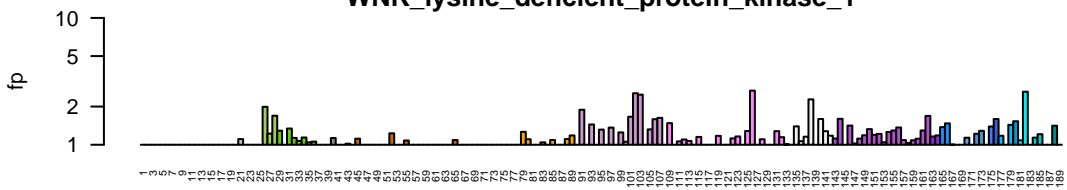


HoiH23 OG_7015
HoiH23_PIH23_002074-RA
C_C_motif_chemokine_receptor_1,trace_amine_associated_receptor_8,adrenoceptor_alpha2



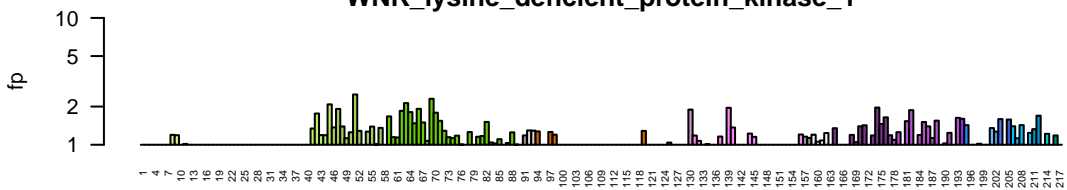


Tadh OG_7856
Tadh_TriadT4059
WNK_lysine_deficient_protein_kinase_1



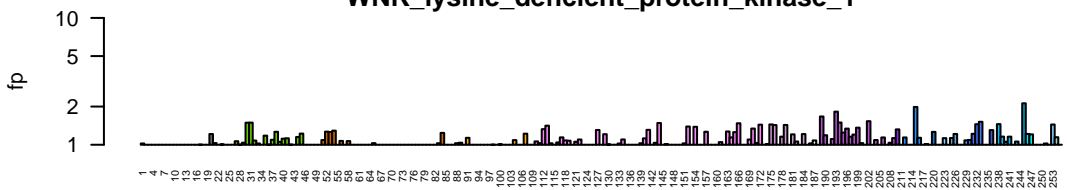
metacells

TrH2 OG_7856
TrH2_TrispH2_007460-RA
WNK_lysine_deficient_protein_kinase_1



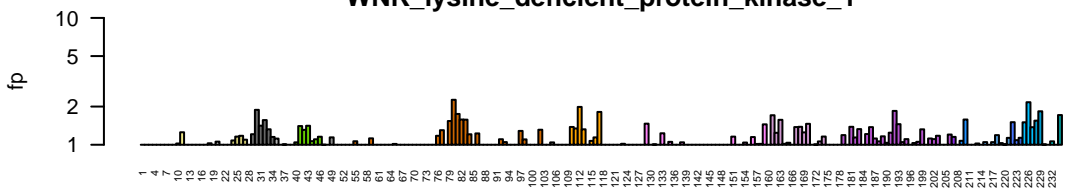
metacells

Hhon OG_7856
Hhon_g08521.t1
WNK_lysine_deficient_protein_kinase_1



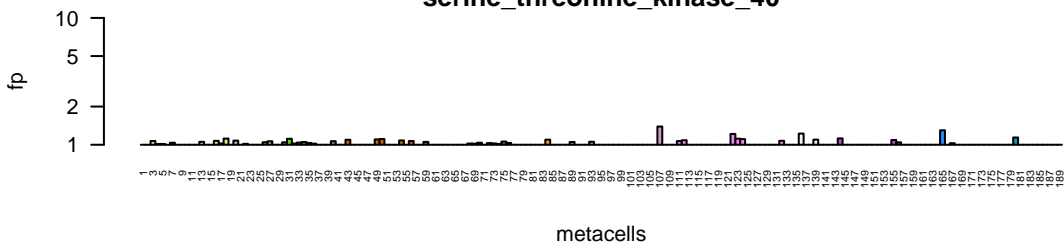
metacells

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HoiH23_PIH23_009337-RA
WNK_lysine_deficient_protein_kinase_1

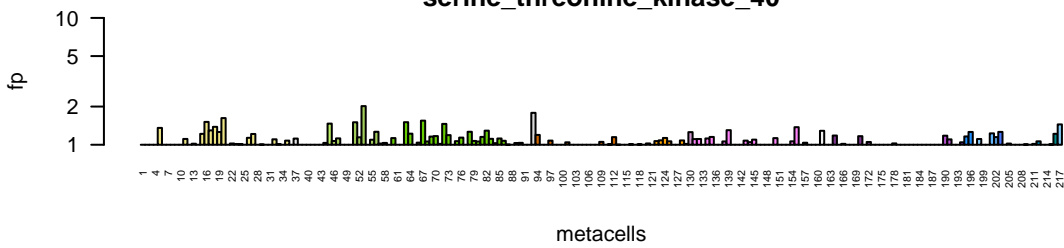


metacells

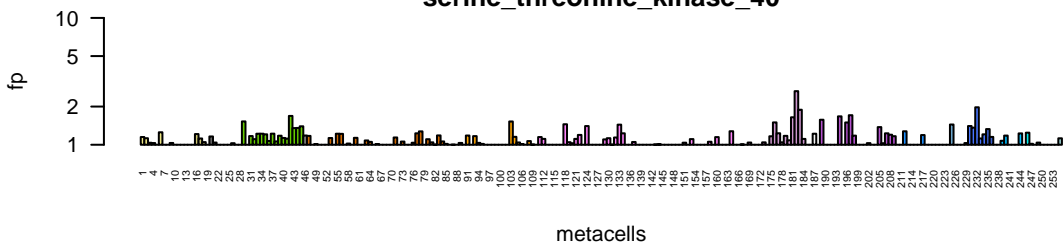
Tadh OG_8034
Tadh_TriadT26082
serine_threonine_kinase_40



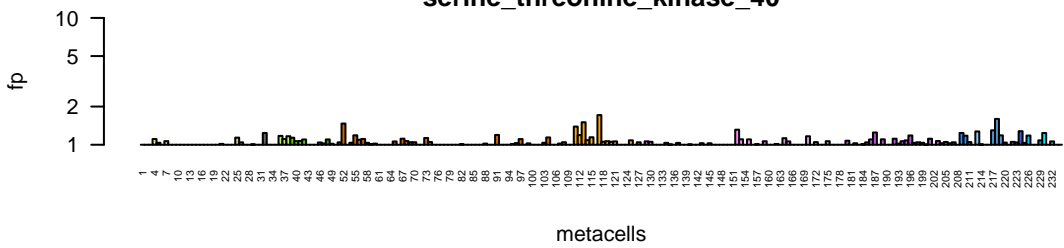
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TrH2_TrispH2_007966-RA
serine_threonine_kinase_40



Hhon OG_8034
Hhon_g09161.t1
serine_threonine_kinase_40



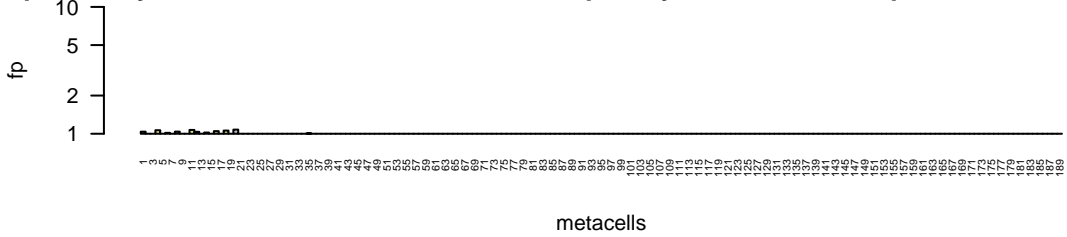
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serine_threonine_kinase_40



Tadh OG_8521

Tadh_TriadT55132

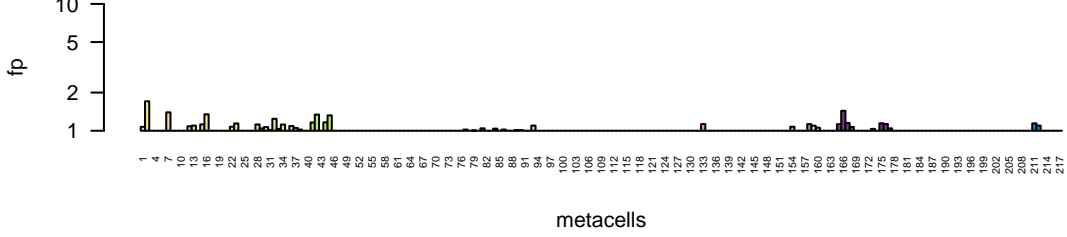
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TrH2 OG_8521

TrH2_TrispH2_011101-RA

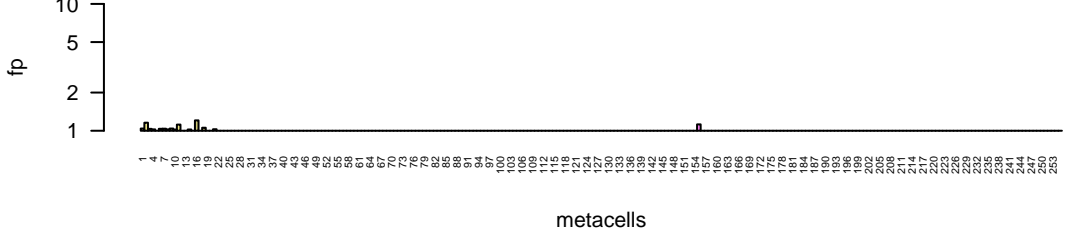
protein_tyrosine_kinase_6,fms_related_receptor_tyrosine_kinase_1,protein_kinase_C_ze



Hhon OG_8521

Hhon_g07209.t1

protein_tyrosine_kinase_6,fms_related_receptor_tyrosine_kinase_1,protein_kinase_C_ze



HoiH23 OG_8521

HoiH23_PIH23_004412-RA

protein_tyrosine_kinase_6,fms_related_receptor_tyrosine_kinase_1,protein_kinase_C_ze

