

A bar chart showing the frequency of metacells (x-axis) across different frequency bins (y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows a distribution of metacells across frequency bins, with a notable peak at metacell 202.

metacells	fp
1	1
4	1
7	1
13	1
16	1
19	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
202	4
205	1
208	1
211	1
214	1
217	1

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacell	fp
1	1
4	1
17	1
13	1
16	1
19	1
25	1
28	1
31	1
34	1
40	1
43	1
46	1
52	1.5
55	1
58	1
61	1
67	1
70	1
73	1
79	1
82	1
85	1
88	1
94	1.2
97	1
100	1
103	1
106	1
109	1
112	1
118	1
121	1
124	1
127	1
133	1
136	1
139	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	1
169	1
172	1
175	1.8
179	1
184	1
187	1
190	1
193	1
199	1.8
202	1.8
205	1.5
211	1
214	1.2
217	1.8

metacell	fp
1	0
4	0
7	0
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	0
34	0
37	0
40	0
43	0
46	0
49	0
52	1
55	0
58	0
61	0
64	0
67	0
70	0
73	0
76	0
79	0
82	0
85	0
88	0
91	0
94	0
97	0
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	0
130	0
133	1
136	0
139	0
142	0
145	0
148	0
151	0
154	1
155	1
156	0
159	0
162	0
165	0
168	0
171	0
174	0
177	0
180	0
183	0
186	0
189	1
192	0
195	0
198	0
200	2
201	2
202	1
203	1
204	1
205	1
206	1
207	1
208	1
209	1
210	1
211	1
214	1
217	1

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacell	fp
1	1
4	1
13	1
16	1.2
17	1
19	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
52	1
55	1
58	1
61	1
67	1
70	1
73	1.1
77	1
79	1
82	1
85	1
88	1
94	1
97	1
100	1.3
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1.1
133	1.2
136	1
139	1
142	1
145	1
148	1
151	1
154	1.3
157	1
160	1
163	1
166	1
169	1
172	1
175	1
179	1
181	1
184	1
187	1
190	1
193	1.5
196	1.8
199	1.5
202	2.2
205	1.5
208	1.2
211	1.2
214	1.2
217	1.5

A bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists indices from 1 to 217. Most metacells have 1 false positive, but metacells 202 and 214 have 2 false positives.

metacell	fp
1	1
4	1
7	1
10	1
13	1
16	1
19	1
22	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
202	2
205	1
208	1
211	1
214	2
217	1

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

fp

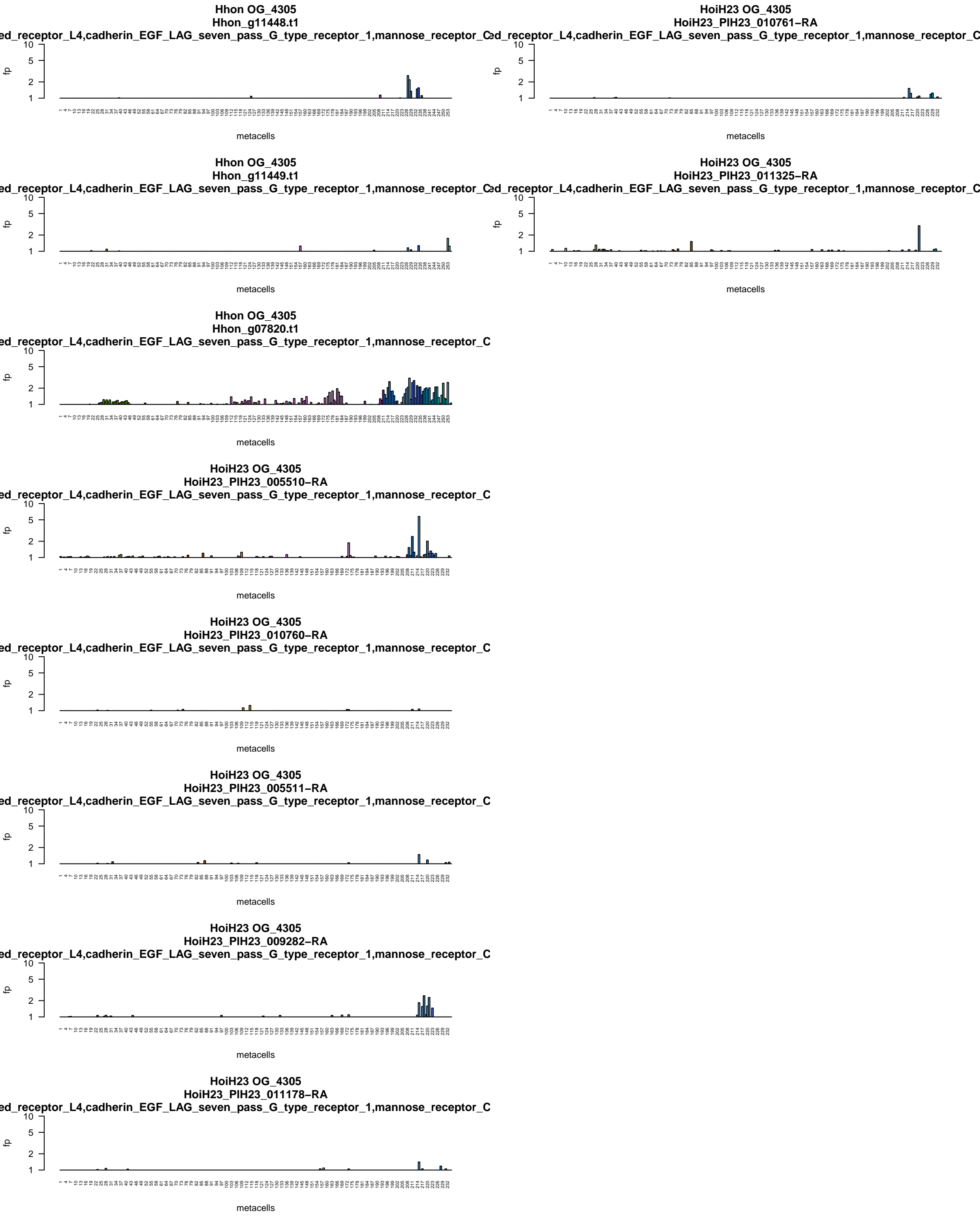
metacells

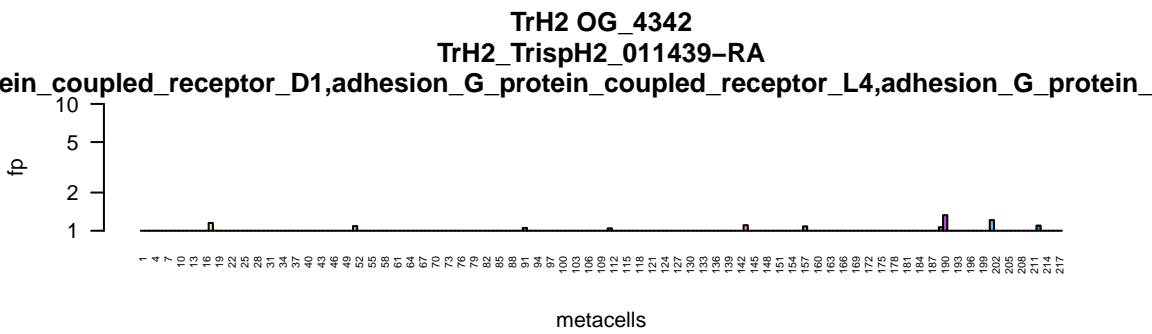
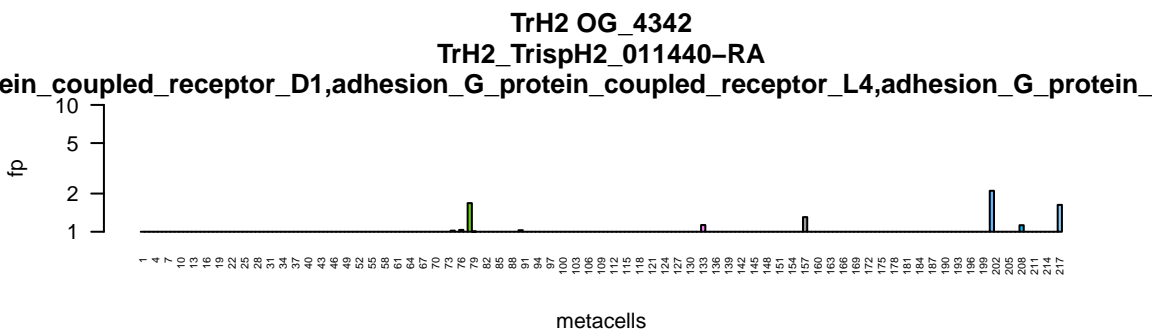
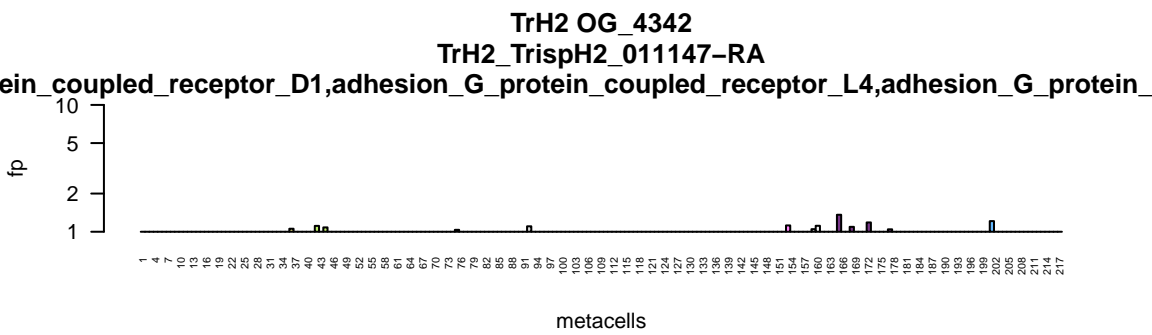
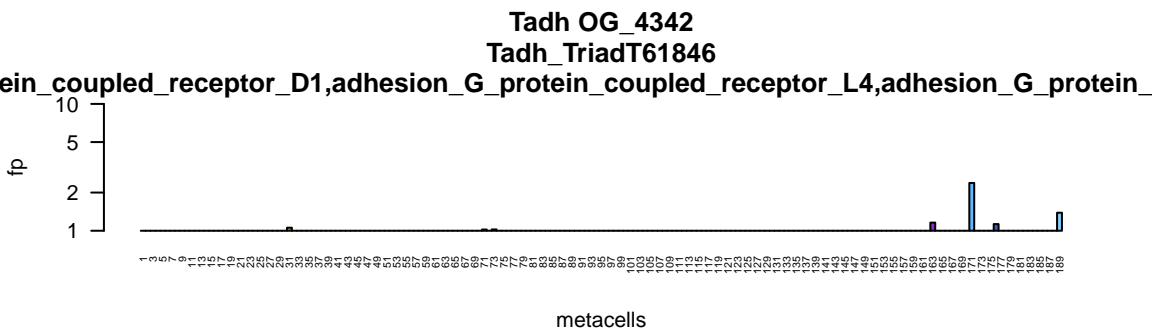
Hhon OG_4305
Hhon_g09074.t1

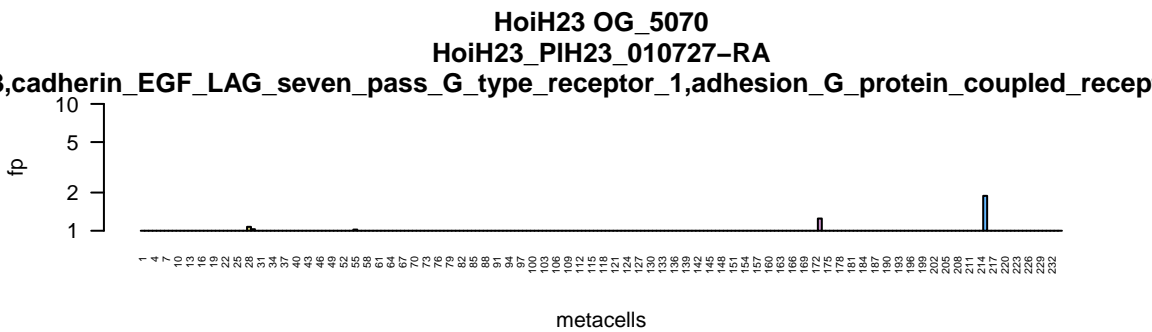
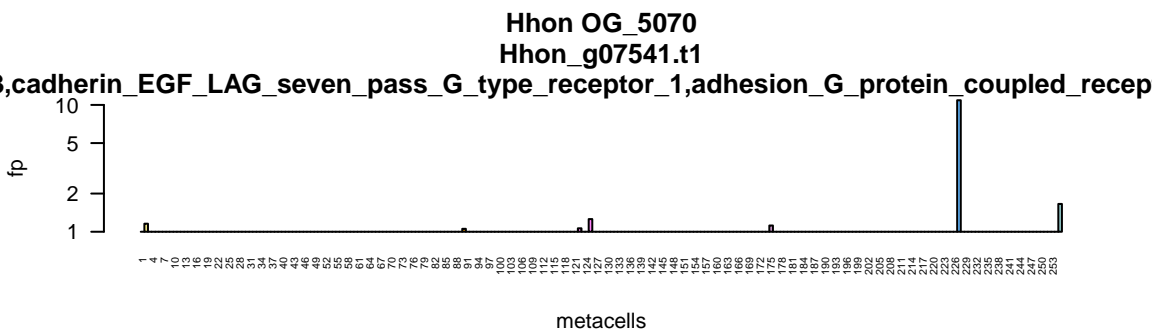
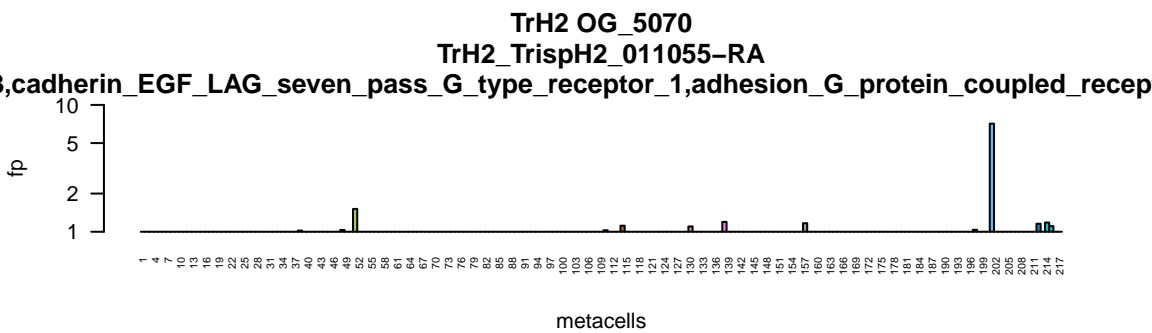
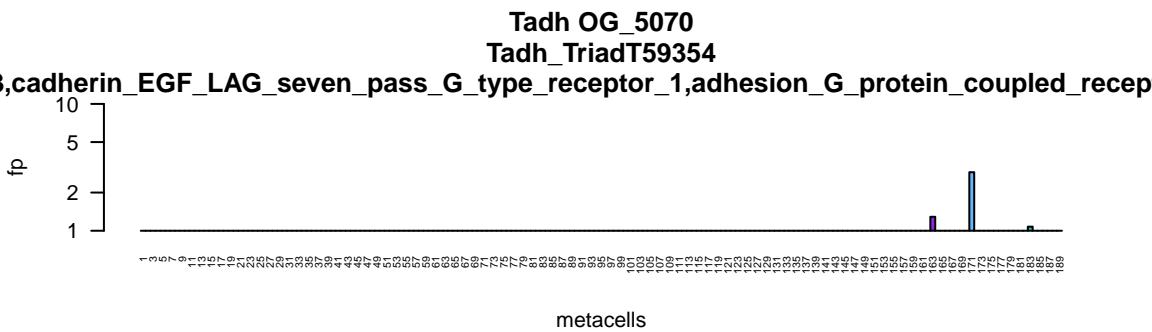
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

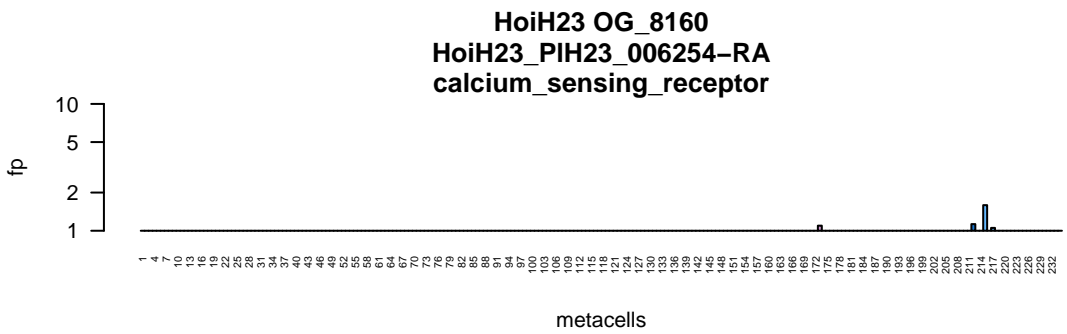
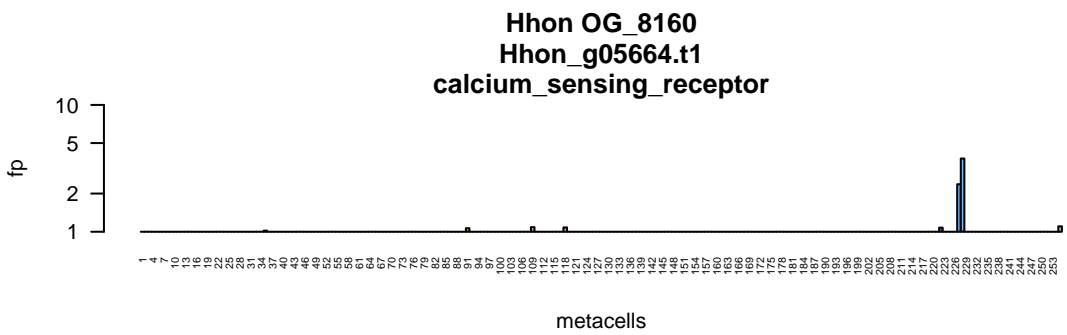
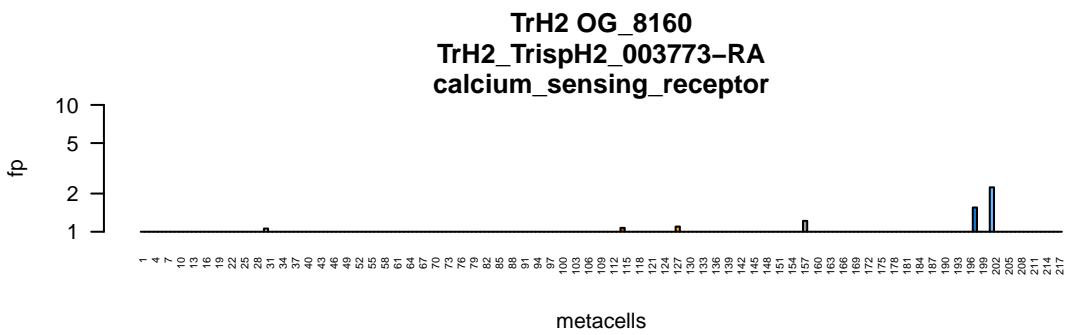
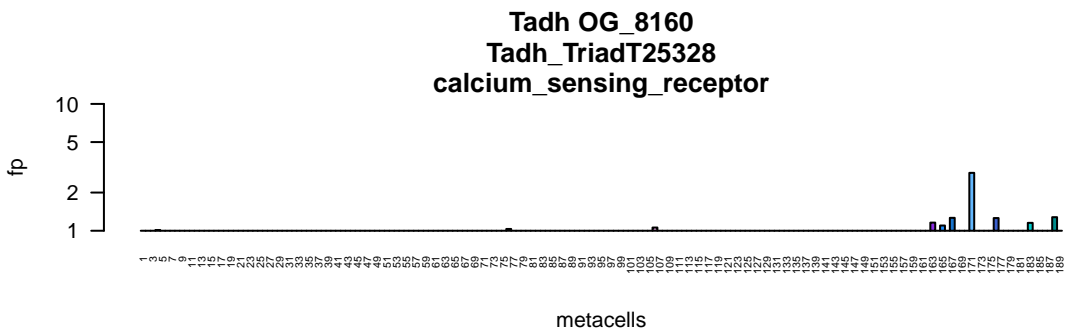
fp

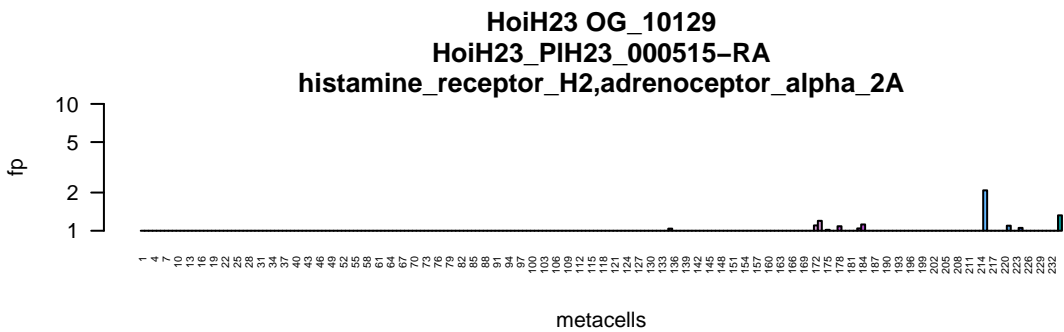
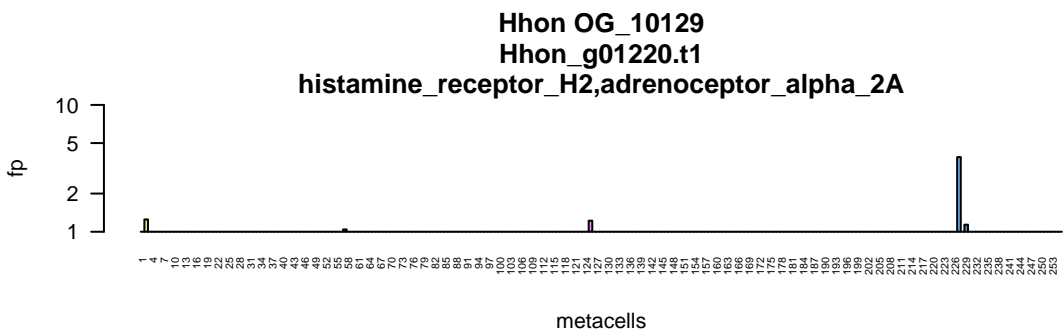
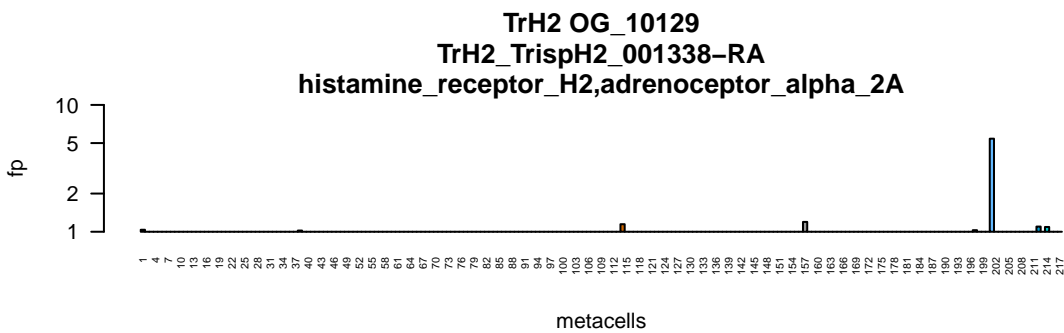
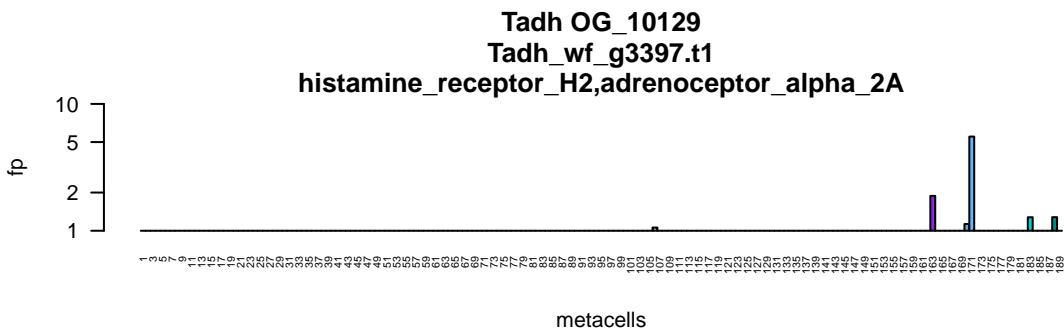
metacells



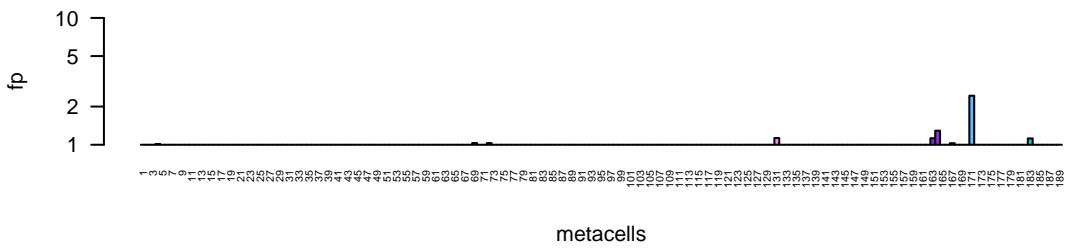




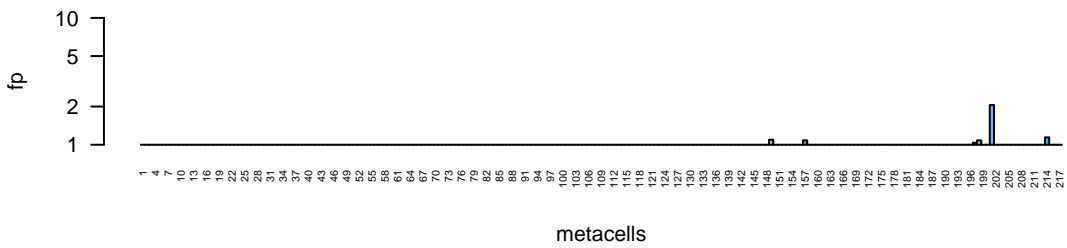




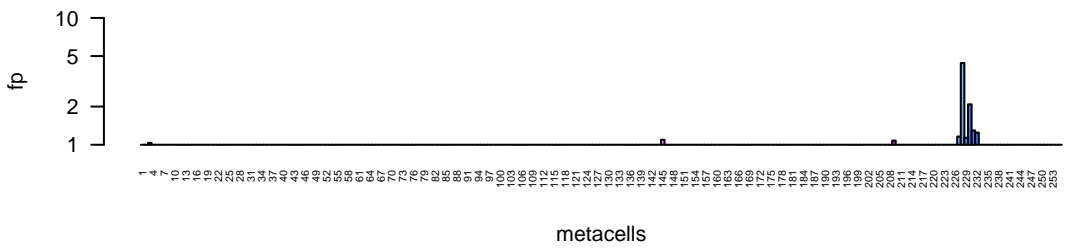
Tadh OG_4120
Tadh_TriadT57620



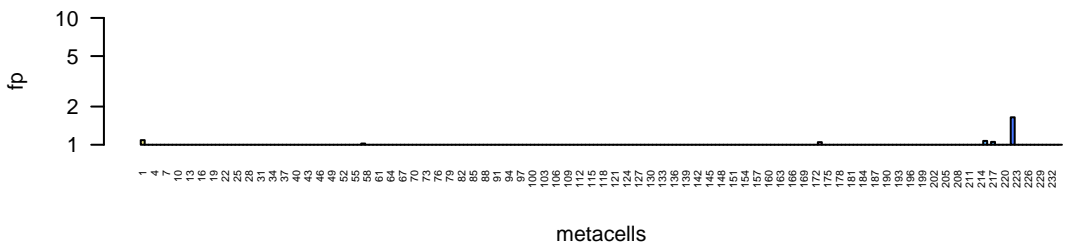
TrH2 OG_4120
TrH2_TrispH2_007435-RA



Hhon OG_4120
Hhon_g03439.t1



HoiH23 OG_4120
HoiH23_PIH23_007154-RA



TadH OG_4214

TrH2 OG_4214

ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

fp

metacells

Tadh OG_4214
Tadh_TriadT30372
 subtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

TrH2 OG_4214
TrH2_TripH2_010333-RA
 subtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

fp

metacells

TadH OG_4214

TrH2 OG_4214

metacells

Two bar charts showing the frequency of metacells for different cell types across two datasets: TadH OG_4214 and Hhon OG_4214. The y-axis is 'fp' (frequency) on a log scale from 1 to 10. The x-axis is 'metacells' with indices from 1 to 253. The left chart is for TadH OG_4214 and the right chart is for Hhon OG_4214. Both charts show a peak in frequency for metacells 223-232, with the right chart showing a much higher peak (fp > 10) for metacell 223.

Figure 1: Frequency of metacells for different cell types across two datasets: TadH OG_4214 and Hhon OG_4214. The y-axis represents frequency (fp) on a log scale from 1 to 10. The x-axis lists metacells from 1 to 253. The left chart is for TadH OG_4214 and the right chart is for Hhon OG_4214. Both charts show a similar distribution of metacells, with a peak at metacell 253.

TrH2 OG_4214
TrH2_TrispH2_010336-RA

ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

fp

metacells

Hhon OG_4214
Hhon_g06800.t1

ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

fp

metacells

ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

metacell	fp
1	1
4	1
10	1
13	1
16	1
22	1
25	1
31	1
34	1
37	1
43	1
46	1
55	1
58	1
59	1
64	1
67	1
73	1
76	1
79	1
85	1
88	1
91	1
97	1
100	1
103	1
106	1
109	1
112	1
118	1
121	1
127	1
130	1
133	1
139	1
142	1
151	1
154	1
160	1
163	1
169	1
172	1
175	1
181	1
184	1
187	1
190	1
193	1
196	1
202	1
205	1
209	1
214	1
217	1
223	2
226	1
229	1
232	3

ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

fp

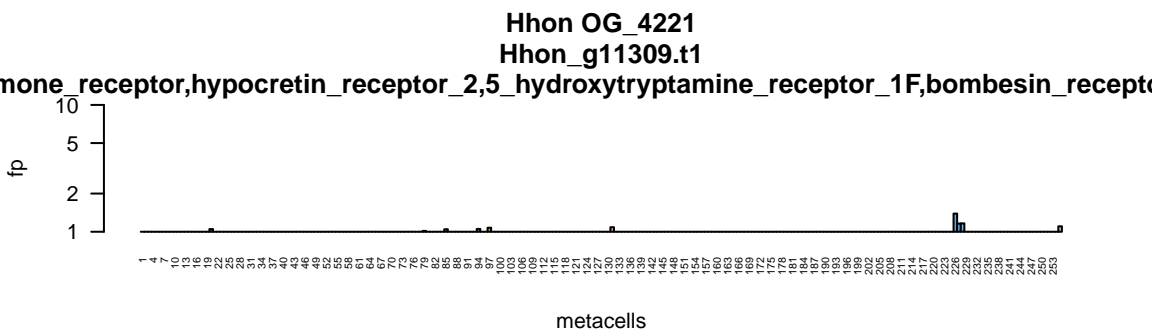
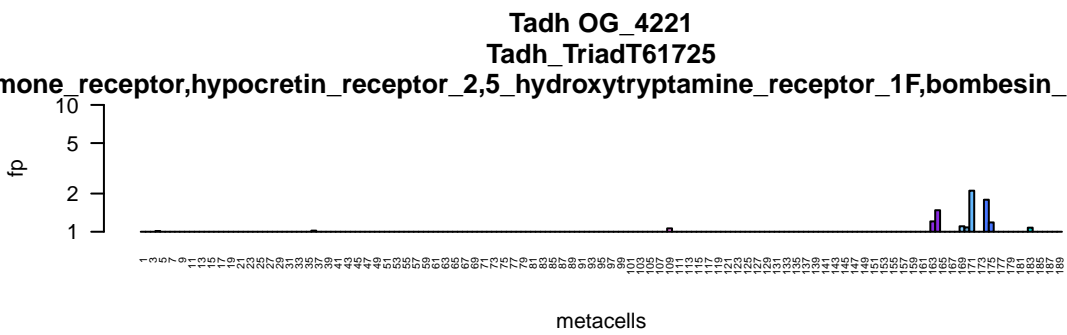
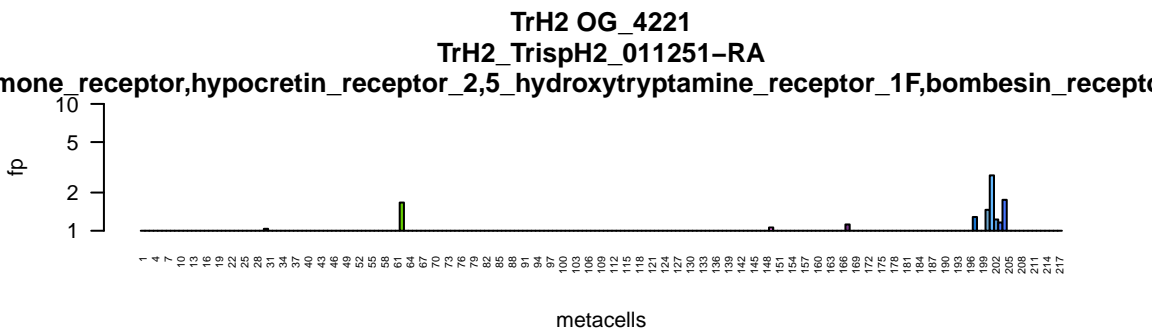
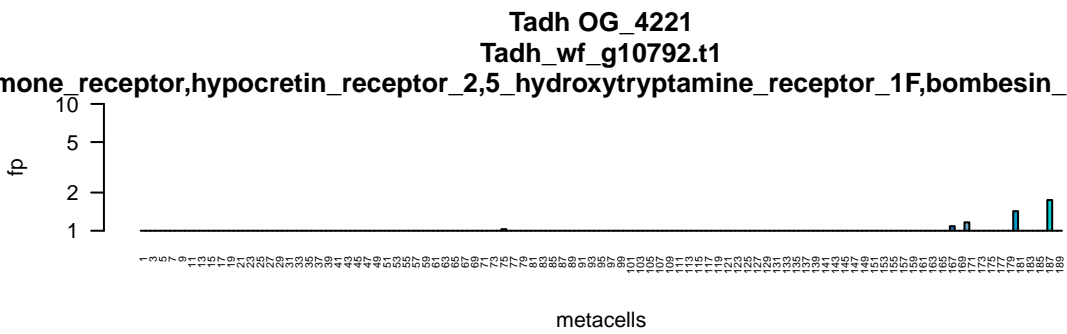
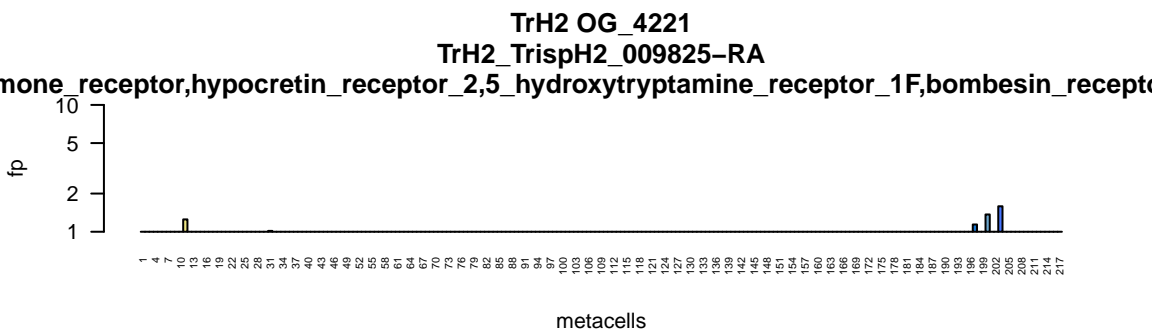
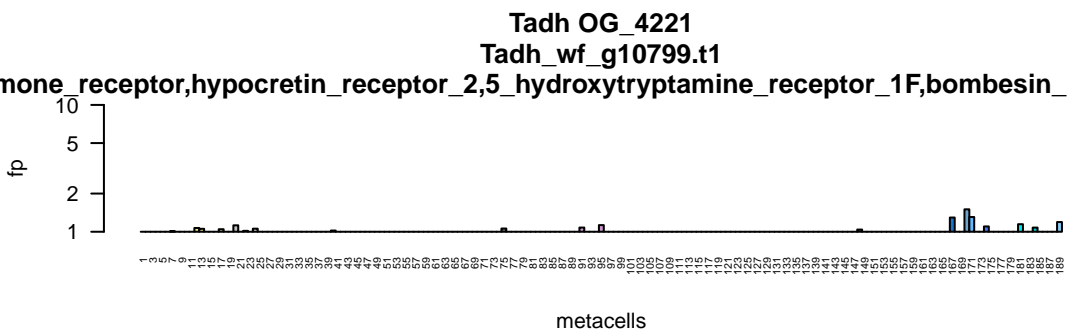
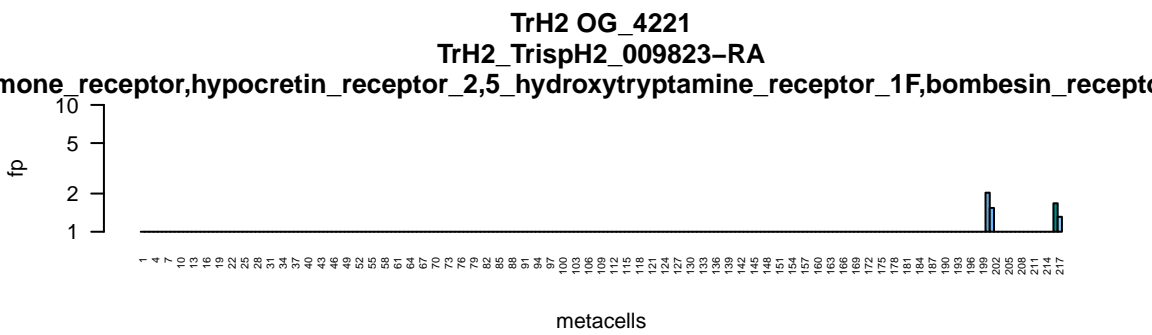
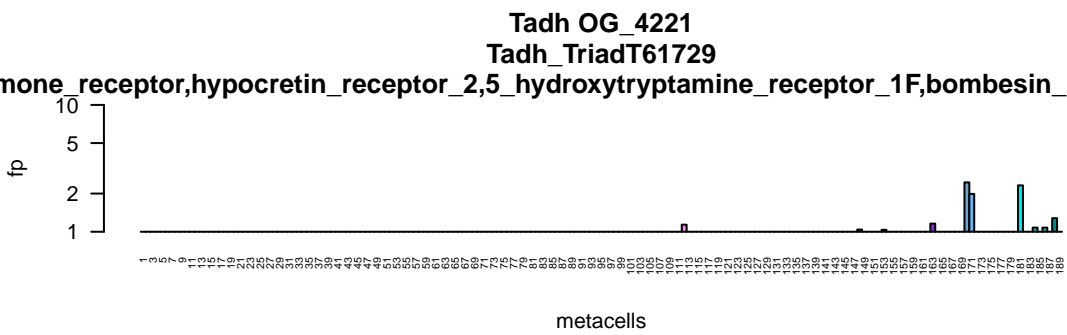
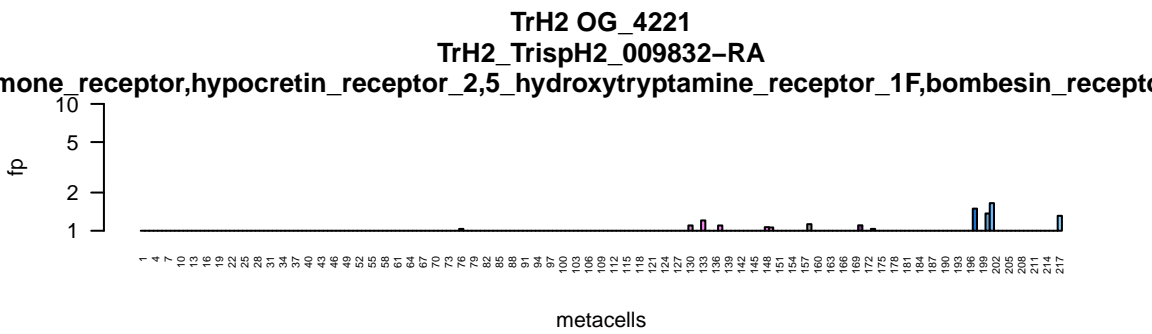
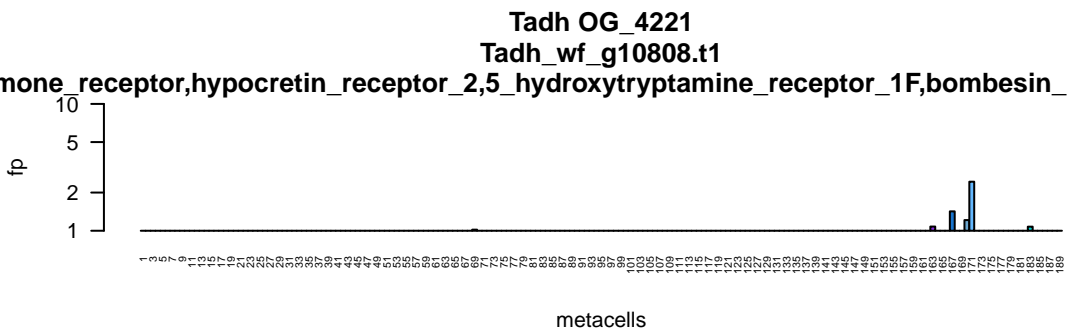
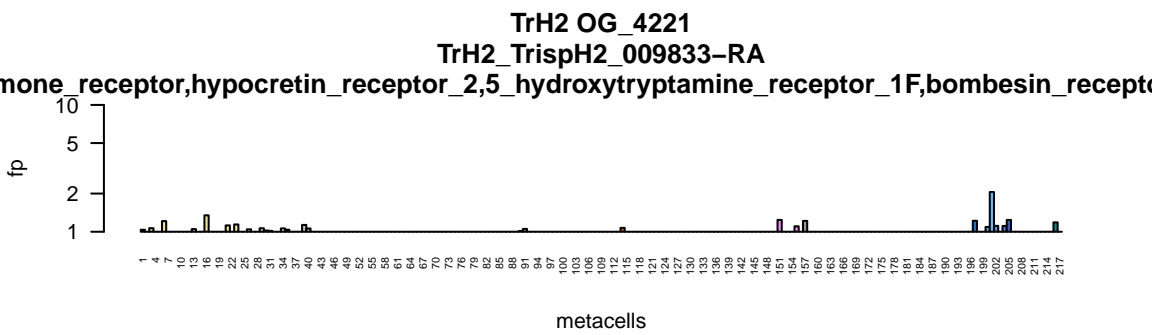
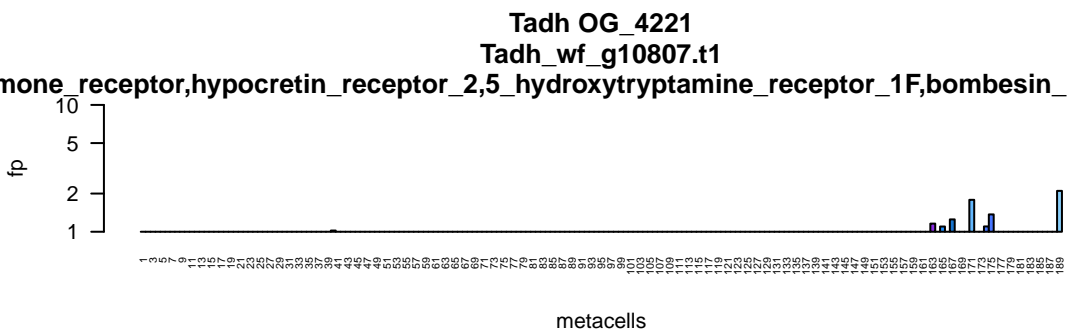
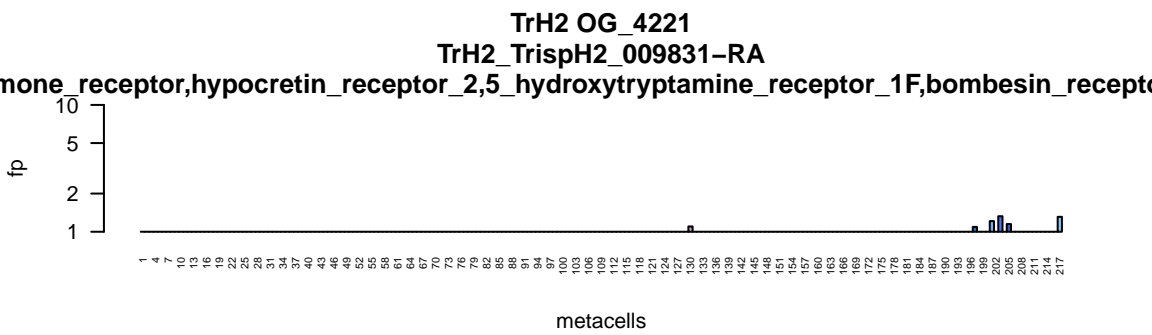
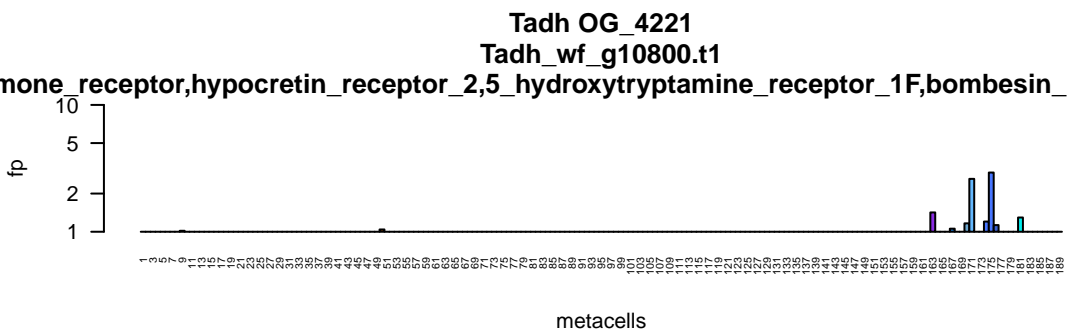
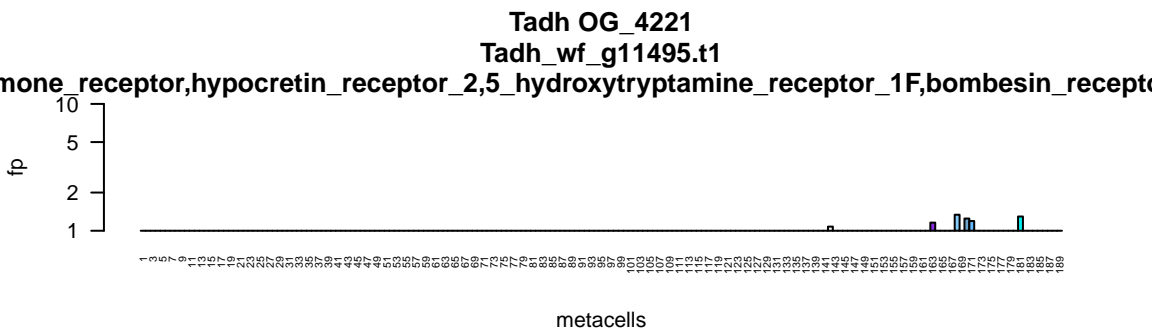
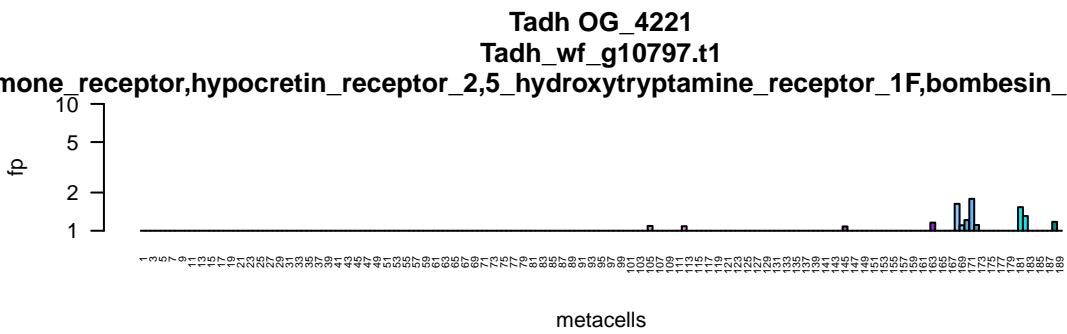
metacells

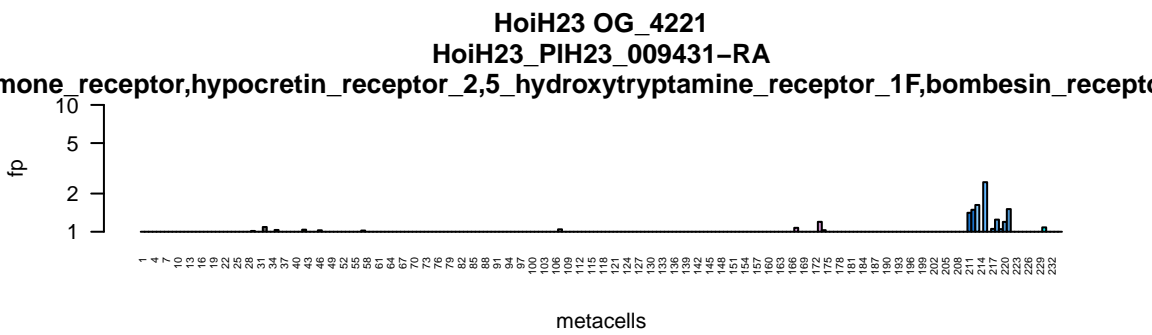
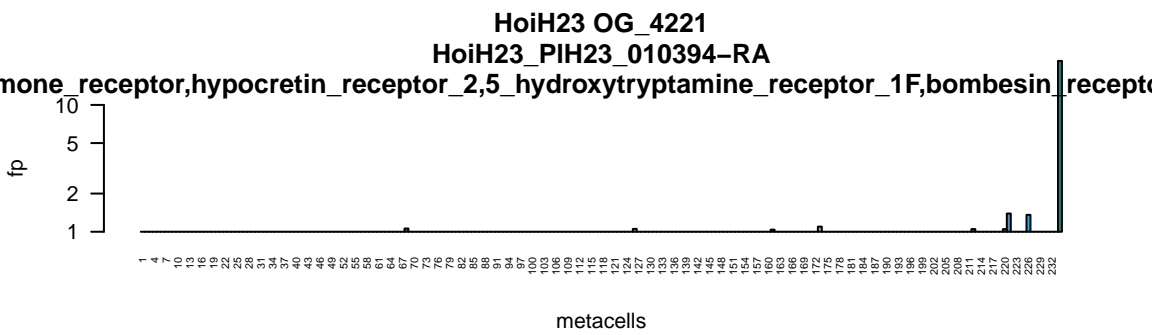
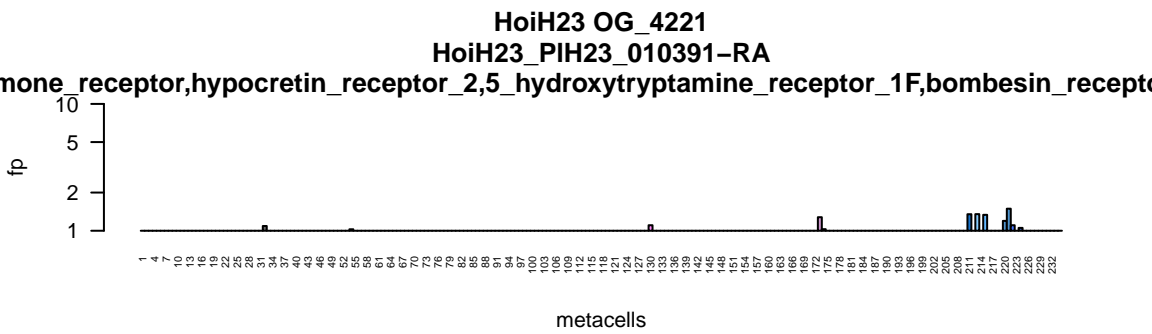
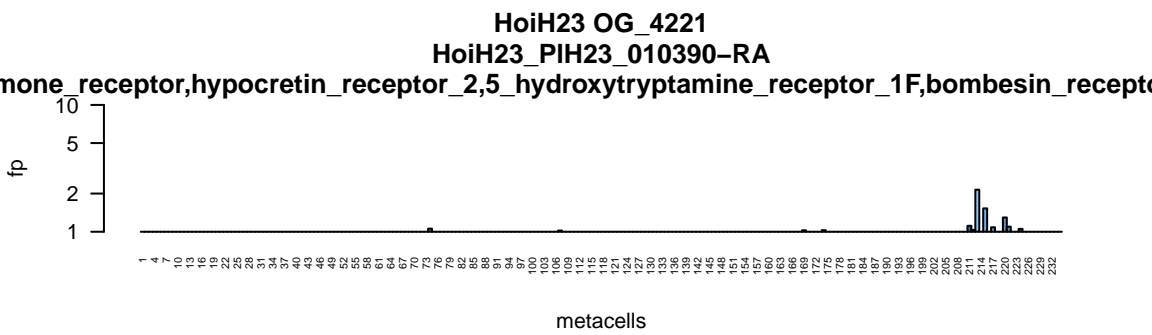
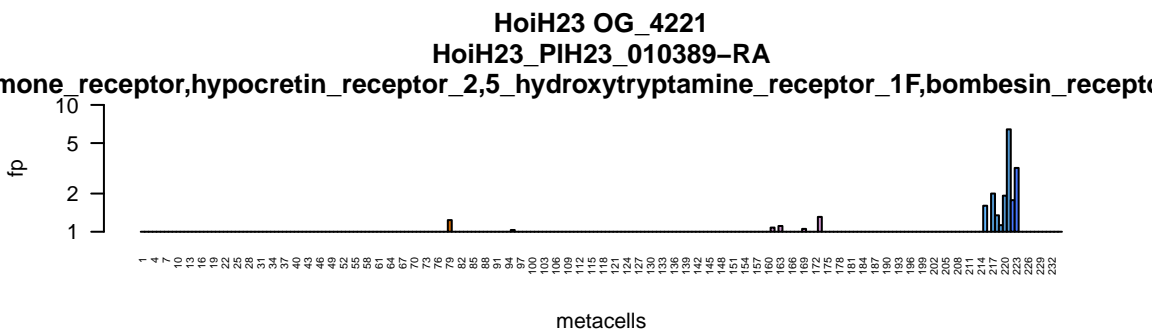
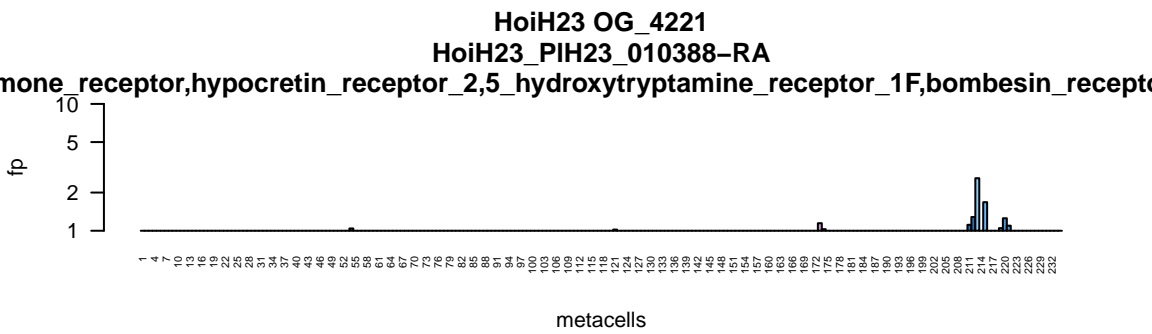
HoiH23 OG_4214
HoiH23_PIH23_009309-RA

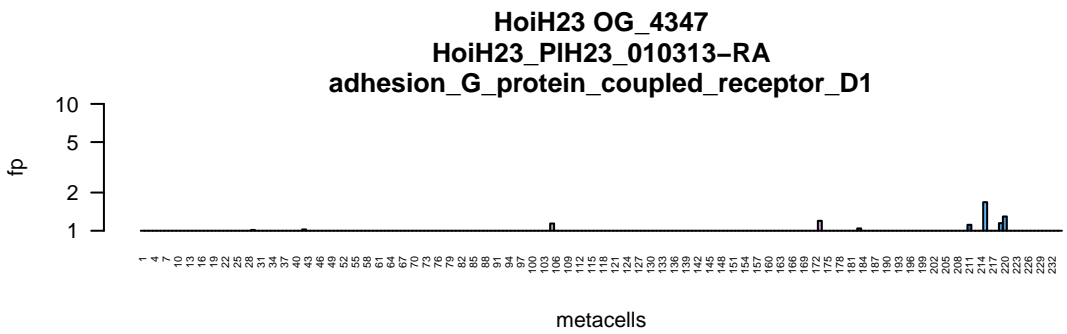
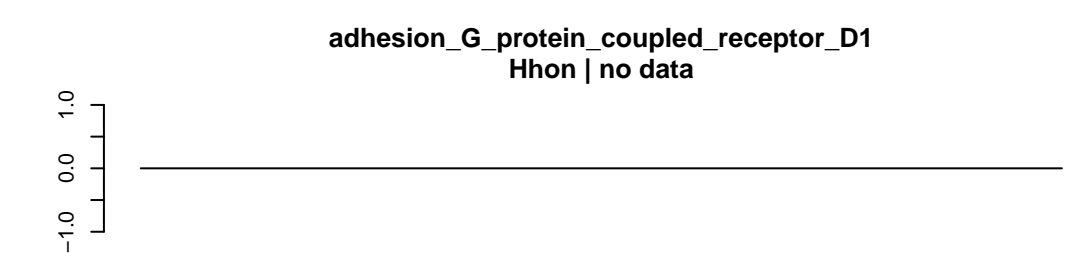
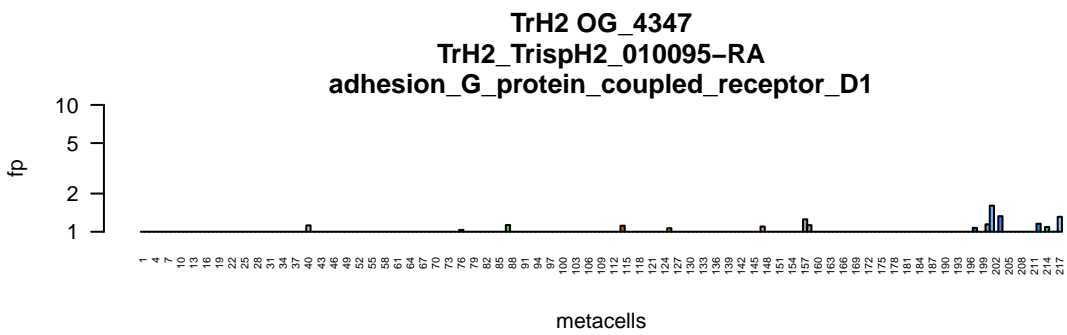
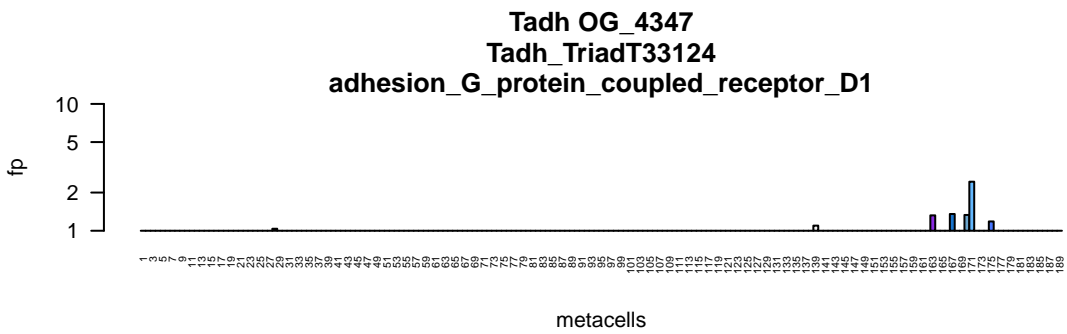
ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

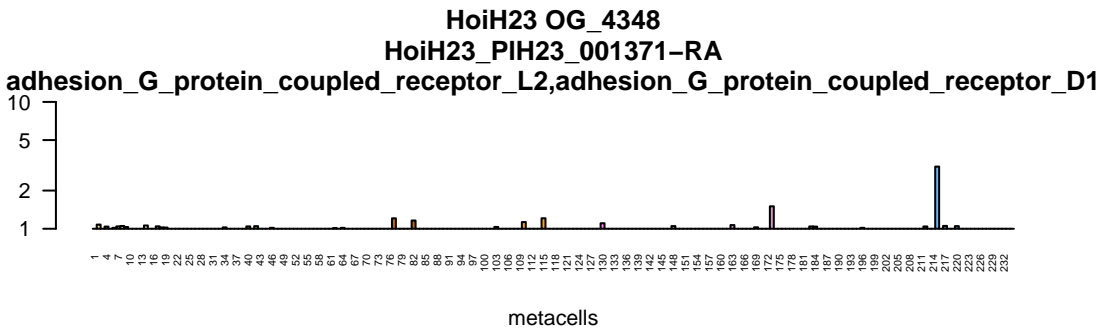
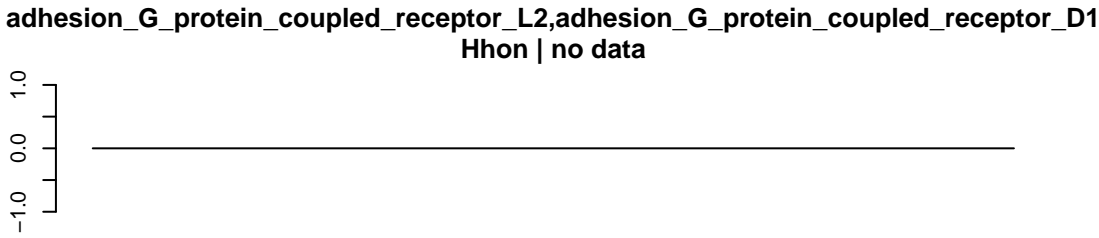
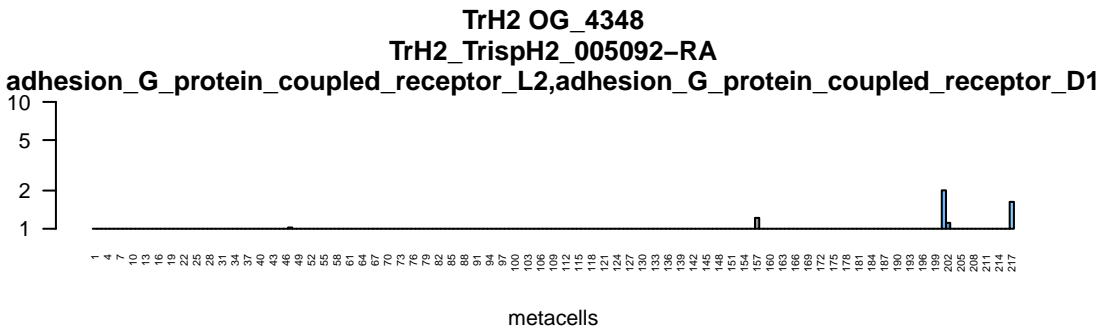
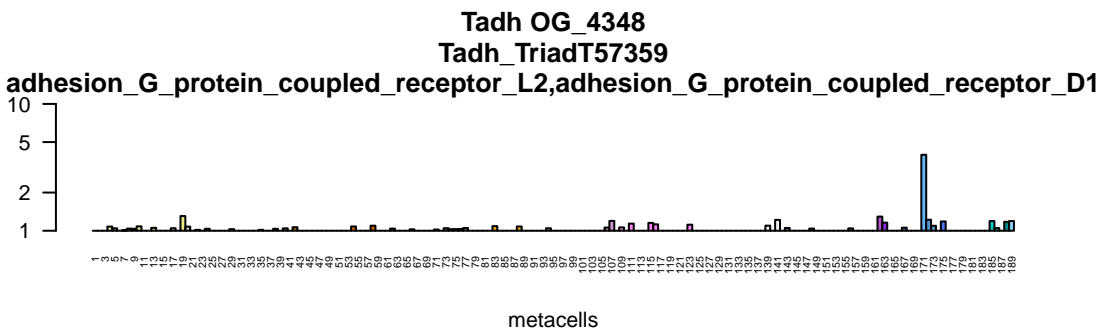
fp

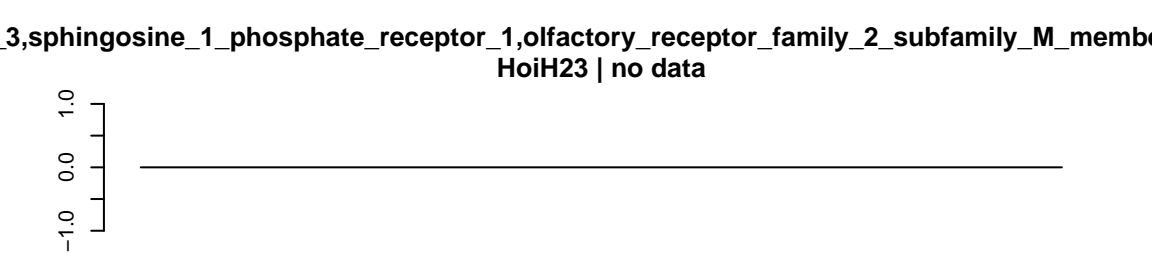
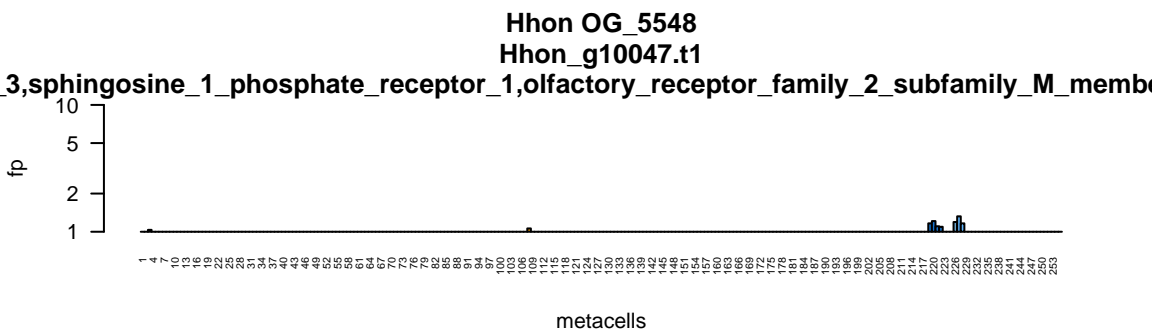
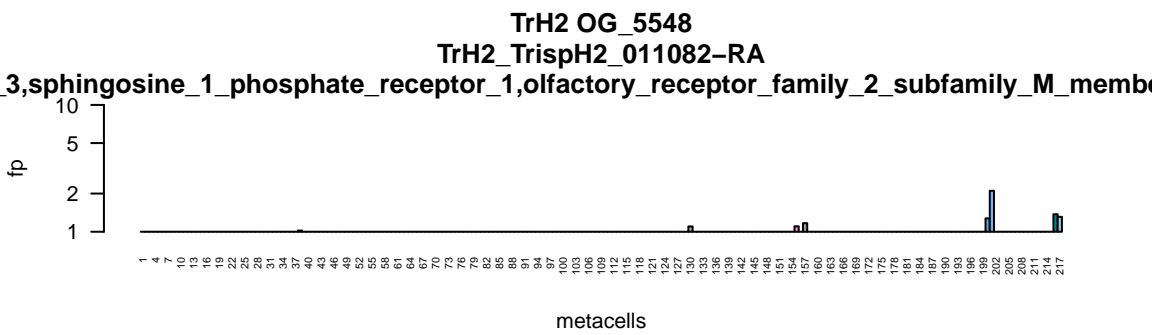
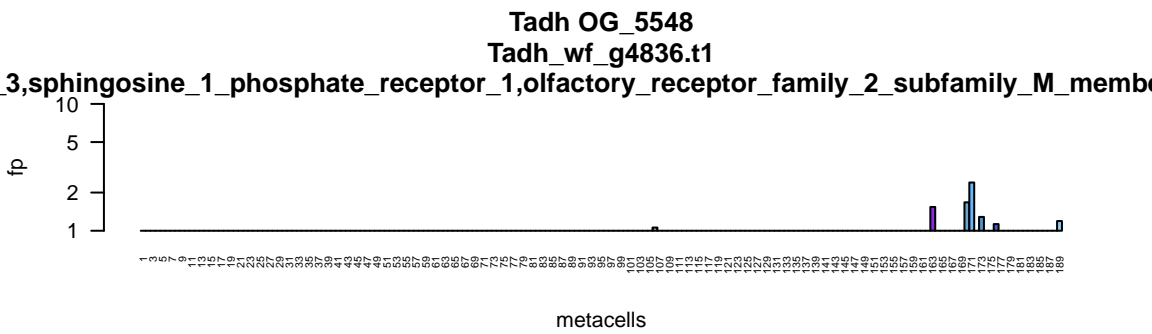
metacells

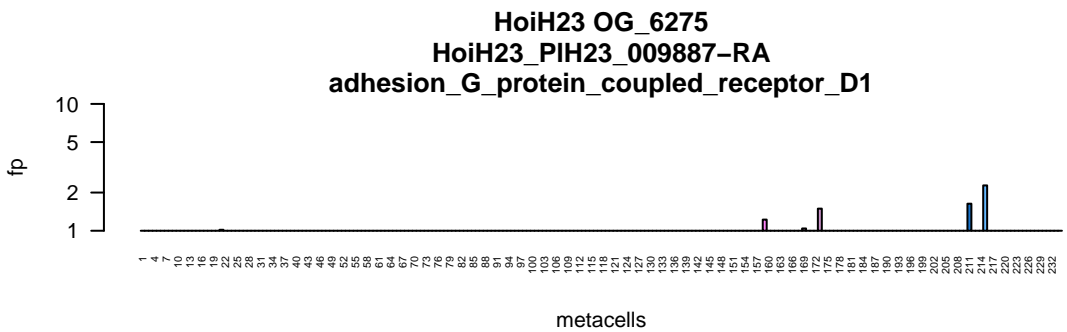
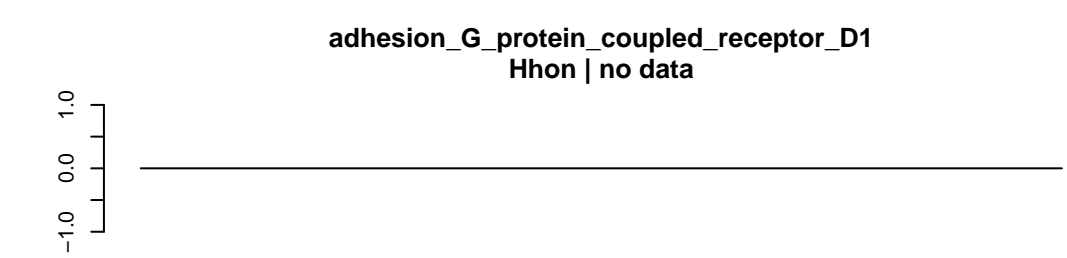
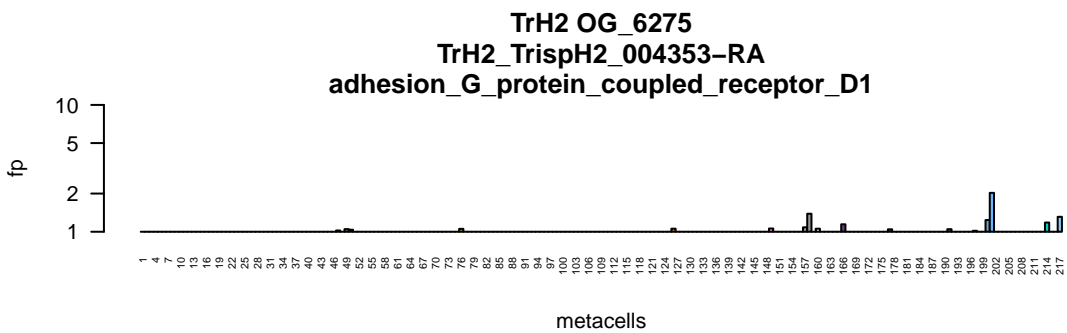
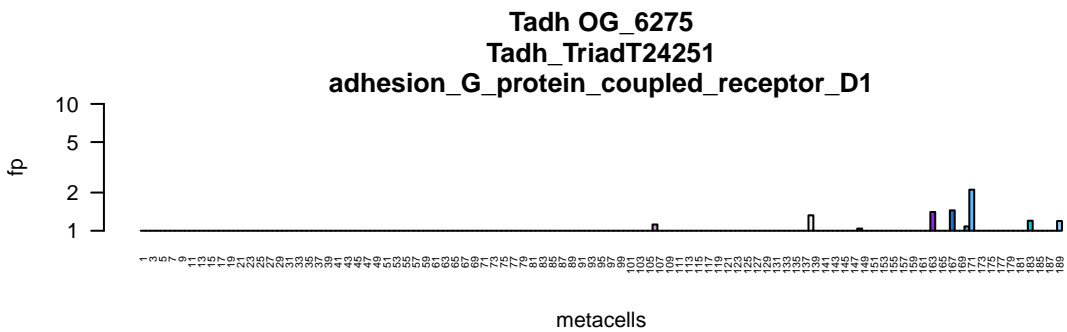


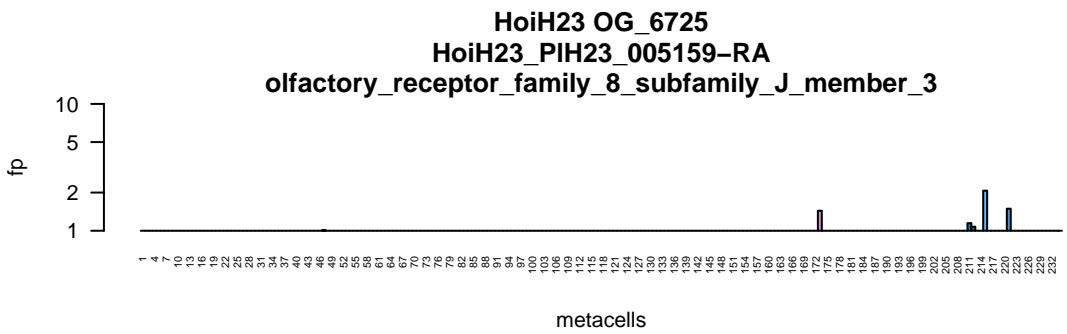
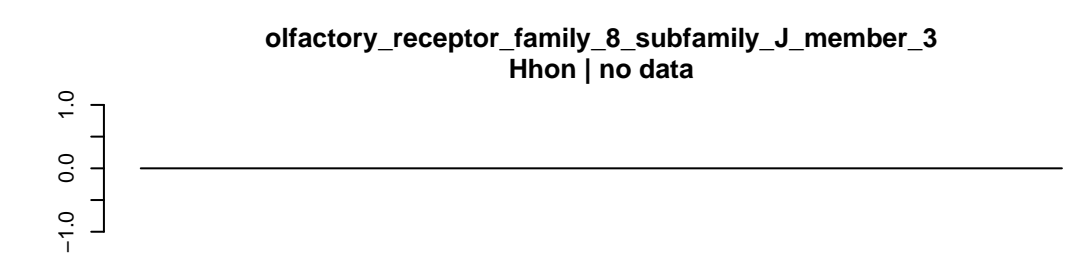
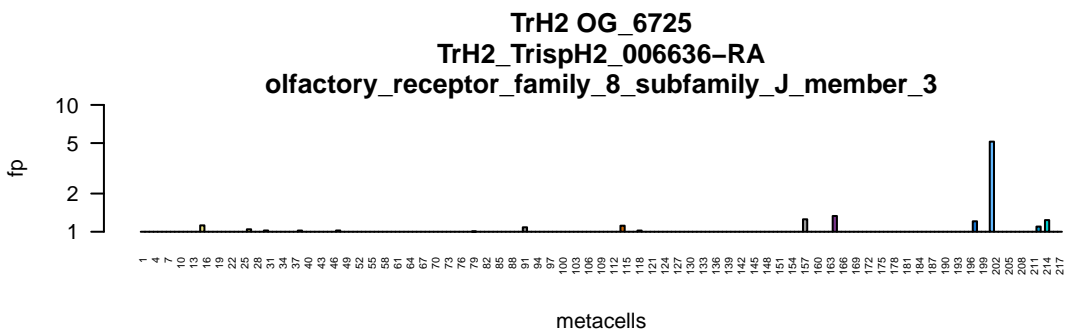
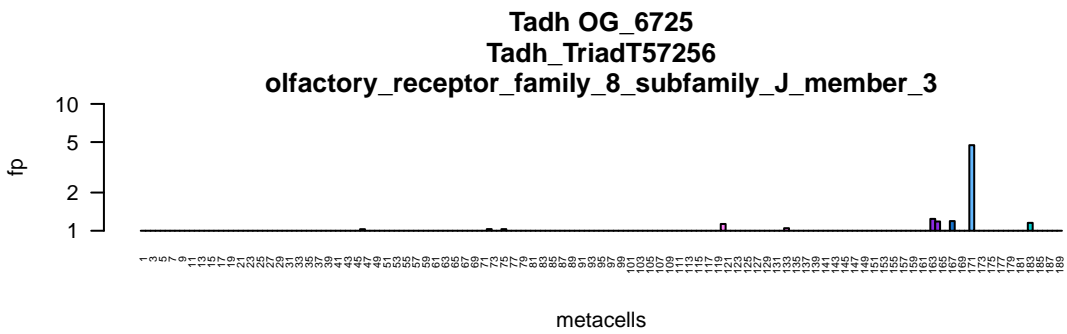


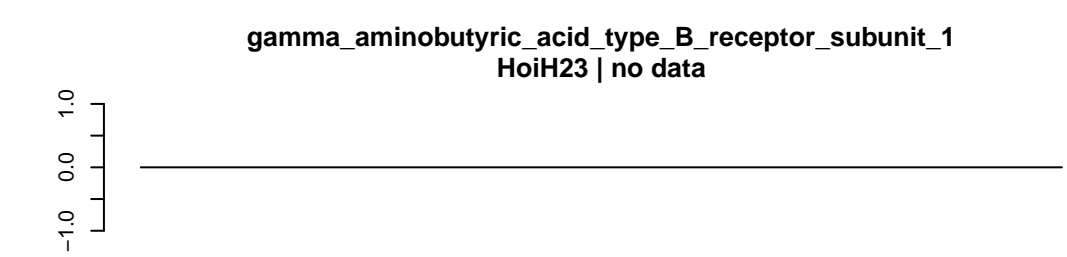
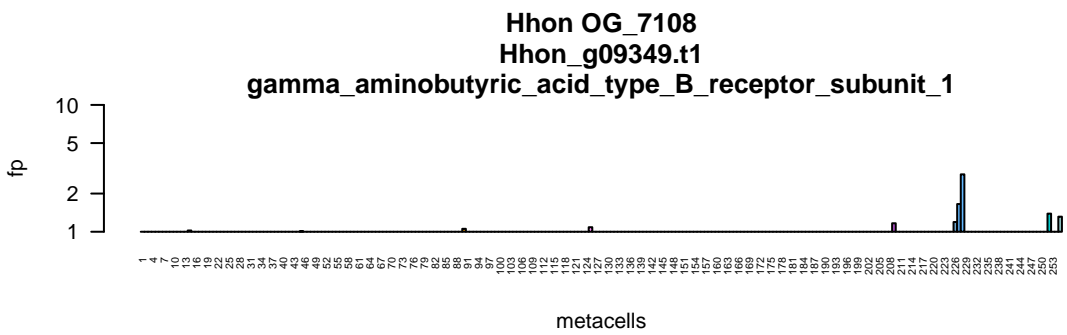
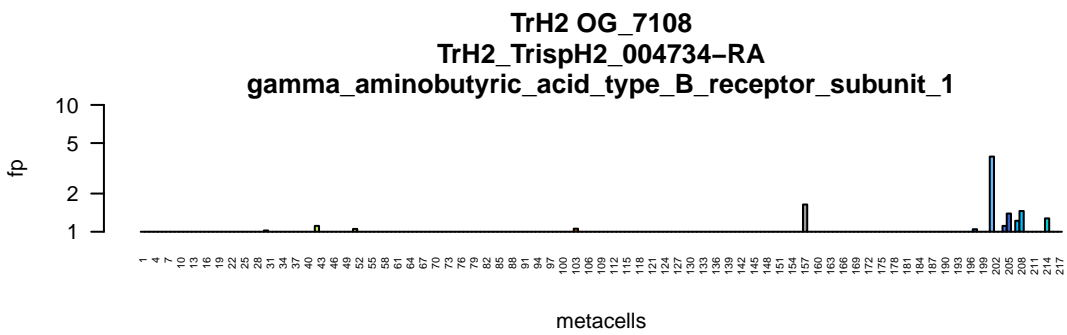
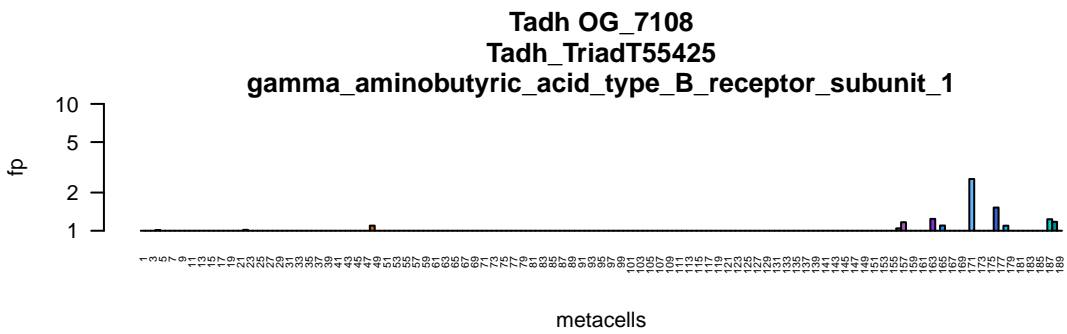


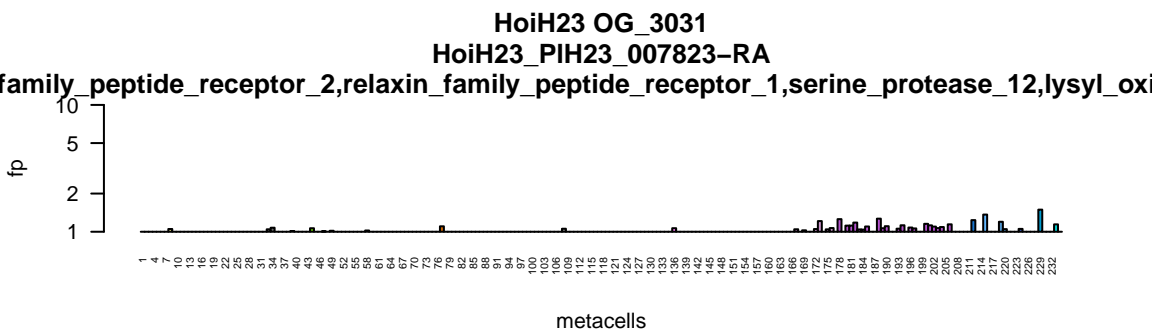
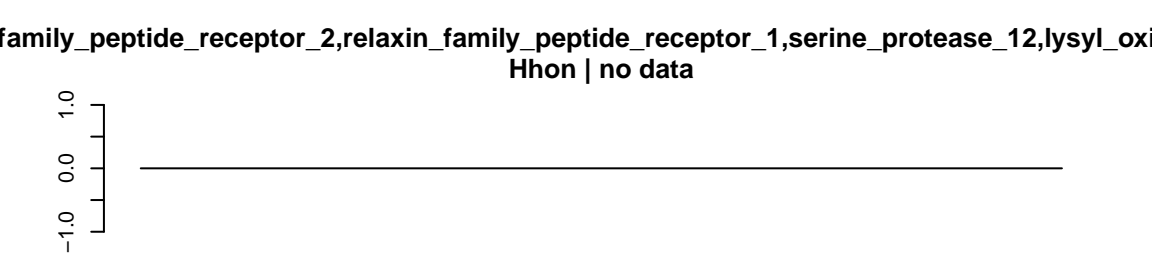
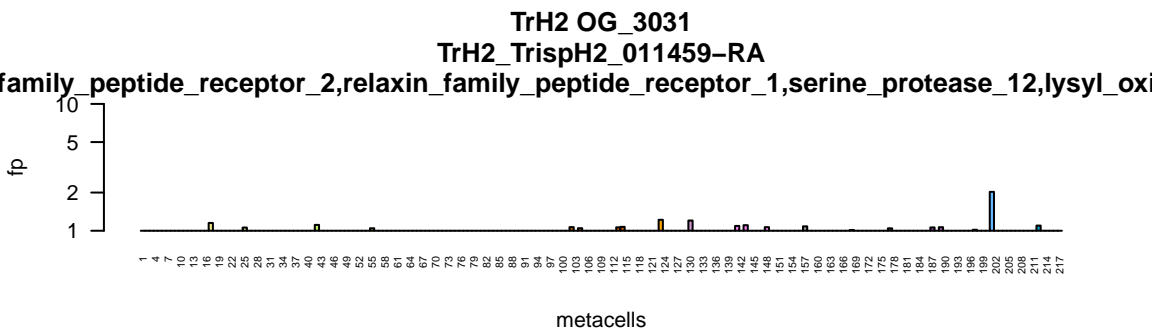
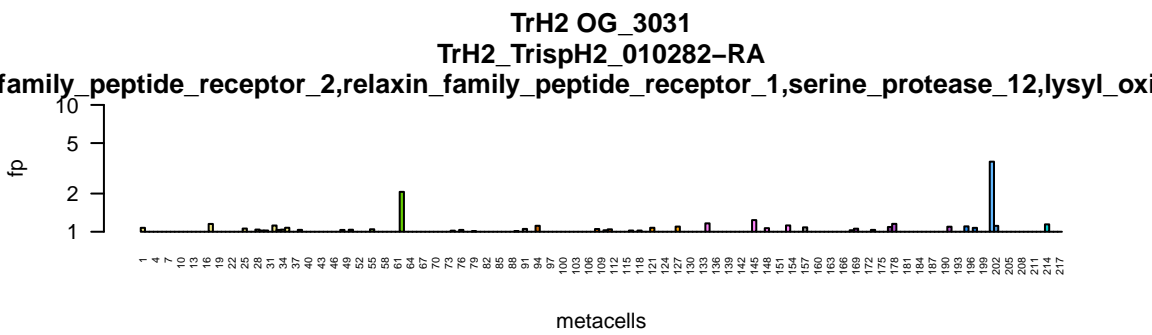
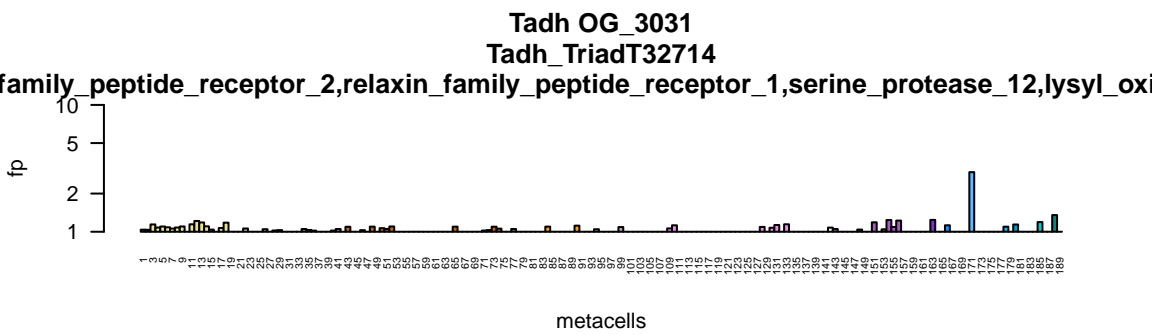
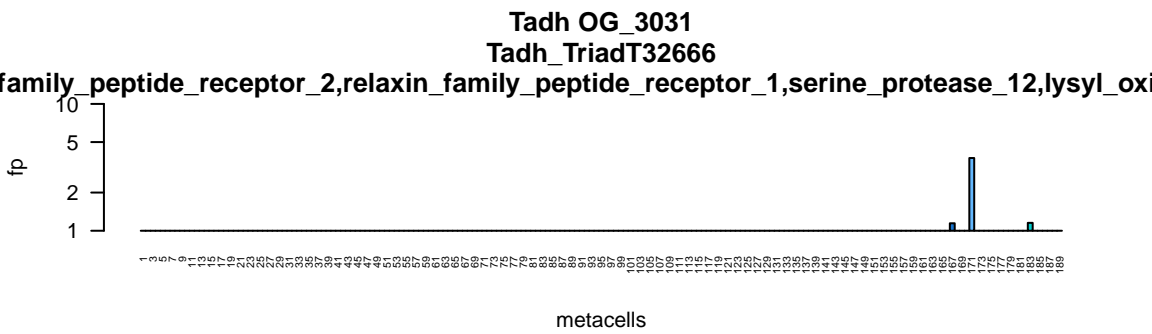


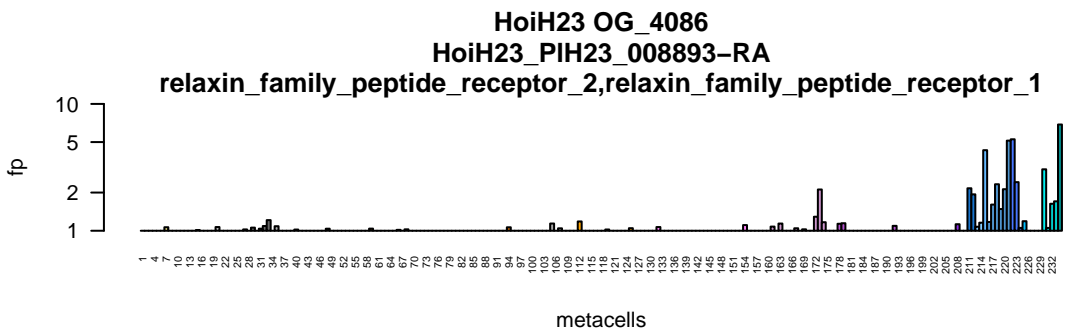
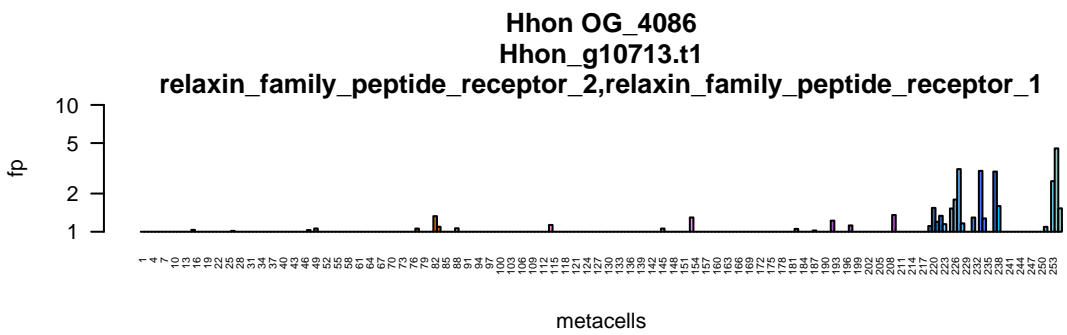
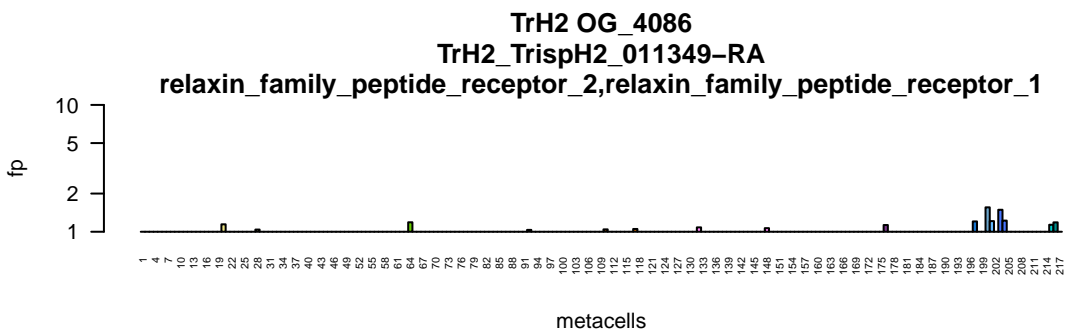
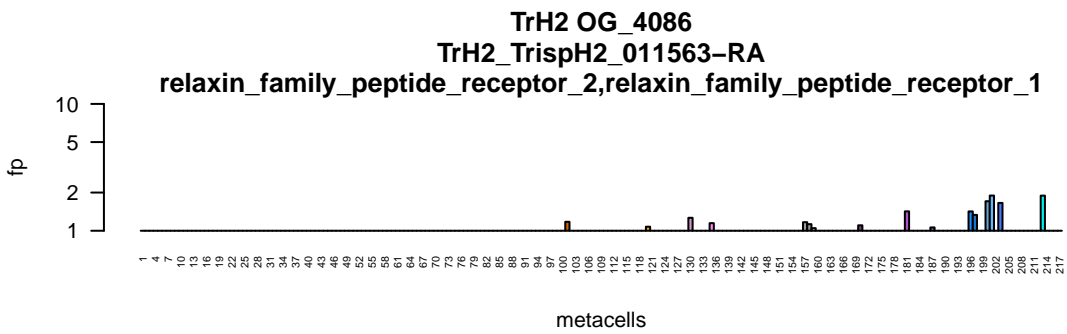
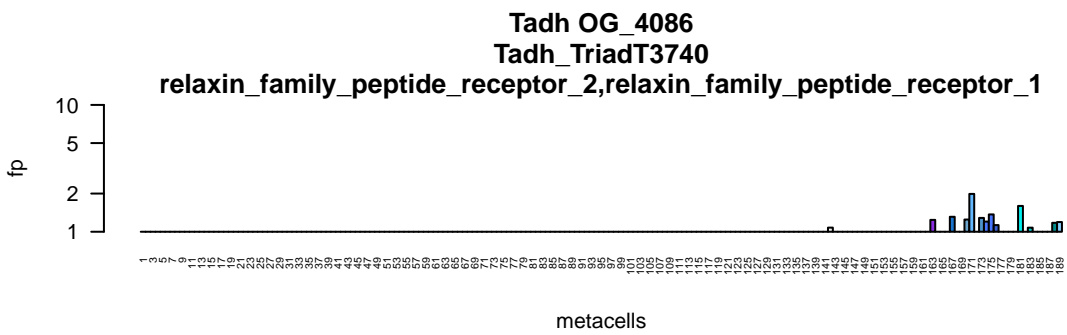
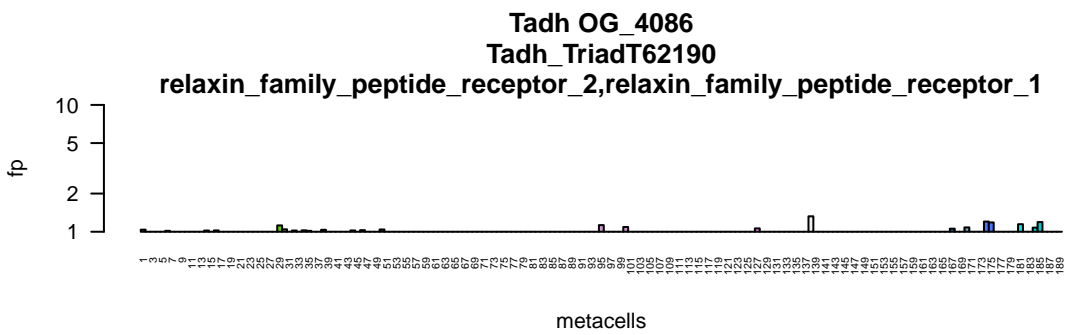
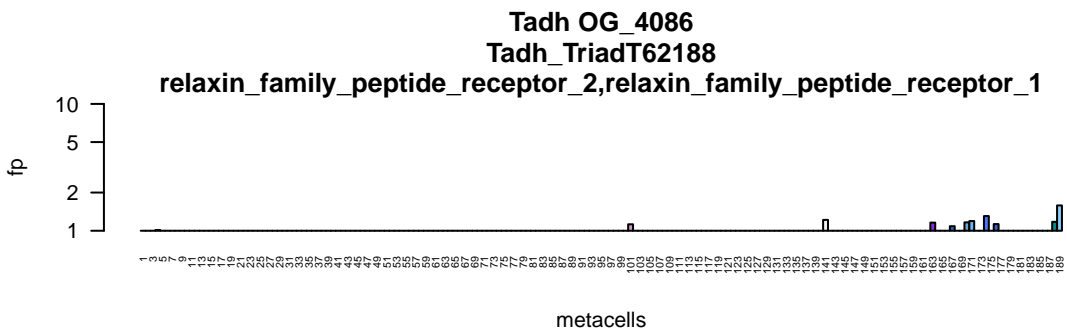


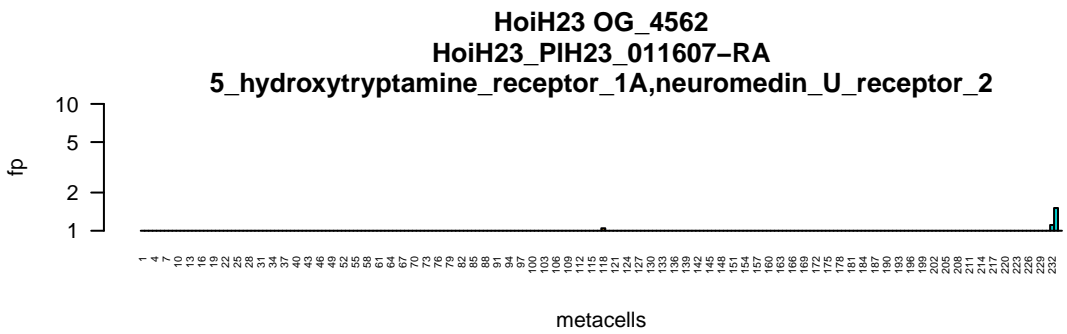
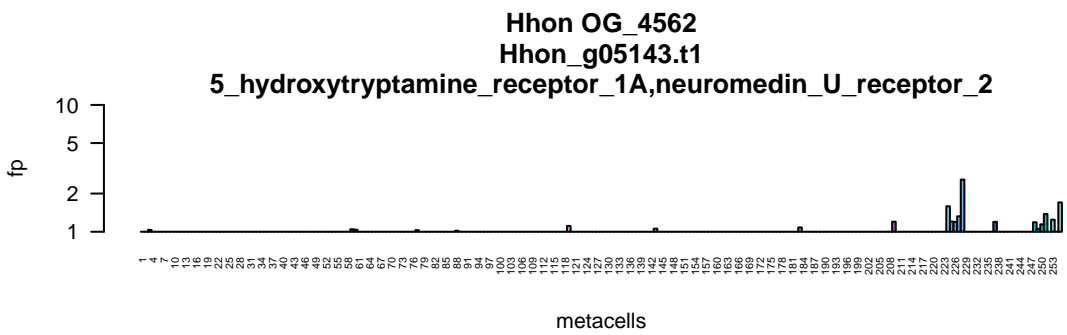
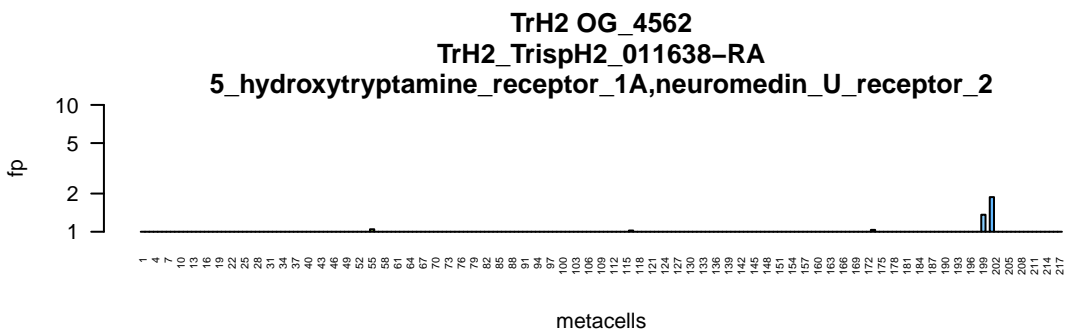
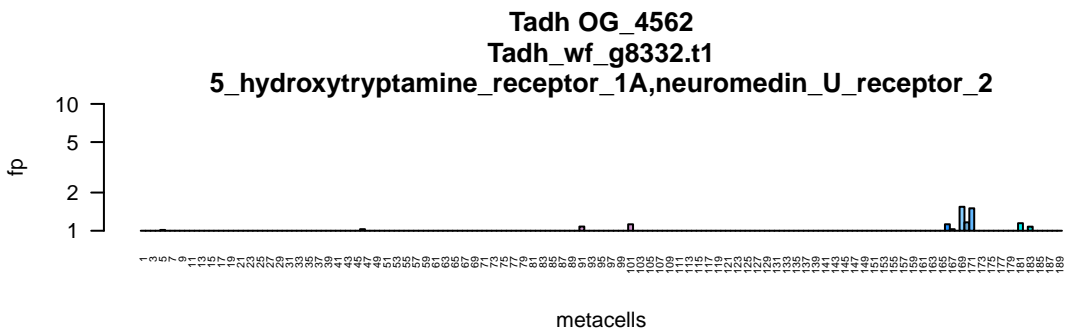


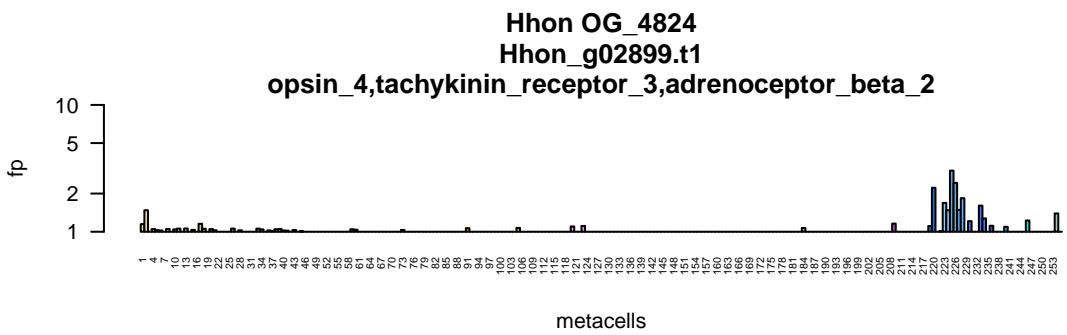
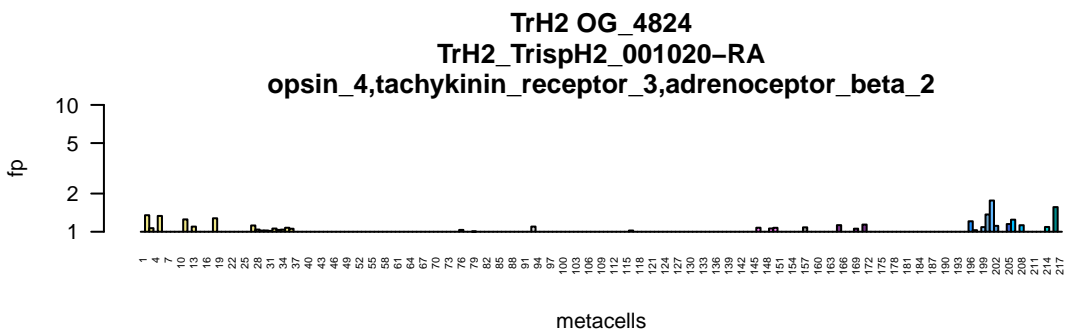
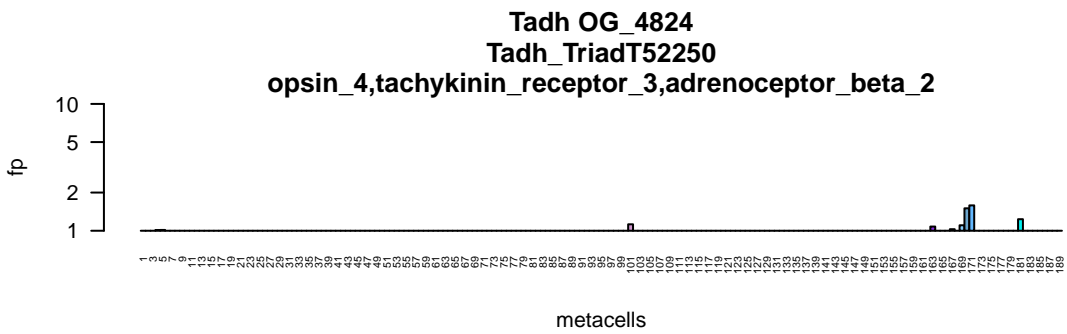


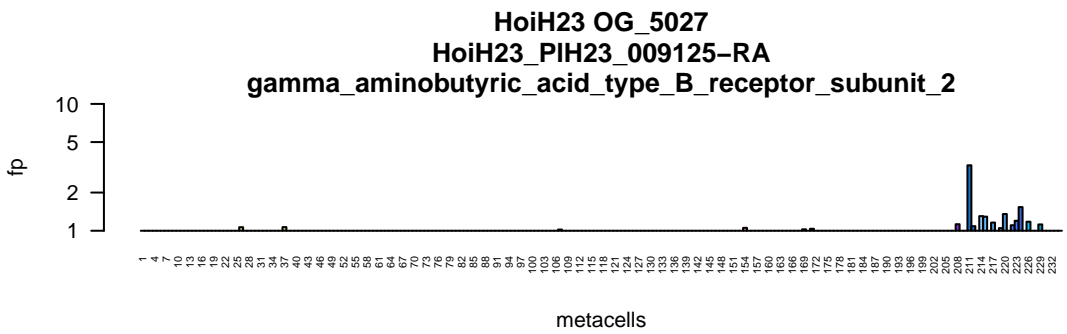
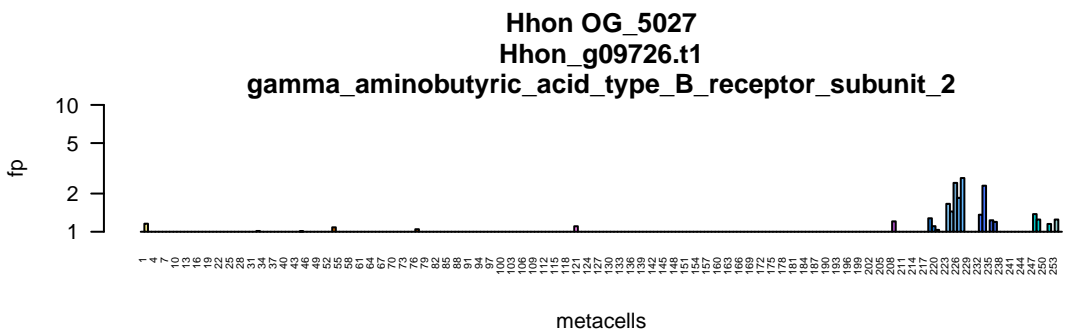
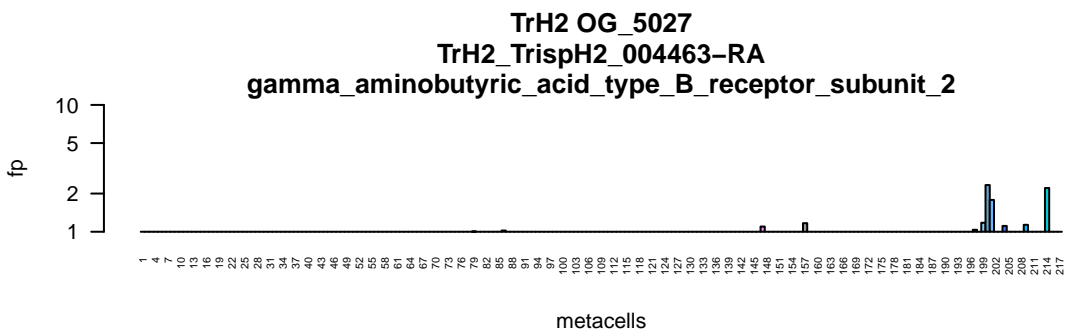
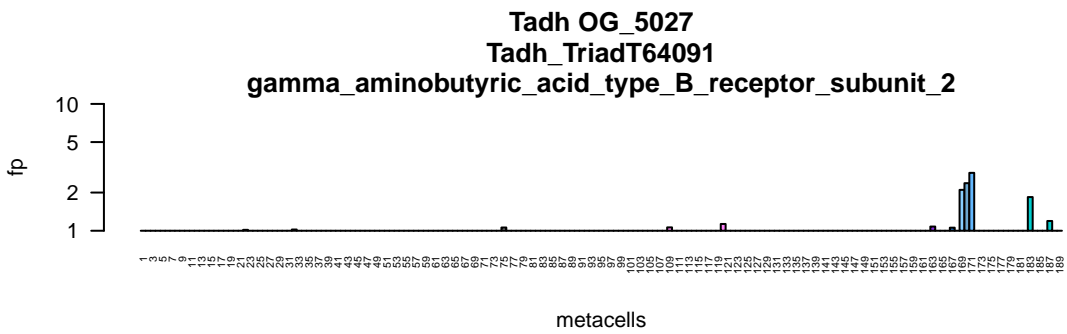


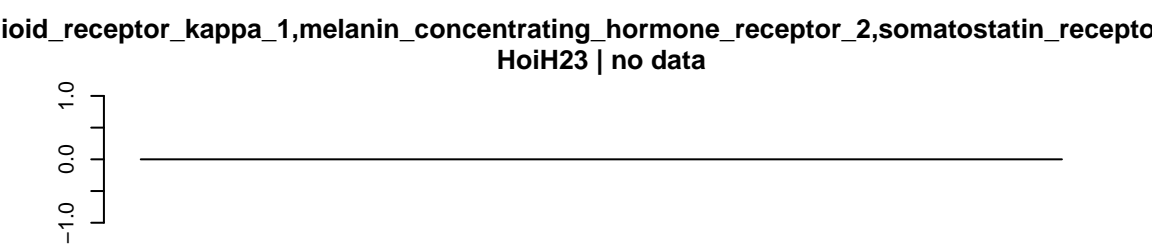
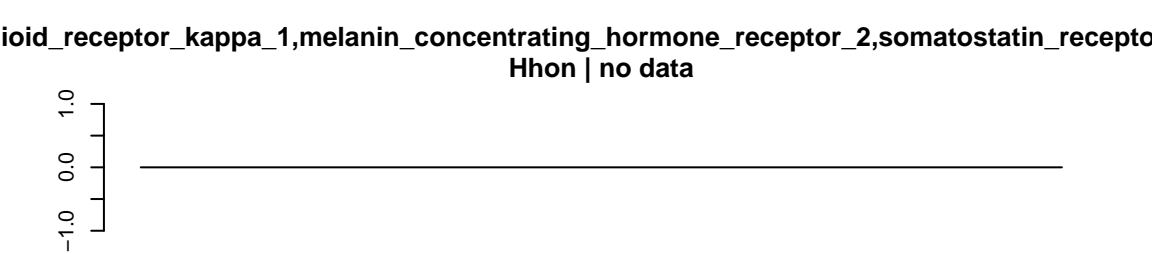
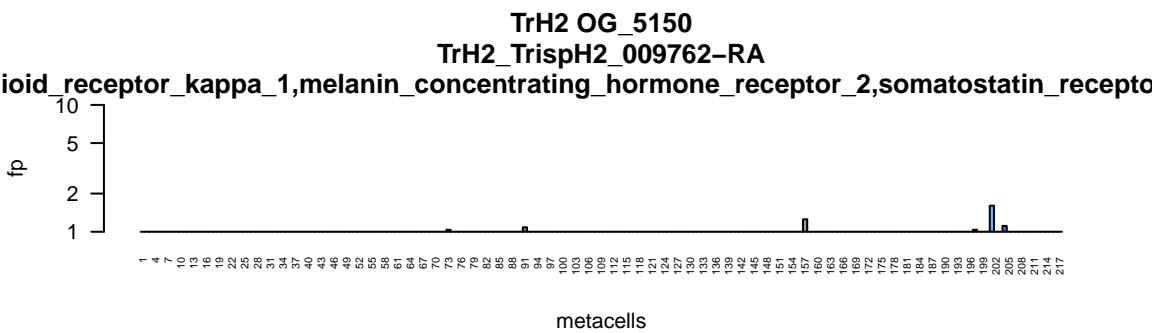
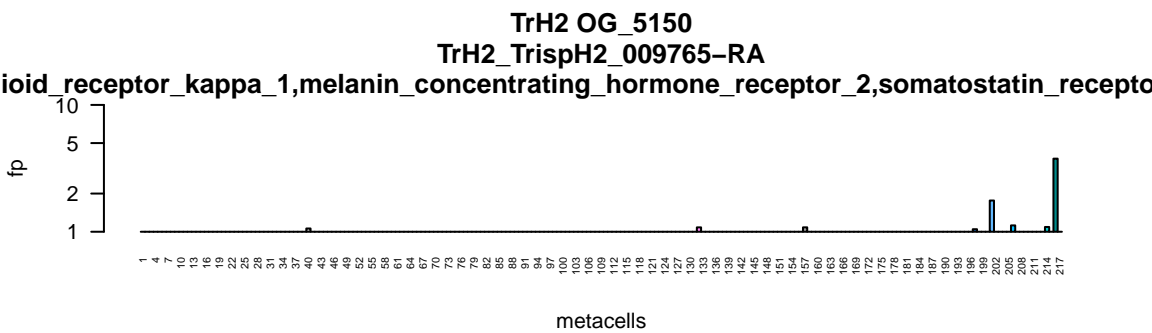
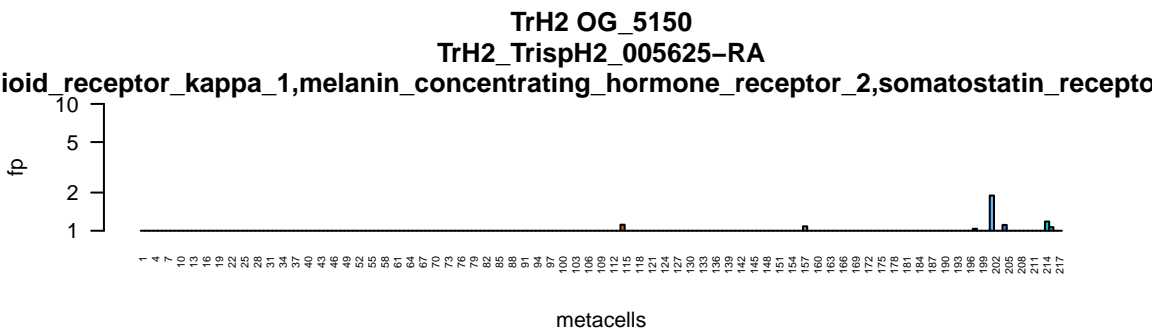
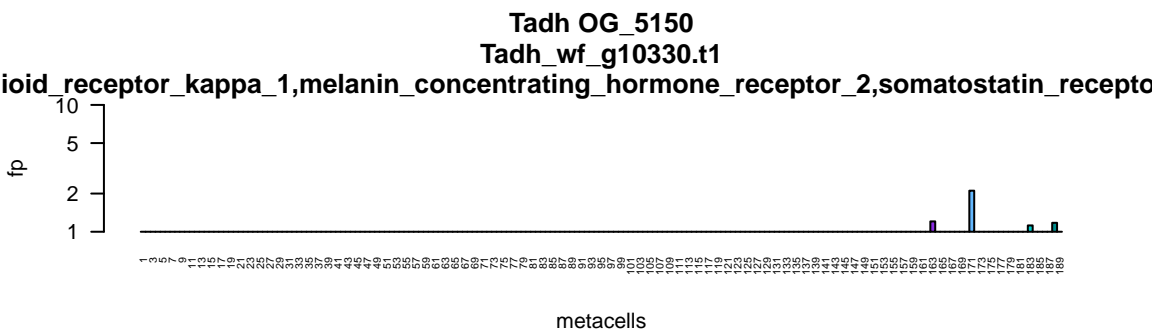
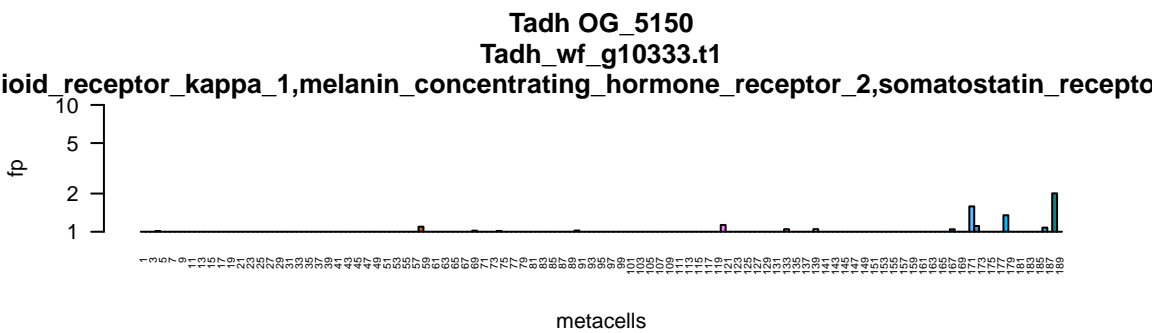
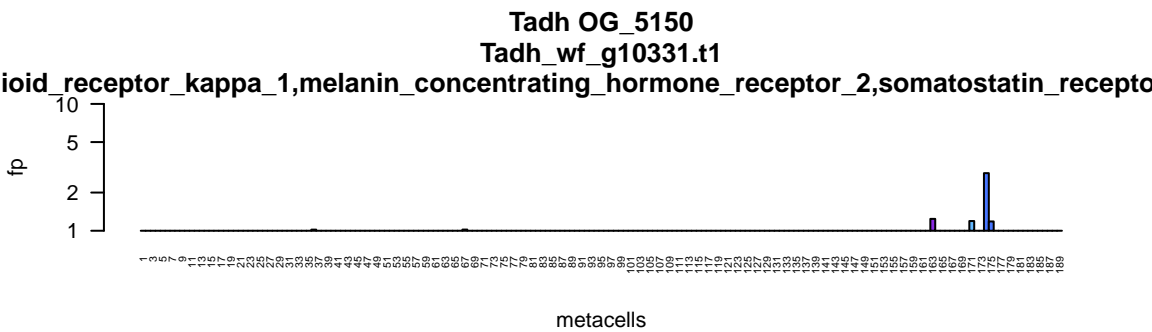


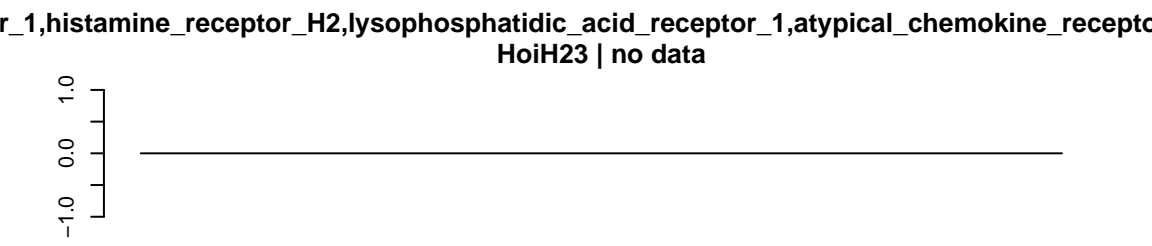
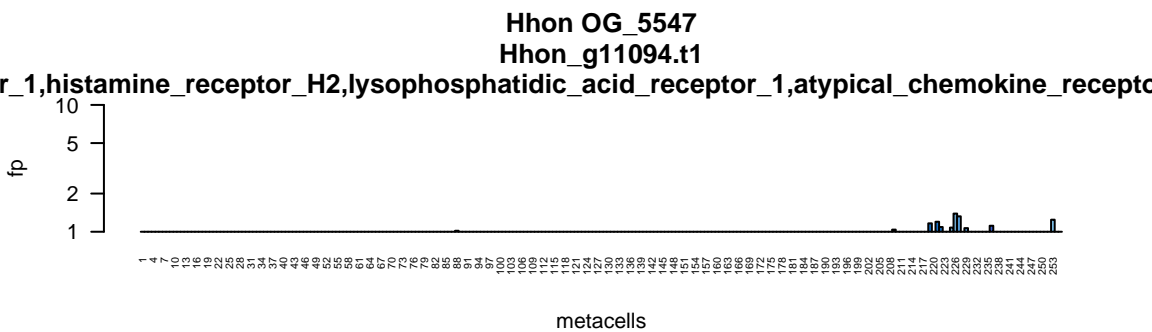
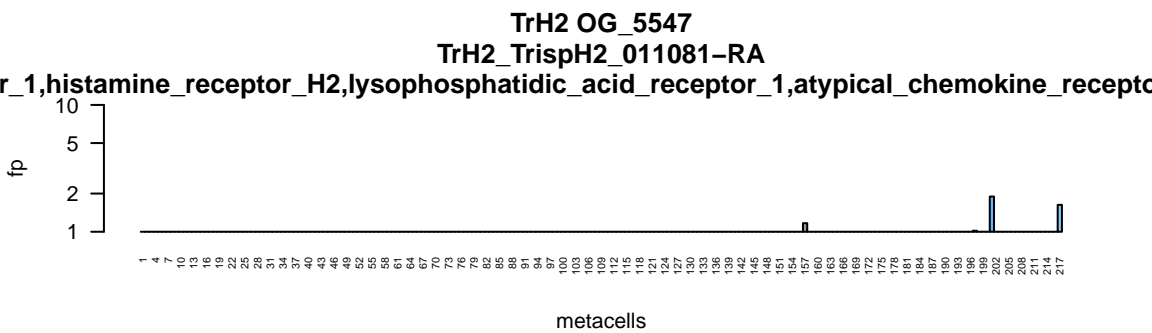
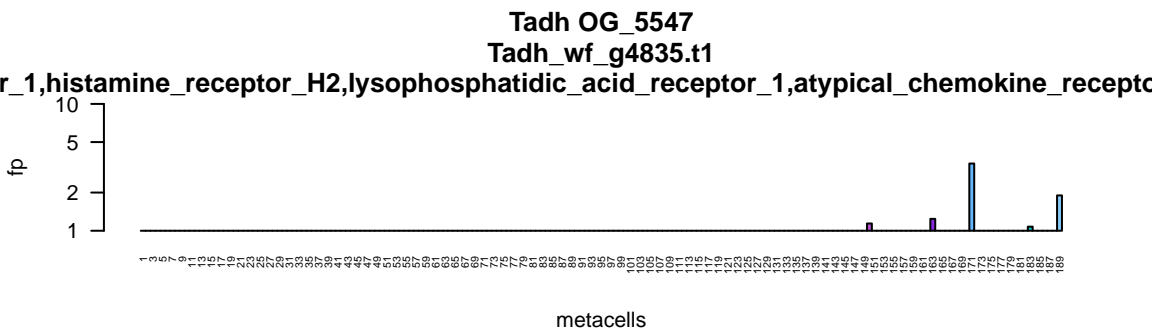








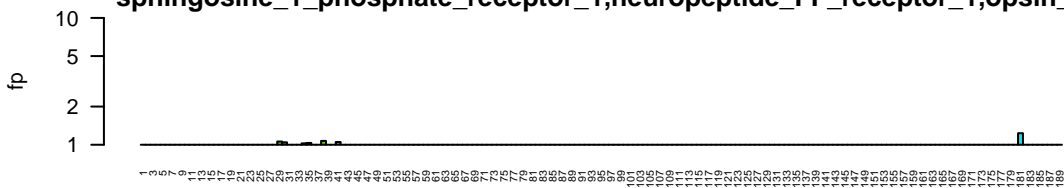




Tadh OG 5789

Tadh_TriadT3838

sphingosine_1_phosphate_receptor_1,neuropeptide_FF_receptor_1,opsin_3

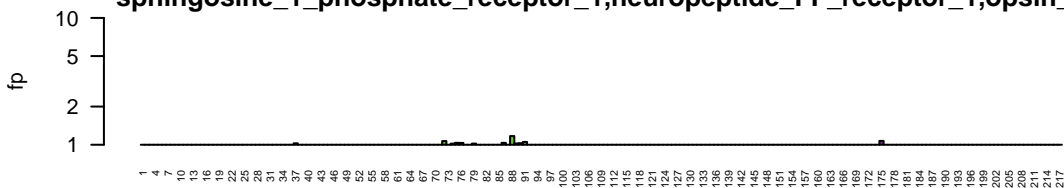


metacells

TrH2 OG_5789

TrH2_TrispH2_011979-RA

sphingosine_1_phosphate_receptor_1,neuropeptide_FF_receptor_1,opsin_3

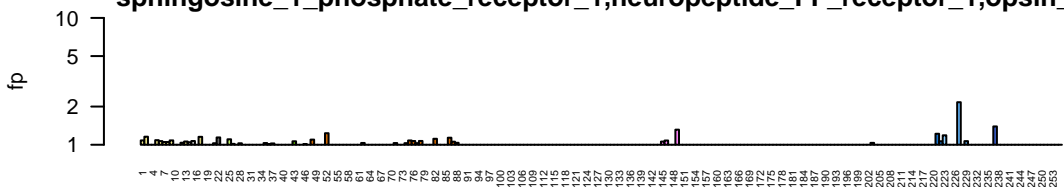


metacells

Hhon OG_5789

Hhon_g11091.t1

sphingosine_1_phosphate_receptor_1,neuropeptide_FF_receptor_1,opsin_3

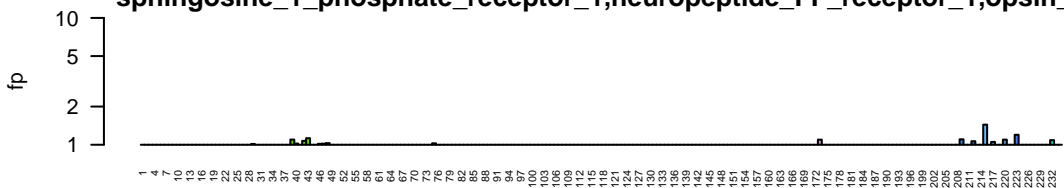


metacells

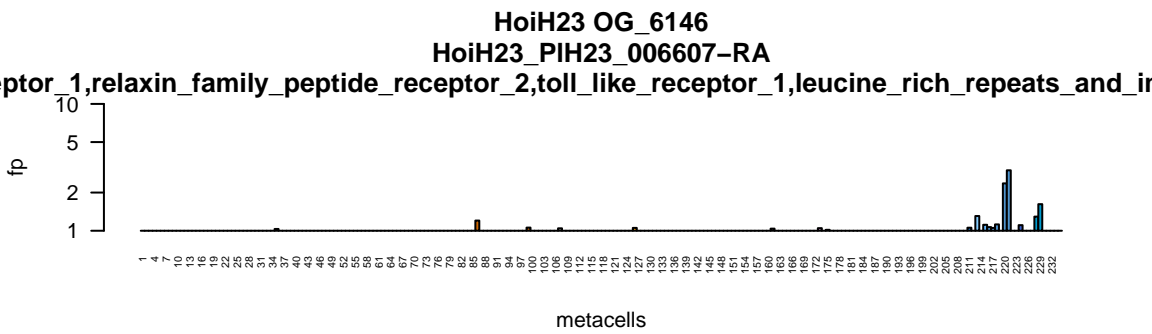
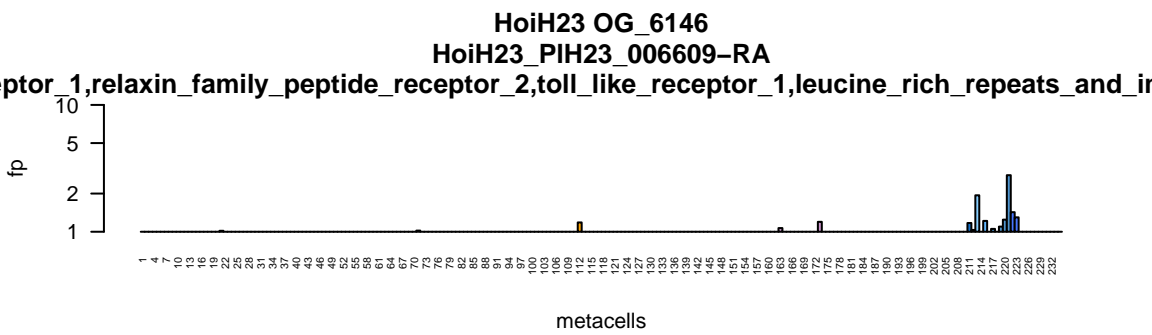
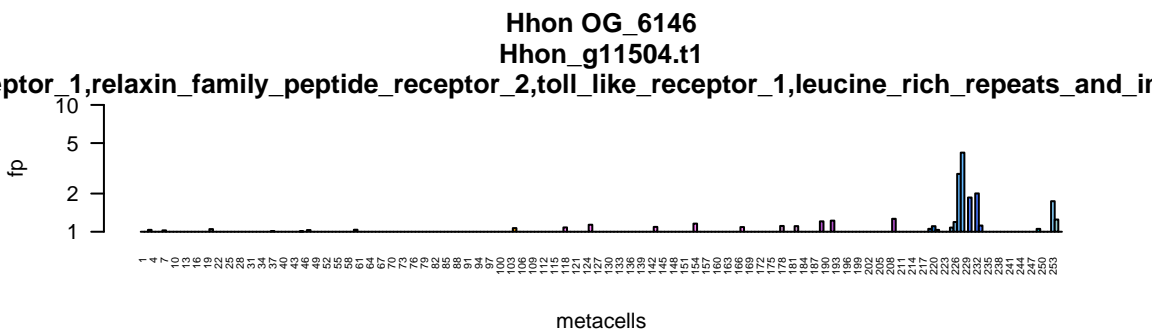
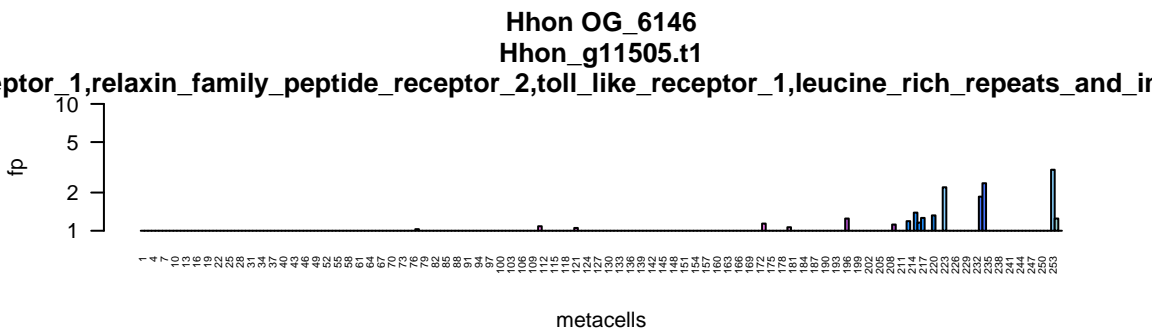
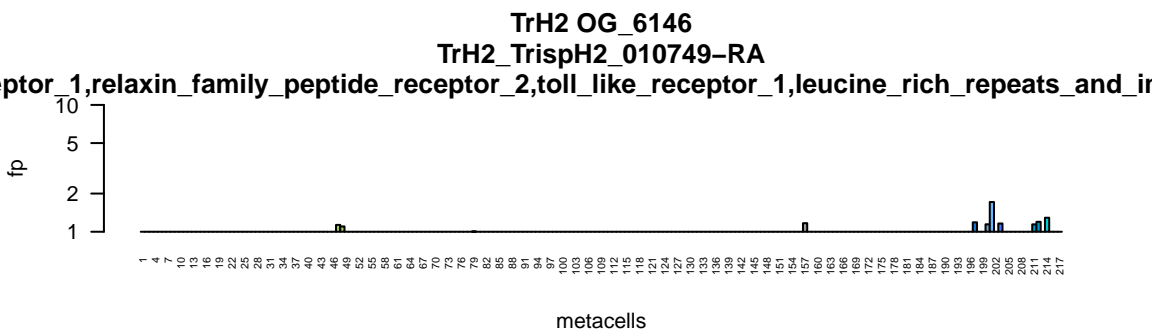
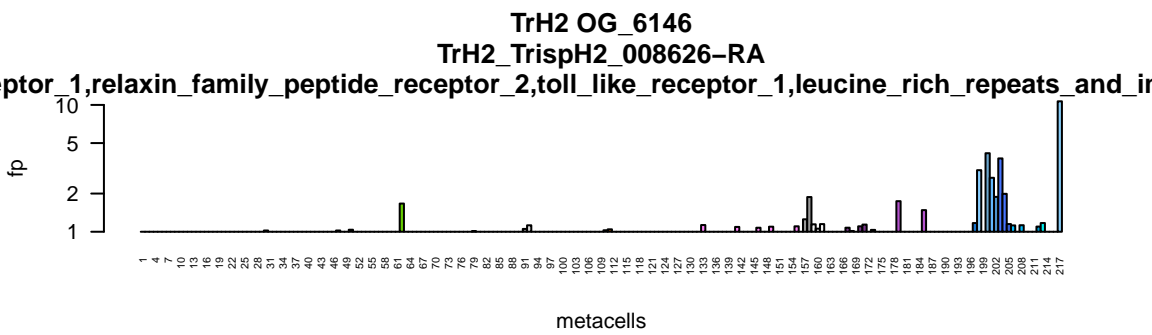
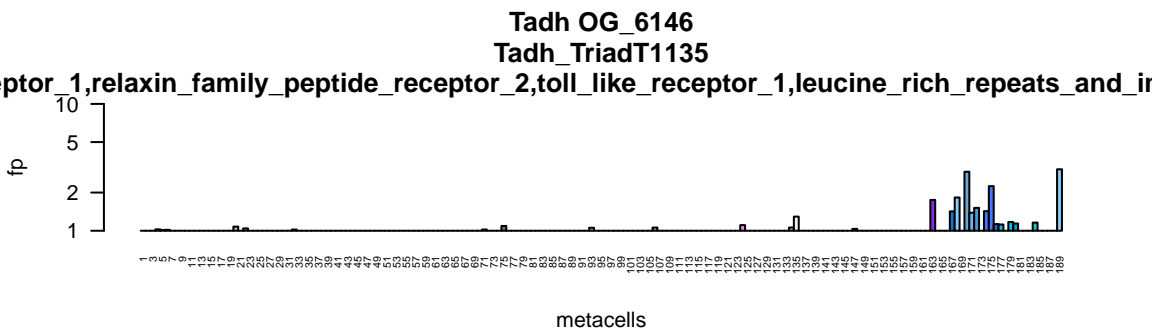
HoiH23 OG_5789

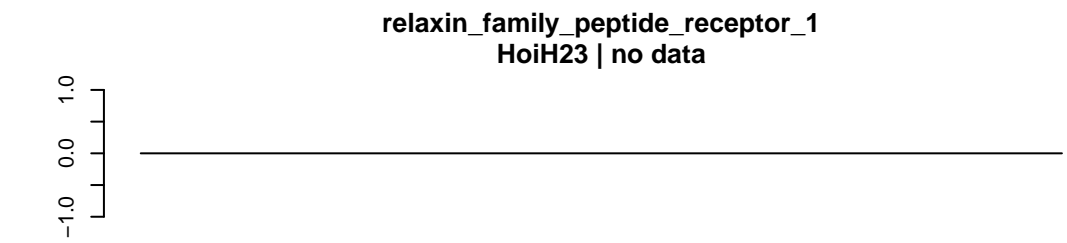
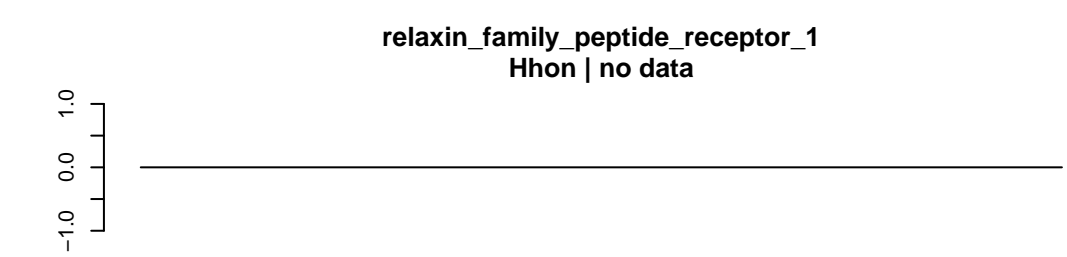
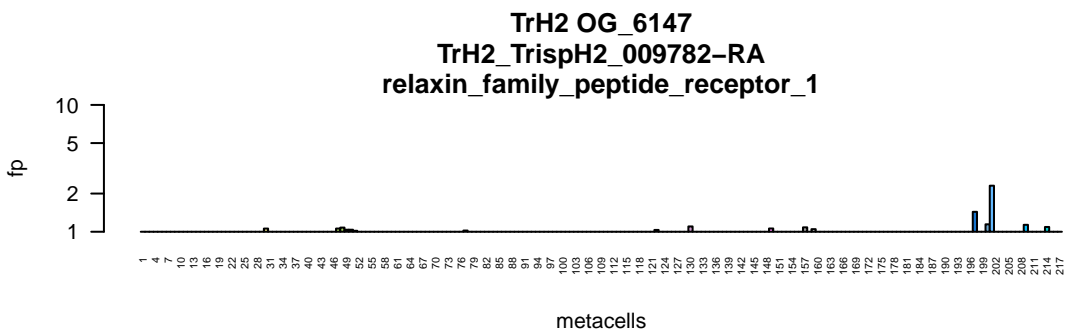
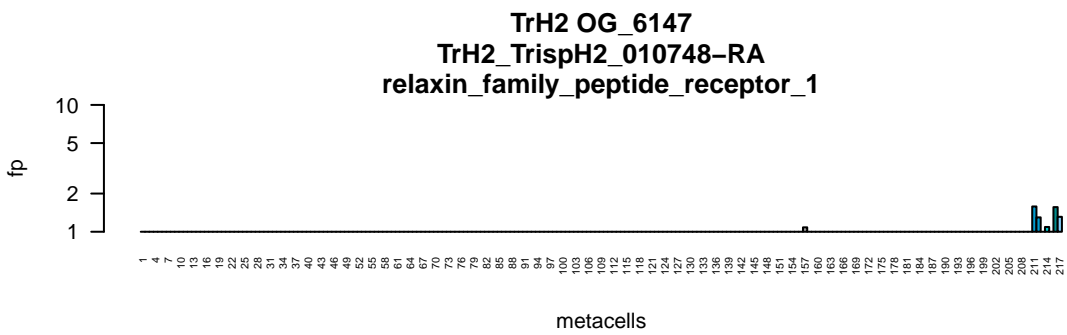
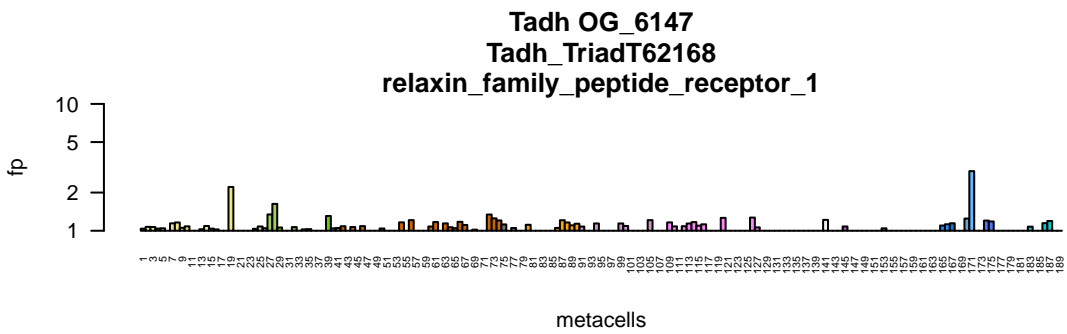
HoiH23_PIH23_011720-RA

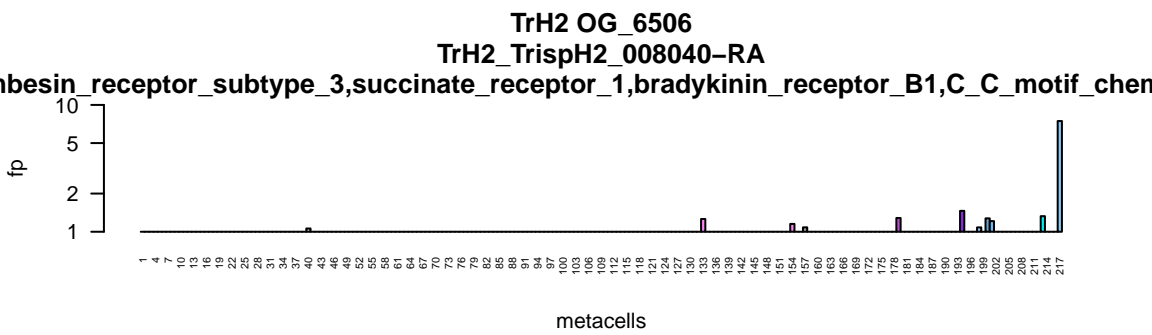
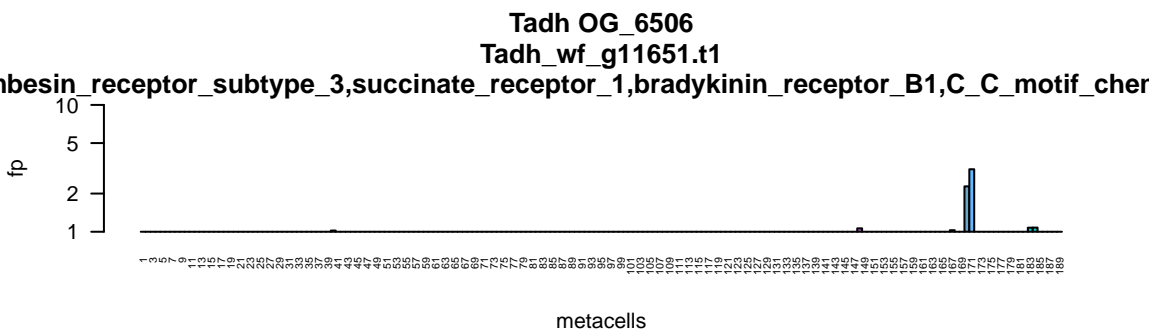
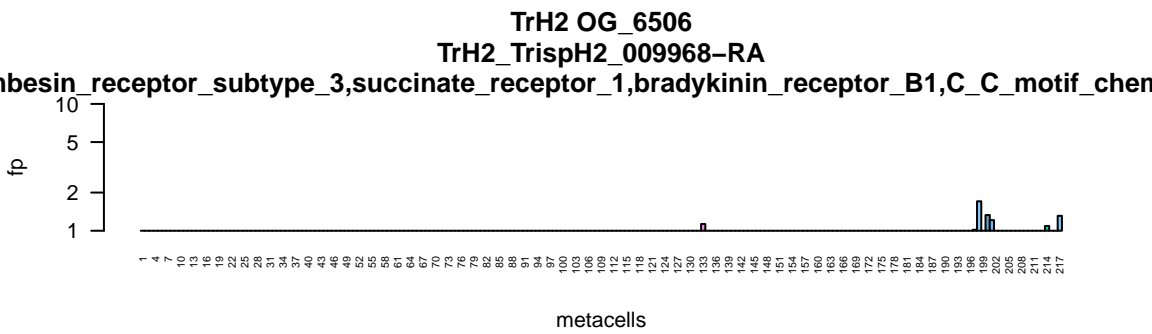
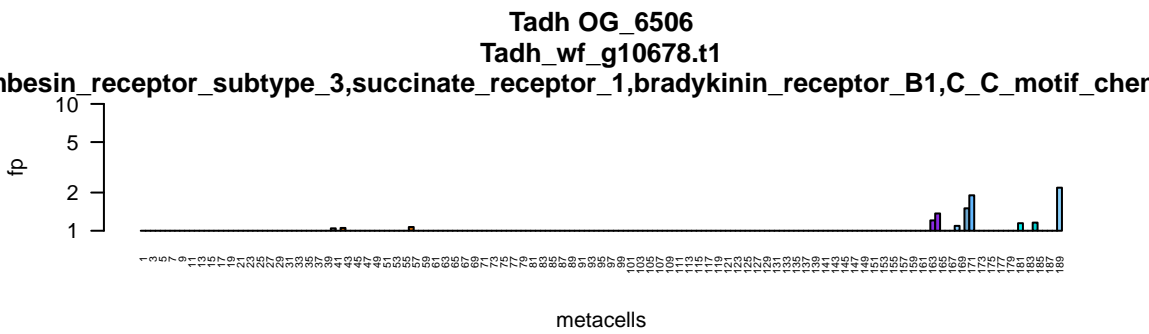
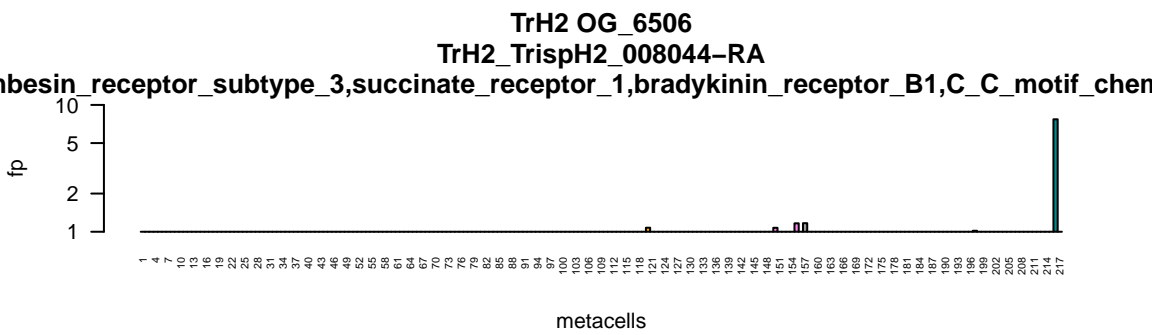
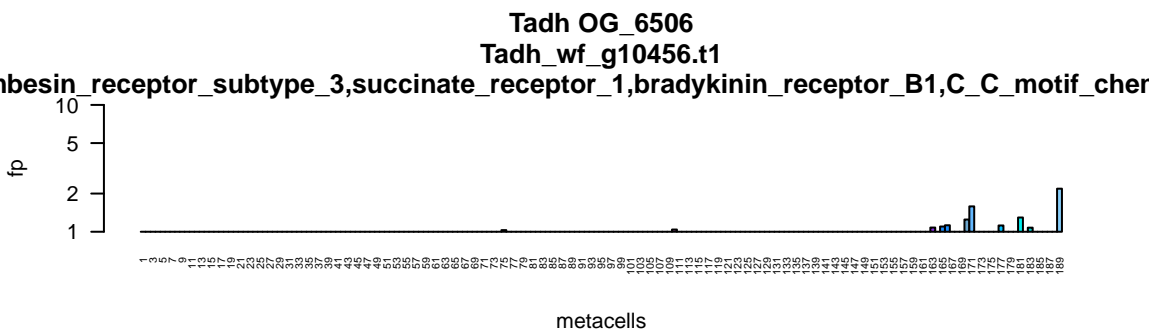
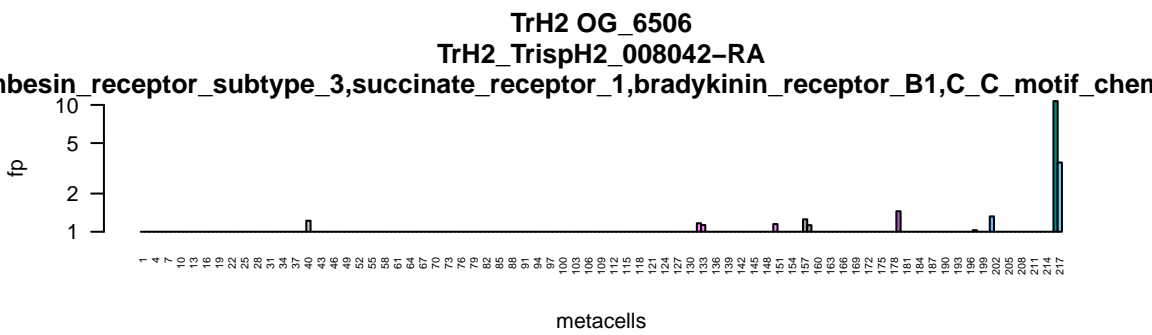
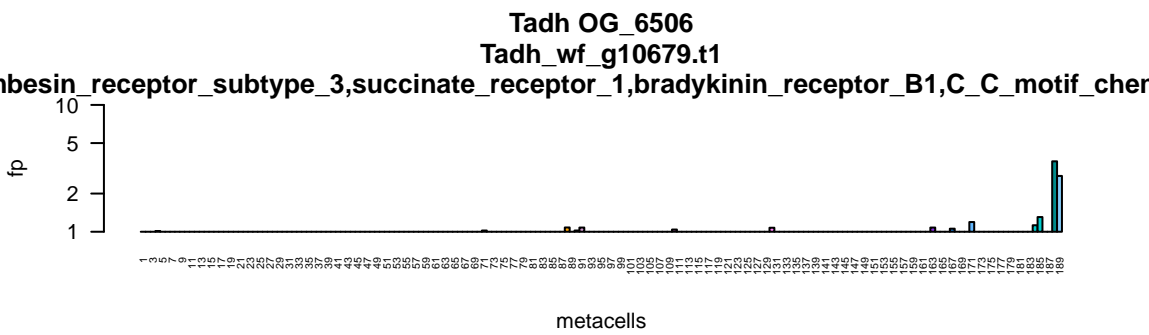
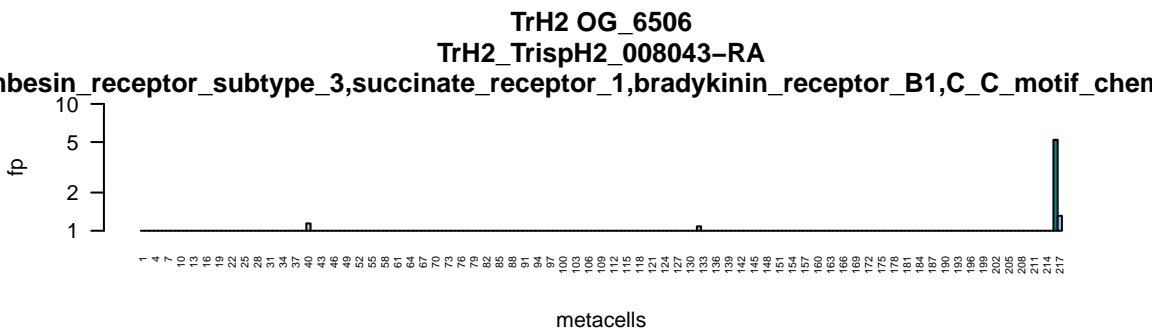
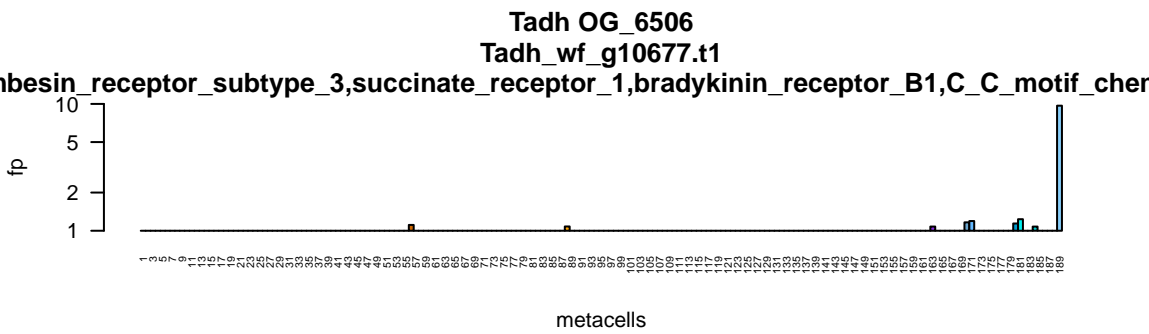
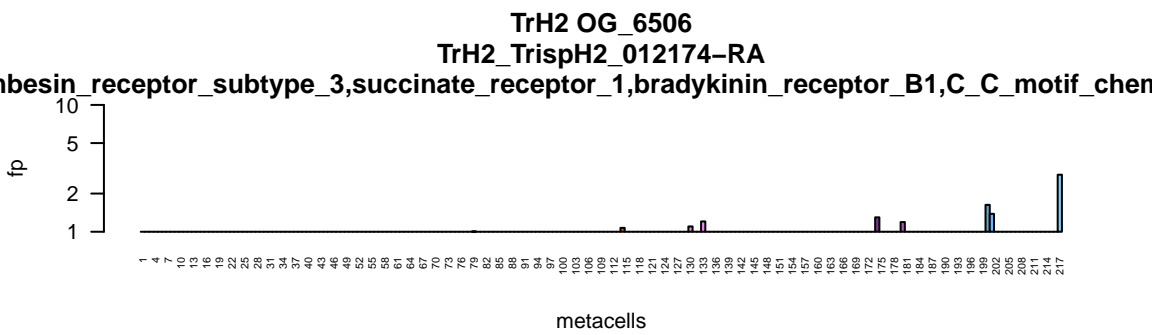
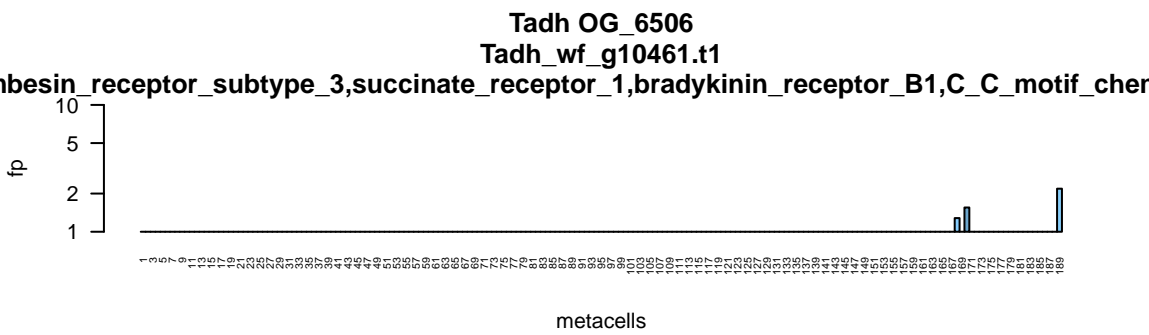
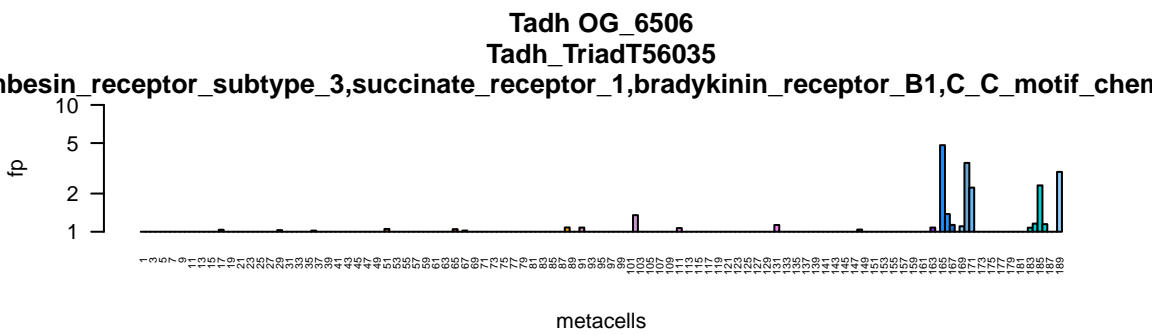
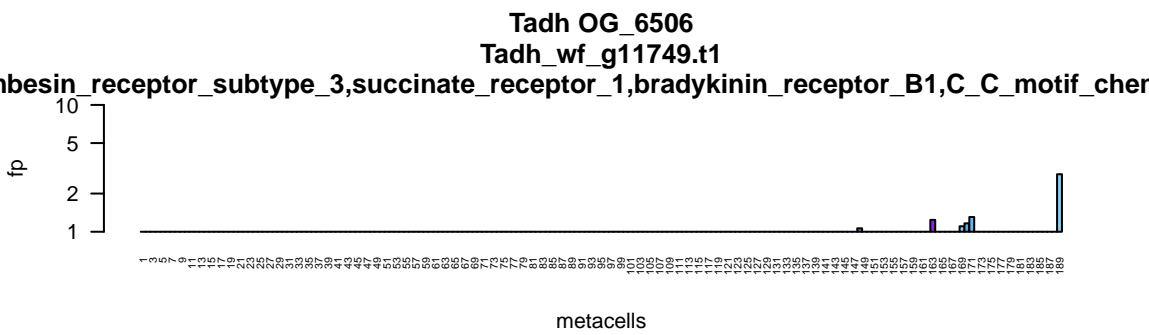
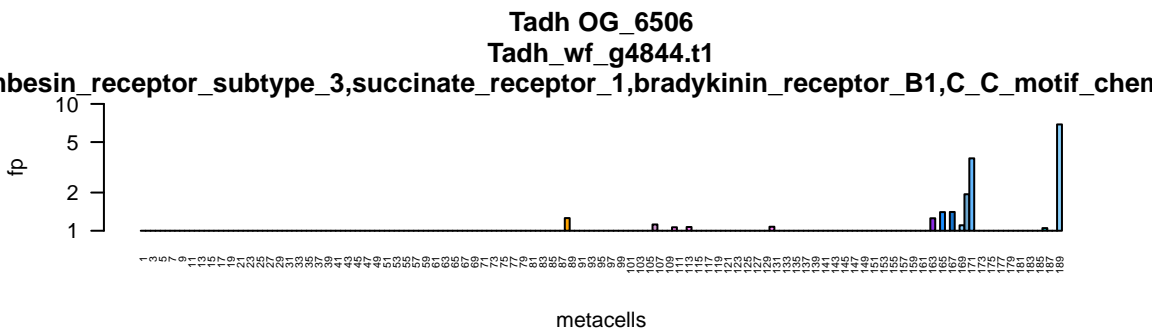
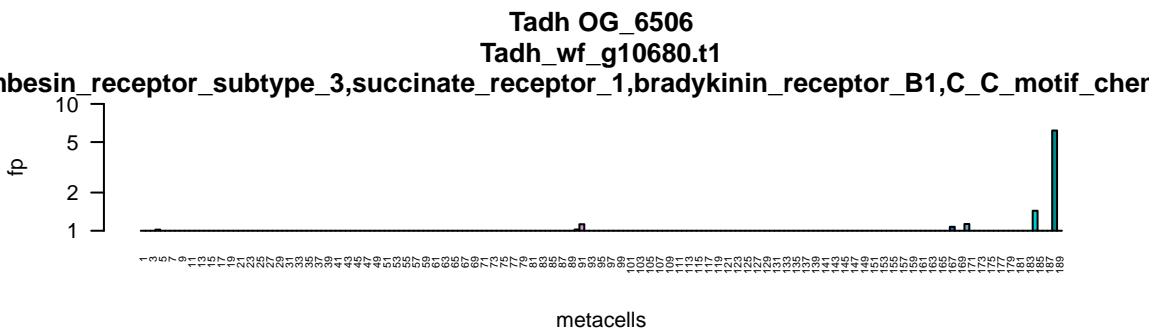
sphingosine_1_phosphate_receptor_1,neuropeptide_FF_receptor_1,opsin_3



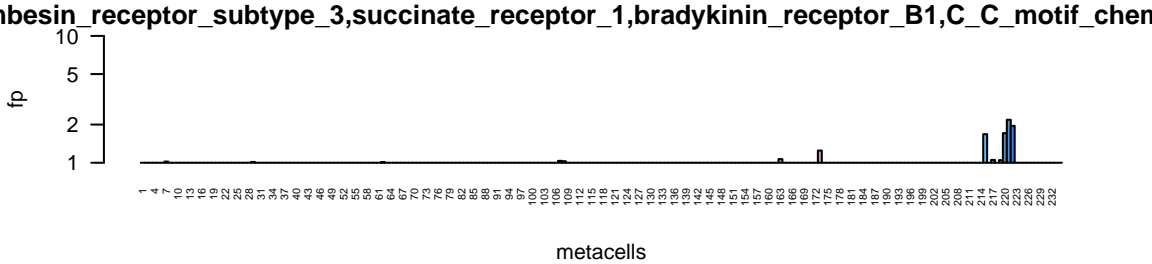
metacells



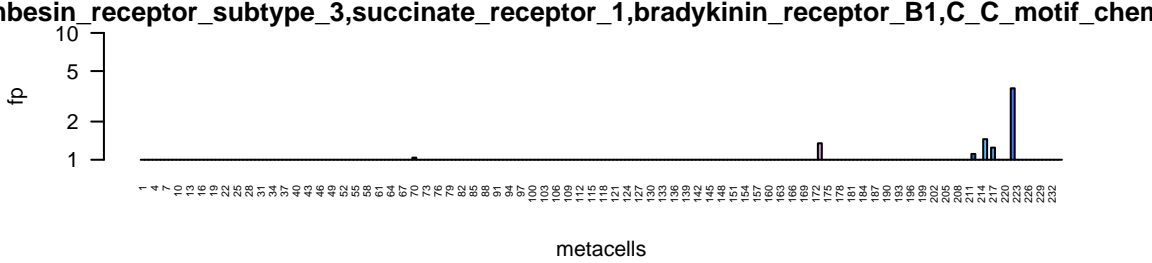




HoiH23 OG_6506
HoiH23_PIH23_008211-RA



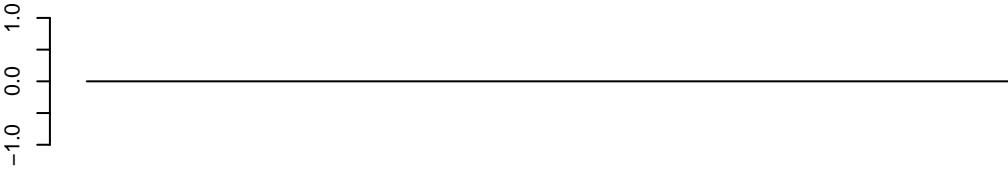
HoiH23 OG_6506
HoiH23_PIH23_012037-RA



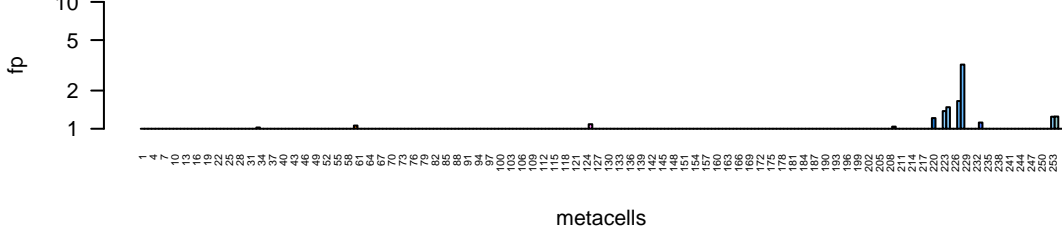
opioid_receptor_kappa_1,opioid_receptor_mu_1
Tadh | no data



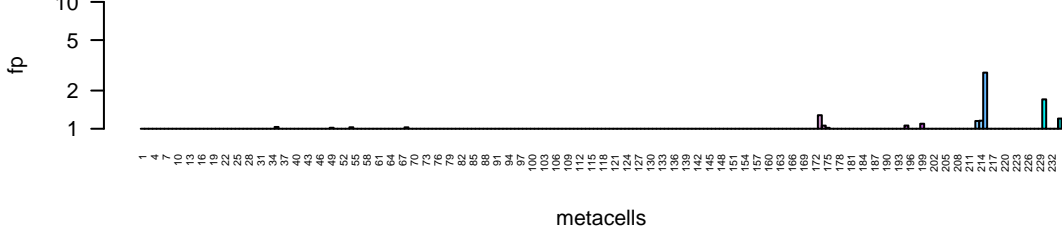
opioid_receptor_kappa_1,opioid_receptor_mu_1
TrH2 | no data

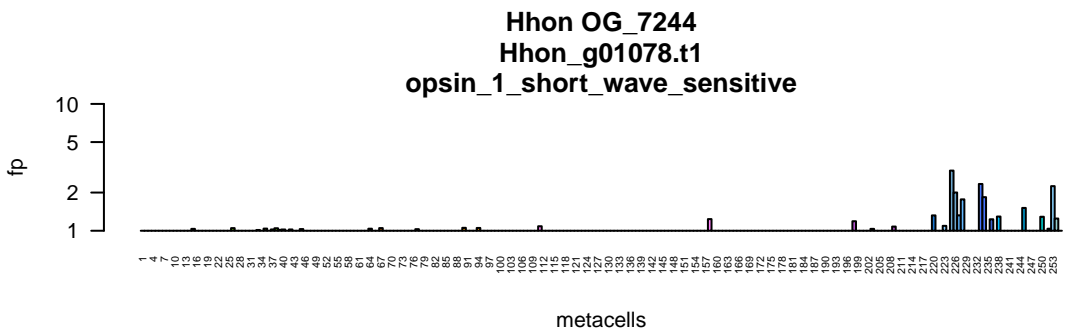
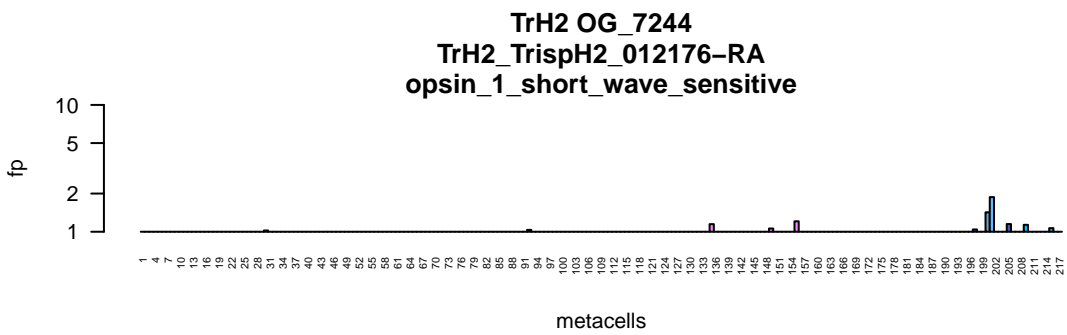
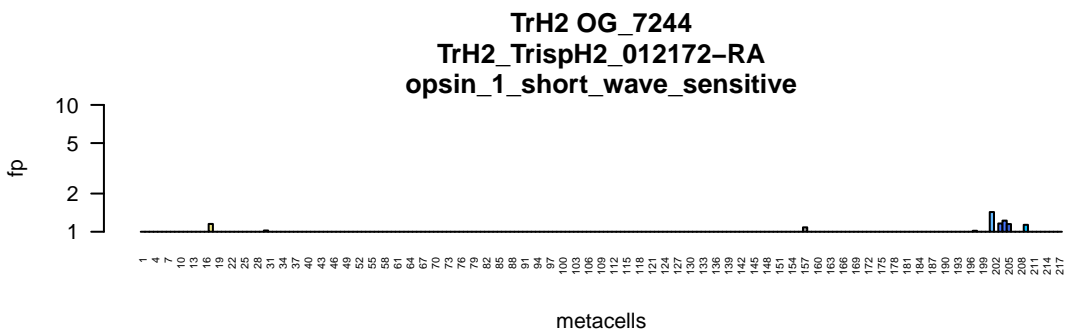
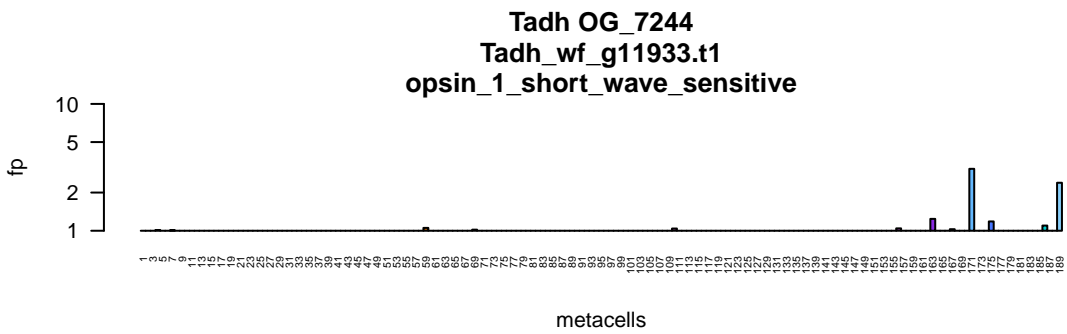


Hhon OG_6769
Hhon_g06796.t1
opioid_receptor_kappa_1,opioid_receptor_mu_1

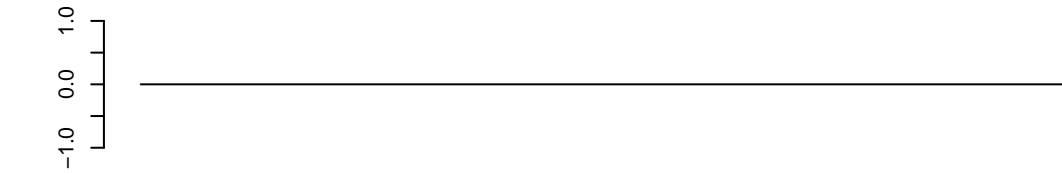


HoiH23 OG_6769
HoiH23_PIH23_003884-RA
opioid_receptor_kappa_1,opioid_receptor_mu_1

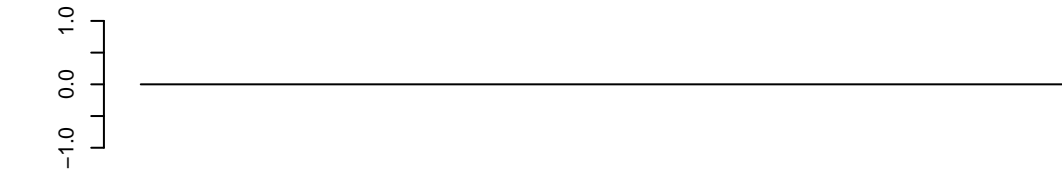




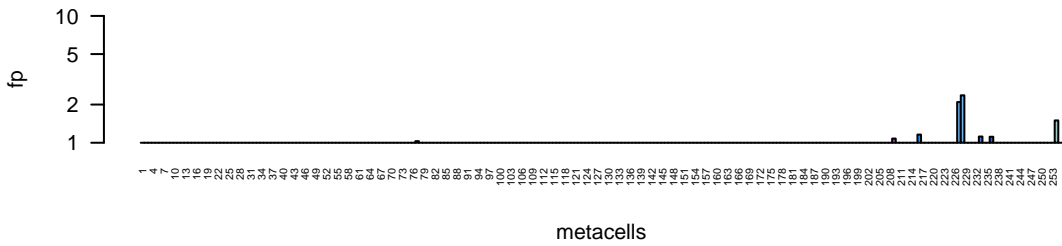
Tadh | no data



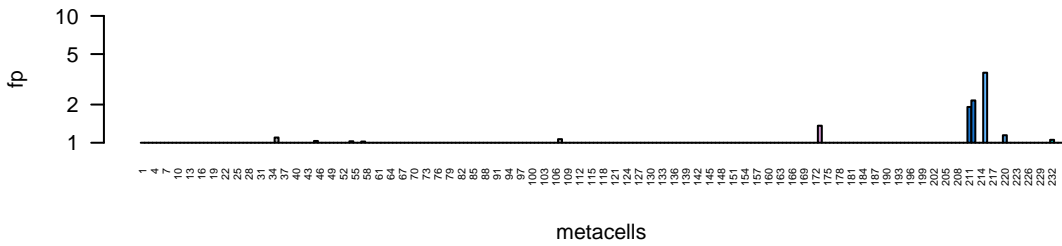
TrH2 | no data



Hhon OG_8093
Hhon_g03455.t1



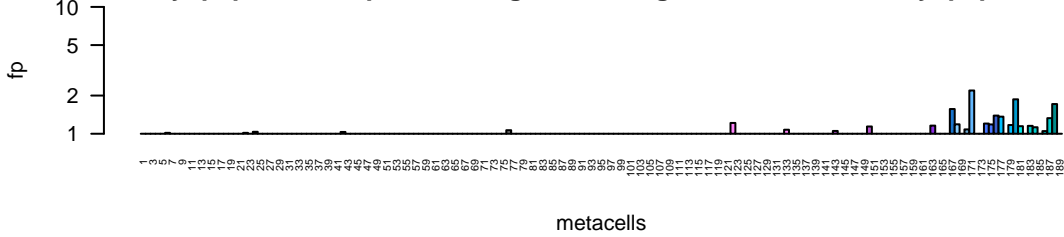
HoiH23 OG_8093
HoiH23_PIH23_007146-RA



Tadh OG_9223

Tadh_TriadT51796

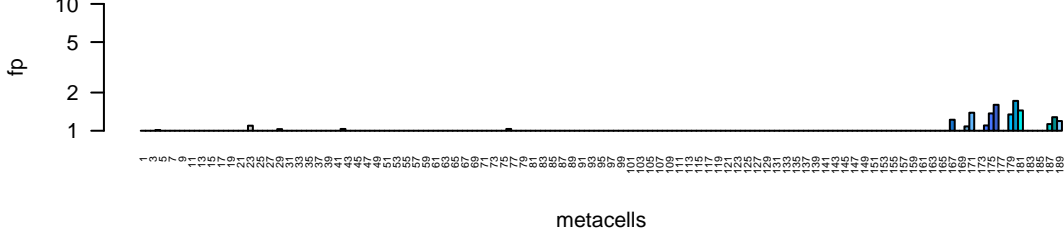
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Tadh OG_9223

Tadh_TriadT51797

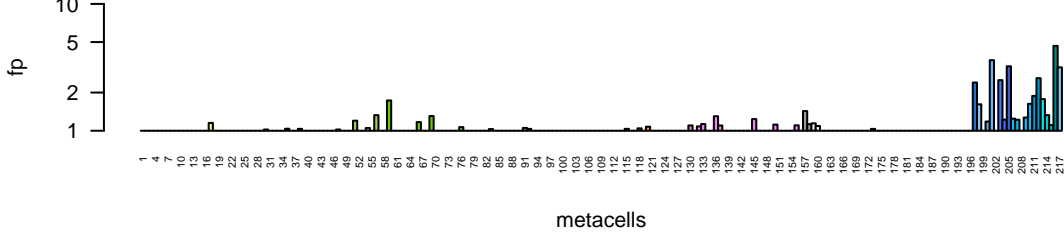
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



TrH2 OG_9223

TrH2_TrispH2_008387-RA

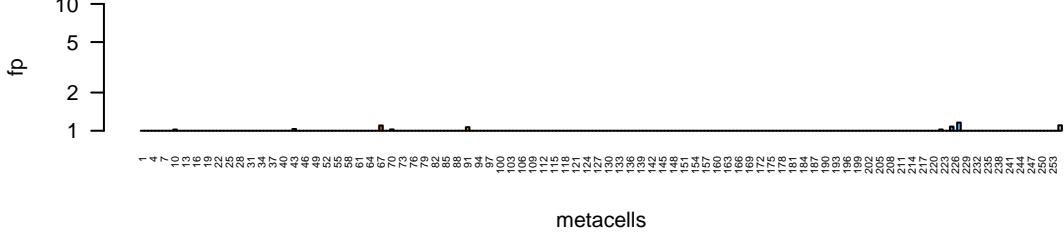
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Hhon OG_9223

Hhon_g11380.t1

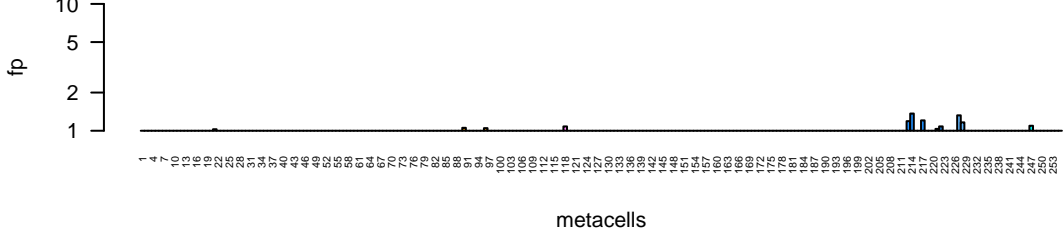
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Hhon OG_9223

Hhon_g01959.t1

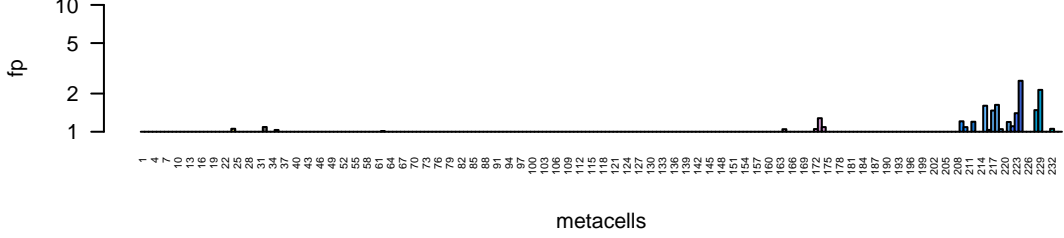
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto

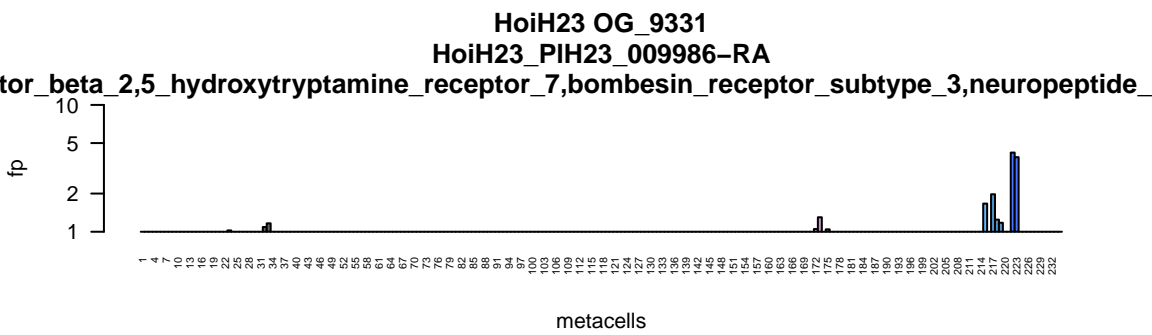
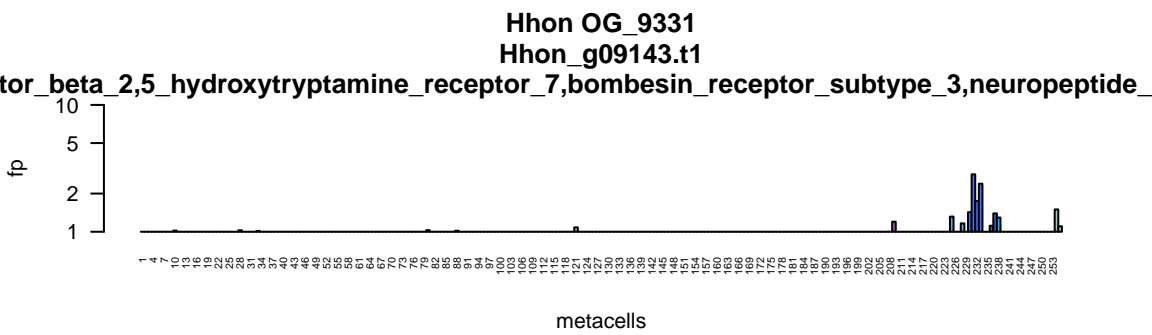
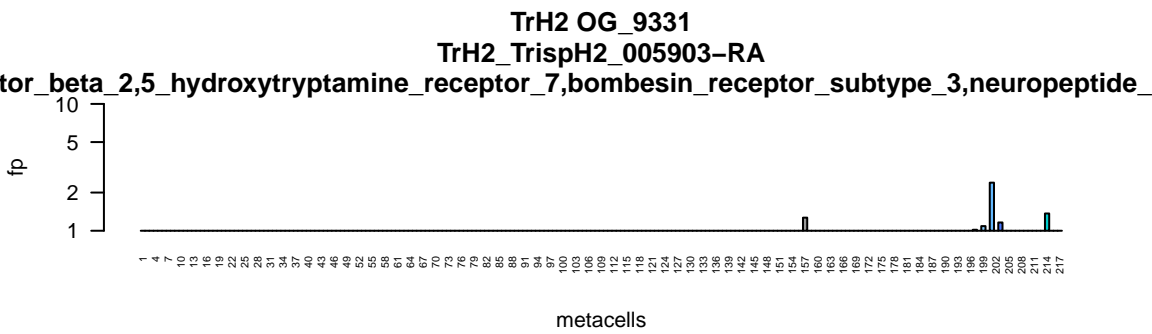
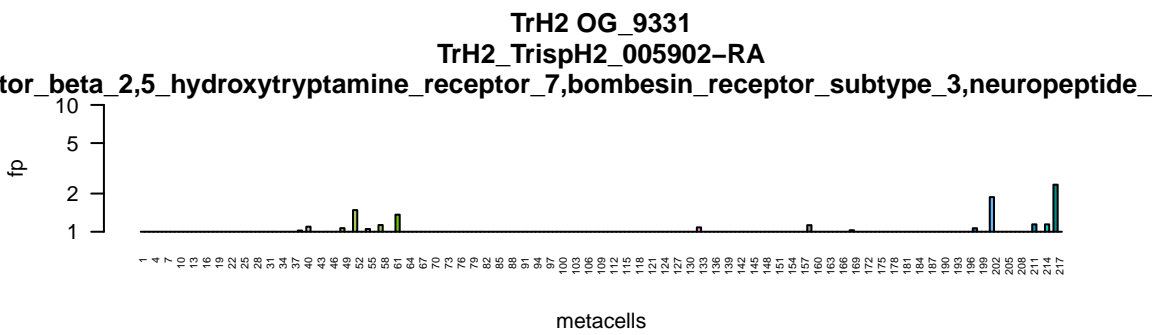
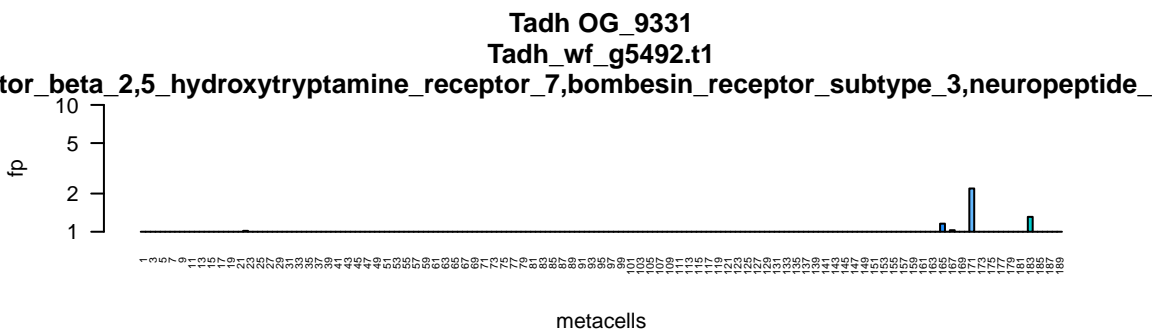
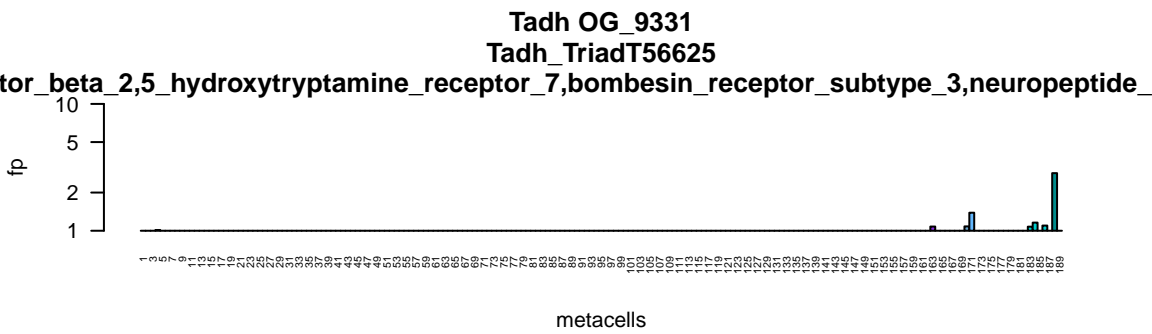


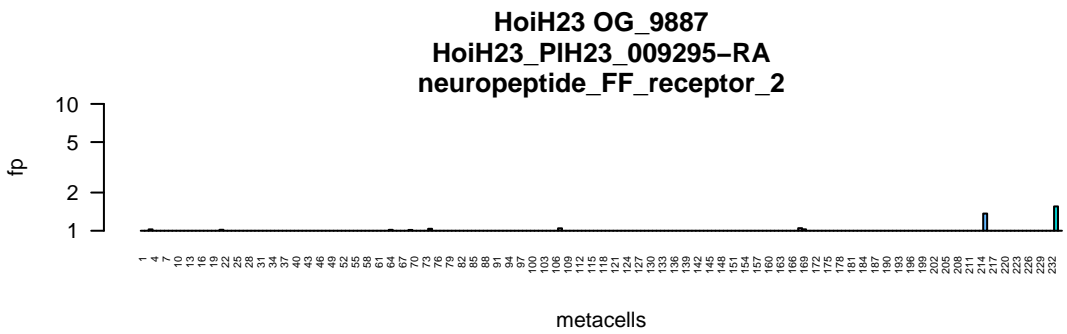
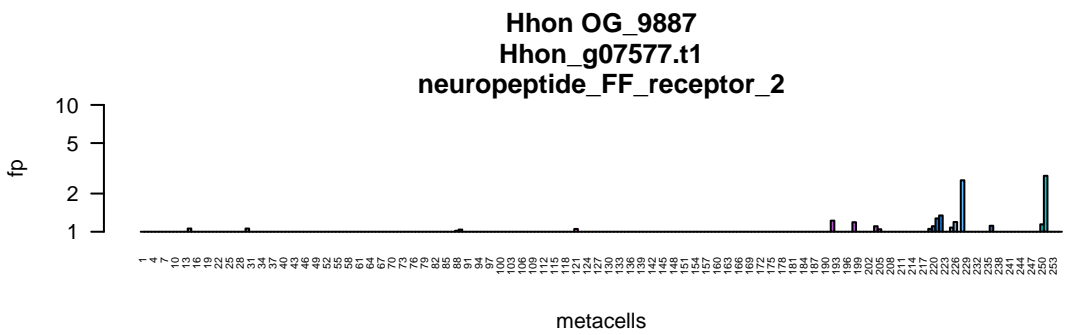
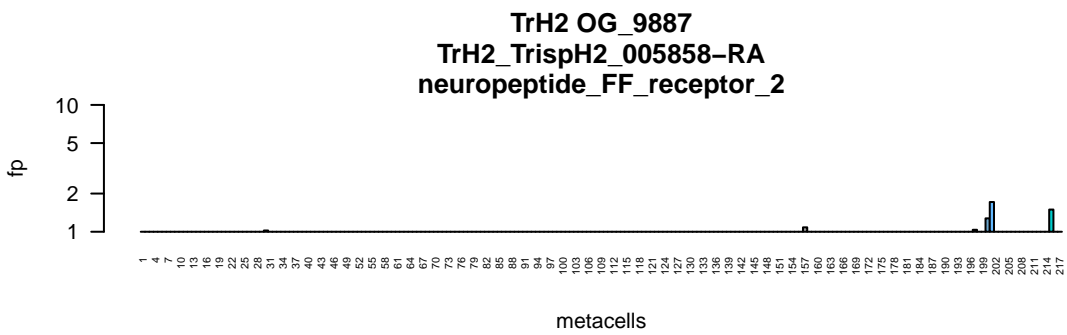
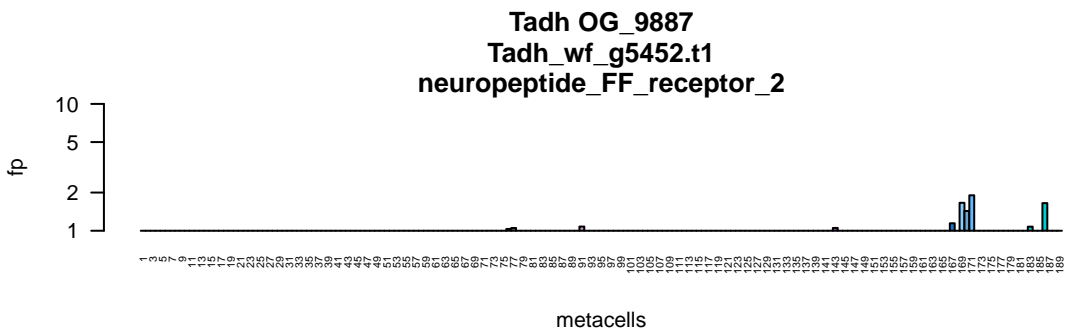
HoiH23 OG_9223

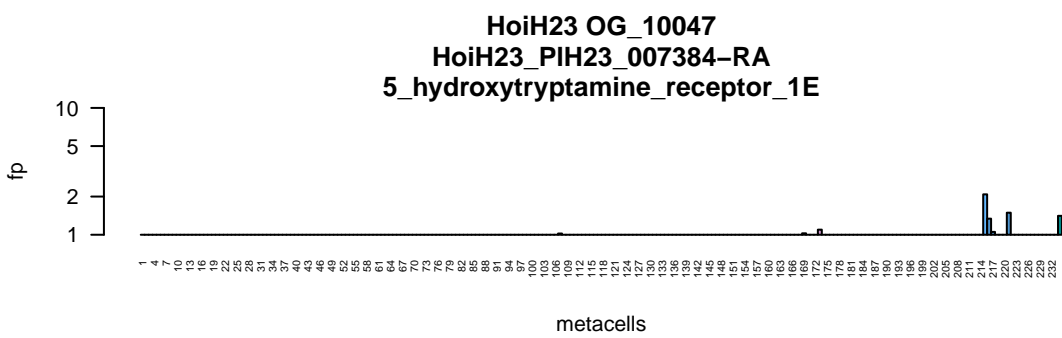
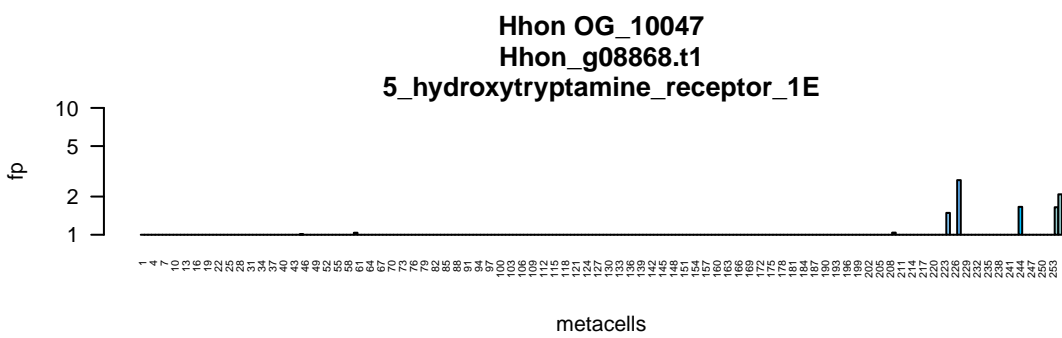
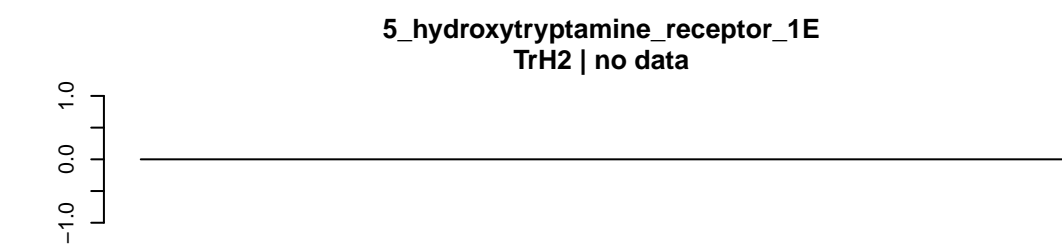
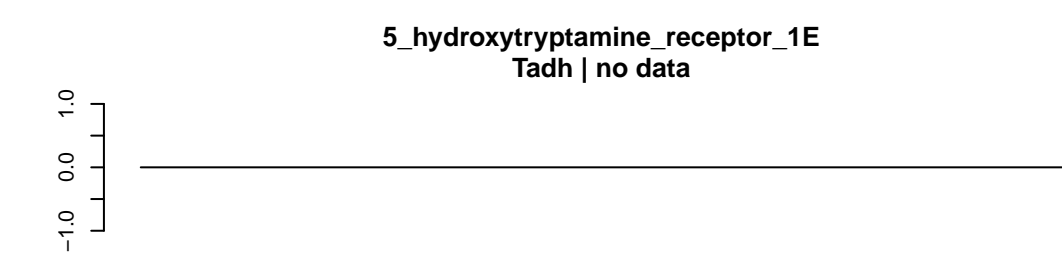
HoiH23_PIH23_004873-RA

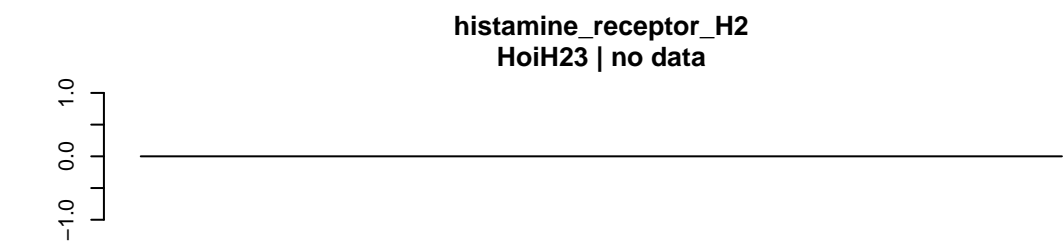
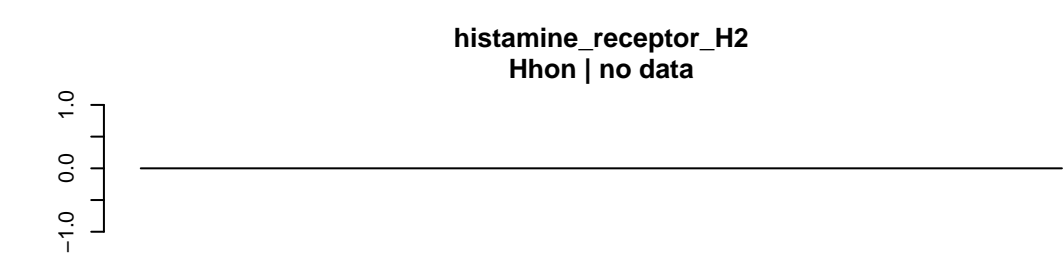
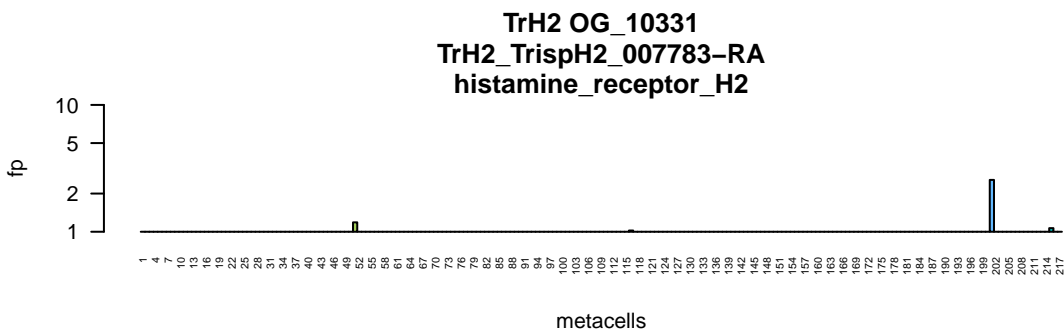
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto

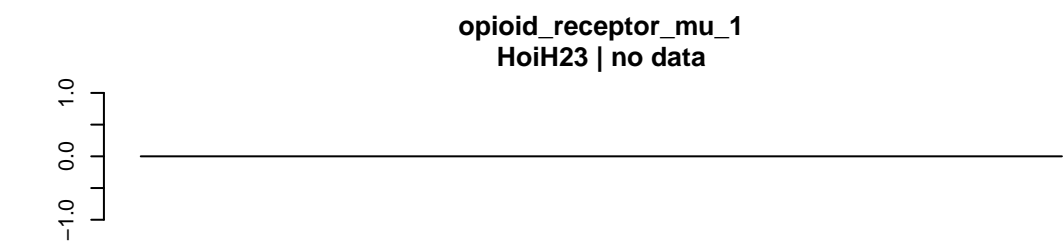
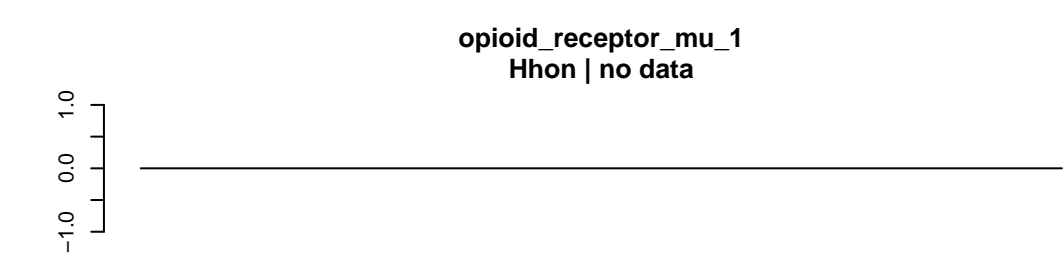
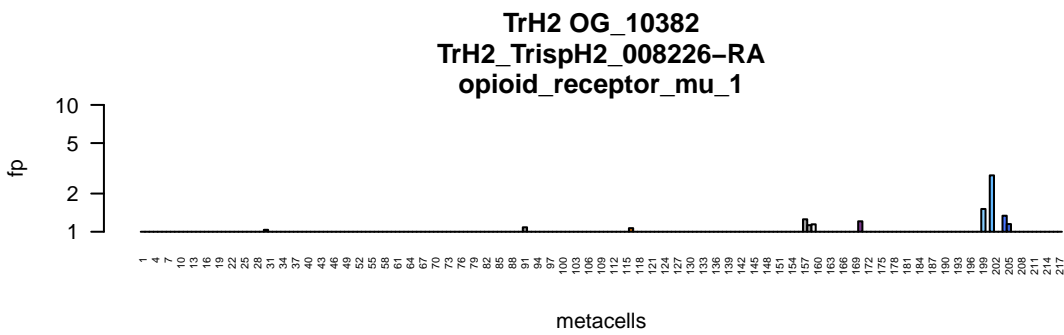
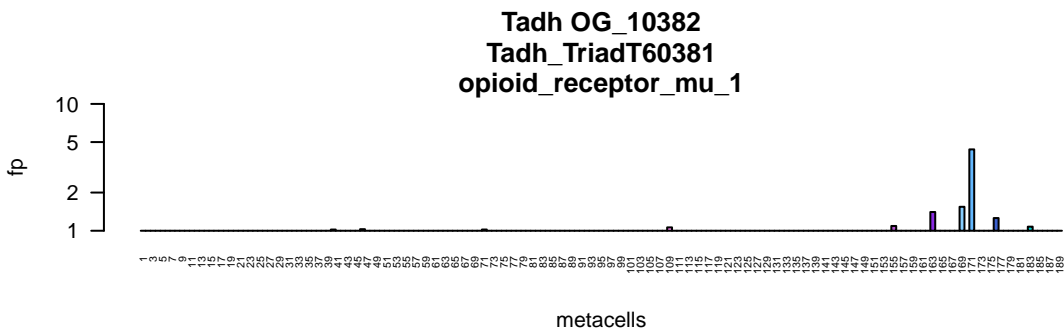


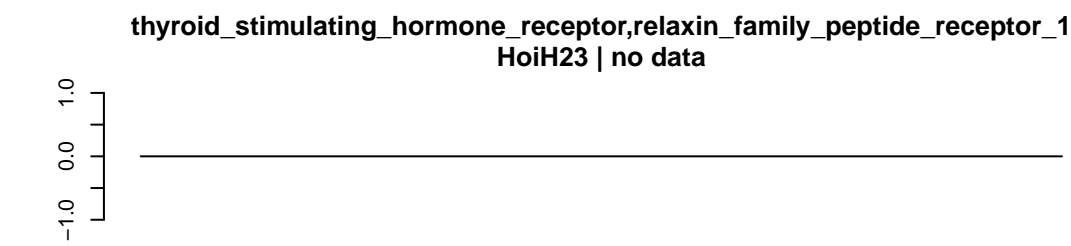
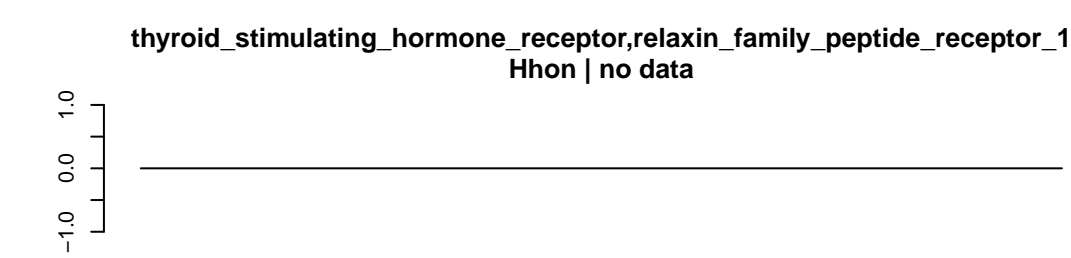
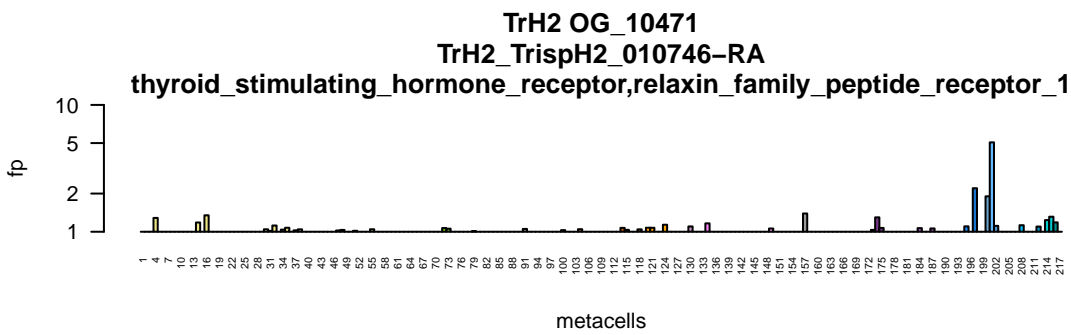
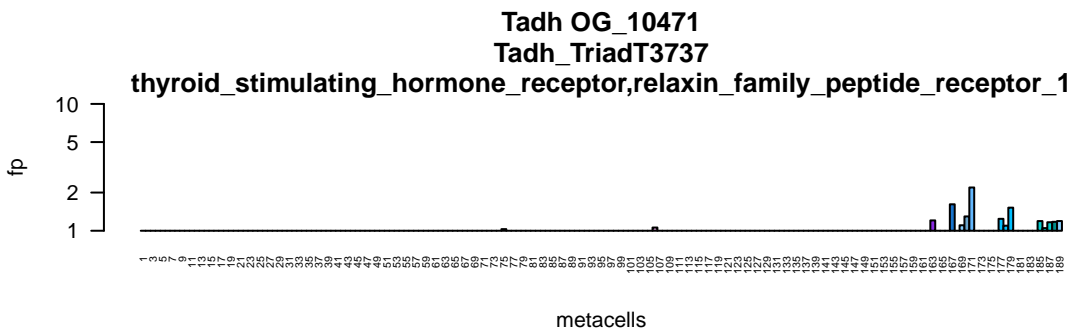


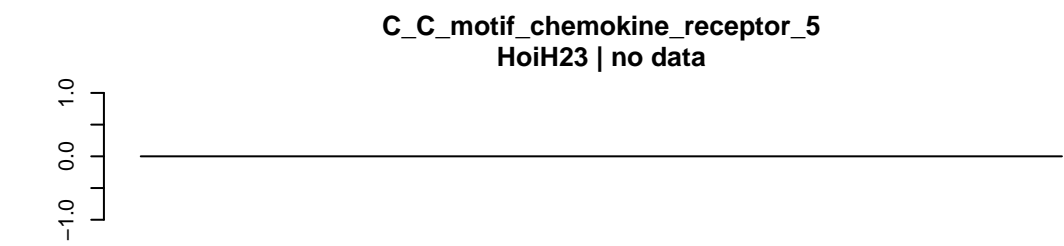
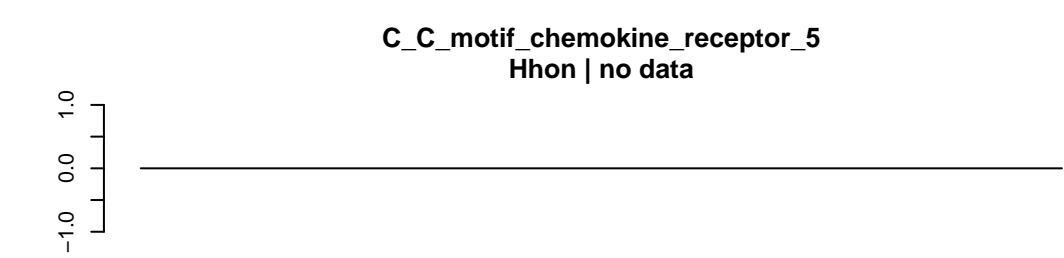
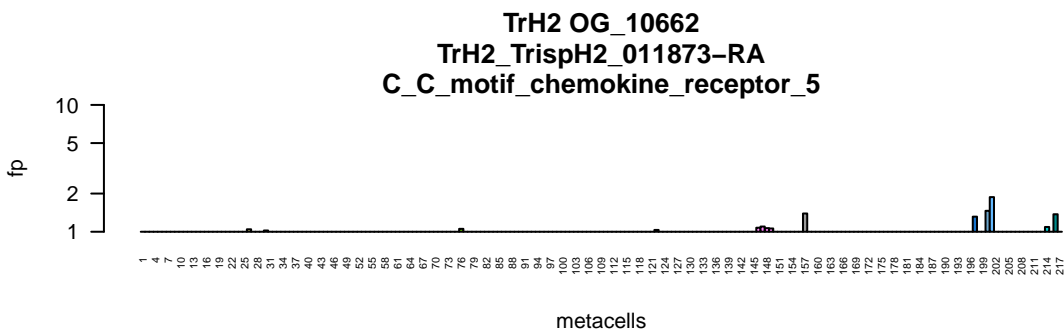
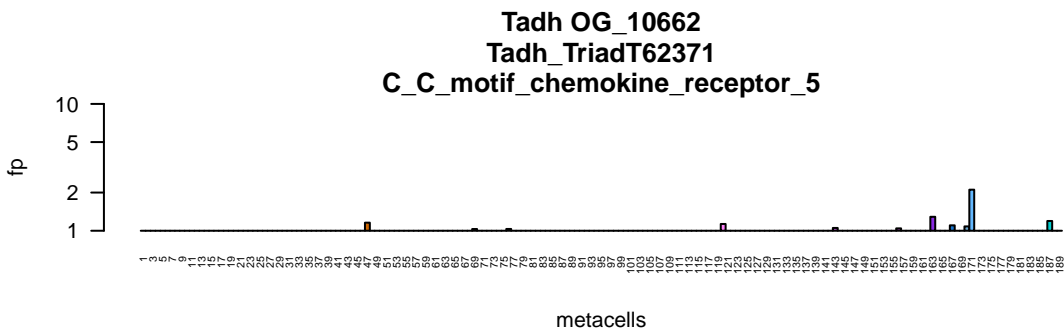


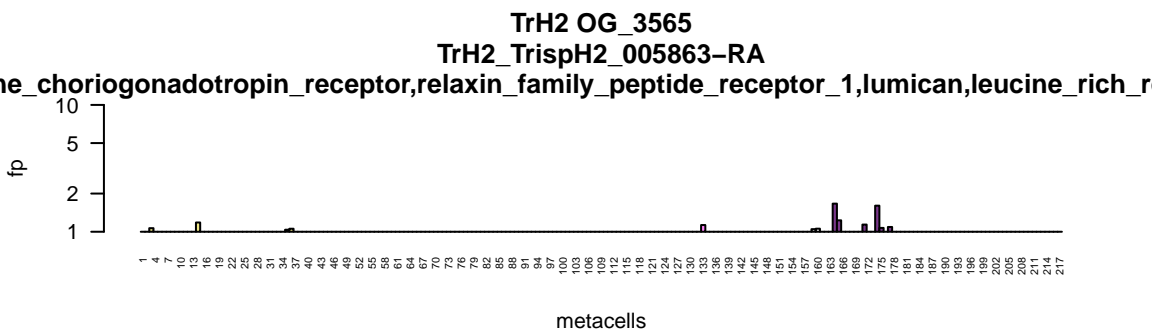
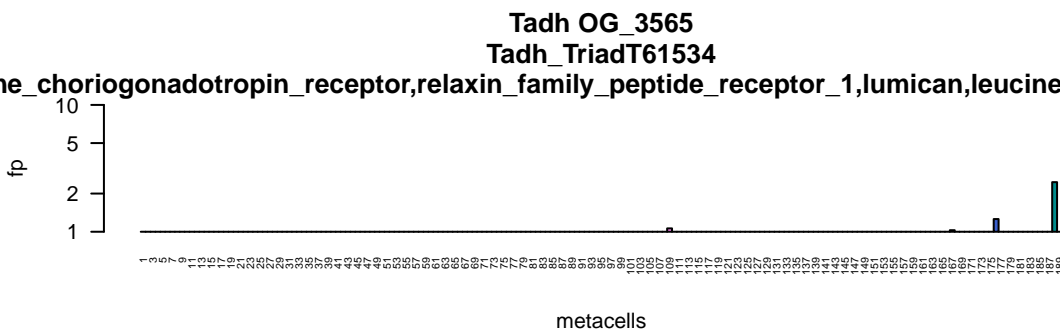
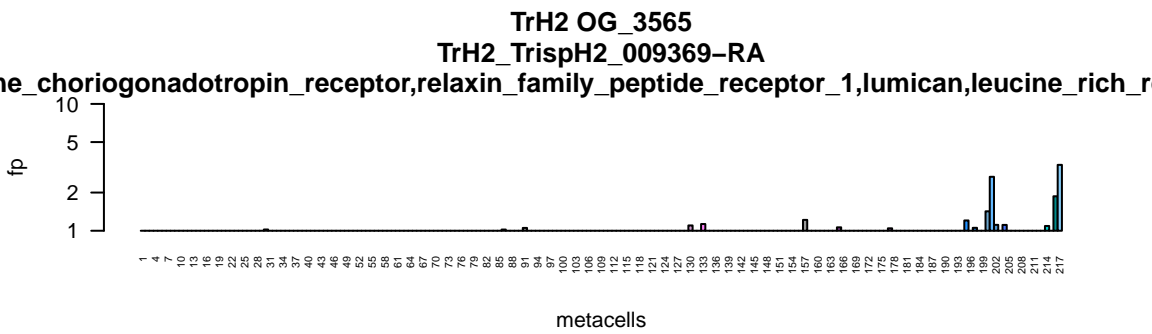
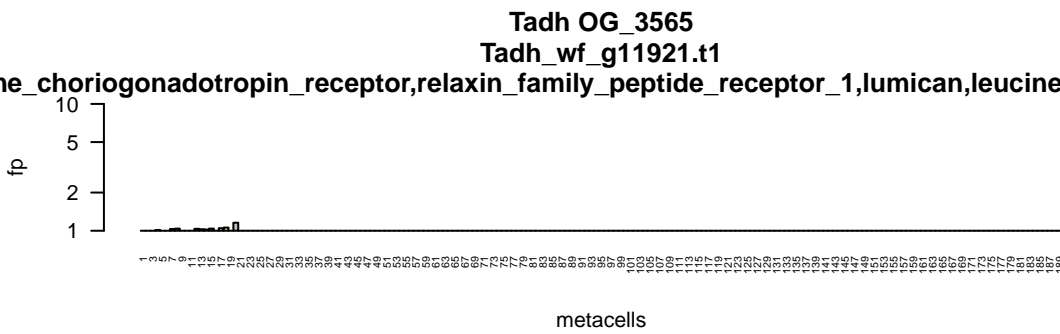
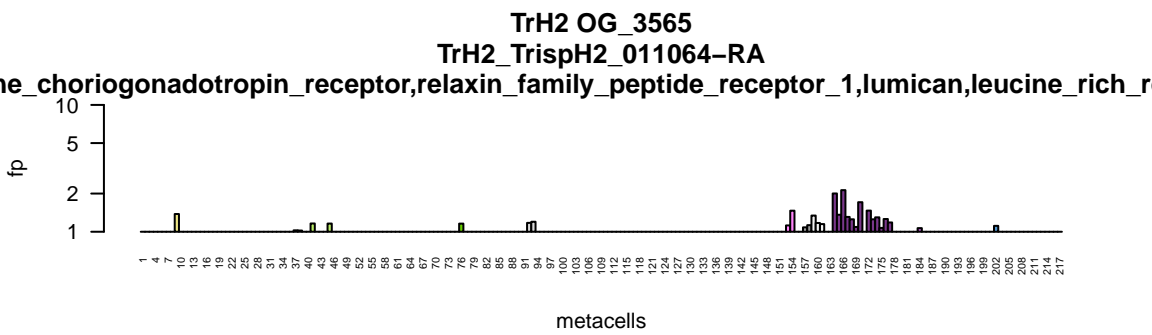
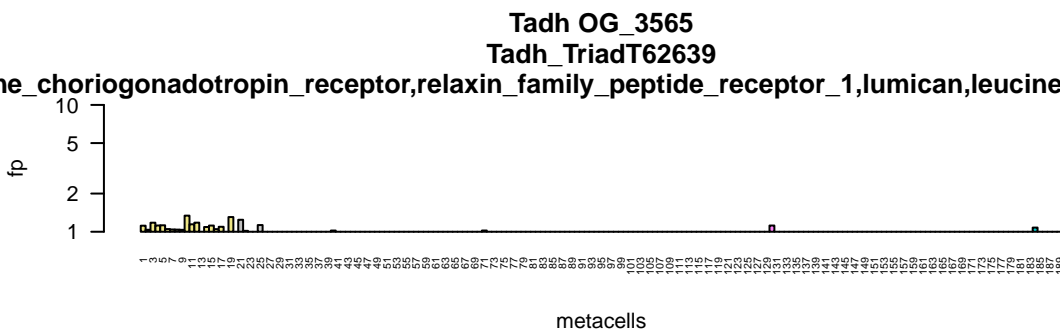
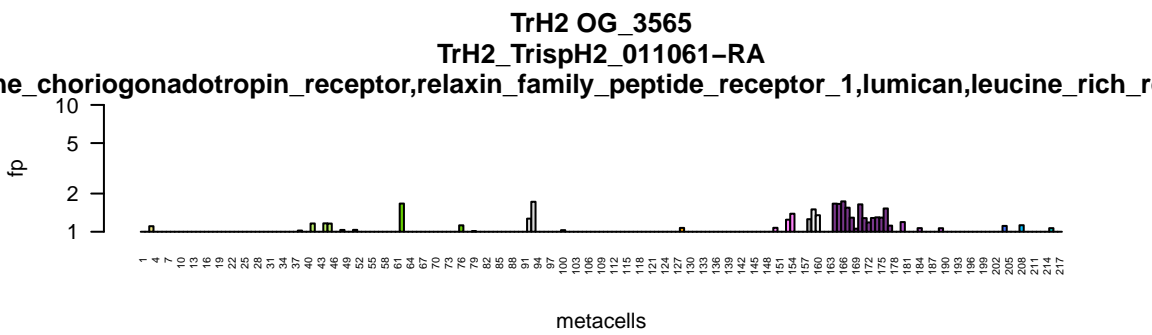
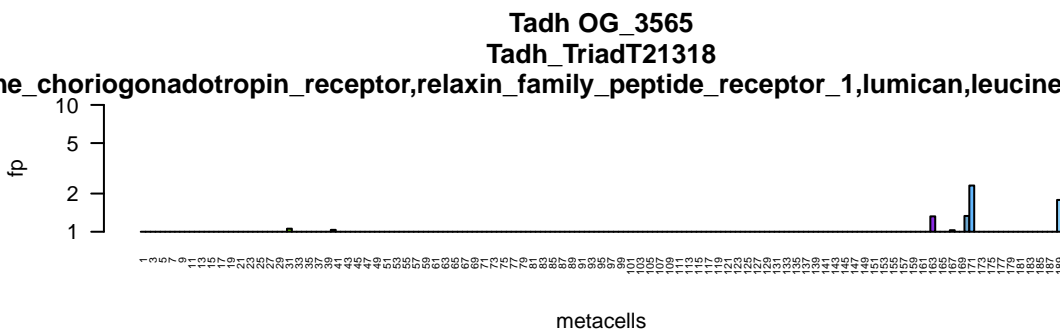
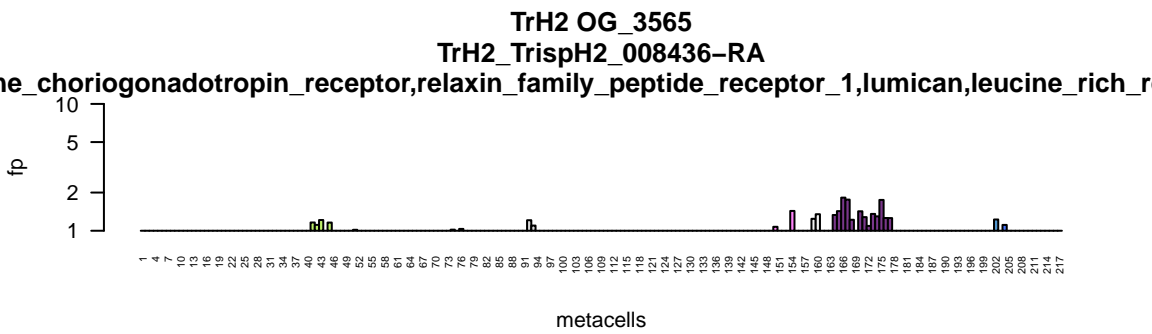
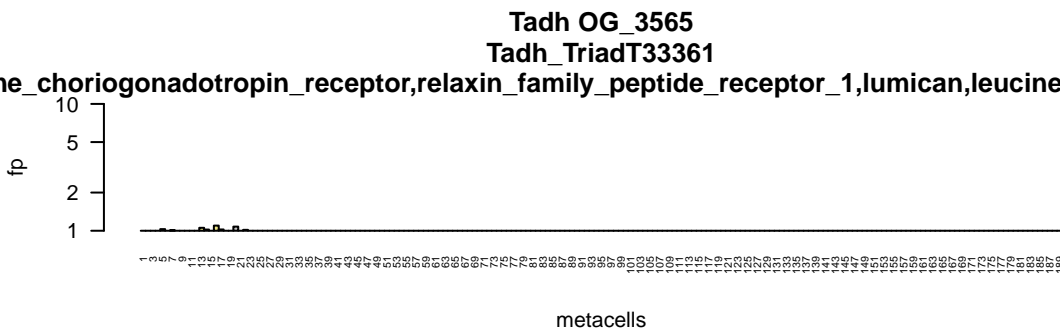
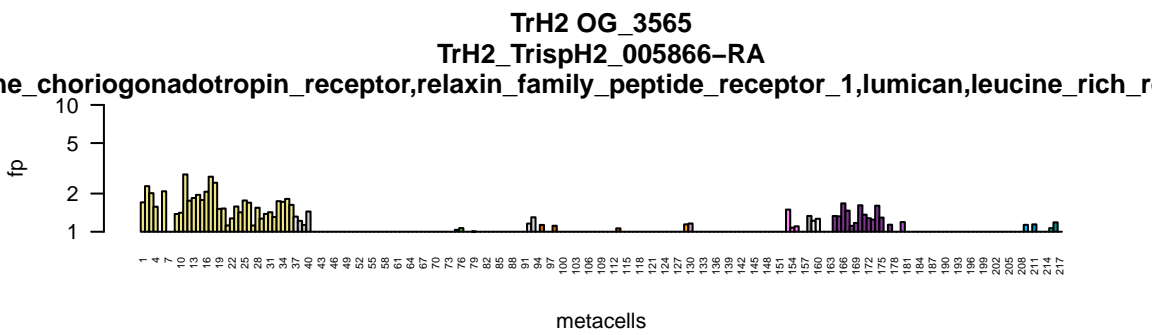
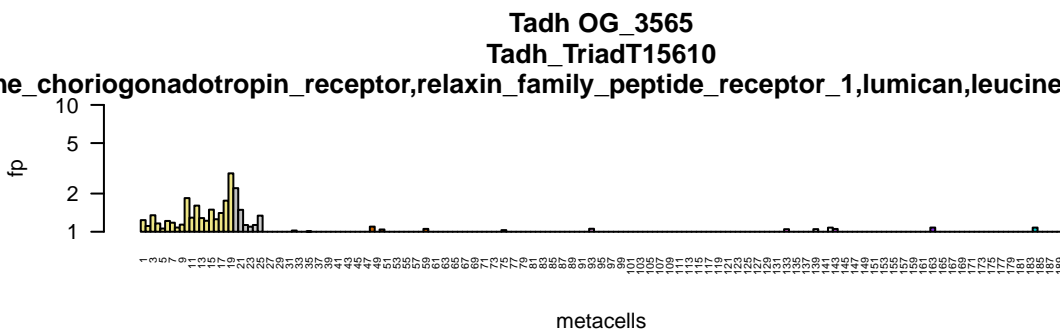
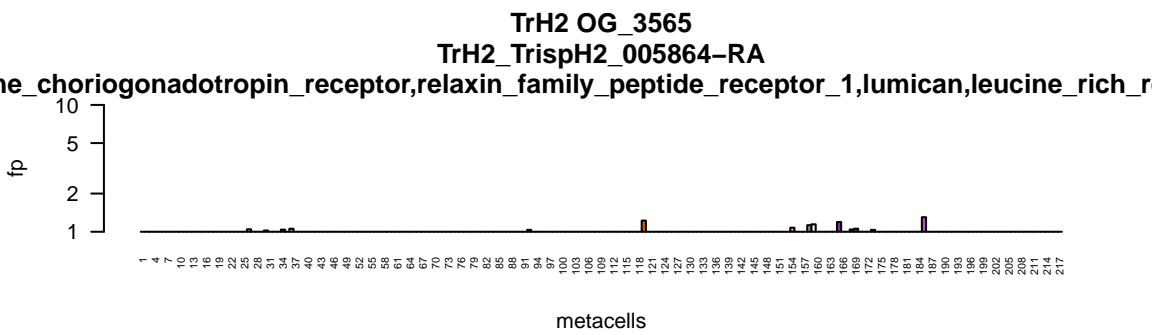
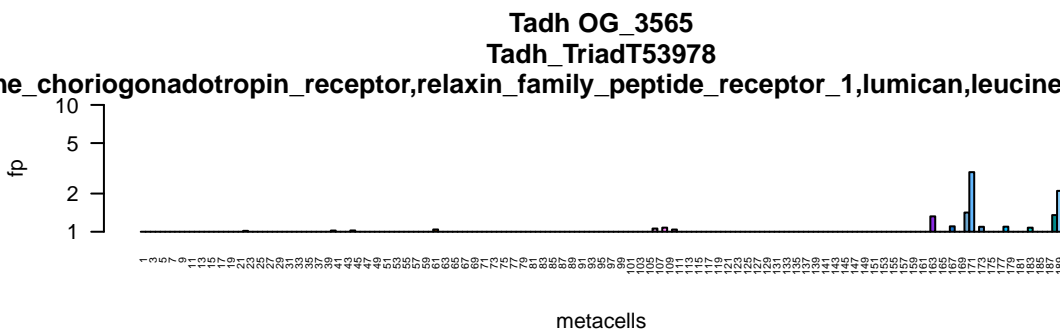
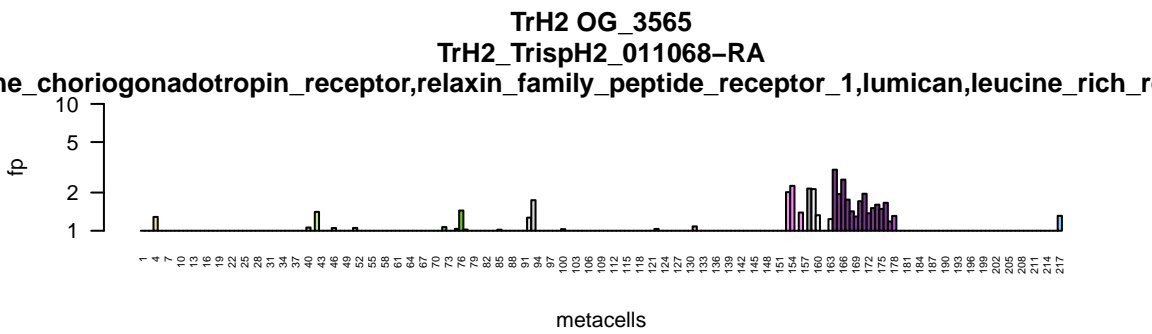
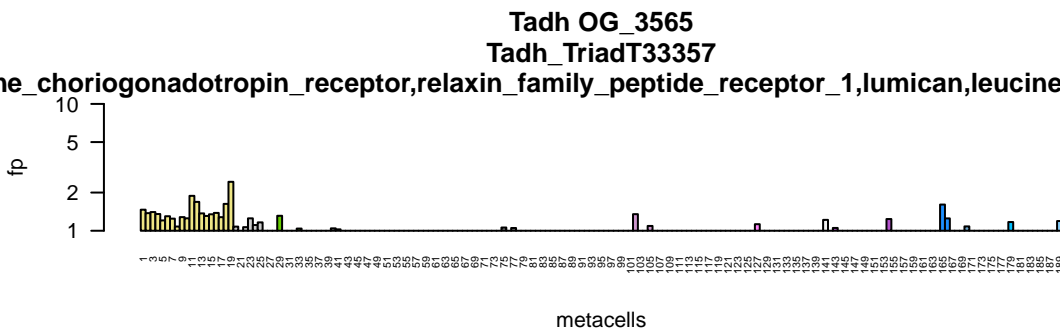


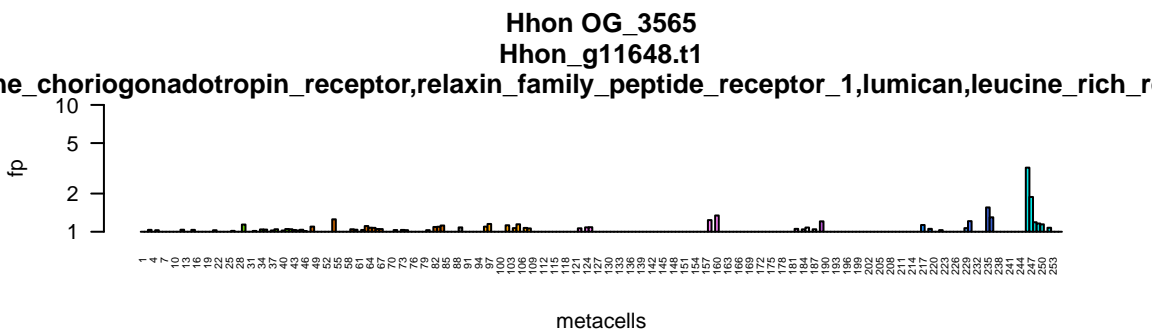
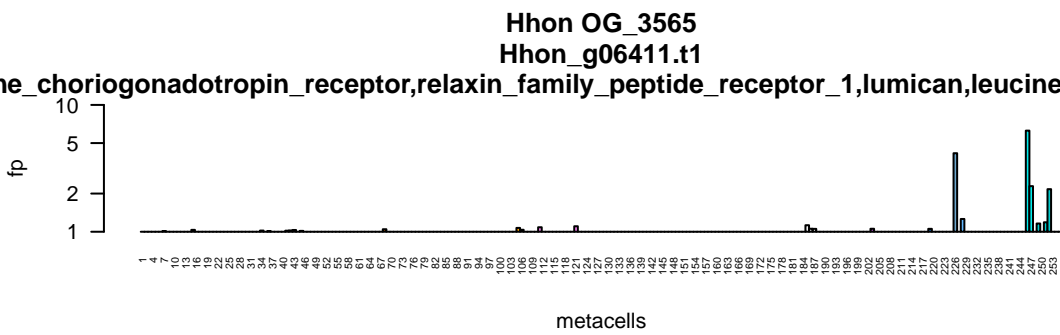
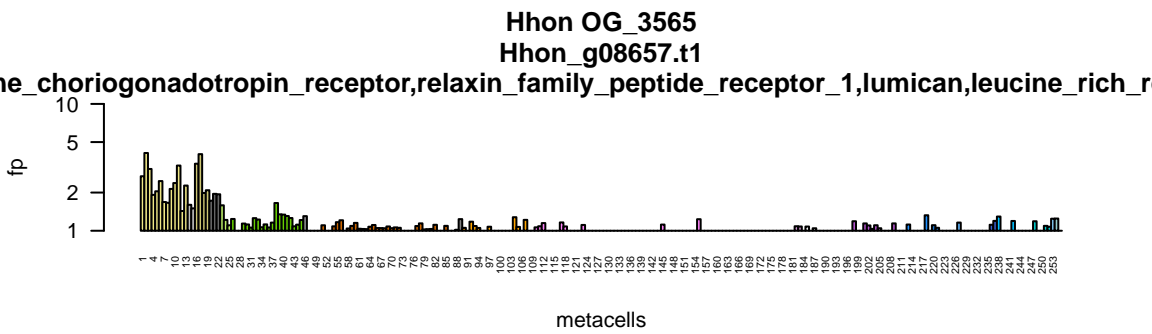
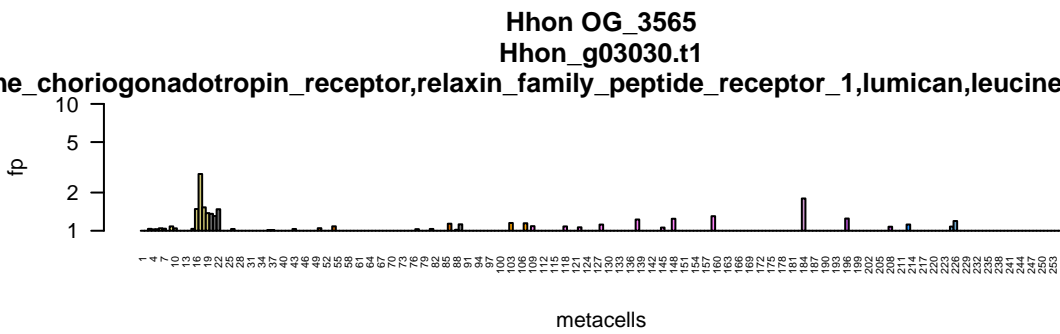
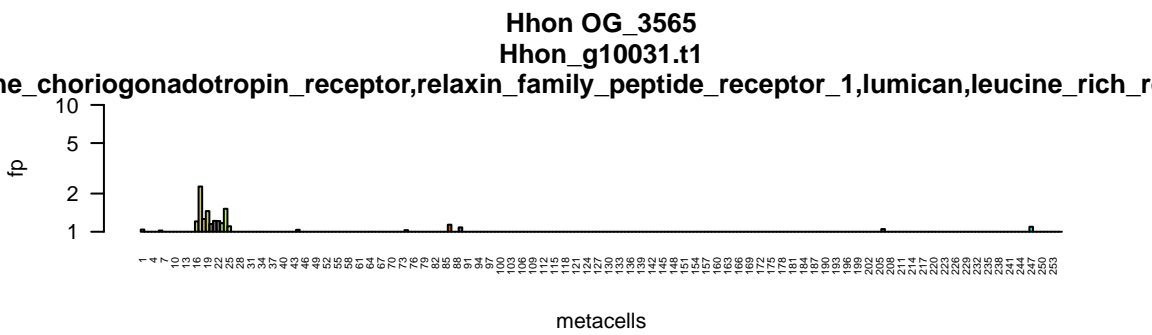
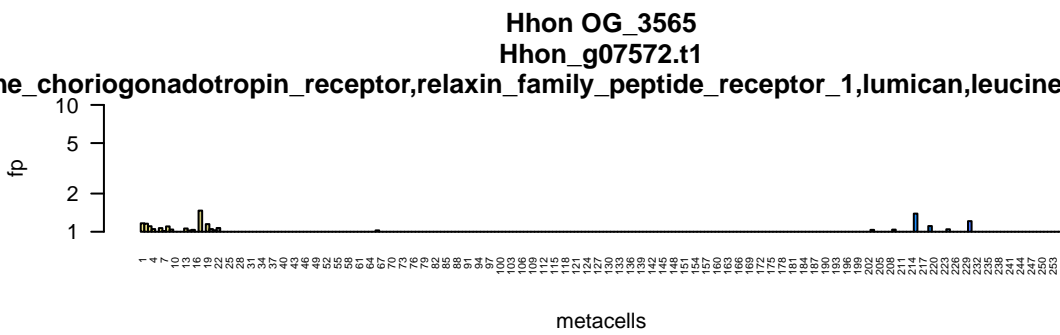
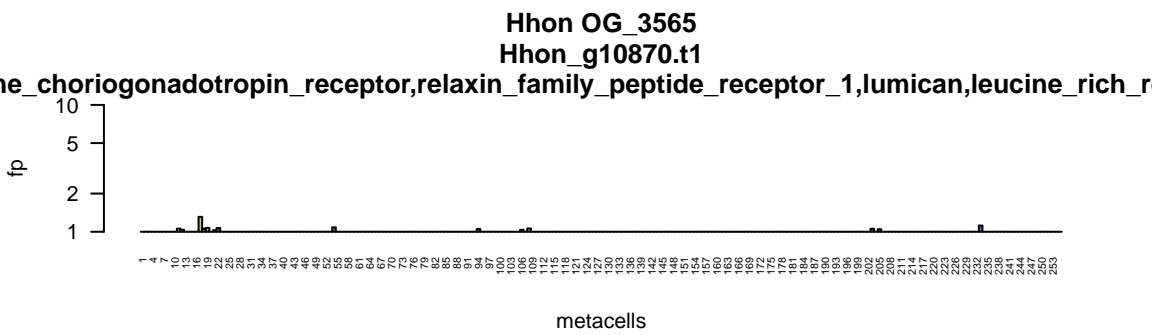
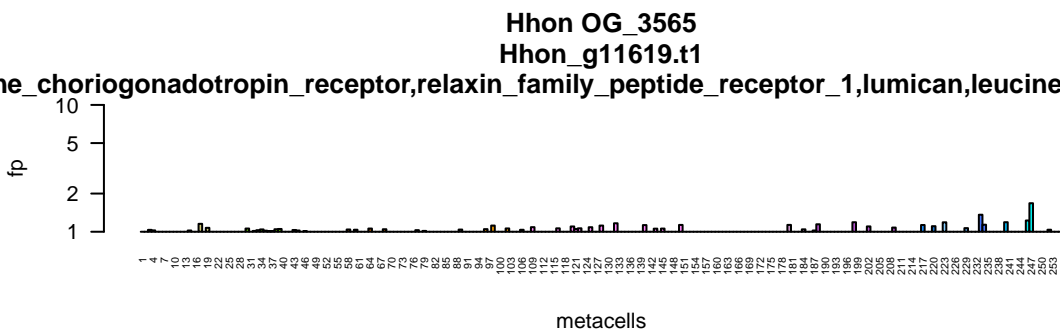
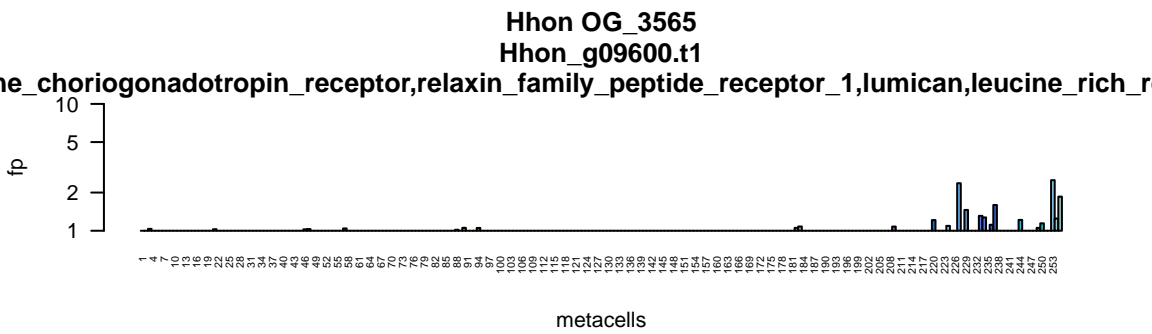
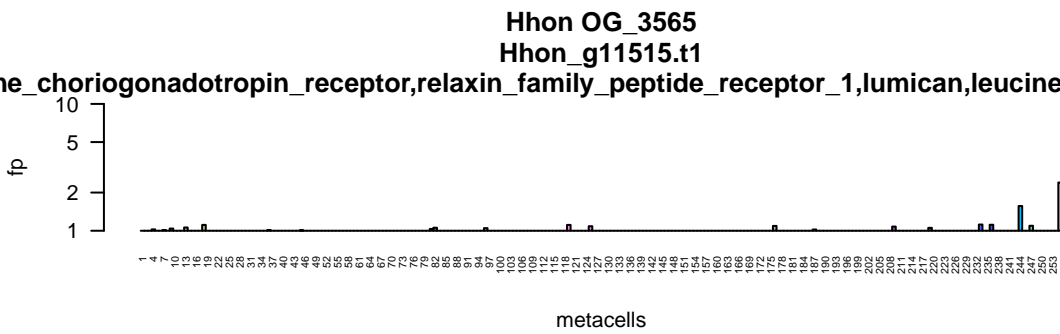
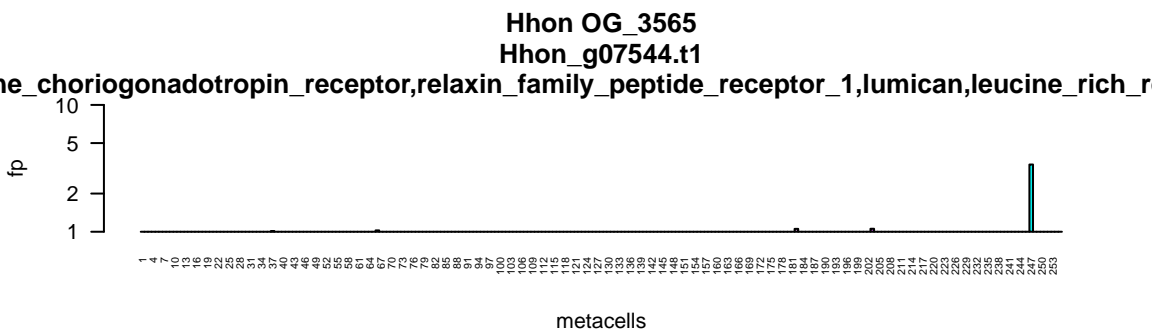
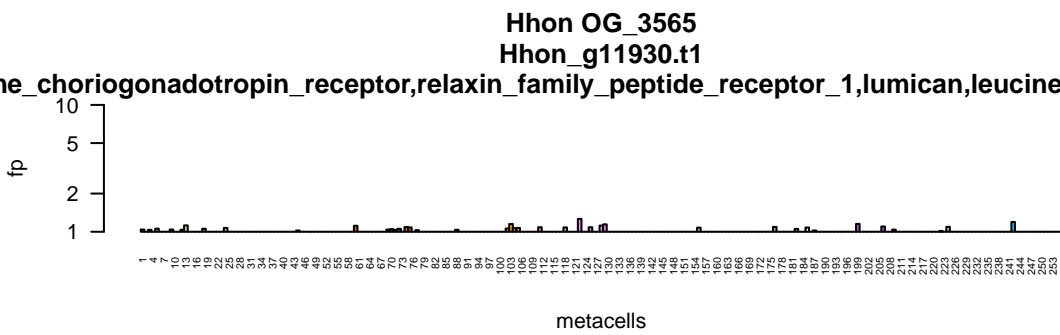
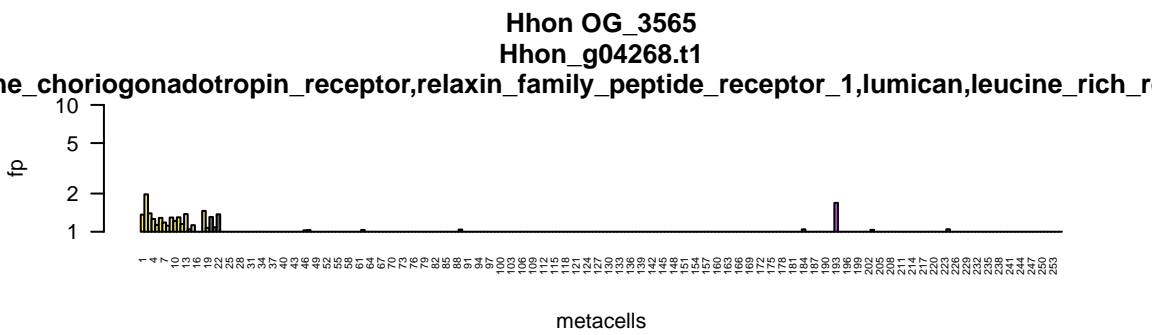
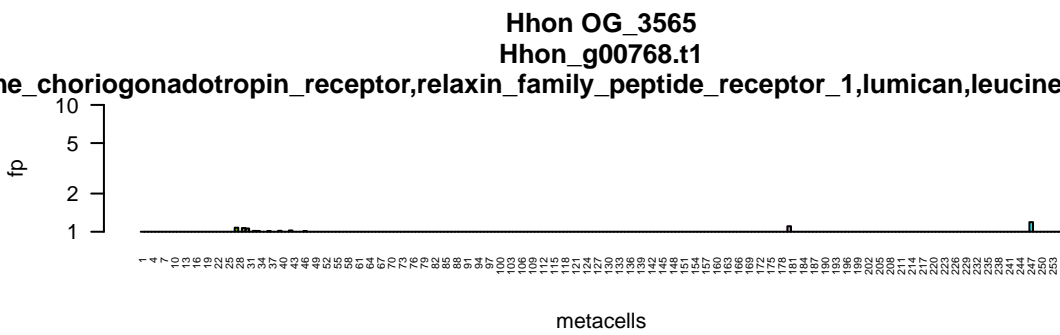
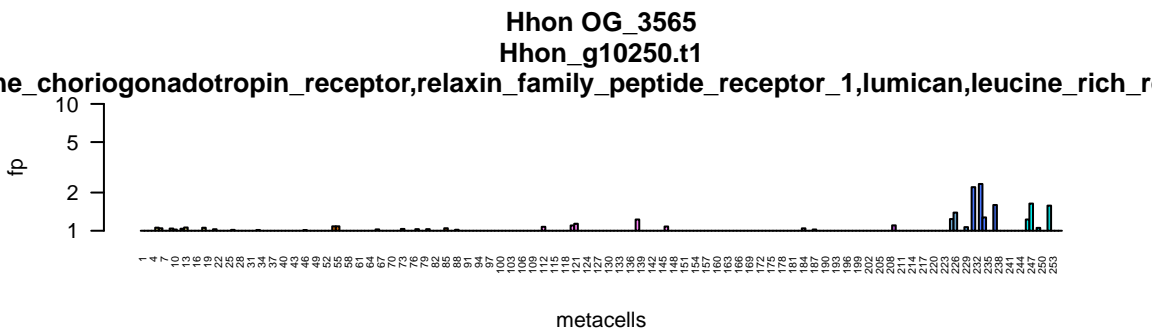
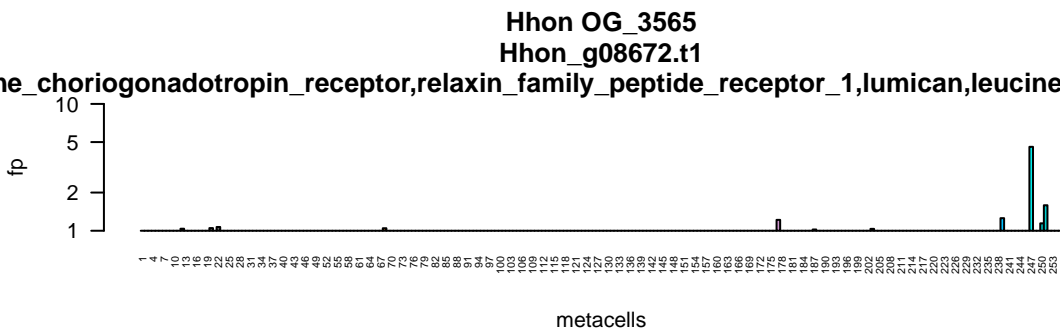


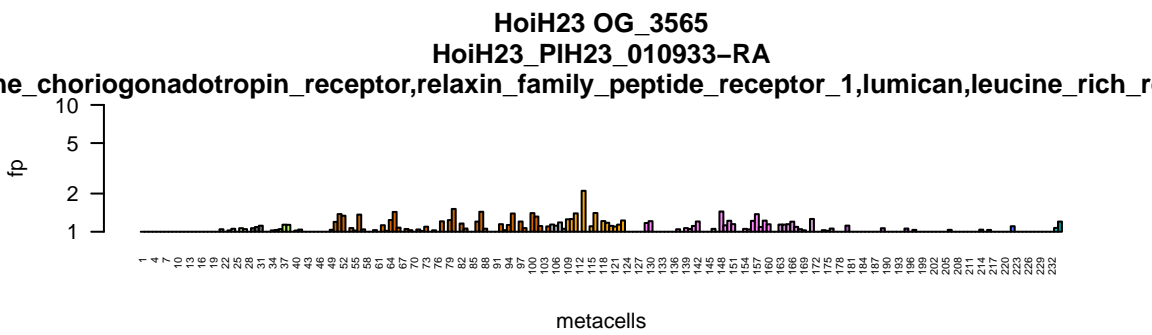
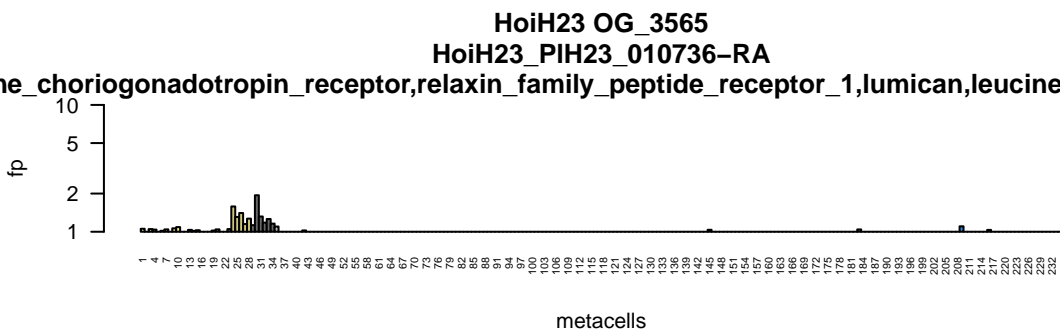
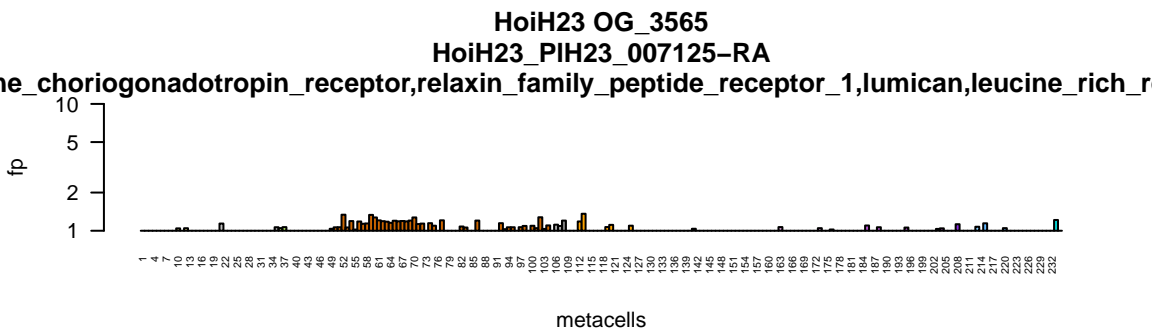
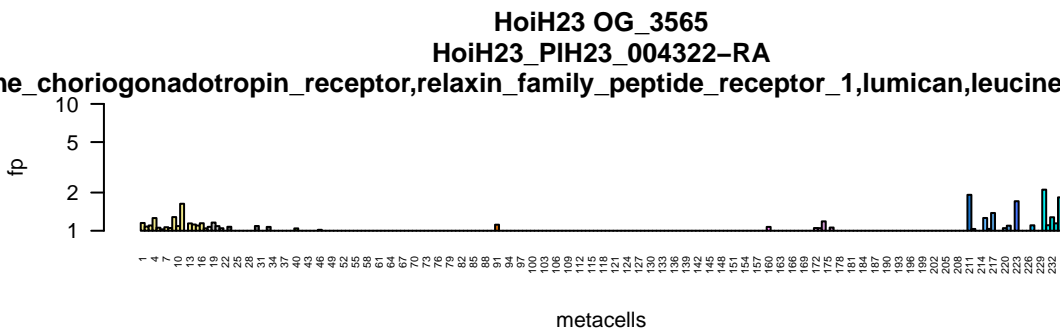
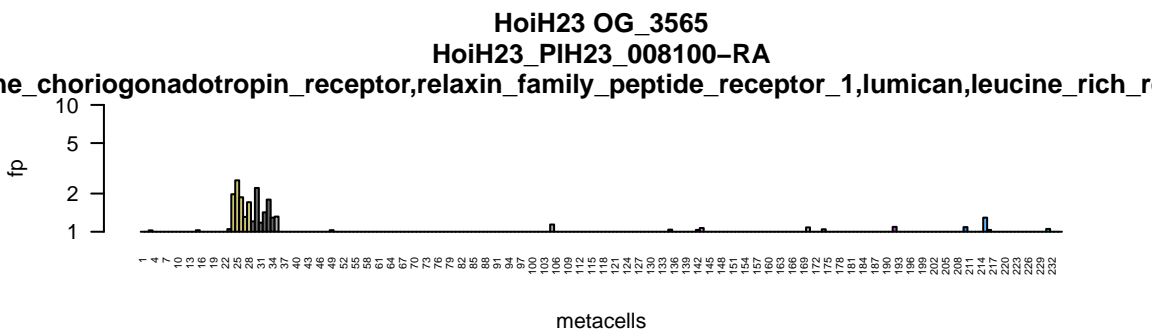
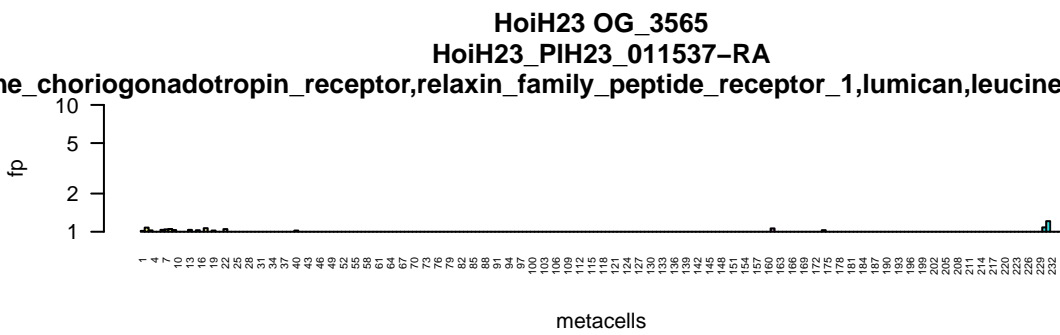
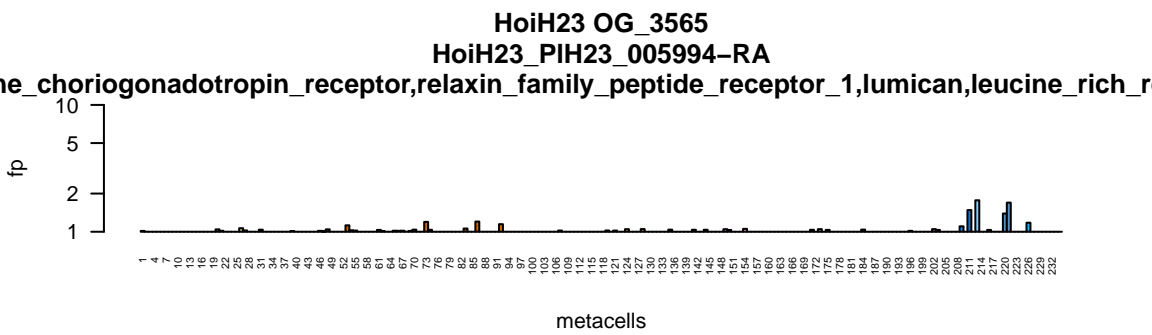
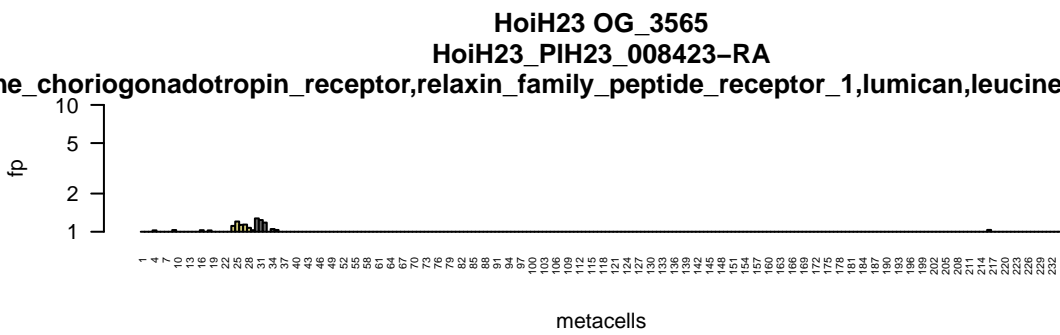
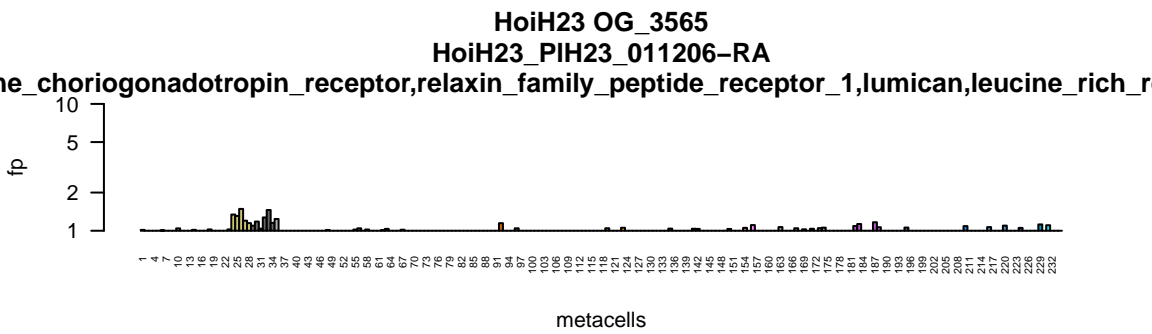
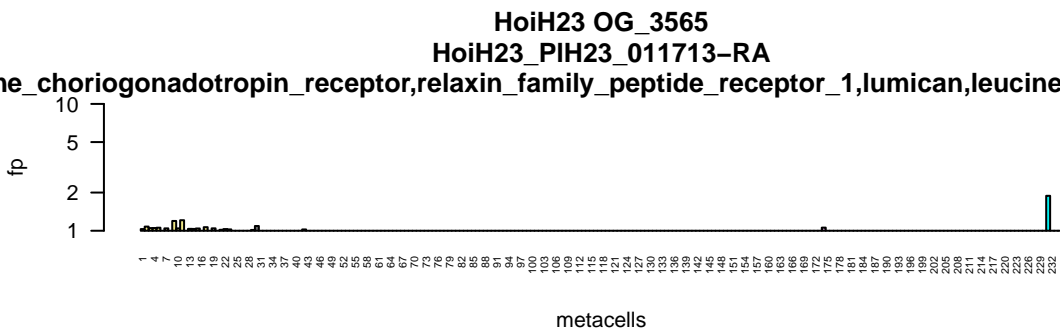
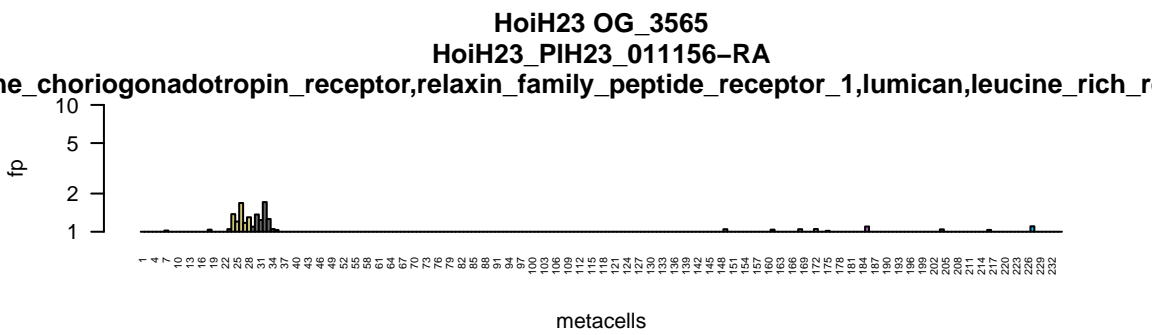
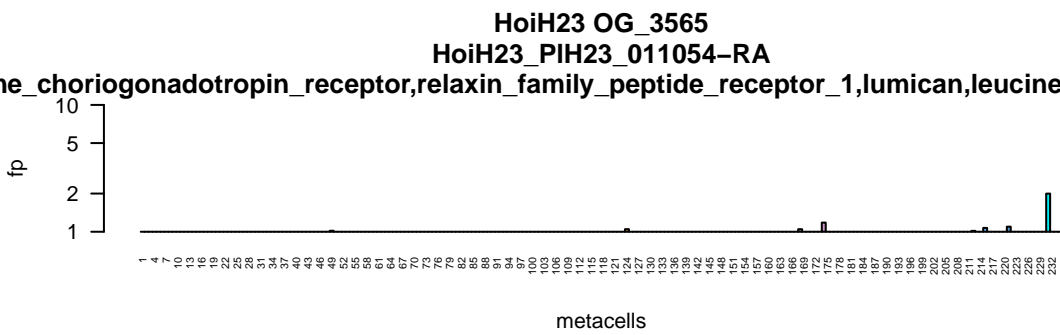
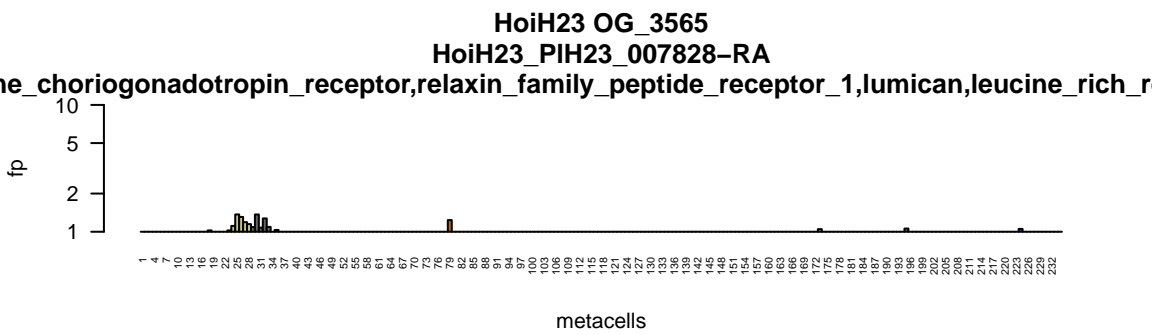
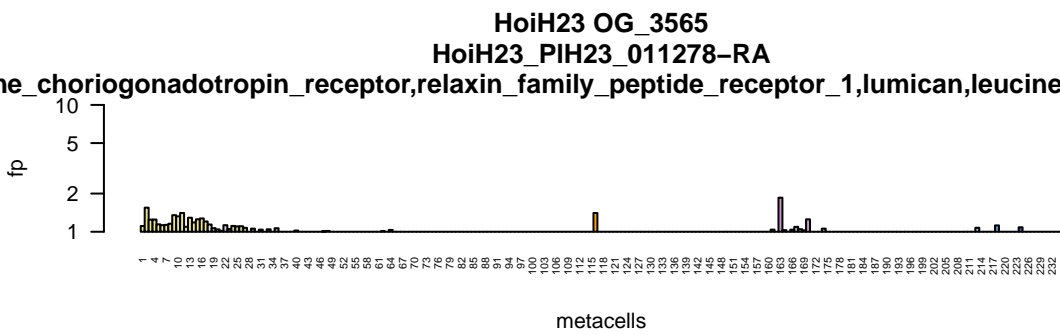
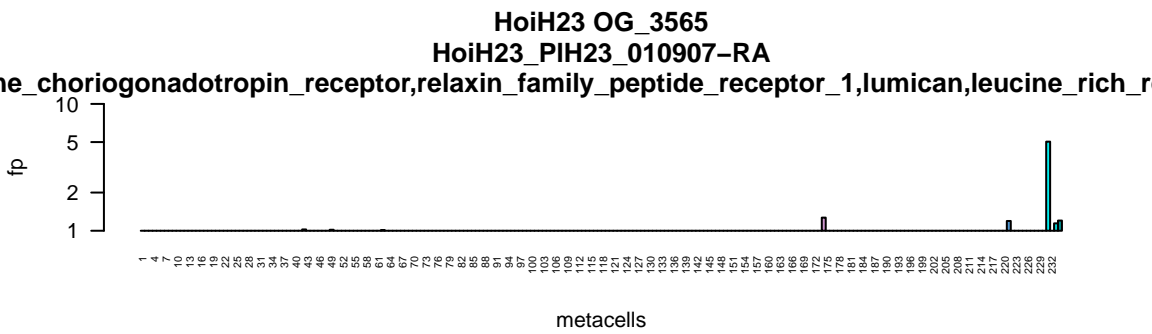
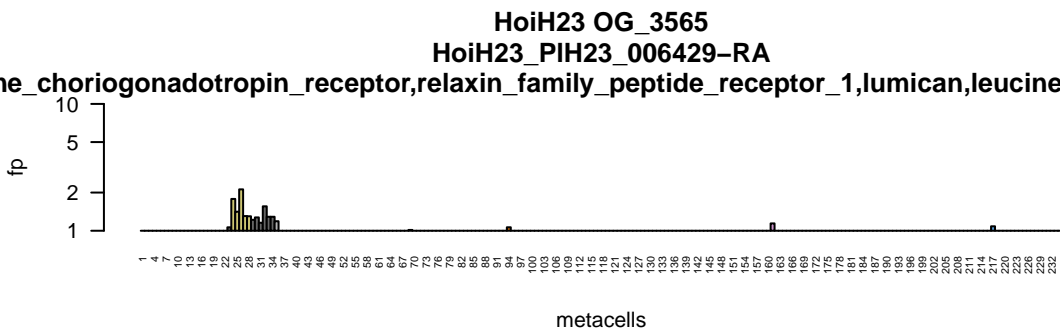


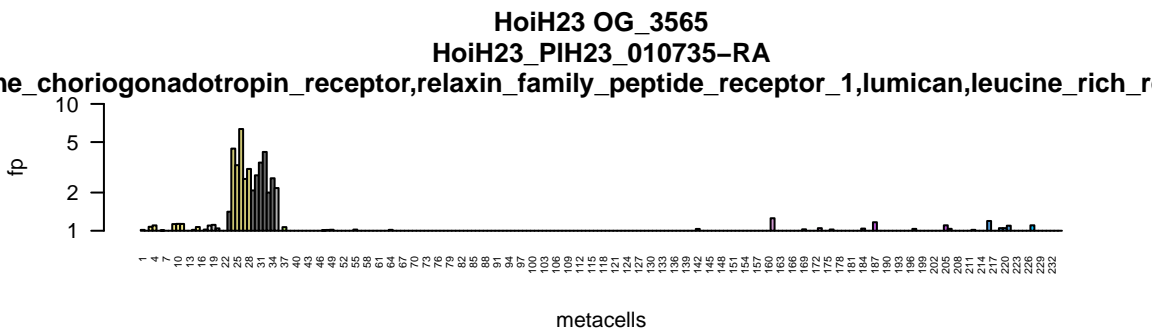
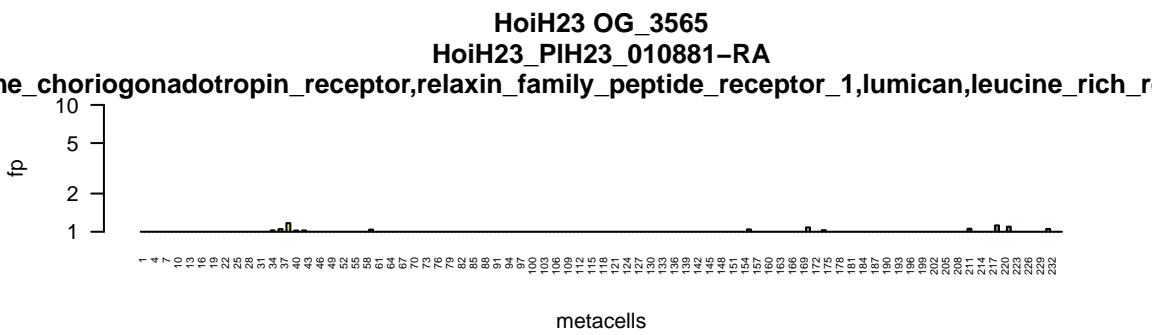
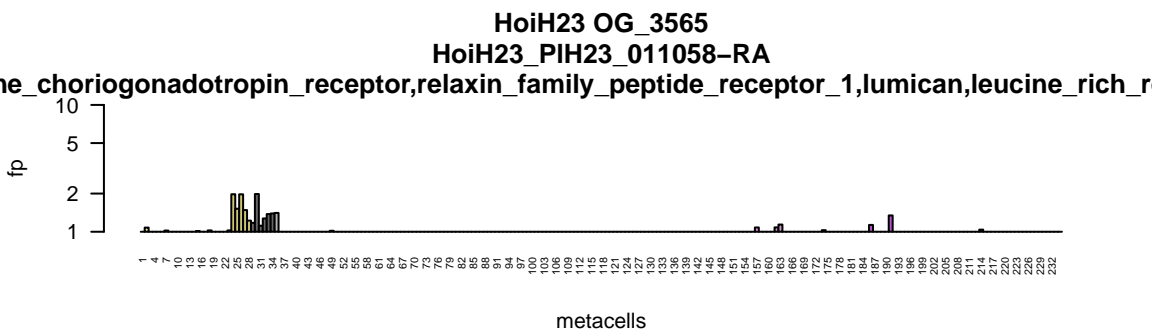
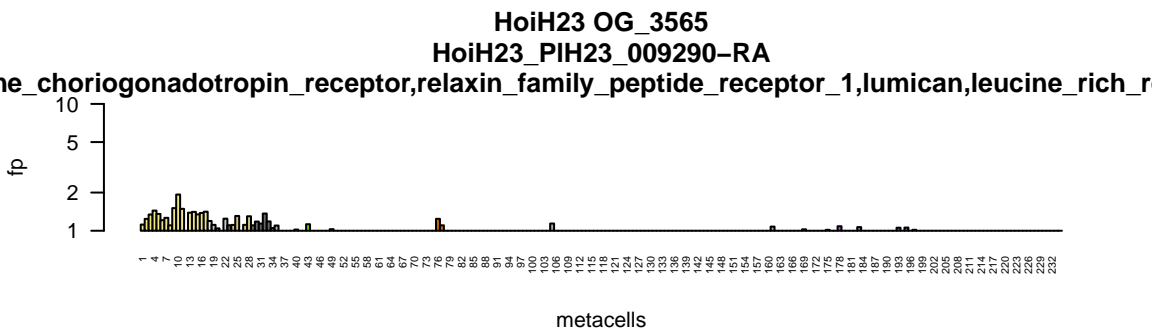


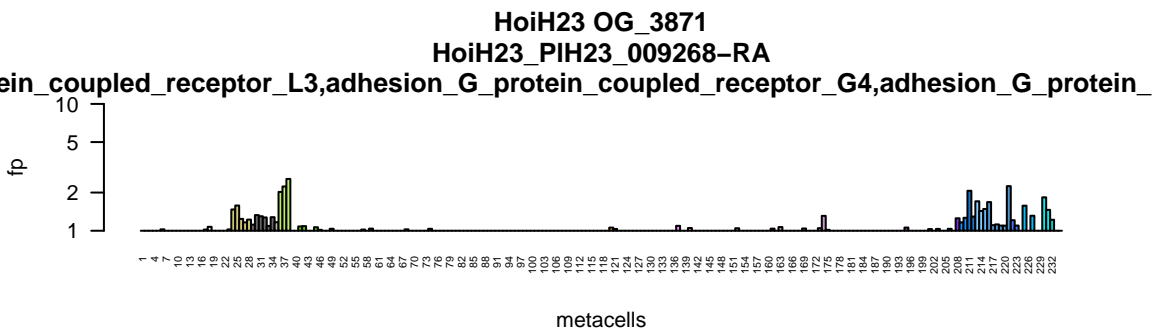
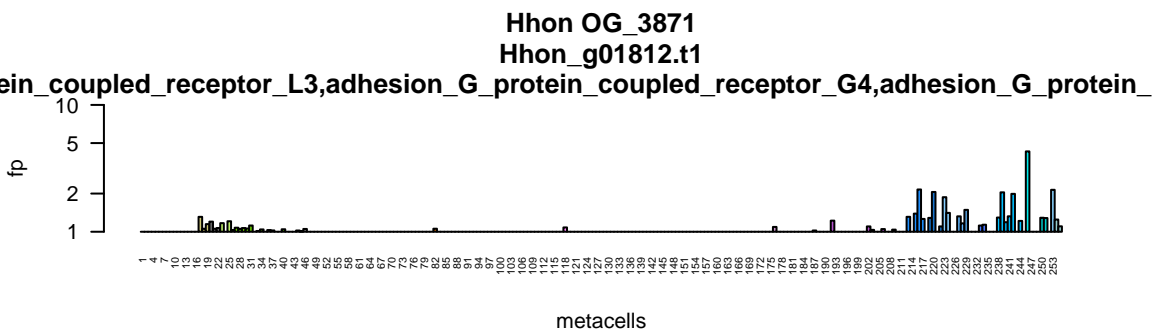
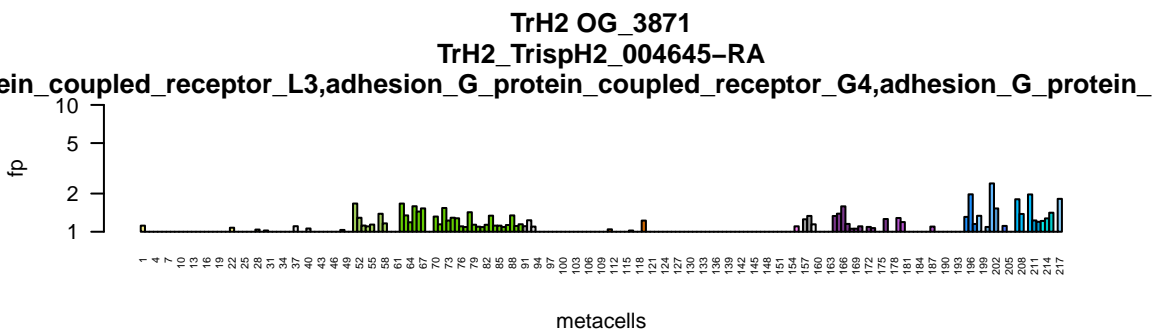
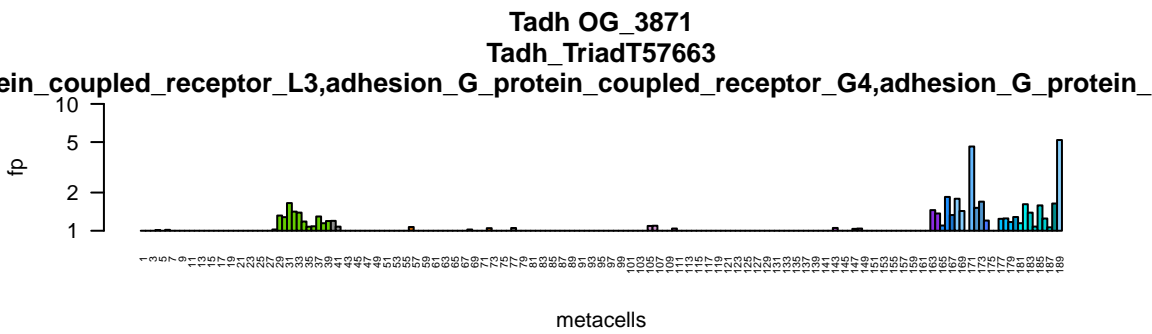


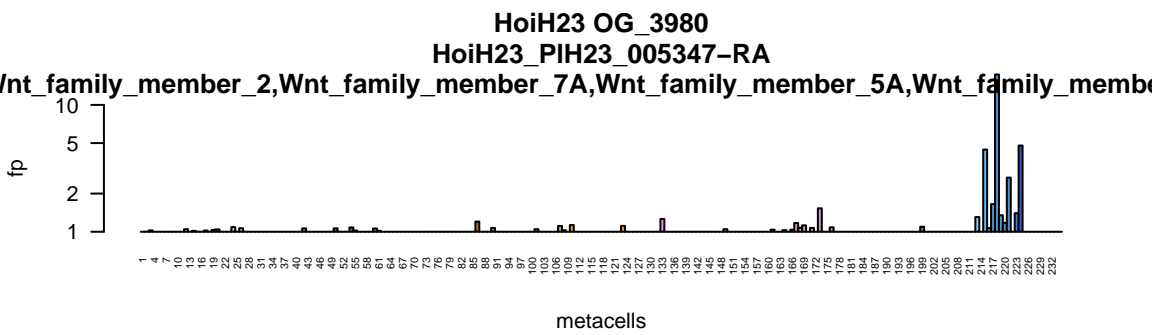
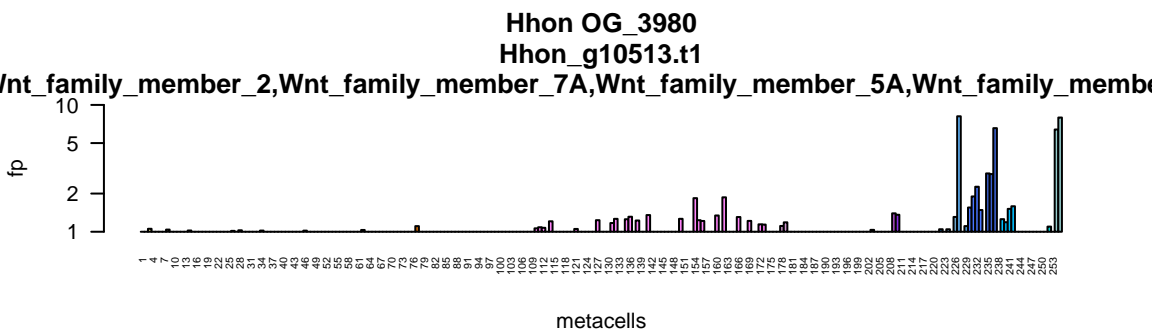
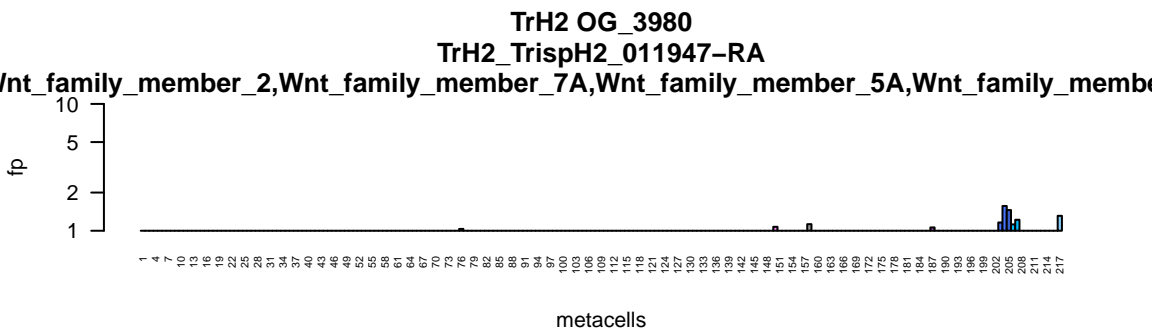
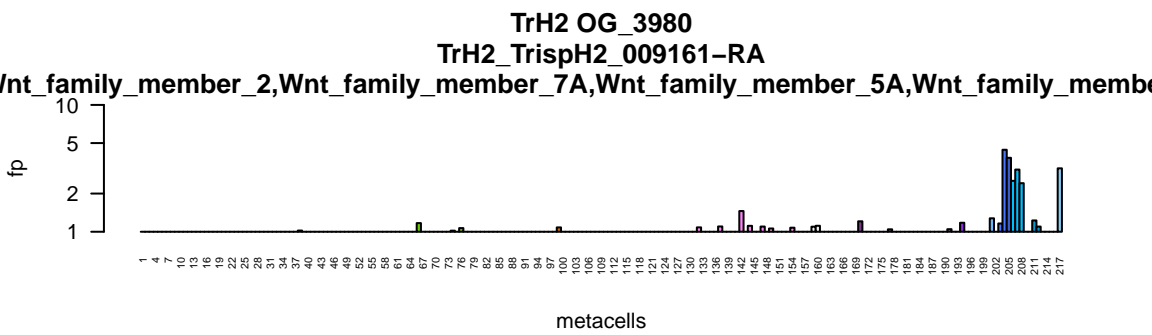
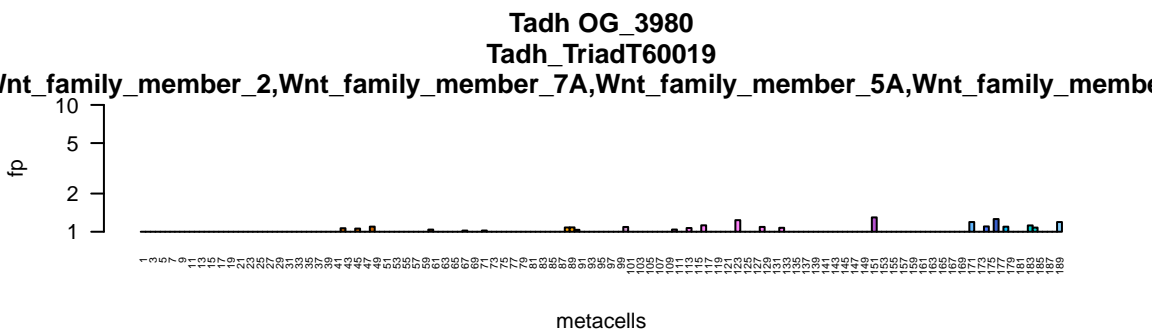
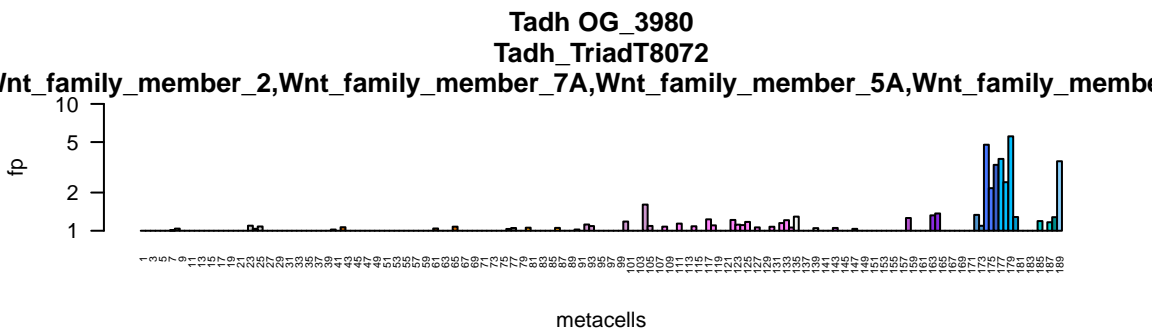


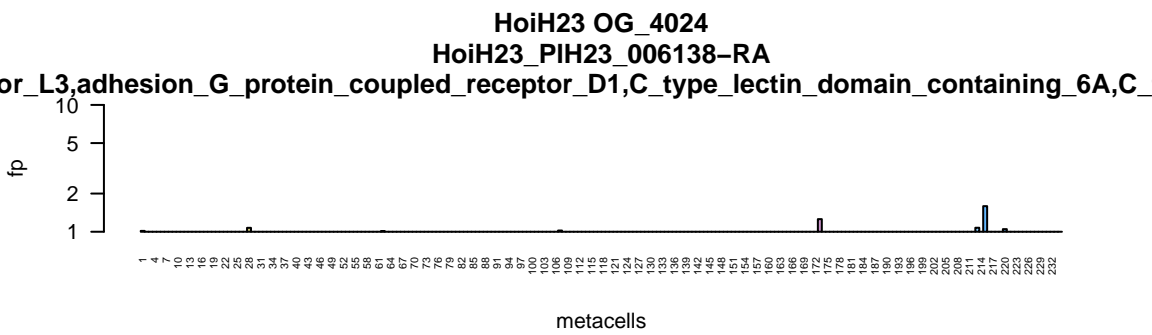
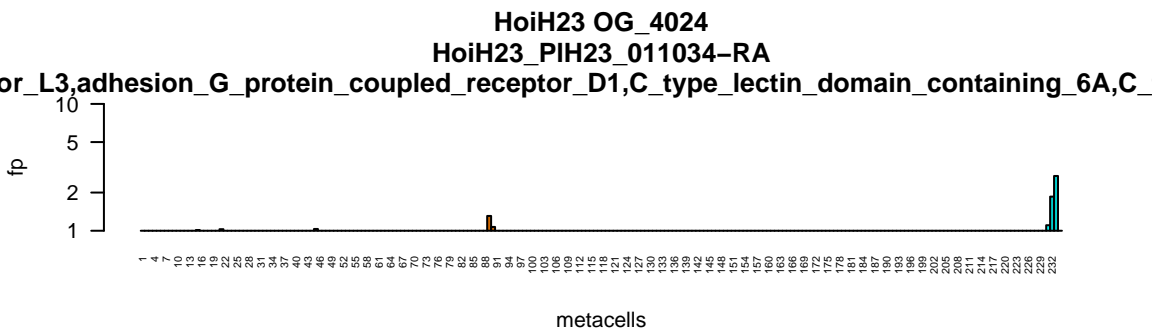
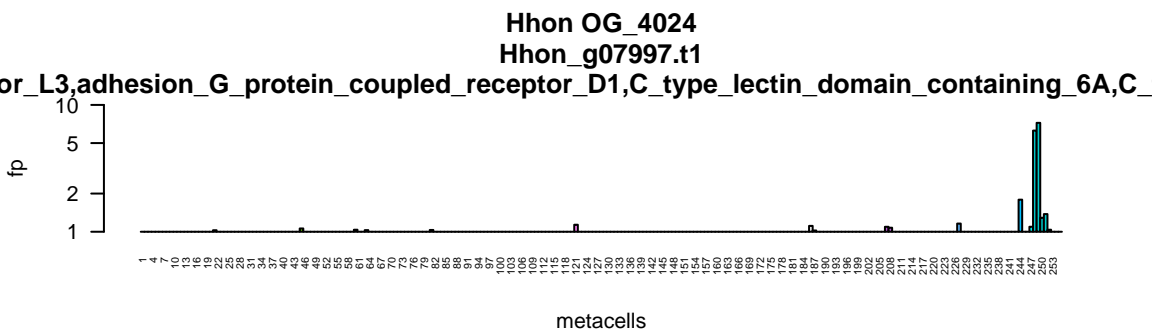
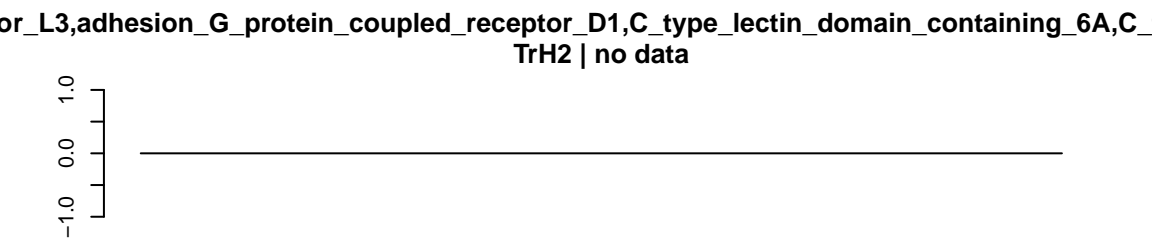
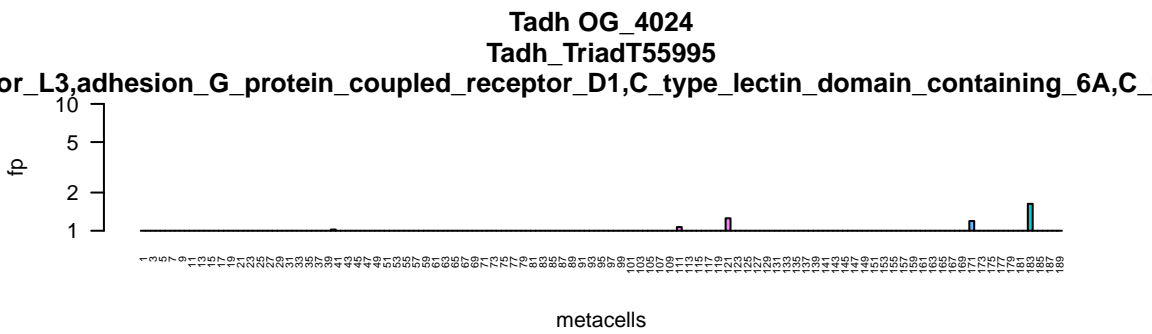


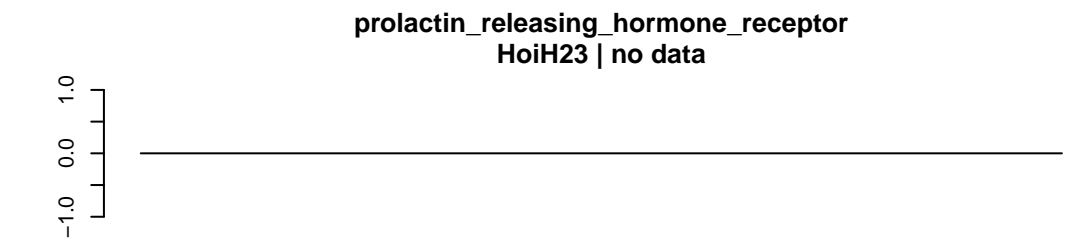
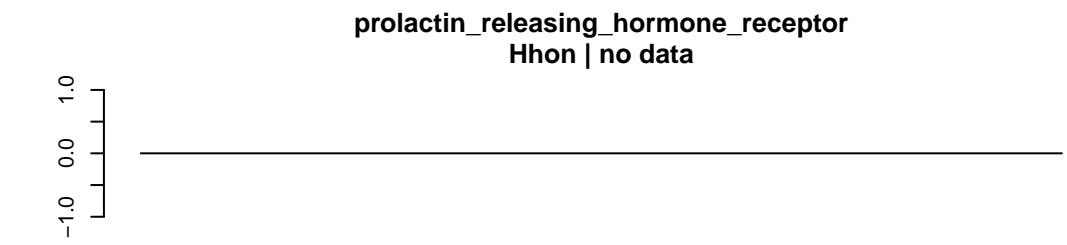
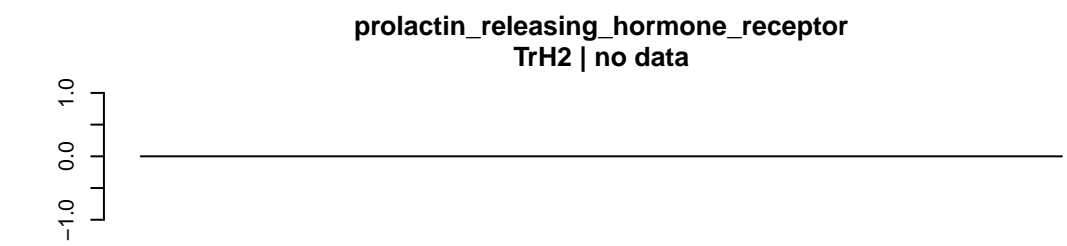
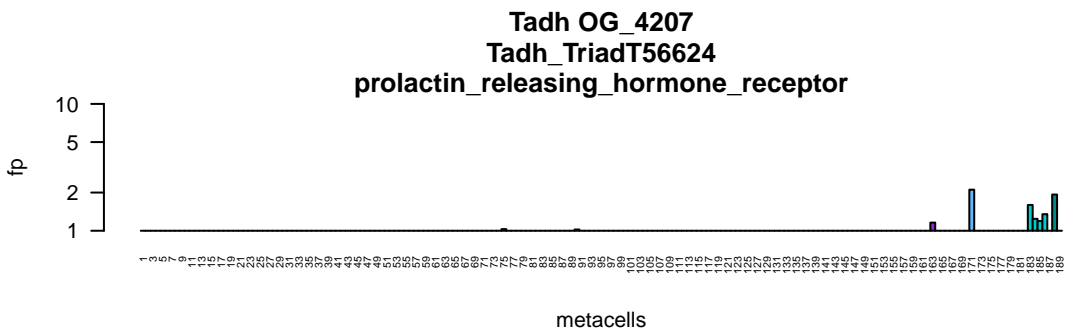


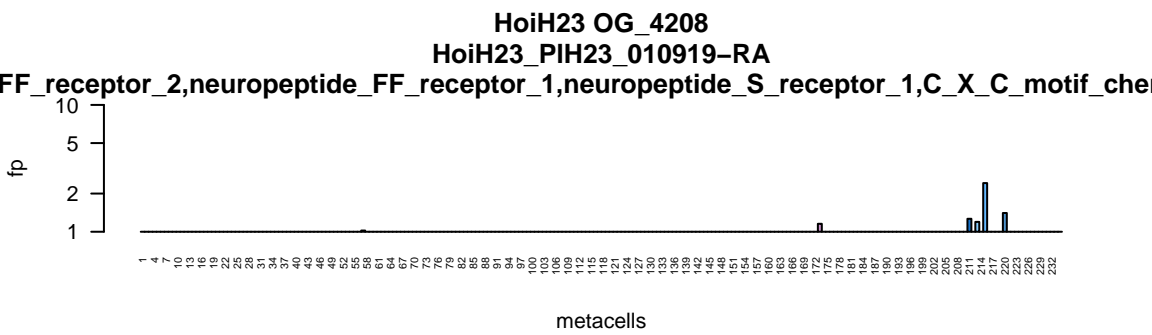
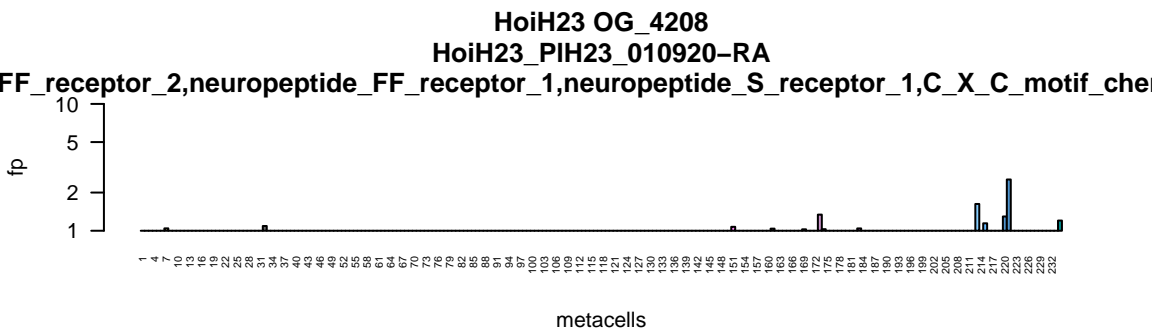
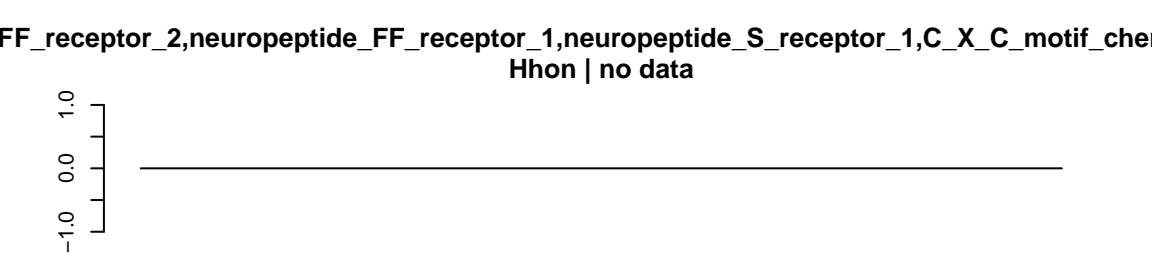
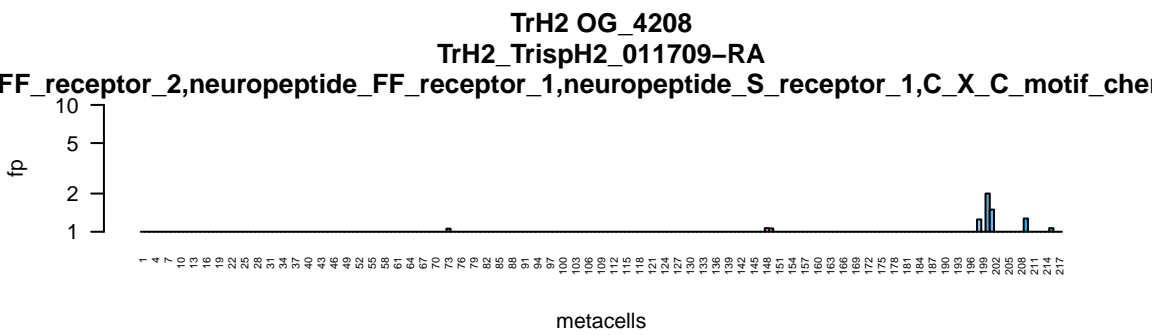
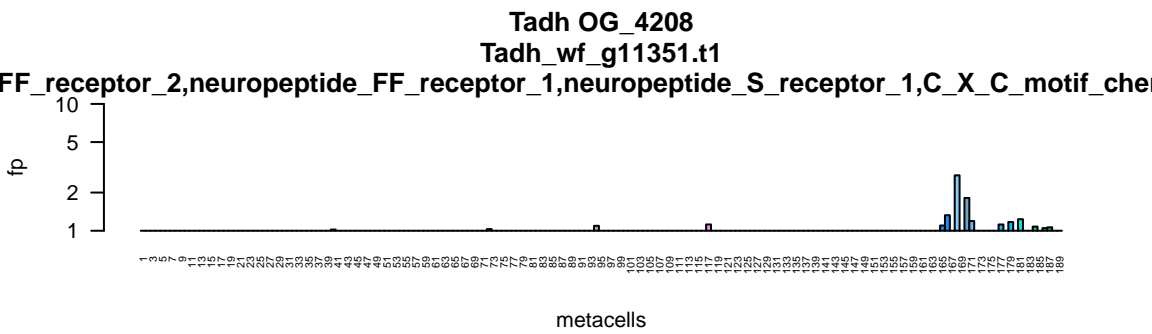


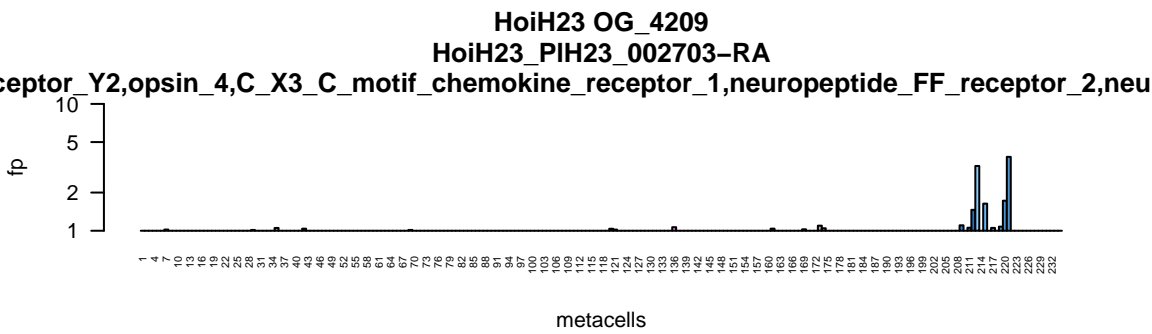
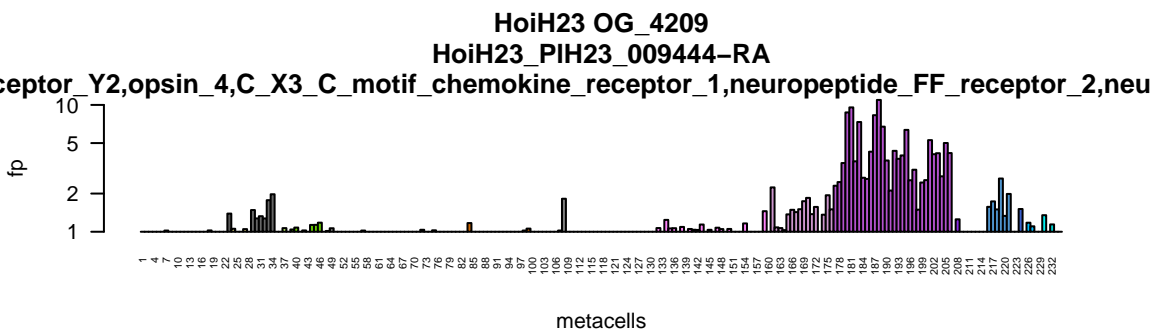
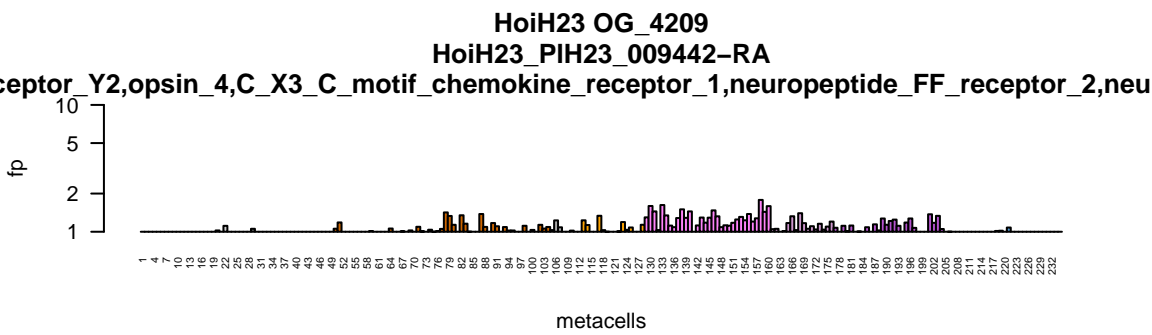
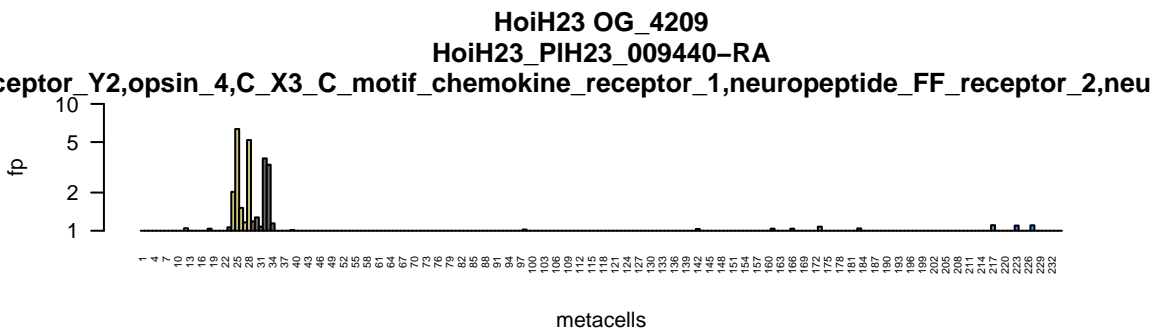
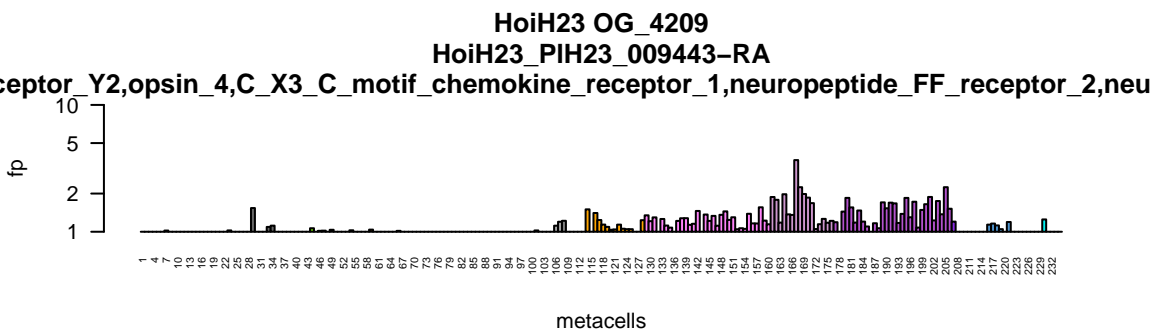
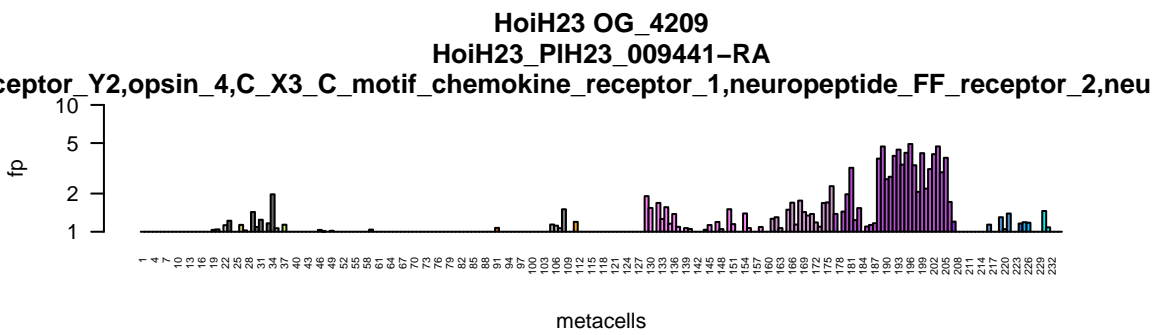
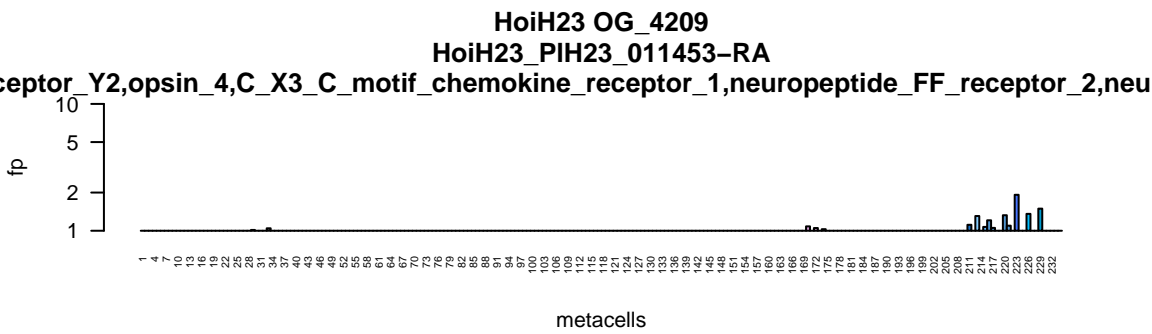


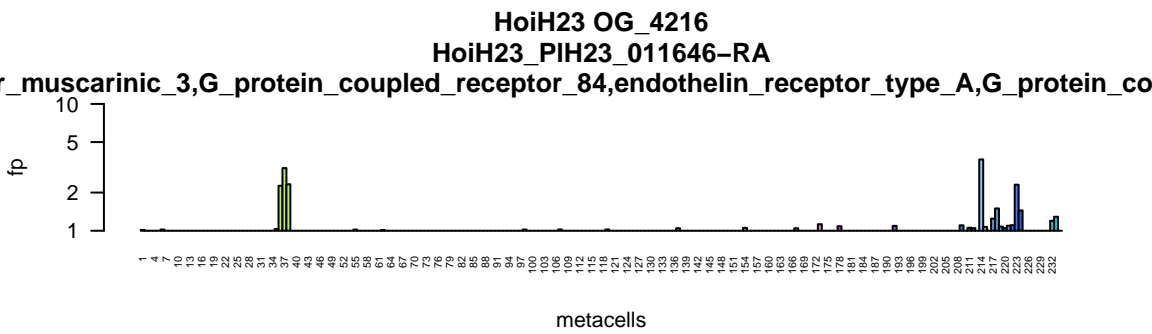
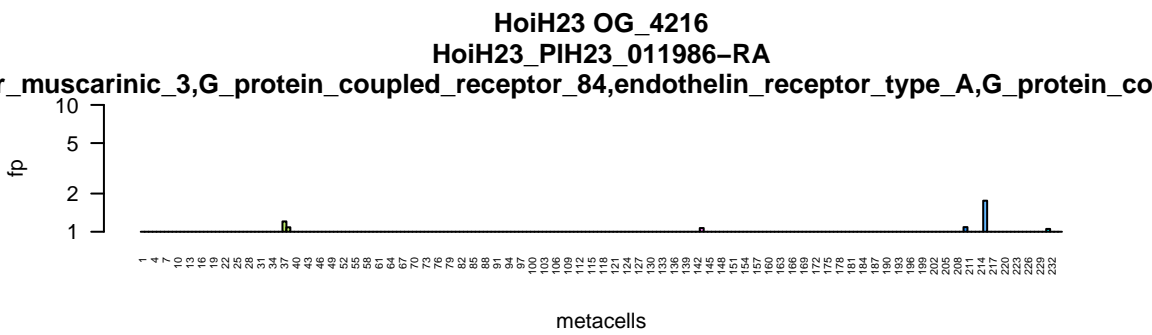
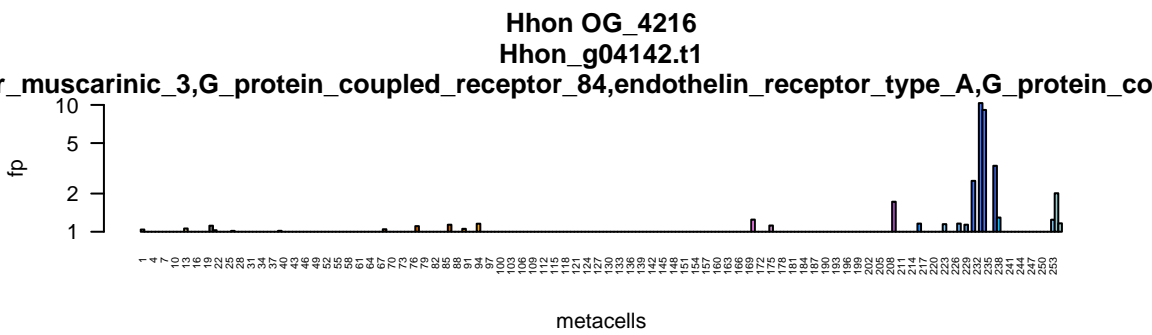
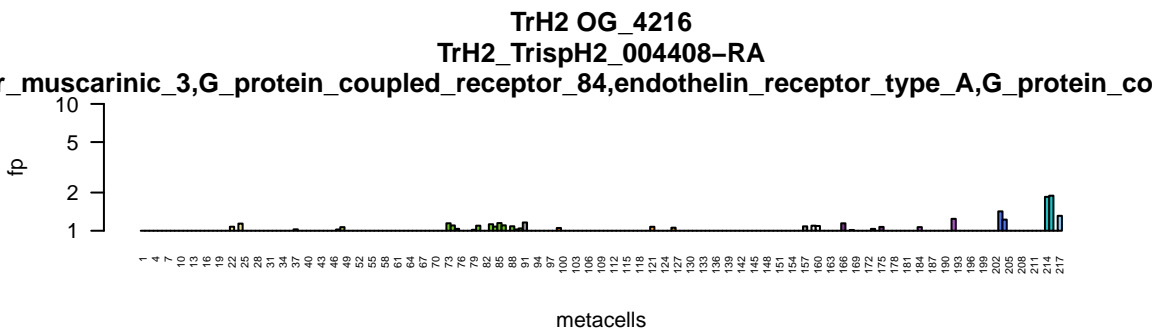
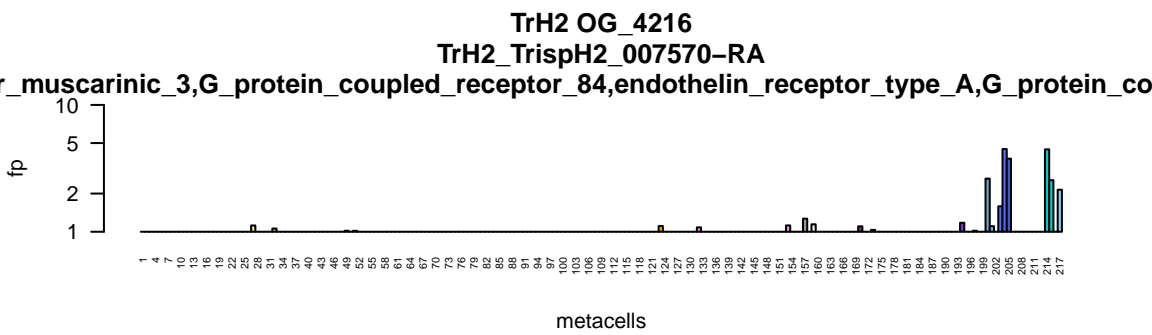
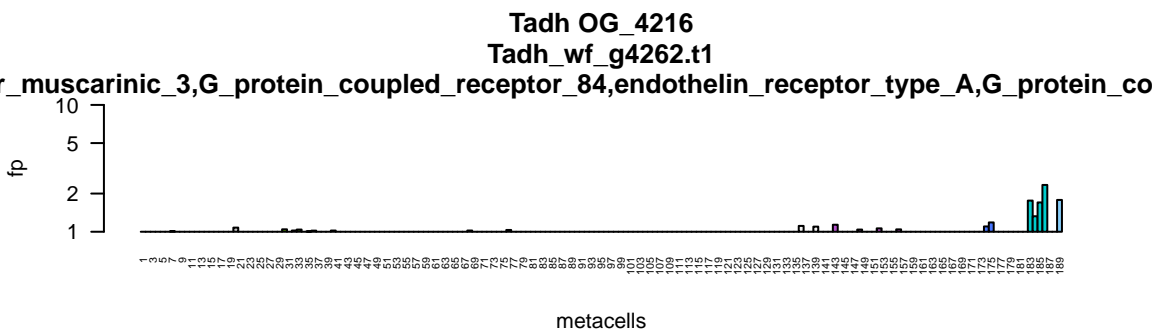
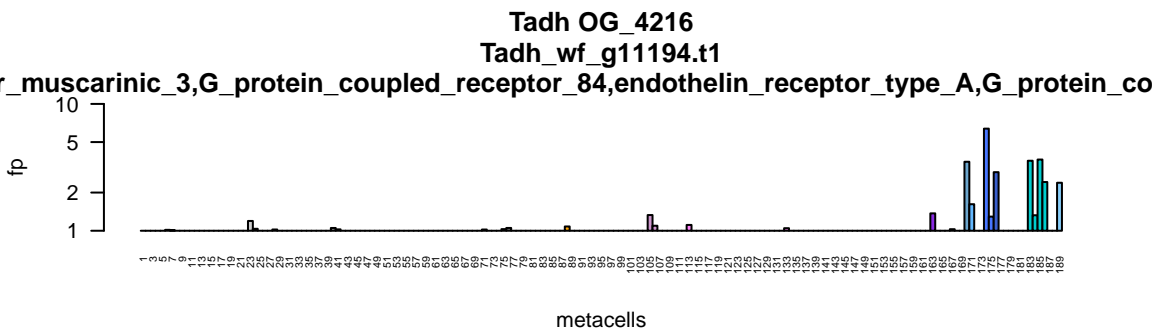




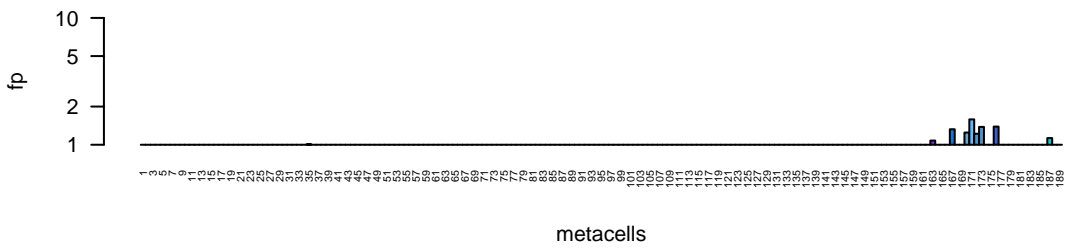






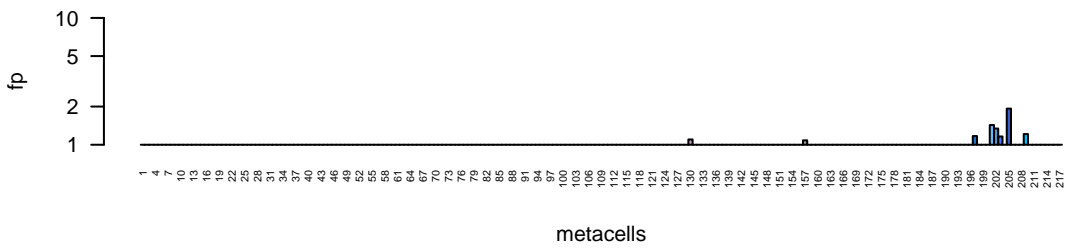


Tadh OG_4270
Tadh_wf_g4875.t1



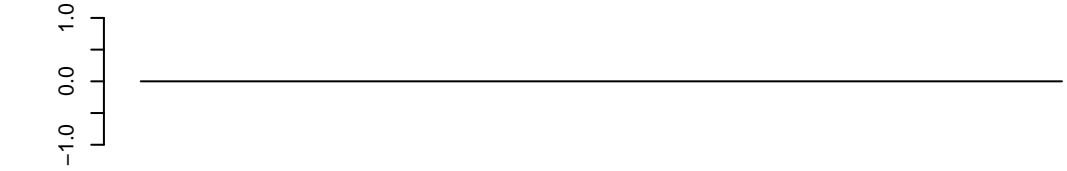
metacells

TrH2 OG_4270
TrH2_TrispH2_008489-RA

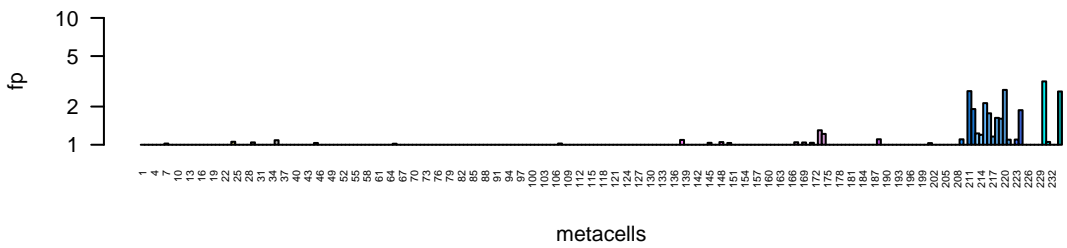


metacells

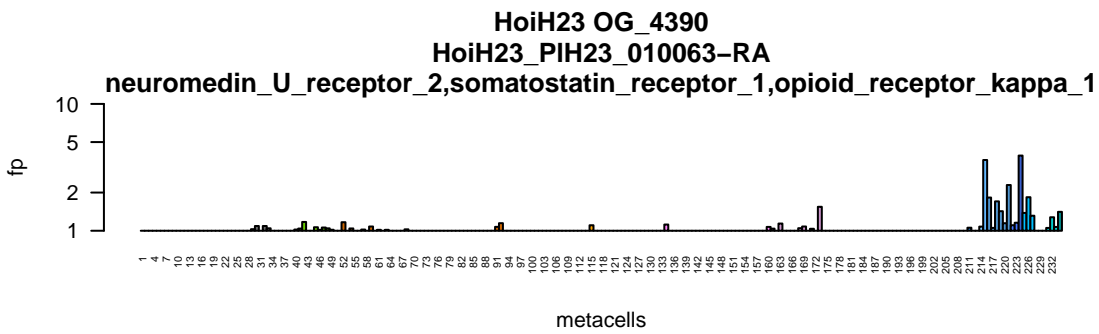
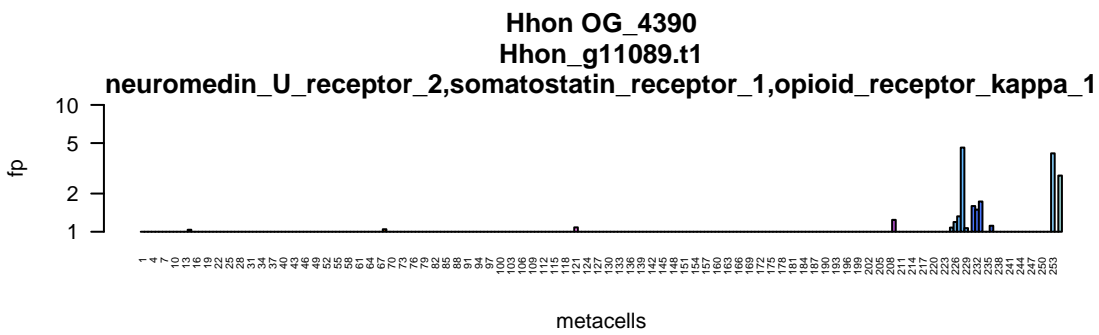
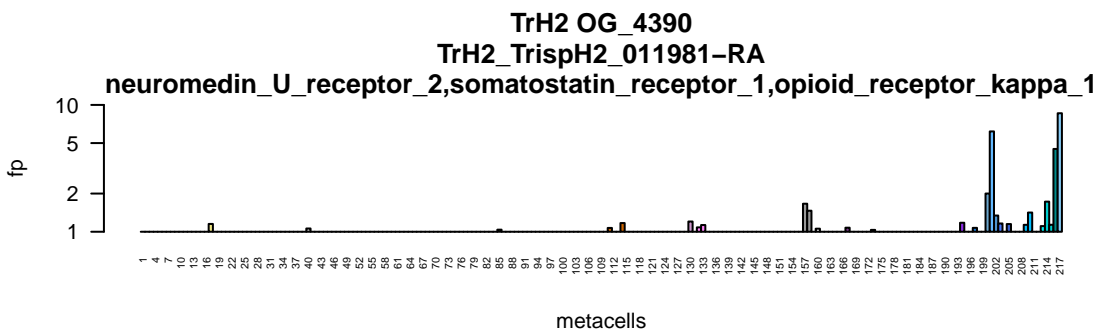
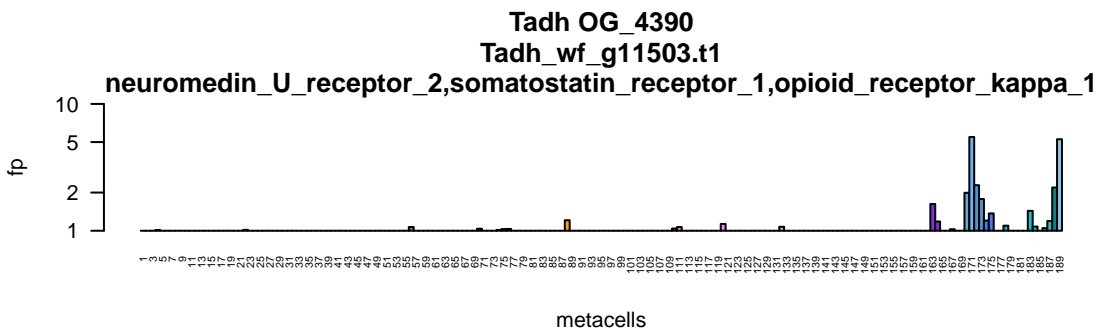
Hhon | no data

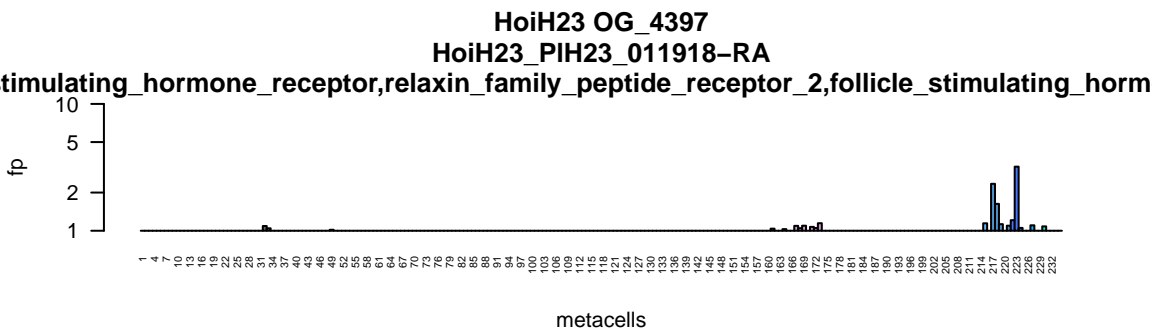
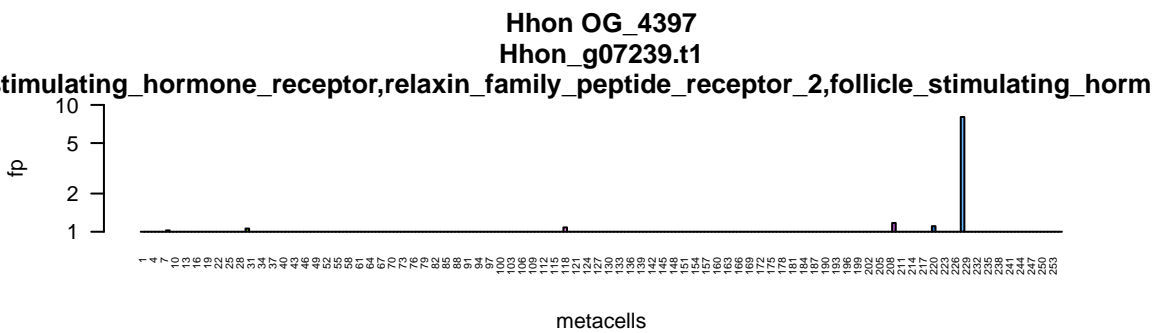
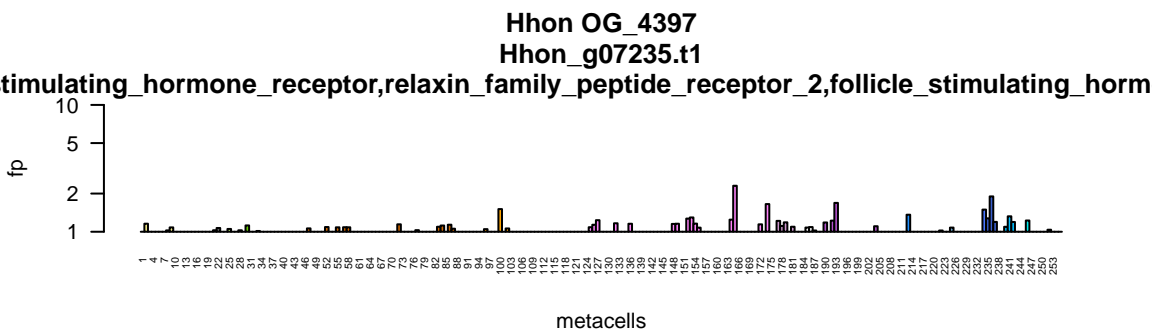
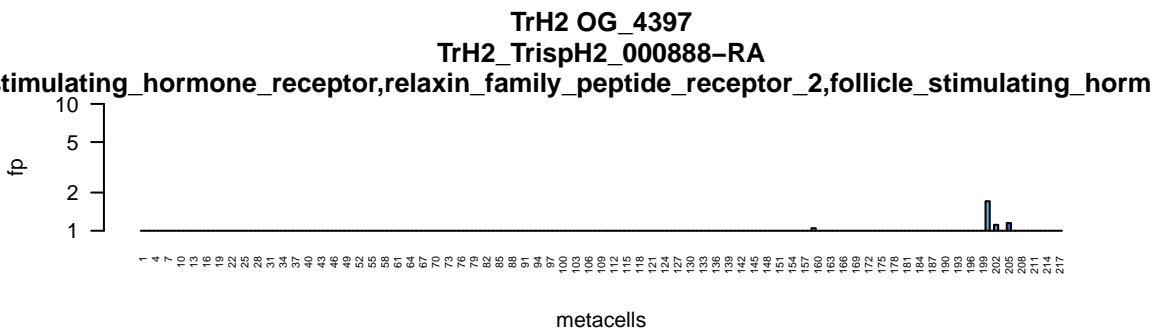
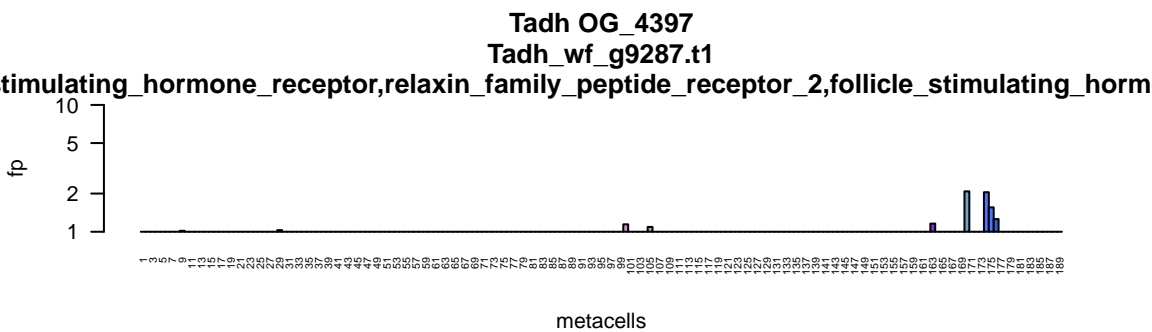
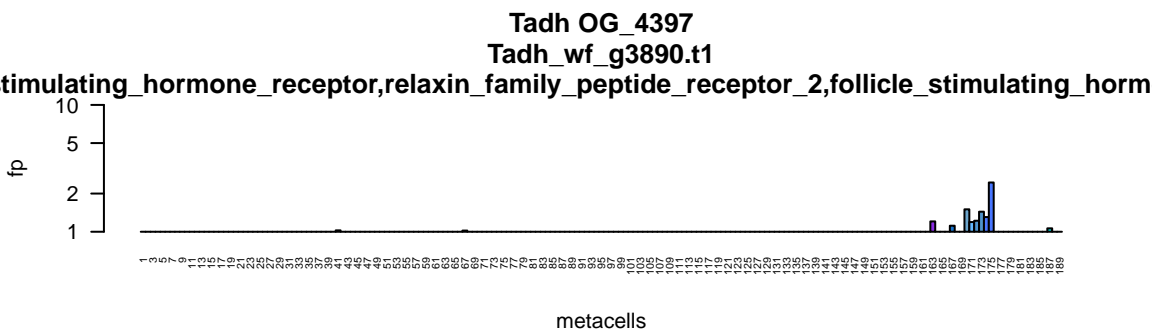
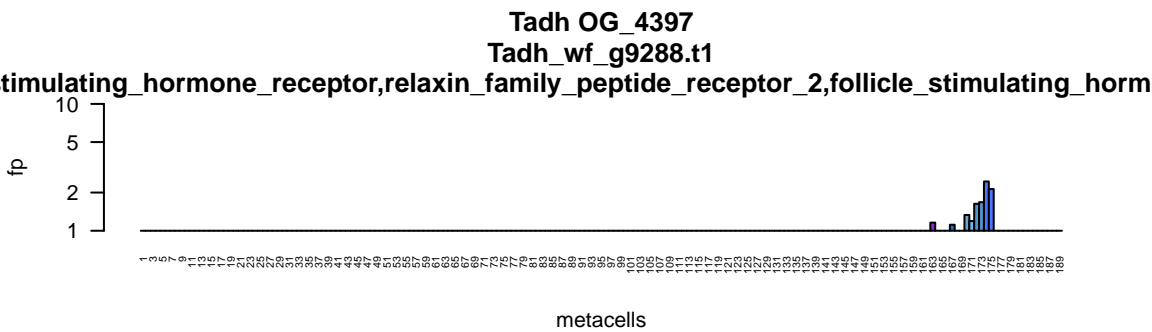


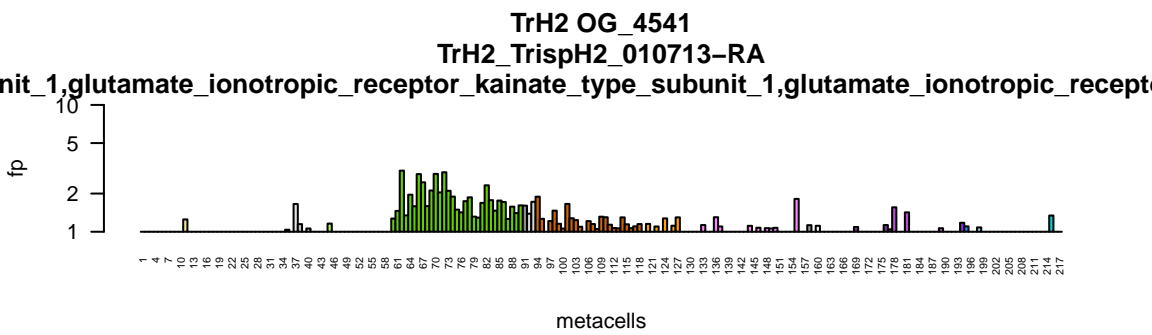
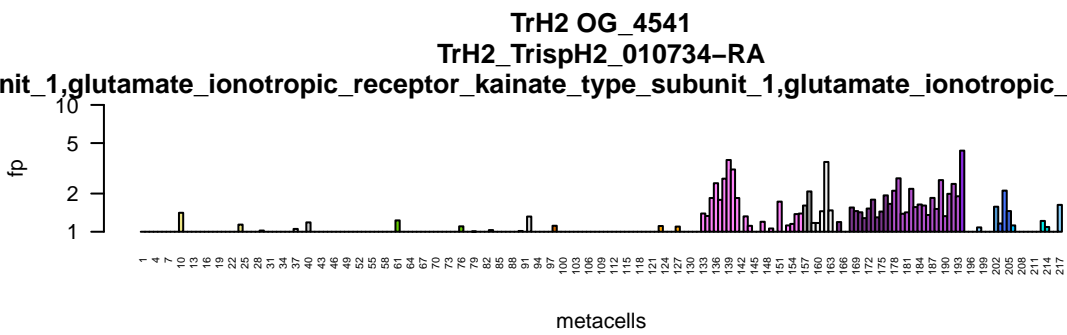
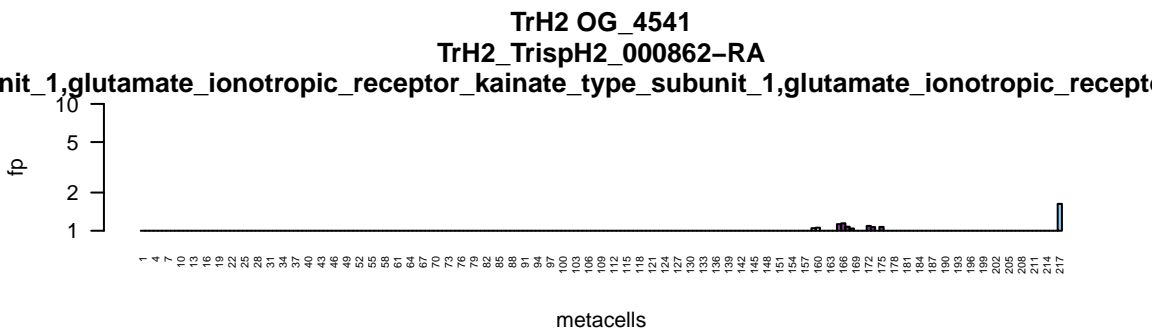
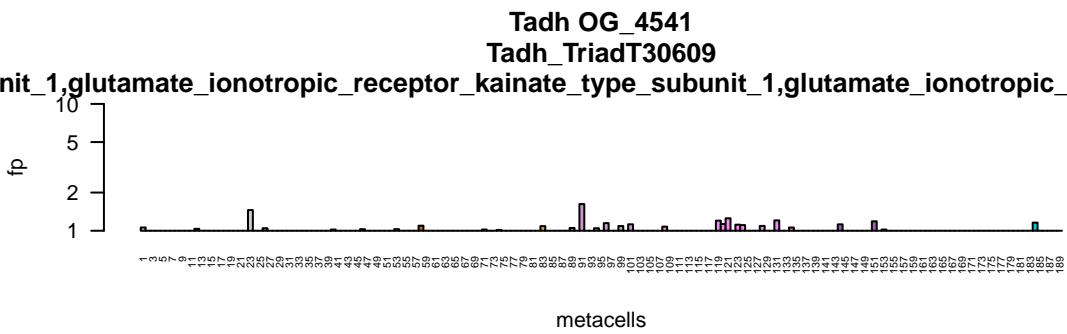
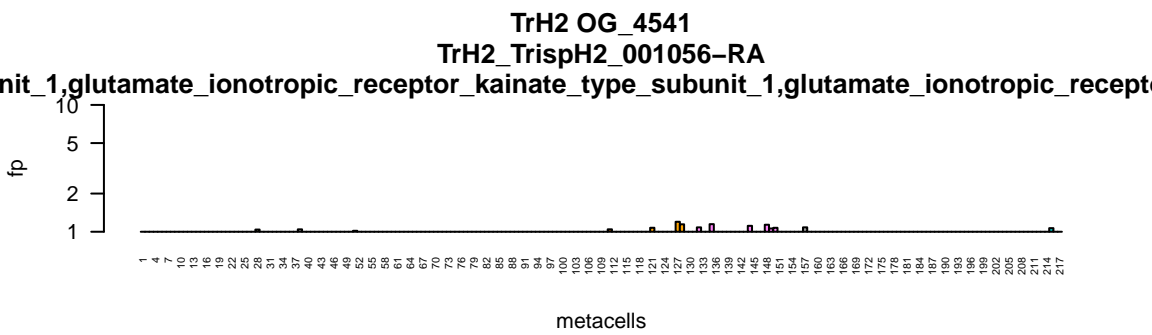
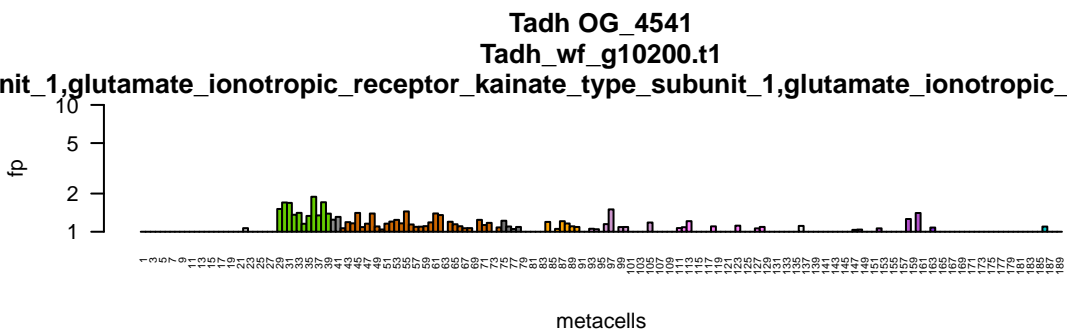
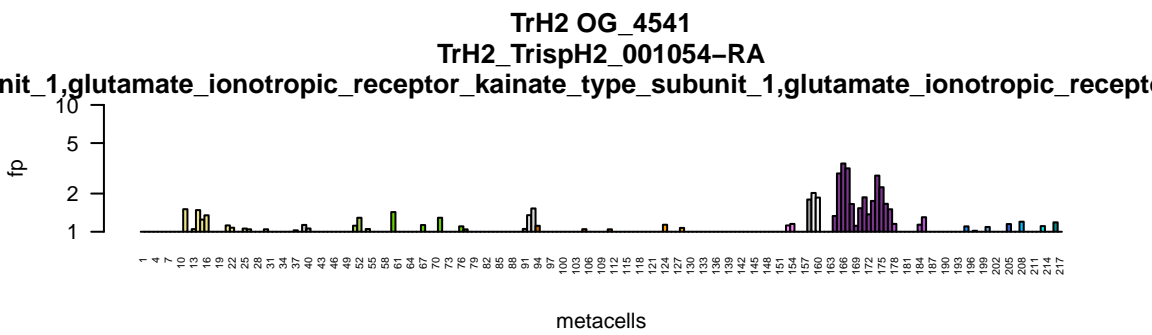
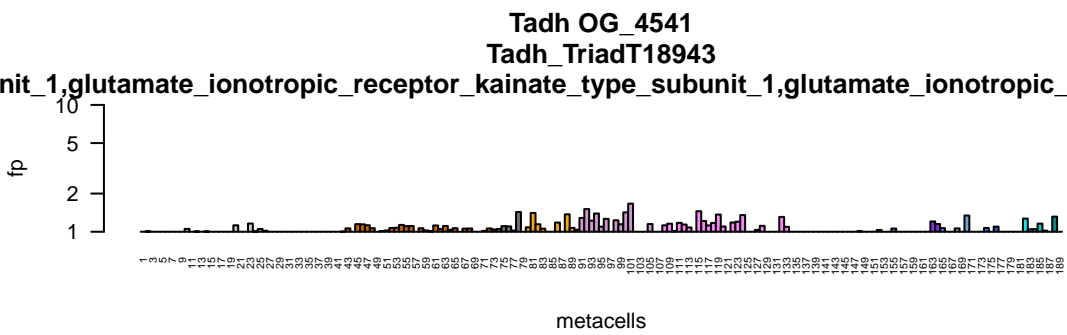
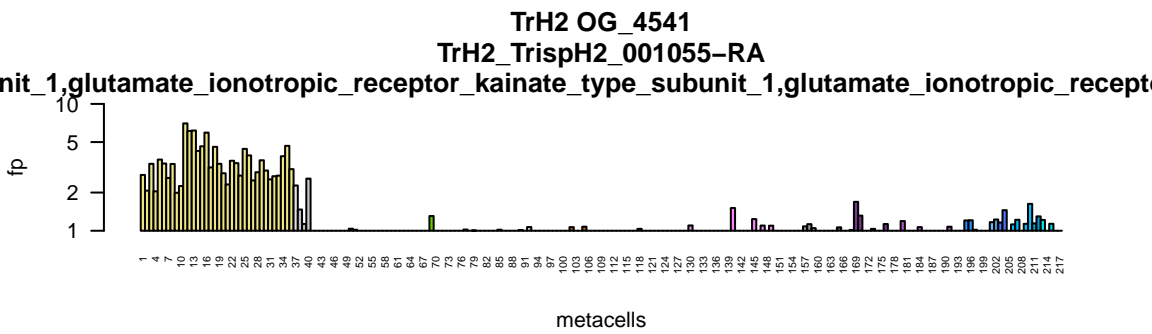
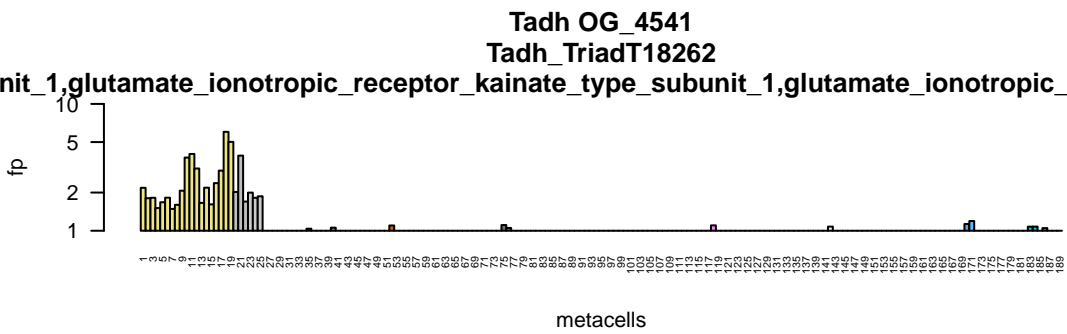
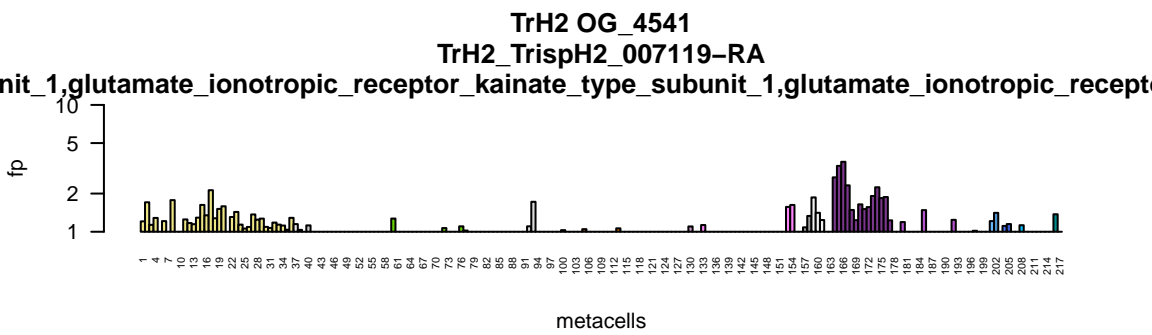
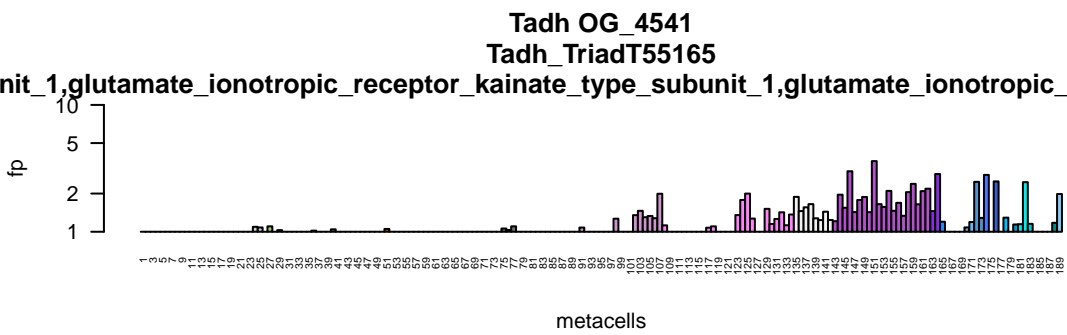
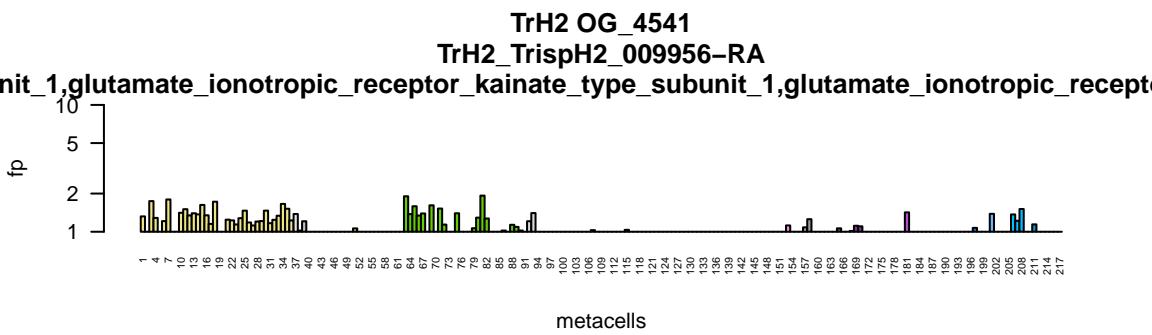
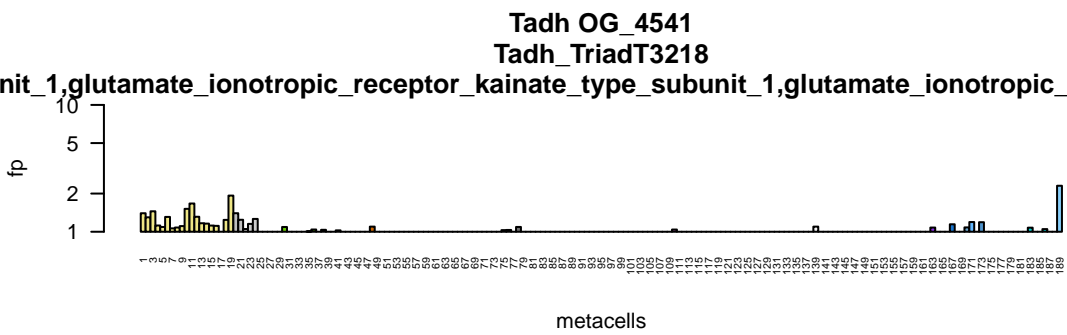
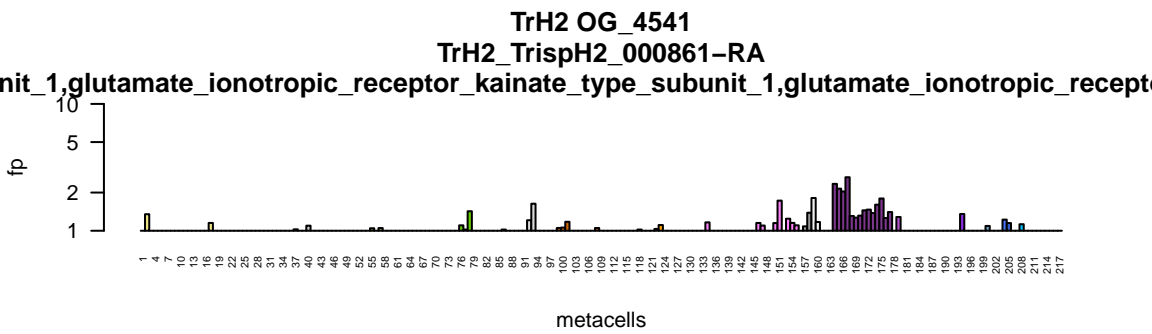
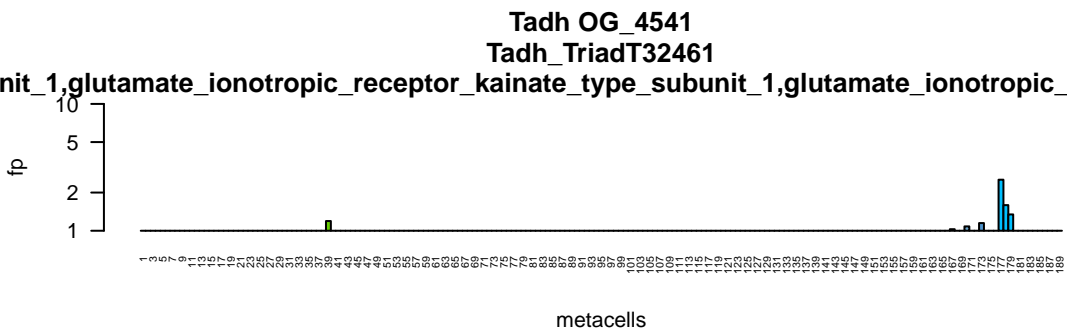
HoiH23 OG_4270
HoiH23_PIH23_009325-RA

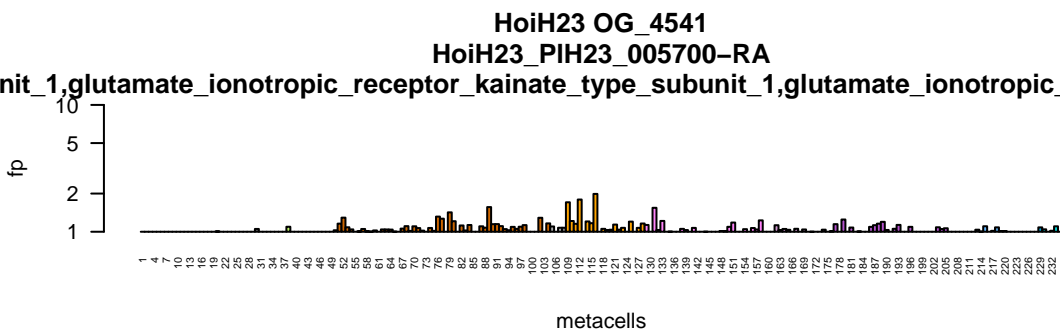
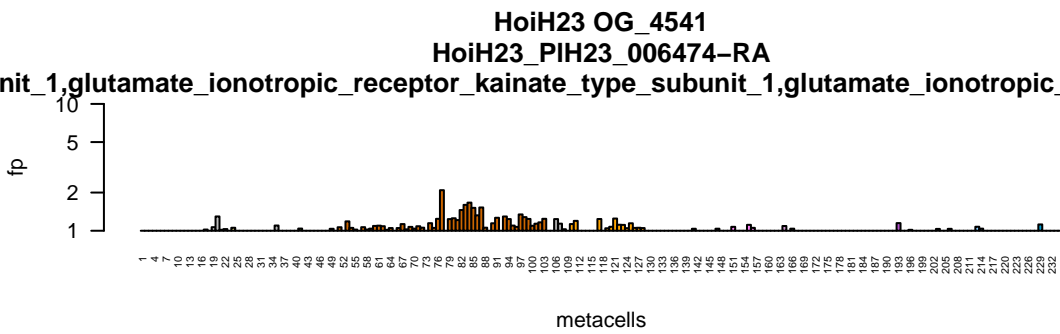
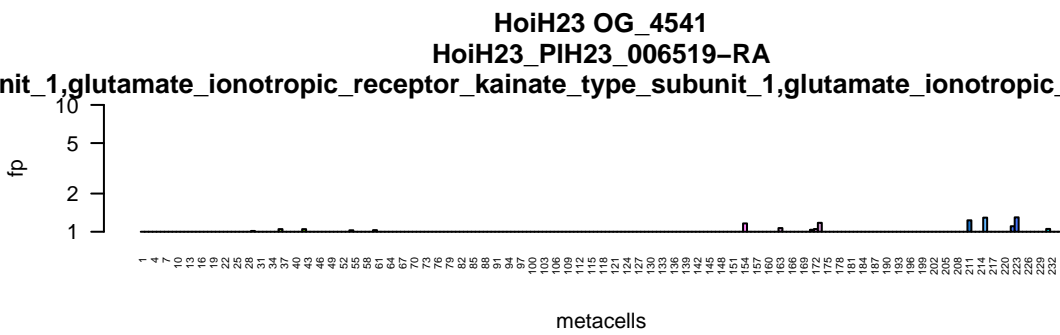
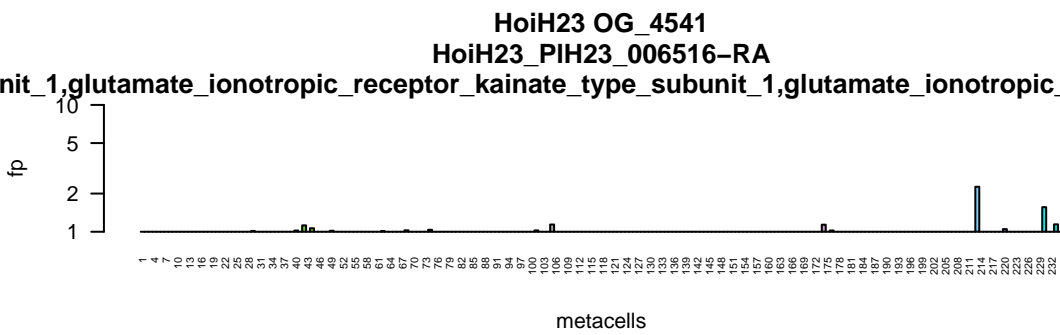
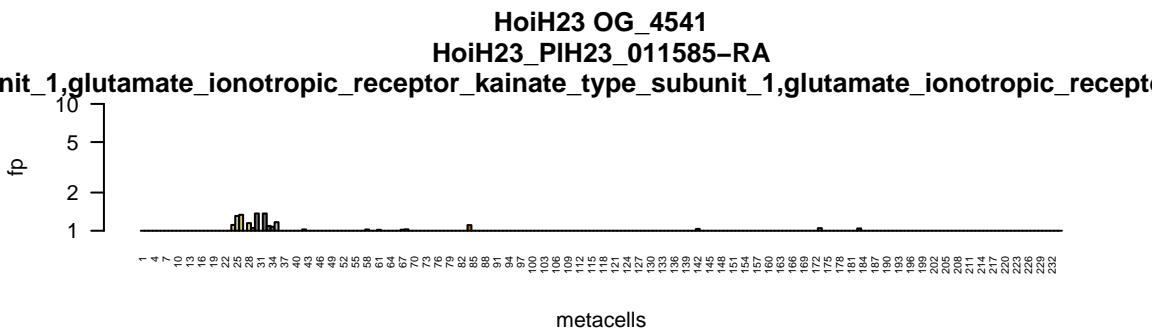
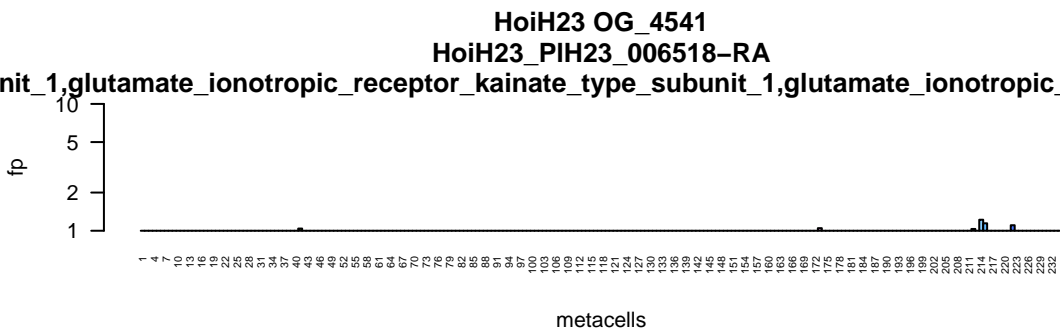
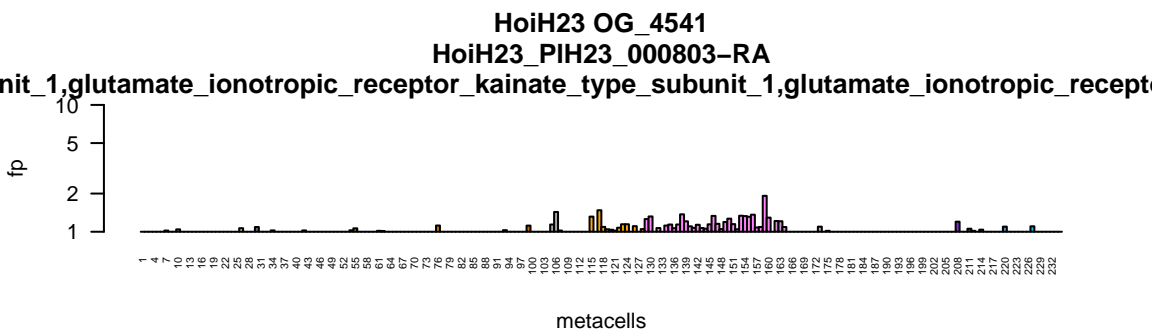
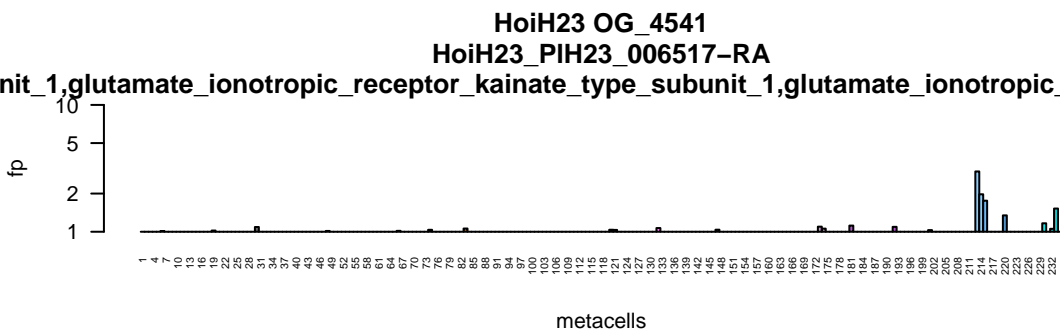
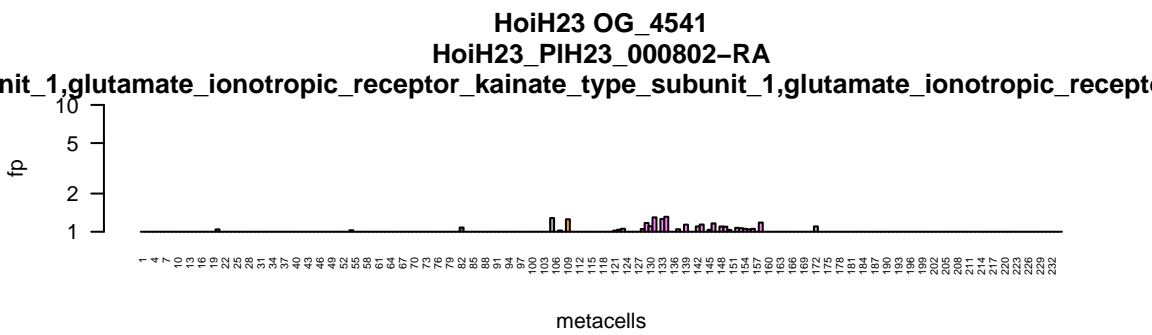
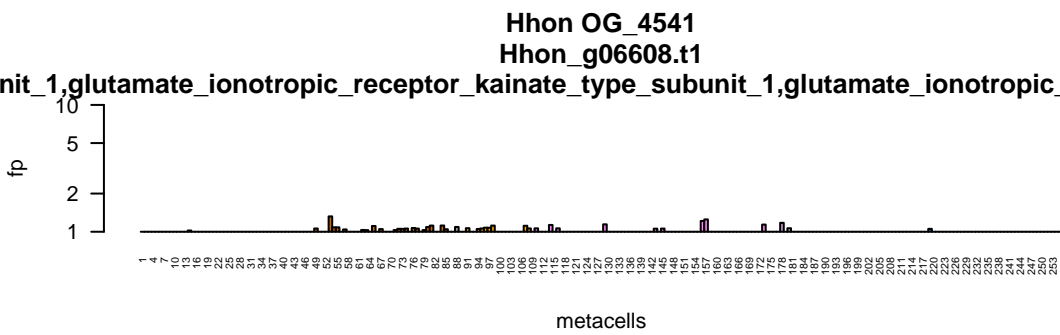
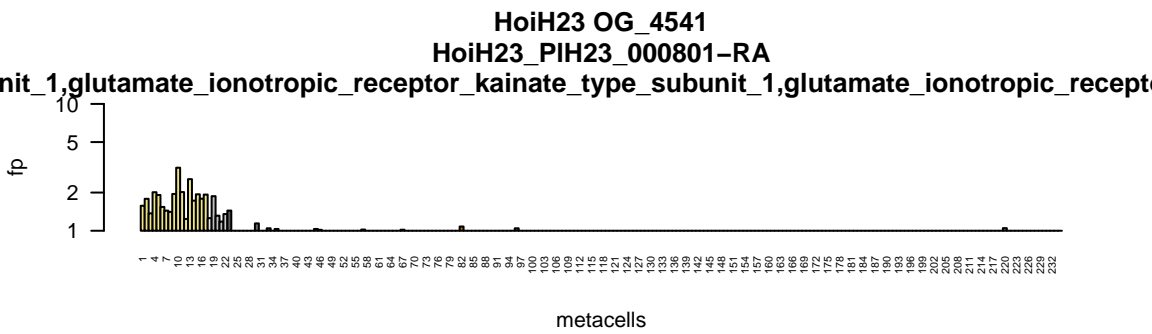
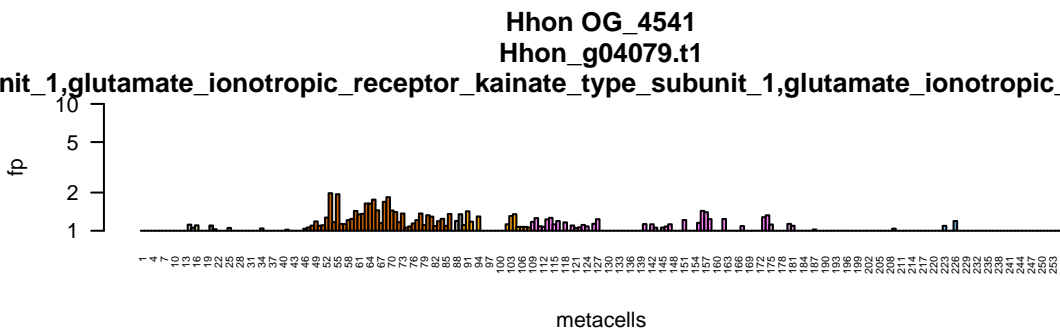


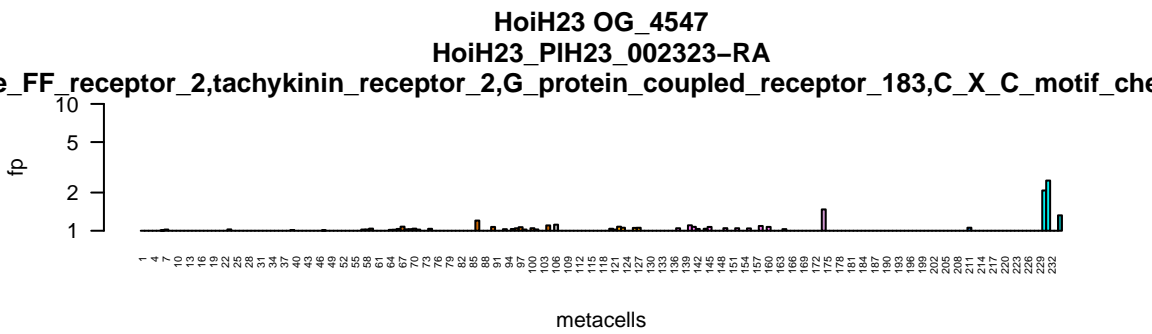
metacells

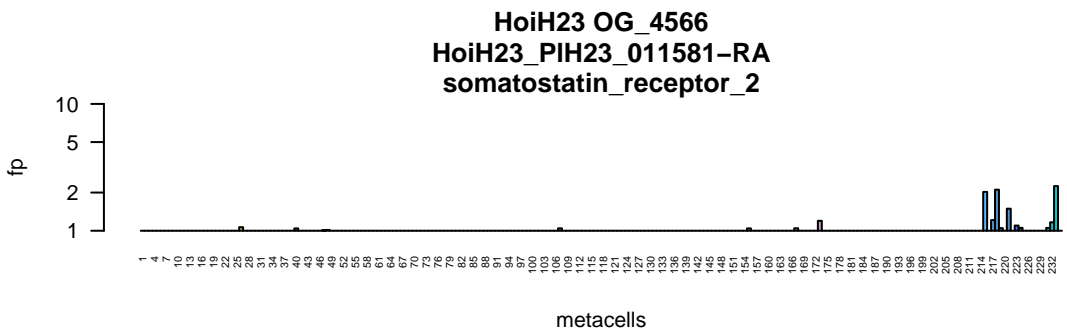
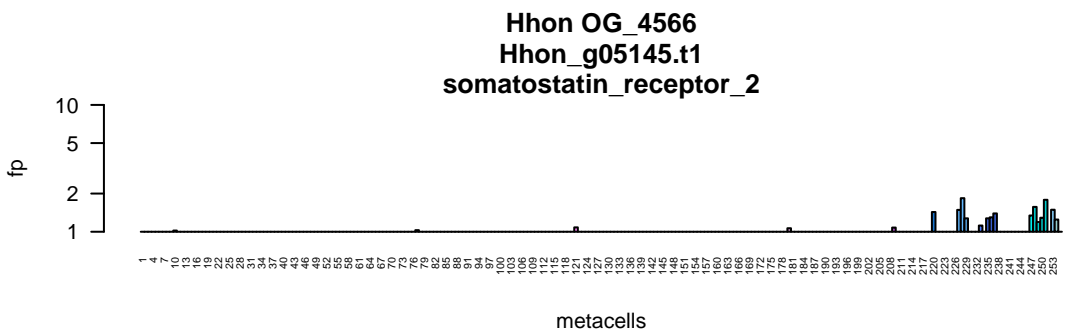
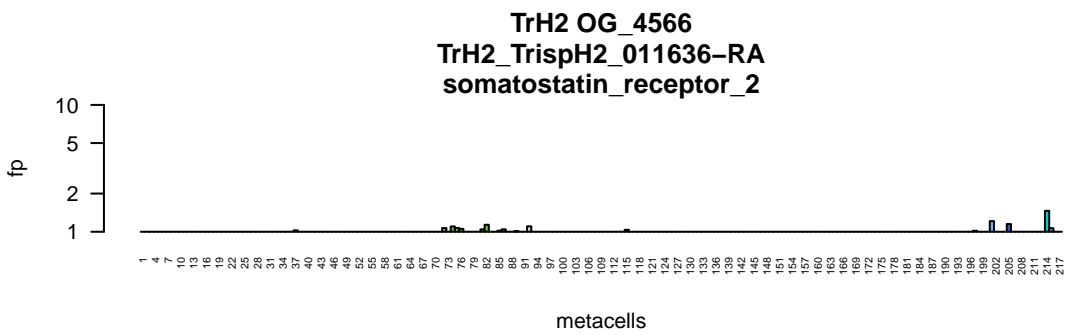
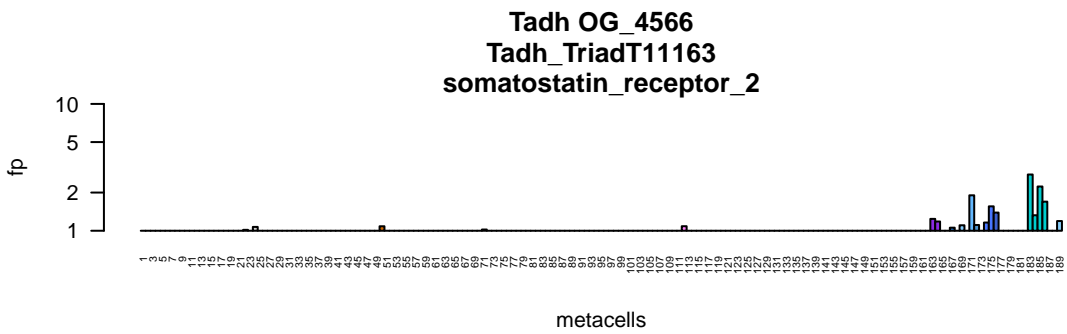




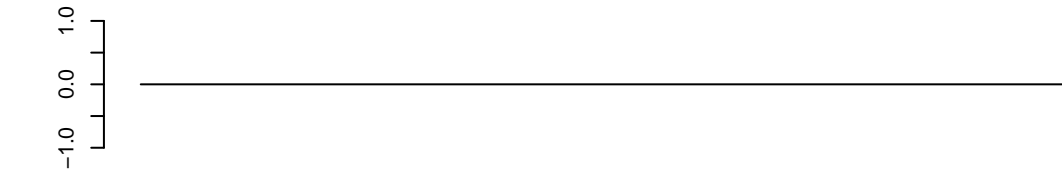




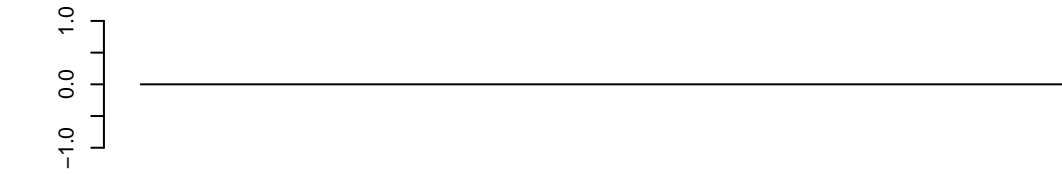




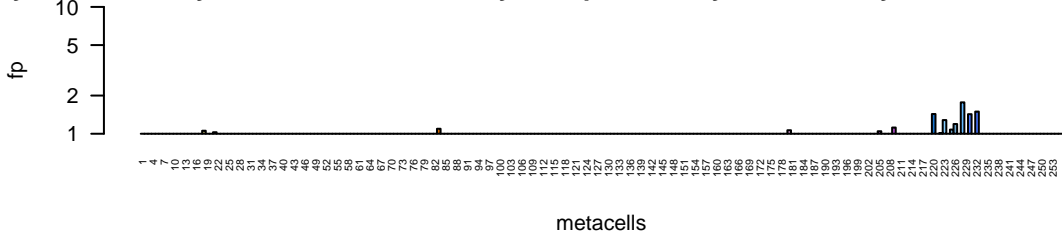
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi
Tadh | no data



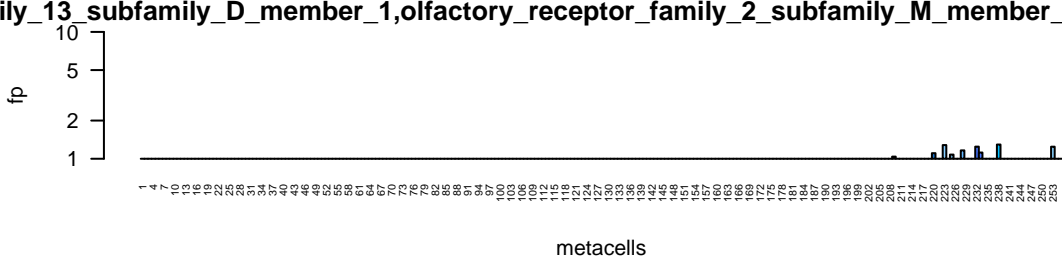
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi
TrH2 | no data



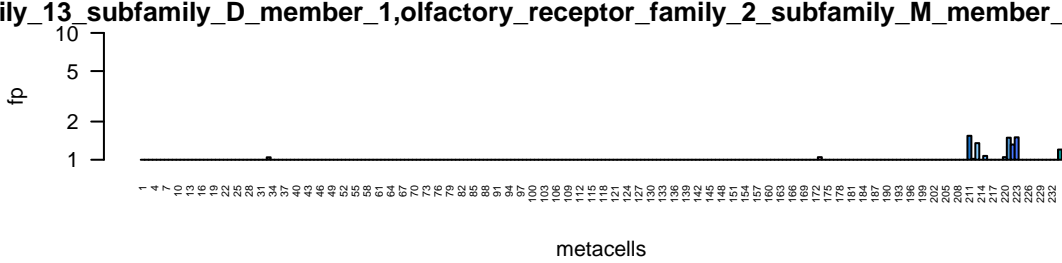
Hhon OG_4788
Hhon_g08016.t1
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi



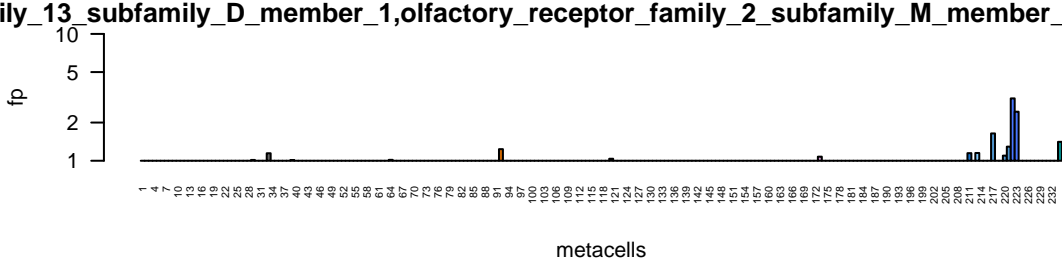
Hhon OG_4788
Hhon_g08015.t1
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi

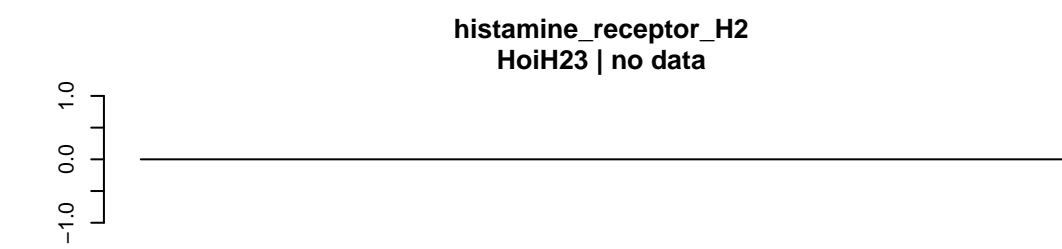
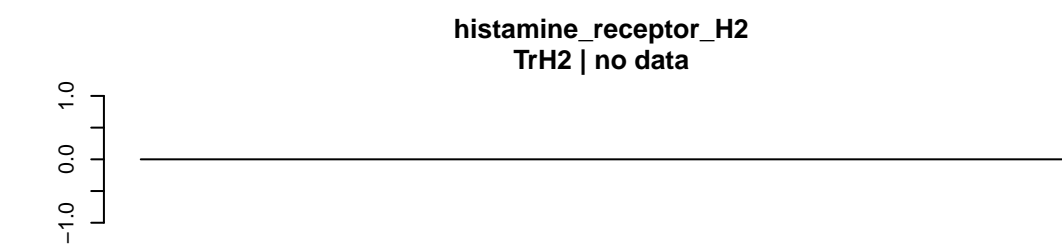
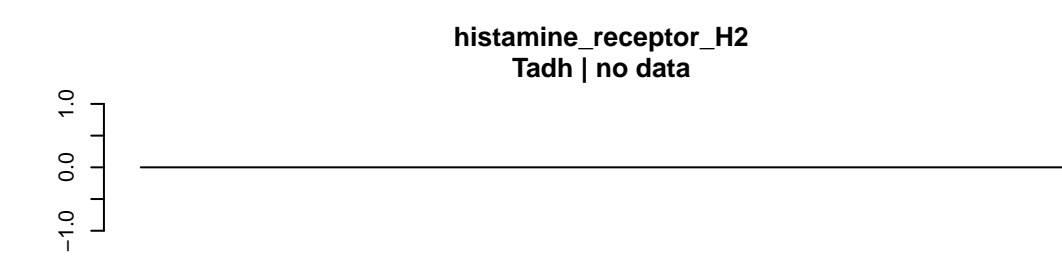


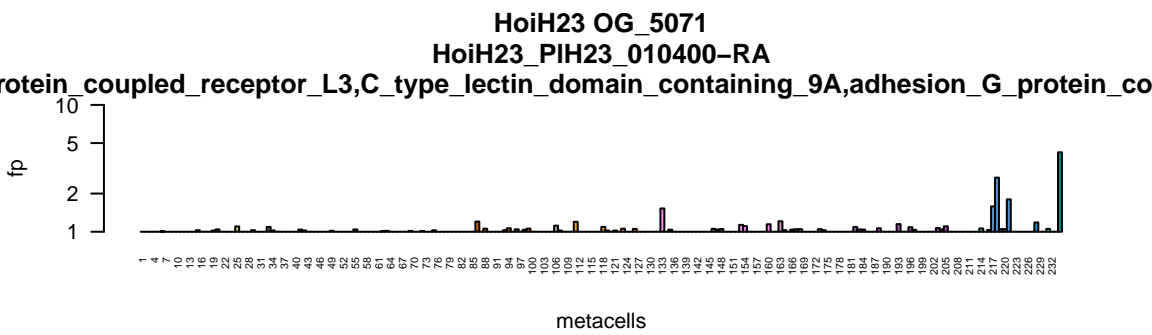
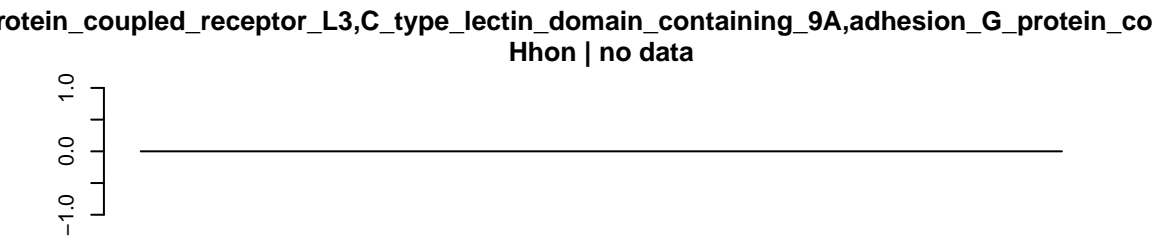
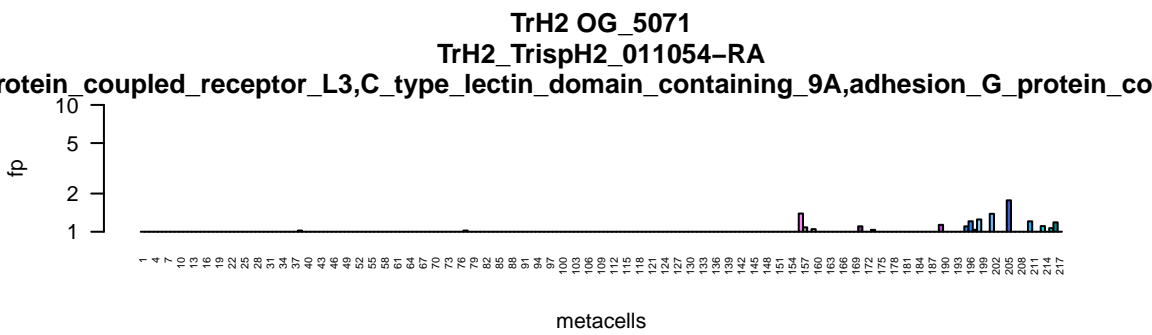
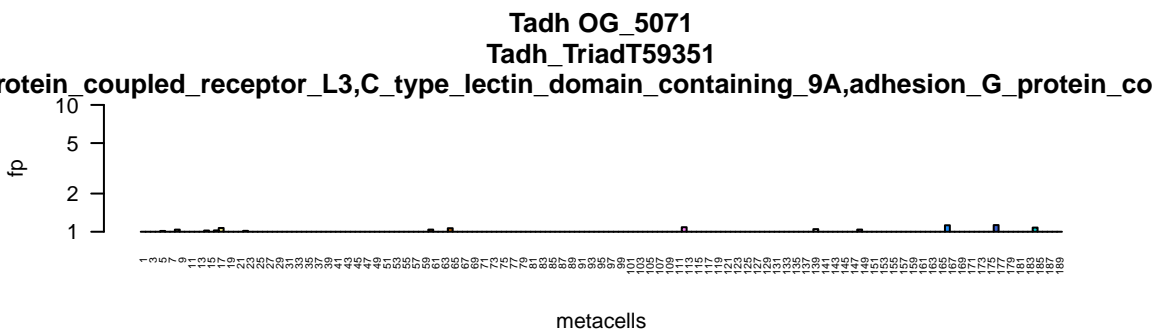
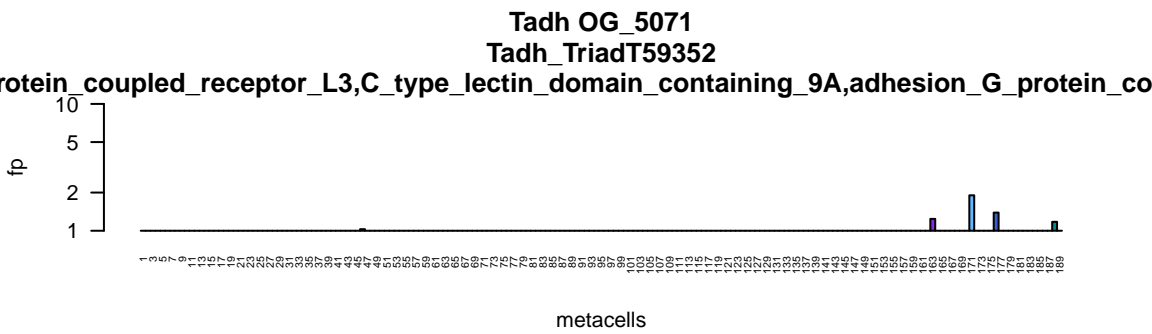
HoiH23 OG_4788
HoiH23_PIH23_005444-RA
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi



HoiH23 OG_4788
HoiH23_PIH23_005445-RA
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi



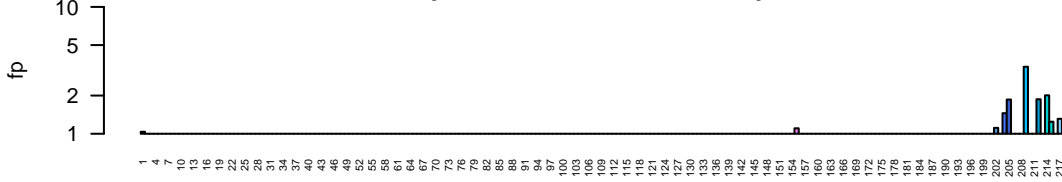




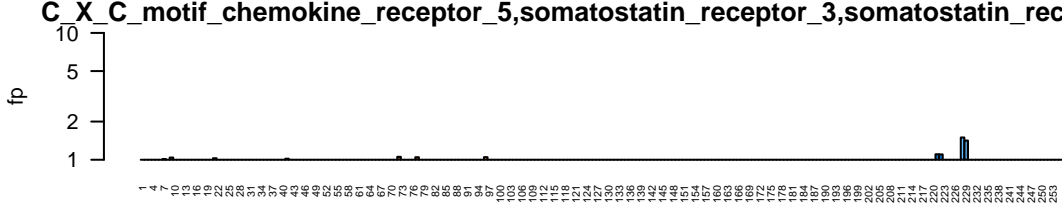
C_X_C_motif_chemokine_receptor_5,somatostatin_receptor_3,somatostatin_receptor_1
Tadh | no data



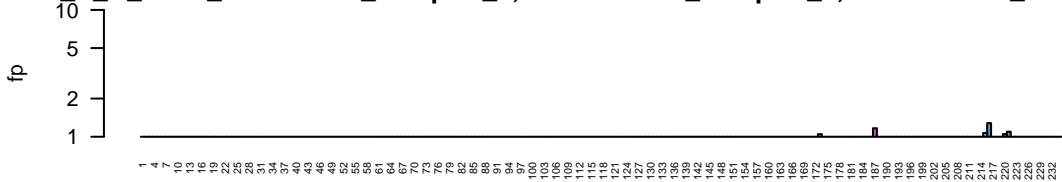
TrH2 OG_5413
TrH2_TrispH2_008283-RA
C_X_C_motif_chemokine_receptor_5,somatostatin_receptor_3,somatostatin_receptor_1

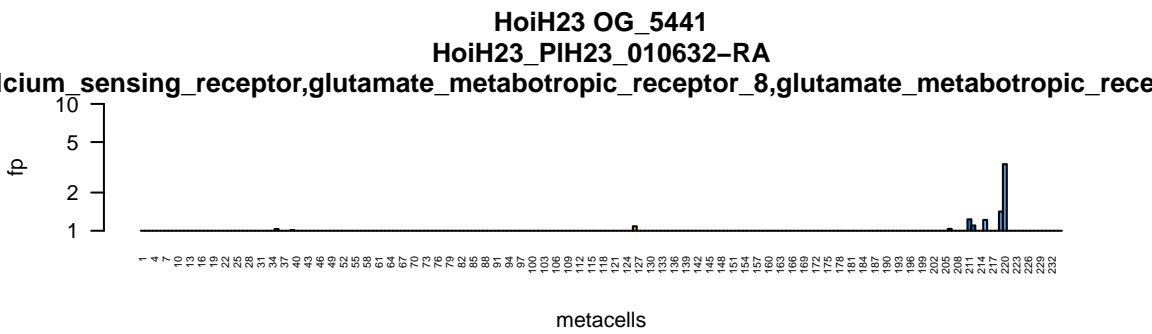
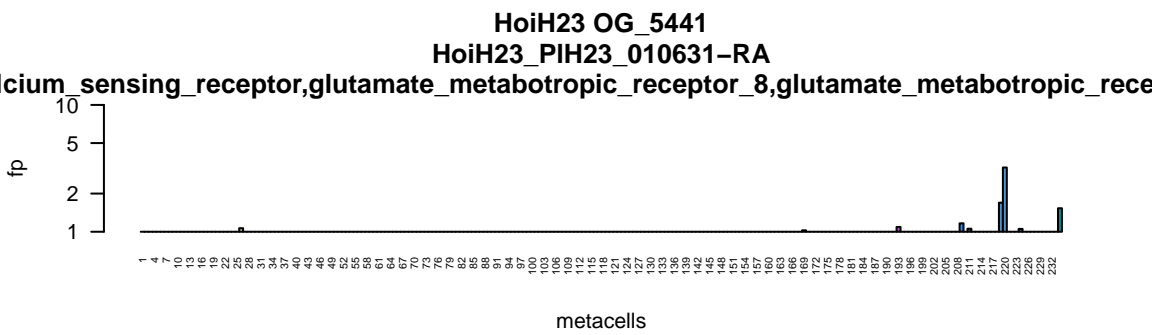
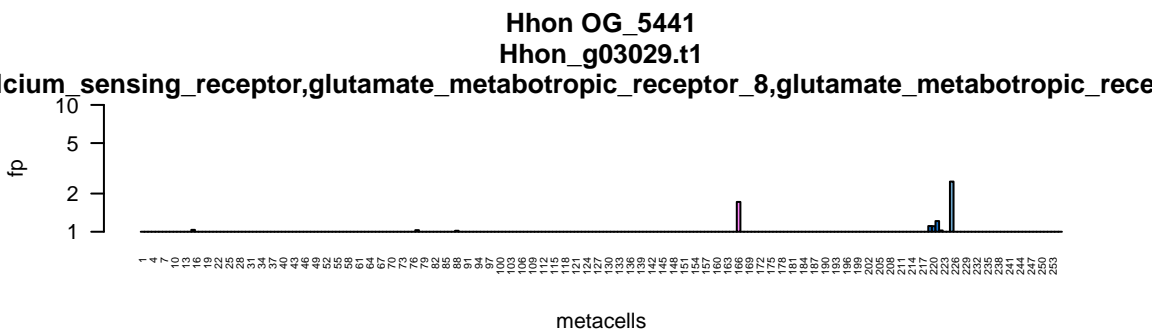
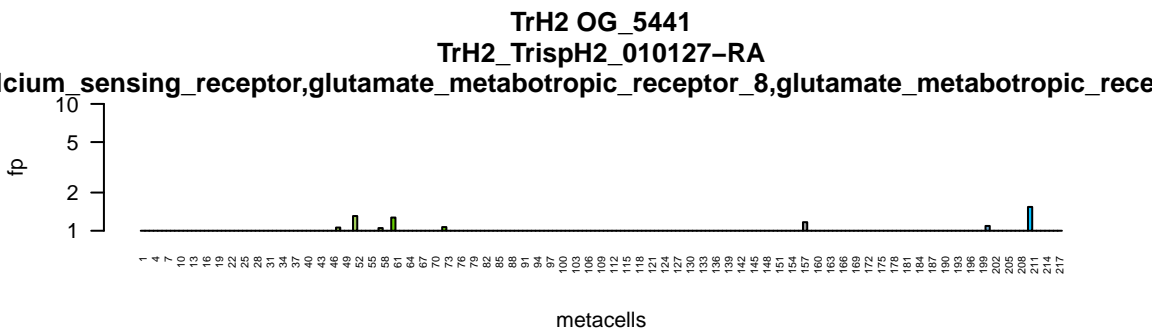
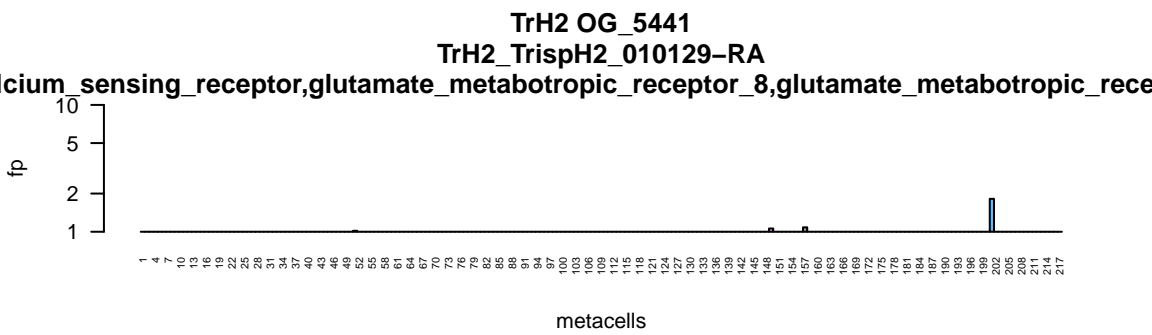
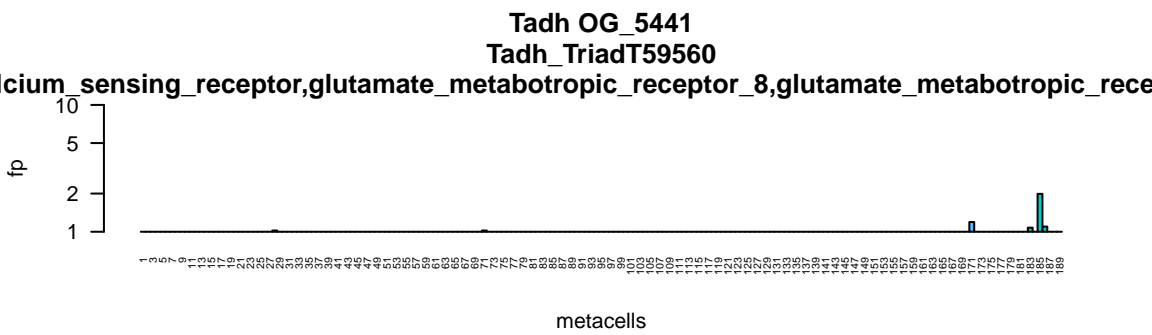
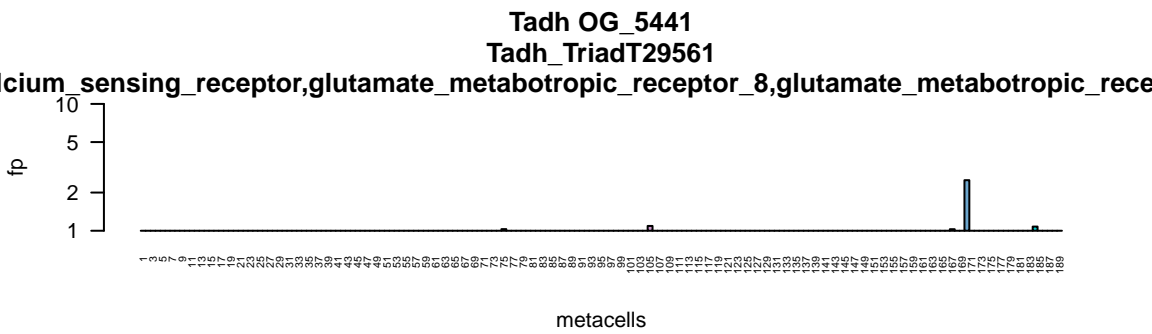


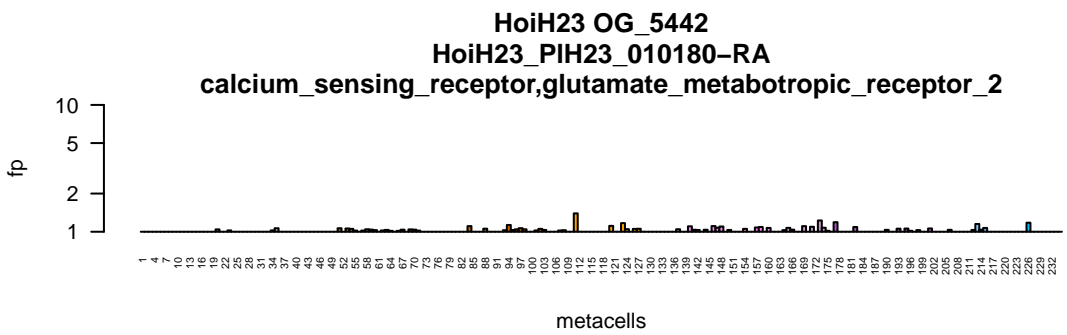
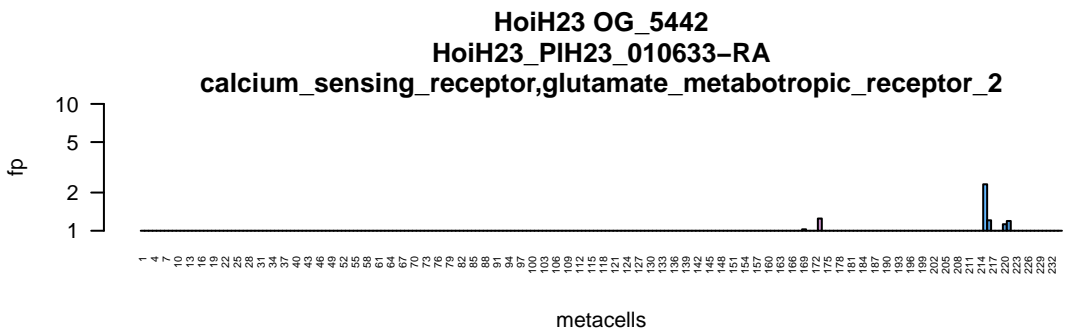
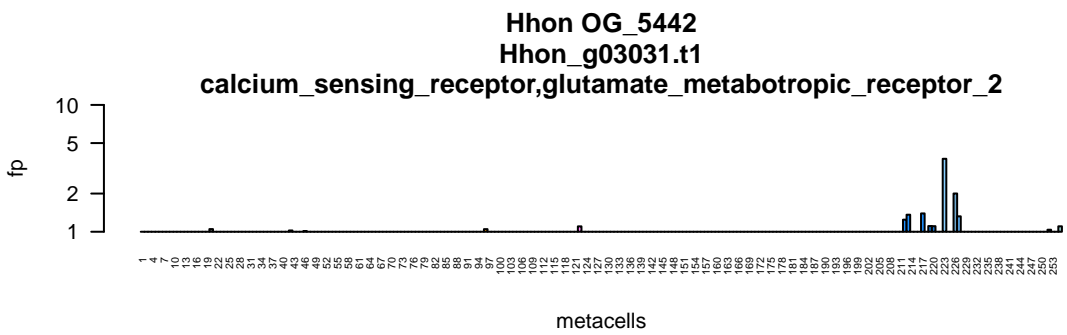
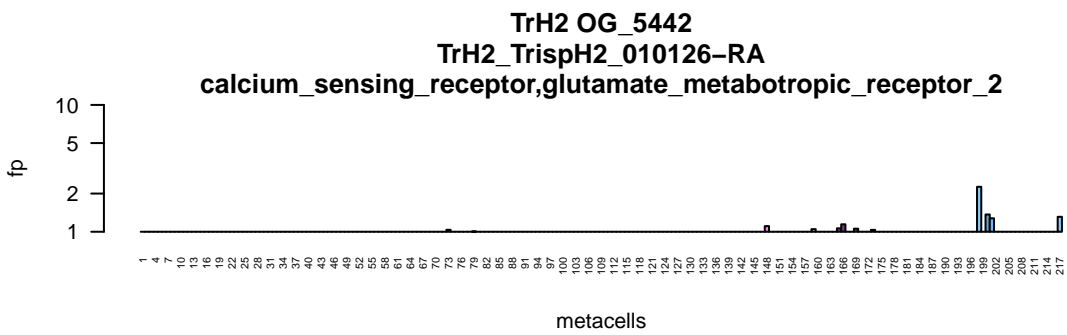
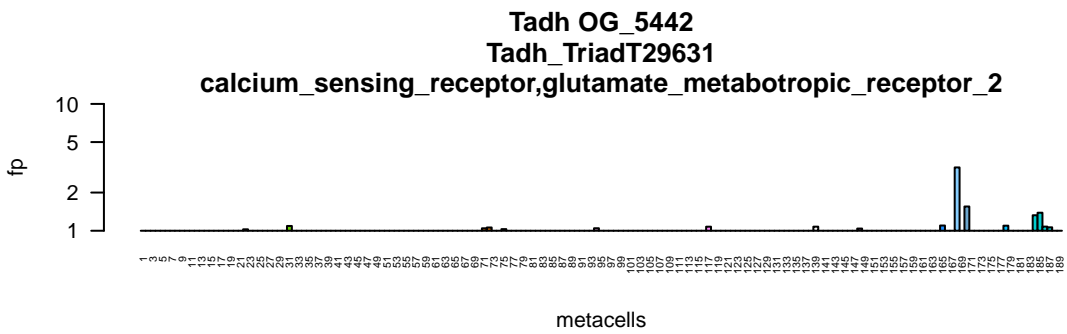
metacells
Hhon OG_5413
Hhon_g11630.t1
C_X_C_motif_chemokine_receptor_5,somatostatin_receptor_3,somatostatin_receptor_1



HoiH23 OG_5413
HoiH23_PIH23_006165-RA
C_X_C_motif_chemokine_receptor_5,somatostatin_receptor_3,somatostatin_receptor_1





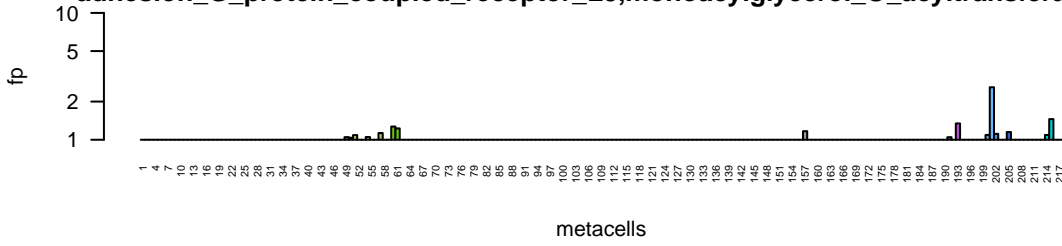


adhesion_G_protein_coupled_receptor_L3,monoacylglycerol_O_acyltransferase_1
Tadh | no data

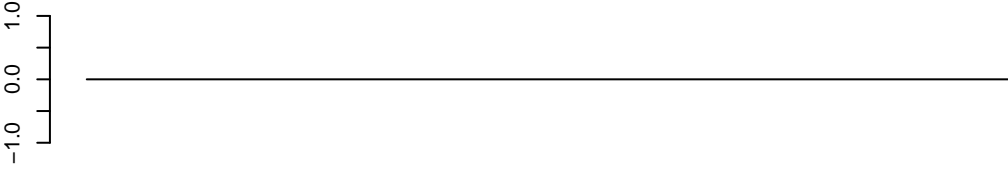


TrH2 OG_5734
TrH2_TrispH2_011056-RA

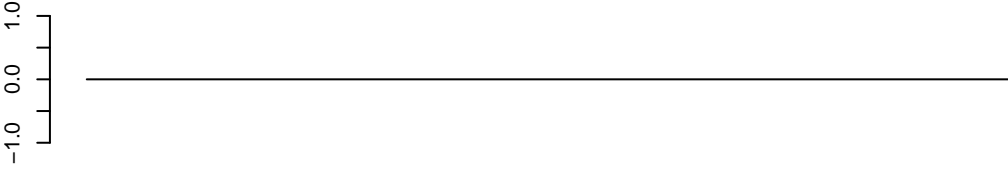
adhesion_G_protein_coupled_receptor_L3,monoacylglycerol_O_acyltransferase_1



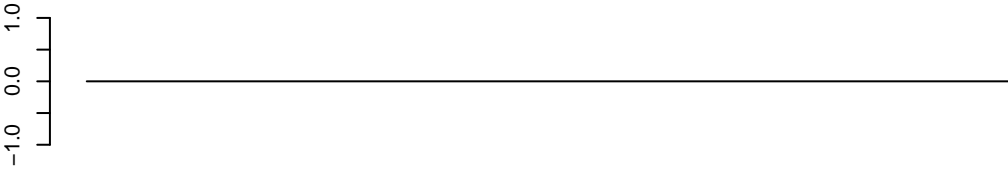
adhesion_G_protein_coupled_receptor_L3,monoacylglycerol_O_acyltransferase_1
Hhon | no data



adhesion_G_protein_coupled_receptor_L3,monoacylglycerol_O_acyltransferase_1
HoiH23 | no data



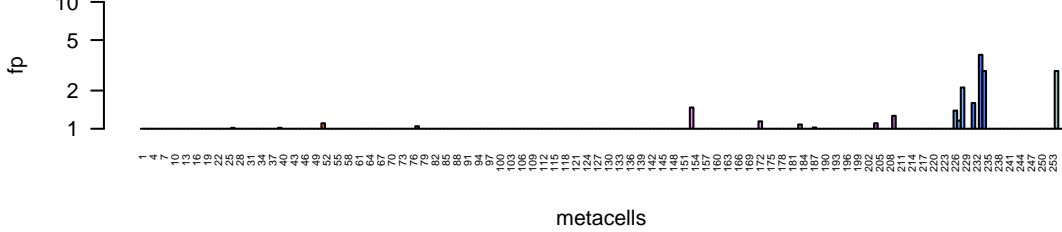
gamma_aminobutyric_acid_type_B_receptor_subunit_2
Tadh | no data



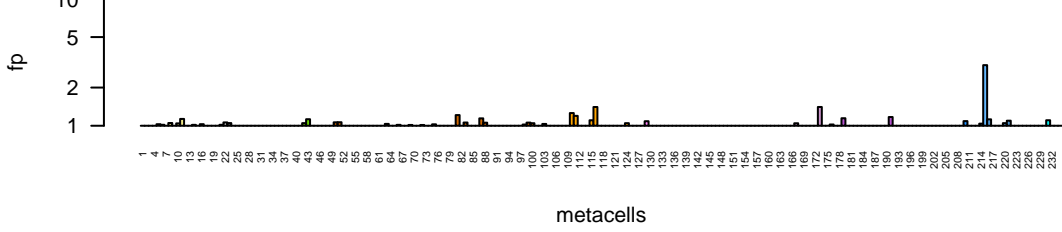
gamma_aminobutyric_acid_type_B_receptor_subunit_2
TrH2 | no data



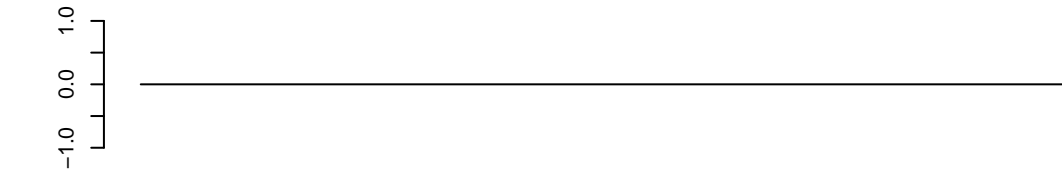
Hhon OG_6010
Hhon_g01250.t1
gamma_aminobutyric_acid_type_B_receptor_subunit_2



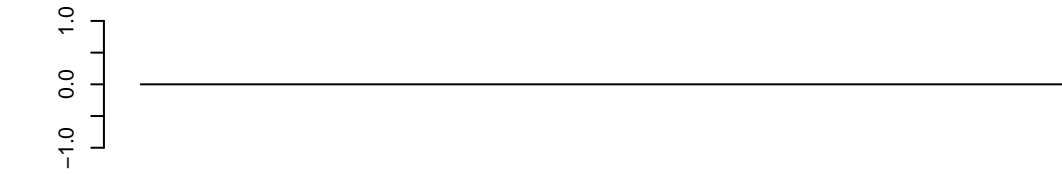
HoiH23 OG_6010
HoiH23_PIH23_000005-RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2



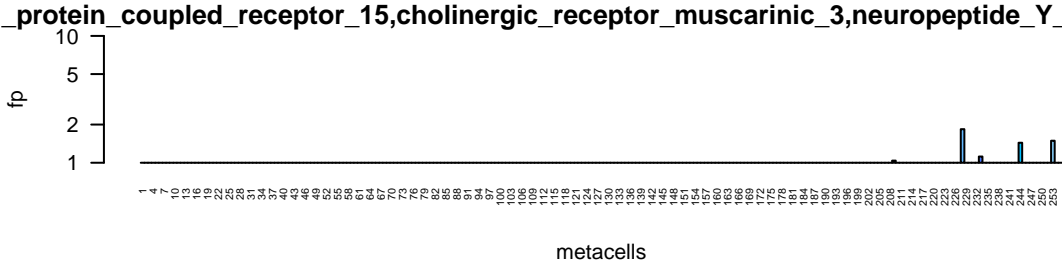
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept
Tadh | no data



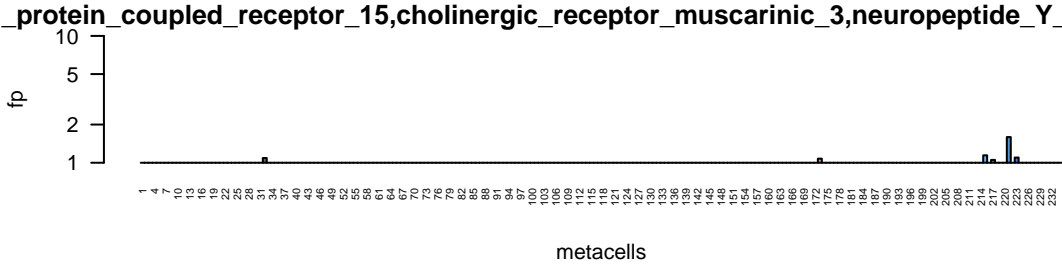
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept
TrH2 | no data

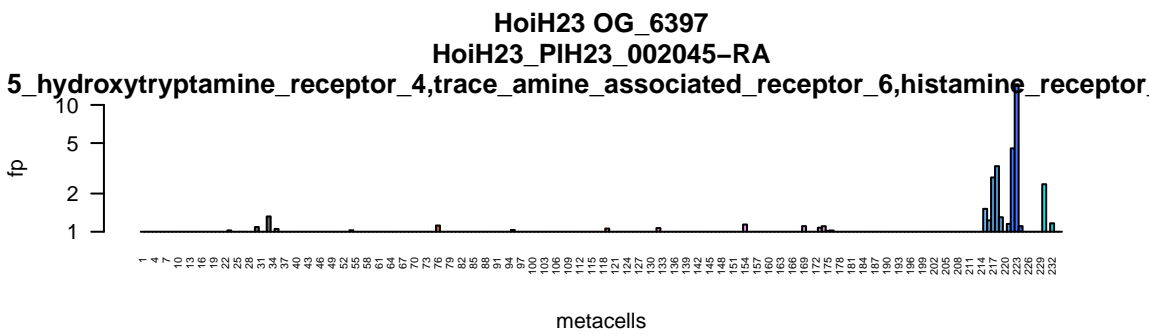
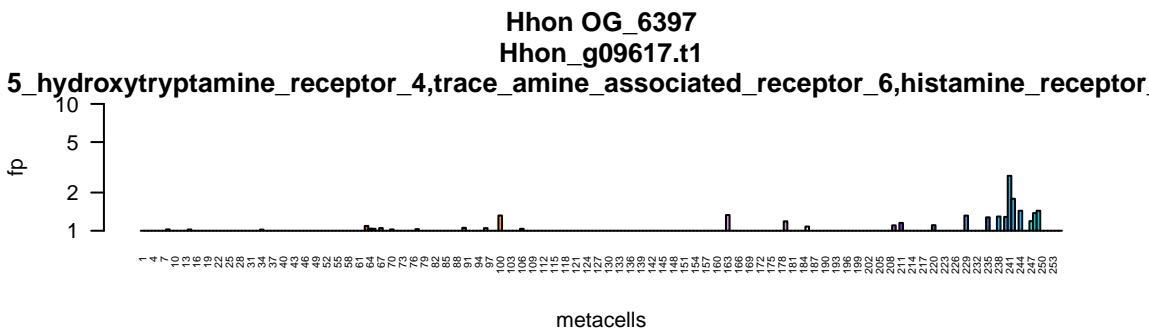
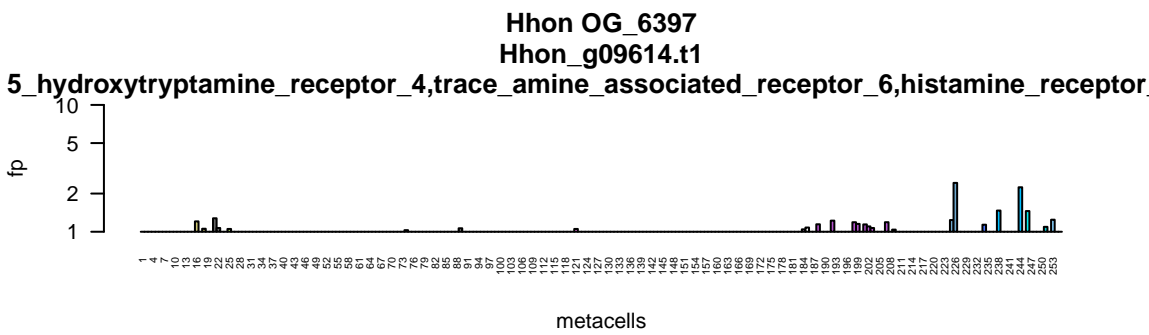
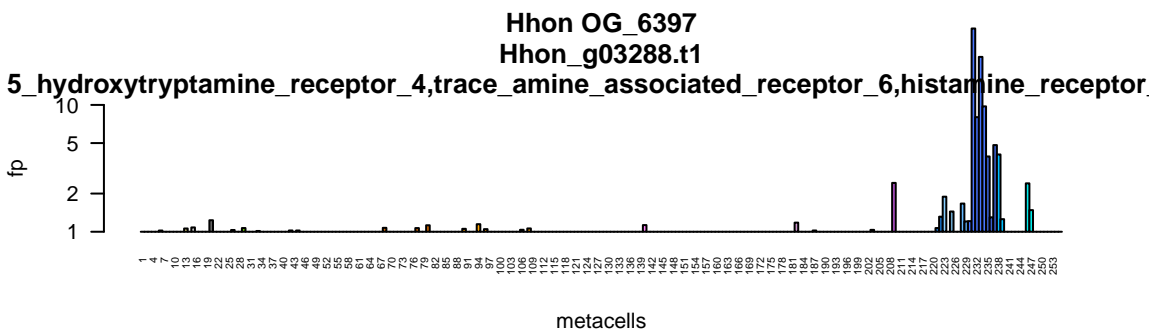
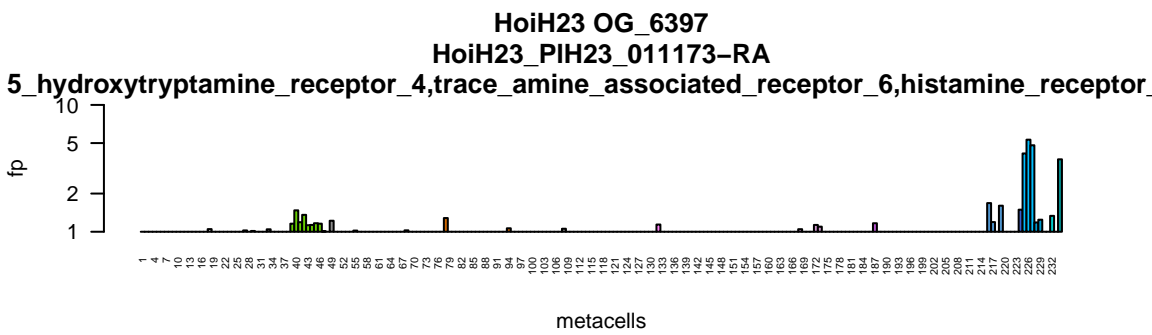
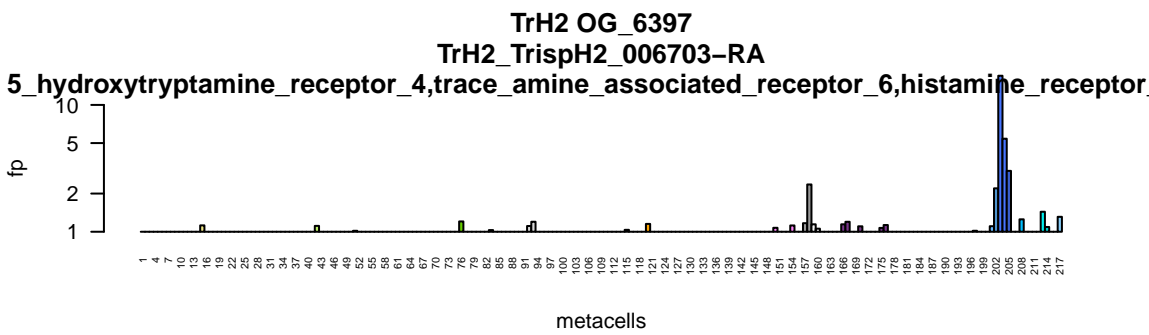
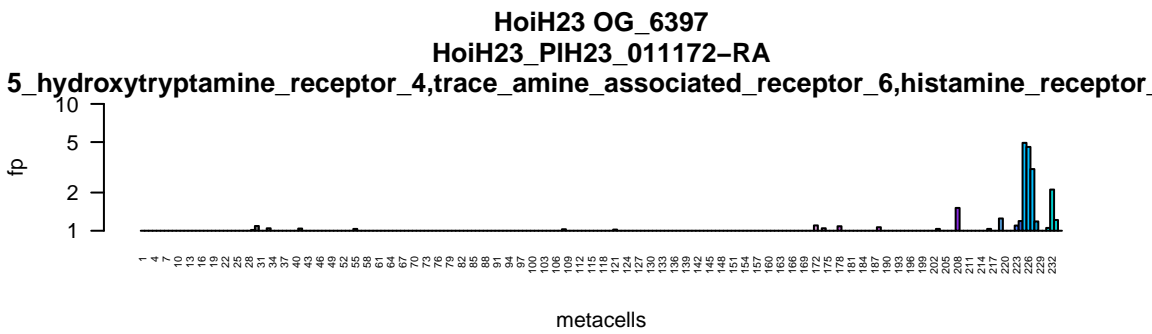
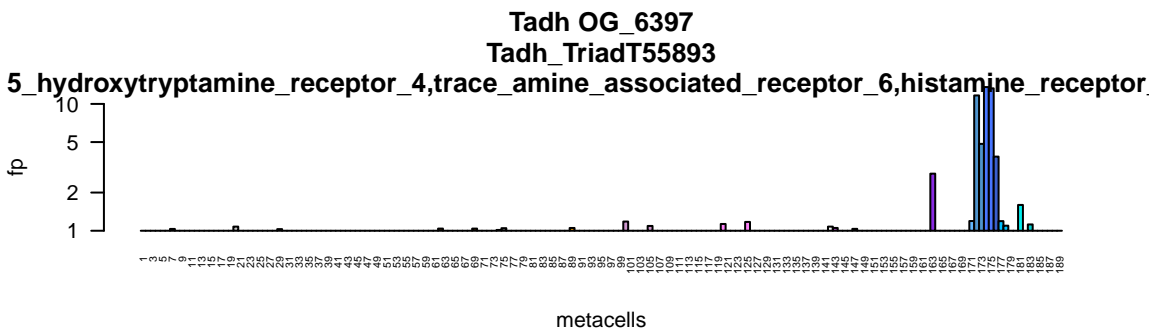


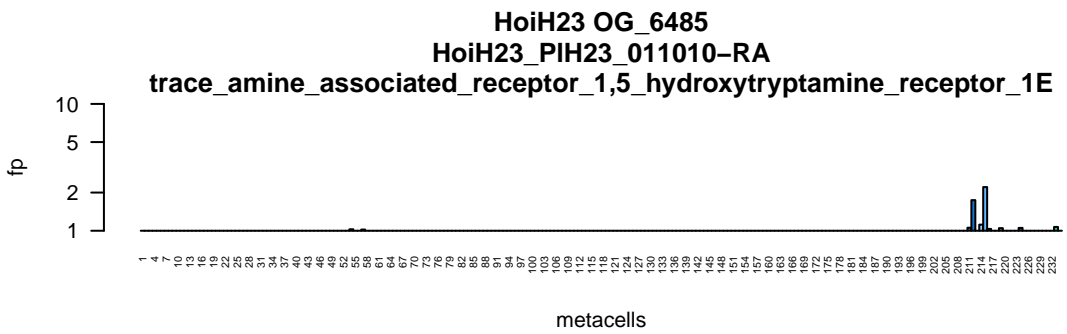
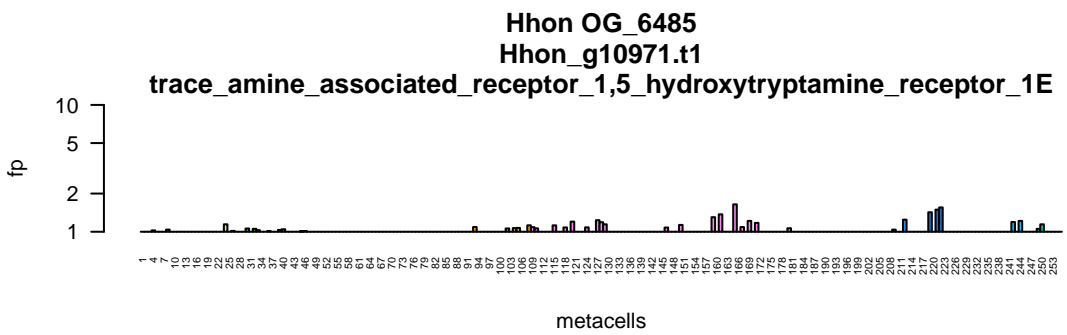
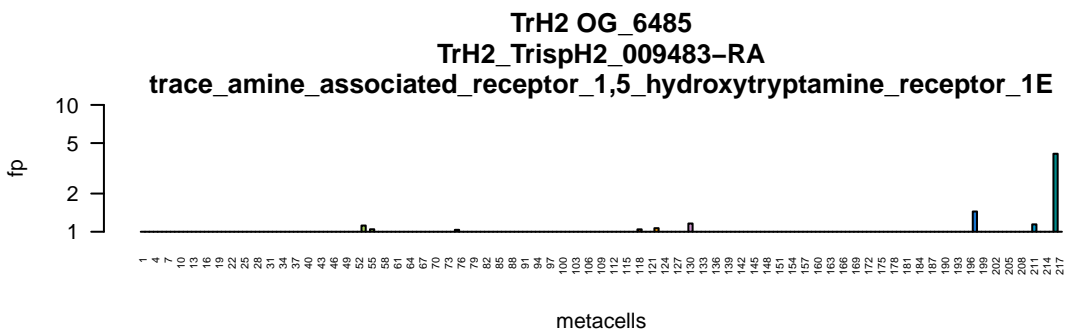
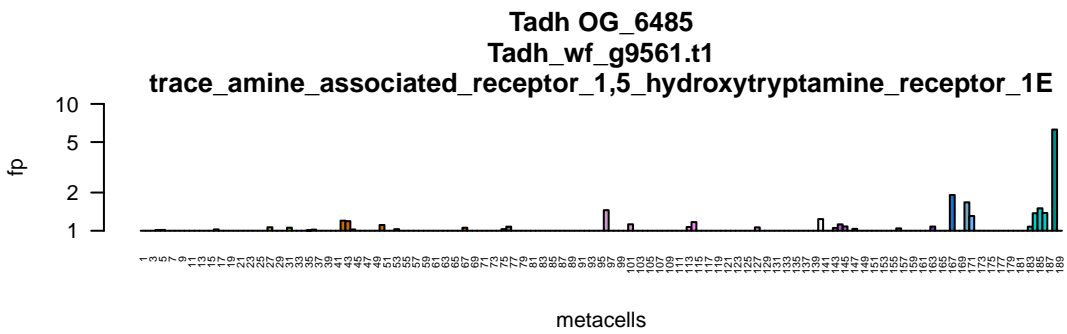
Hhon OG_6355
Hhon_g11199.t1
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept

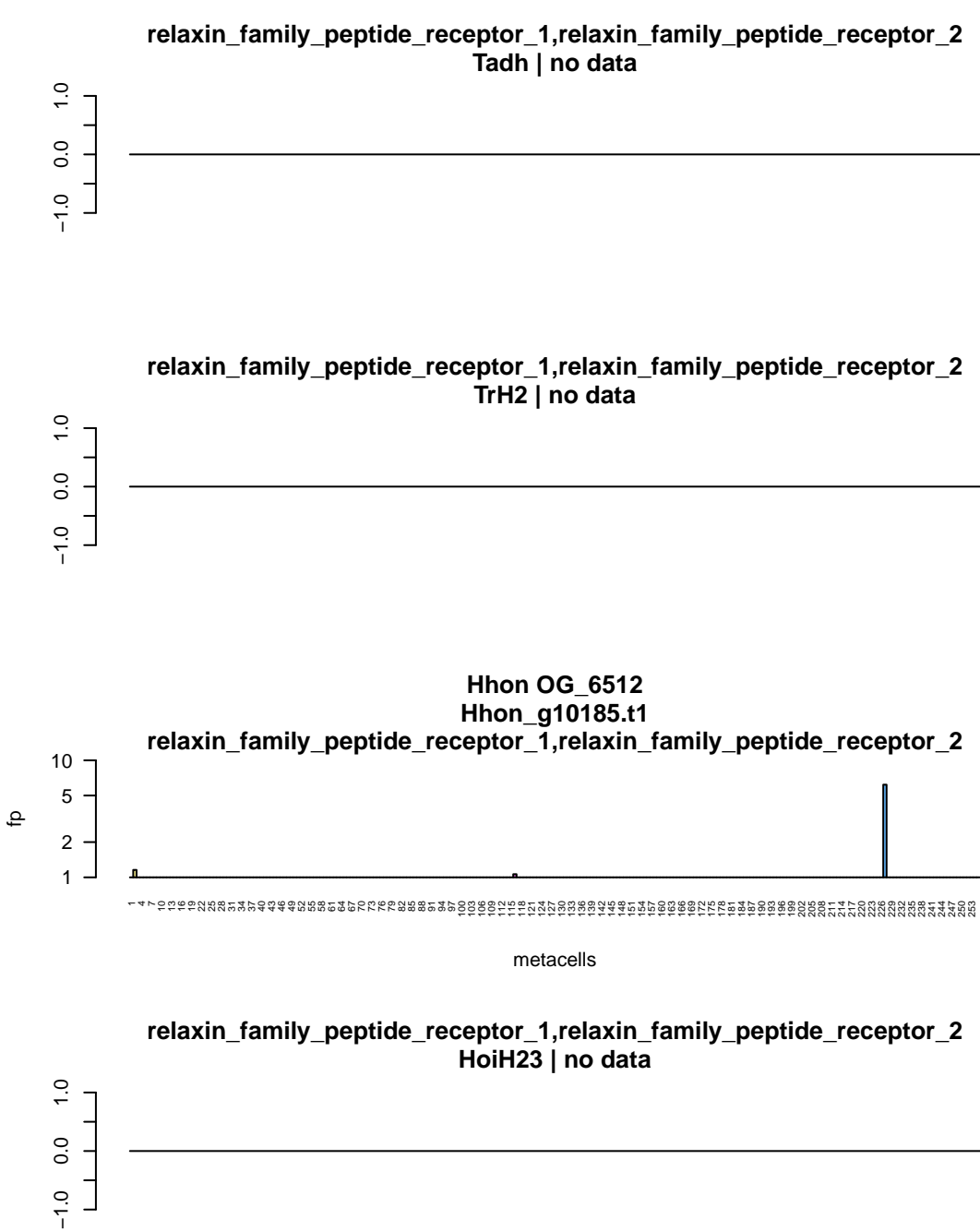


HoiH23 OG_6355
HoiH23_PIH23_012003-RA
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept









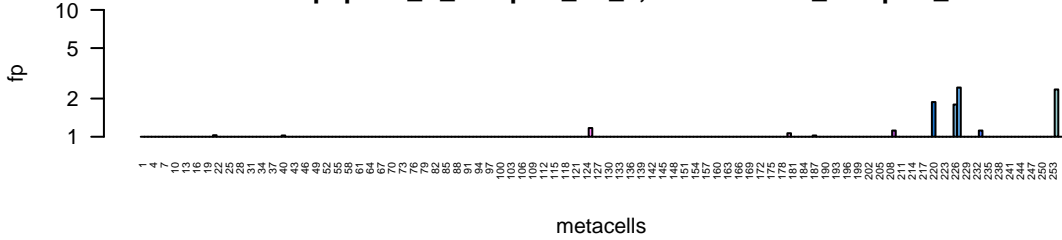
neuropeptide_Y_receptor_Y4_2,somatostatin_receptor_1
Tadh | no data



neuropeptide_Y_receptor_Y4_2,somatostatin_receptor_1
TrH2 | no data



Hhon OG_6729
Hhon_g05494.t1
neuropeptide_Y_receptor_Y4_2,somatostatin_receptor_1



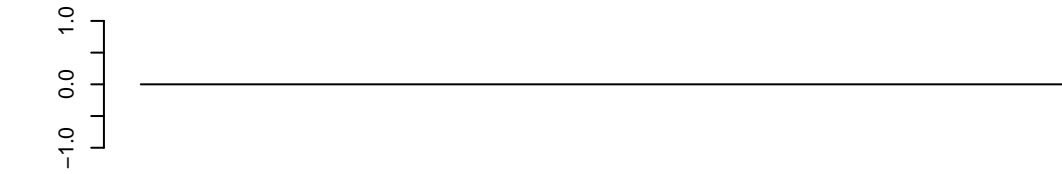
neuropeptide_Y_receptor_Y4_2,somatostatin_receptor_1
HoiH23 | no data



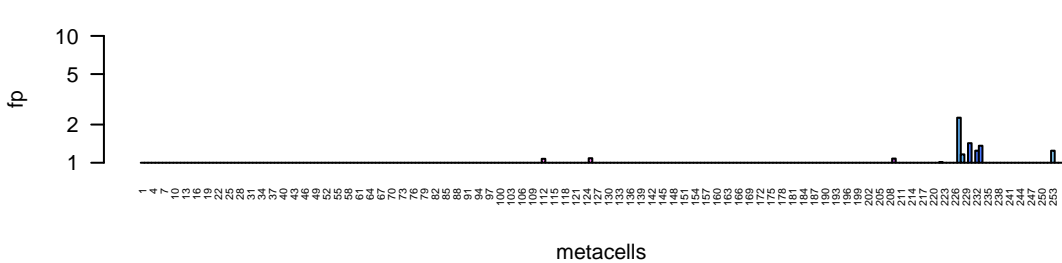
Tadh | no data



TrH2 | no data

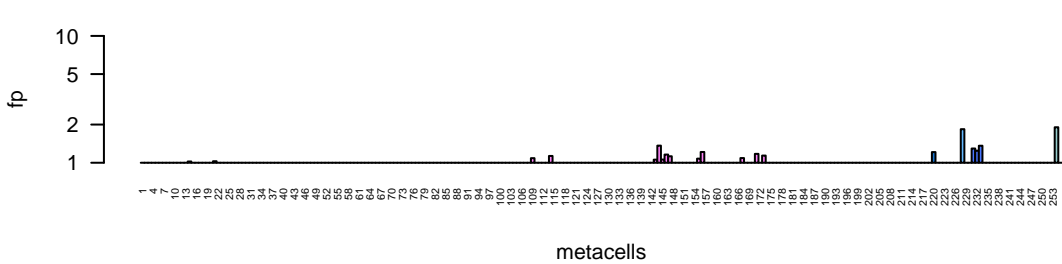


Hhon OG_6868
Hhon_g08412.t1



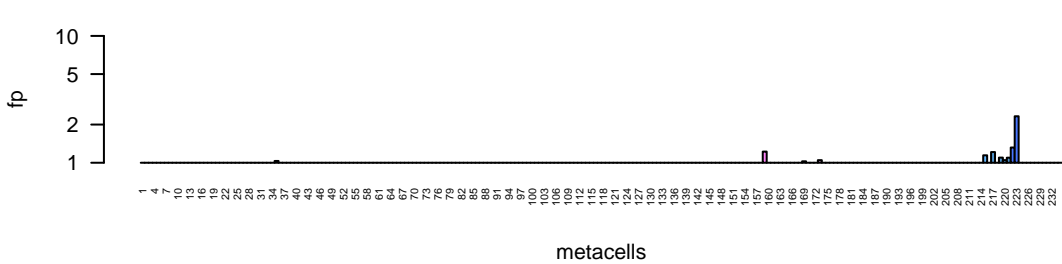
metacells

Hhon OG_6868
Hhon_g07653.t1



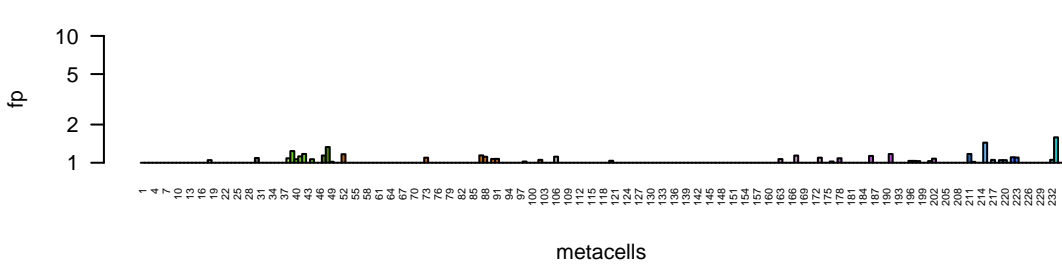
metacells

HoiH23 OG_6868
HoiH23_PIH23_011708-RA

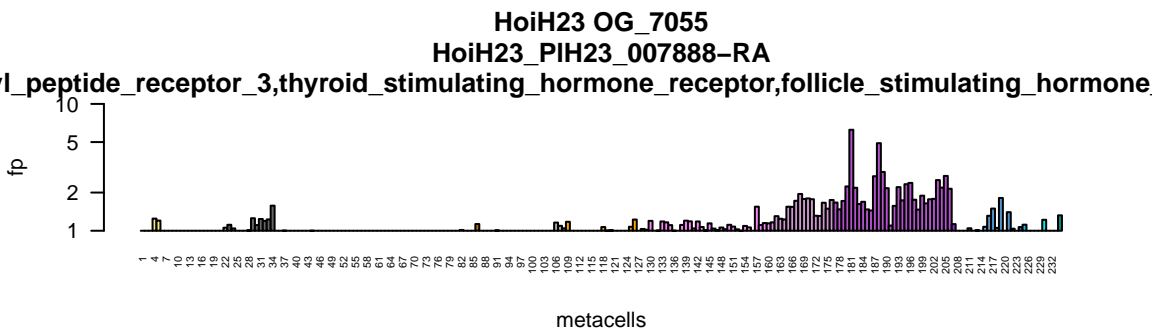
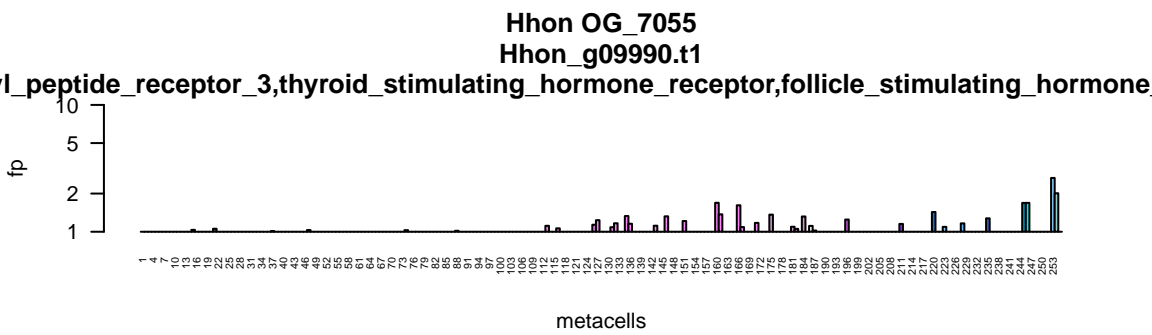
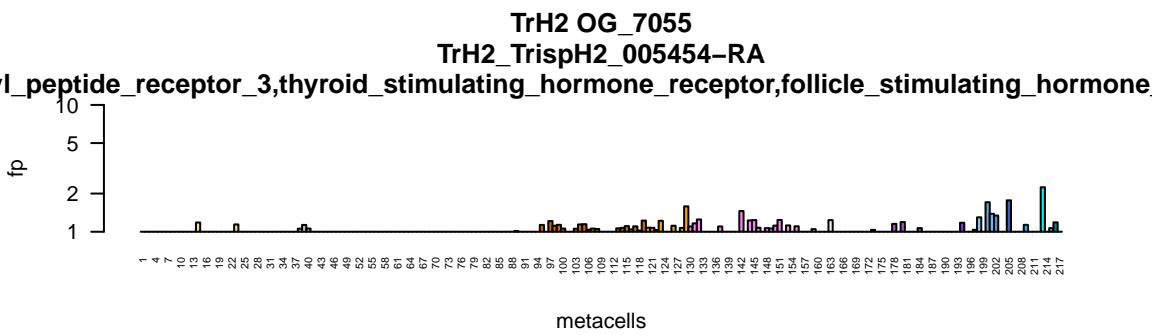
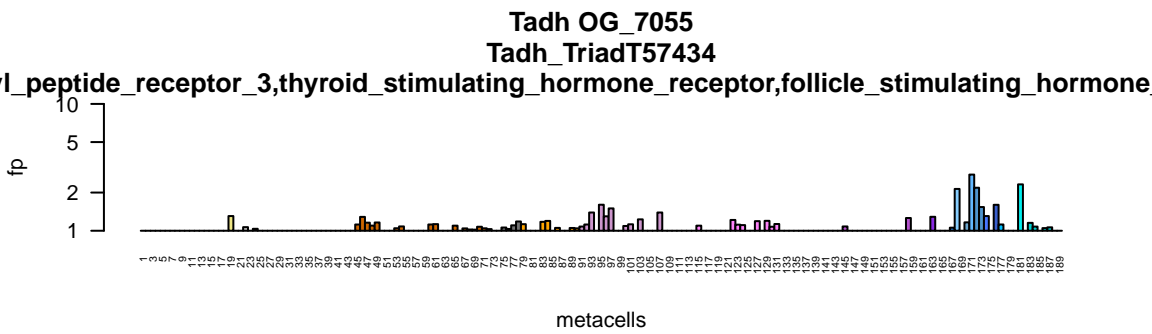


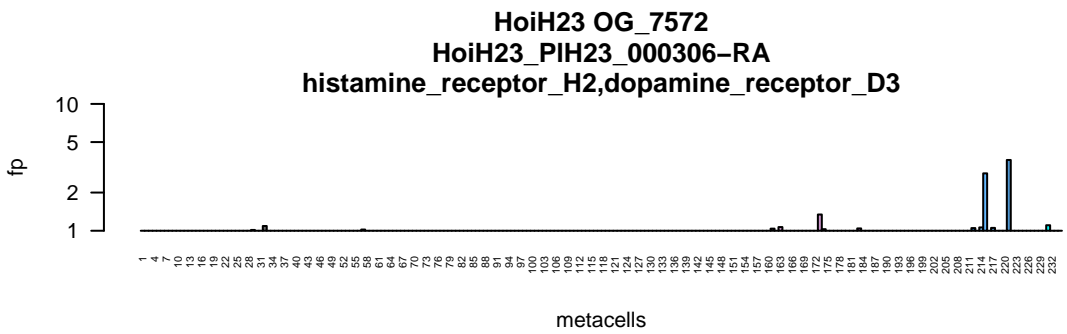
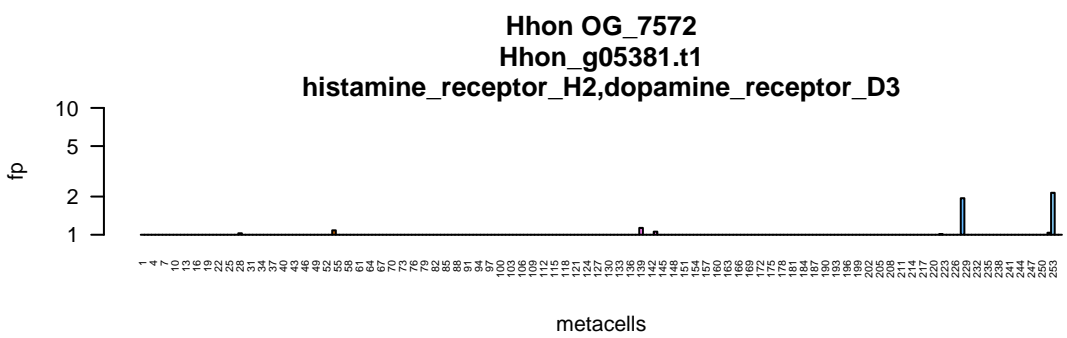
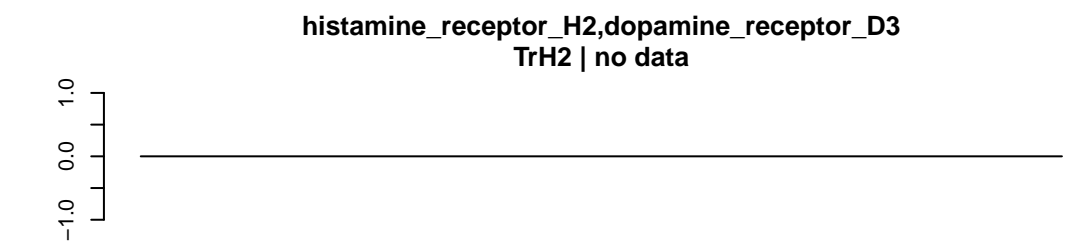
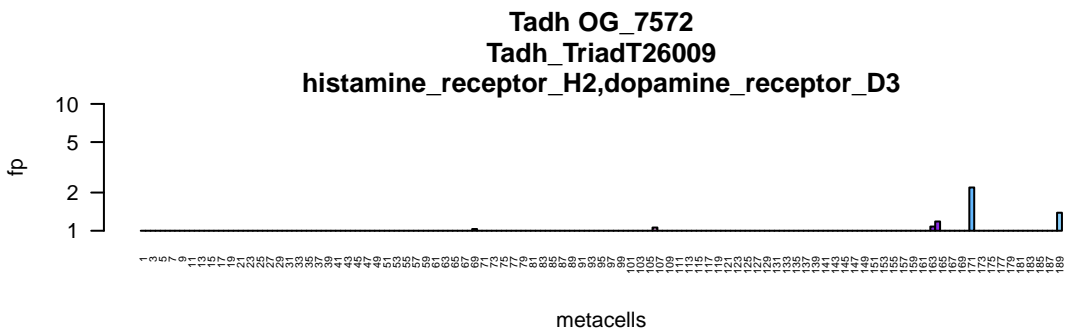
metacells

HoiH23 OG_6868
HoiH23_PIH23_011709-RA

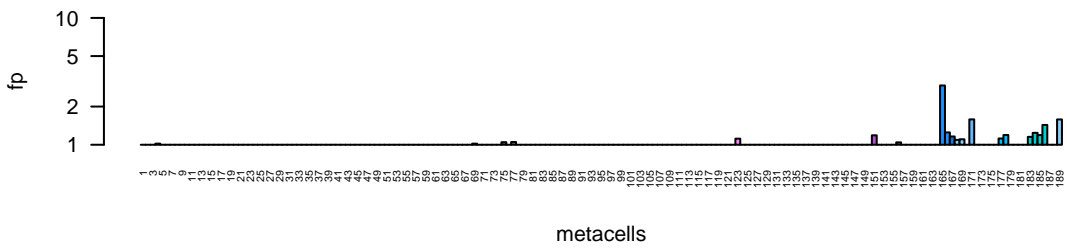


metacells

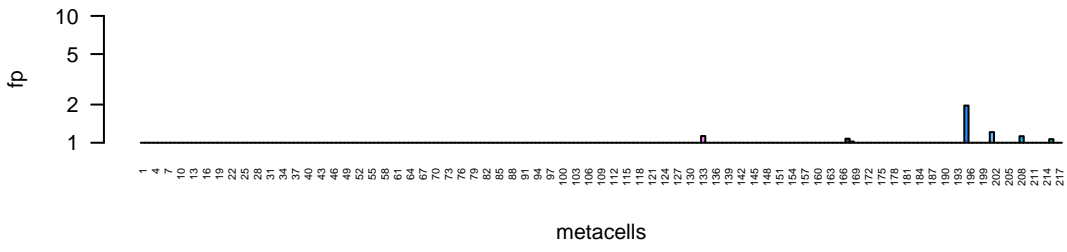




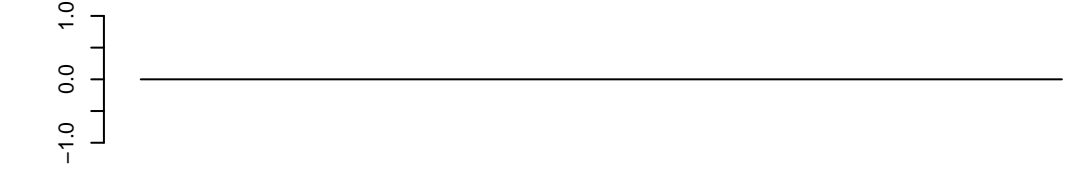
Tadh OG_7681
Tadh_wf_g5016.t1



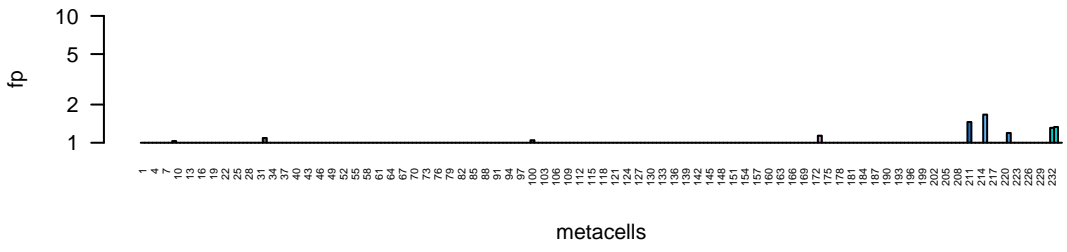
TrH2 OG_7681
TrH2_TrispH2_005499-RA

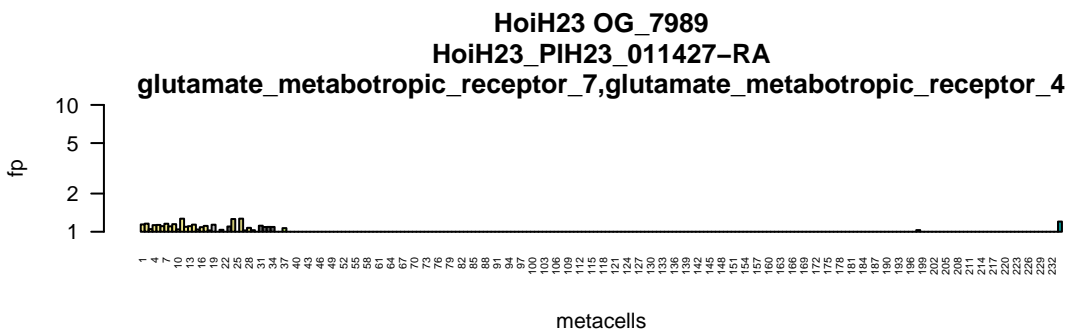
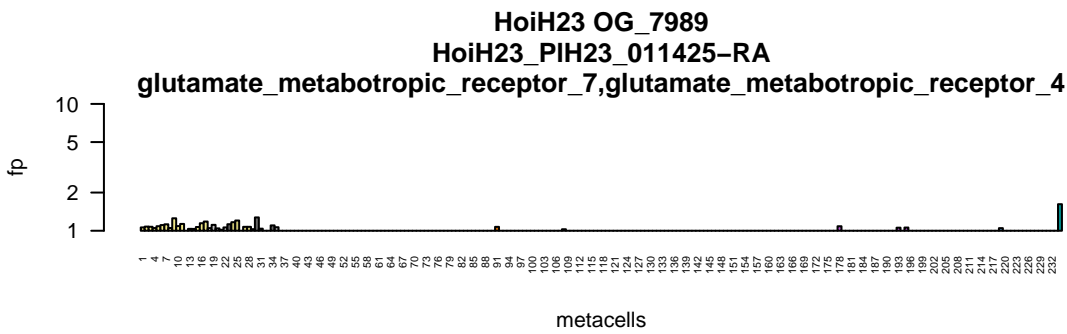
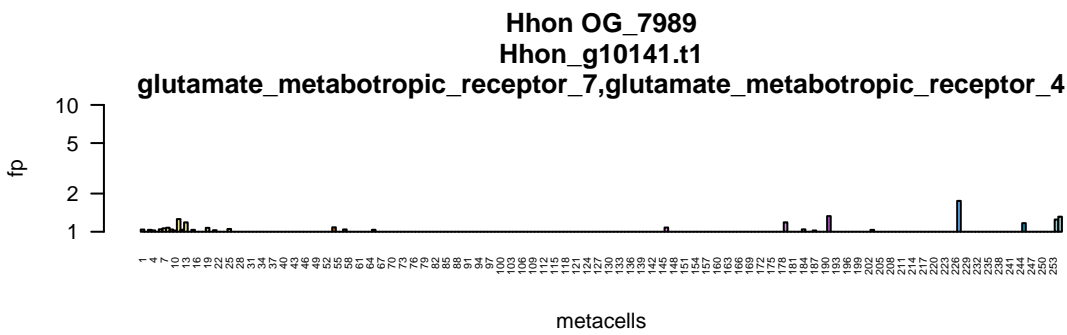
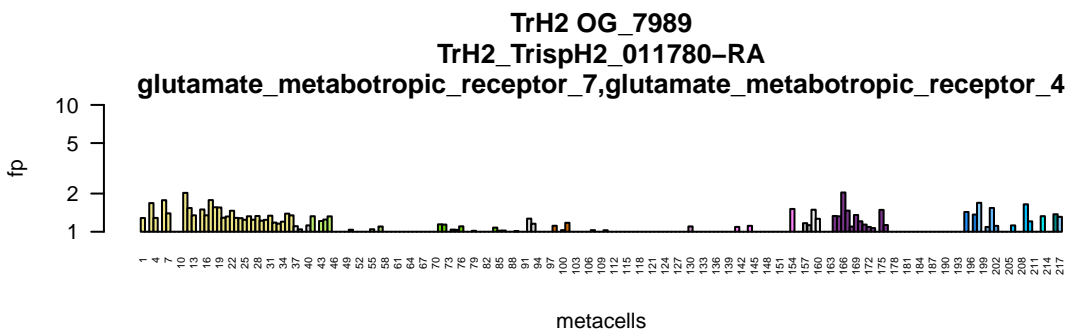
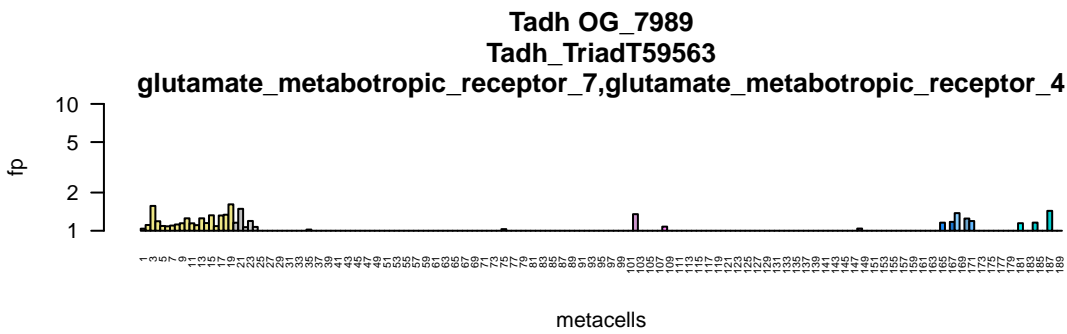


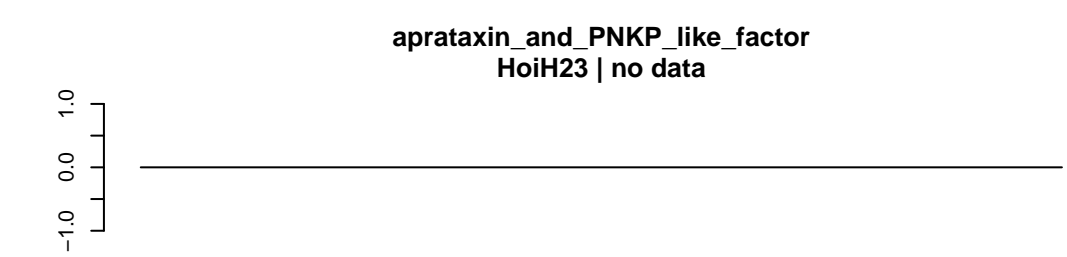
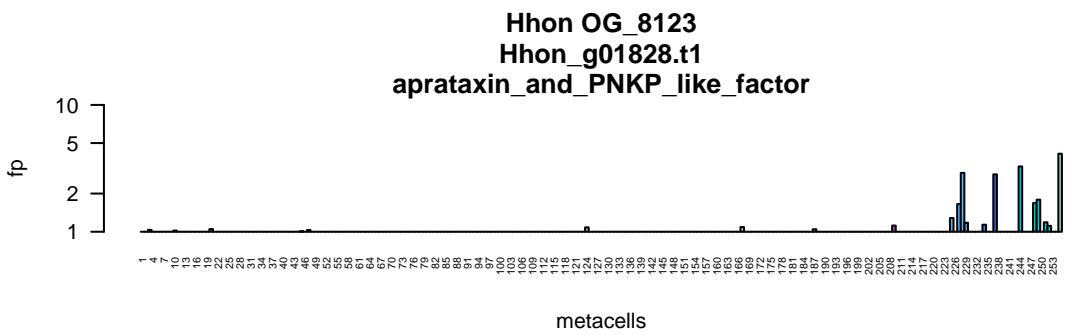
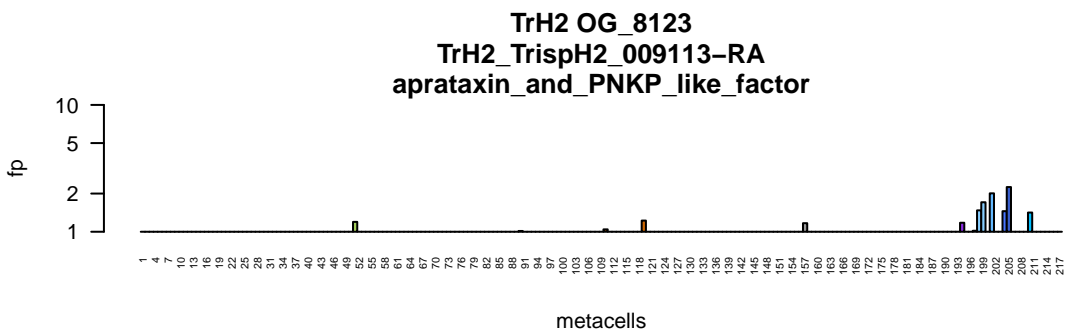
Hhon | no data

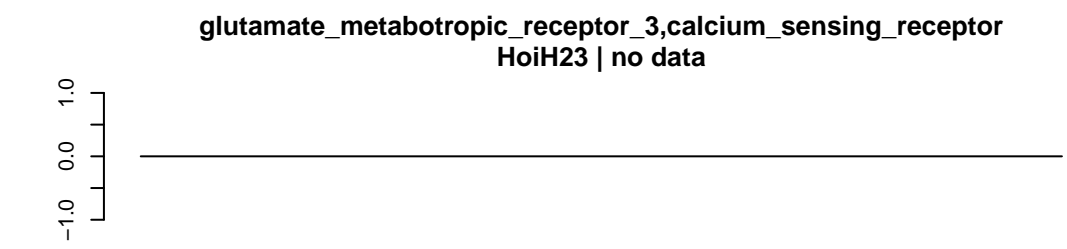
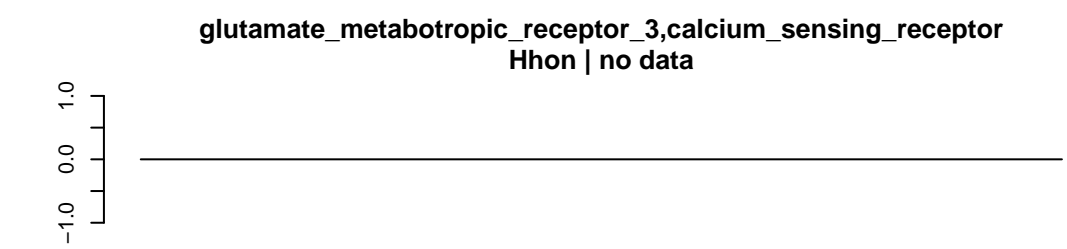
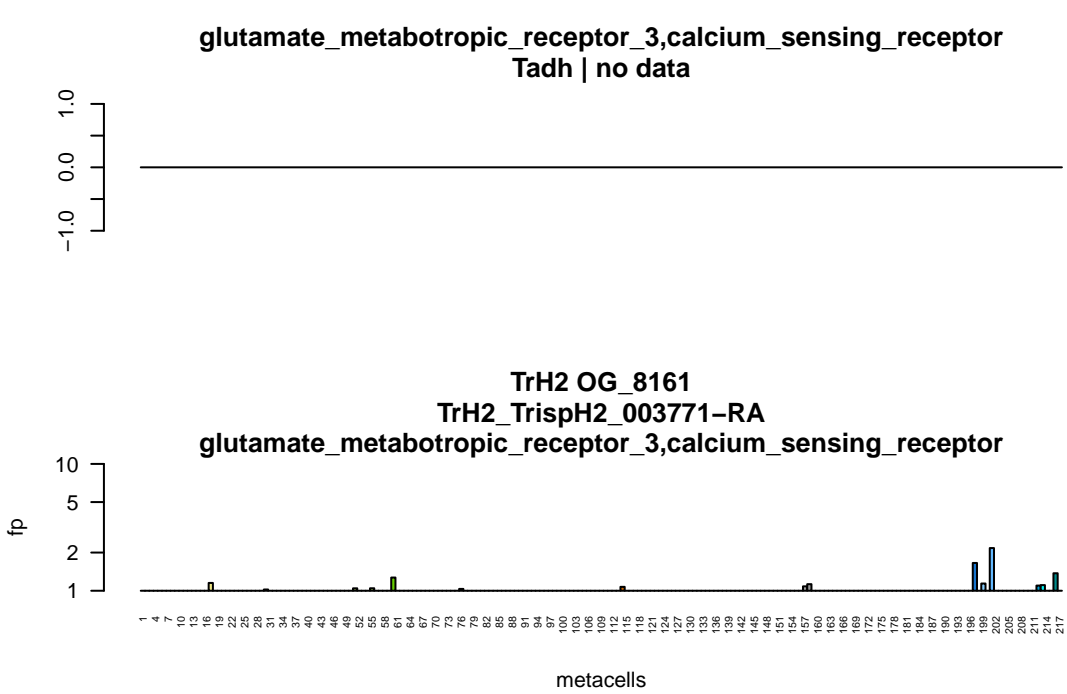


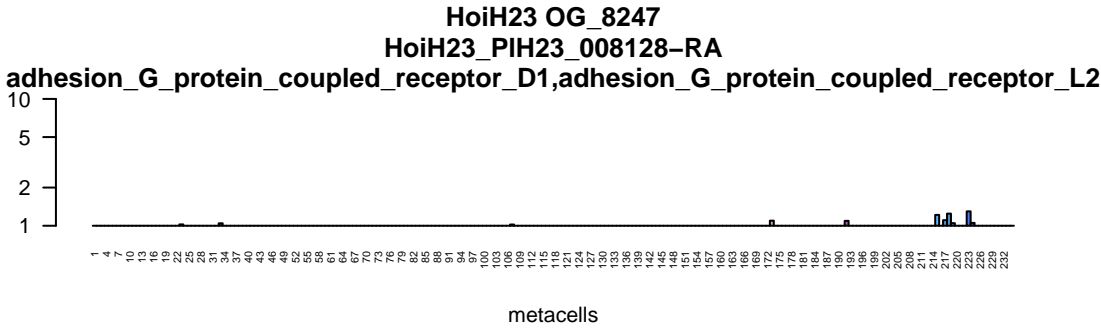
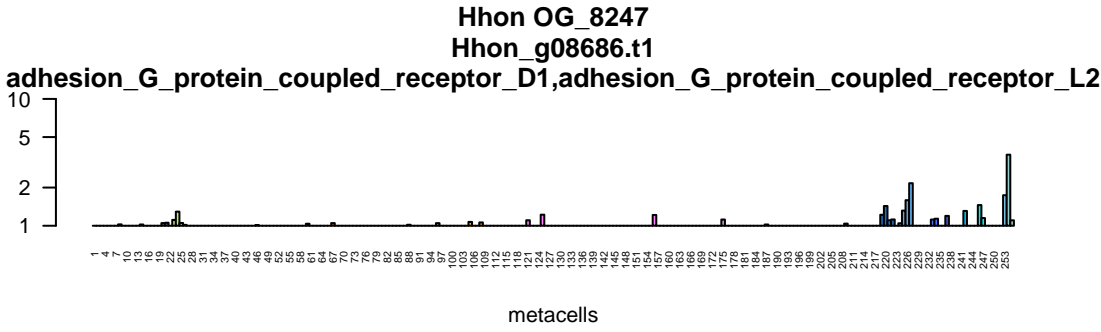
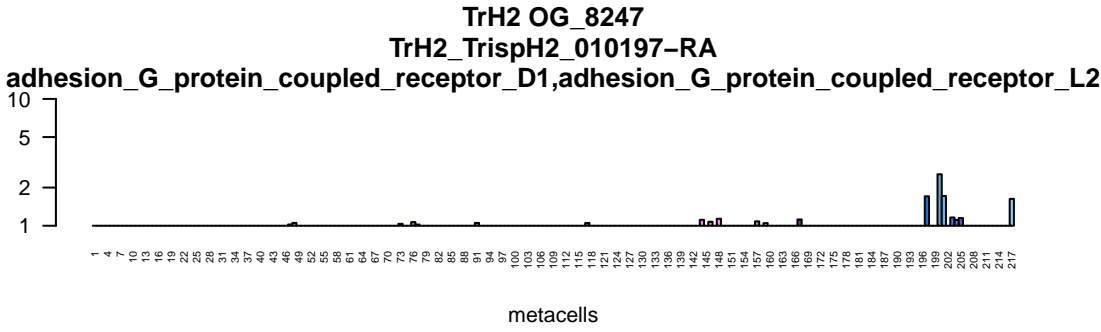
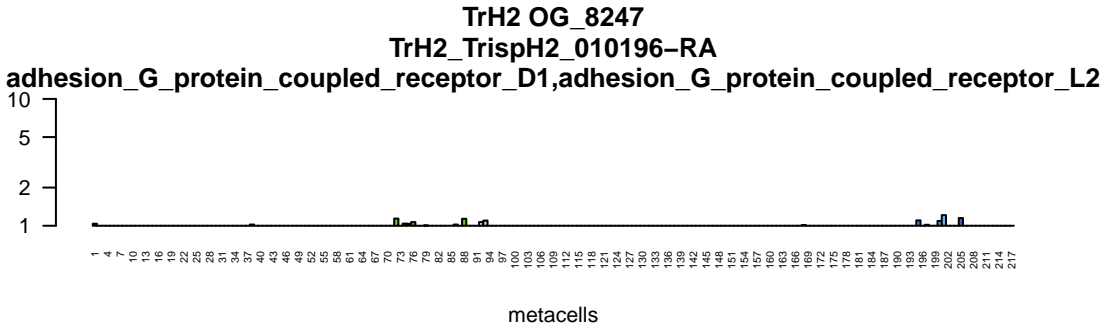
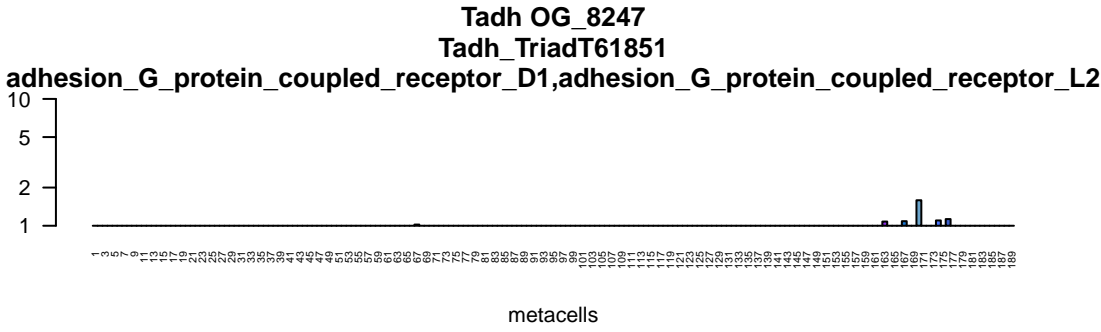
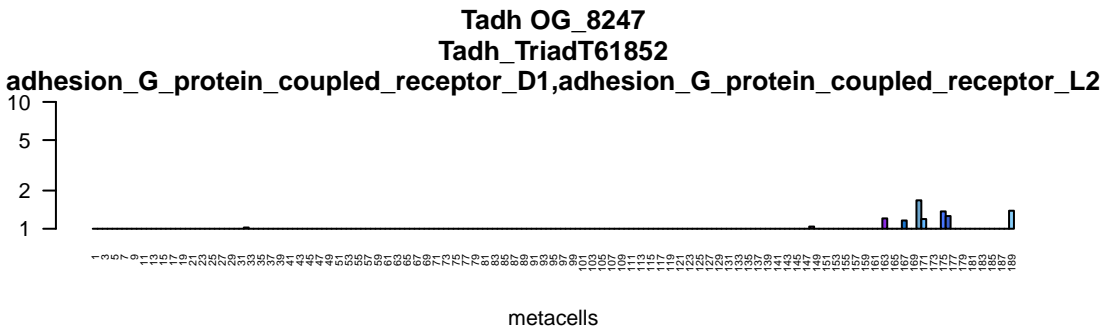
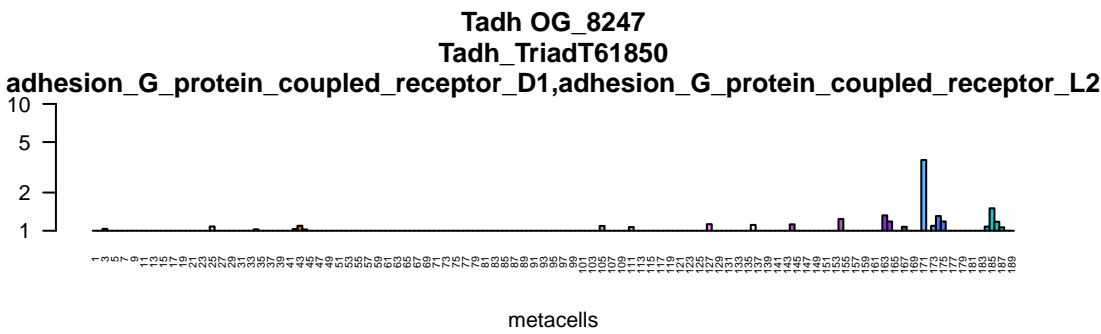
HoiH23 OG_7681
HoiH23_PIH23_006762-RA

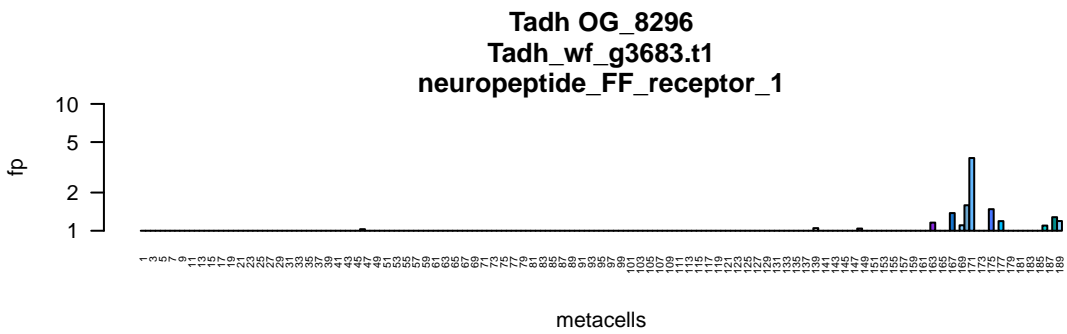




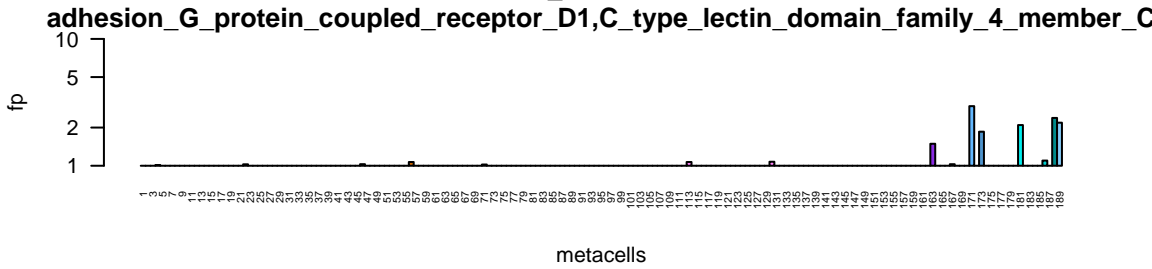




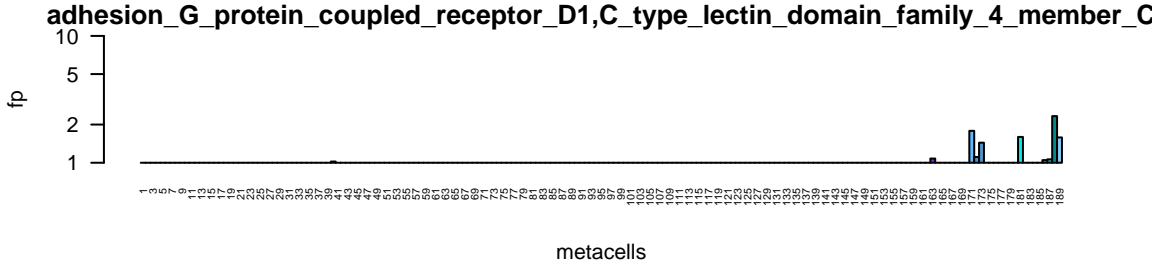




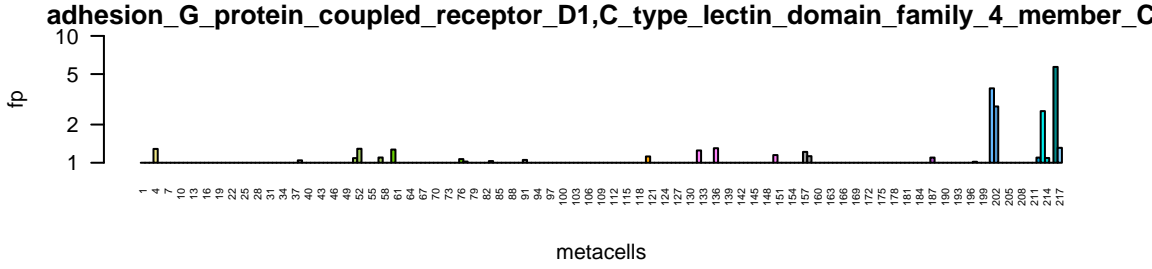
Tadh OG_8454
Tadh_TriadT58102



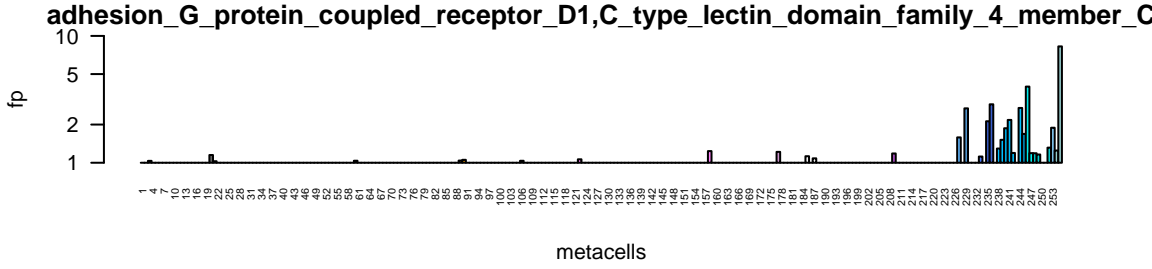
Tadh OG_8454
Tadh_TriadT58103



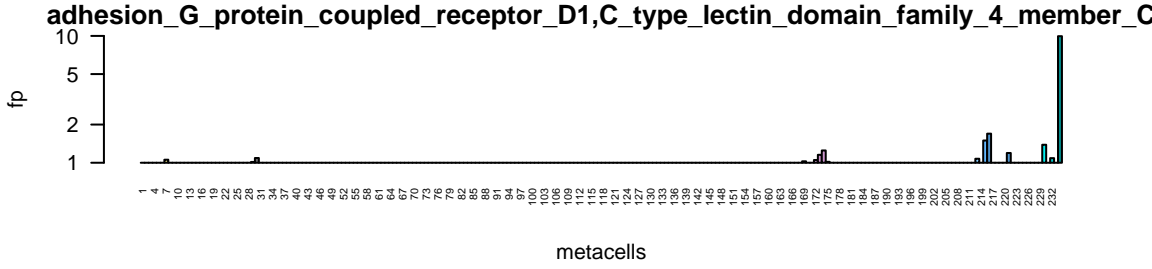
TrH2 OG_8454
TrH2_TrispH2_000572-RA

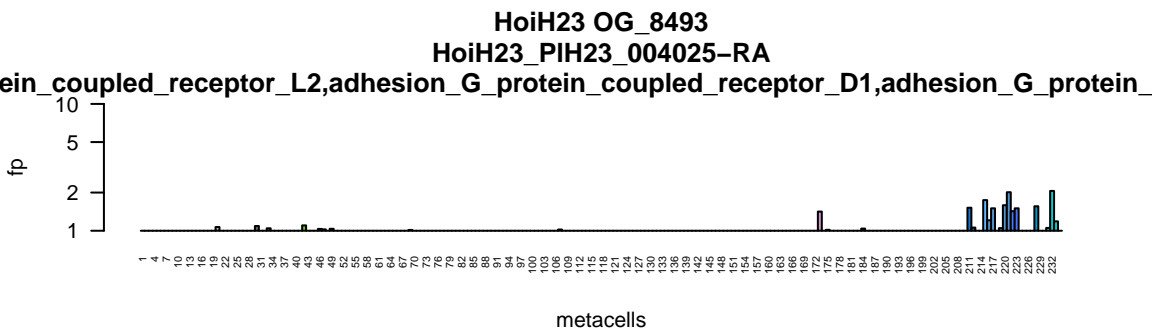
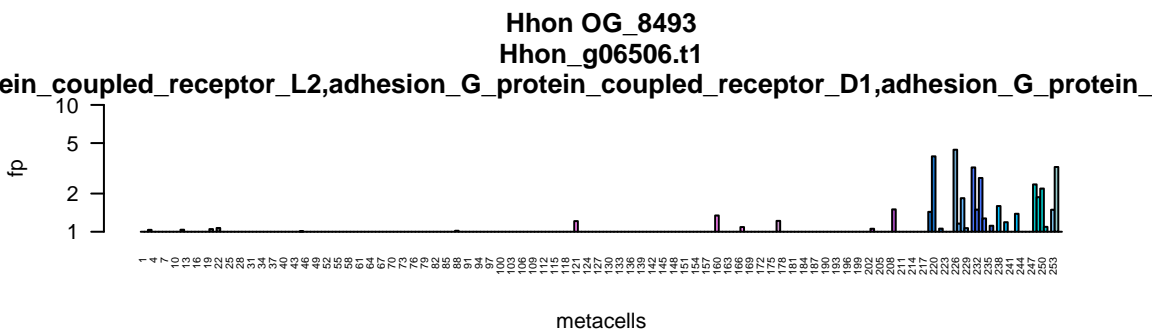
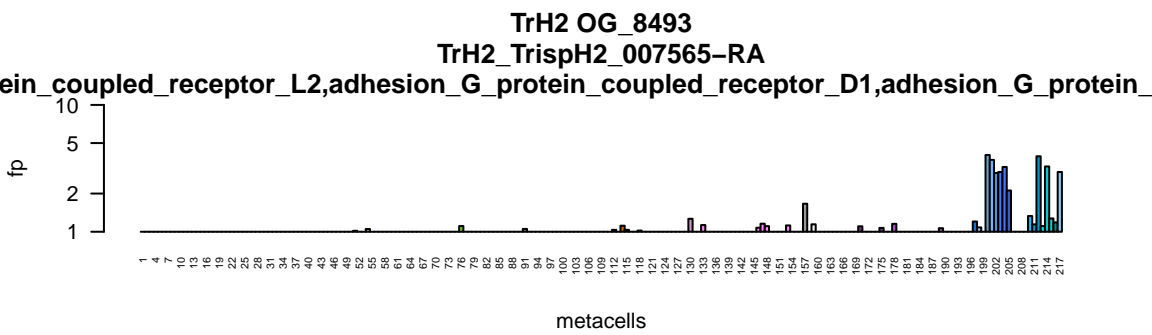
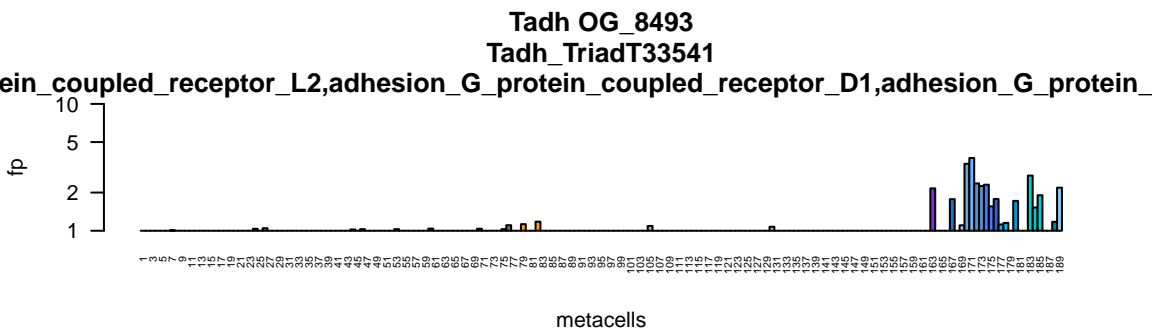


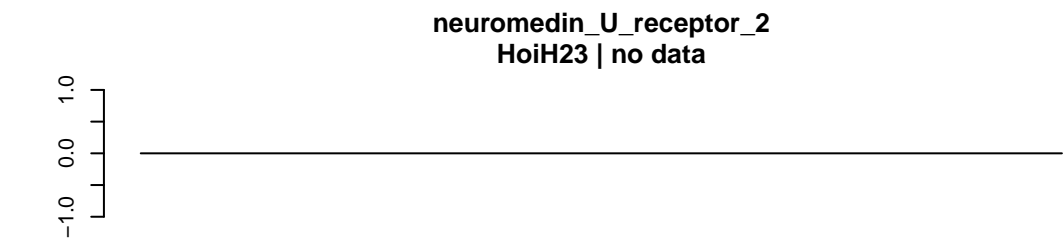
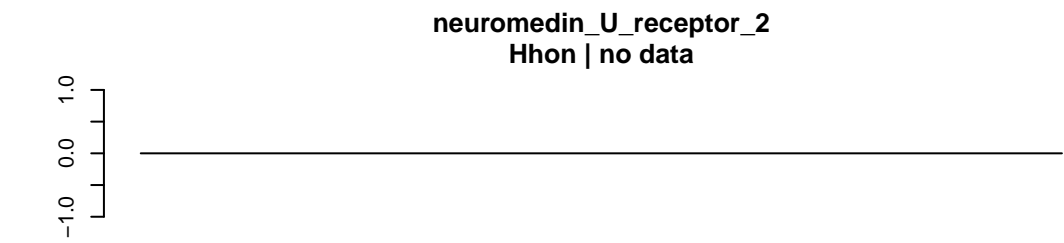
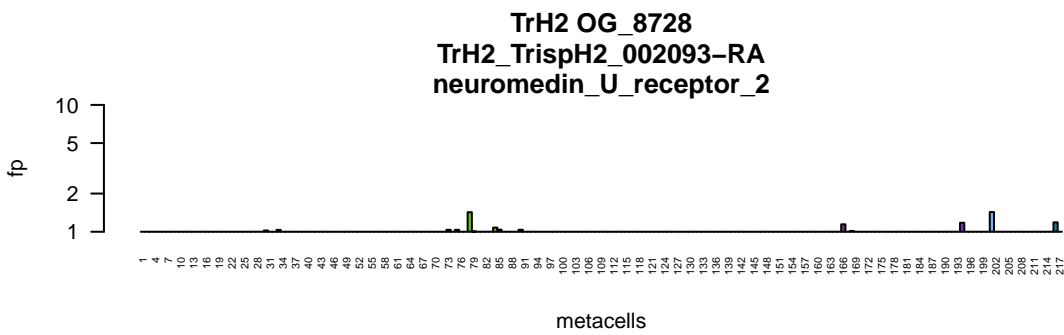
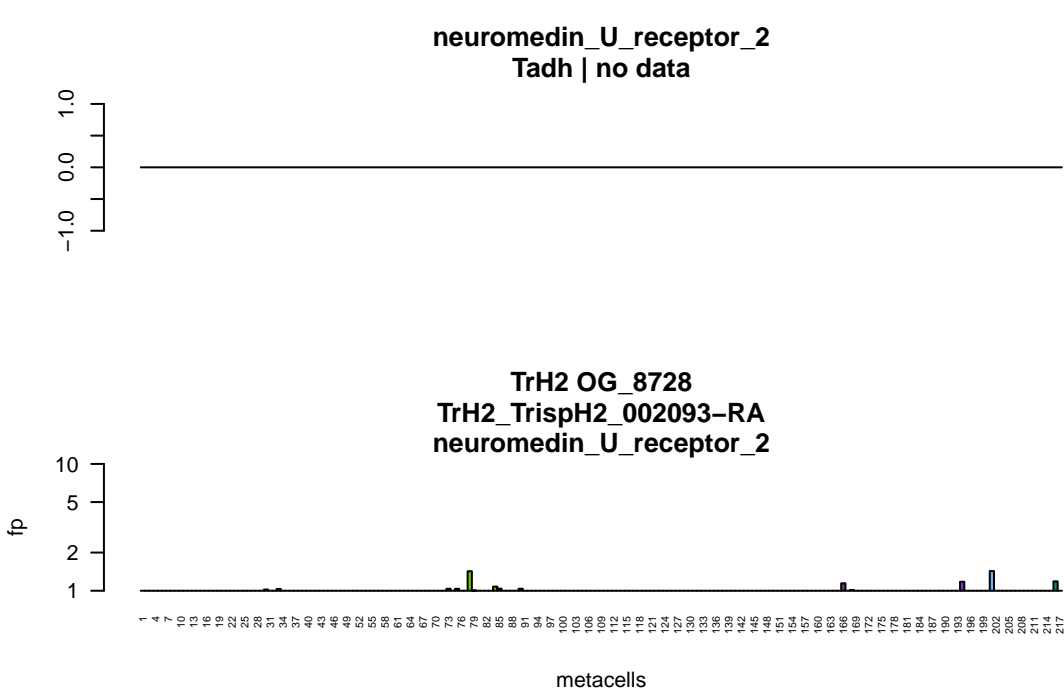
Hhon OG_8454
Hhon_g00116.t1

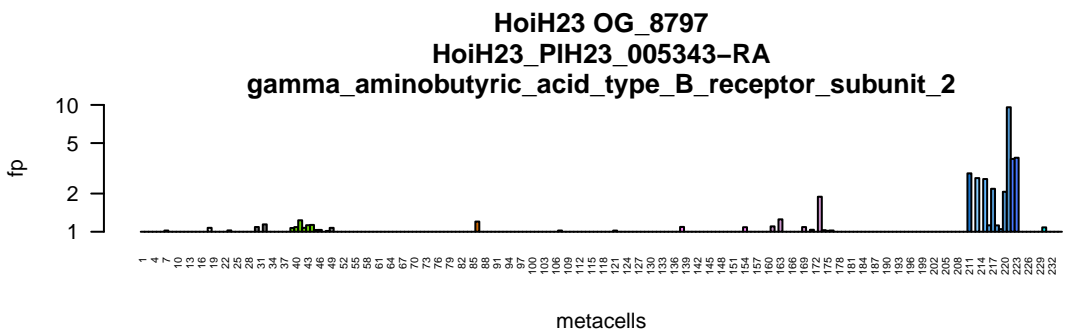
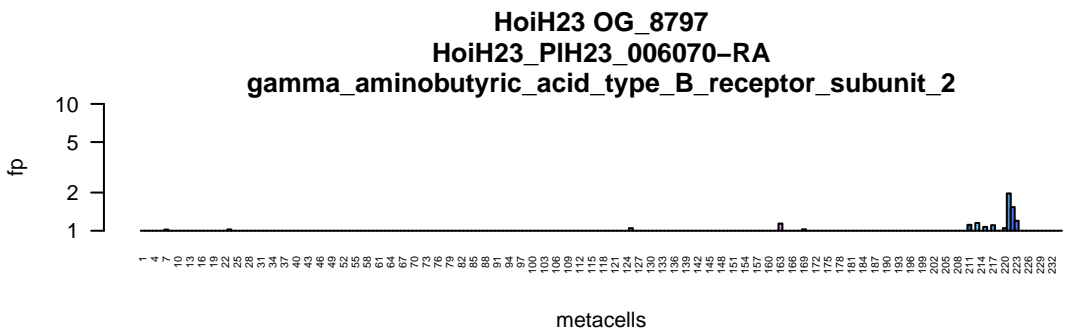
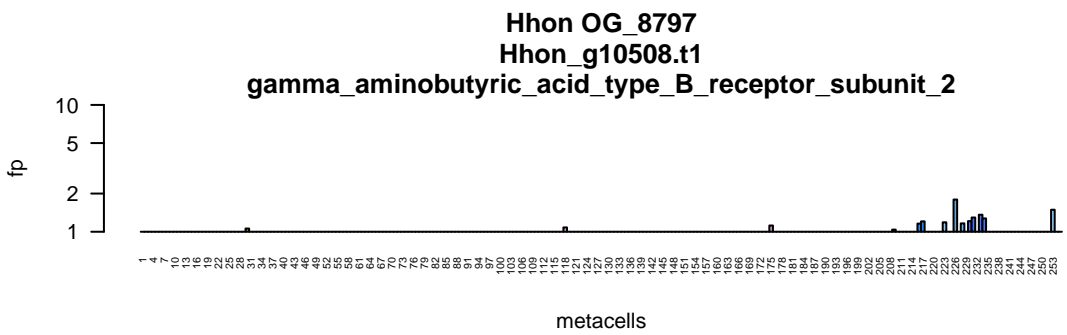
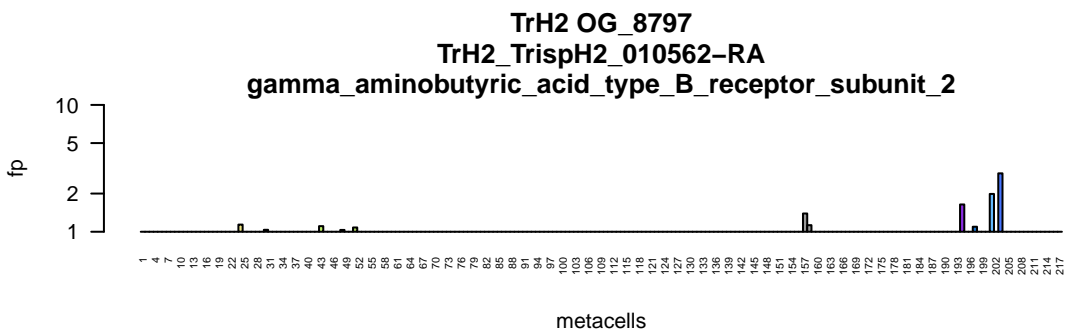
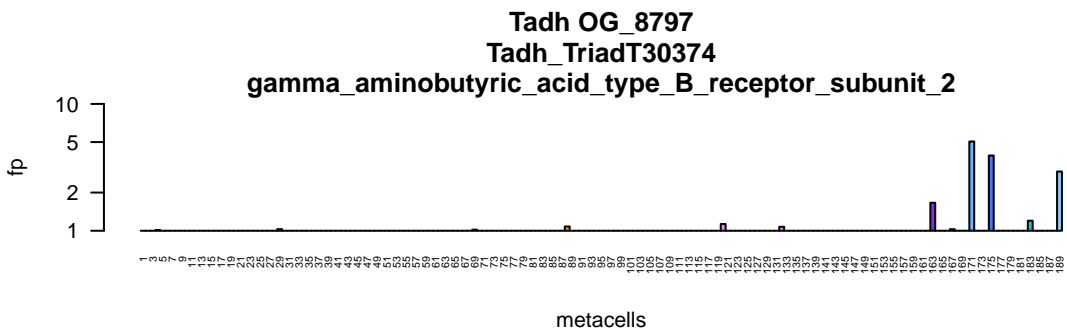


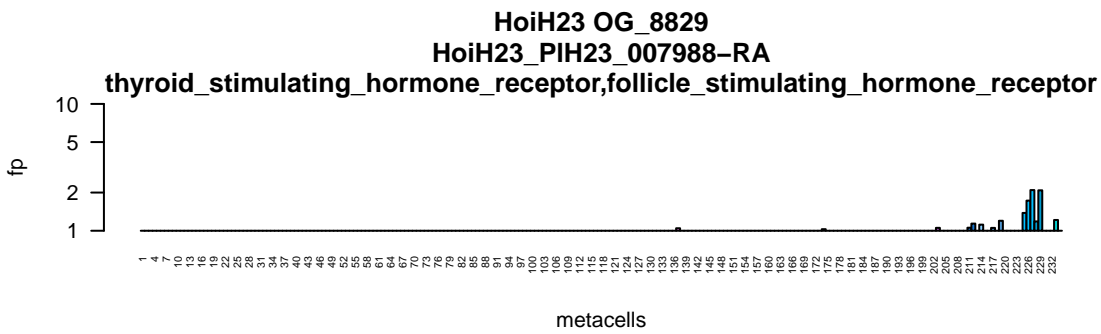
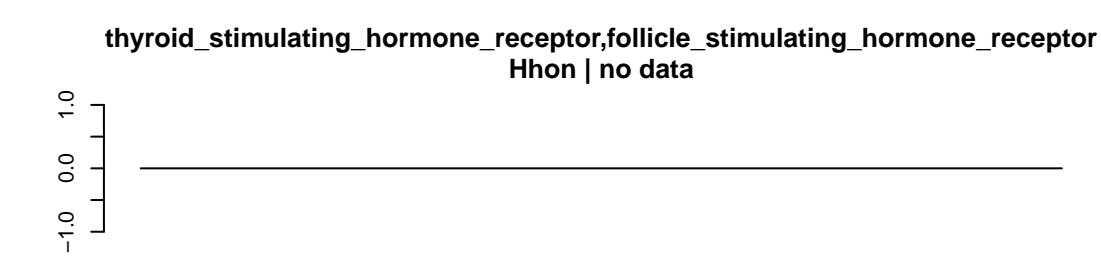
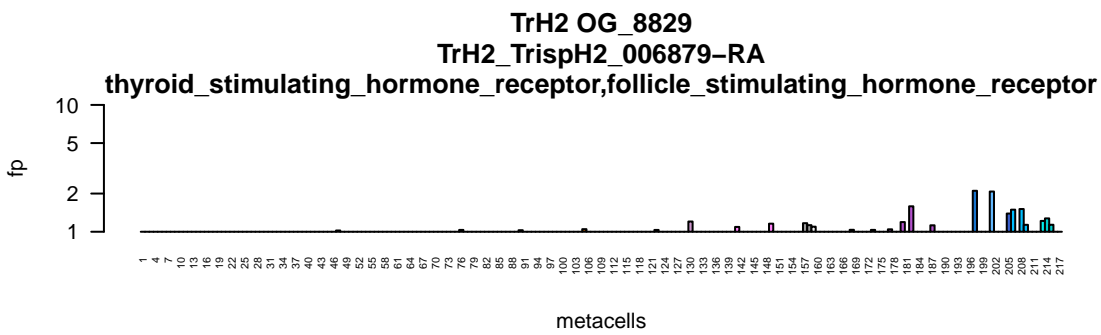
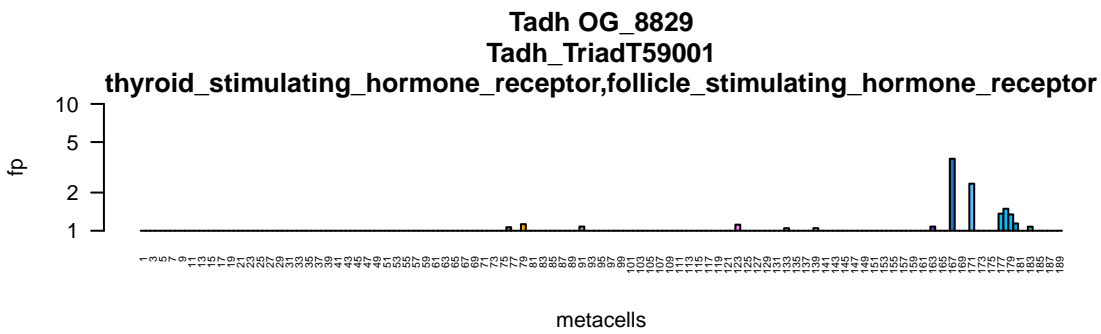
HoiH23 OG_8454
HoiH23_PIH23_002839-RA





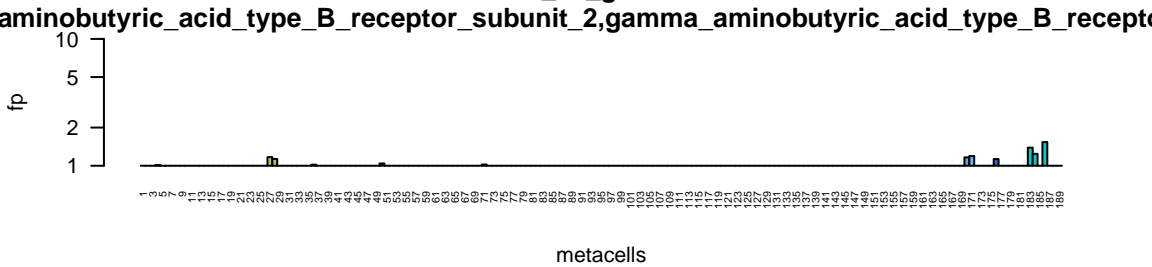






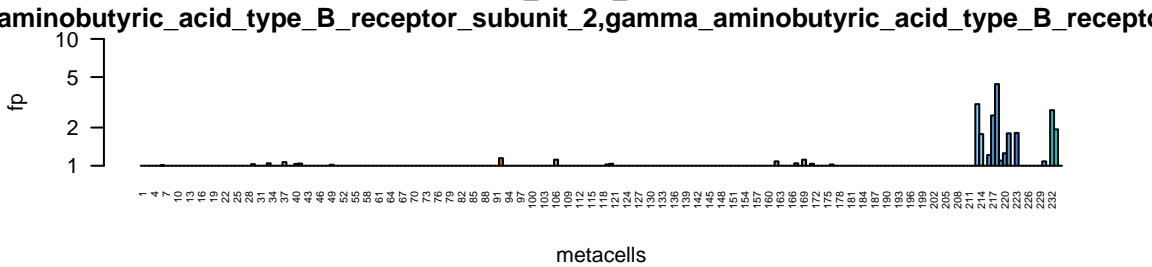
Tadh OG 8960

Tadh_wf_g11896.t1



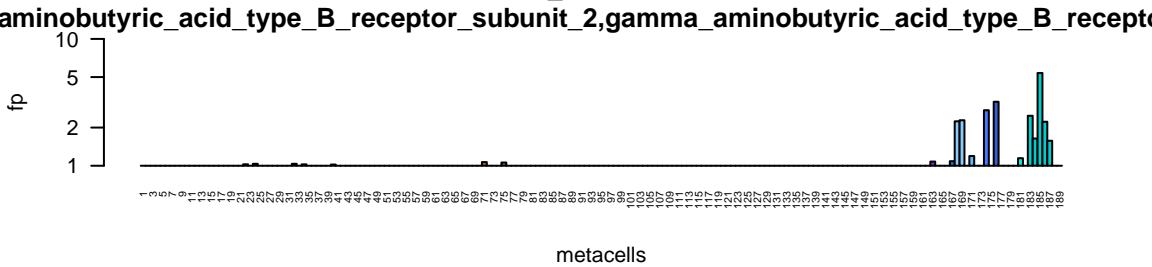
HoiH23 OG 8960

HoiH23_PIH23_000719-RA



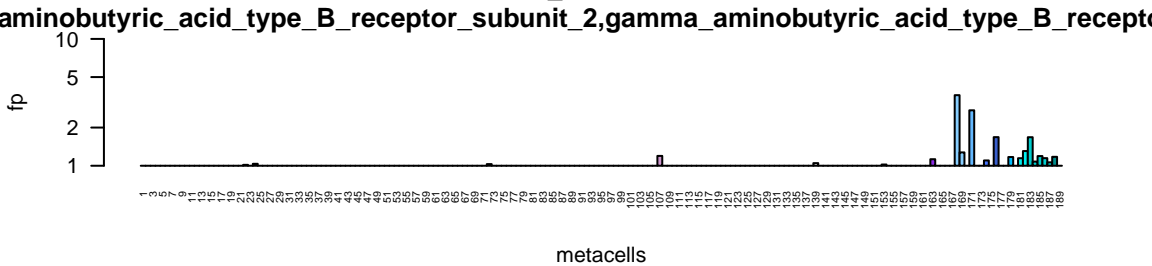
Tadh OG_8960

Tadh_TriadT52578



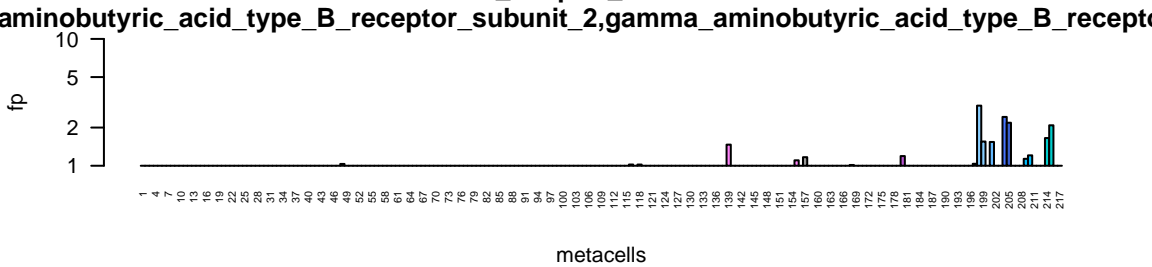
Tadh OG_8960

Tadh_TriadT52579



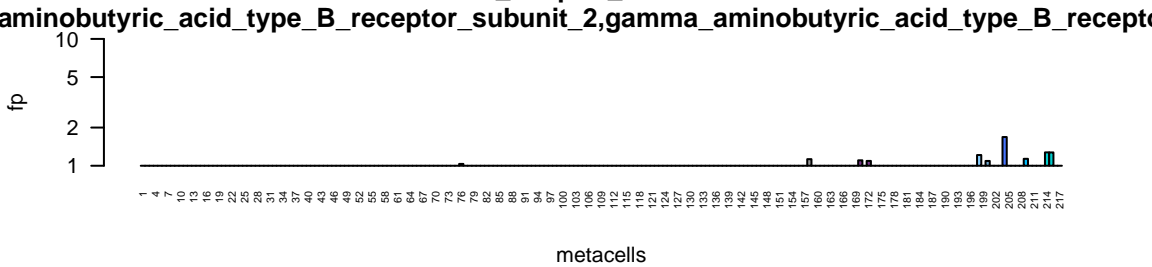
TrH2 OG_8960

TrH2_TrispH2_012100-RA



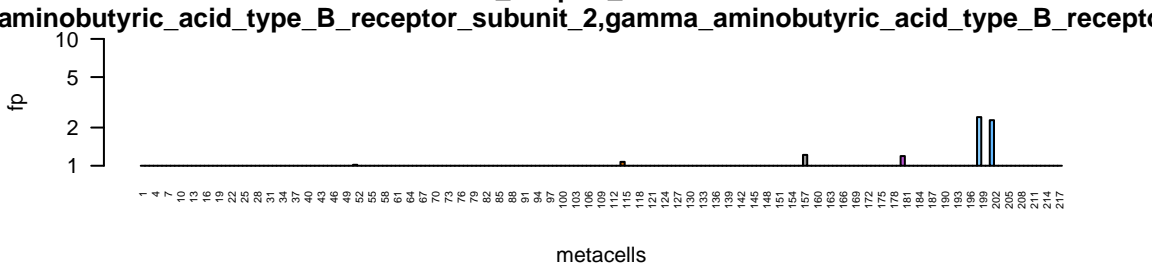
TrH2 OG_8960

TrH2_TrispH2_000234-RA



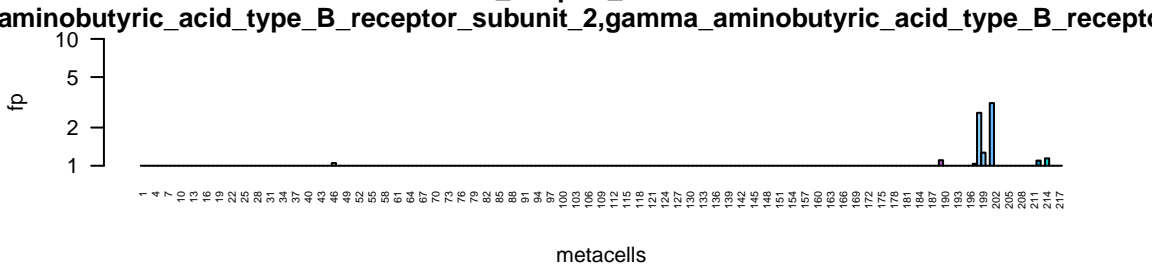
TrH2 OG_8960

TrH2_TrispH2_011964-RA



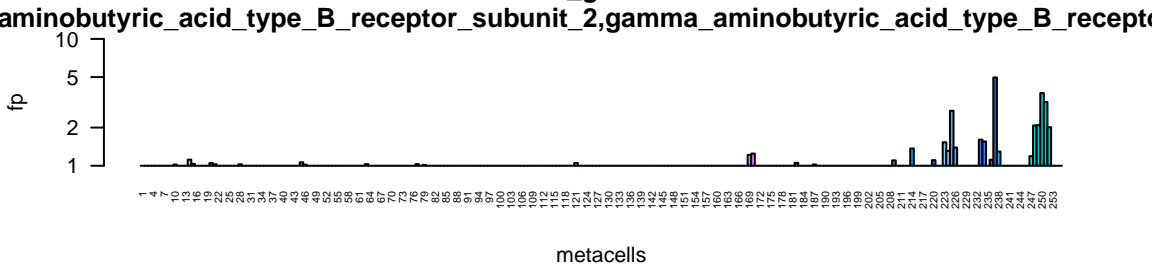
TrH2 OG_8960

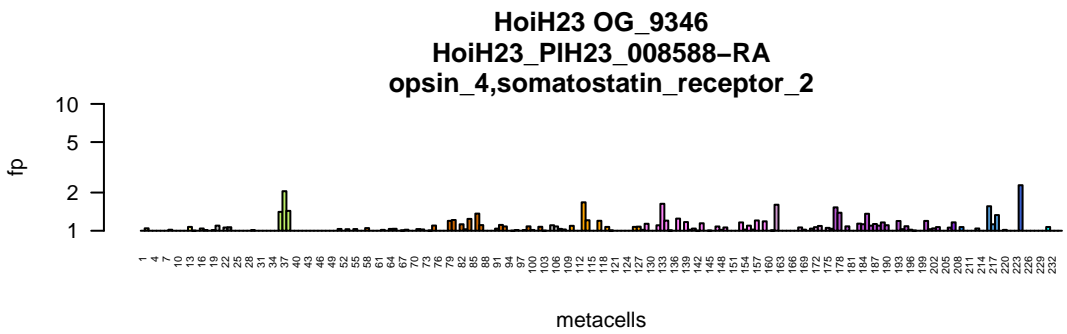
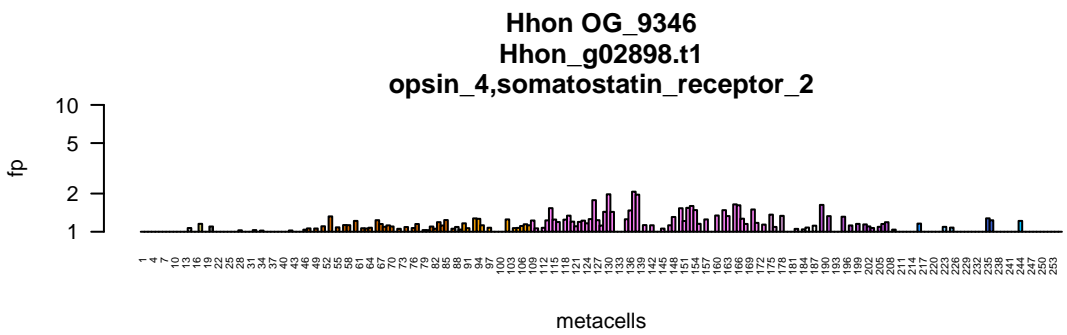
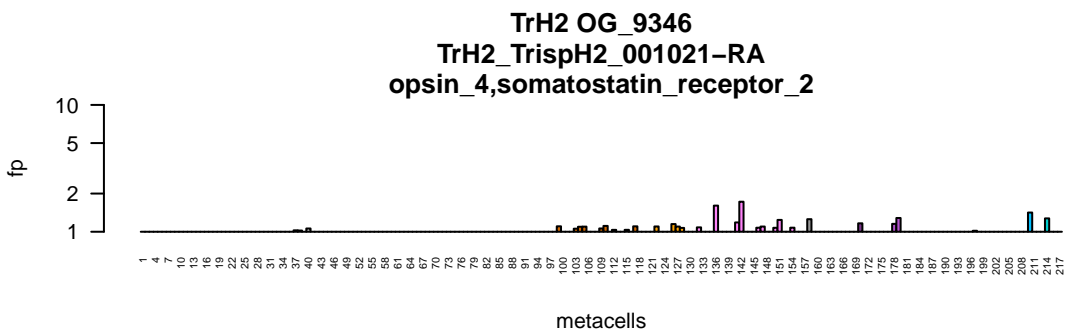
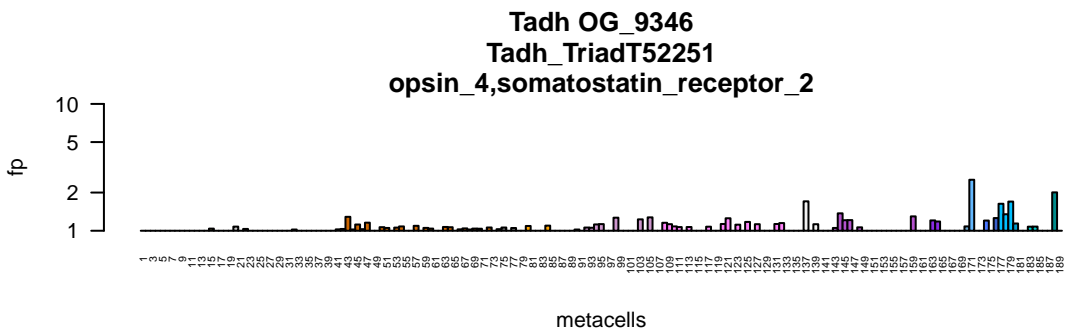
TrH2_TrispH2_011847-RA

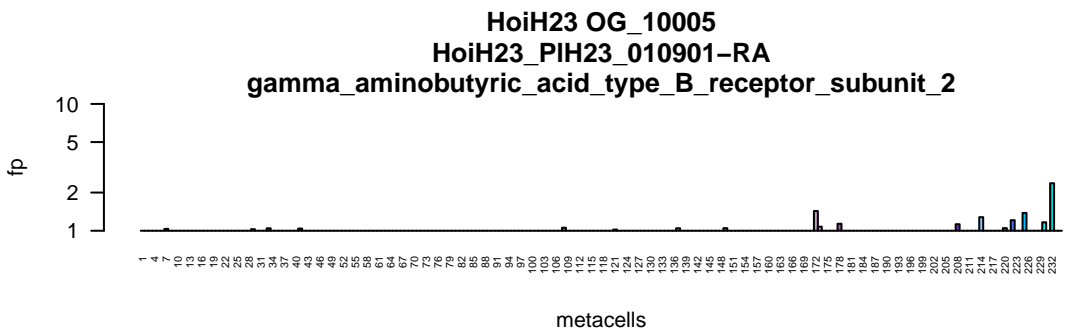


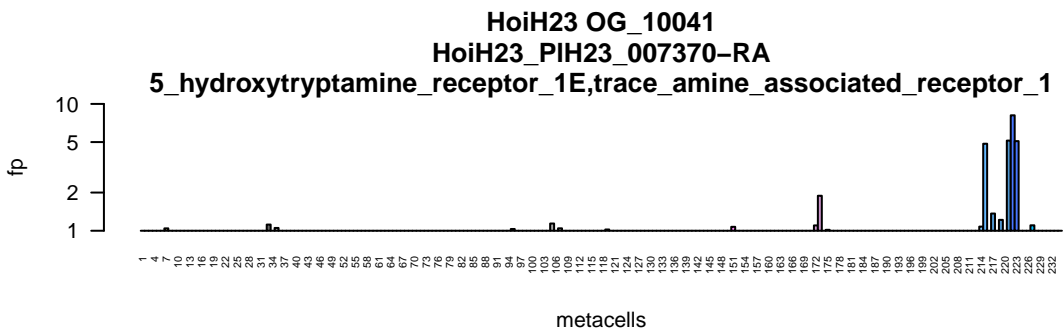
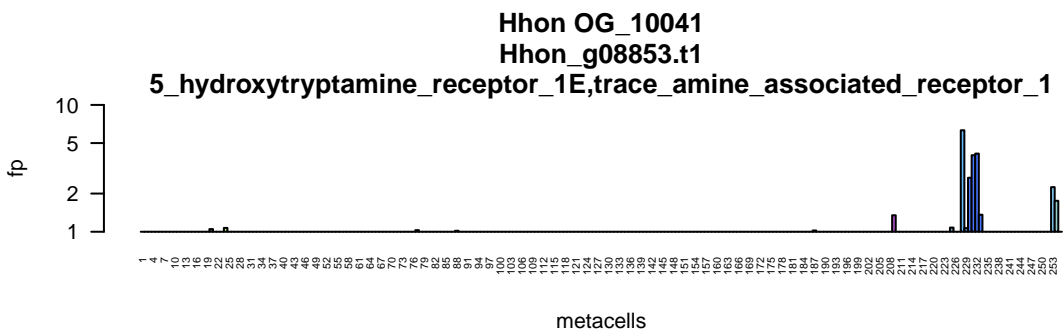
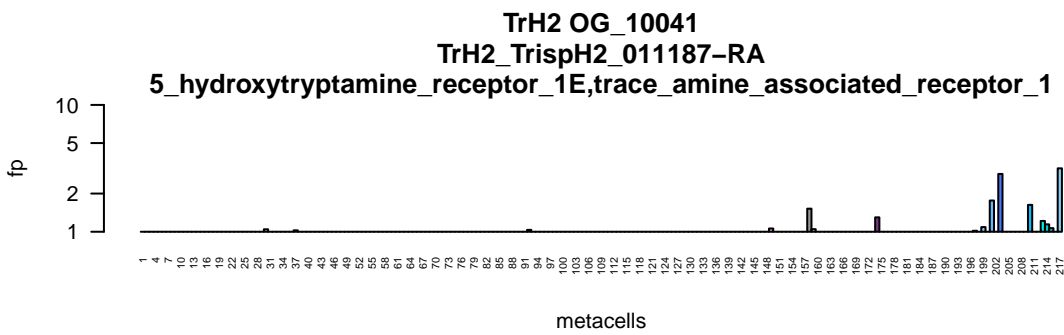
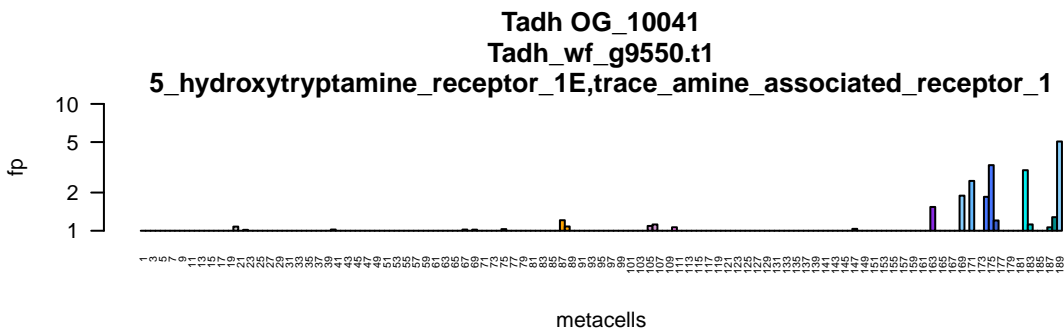
Hhon OG_8960

Hhon_g05296.t1

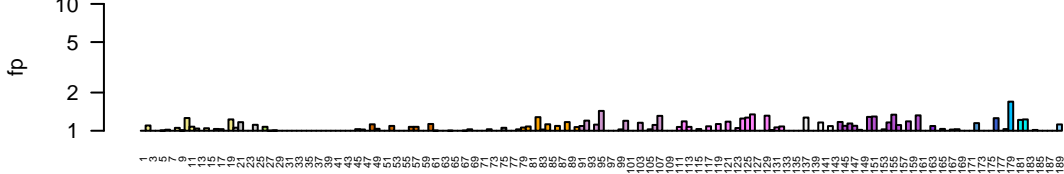






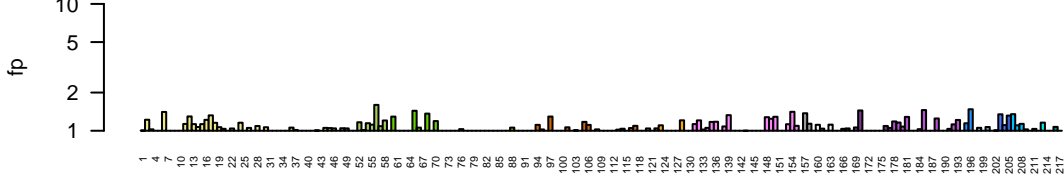


Tadh OG_10052
Tadh_TriadT3190
activin_A_receptor_type_2B



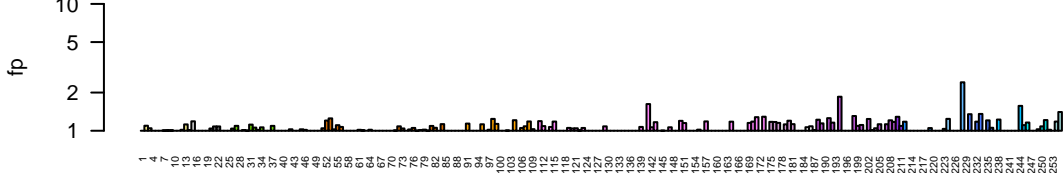
metacells

TrH2 OG_10052
TrH2_TrispH2_008797-RA
activin_A_receptor_type_2B



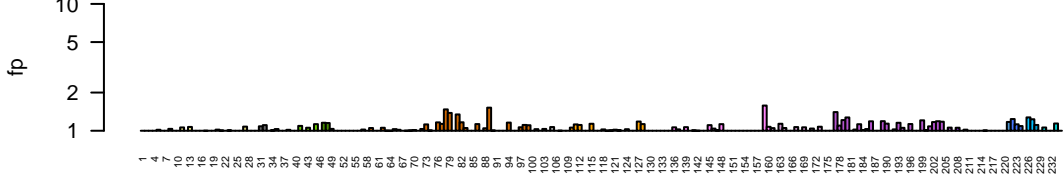
metacells

Hhon OG_10052
Hhon_g08874.t1
activin_A_receptor_type_2B



metacells

HoiH23 OG_10052
HoiH23_PIH23_007390-RA
activin_A_receptor_type_2B



metacells

