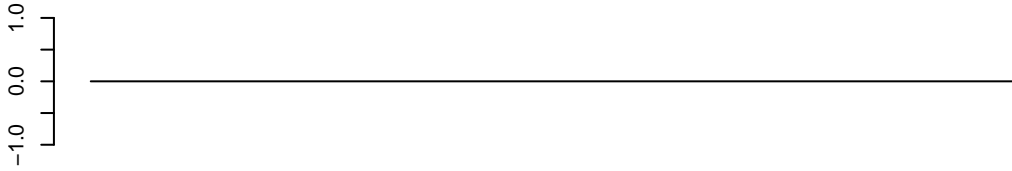
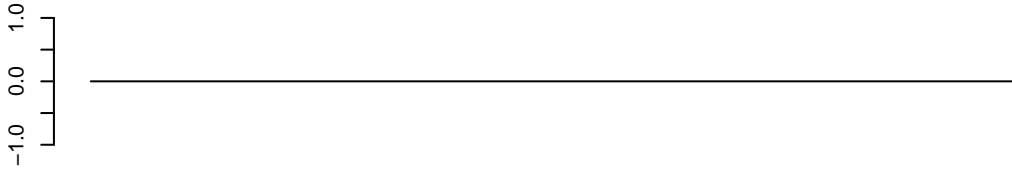


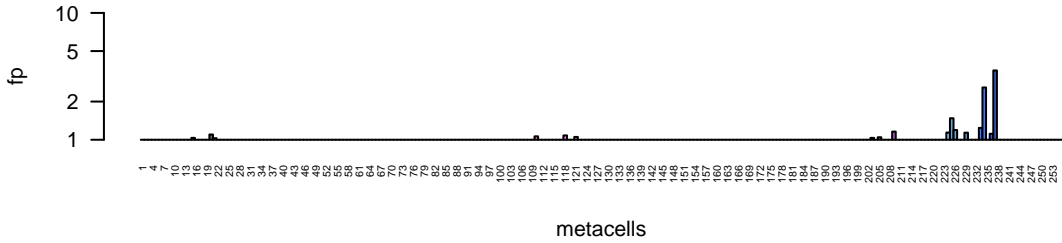
Tadh | no data



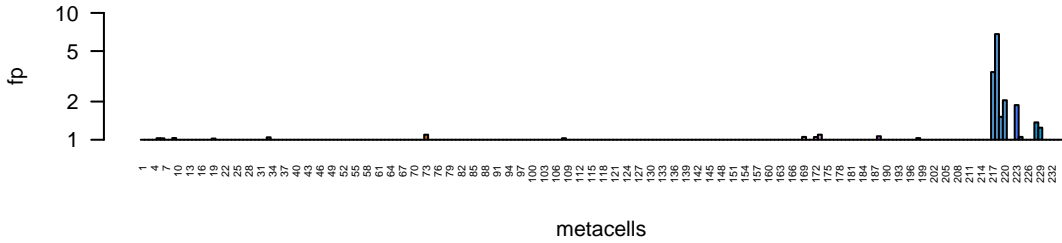
TrH2 | no data



Hhon OG_6709
Hhon_g01989.t1



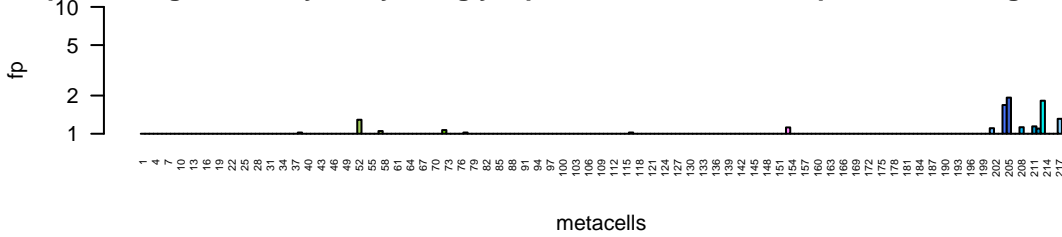
HoiH23 OG_6709
HoiH23_PIH23_003664-RA



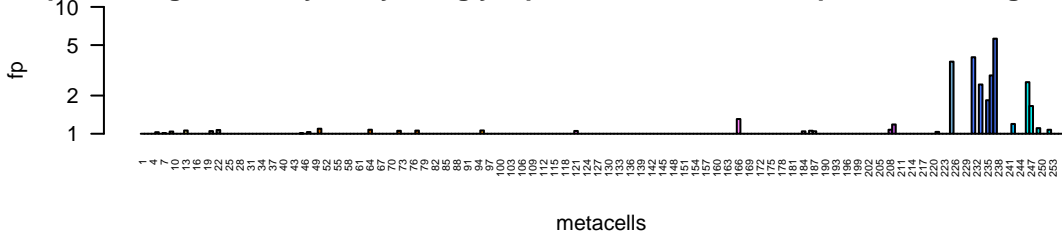
receptor_1,oligodendrocyte_myelin_glycoprotein,leucine_rich_repeat_containing_40,relaxir
Tadh | no data



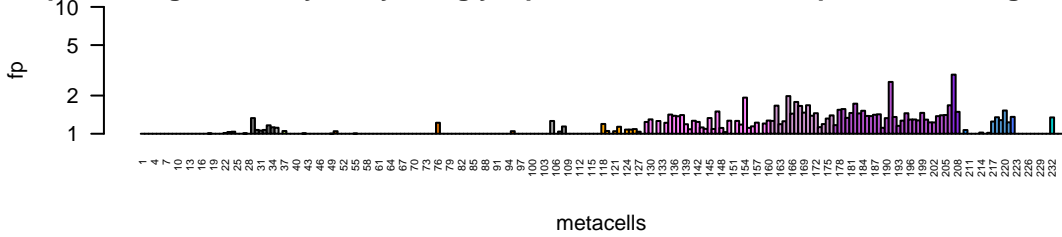
TrH2 OG_6800
TrH2_TrispH2_006274-RA
receptor_1,oligodendrocyte_myelin_glycoprotein,leucine_rich_repeat_containing_40,relaxir



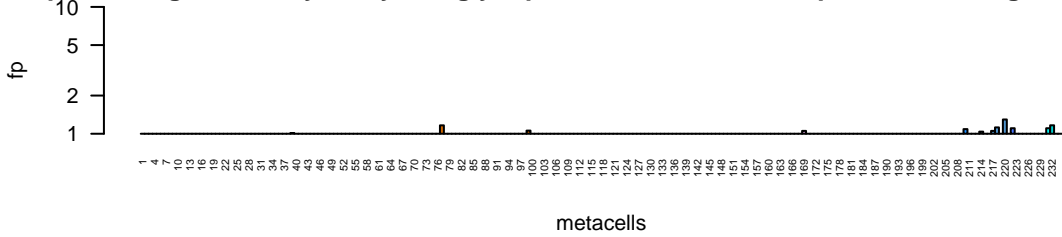
Hhon OG_6800
Hhon_g04761.t1
receptor_1,oligodendrocyte_myelin_glycoprotein,leucine_rich_repeat_containing_40,relaxir



HoiH23 OG_6800
HoiH23_PIH23_001555-RA
receptor_1,oligodendrocyte_myelin_glycoprotein,leucine_rich_repeat_containing_40,relaxir



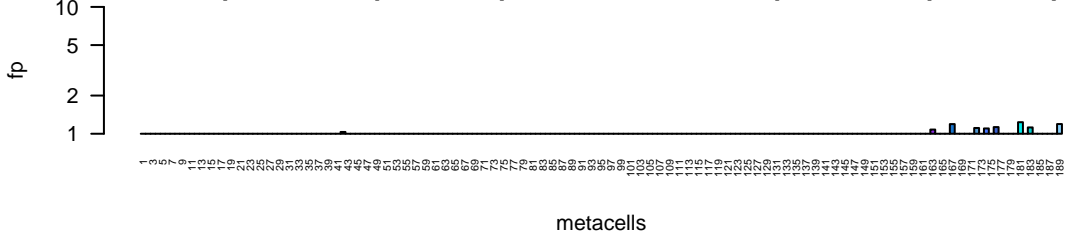
HoiH23 OG_6800
HoiH23_PIH23_001554-RA
receptor_1,oligodendrocyte_myelin_glycoprotein,leucine_rich_repeat_containing_40,relaxir



Tadh OG_7054

Tadh_TriadT59349

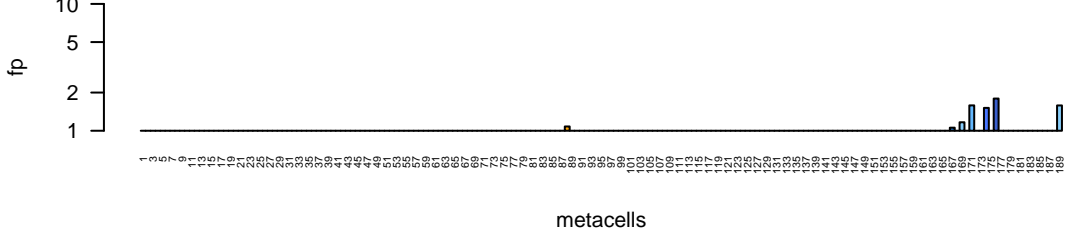
adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3



Tadh OG_7054

Tadh_TriadT59345

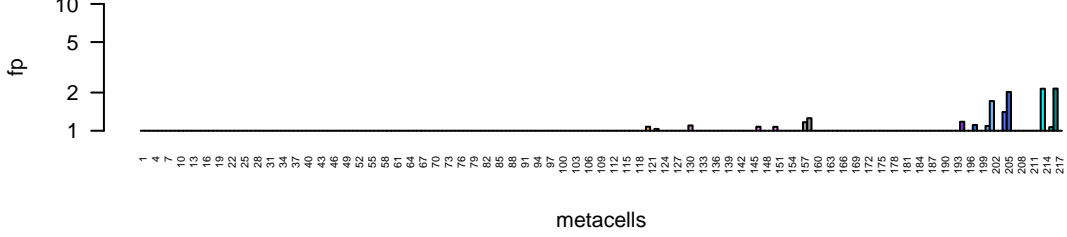
adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3



TrH2 OG_7054

TrH2_TrispH2_011841-RA

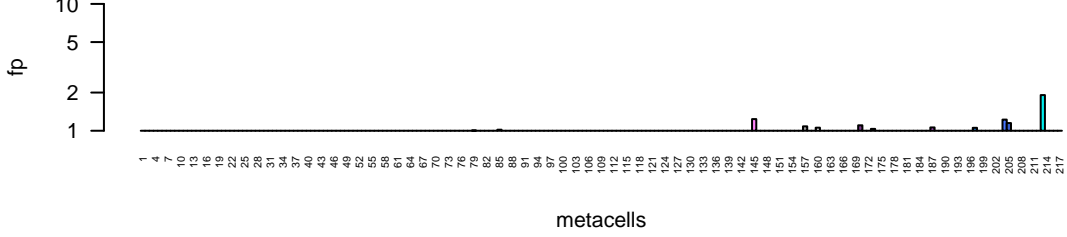
adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3



TrH2 OG_7054

TrH2_TrispH2_010220-RA

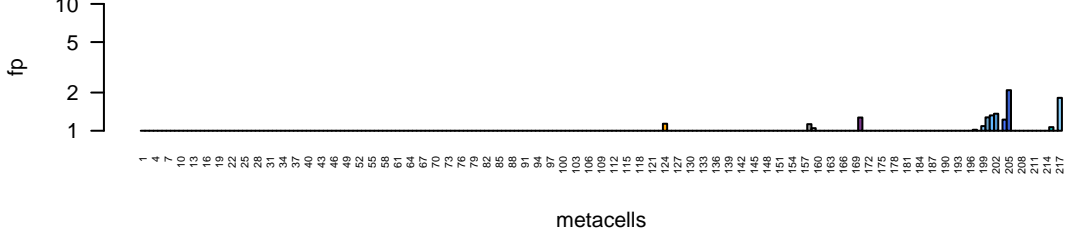
adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3



TrH2 OG_7054

TrH2_TrispH2_010222-RA

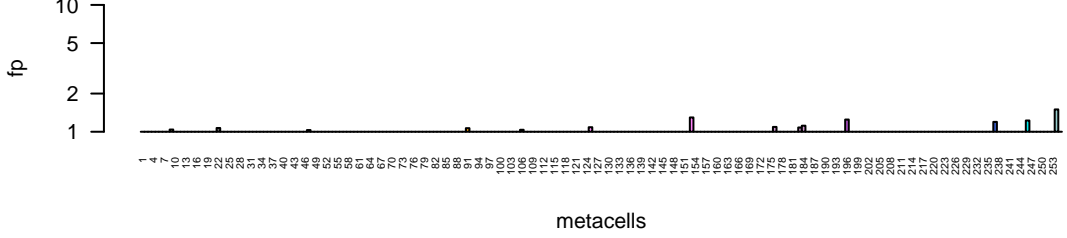
adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3



Hhon OG_7054

Hhon_g10892.t1

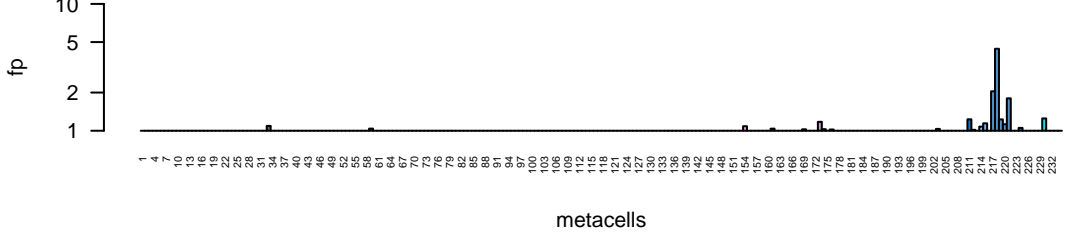
adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3



HoiH23 OG_7054

HoiH23_PIH23_010401-RA

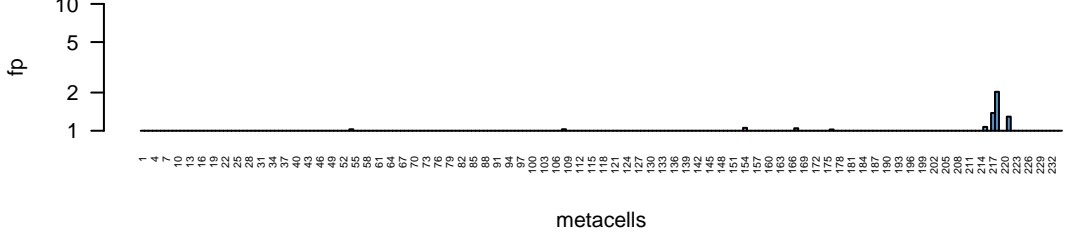
adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3

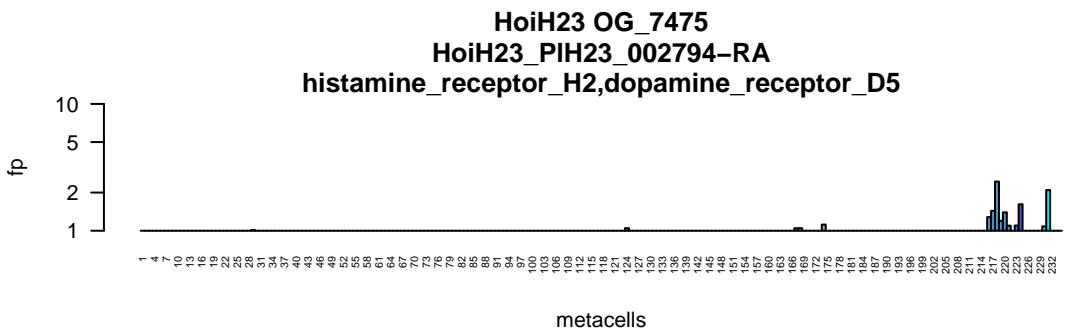
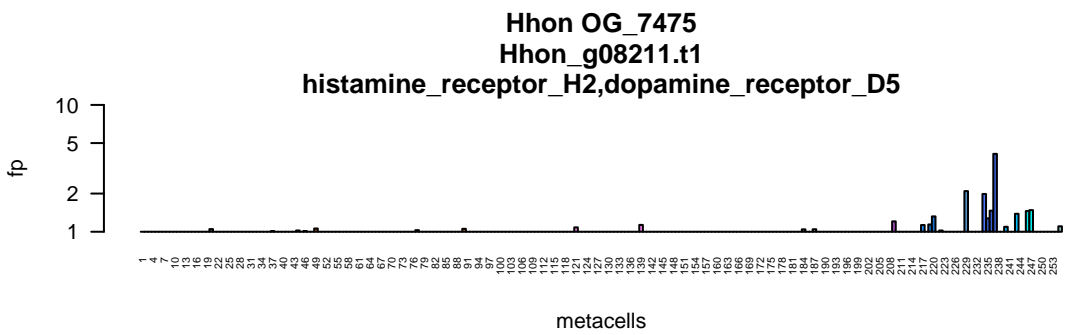
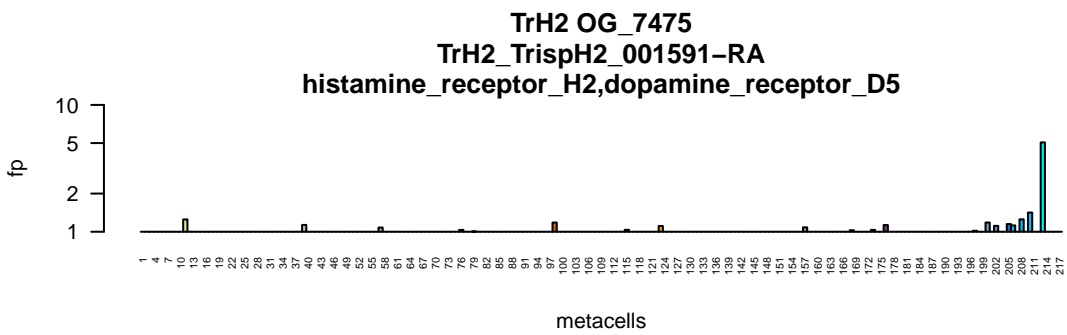
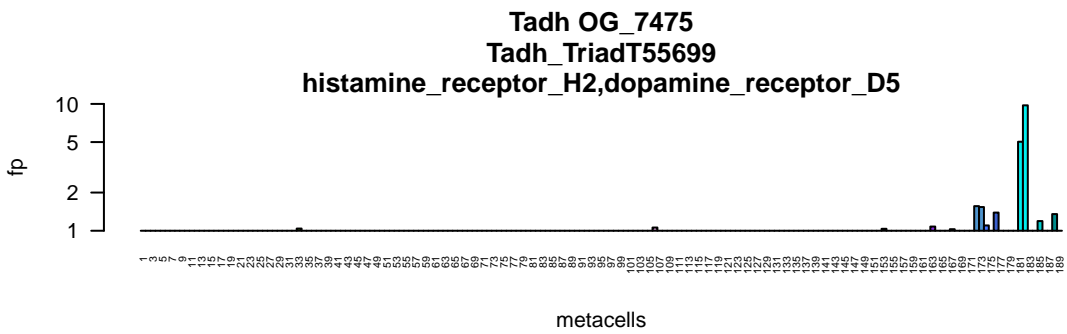


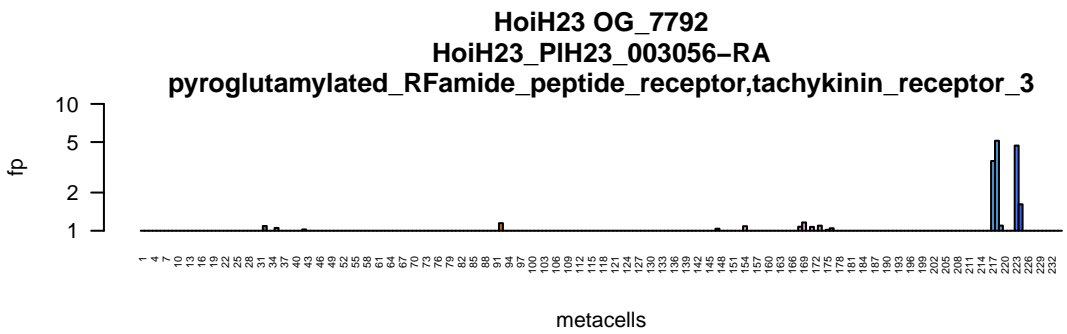
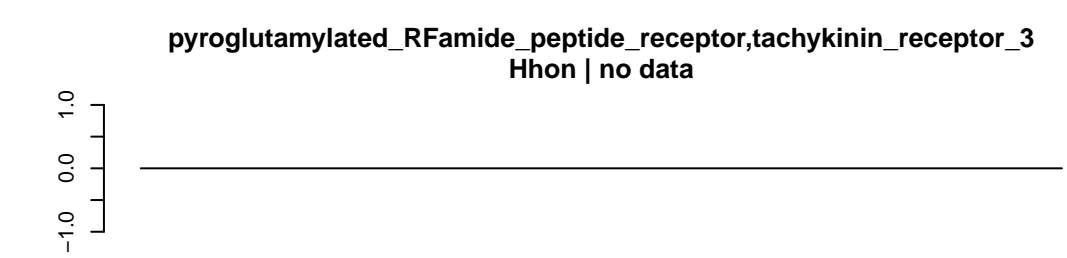
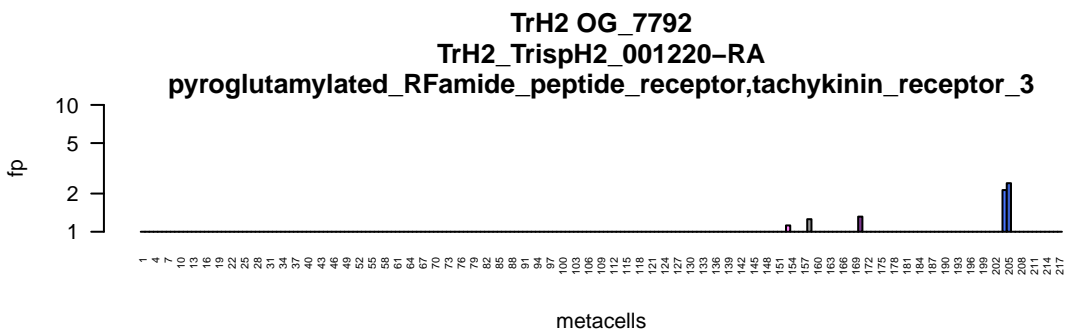
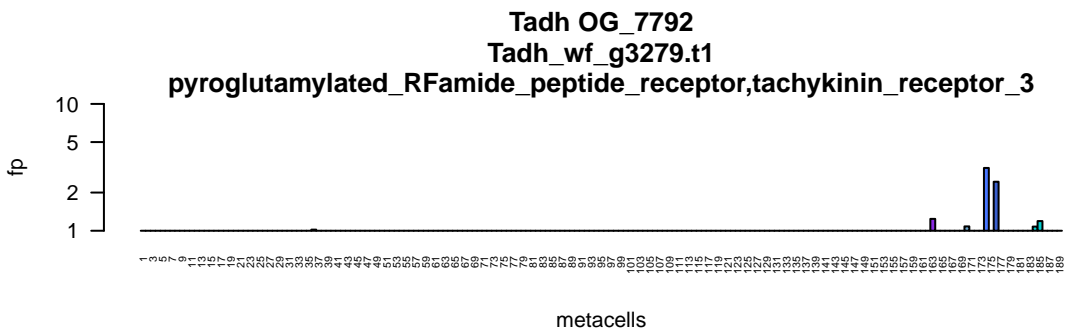
HoiH23 OG_7054

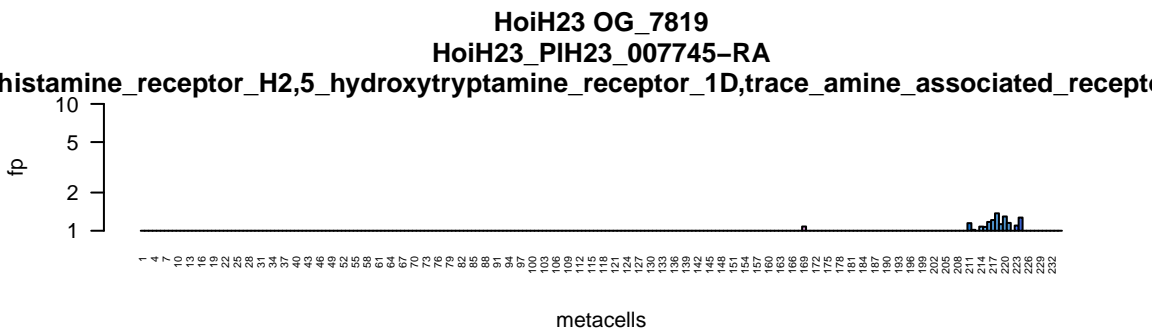
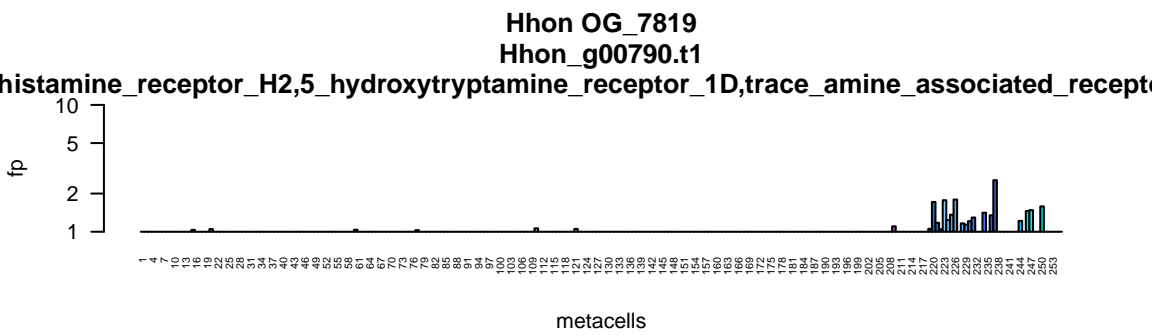
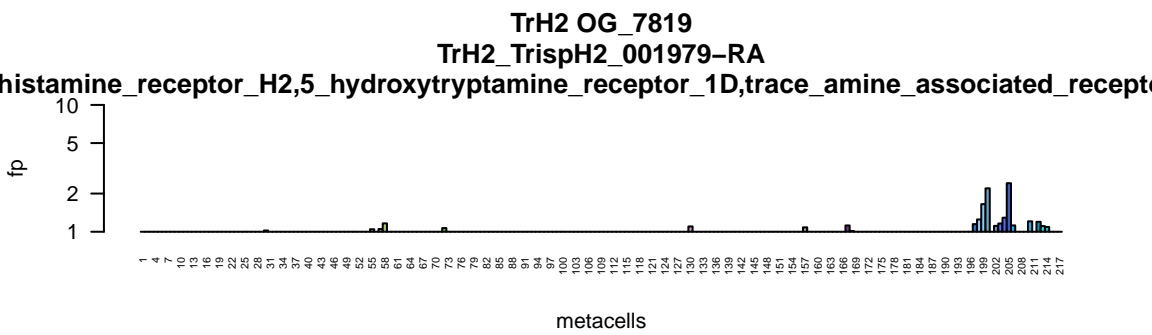
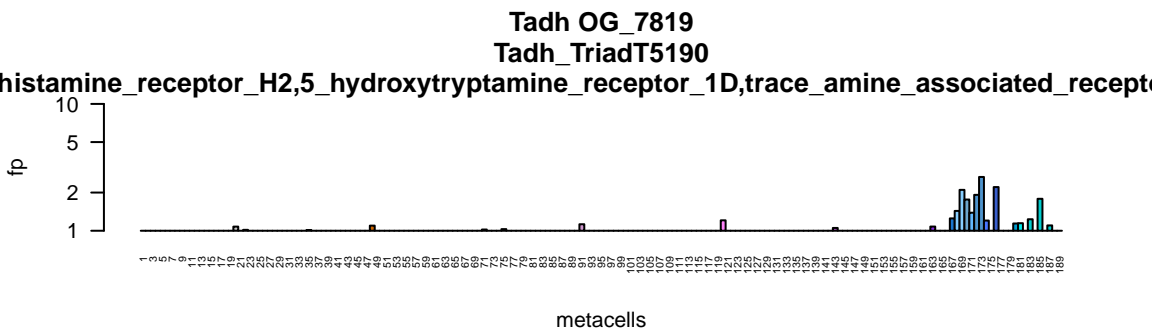
HoiH23_PIH23_006966-RA

adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3



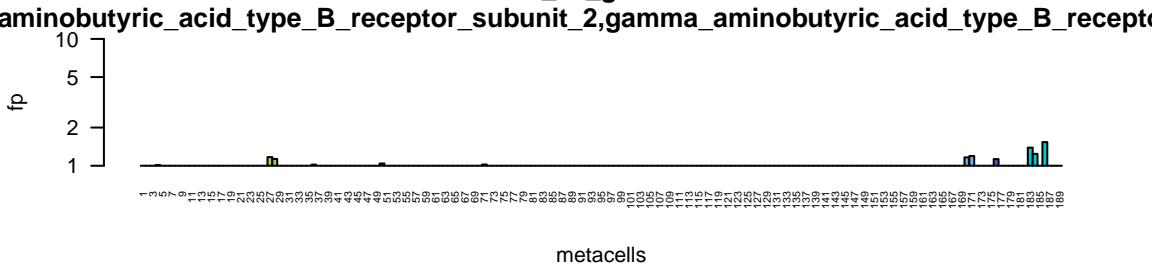






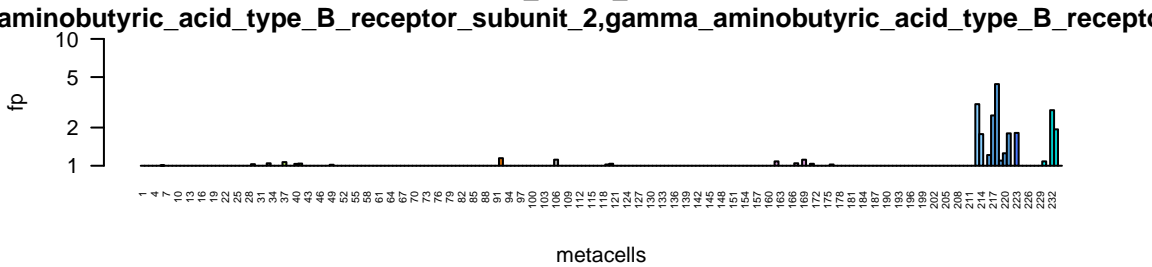
Tadh OG 8960

Tadh_wf_g11896.t1



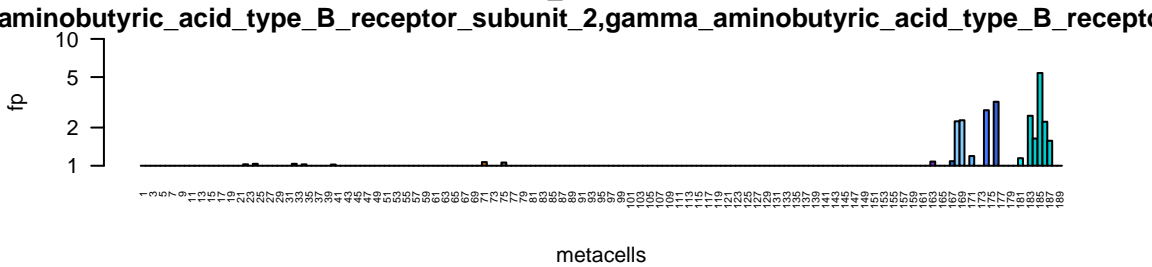
HoiH23 OG 8960

HoiH23_PIH23_000719-RA



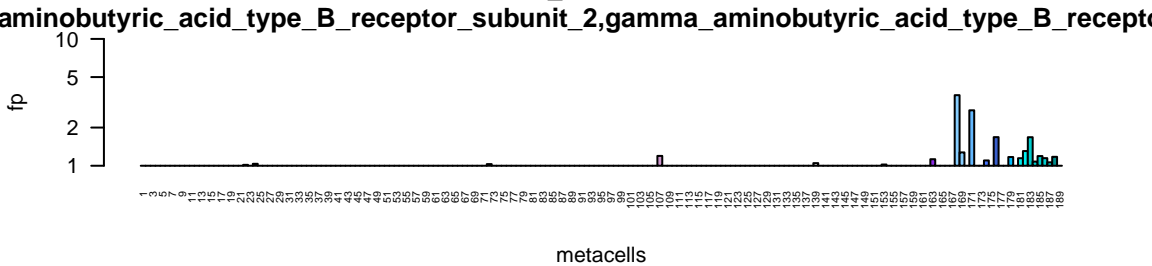
Tadh OG_8960

Tadh_TriadT52578



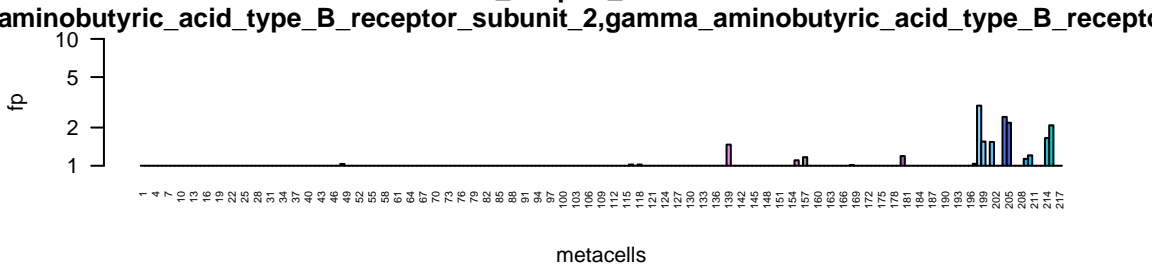
Tadh OG_8960

Tadh_TriadT52579



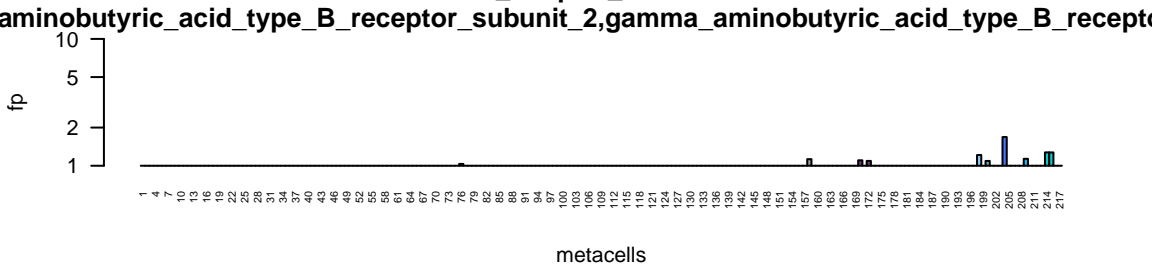
TrH2 OG_8960

TrH2_TrispH2_012100-RA



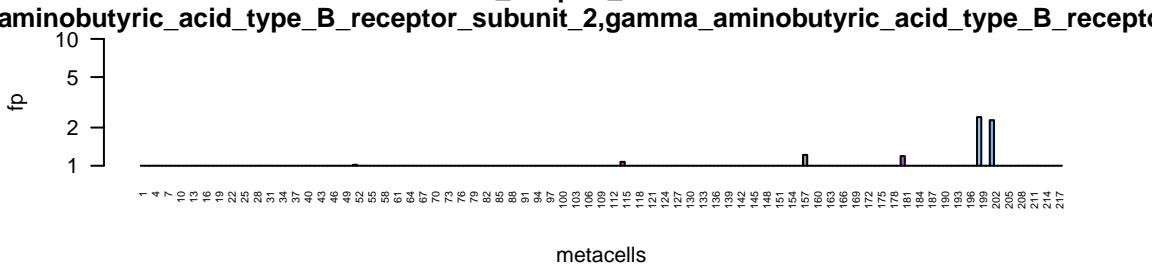
TrH2 OG_8960

TrH2_TrispH2_000234-RA



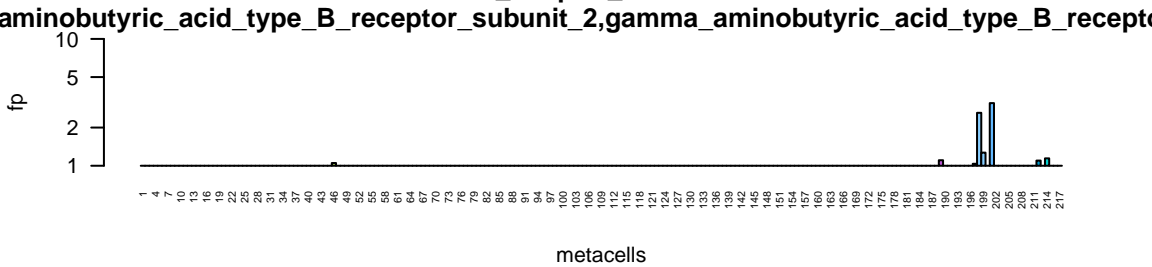
TrH2 OG_8960

TrH2_TrispH2_011964-RA

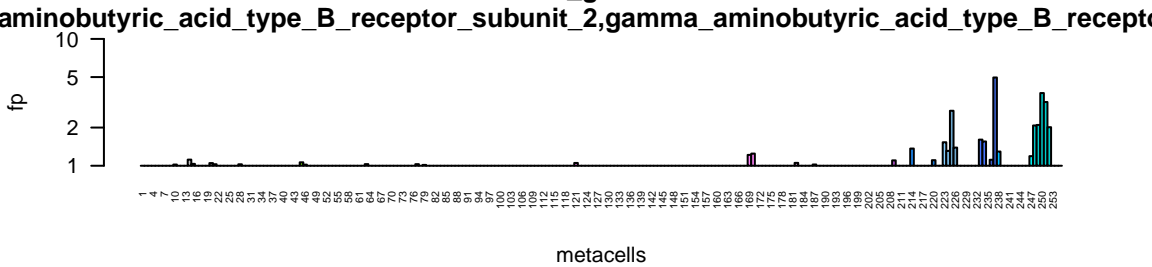


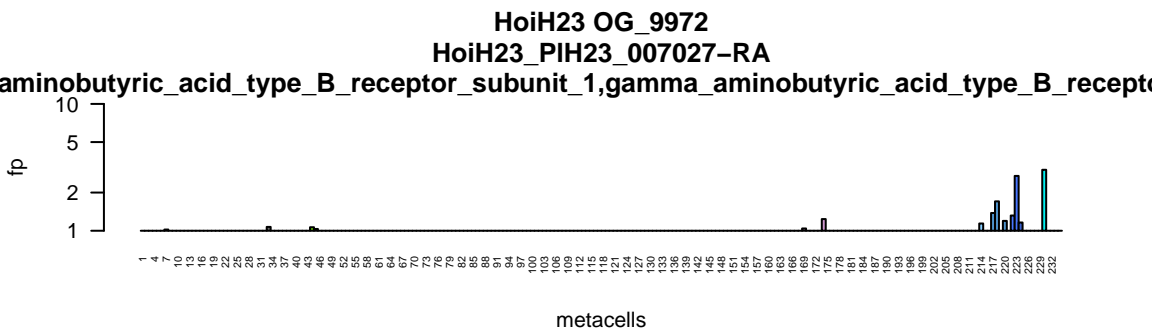
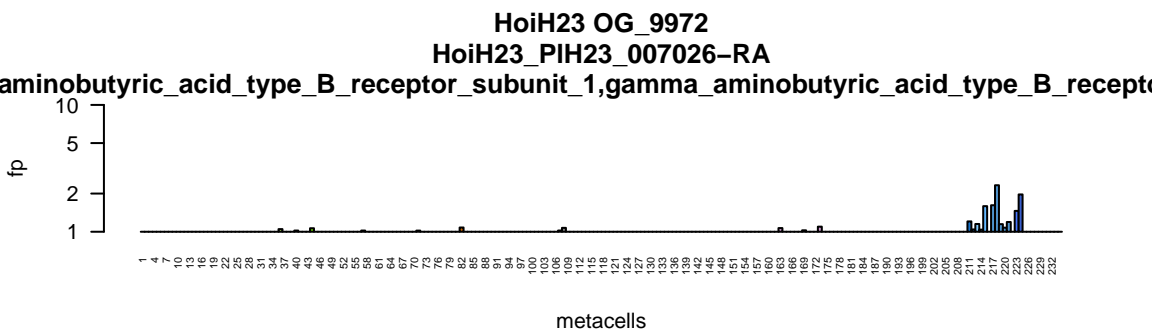
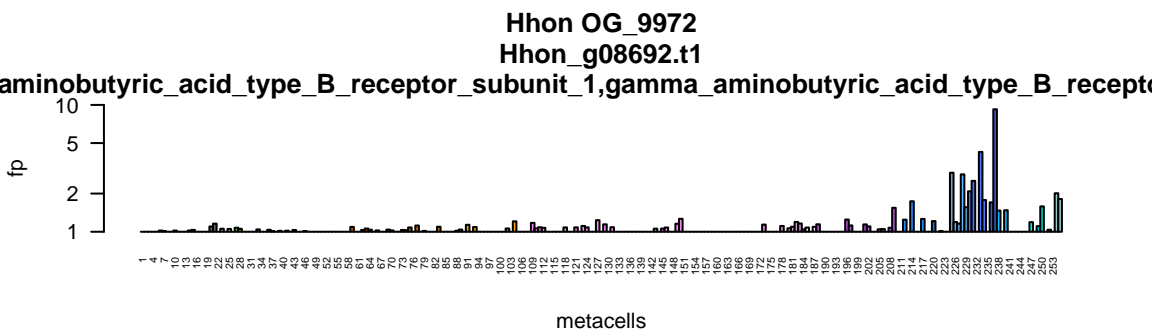
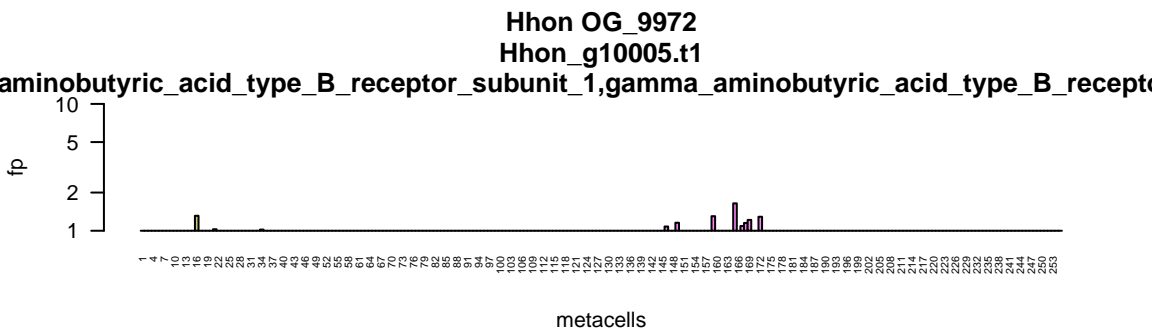
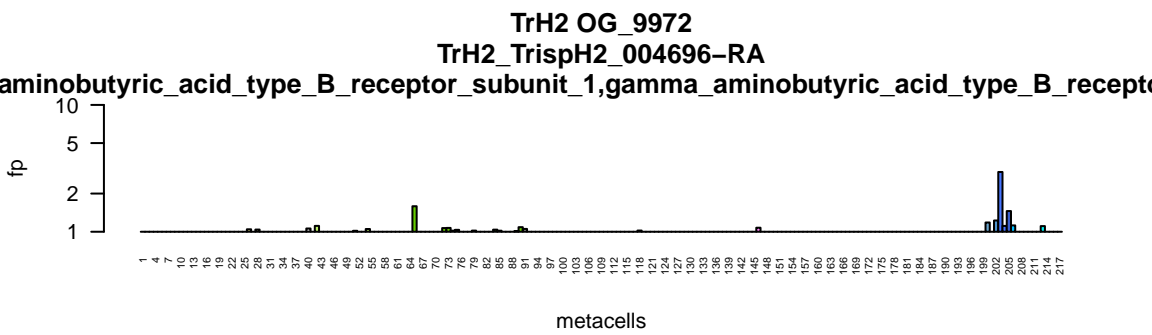
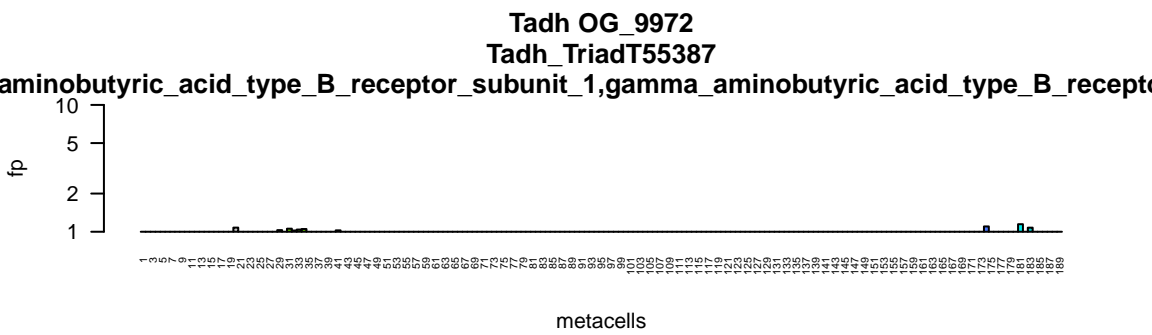
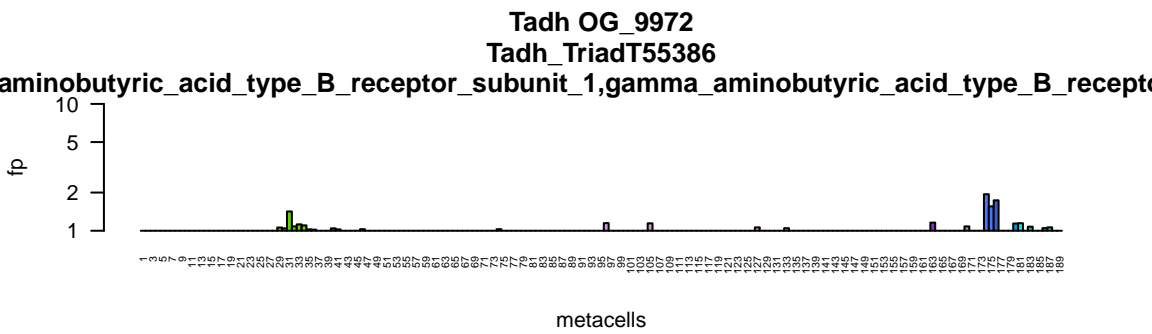
TrH2 OG_8960

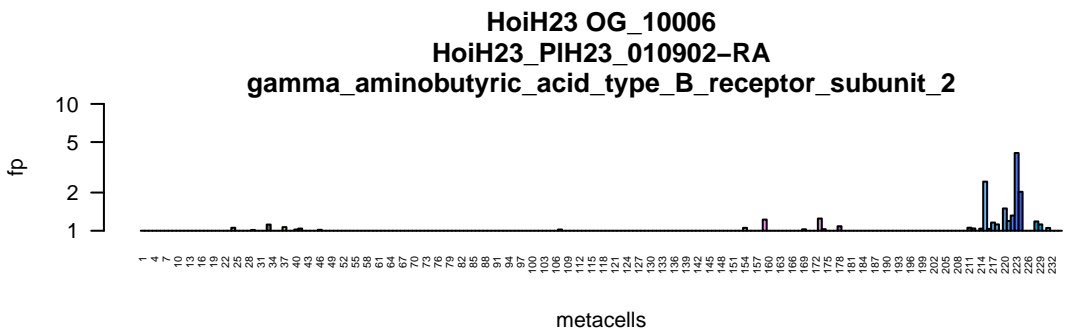
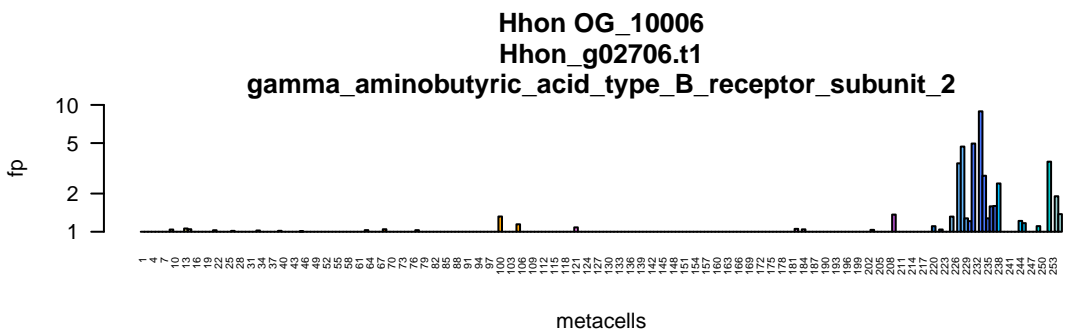
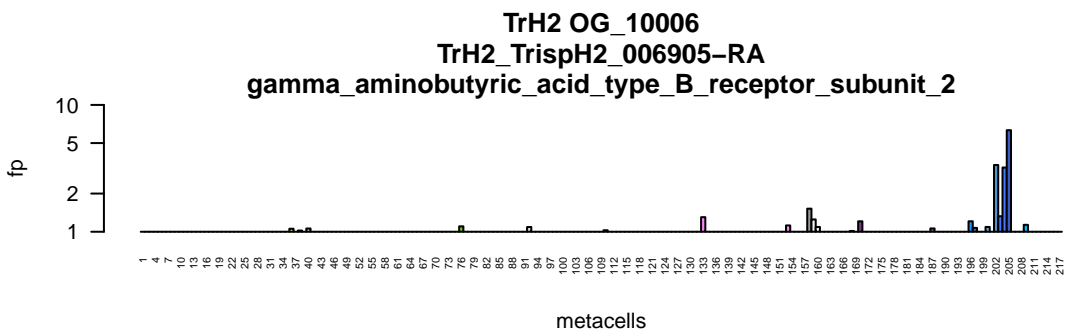
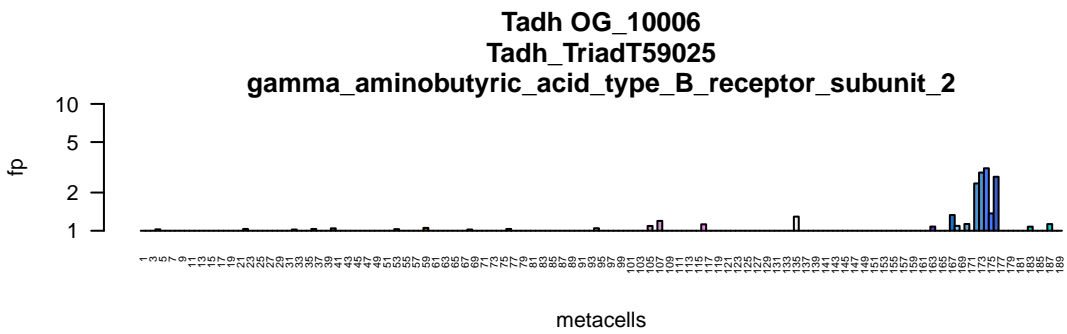
TrH2_TrispH2_011847-RA

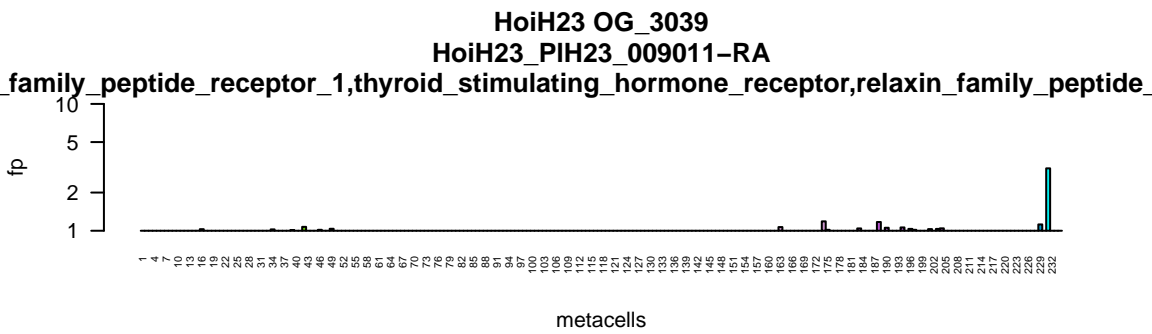
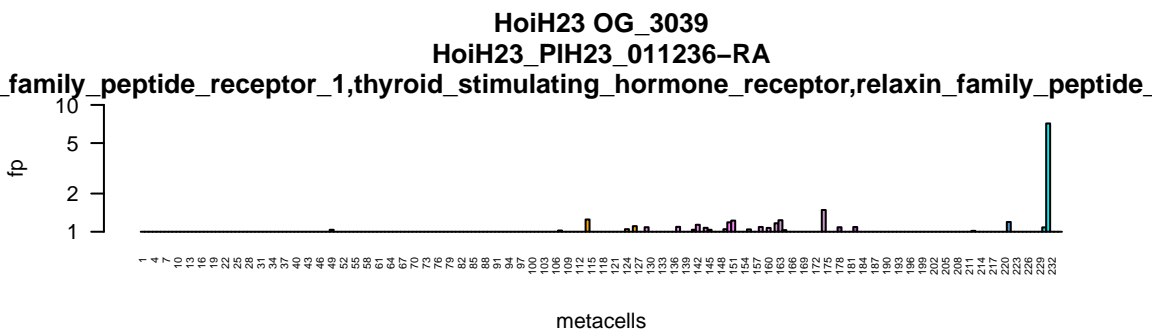
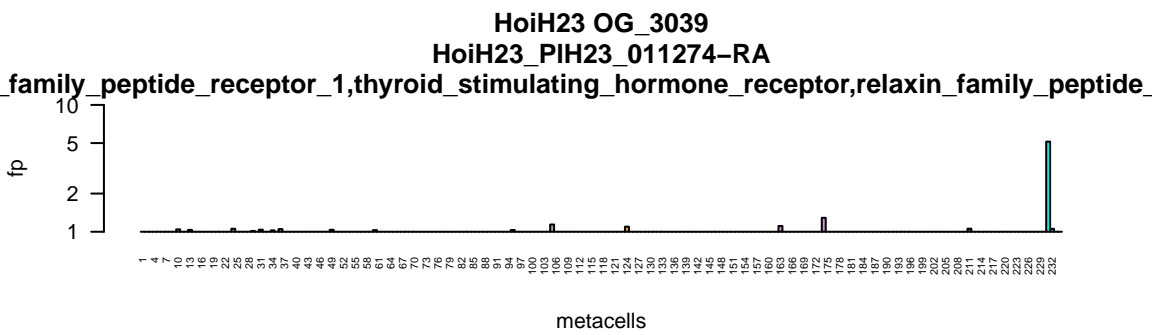
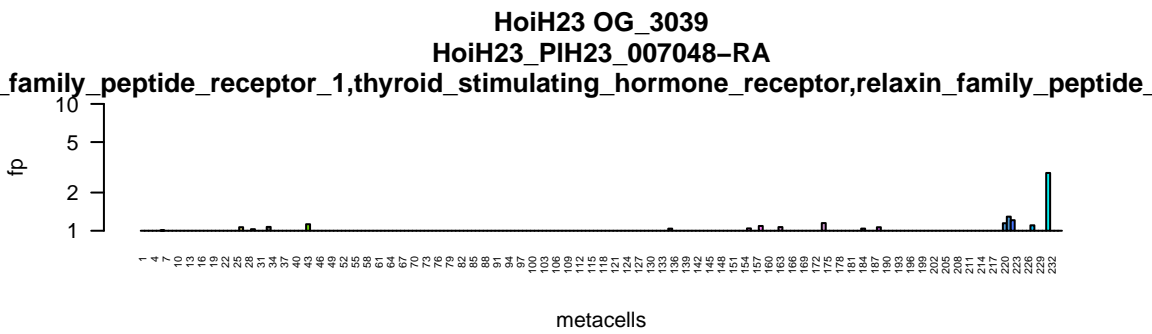
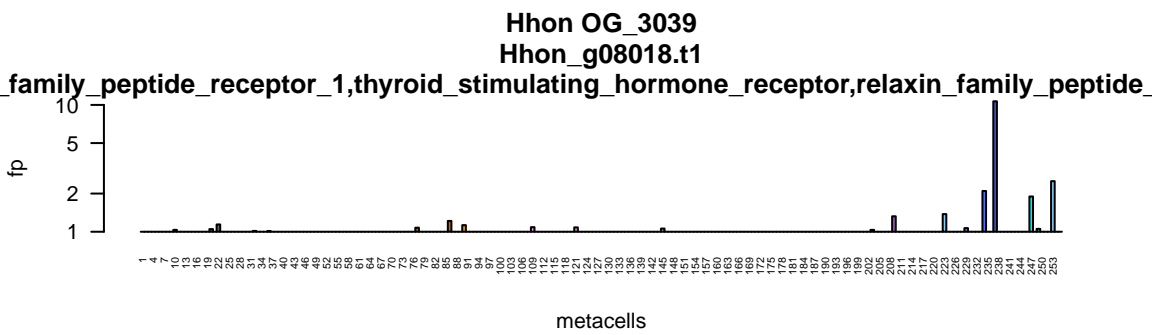
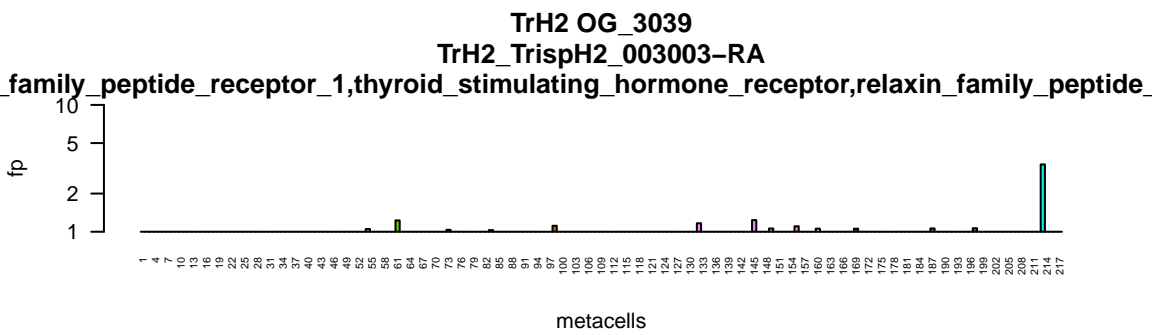
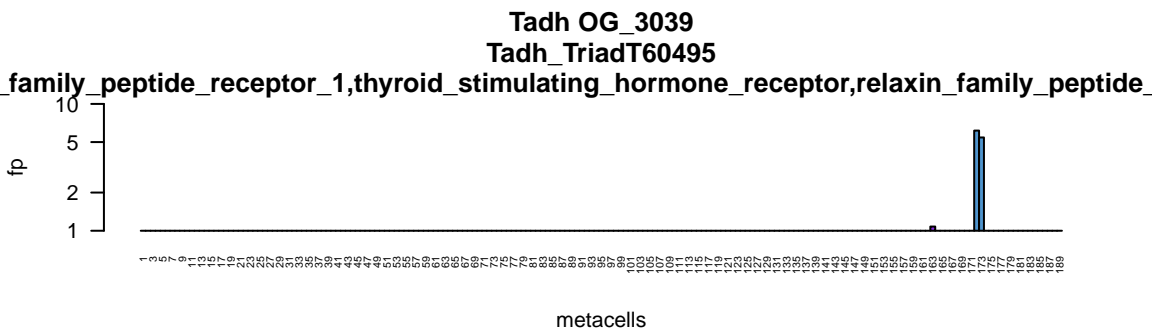
**Hhon OG_8960**

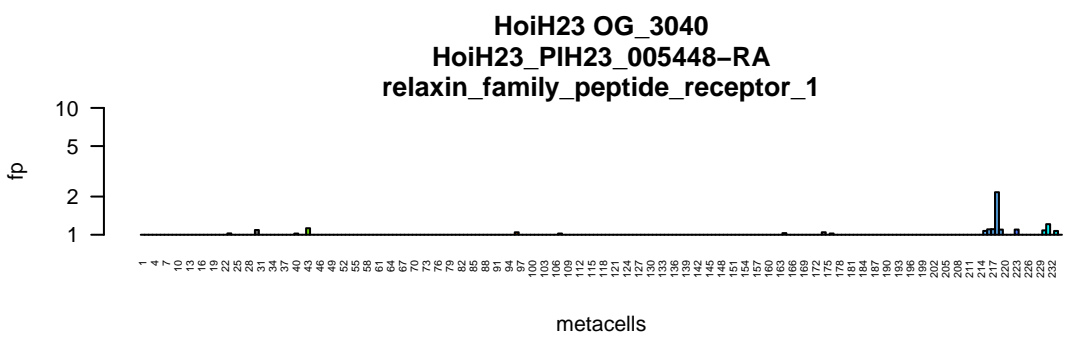
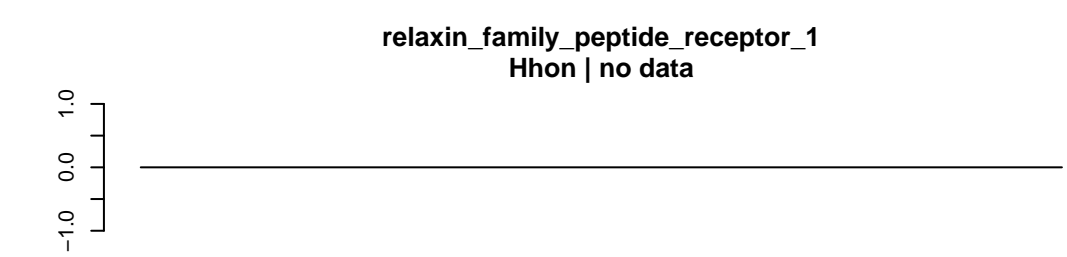
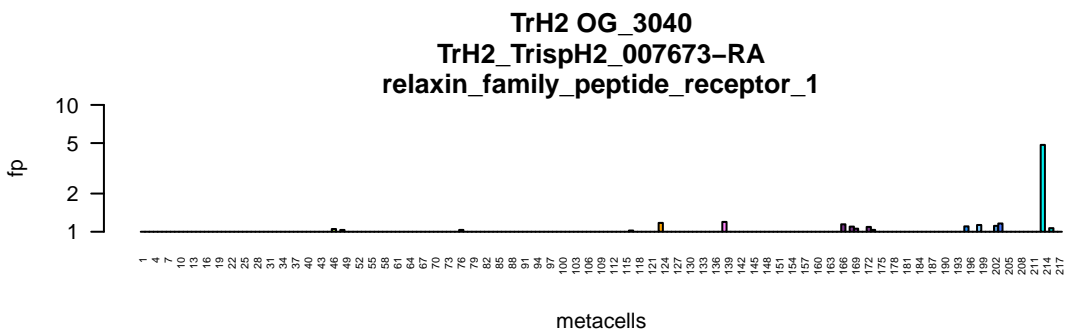
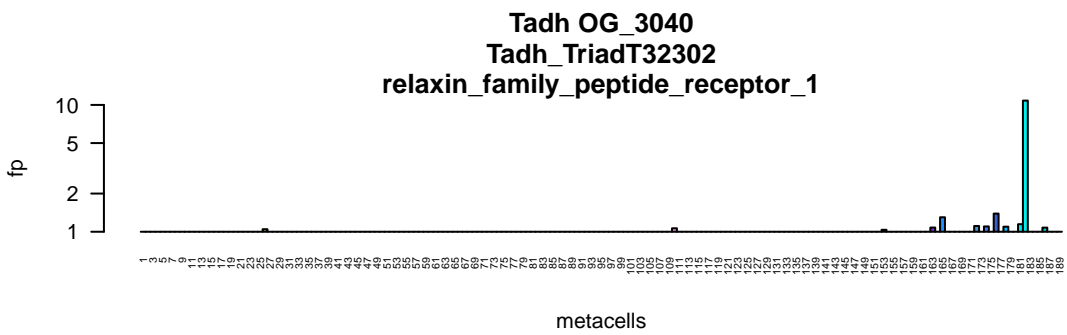
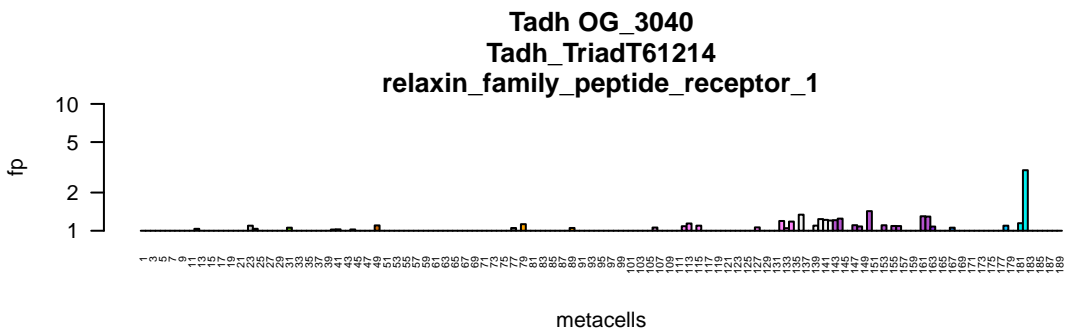
Hhon_g05296.t1

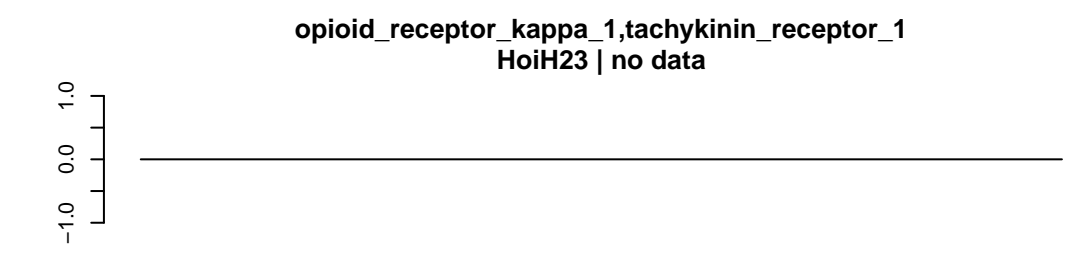
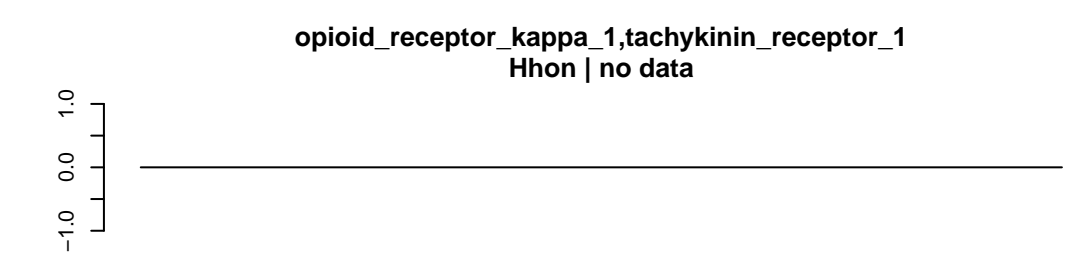
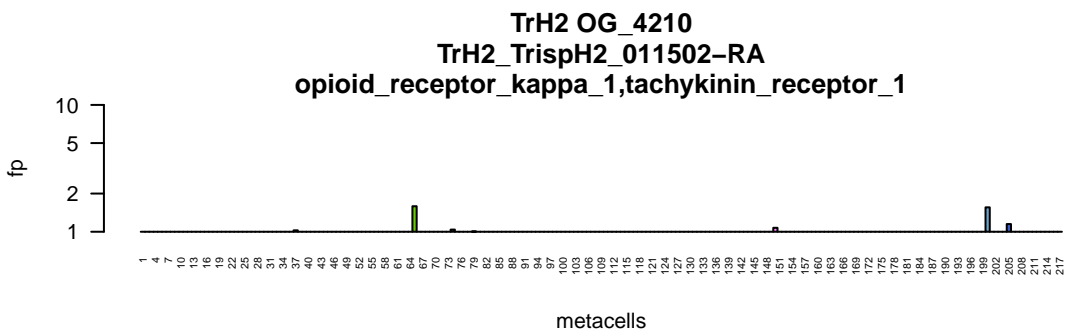
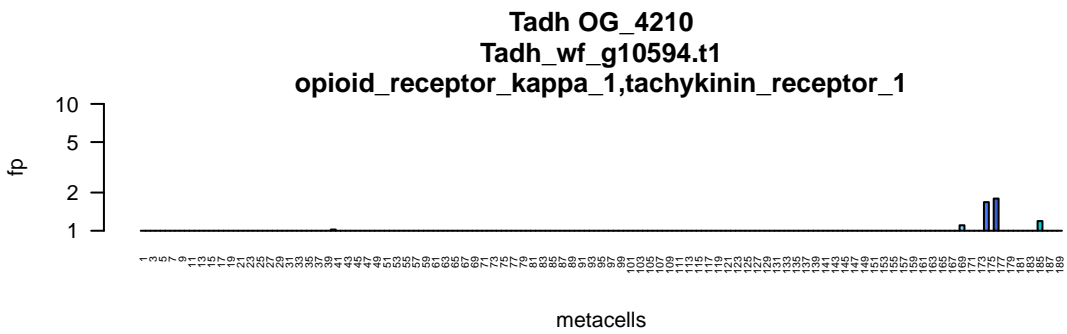


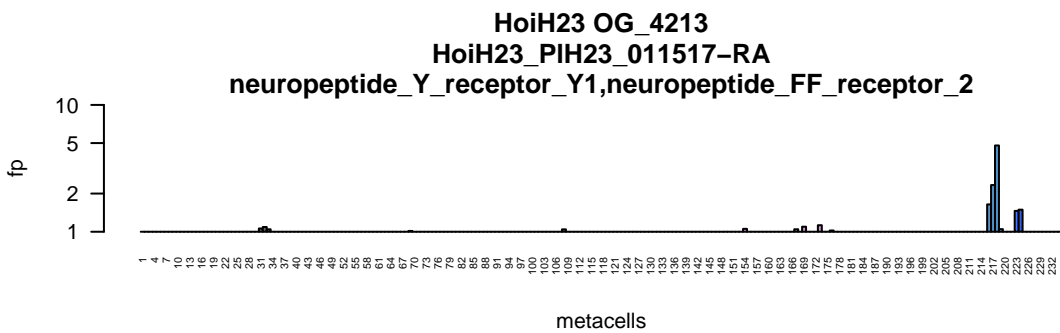
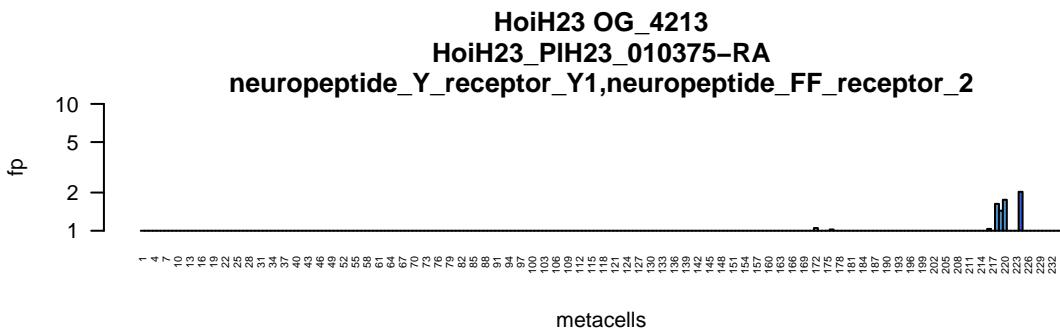
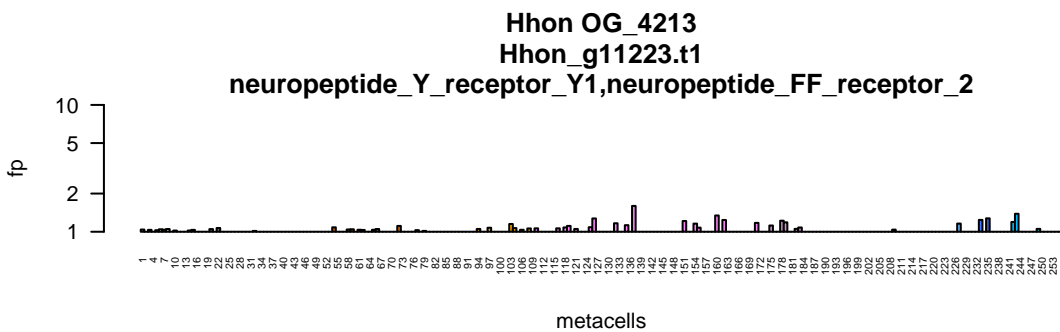
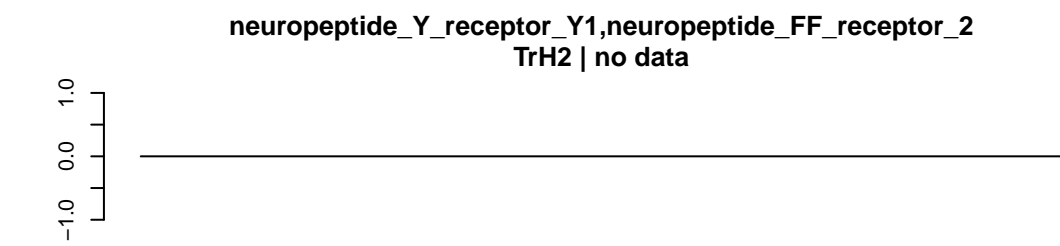
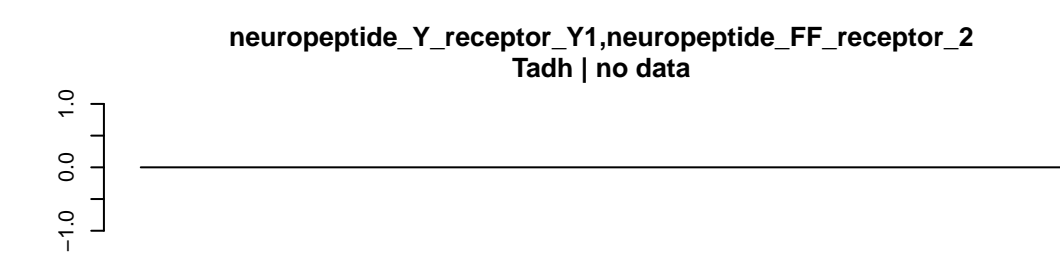


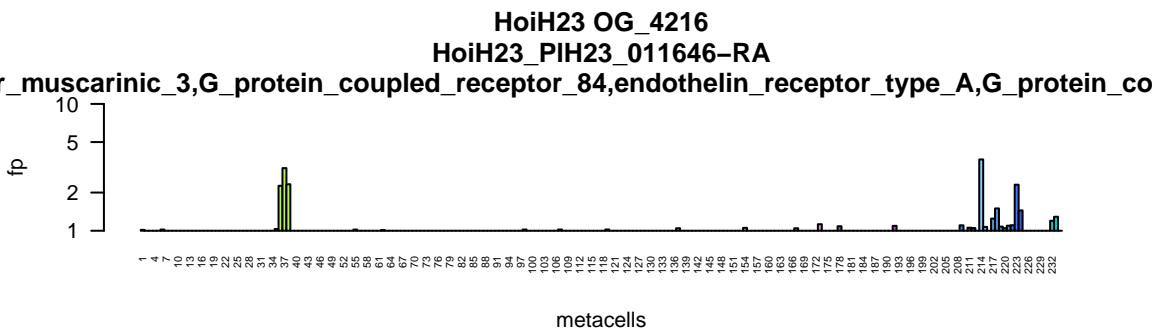
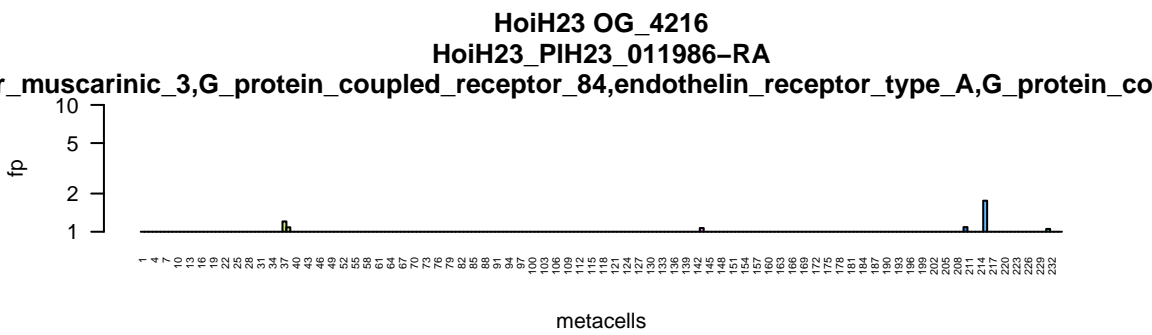
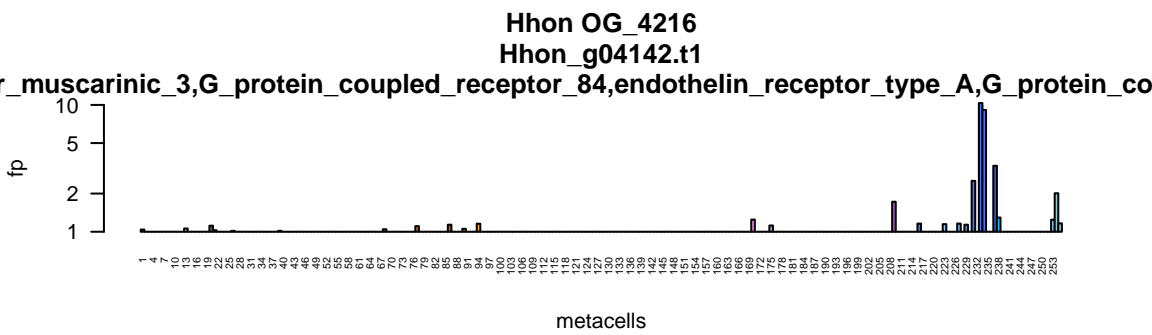
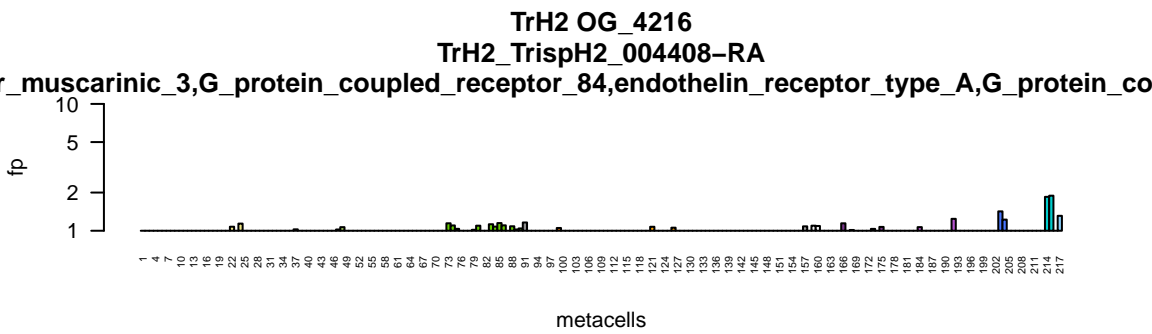
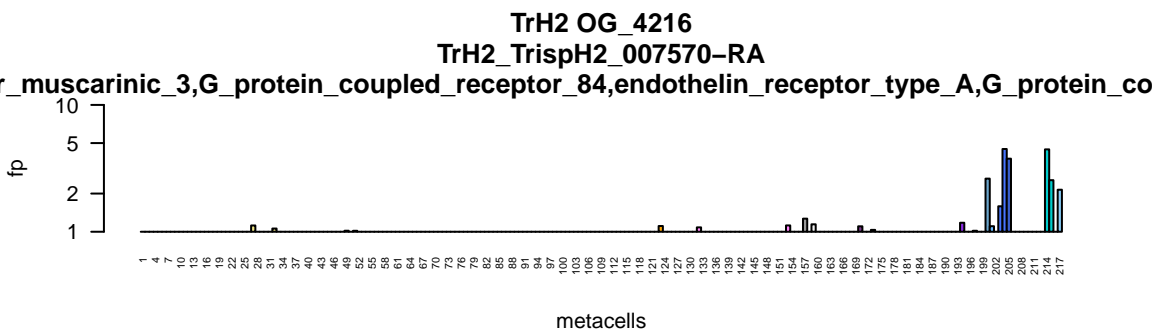
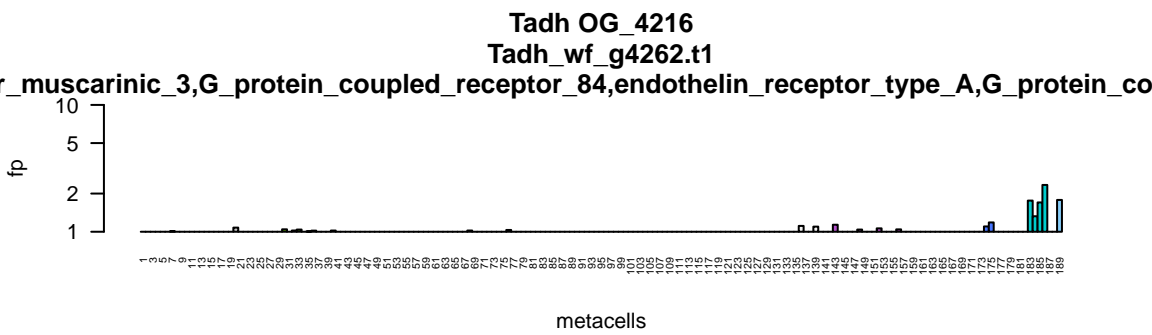
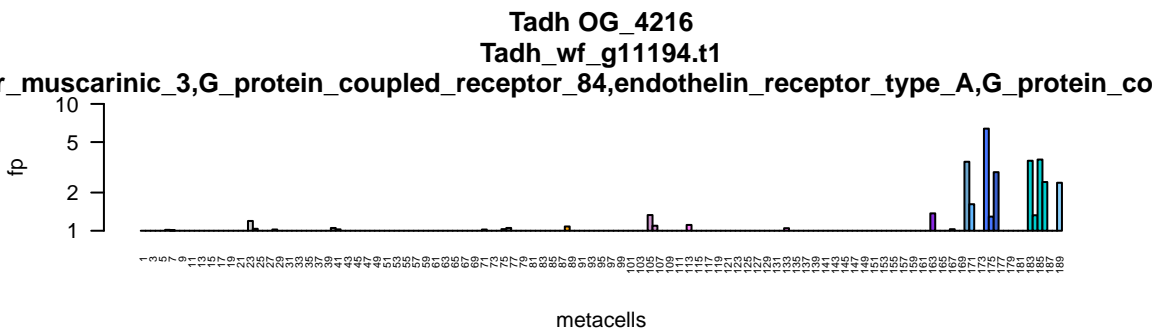


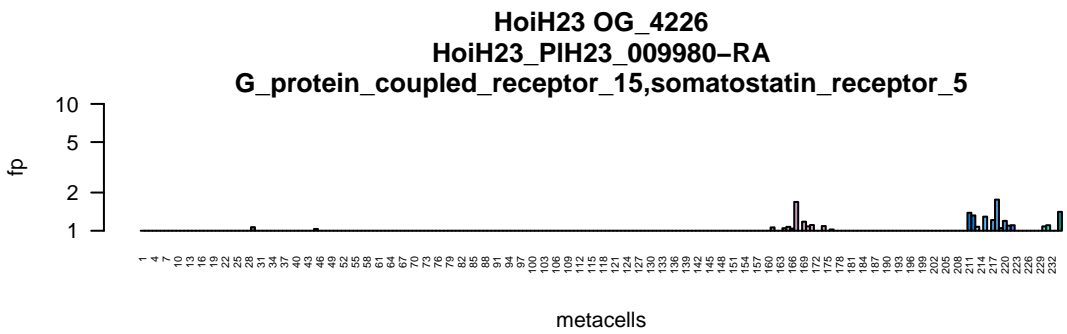
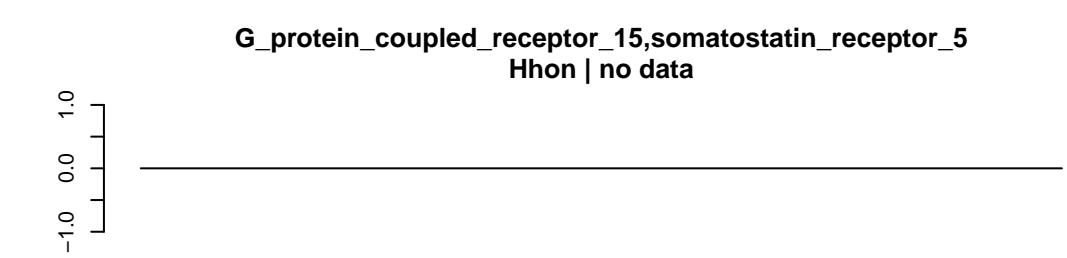
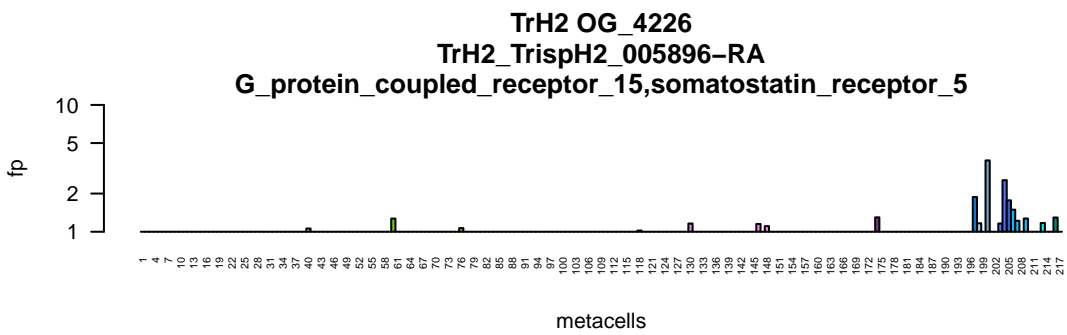
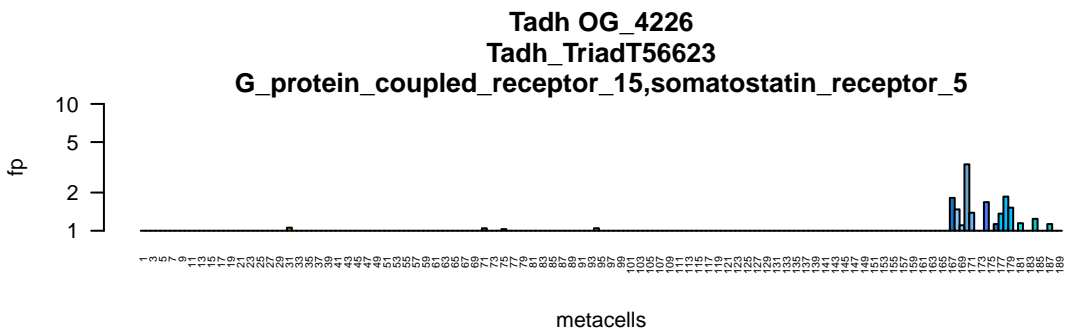












Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists indices from 1 to 217. Most metacells have 0 false positives, but some have 1 or 2. Metacell 202 has the highest number of false positives, with 4.

metacell	fp
1	0
4	0
13	1
16	1
19	1
25	1
28	0
31	0
34	0
37	0
40	1
43	1
46	0
49	1
52	1
55	1
58	1
61	0
64	0
67	1
70	0
73	0
76	0
79	0
82	0
85	0
88	0
91	1
94	1
97	0
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	0
130	1
133	0
136	0
139	0
142	1
145	0
148	1
151	1
154	0
157	0
160	0
163	1
166	0
169	0
172	0
175	0
178	0
181	0
184	0
187	0
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
7	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	1.2
58	1
61	1
67	1
70	1
73	1
76	1
79	1
85	1
86	1
89	1
94	1
97	1.2
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1.2
130	1
133	1
136	1
139	1
142	1
145	1.2
148	1
151	1.2
154	1
157	1.2
160	1.2
163	1
166	1
169	1
172	1
175	1.5
178	1
181	1
184	1
187	1.2
190	1
193	1
196	1
199	2
202	2
205	2
208	1
211	1
214	1.2
217	2

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

fp

metacells

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacells	fp
1	1
4	1
13	1
16	1
17	1
25	1
26	1
28	1
31	1
34	1
40	1
43	1
46	1
52	1
55	1
58	1
61	1
67	1
70	1
73	1
75	1
79	1
85	1
86	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
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	1
205	1
208	1
211	1
214	1
217	1

metacells	fp
1	1
4	1
7	1
13	1
16	1
19	1
25	1
28	1
31	1
34	1
40	1
43	1
46	1
52	1
55	1
59	1
61	1
67	1
70	1
73	1
76	1
79	1
85	1
86	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	2
202	2
205	1
208	1
211	1
214	1
217	2

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

fp

metacells

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacells	fp
1	1
4	1
10	1
13	1
19	1
22	1
28	1
31	1
37	1
43	1
46	1
52	1
55	1
61	1
64	1
70	1
73	1
79	1
82	1
88	1
91	1
97	1
100	1
106	1
109	1
115	1
119	1
127	1
124	1
133	1
136	1
142	1
146	1
151	1
154	1
160	1
163	1
169	1
175	1
178	1
184	1
187	1
193	1
196	1
202	1
205	1
211	1
214	1
220	1
223	1
229	1
232	1
238	1
241	1
247	1
250	1
253	1

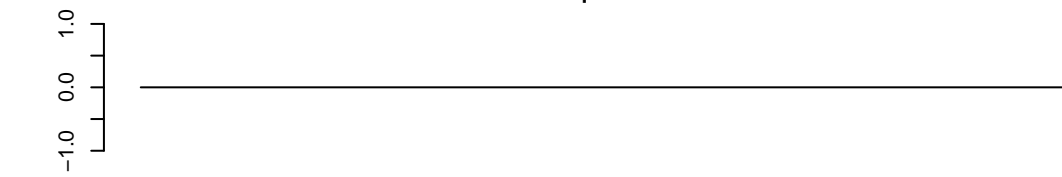
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacells	fp
1	1
4	1
10	1
13	1
19	1
22	1
28	1
31	1
37	1
40	1
46	1
52	1
55	1
64	1
68	1
70	1
73	1
79	1
88	1
91	1
97	1
100	1
106	1
109	1
115	1
118	1
127	1
124	1
133	1
142	1
145	1
151	1
154	1
163	1
169	1
172	1
178	1
184	1
187	1
193	1
196	1
202	1
205	1
211	1
214	1
220	1
223	1
229	1
232	1
238	1
241	1
247	1
250	1
253	2

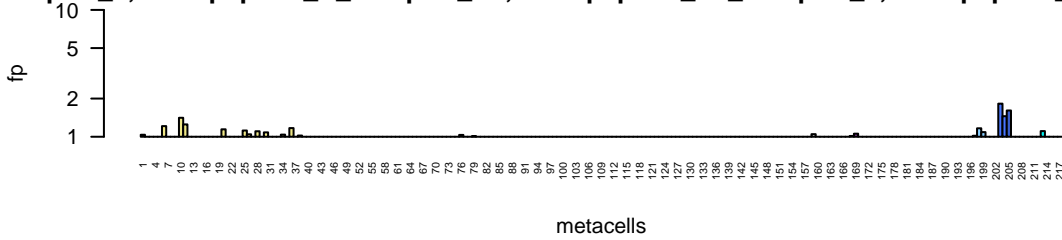
Bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and lists indices from 1 to 232. The y-axis is labeled 'fp' and has a logarithmic scale with values 1, 2, 5, and 10. Most metacells have a frequency of 1. Metacells 214, 223, and 232 show higher frequencies, with 214 reaching approximately 4.

A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 232. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a few outliers reaching up to 4. The bars are colored in a light blue/grey shade.

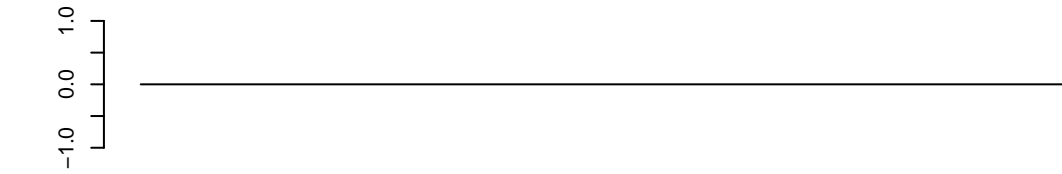
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec
Tadh | no data



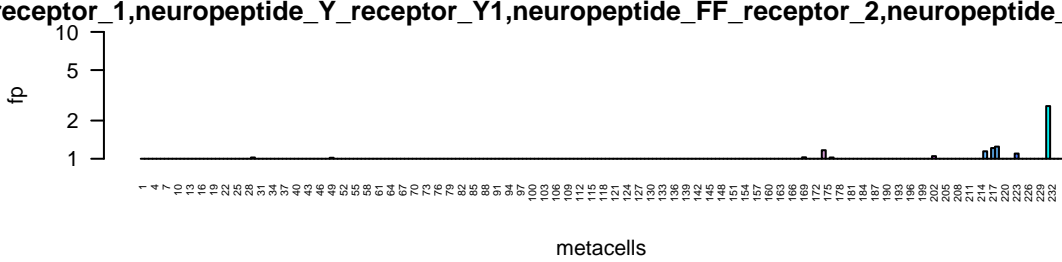
TrH2 OG_4548
TrH2_TrispH2_007473-RA
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec



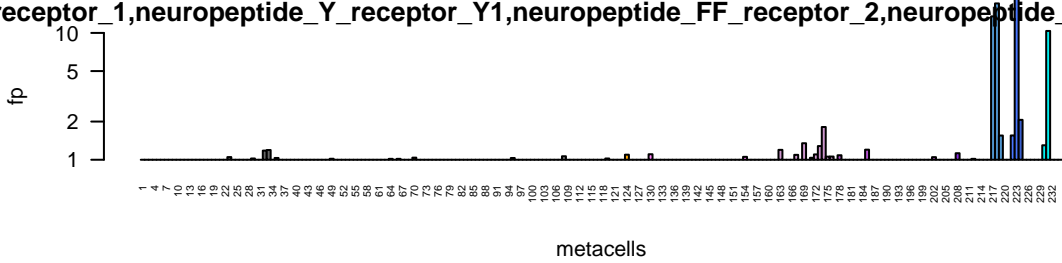
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec
Hhon | no data



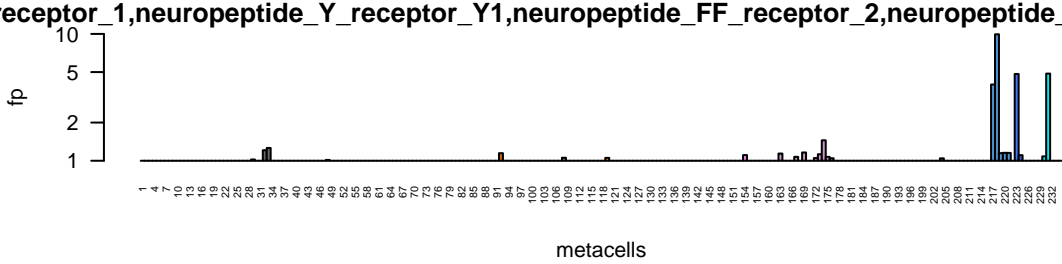
HoiH23 OG_4548
HoiH23_PIH23_006484-RA
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec



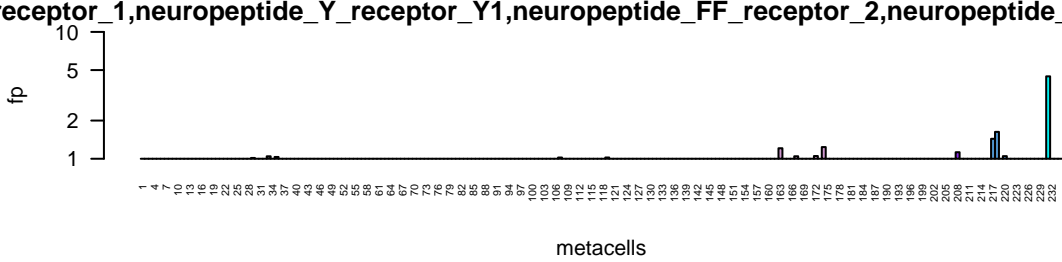
HoiH23 OG_4548
HoiH23_PIH23_011478-RA
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec



HoiH23 OG_4548
HoiH23_PIH23_011479-RA
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec



HoiH23 OG_4548
HoiH23_PIH23_011481-RA
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec



TrH2 OG_4762
TrH2_TrispH2_010632-RA
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

fp

metacells

Tadh OG_4762
Tadh_wf_g8464.t1
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chen

metacells	Tadh OG_4762	Tadh_wf_g8464.t1	hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chen
1	1.0	1.0	1.0
2	1.0	1.0	1.0
3	1.0	1.0	1.0
4	1.0	1.0	1.0
5	1.0	1.0	1.0
6	1.0	1.0	1.0
7	1.0	1.0	1.0
8	1.0	1.0	1.0
9	1.0	1.0	1.0
10	1.0	1.0	1.0
11	1.0	1.0	1.0
12	1.0	1.0	1.0
13	1.0	1.0	1.0
14	1.0	1.0	1.0
15	1.0	1.0	1.0
16	1.0	1.0	1.0
17	1.0	1.0	1.0
18	1.0	1.0	1.0
19	1.0	1.0	1.0
20	1.0	1.0	1.0
21	1.0	1.0	1.0
22	1.0	1.0	1.0
23	1.0	1.0	1.0
24	1.0	1.0	1.0
25	1.0	1.0	1.0
26	1.0	1.0	1.0
27	1.0	1.0	1.0
28	1.0	1.0	1.0
29	1.0	1.0	1.0
30	1.0	1.0	1.0
31	1.0	1.0	1.0
32	1.0	1.0	1.0
33	1.0	1.0	1.0
34	1.0	1.0	1.0
35	1.0	1.0	1.0
36	1.0	1.0	1.0
37	1.0	1.0	1.0
38	1.0	1.0	1.0
39	1.0	1.0	1.0
40	1.0	1.0	1.0
41	1.0	1.0	1.0
42	1.0	1.0	1.0
43	1.0	1.0	1.0
44	1.0	1.0	1.0
45	1.0	1.0	1.0
46	1.0	1.0	1.0
47	1.0	1.0	1.0
48	1.0	1.0	1.0
49	1.0	1.0	1.0
50	1.0	1.0	1.0
51	1.0	1.0	1.0
52	1.0	1.0	1.0
53	1.0	1.0	1.0
54	1.0	1.0	1.0
55	1.0	1.0	1.0
56	1.0	1.0	1.0
57	1.0	1.0	1.0
58	1.0	1.0	1.0
59	1.0	1.0	1.0
60	1.0	1.0	1.0
61	1.0	1.0	1.0
62	1.0	1.0	1.0
63	1.0	1.0	1.0
64	1.0	1.0	1.0
65	1.0	1.0	1.0
66	1.0	1.0	1.0
67	1.0	1.0	1.0
68	1.0	1.0	1.0
69	1.0	1.0	1.0
70	1.0	1.0	1.0
71	1.0	1.0	1.0
72	1.0	1.0	1.0
73	1.0	1.0	1.0
74	1.0	1.0	1.0
75	1.0	1.0	1.0
76	1.0	1.0	1.0
77	1.0	1.0	1.0
78	1.0	1.0	1.0
79	1.0	1.0	1.0
80	1.0	1.0	1.0
81	1.0	1.0	1.0
82	1.0	1.0	1.0
83	1.0	1.0	1.0
84	1.0	1.0	1.0
85	1.0	1.0	1.0
86	1.0	1.0	1.0
87	1.0	1.0	1.0
88	1.0	1.0	1.0
89	1.0	1.0	1.0
90	1.0	1.0	1.0
91	1.0	1.0	1.0
92	1.0	1.0	1.0
93	1.0	1.0	1.0
94	1.0	1.0	1.0
95	1.0	1.0	1.0
96	1.0	1.0	1.0
97	1.0	1.0	1.0
98	1.0	1.0	1.0
99	1.0	1.0	1.0
100	1.0	1.0	1.0
101	1.0	1.0	1.0
102	1.0	1.0	1.0
103	1.0	1.0	1.0
104	1.0	1.0	1.0
105	1.0	1.0	1.0
106	1.0	1.0	1.0
107	1.0	1.0	1.0
1			

Tadh OG_4762
Tadh_TriadT59534
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chen

fp

metacells

TrH2 OG_4762
TrH2_TrispH2_010634-RA
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chen

metacells	fp
1	1
4	1
10	1
13	1
16	1
19	1
22	1
25	1
28	1
31	1
37	1
40	1
43	1
46	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	1
205	1
208	1.2
211	1.1
214	1.8
217	2.2

TrH2 OG_4762
TrH2_TrispH2_010271-RA

hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

metacells

Hhon OG_4762
Hhon_g10761.t1

hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

fp

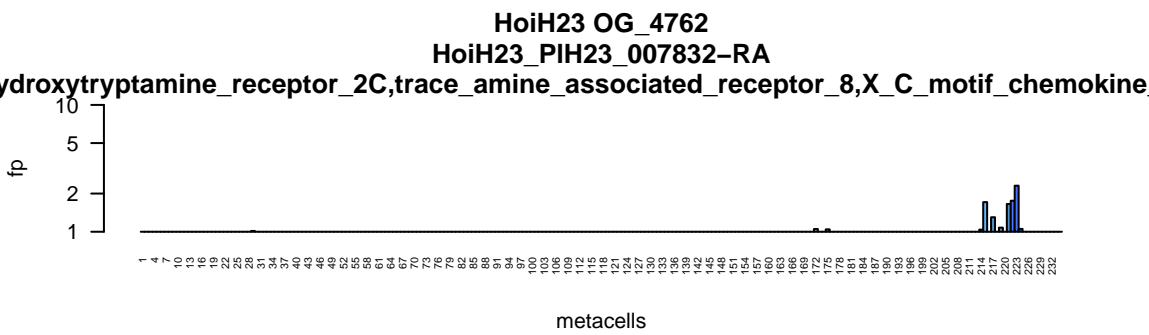
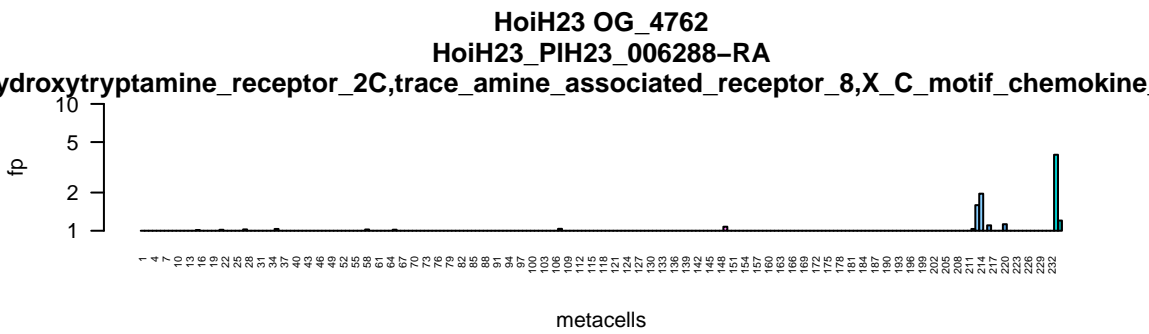
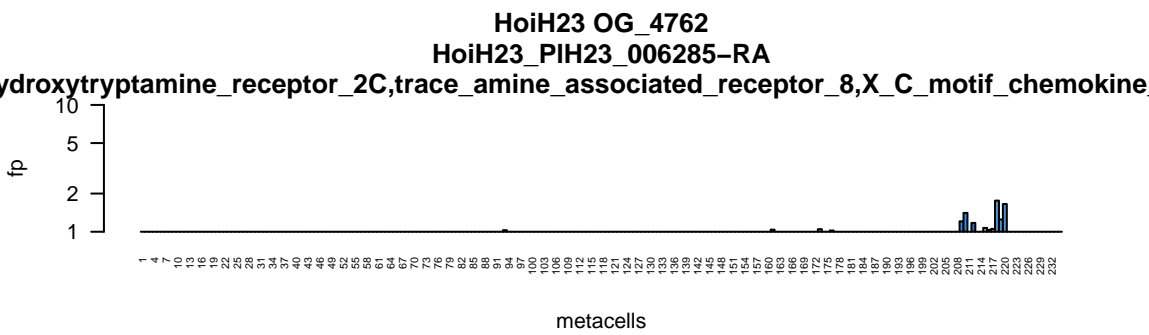
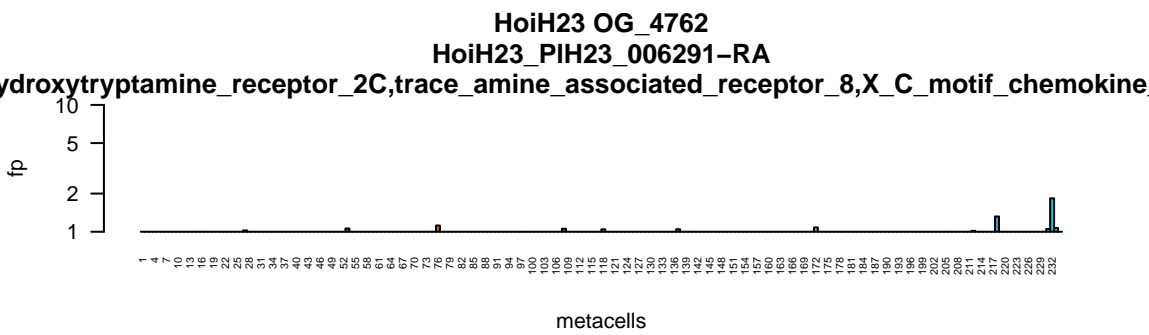
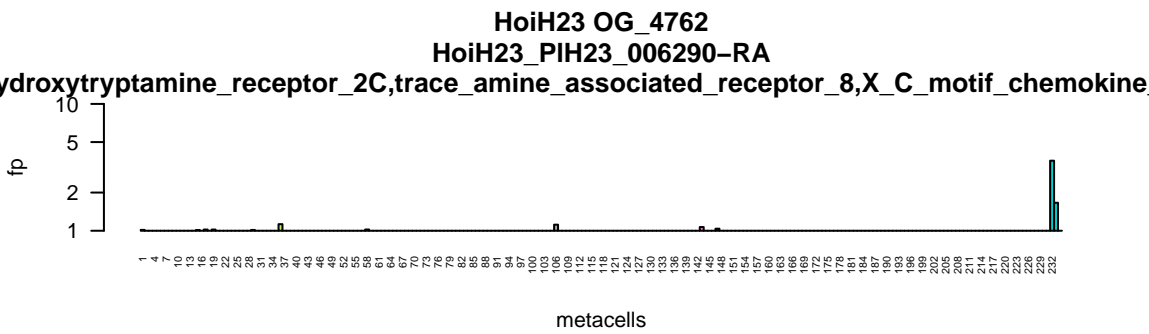
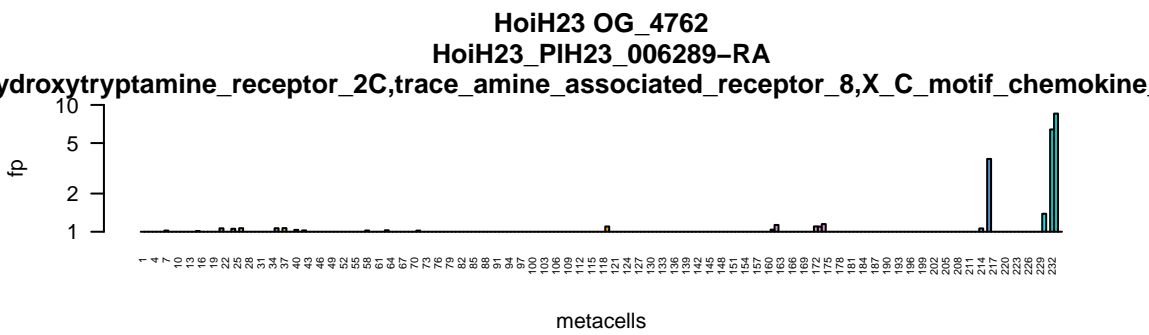
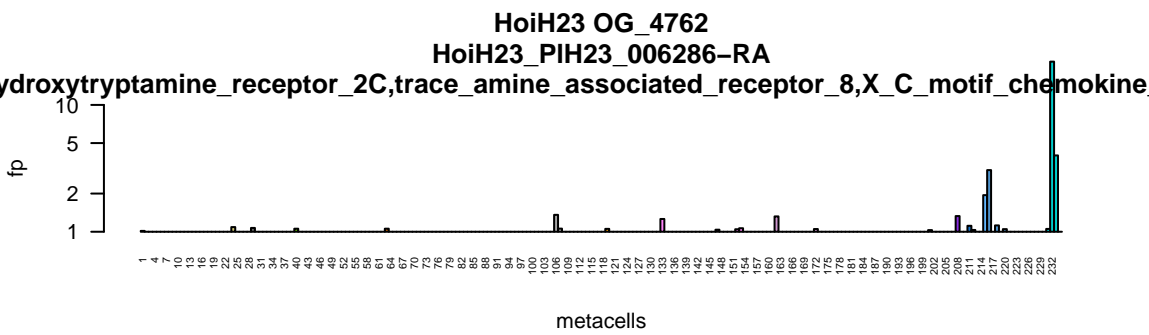
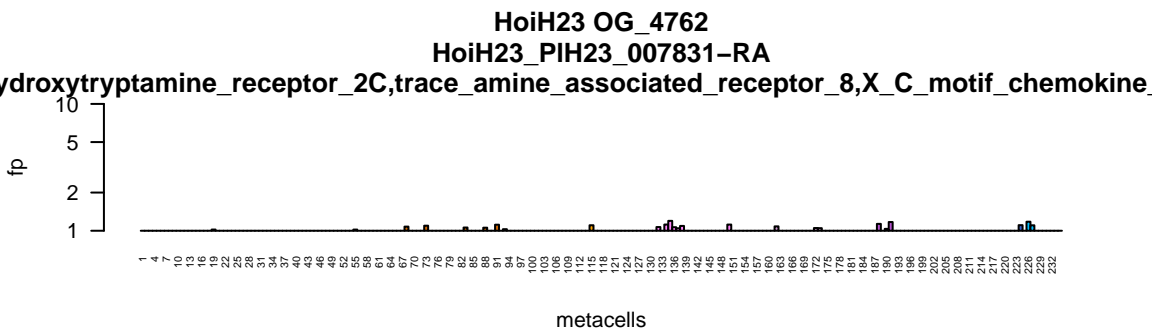
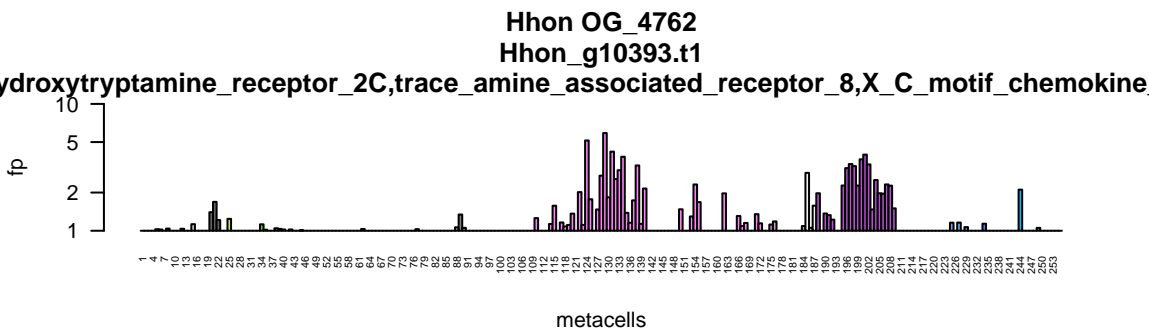
metacells

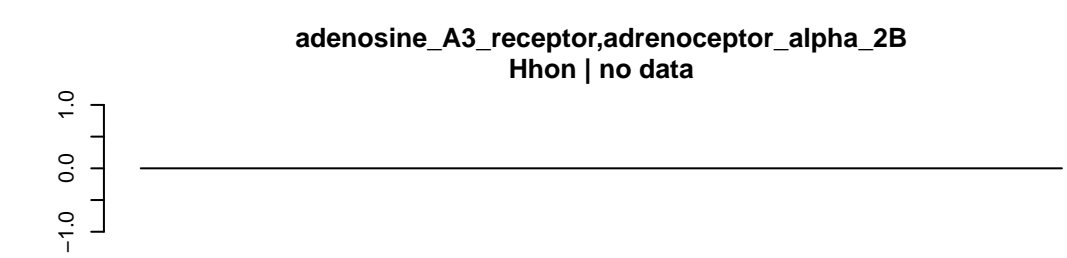
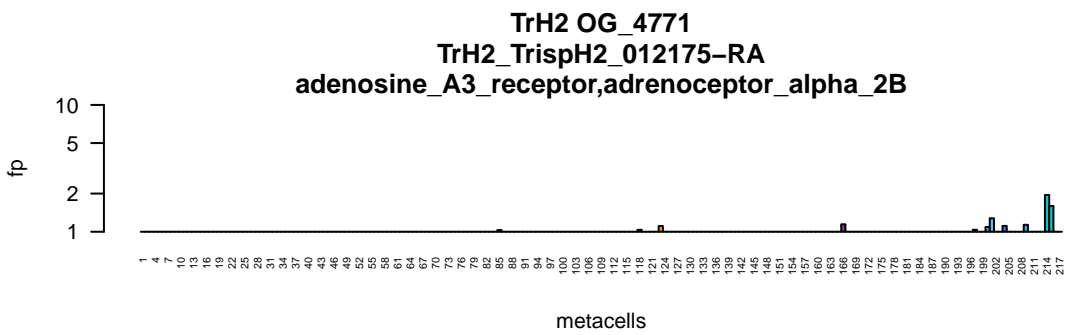
Hhon OG_4762
Hhon_g01703.t1

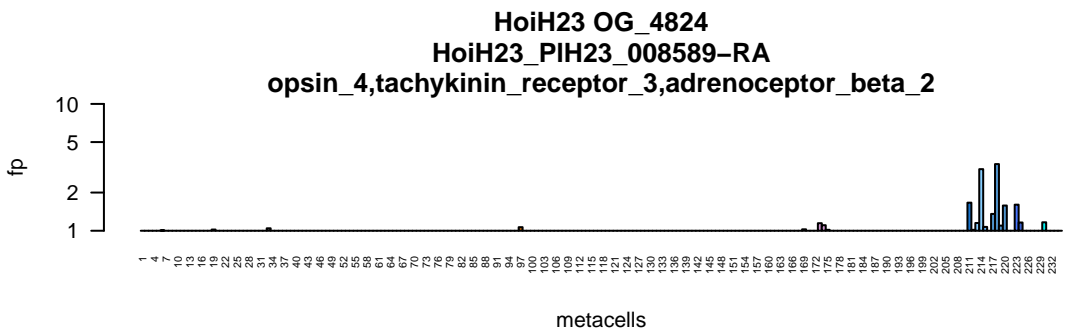
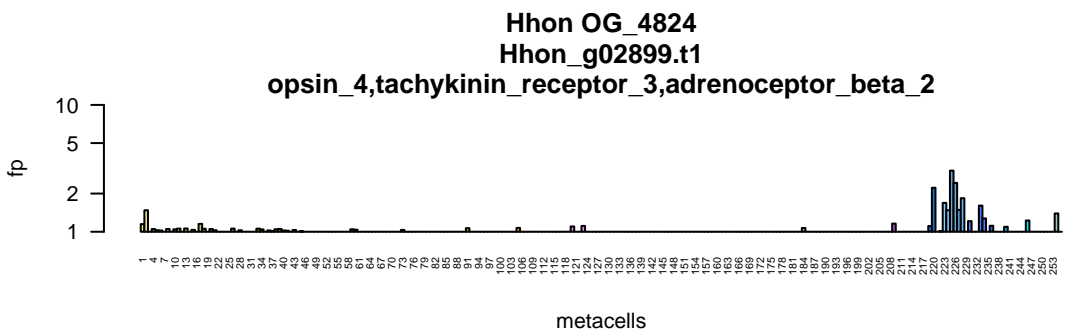
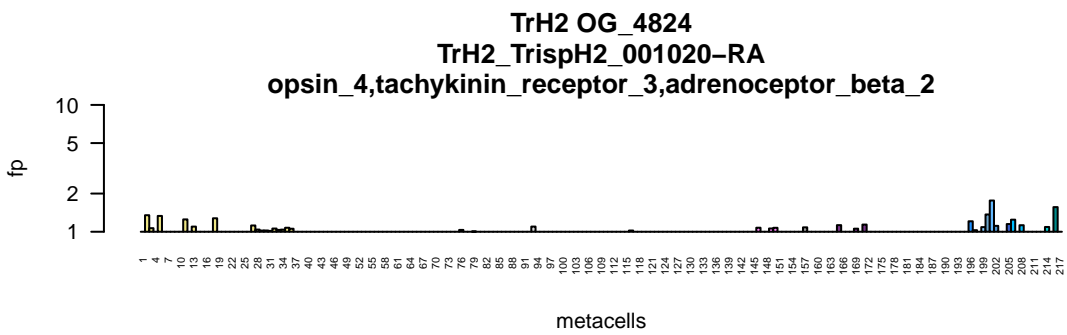
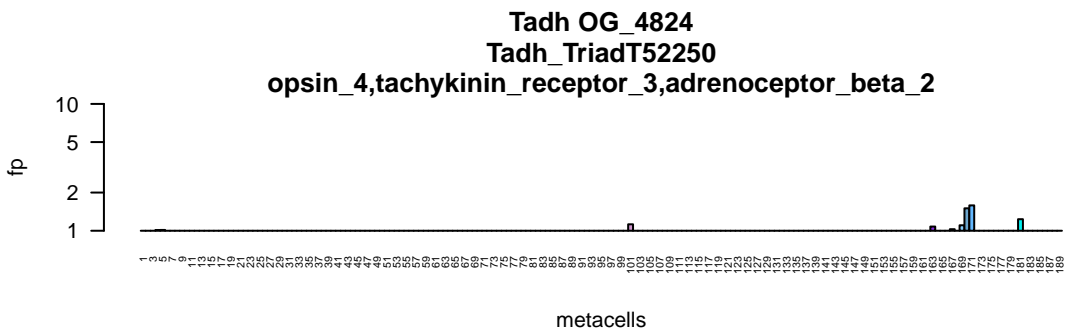
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

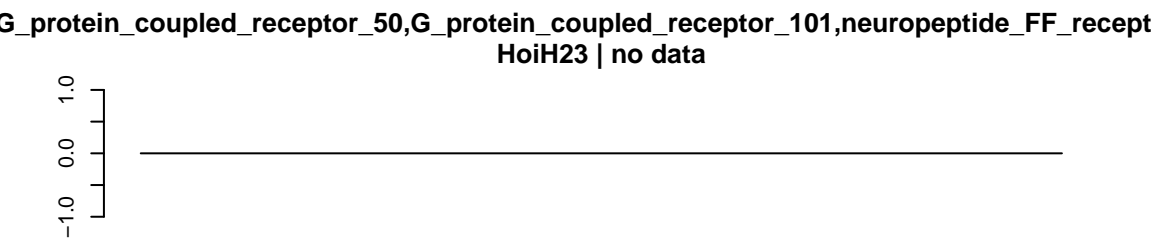
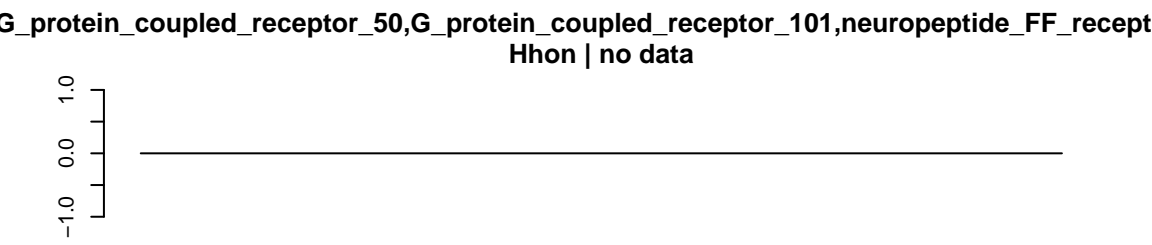
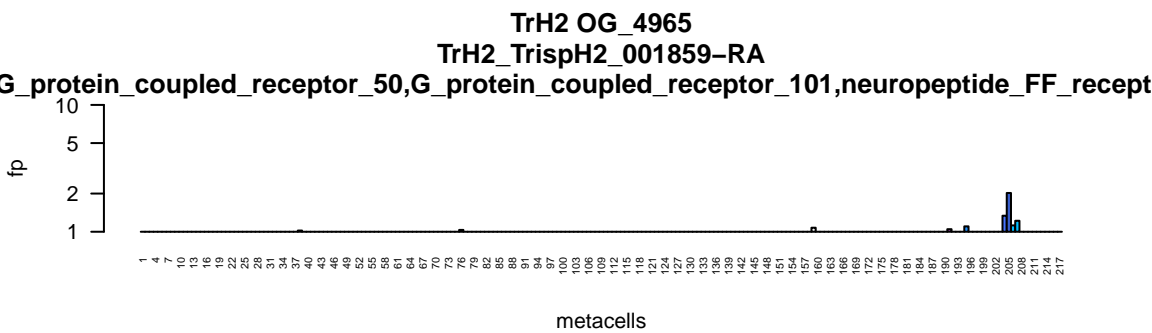
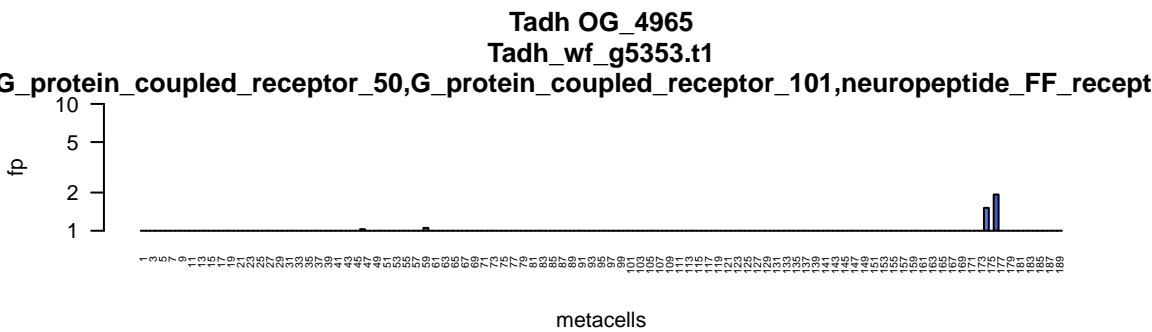
fp

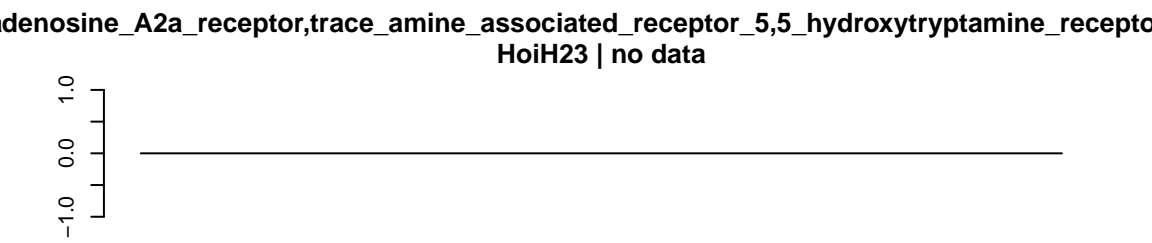
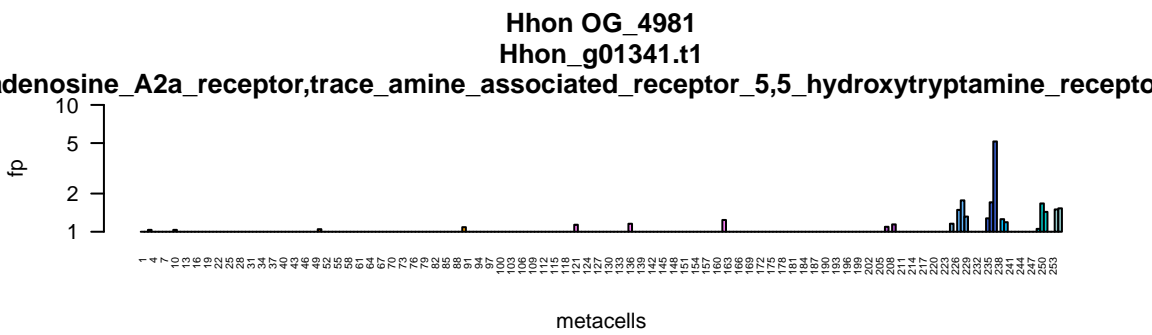
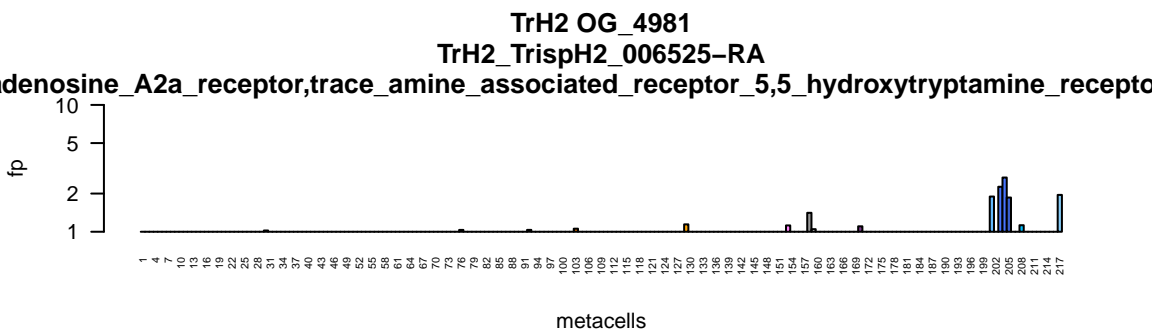
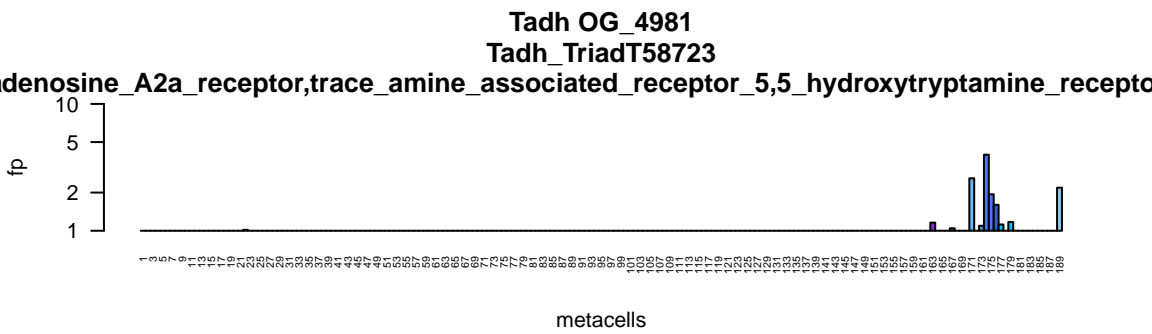
metacells

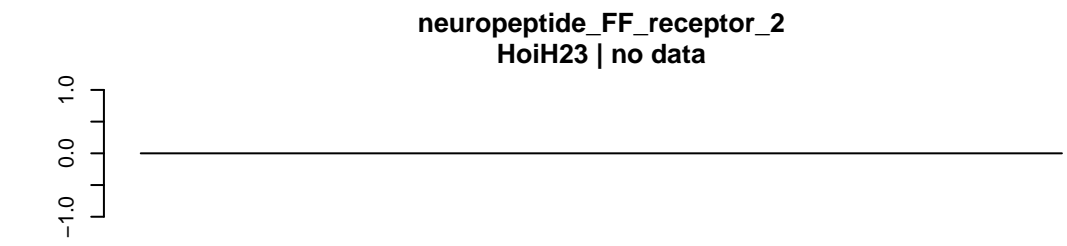
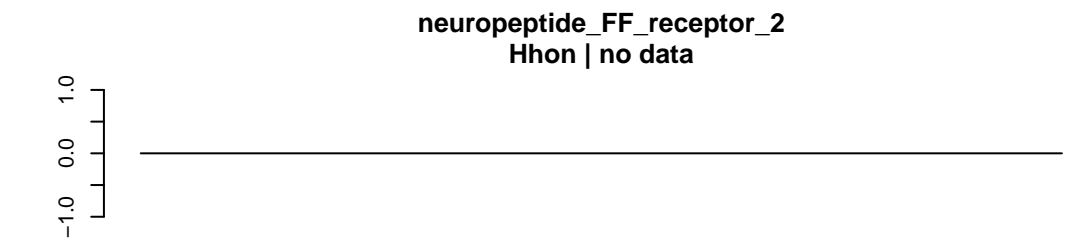
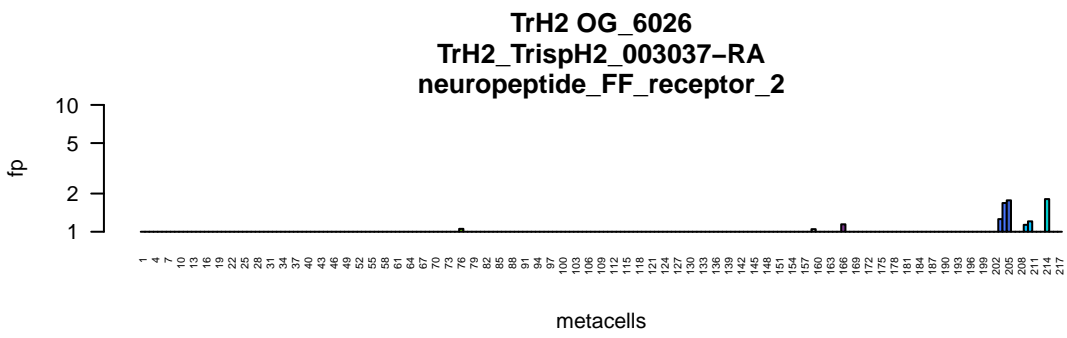
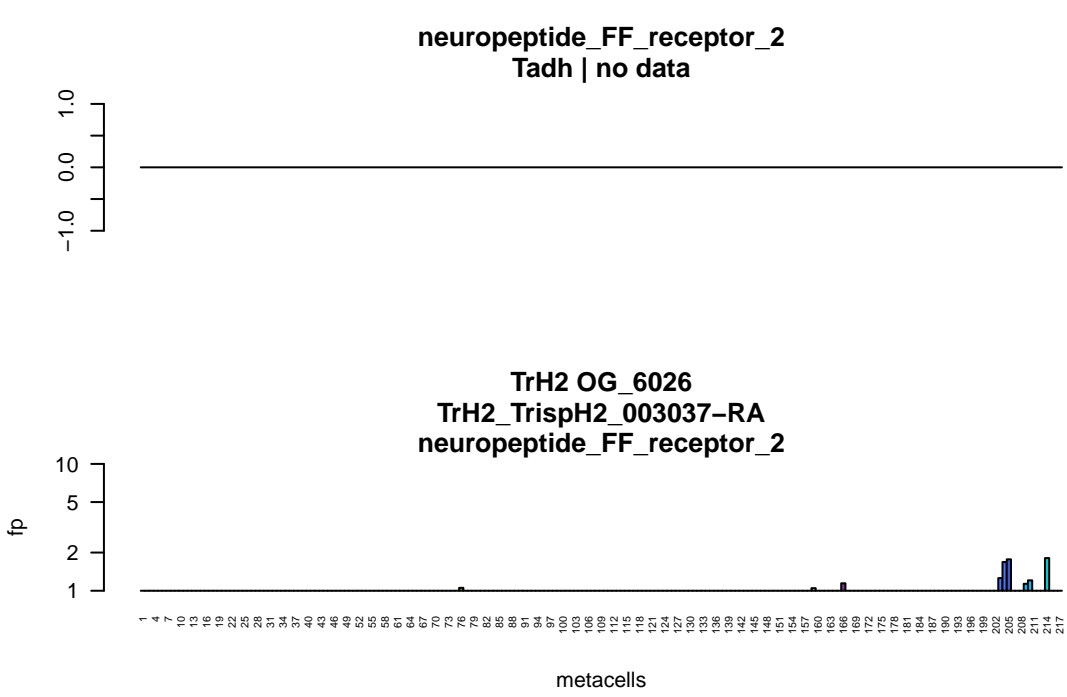


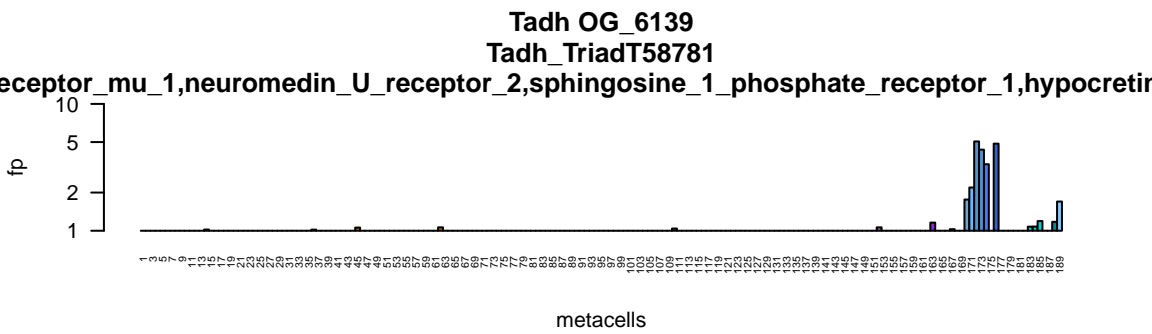


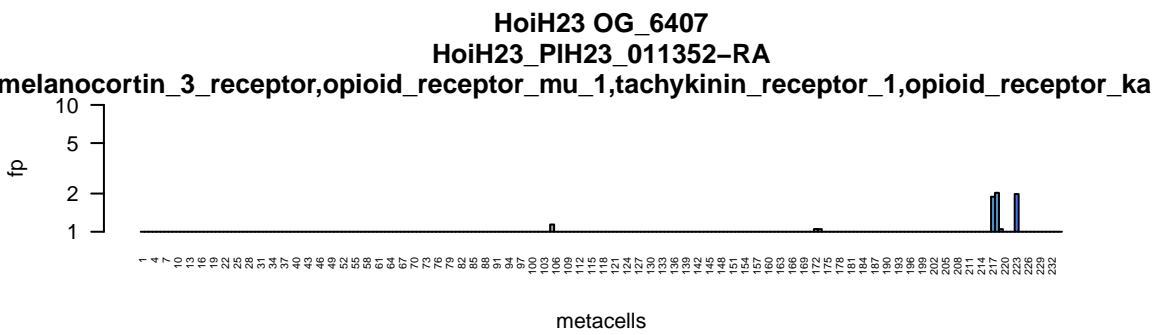
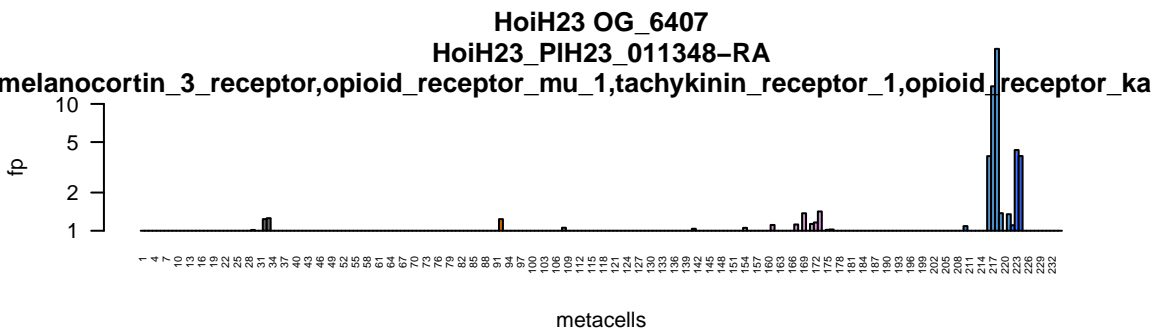
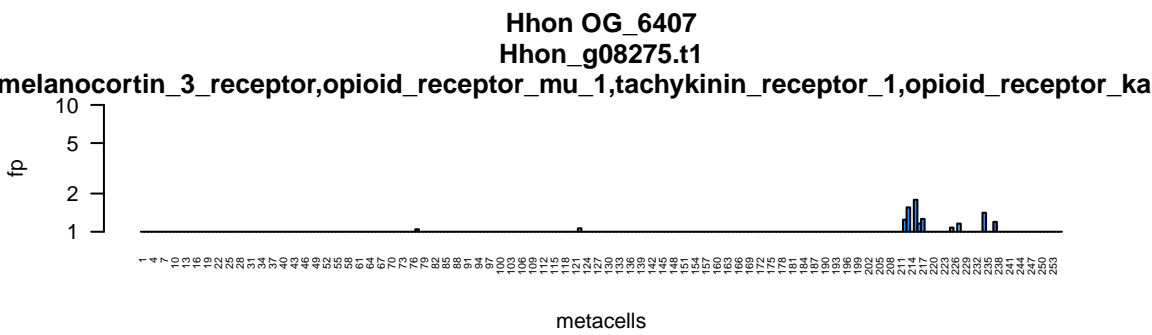
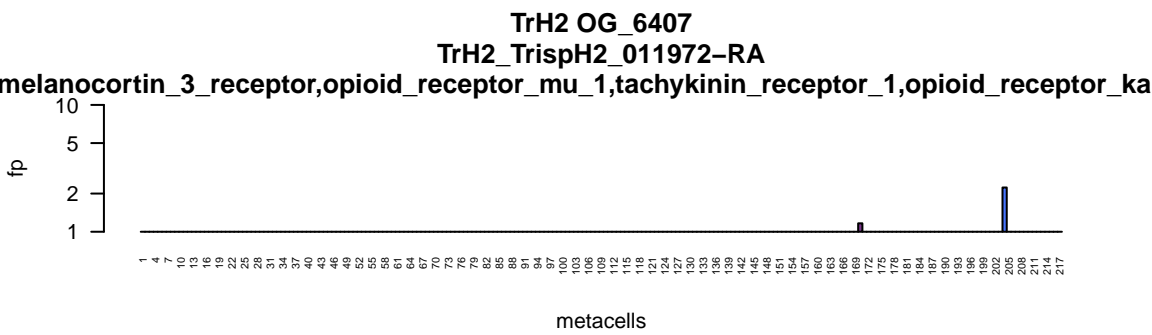
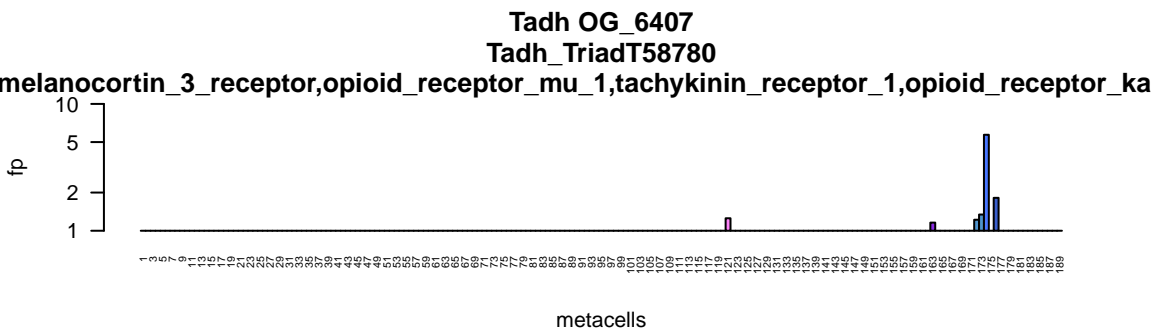








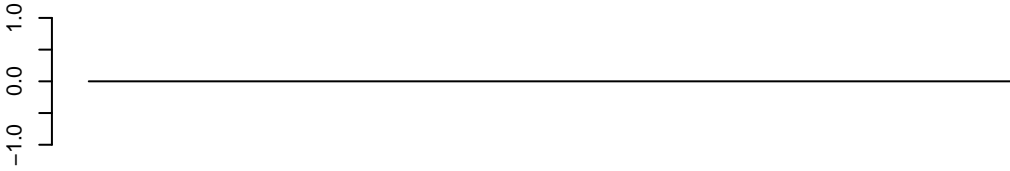




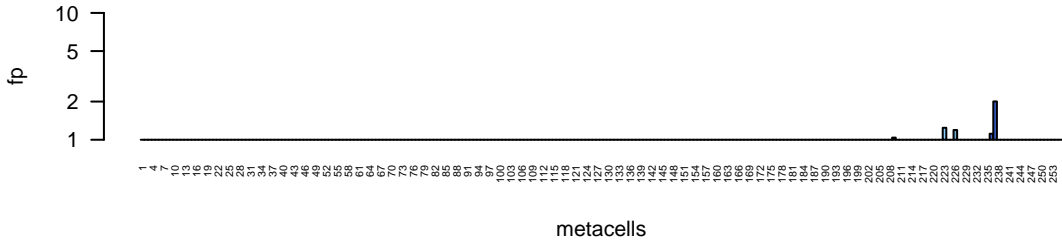
Tadh | no data



TrH2 | no data

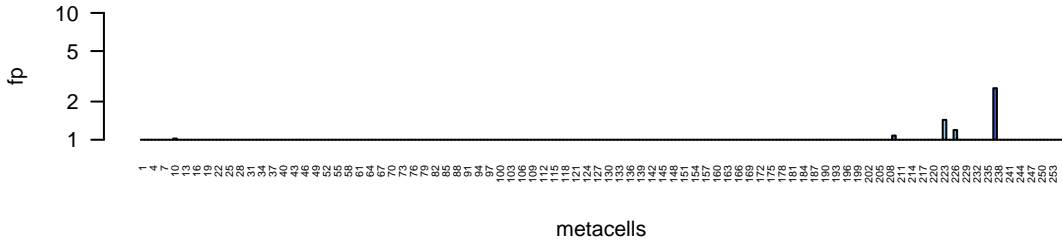


Hhon OG_6424
Hhon_g11706.t1



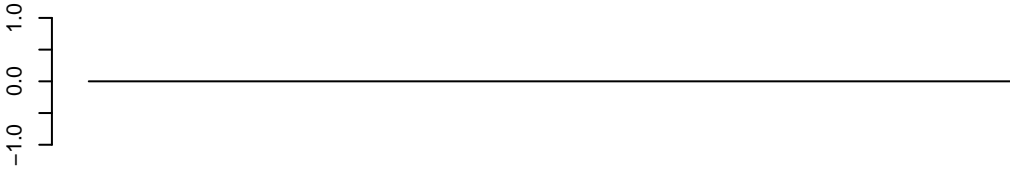
metacells

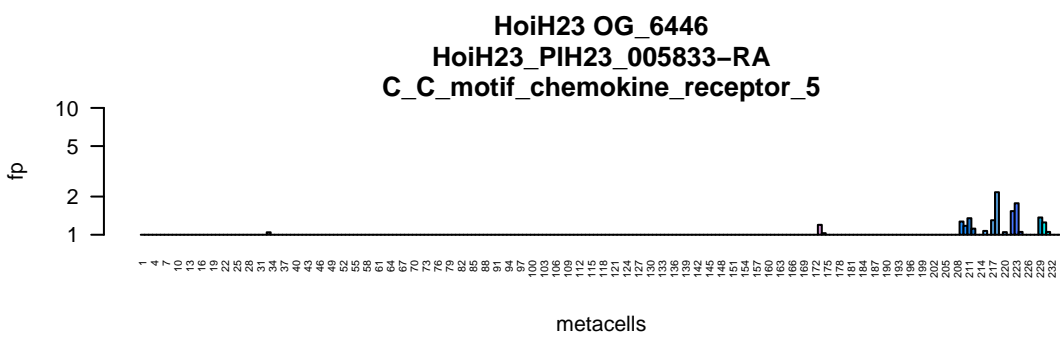
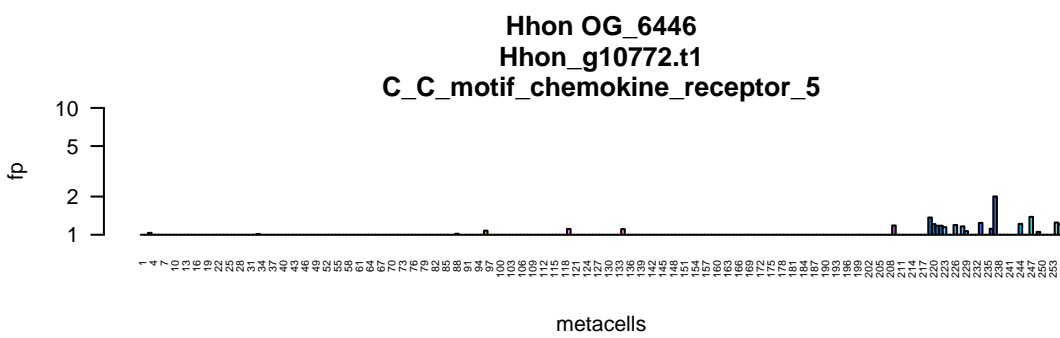
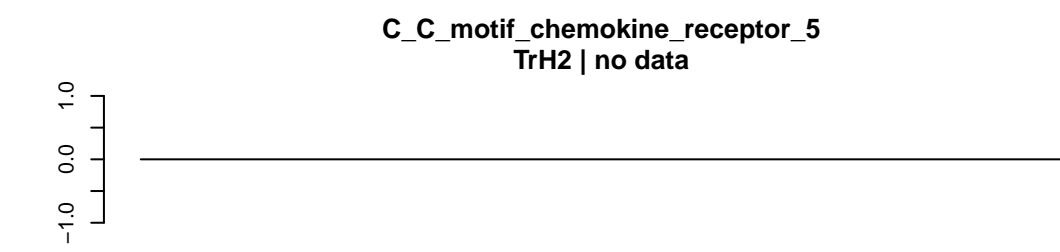
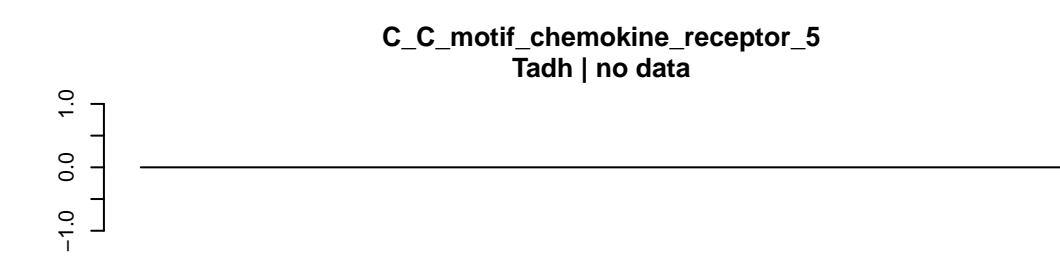
Hhon OG_6424
Hhon_g01935.t1

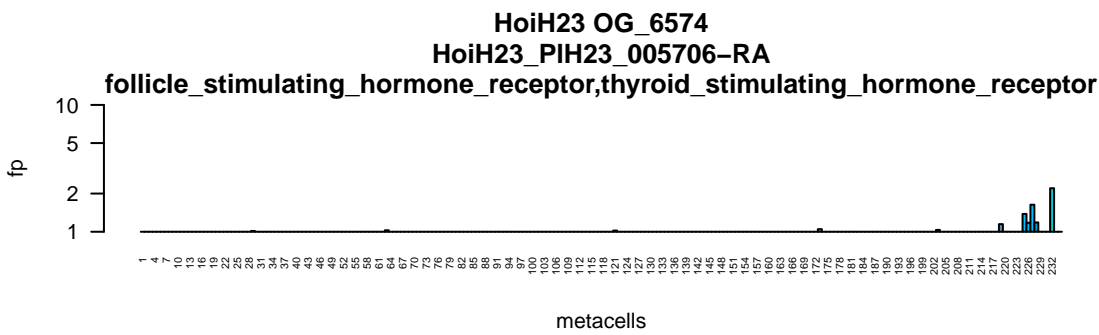
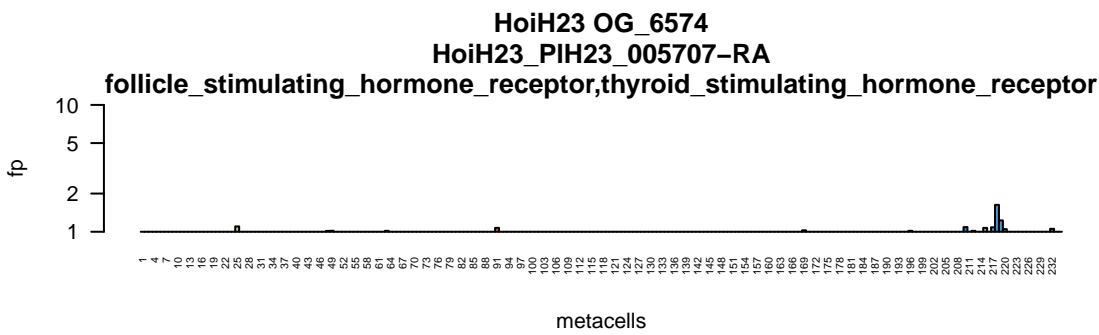
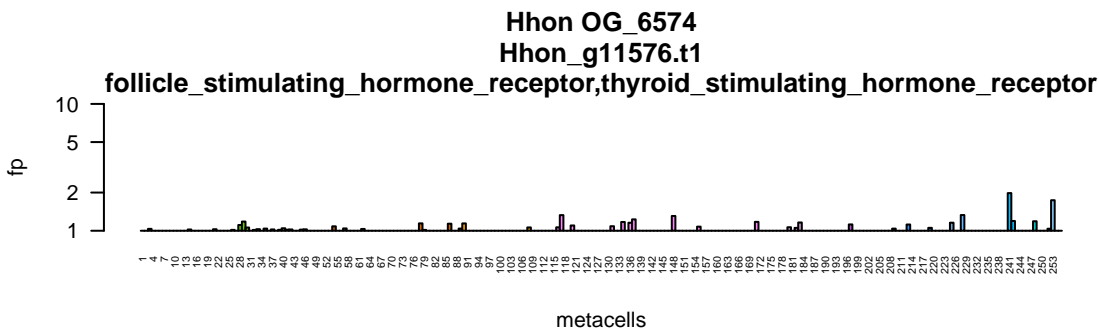
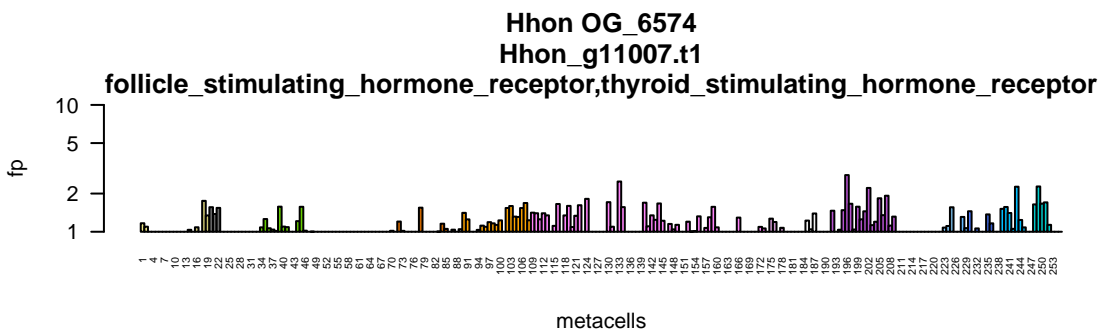
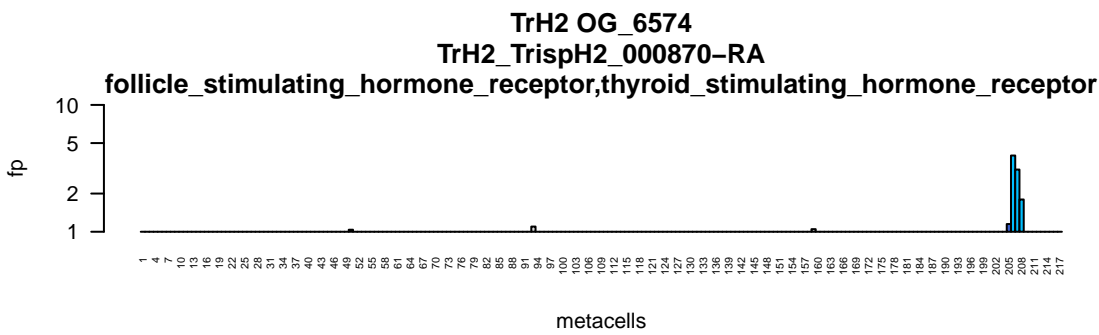
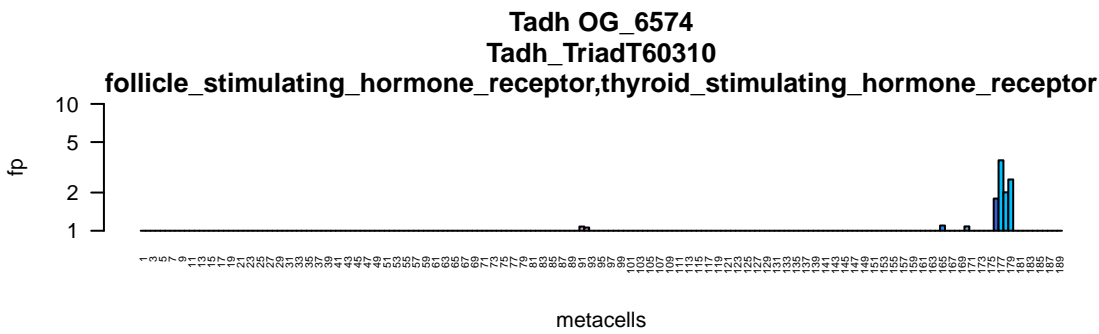


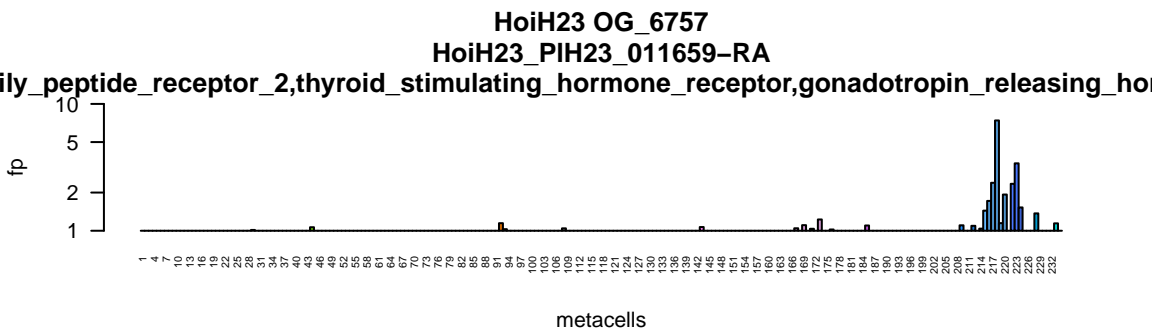
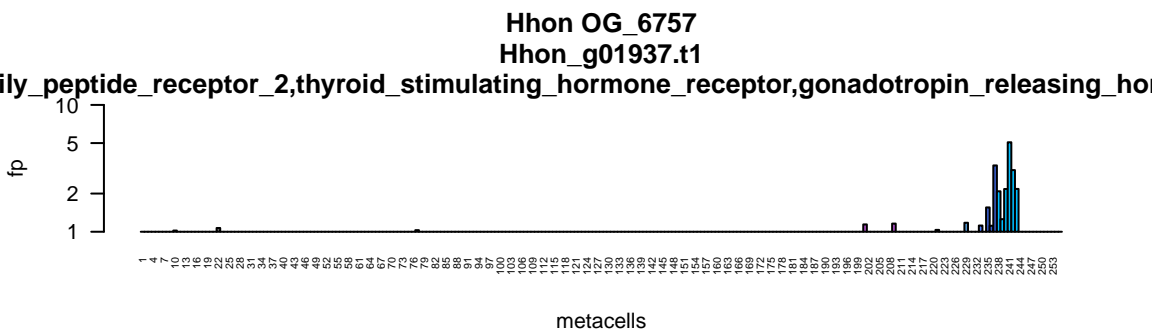
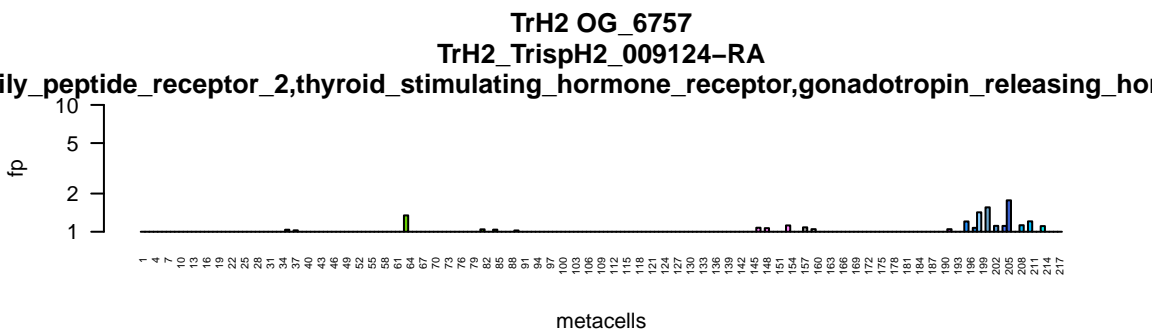
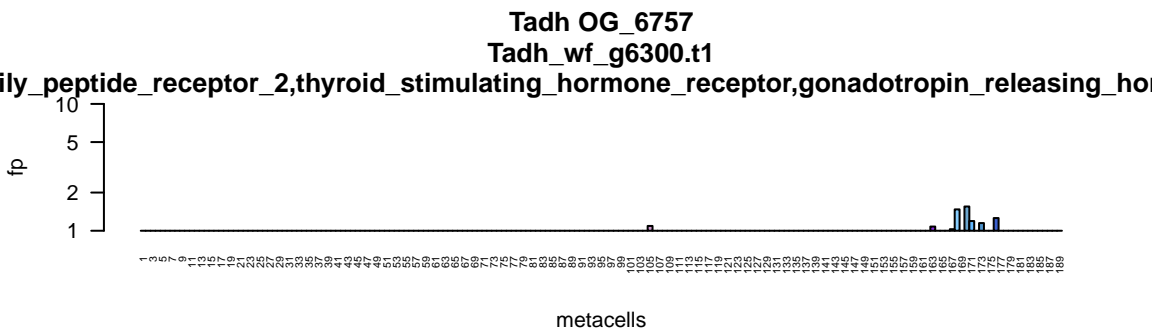
metacells

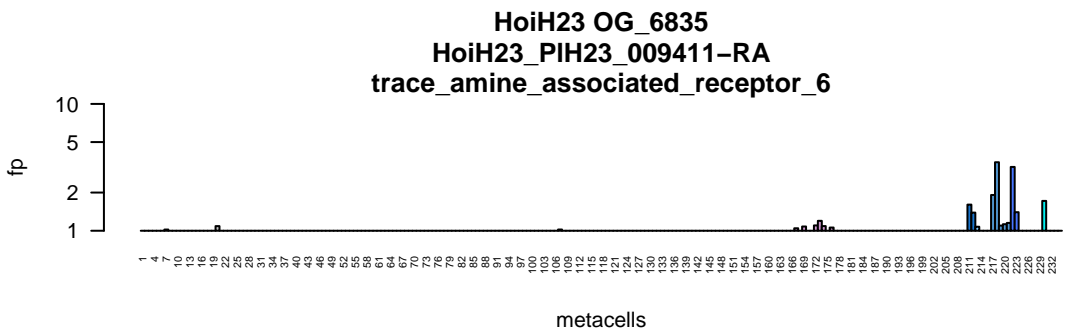
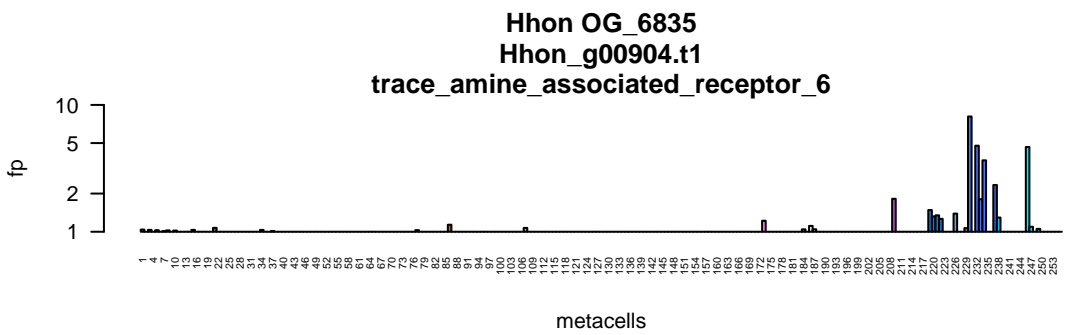
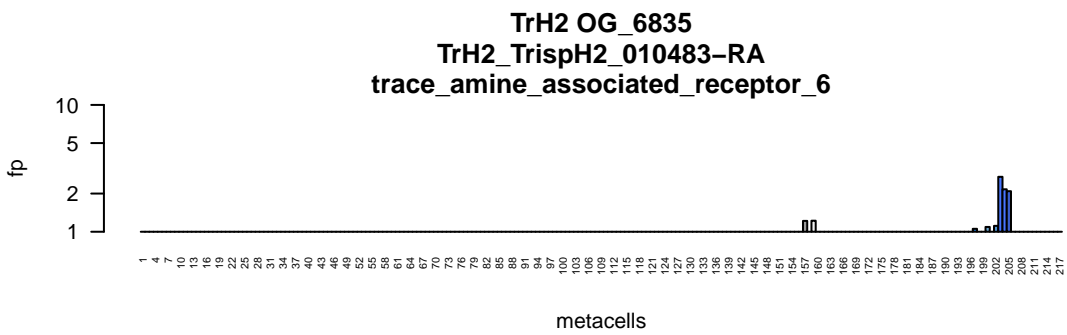
HoiH23 | no data

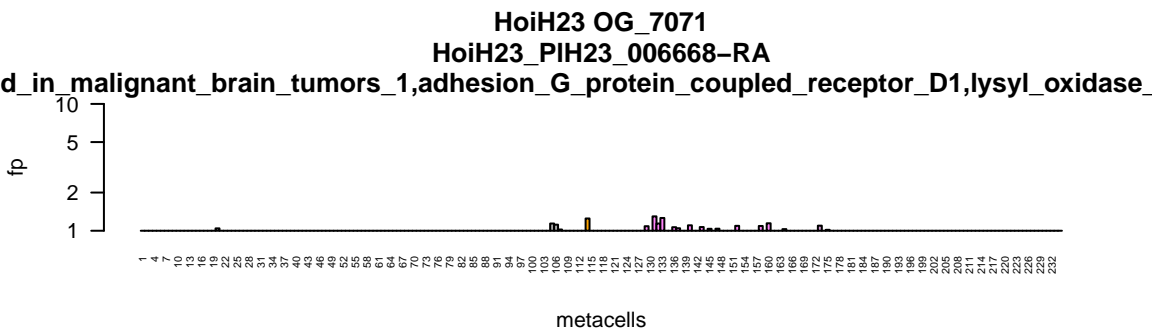
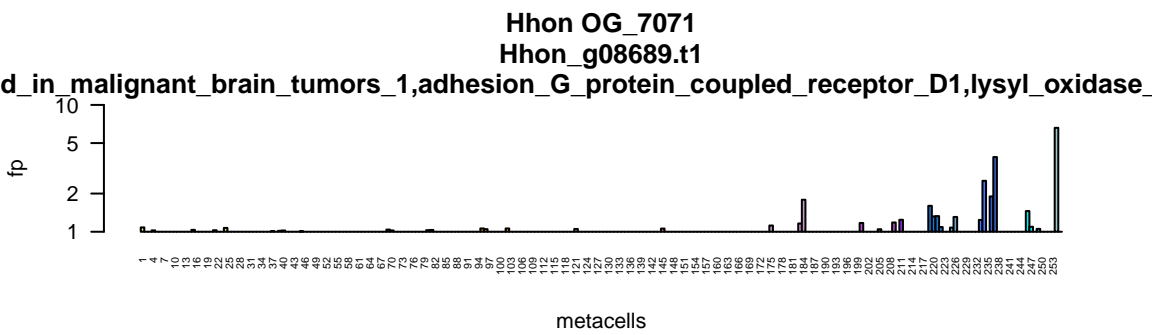
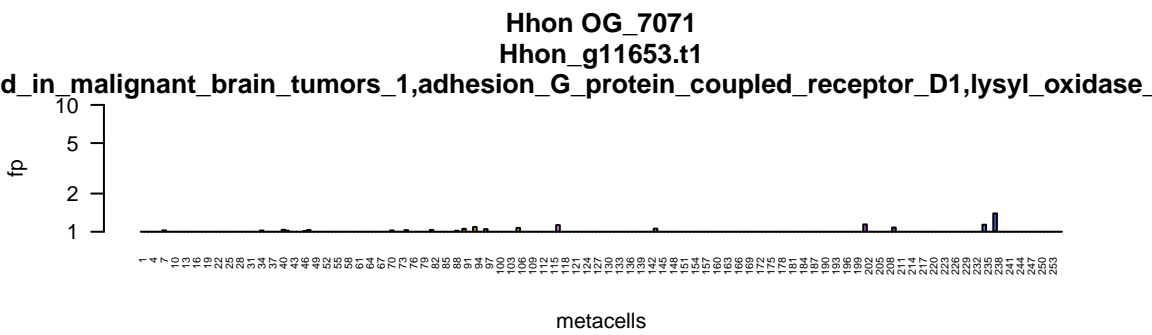
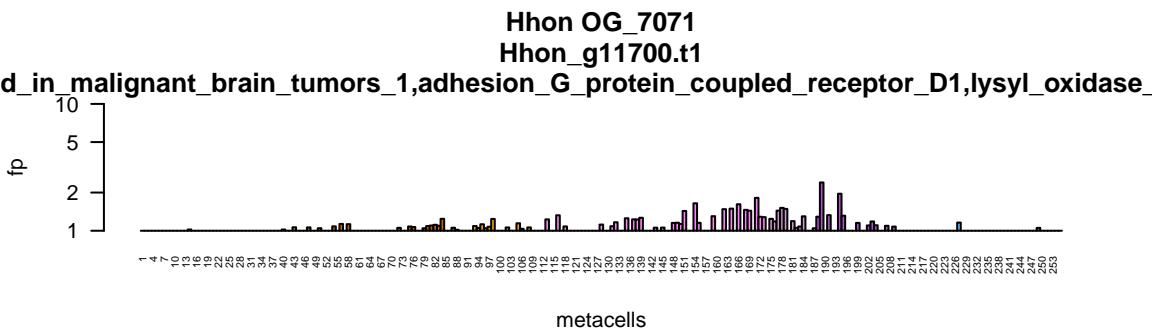
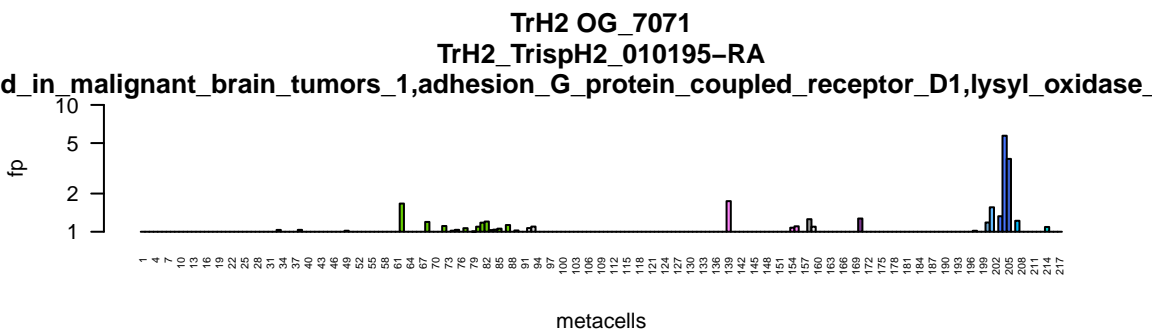
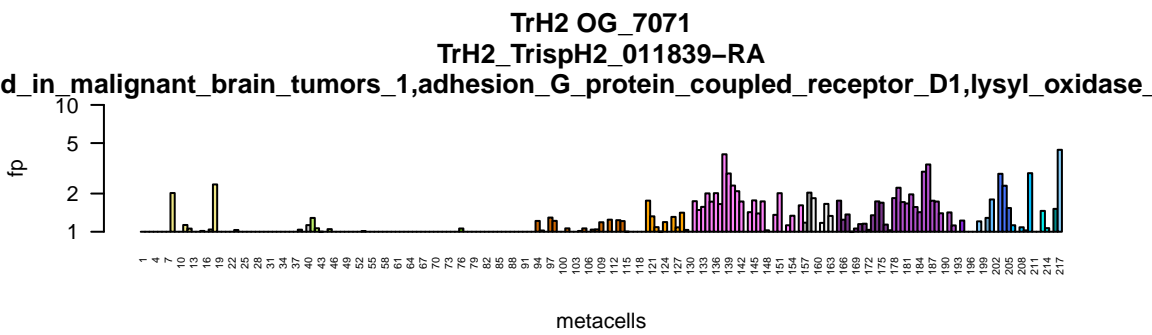
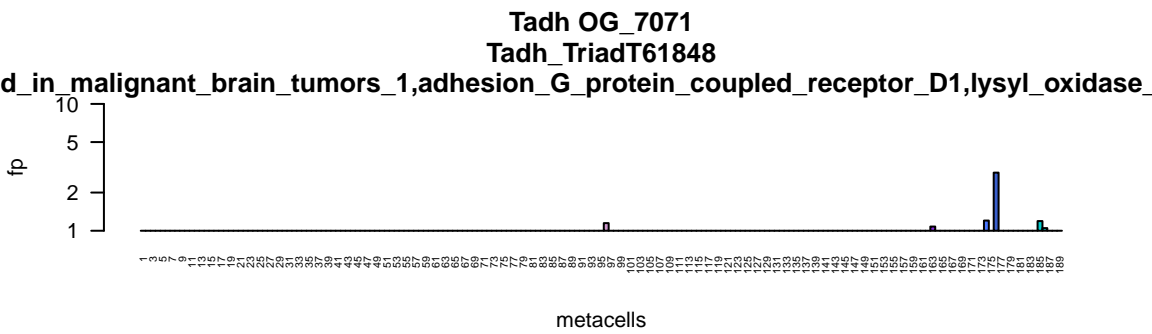






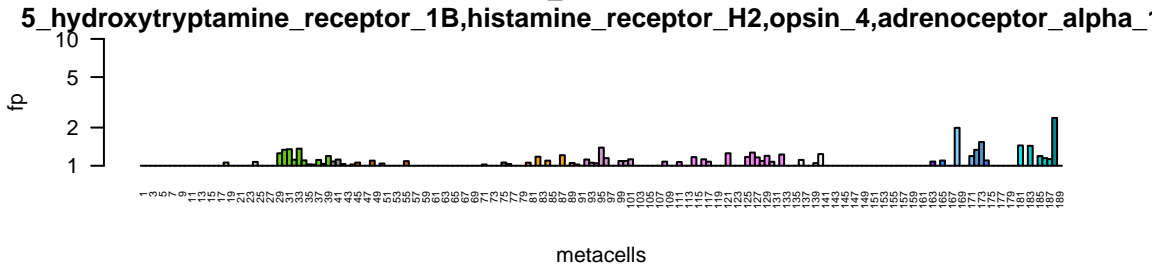






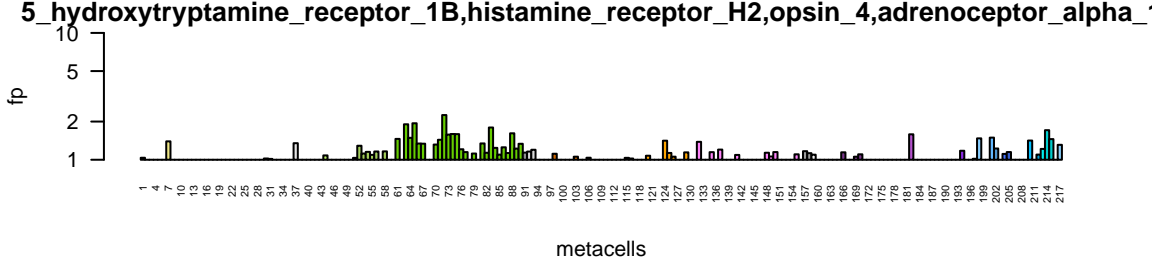
Tadh OG_7573

Tadh_TriadT26040



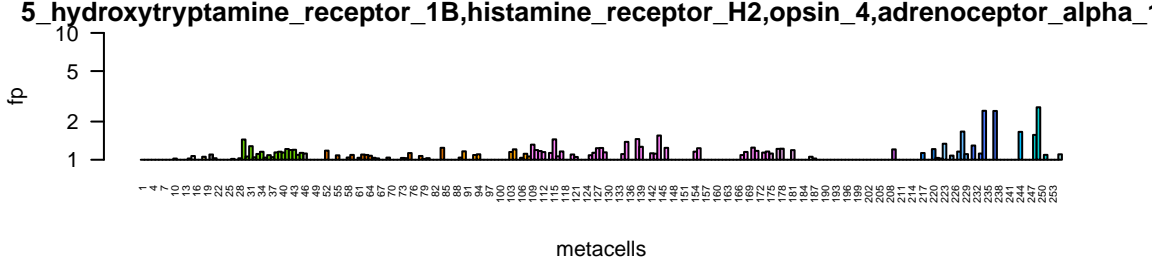
TrH2 OG_7573

TrH2_TrispH2_008979-RA



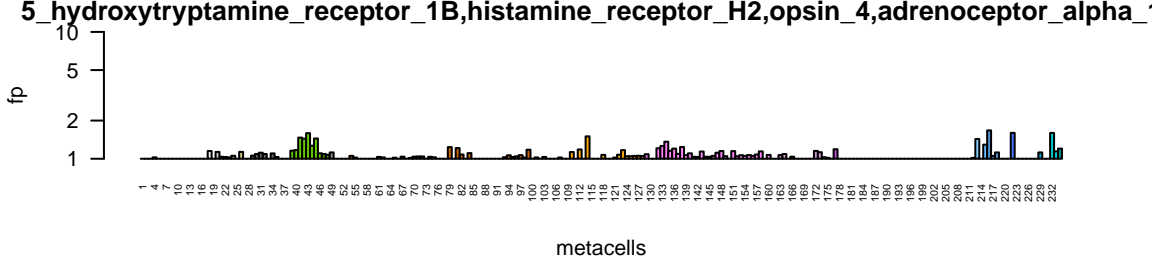
Hhon OG_7573

Hhon_g05380.t1

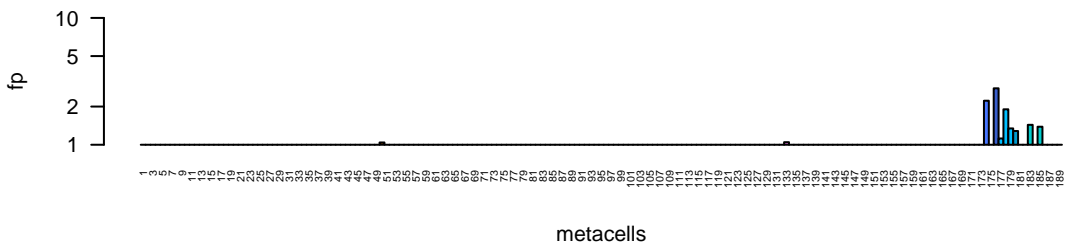


HoiH23 OG_7573

HoiH23_PIH23_000307-RA

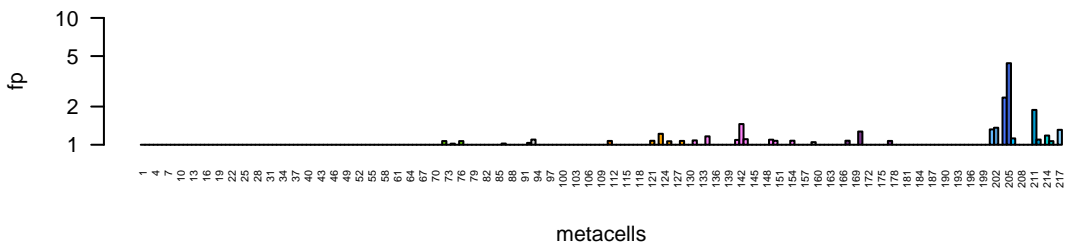


Tadh OG_8094
Tadh_wf_g6497.t1



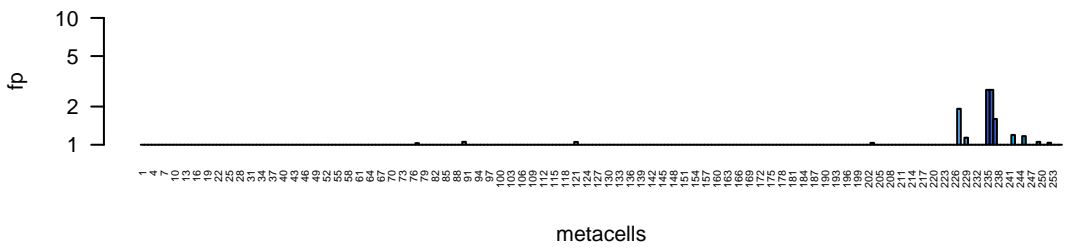
metacells

TrH2 OG_8094
TrH2_TrispH2_011737-RA



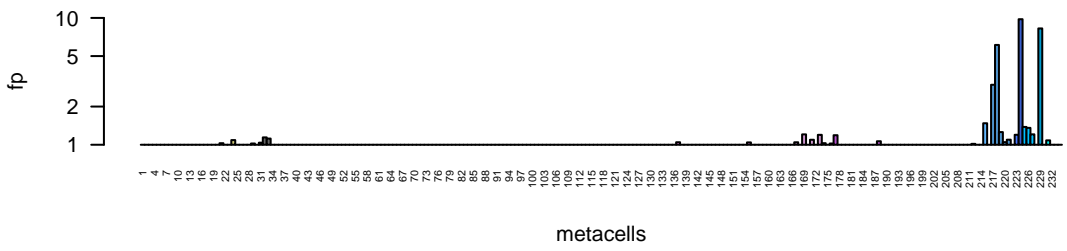
metacells

Hhon OG_8094
Hhon_g08752.t1



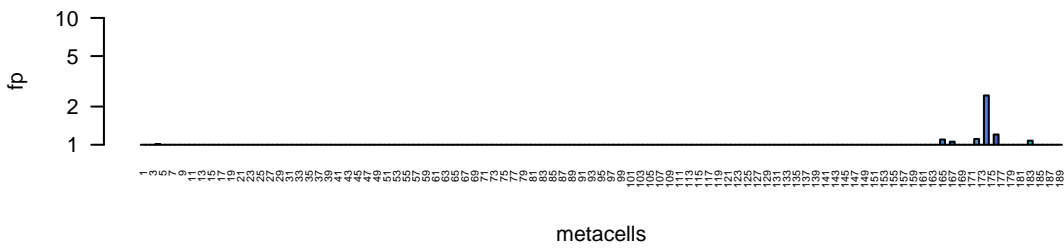
metacells

HoiH23 OG_8094
HoiH23_PIH23_007147-RA

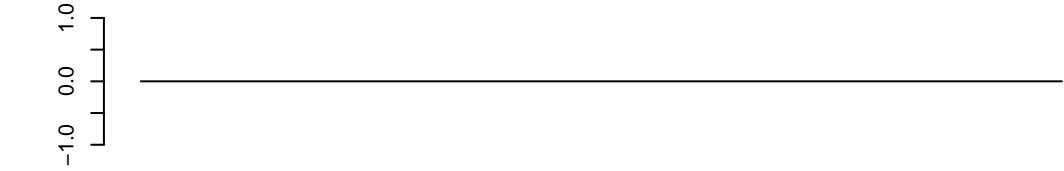


metacells

Tadh OG_8095
Tadh_TriadT57627



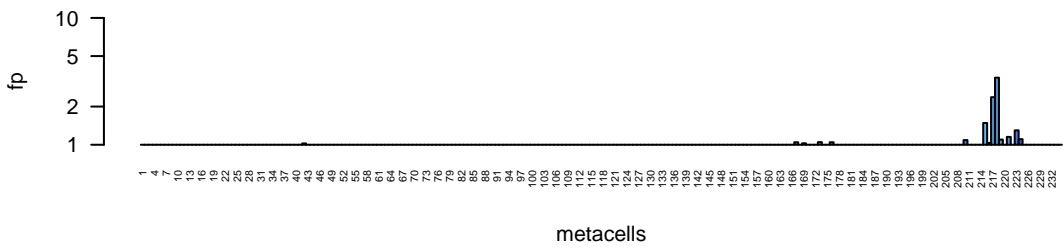
TrH2 | no data



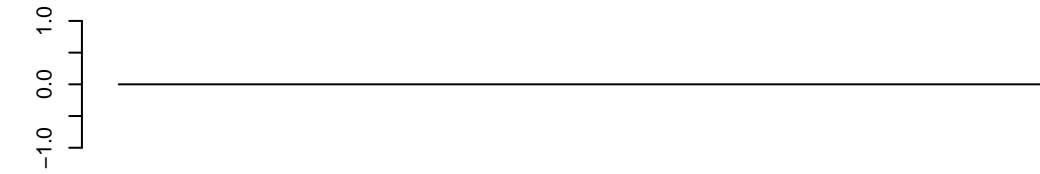
Hhon | no data



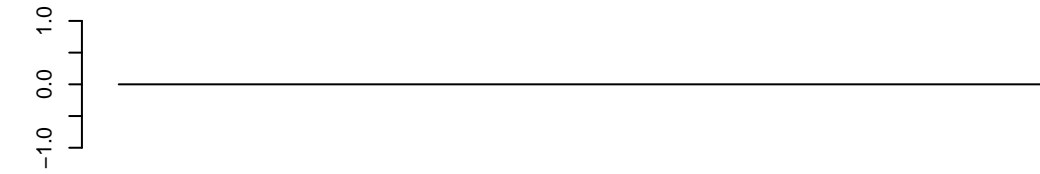
HoiH23 OG_8095
HoiH23_PIH23_007148-RA



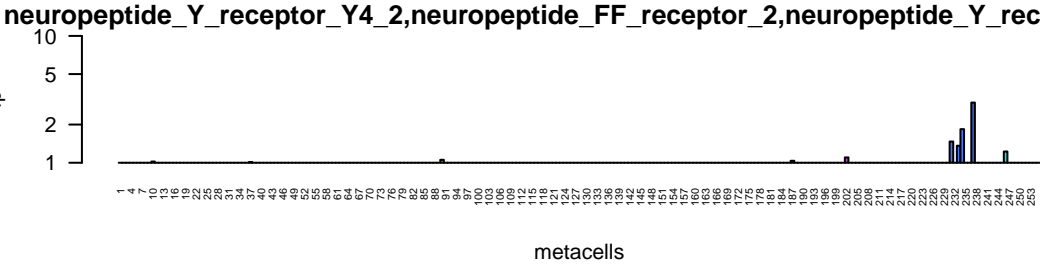
neuropeptide_Y_receptor_Y4_2,neuropeptide_FF_receptor_2,neuropeptide_Y_receptor_\
Tadh | no data



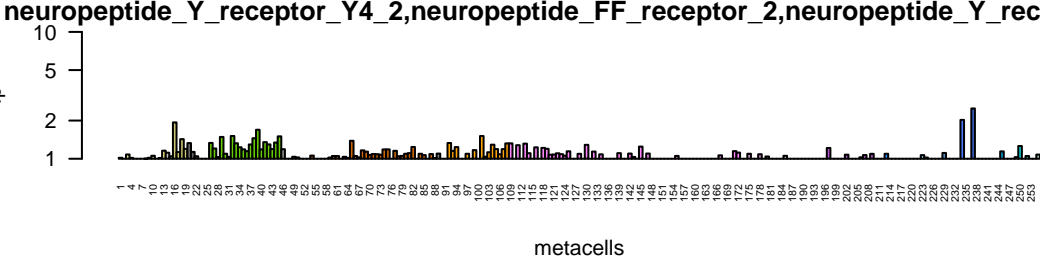
neuropeptide_Y_receptor_Y4_2,neuropeptide_FF_receptor_2,neuropeptide_Y_receptor_\
TrH2 | no data



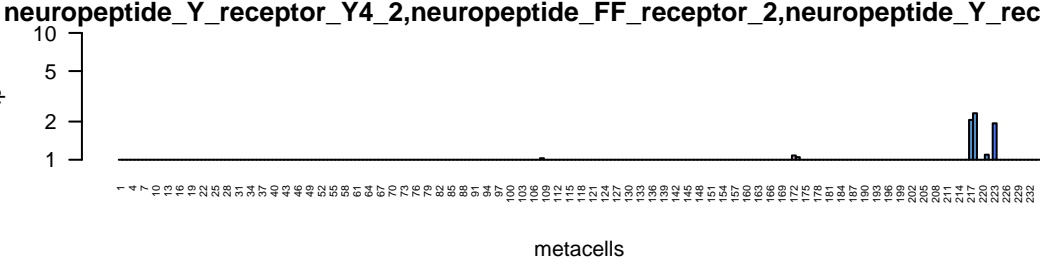
Hhon OG_8289
Hhon_g07093.t1
neuropeptide_Y_receptor_Y4_2,neuropeptide_FF_receptor_2,neuropeptide_Y_receptor_\
metacells

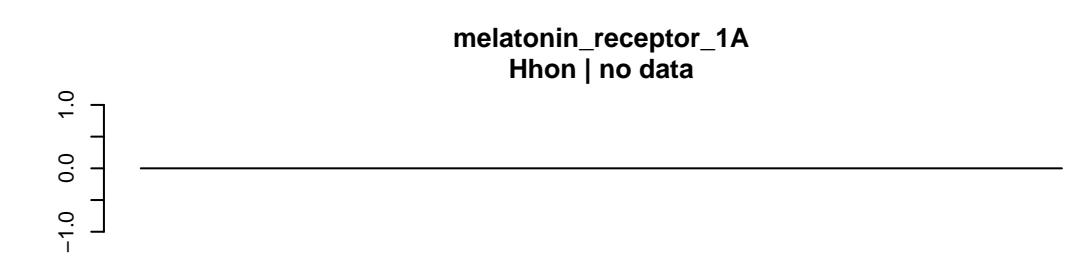
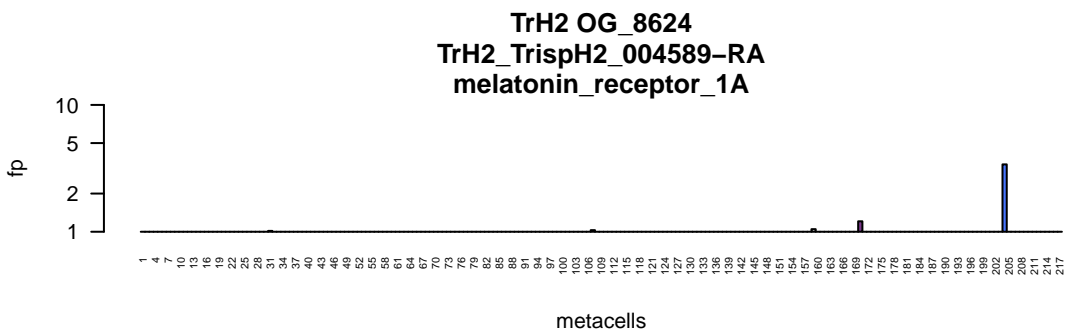
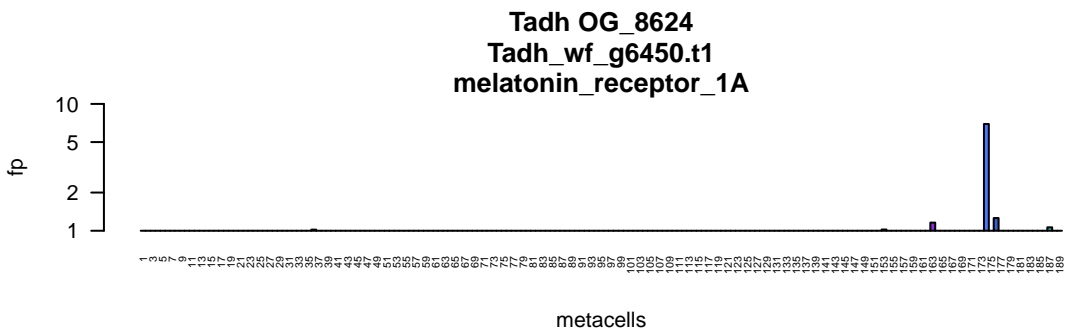


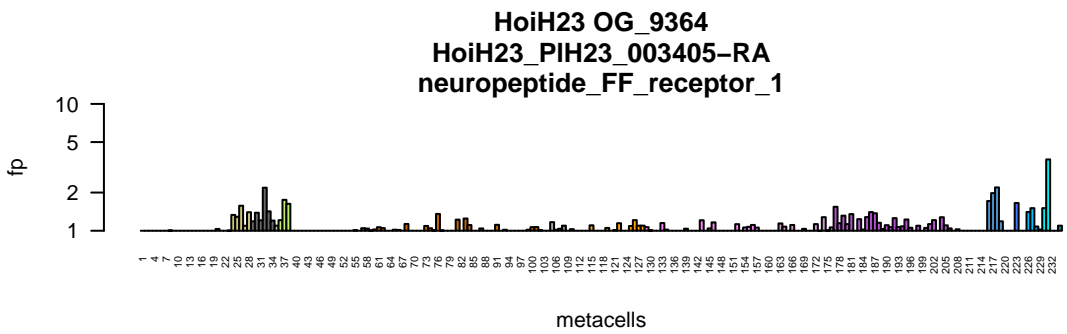
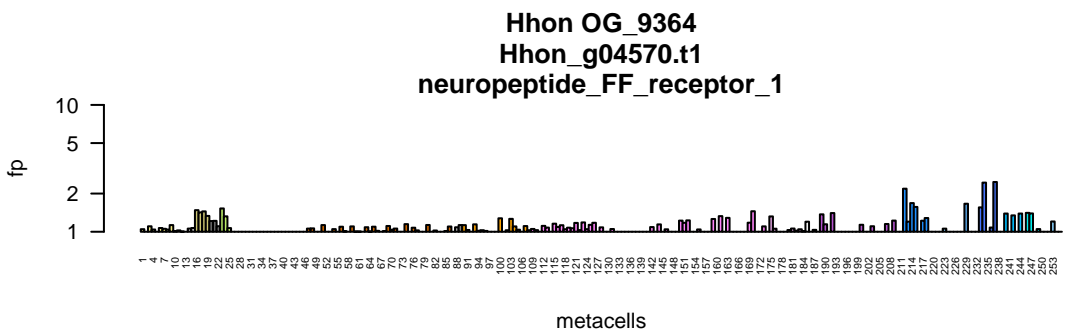
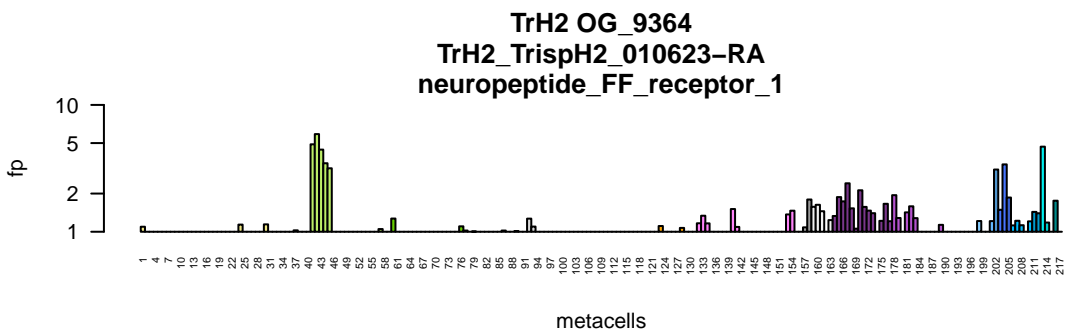
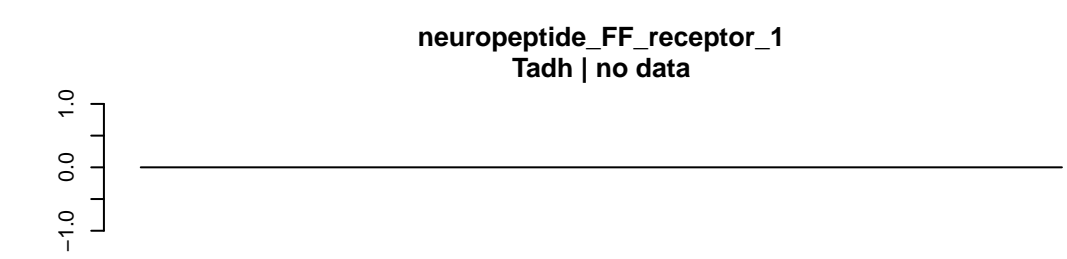
Hhon OG_8289
Hhon_g11435.t1
neuropeptide_Y_receptor_Y4_2,neuropeptide_FF_receptor_2,neuropeptide_Y_receptor_\
metacells



HoiH23 OG_8289
HoiH23_PIH23_002693-RA
neuropeptide_Y_receptor_Y4_2,neuropeptide_FF_receptor_2,neuropeptide_Y_receptor_\
metacells



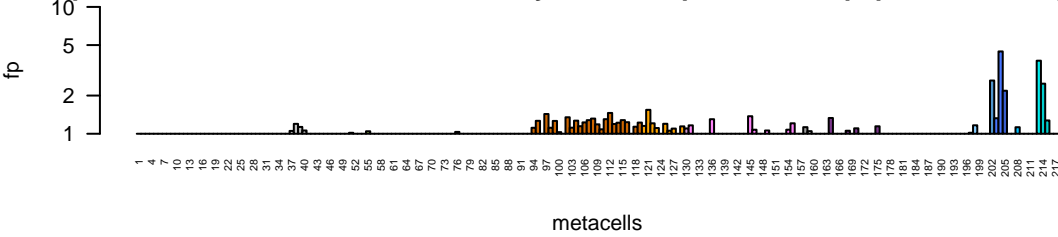




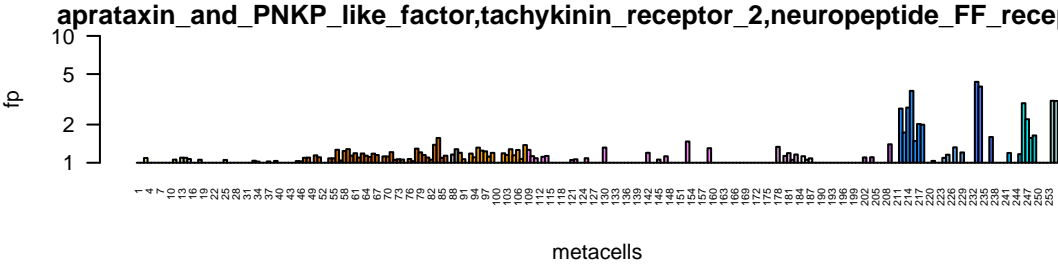
apratatin_and_PNKP_like_factor,tachykinin_receptor_2,neuropeptide_FF_receptor_1
Tadh | no data



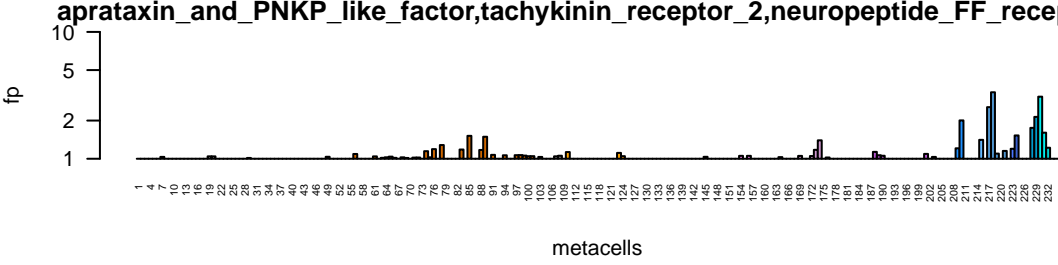
TrH2 OG_9368
TrH2_TrispH2_010619-RA
apratatin_and_PNKP_like_factor,tachykinin_receptor_2,neuropeptide_FF_receptor_1

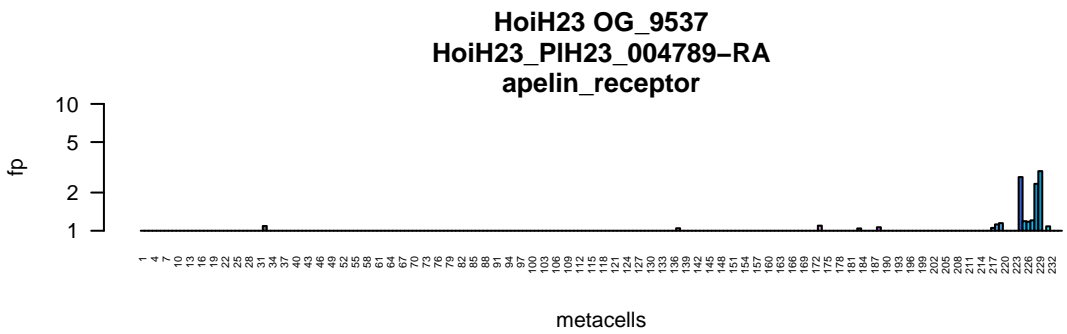
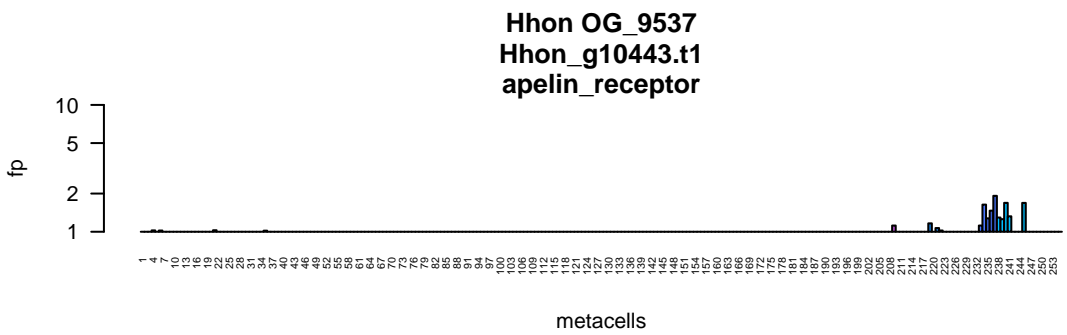
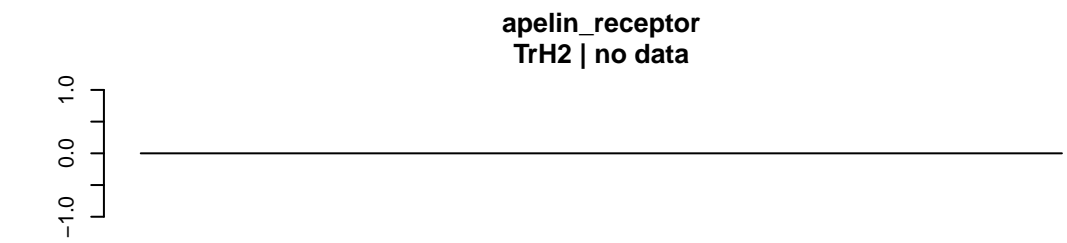
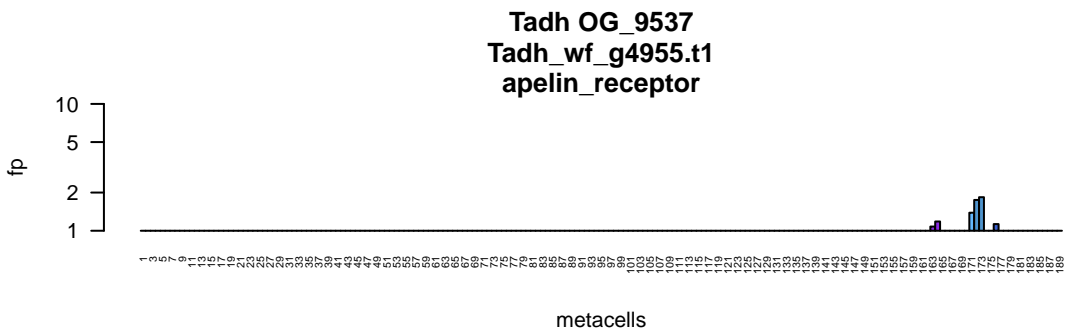


Hhon OG_9368
Hhon_g04575.t1
apratatin_and_PNKP_like_factor,tachykinin_receptor_2,neuropeptide_FF_receptor_1



HoiH23 OG_9368
HoiH23_PIH23_003409-RA
apratatin_and_PNKP_like_factor,tachykinin_receptor_2,neuropeptide_FF_receptor_1

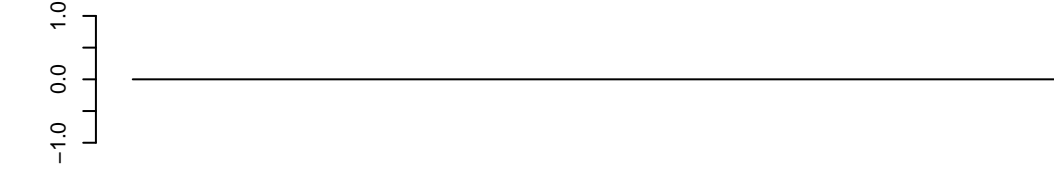




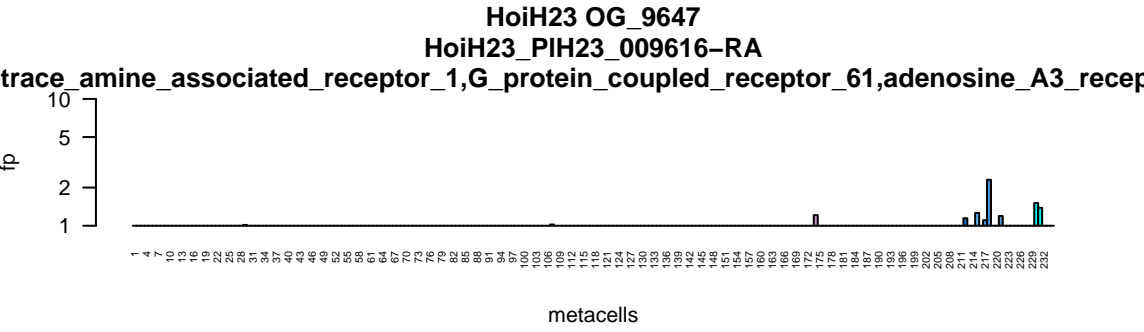
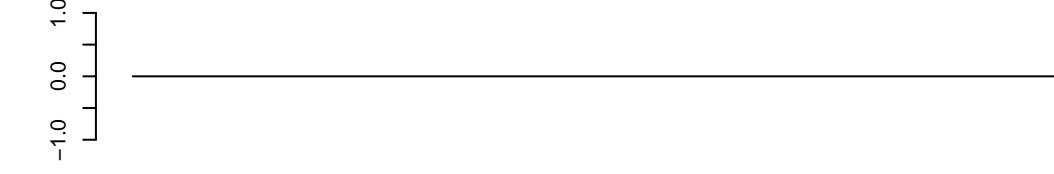
trace_amine_associated_receptor_1,G_protein_coupled_receptor_61,adenosine_A3_receptor
Tadh | no data



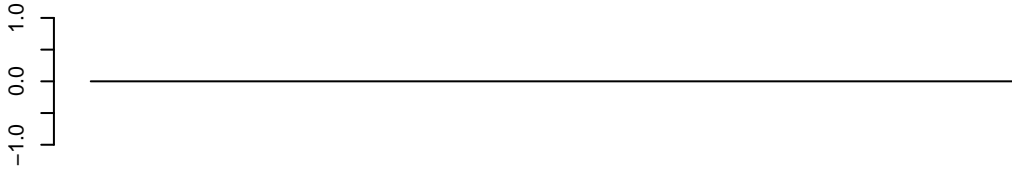
trace_amine_associated_receptor_1,G_protein_coupled_receptor_61,adenosine_A3_receptor
TrH2 | no data



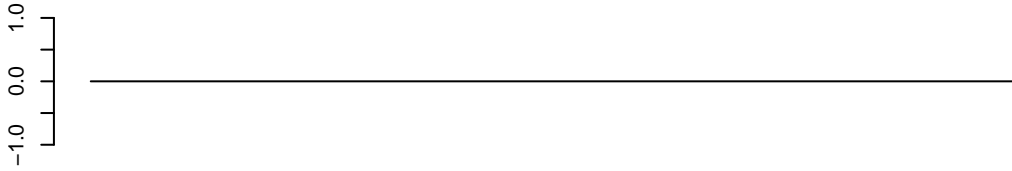
trace_amine_associated_receptor_1,G_protein_coupled_receptor_61,adenosine_A3_receptor
Hhon | no data



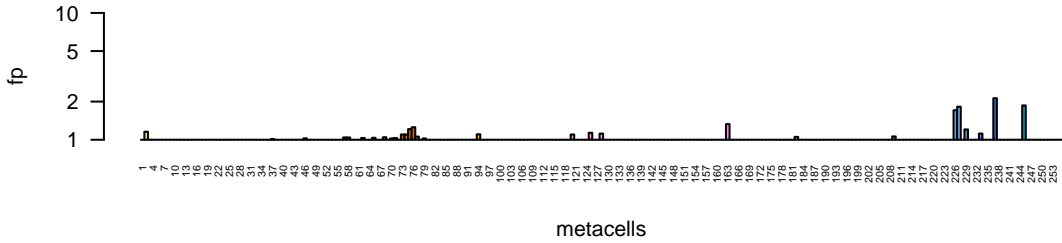
Tadh | no data



TrH2 | no data



Hhon OG_10393
Hhon_g01936.t1



HoiH23 | no data

