

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

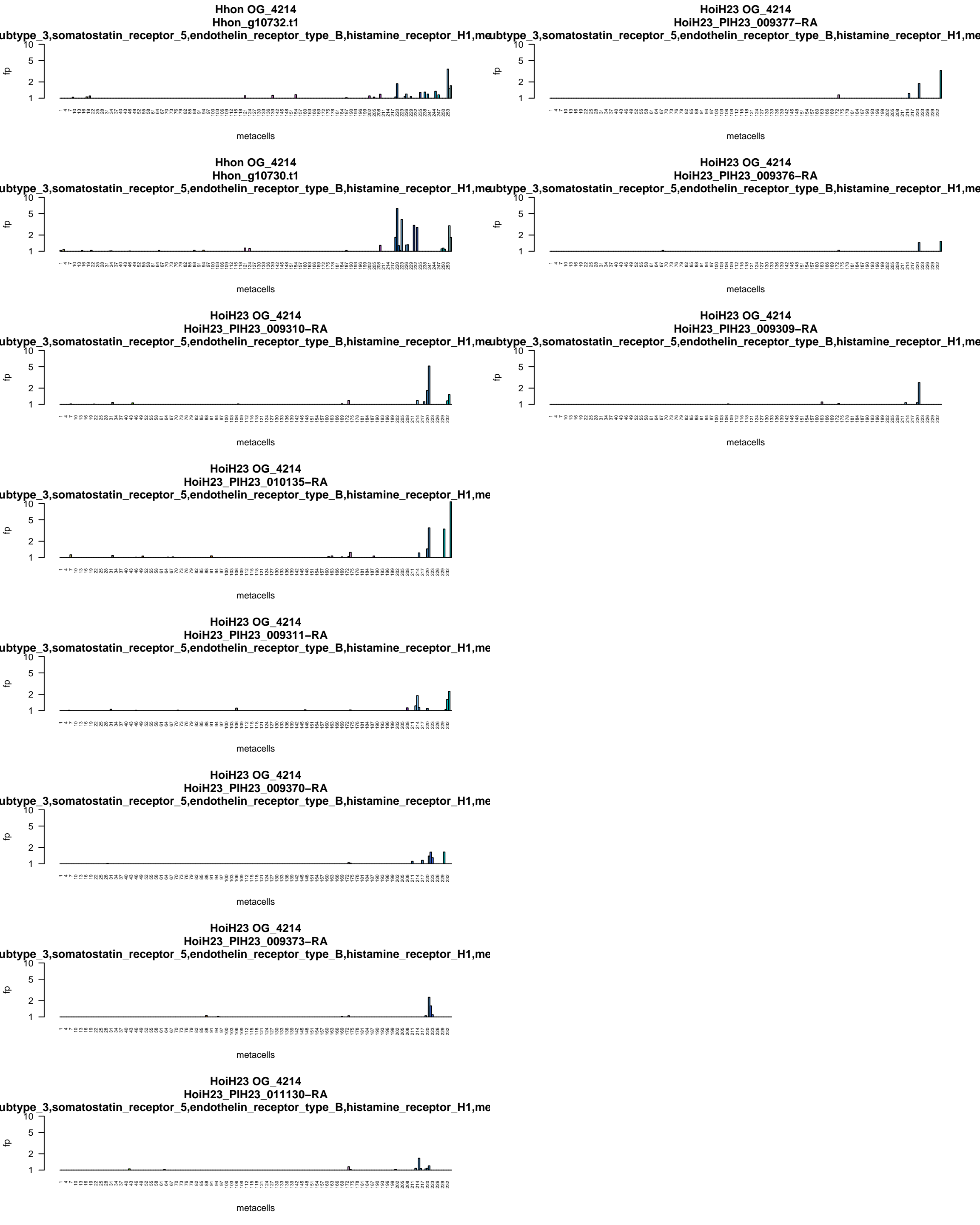
metacells

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



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
88	1
91	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

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
58	1
61	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	2.5
202	2.2
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

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

fp

metacells

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

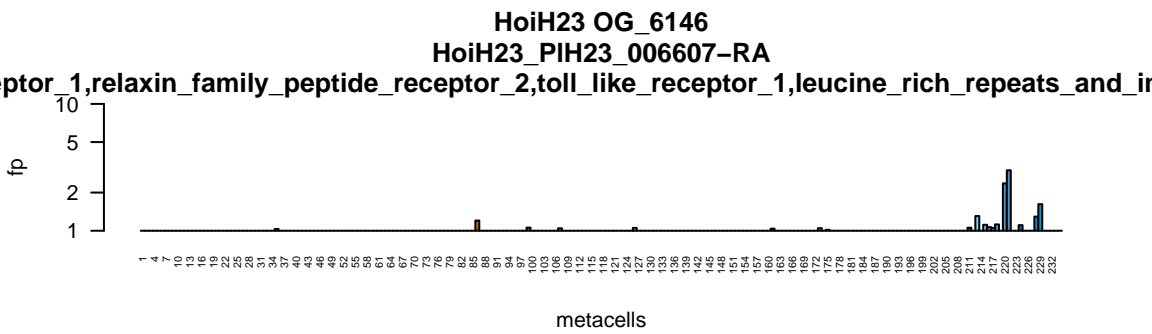
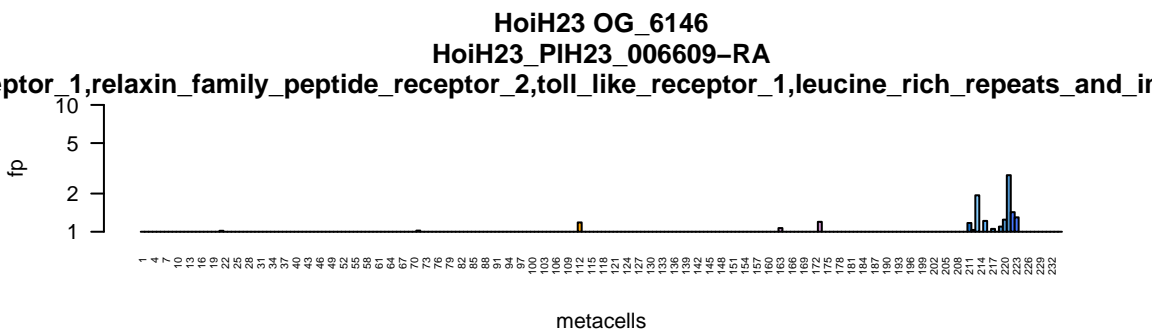
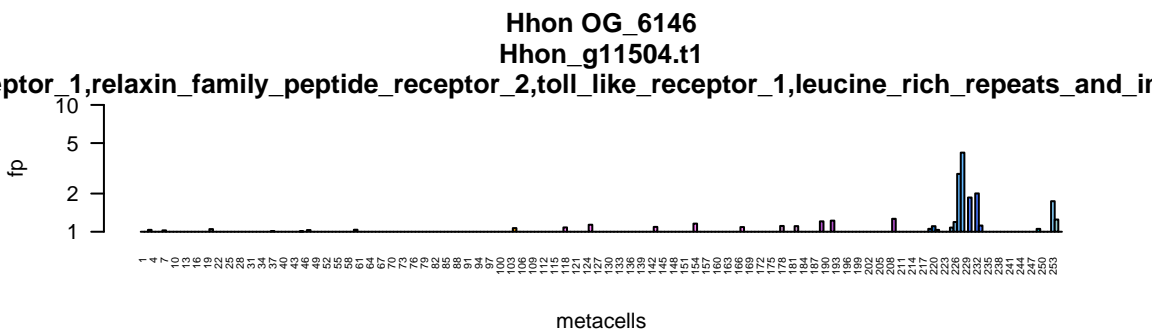
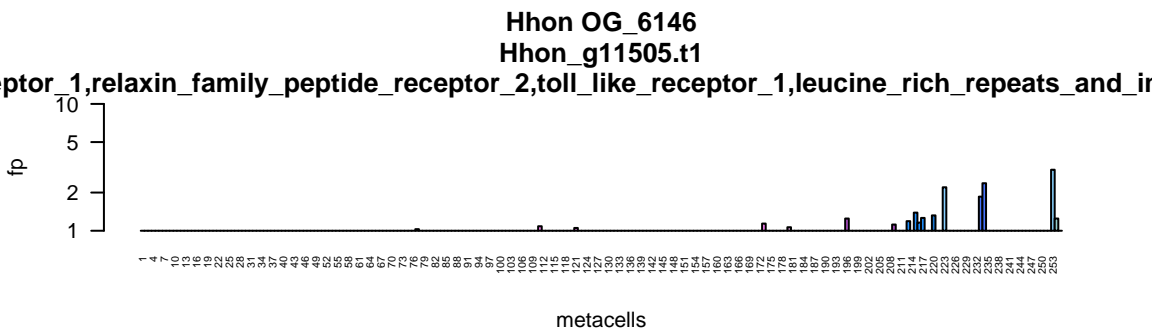
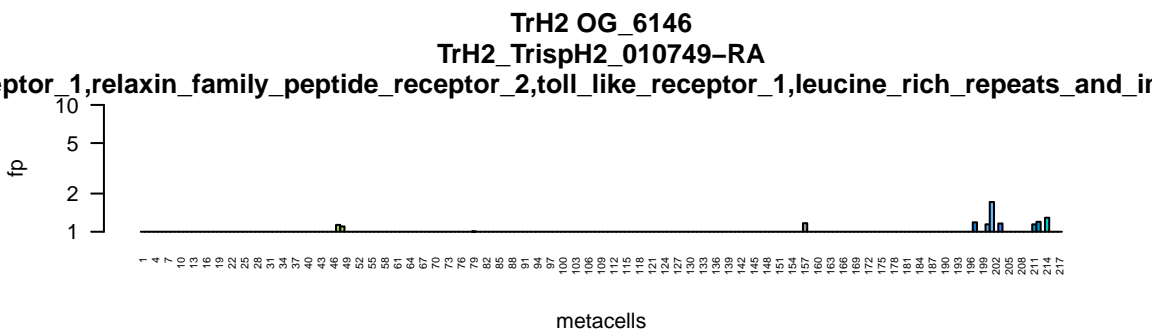
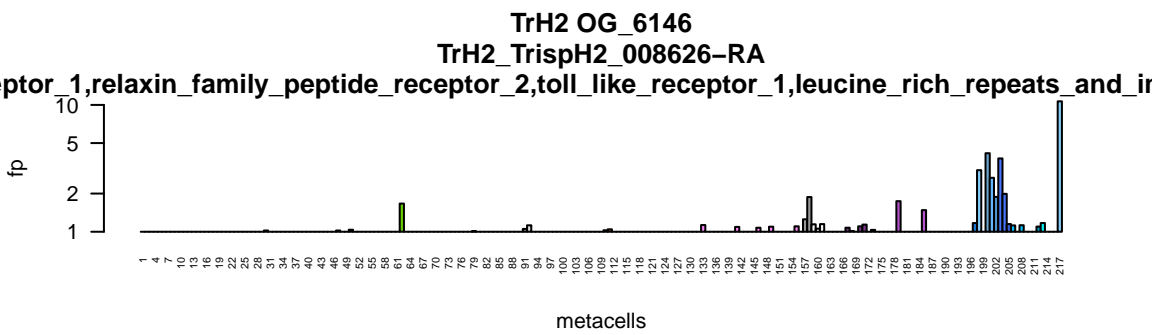
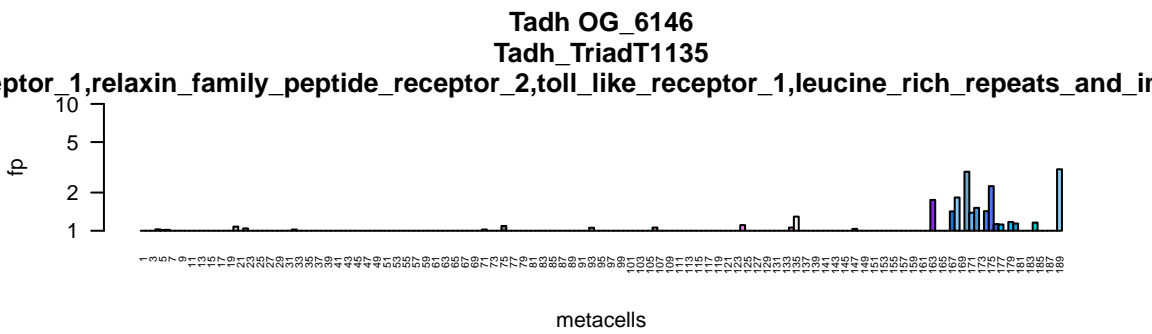
metacells	fp
1	1
4	1
10	1
13	1
19	1
22	1
28	1
31	1
37	1
40	1
46	1
52	1
55	1
64	1
68	1
70	1
73	1
79	1
88	1
91	1
97	1
100	1
106	1
109	1
115	1
118	1
127	1
124	1
133	1
142	1
145	1
151	1
154	1
163	1
169	1
172	1
178	1
184	1
187	1
193	1
196	1
202	1
205	1
211	1
214	1
220	1
223	1
229	1
232	1
238	1
241	1
247	1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

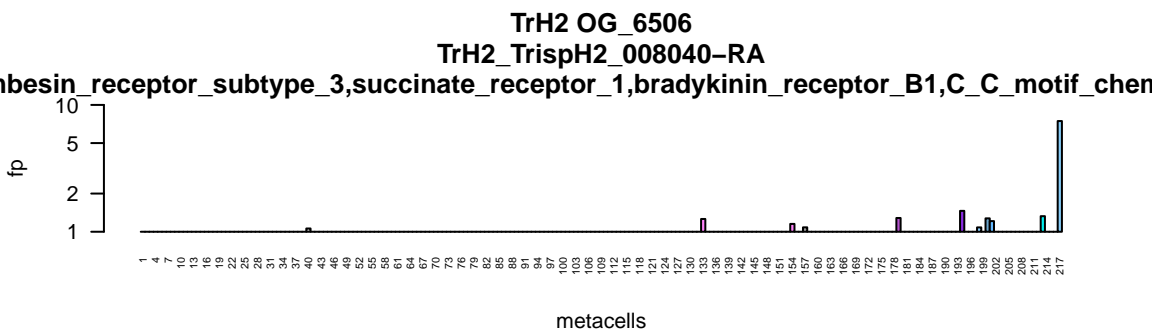
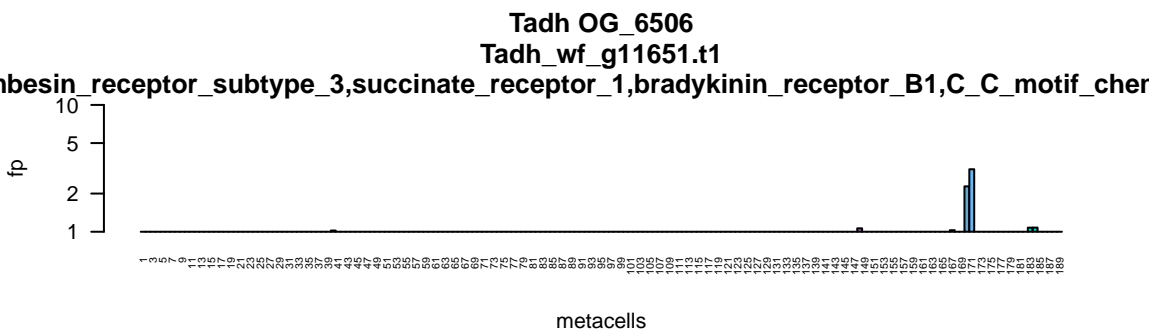
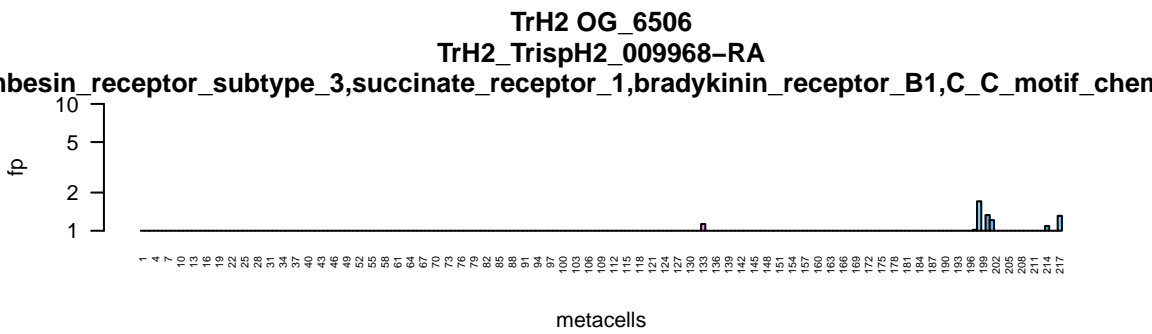
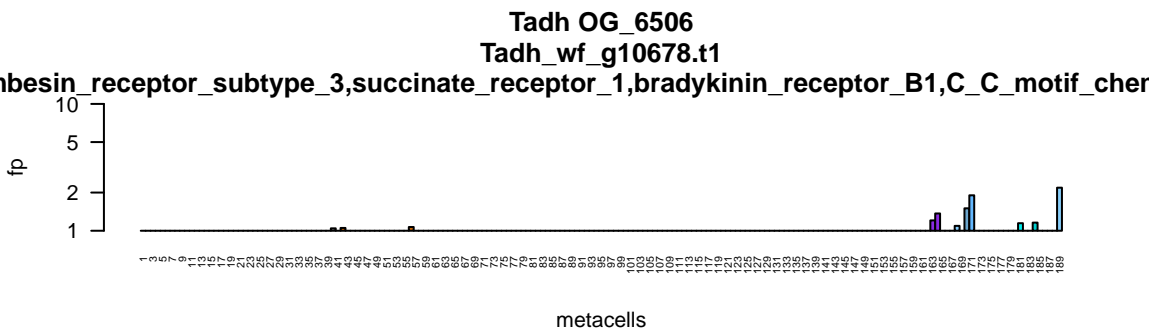
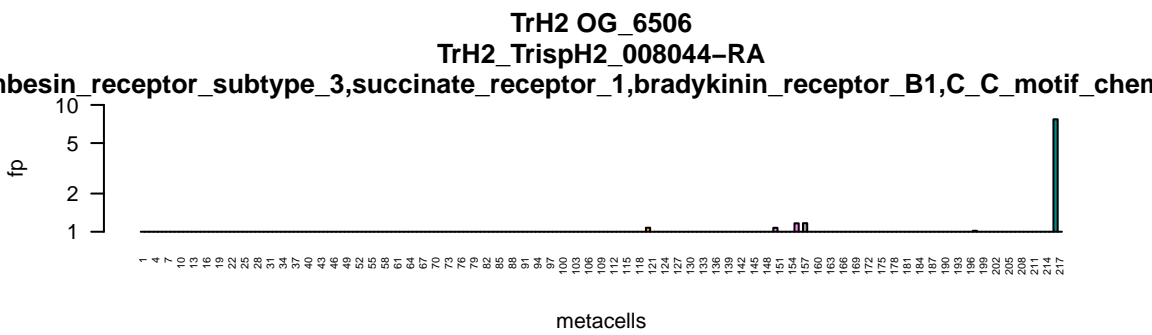
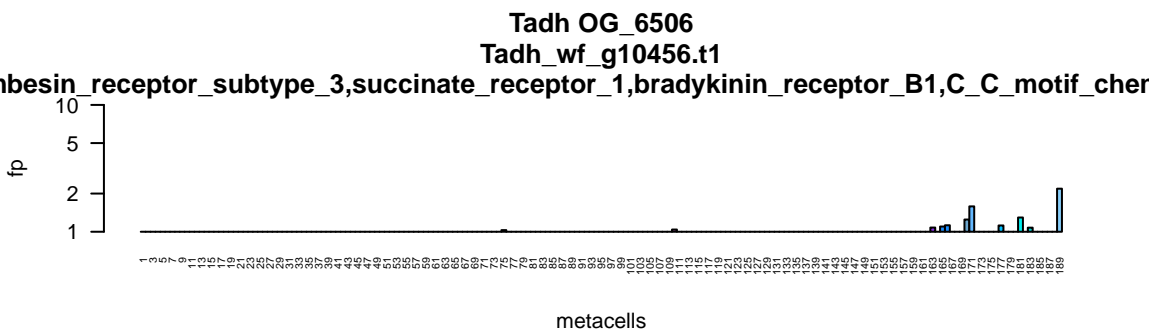
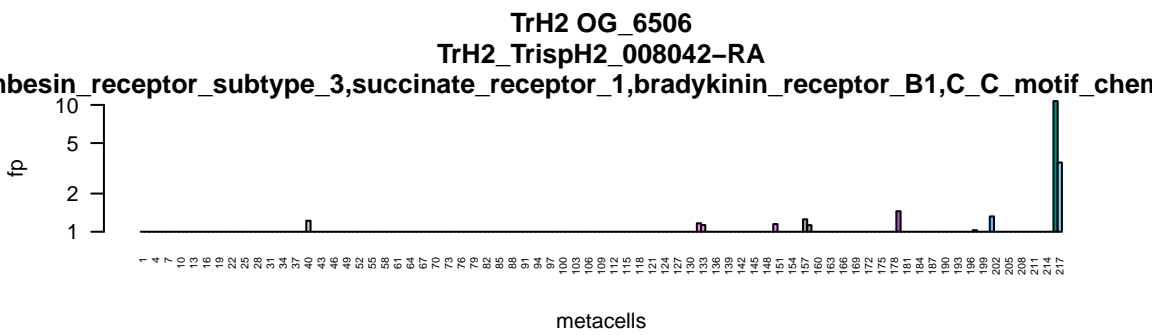
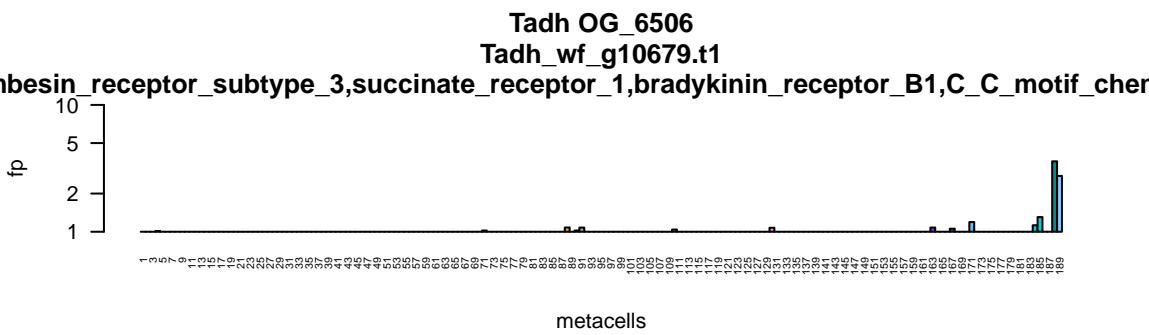
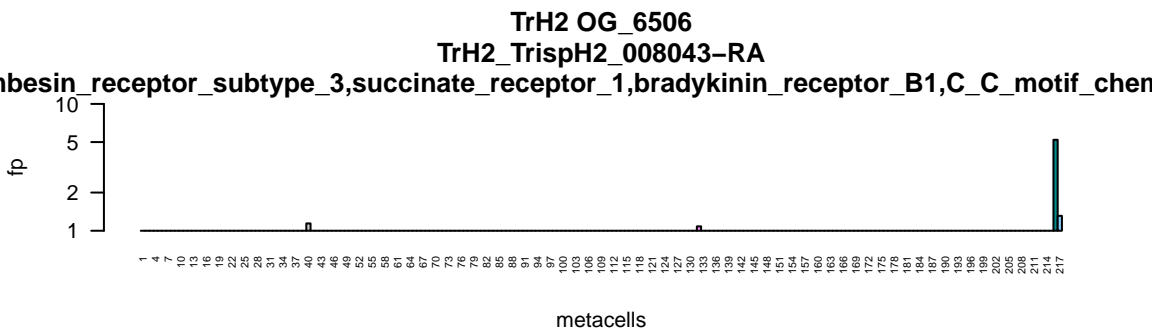
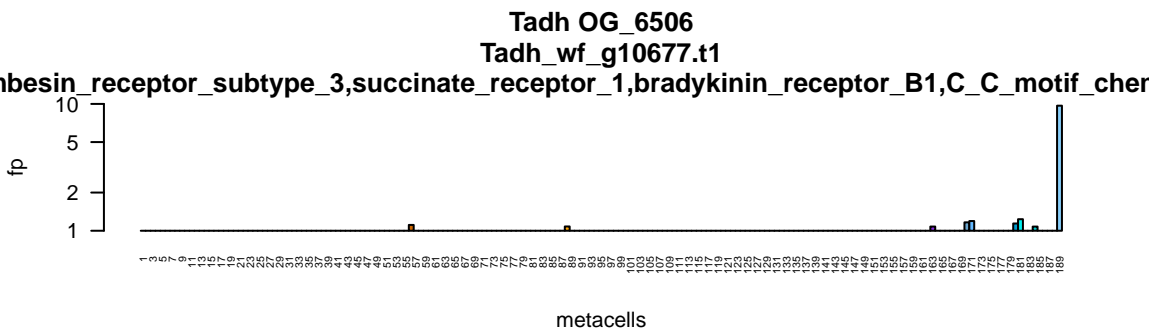
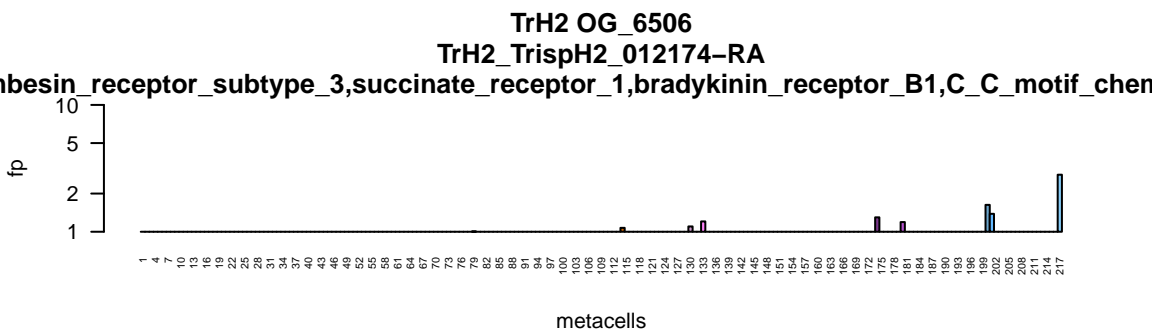
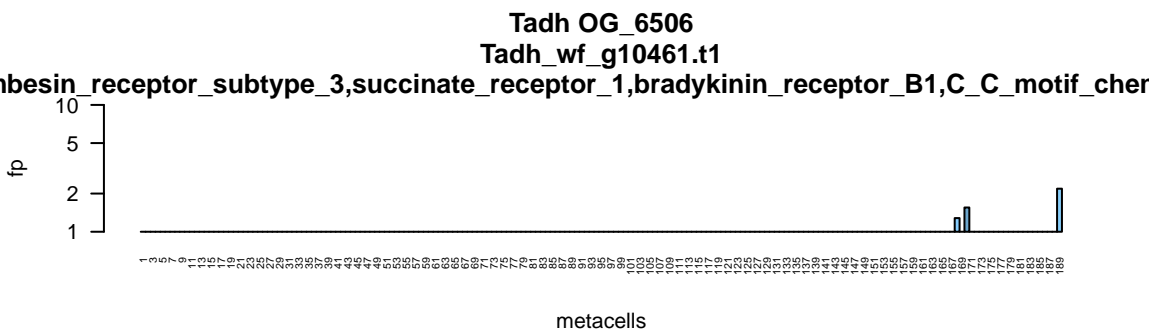
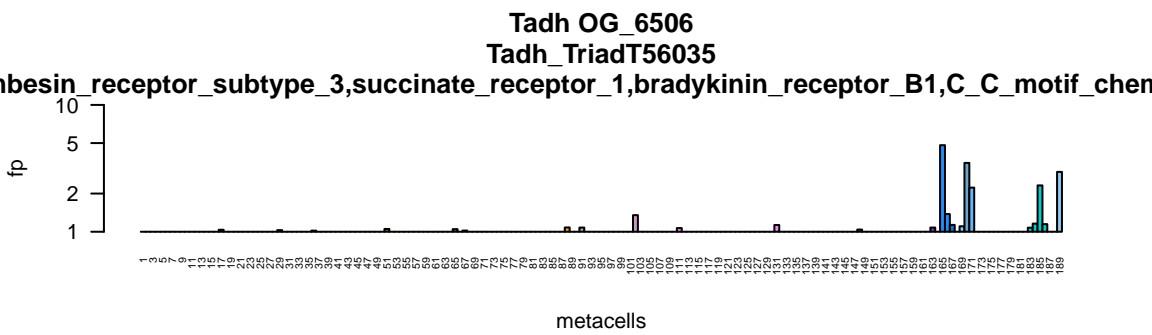
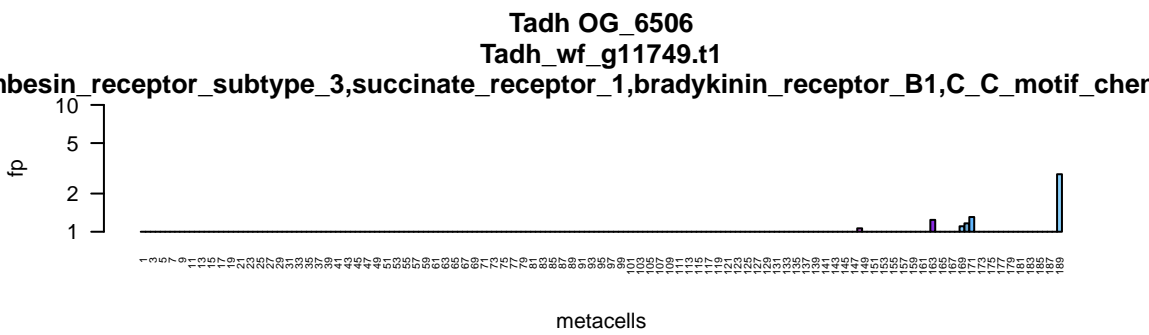
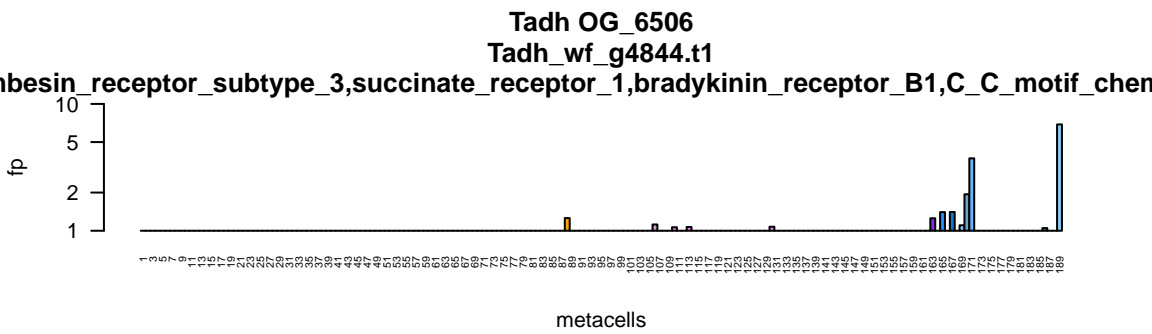
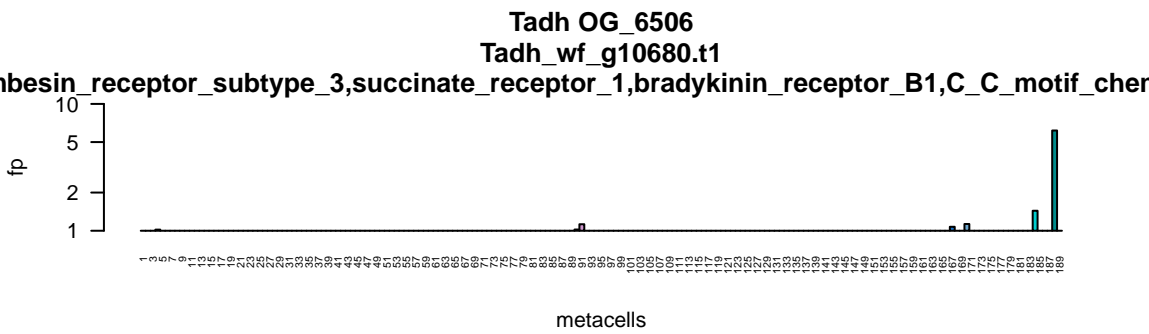
A bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 232. Most metacells have a false positive count of 1, with a few having counts of 2 or 3.

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
52	1
55	1
58	1
64	1
67	1
73	1
76	1
82	1
85	1
88	1
94	1
97	1
103	1
106	1
109	1
115	1
118	1
124	1
127	1
130	1
136	1
139	1
142	1
145	1
151	1
157	1
160	1
163	1
166	1
169	1
172	1
178	1
181	1
184	1
187	1
190	1
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202	1
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211	1
214	2
215	3
220	1
223	1
229	1
232	2

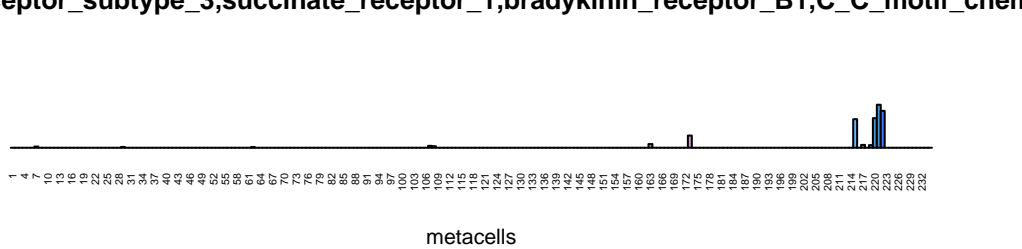
A bar chart showing the frequency of metacells (x-axis) versus the frequency of pairs (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 232. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a few outliers reaching up to 4.

metacells	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
52	1
55	1
58	1
64	1
67	1
73	1
76	1
82	1
85	4
88	1
94	1
97	1
103	1
106	1
109	1
115	1
118	1
124	1
127	1
130	1
136	1
139	1
142	1
145	1
148	1
151	1
157	1
160	1
163	1
166	1
169	1
172	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
220	4
223	1
229	1
232	1

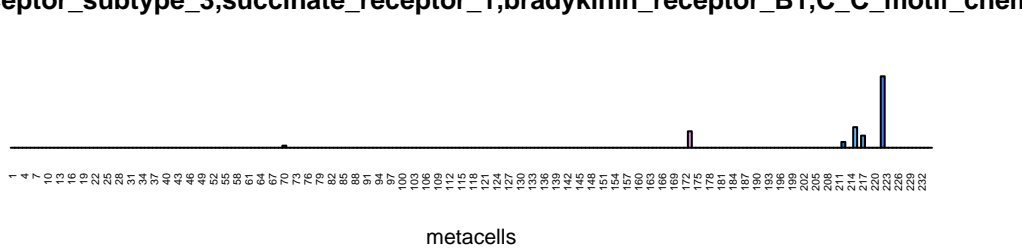


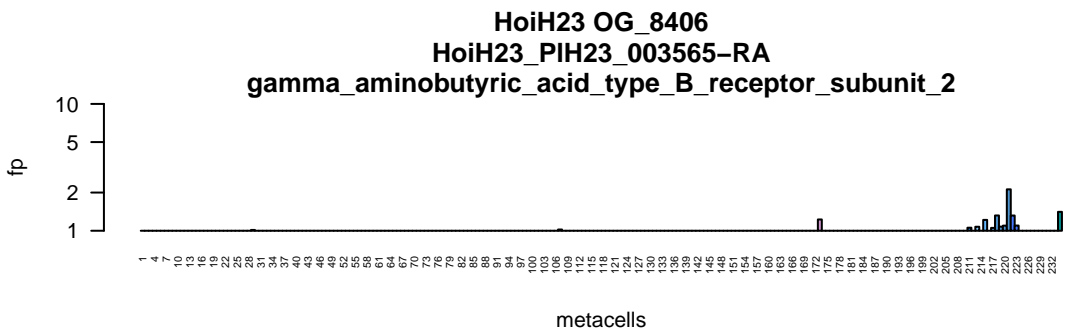
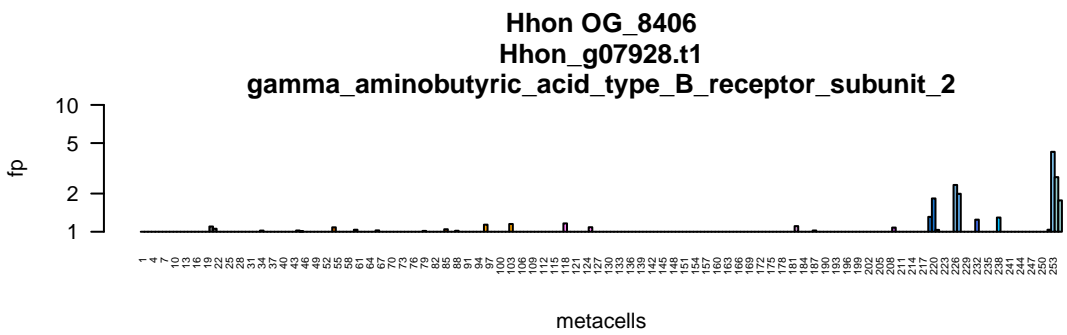
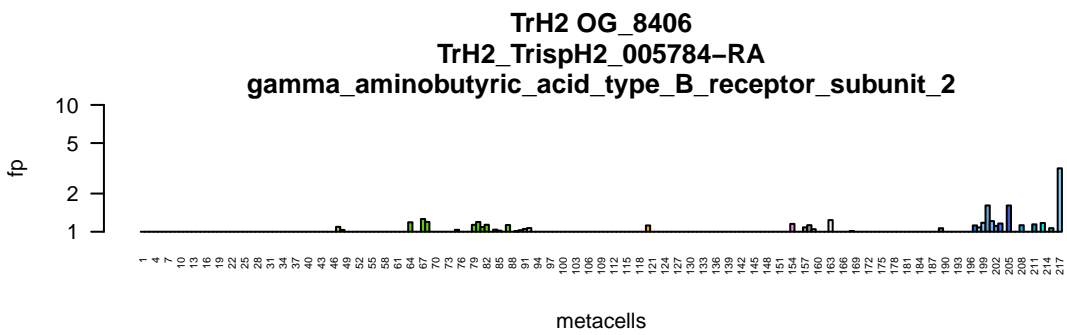
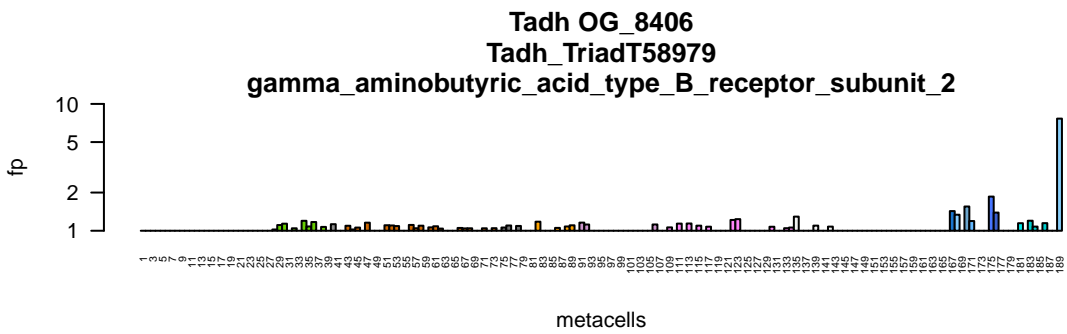


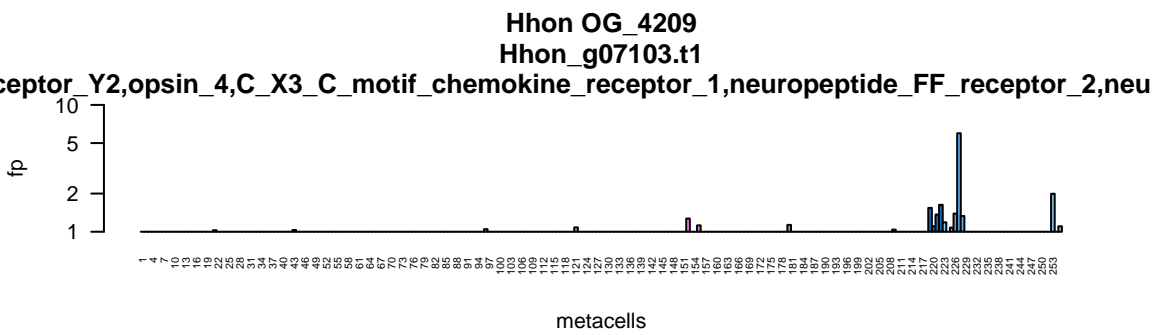
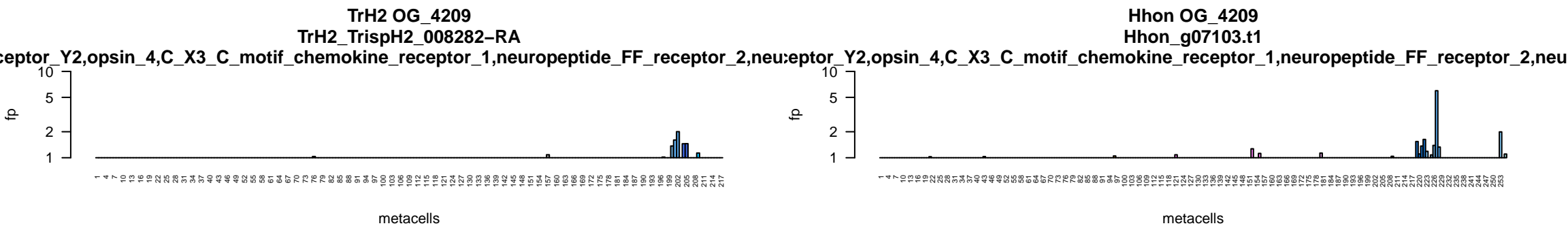
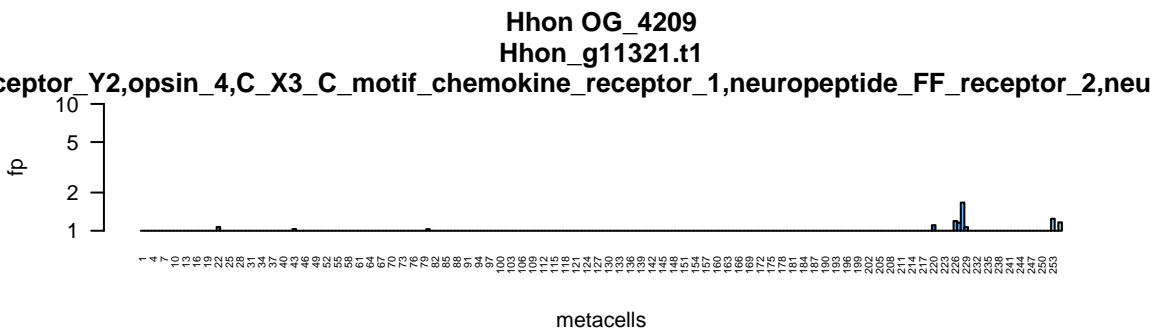
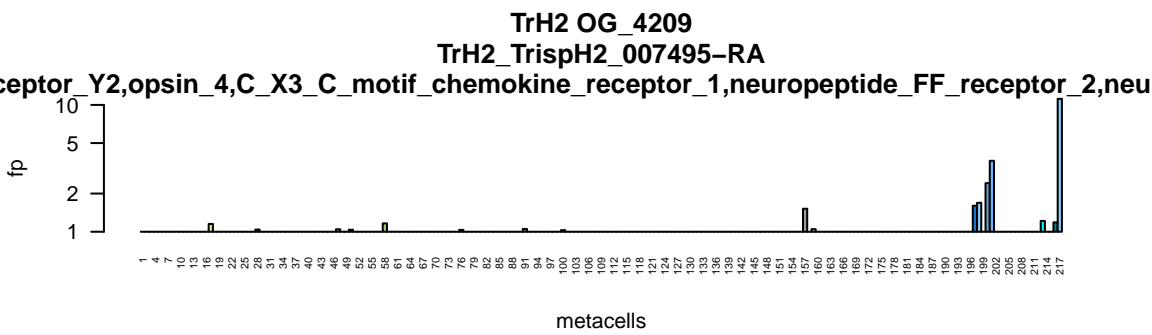
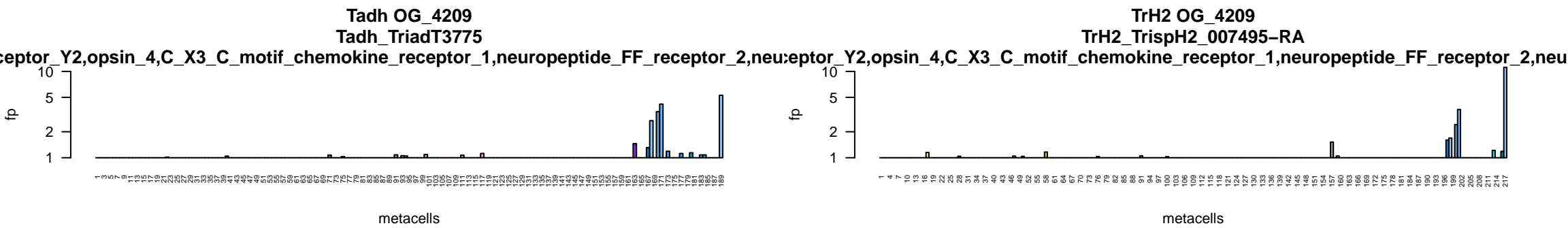
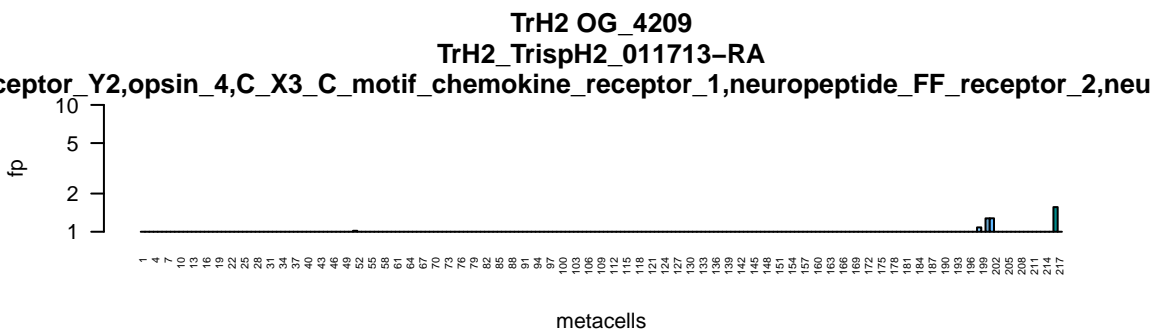
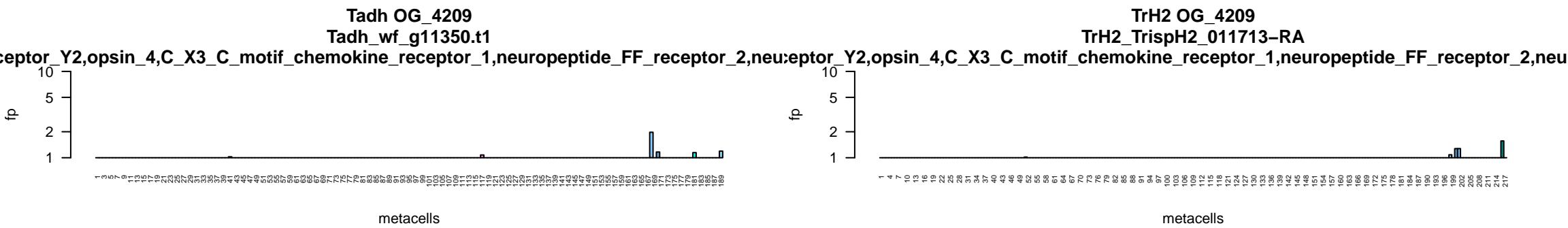
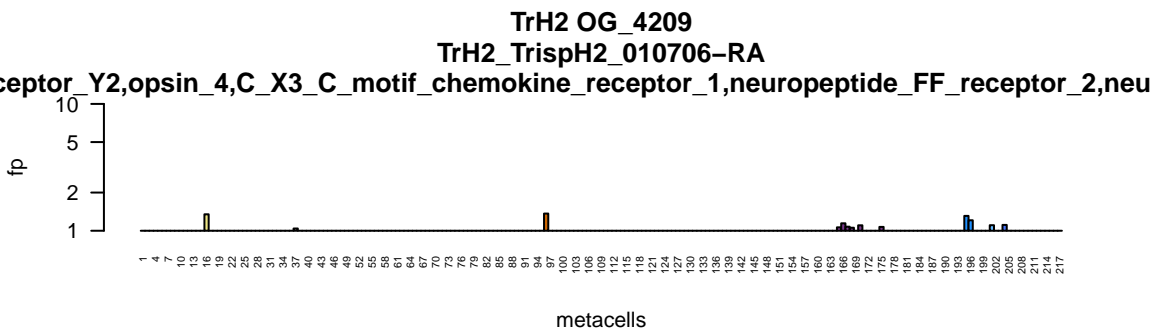
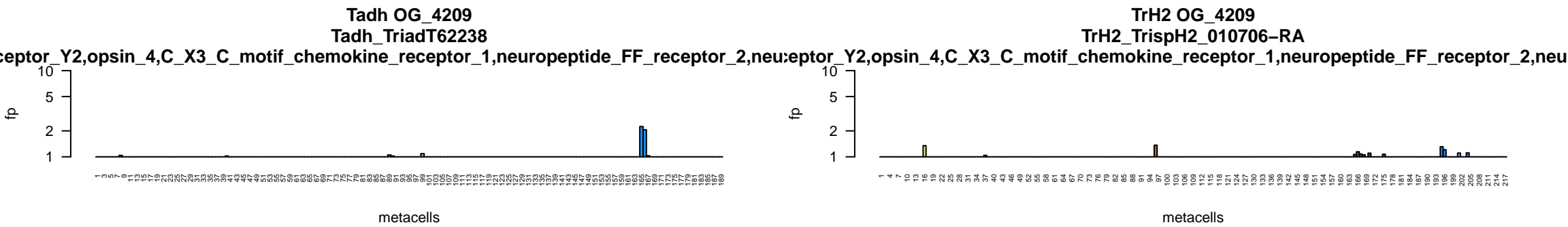
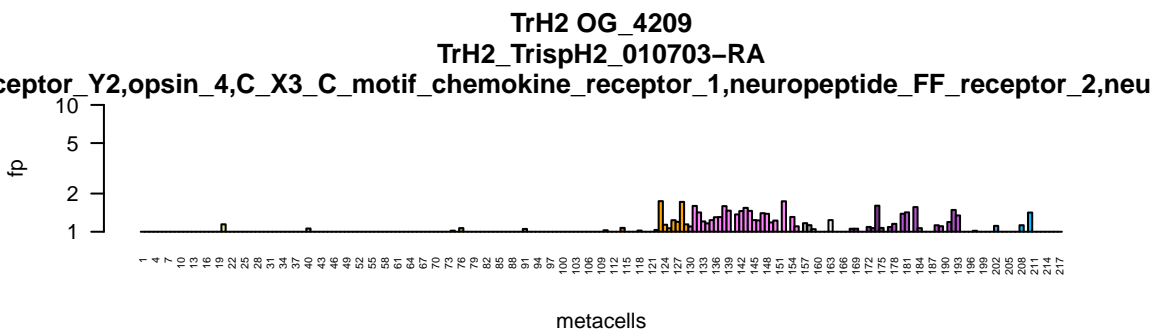
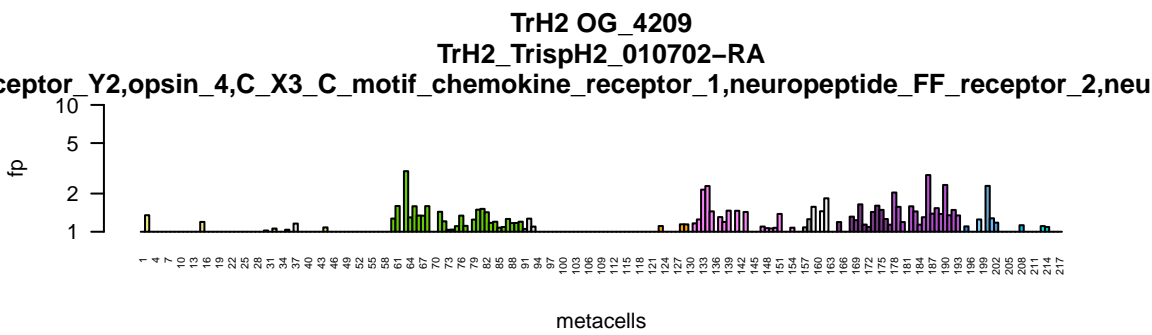
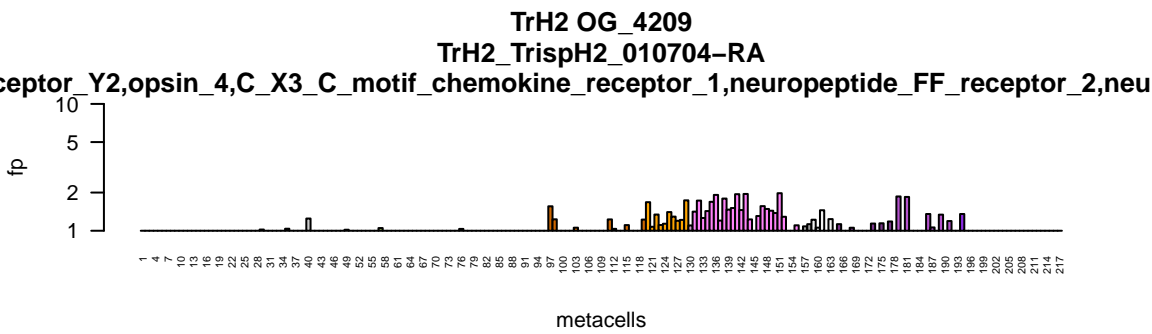
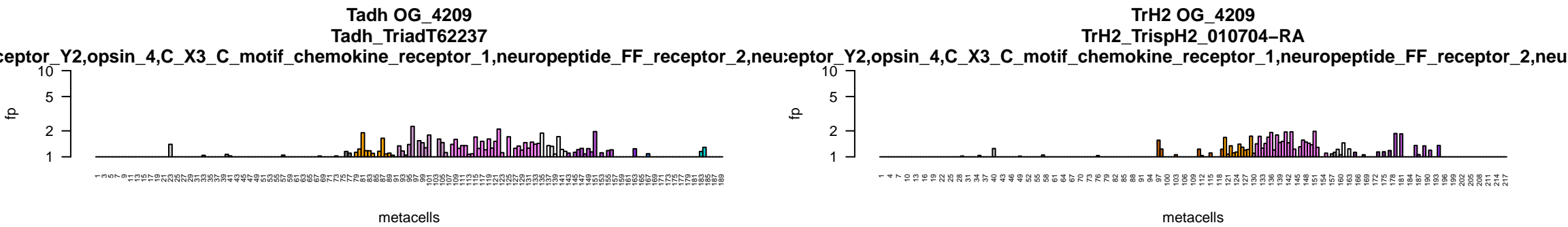
HoiH23 OG_6506
HoiH23_PIH23_008211-RA

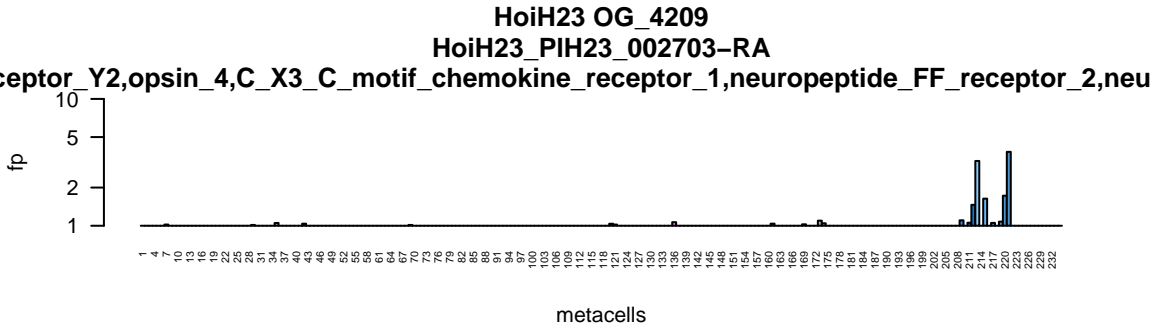
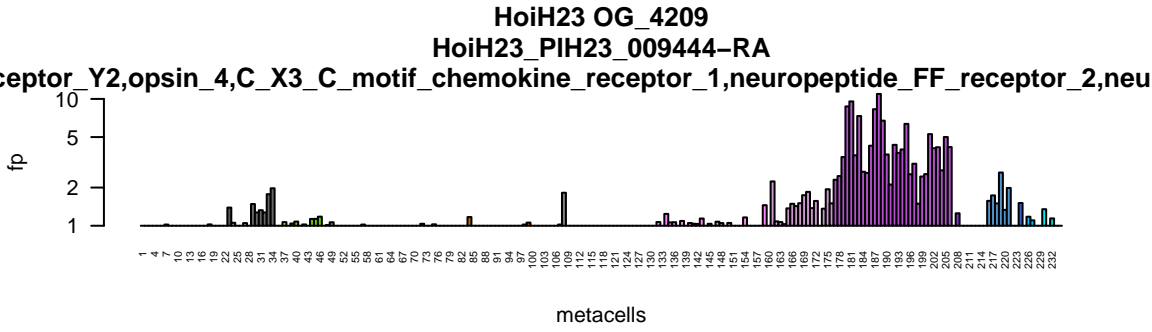
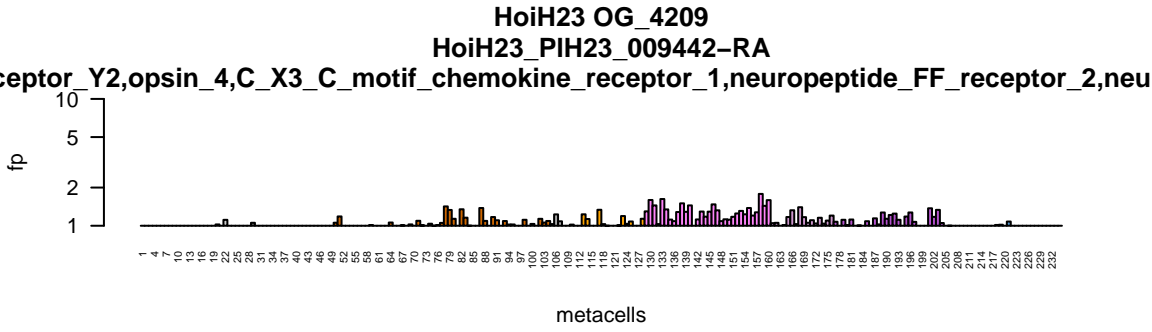
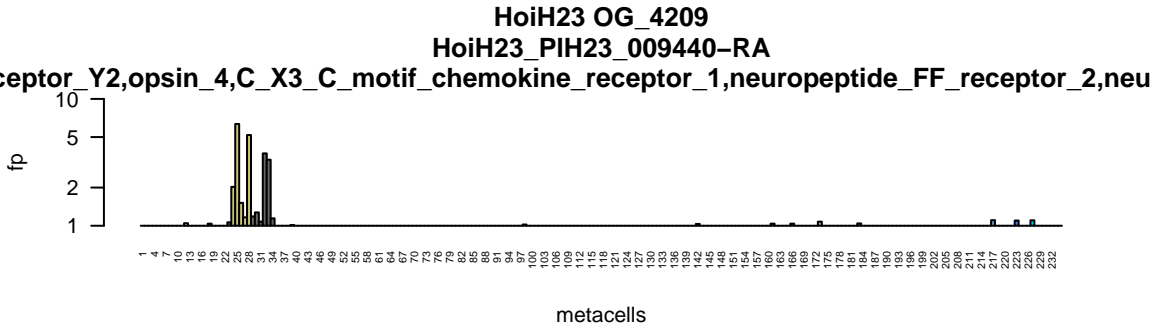
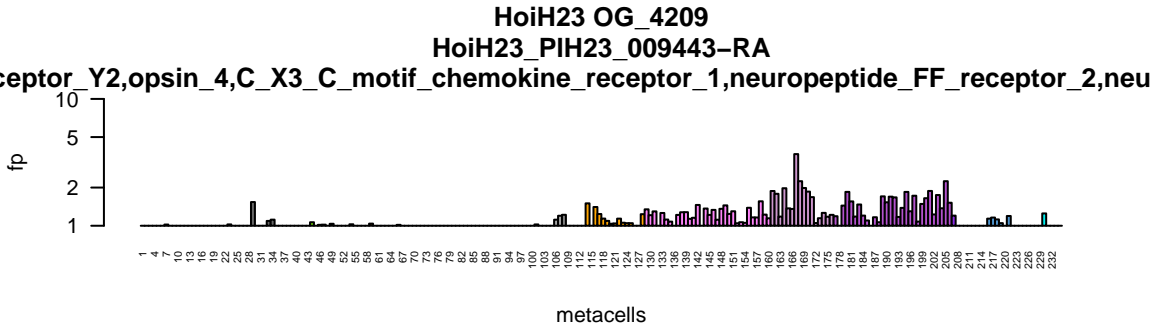
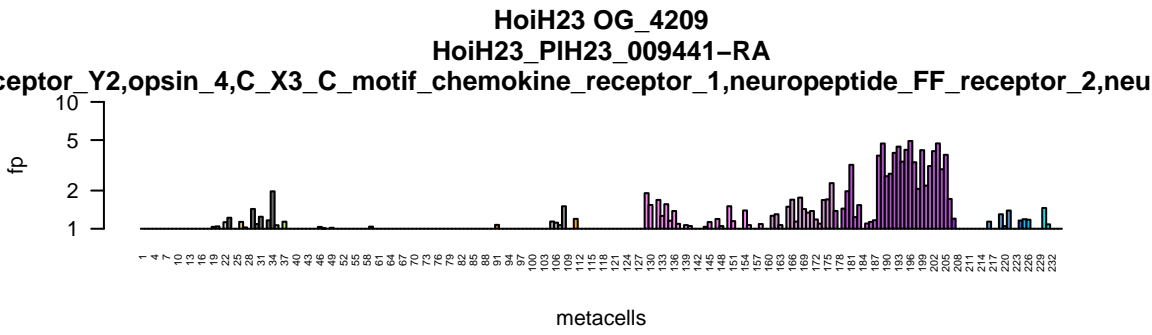
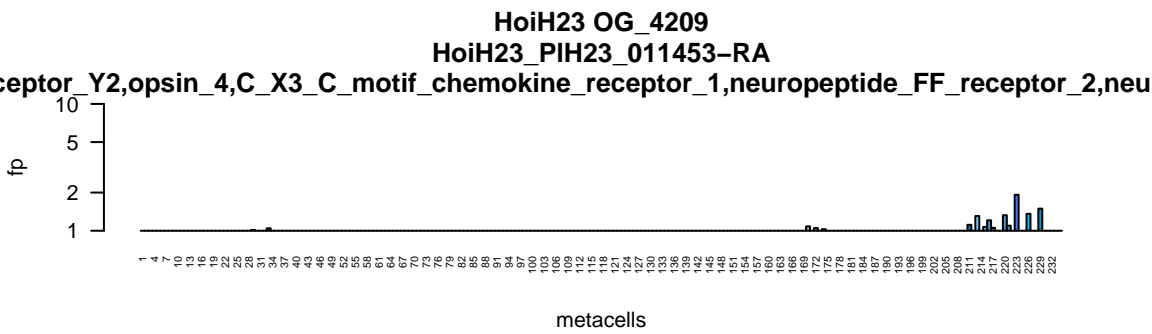


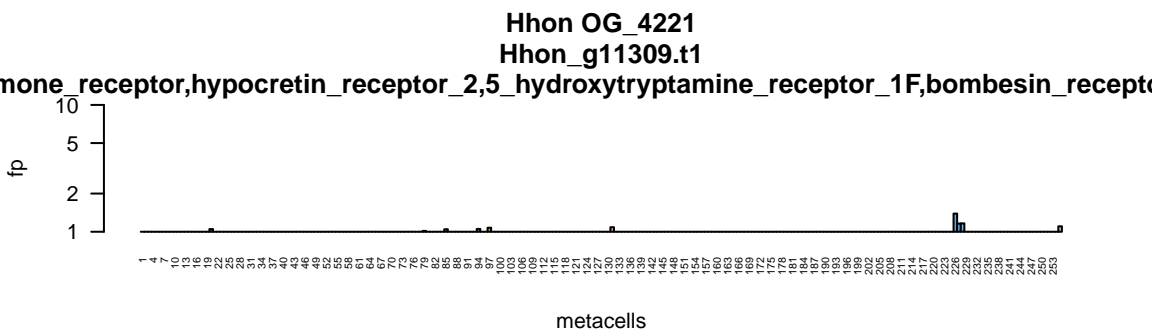
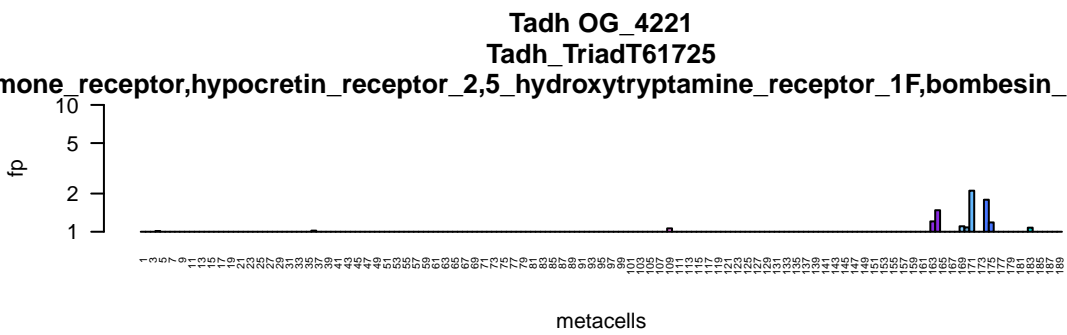
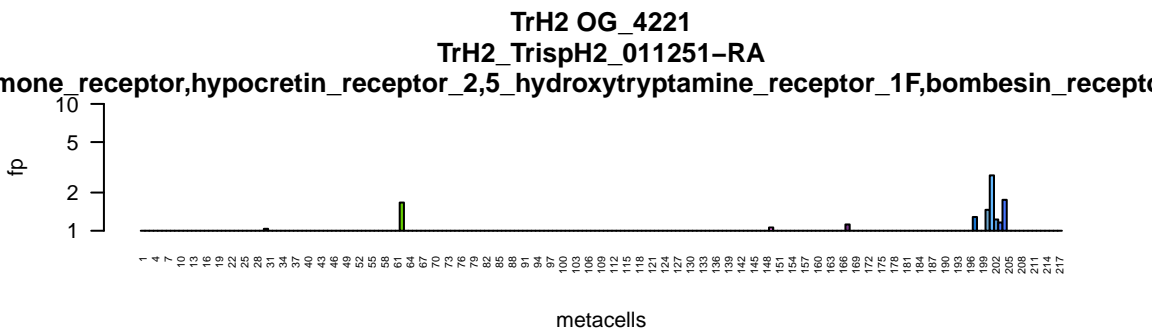
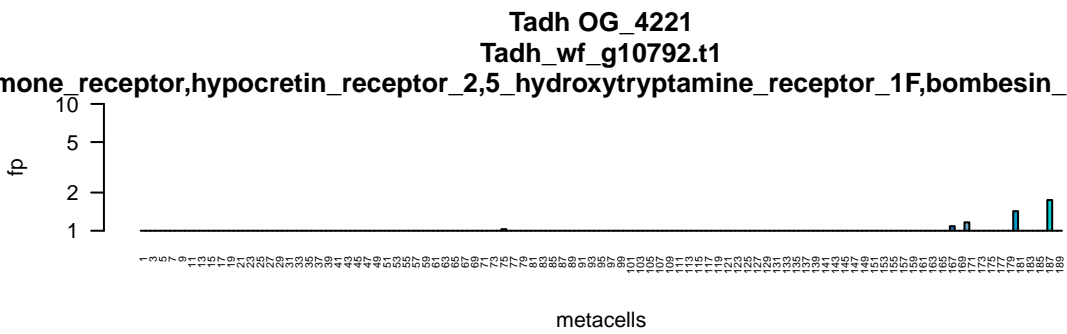
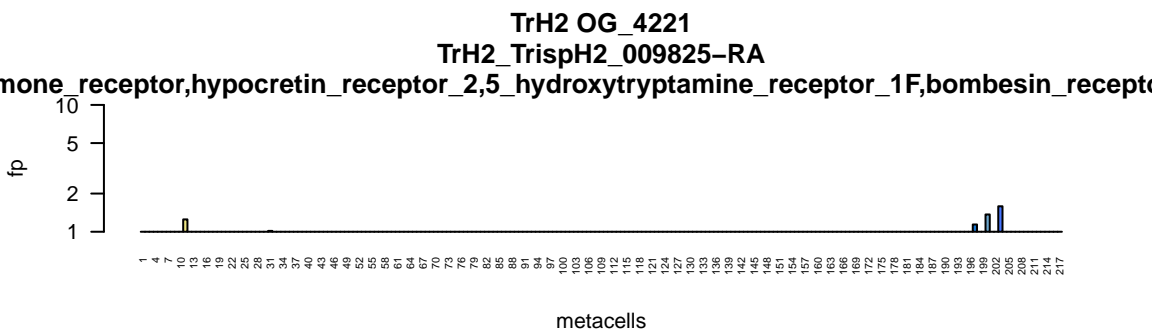
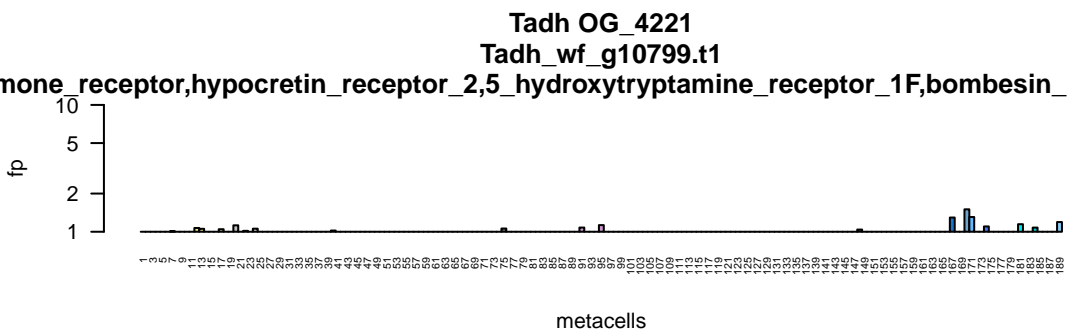
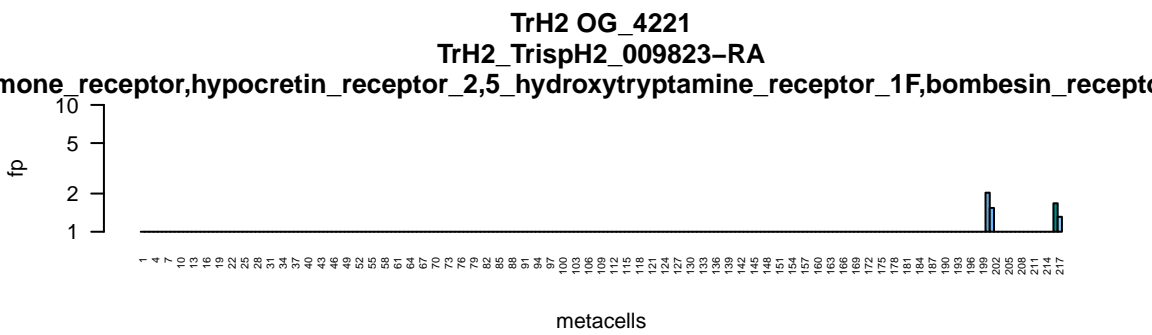
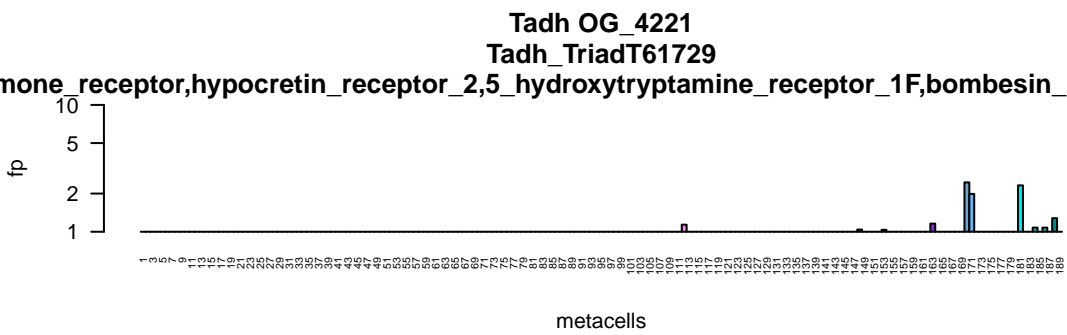
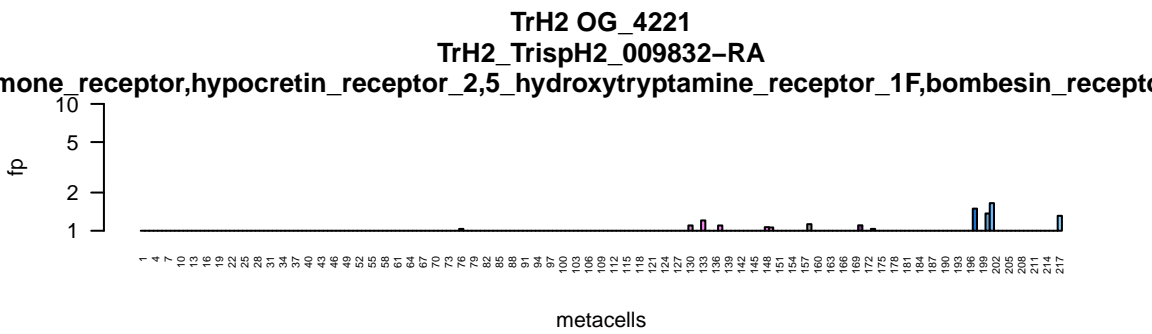
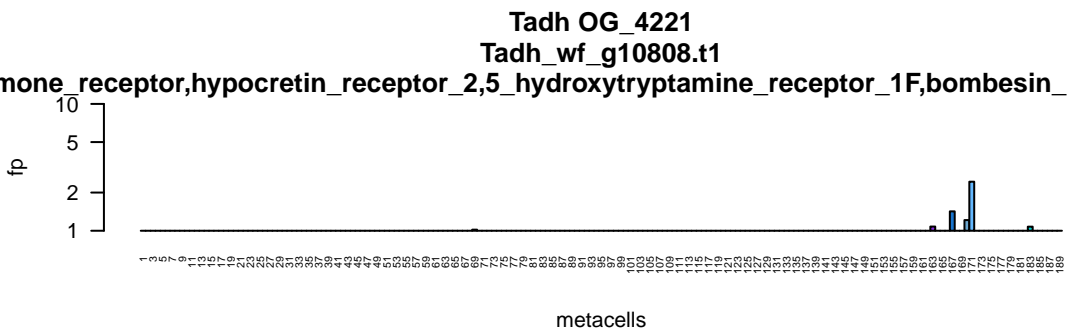
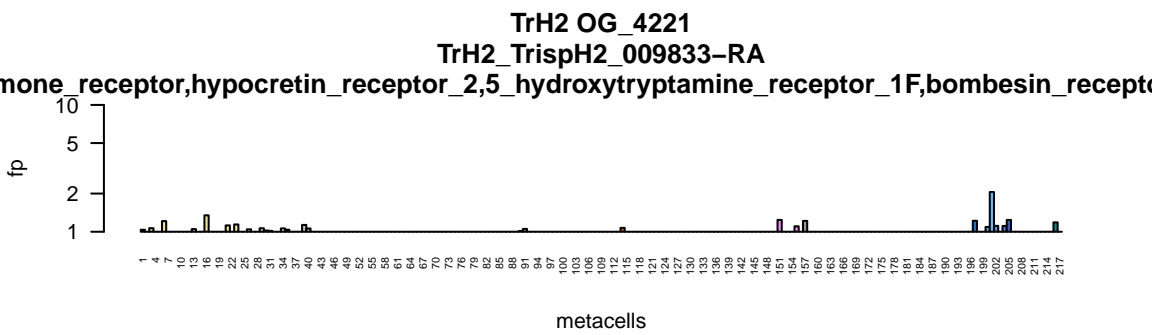
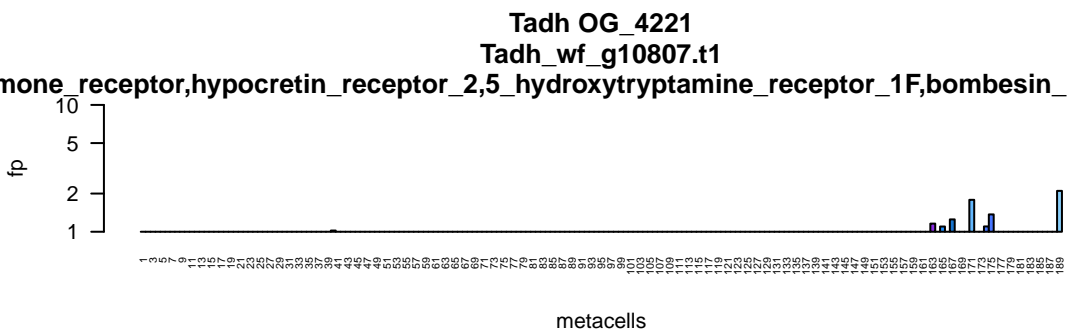
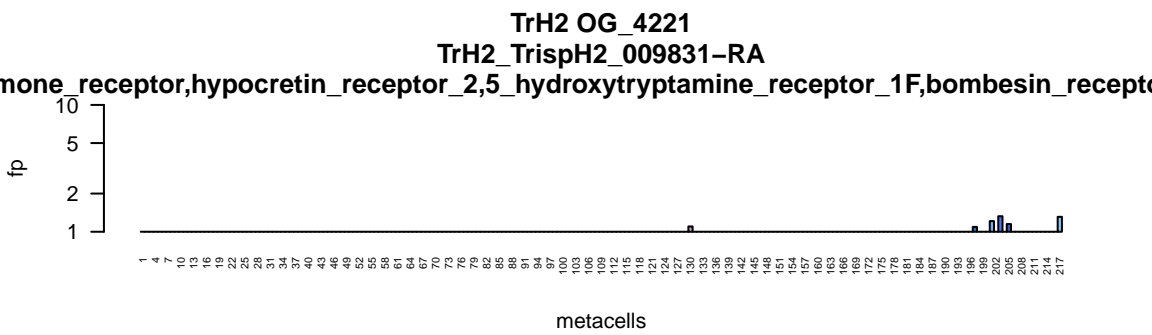
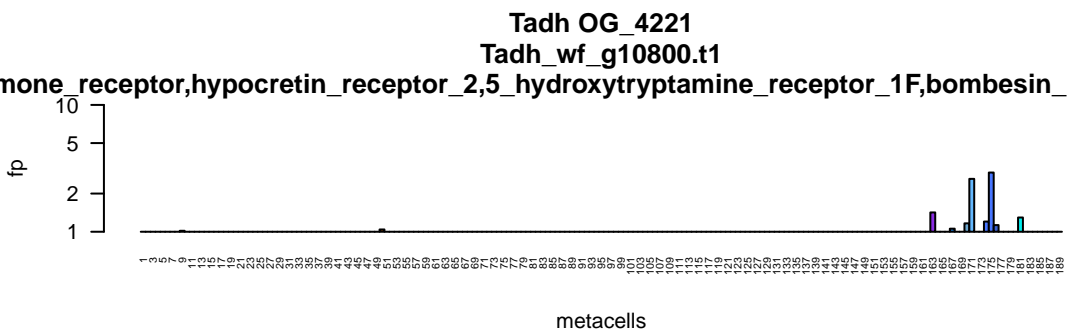
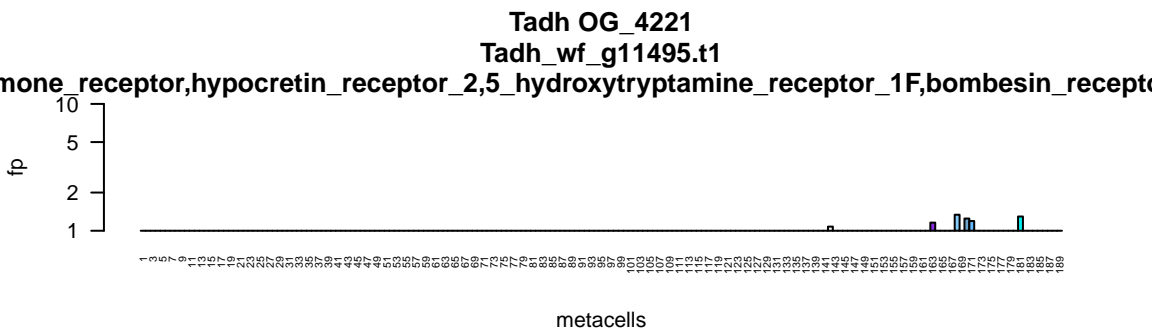
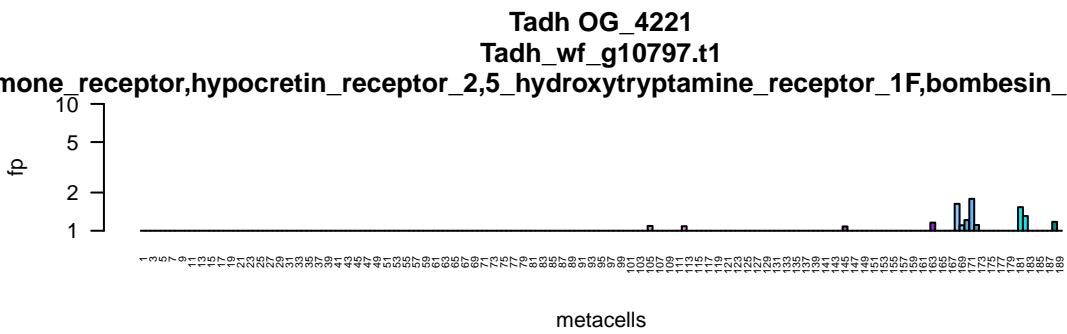
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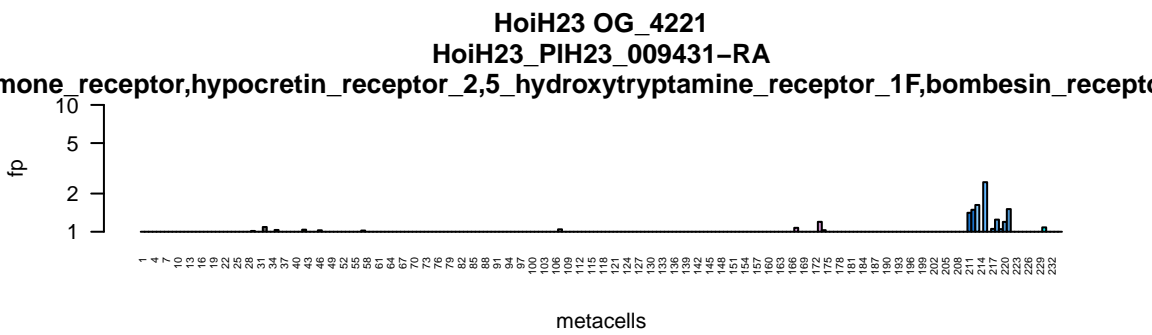
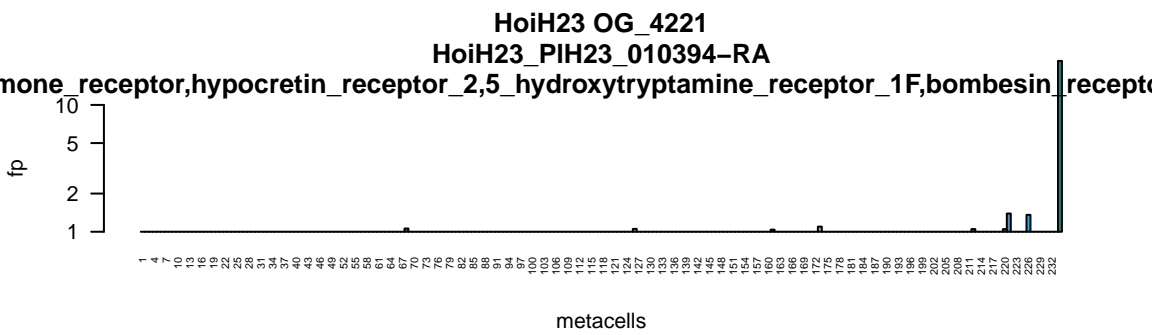
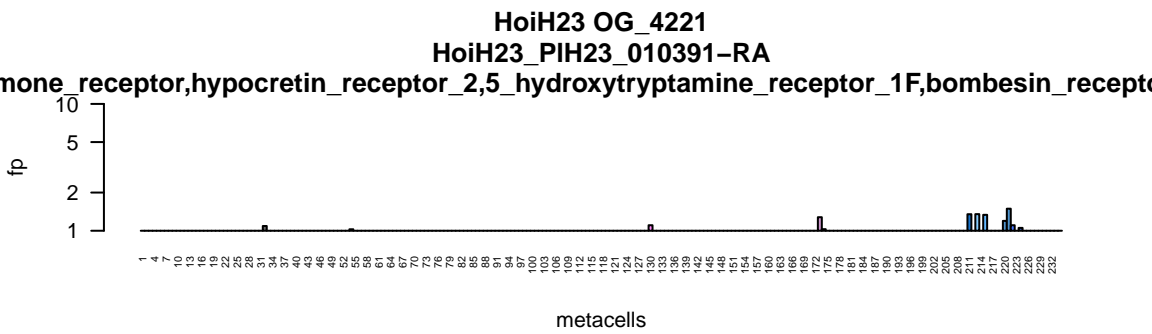
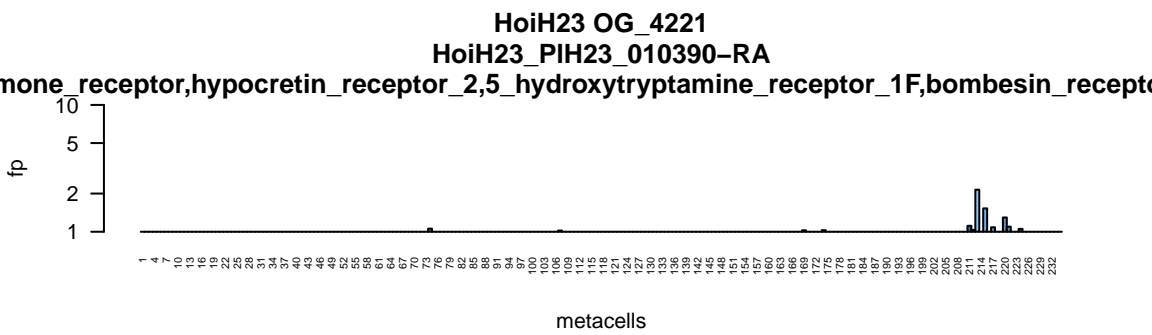
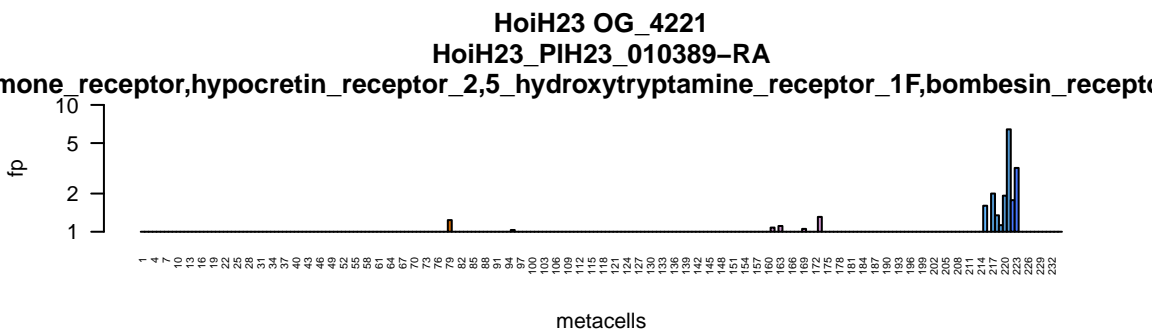
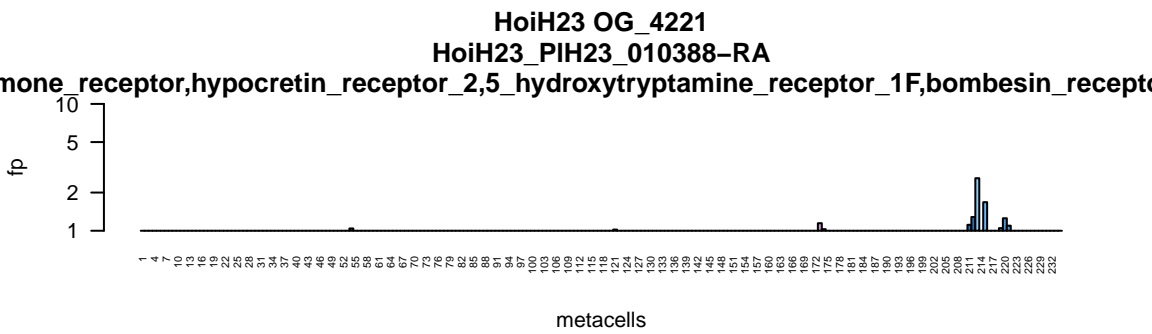


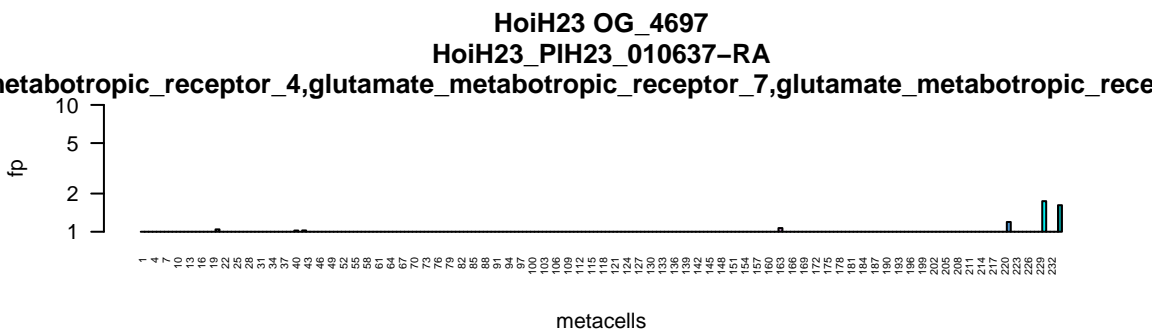
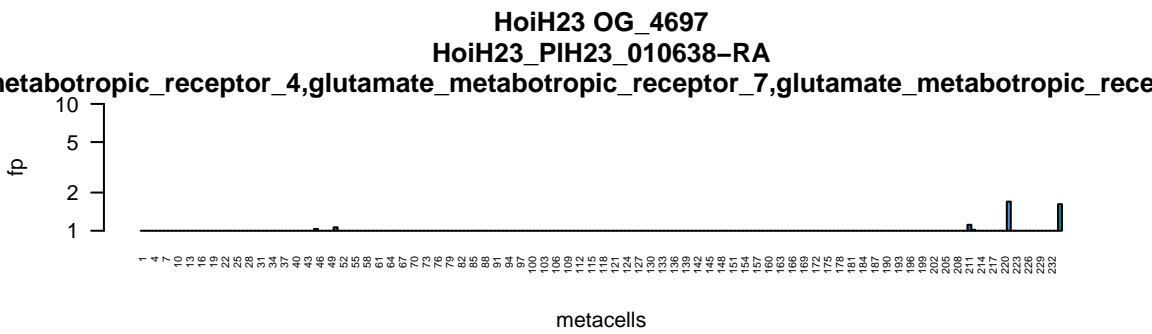
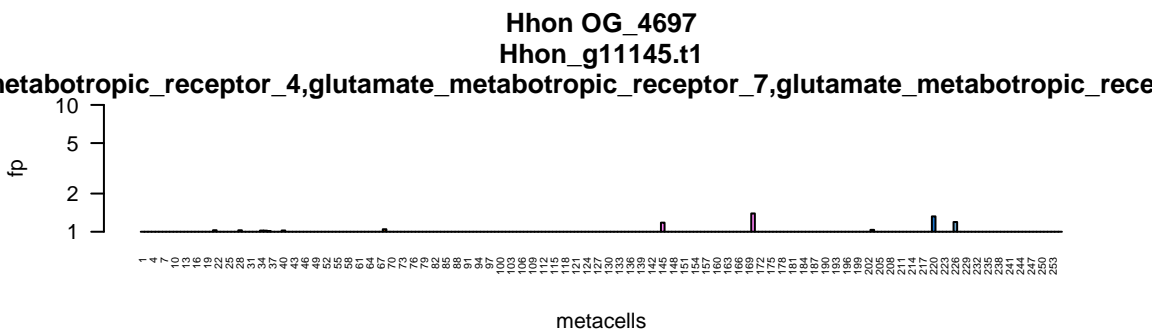
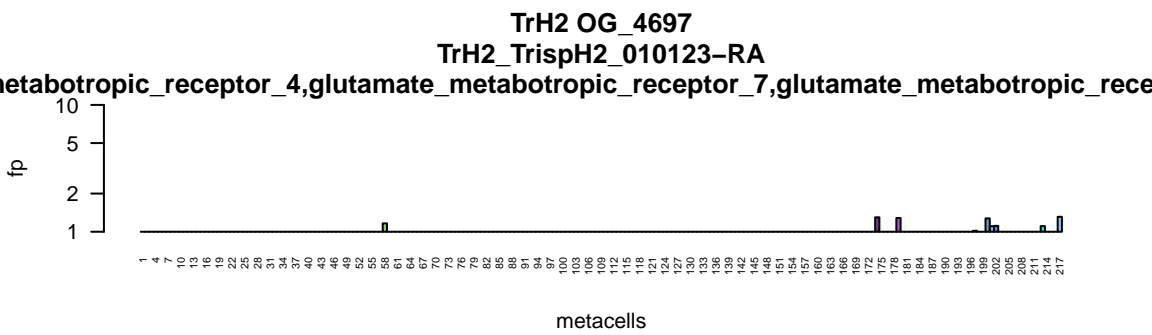
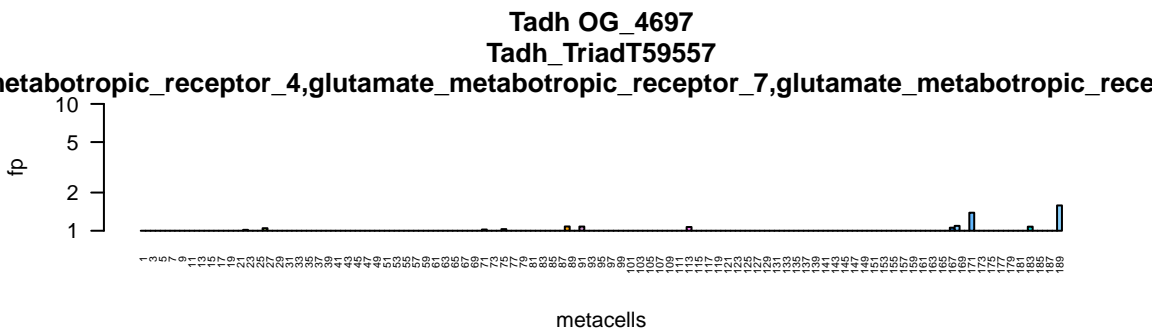


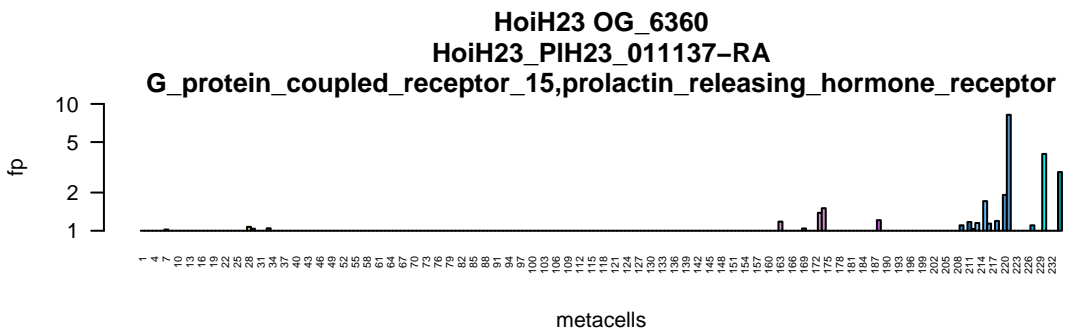
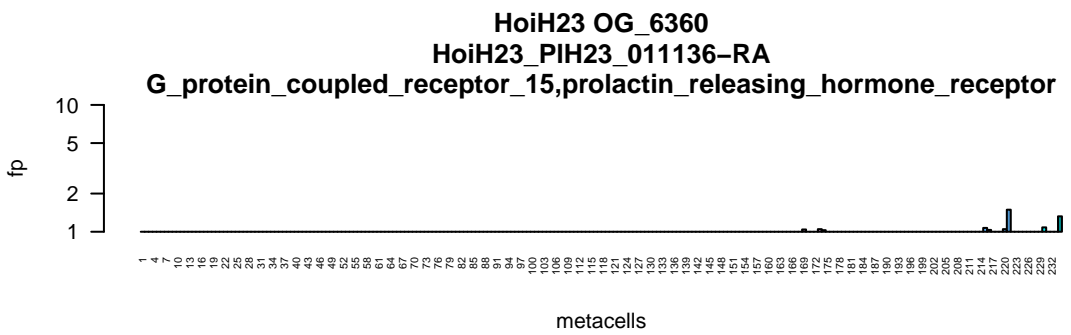
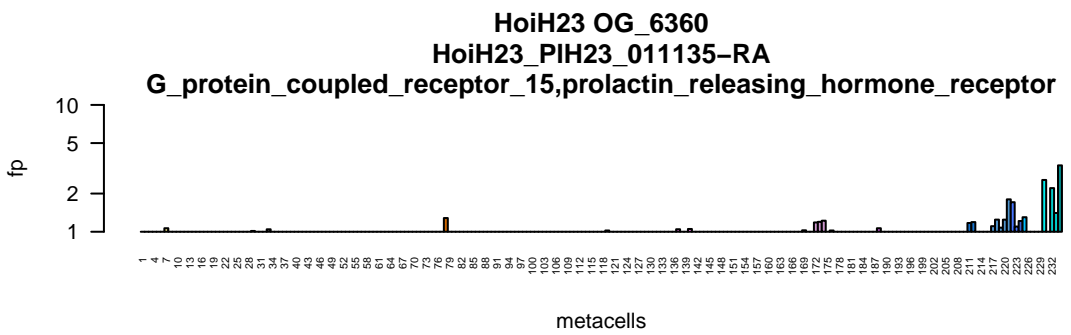
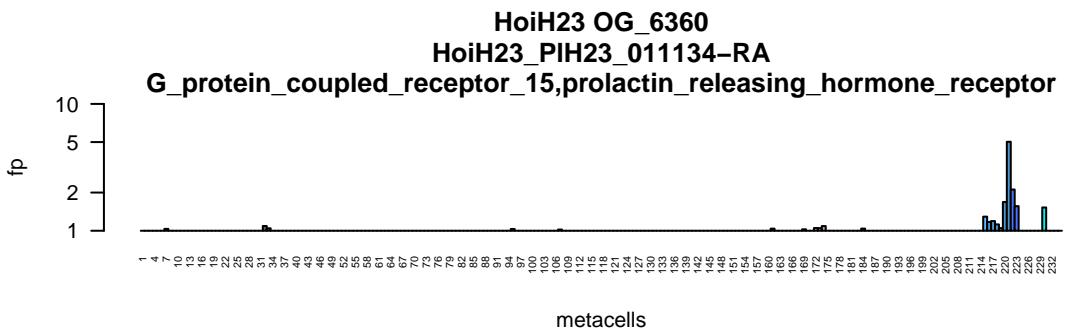
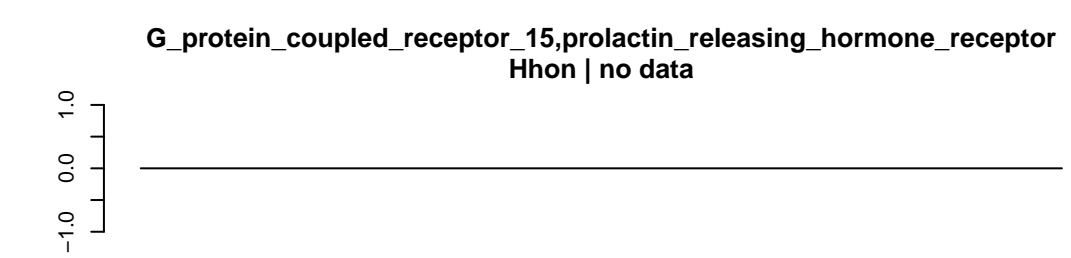
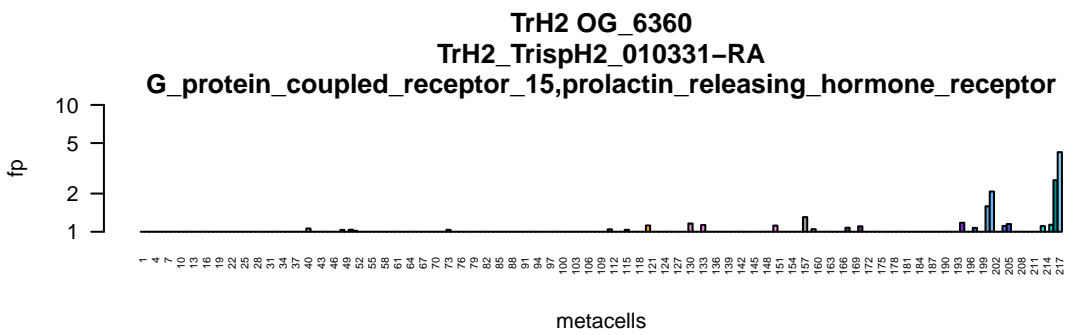
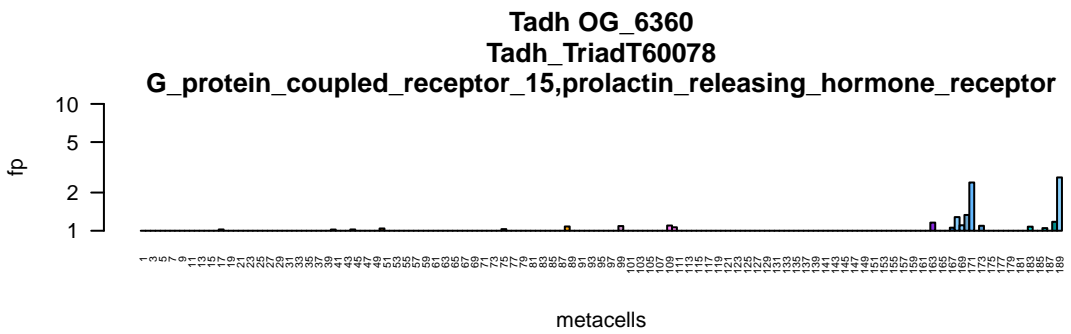


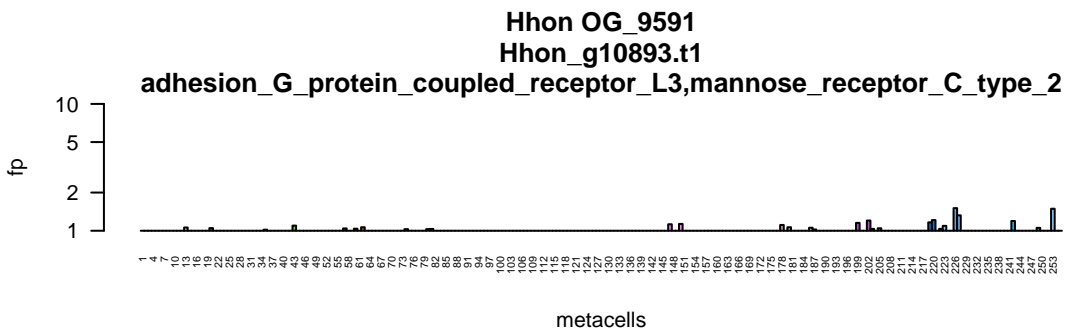
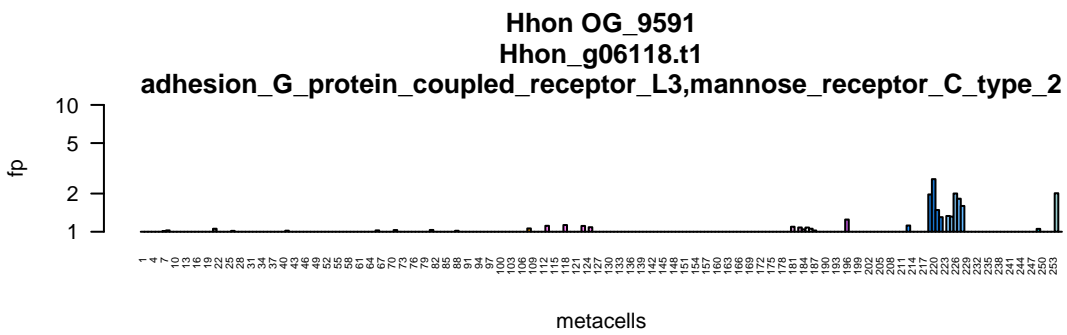
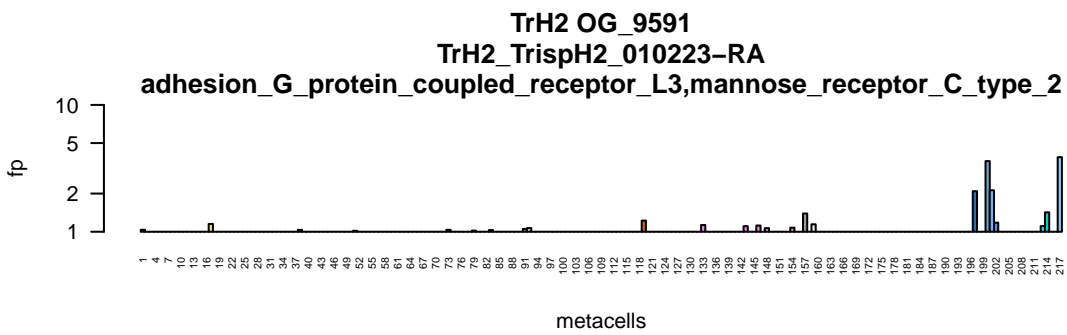
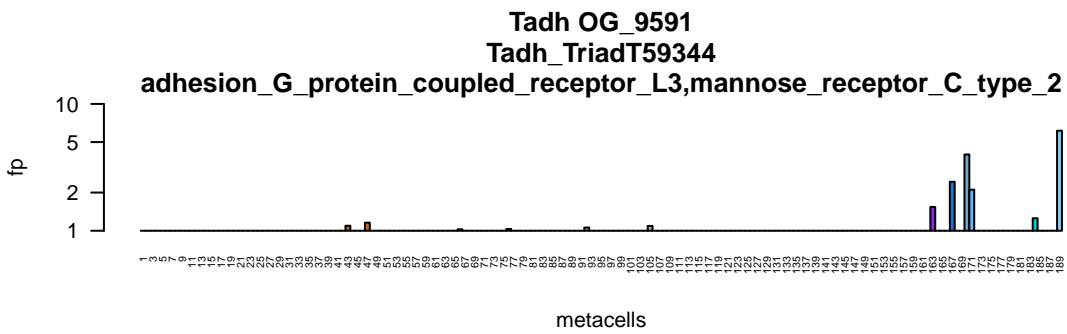


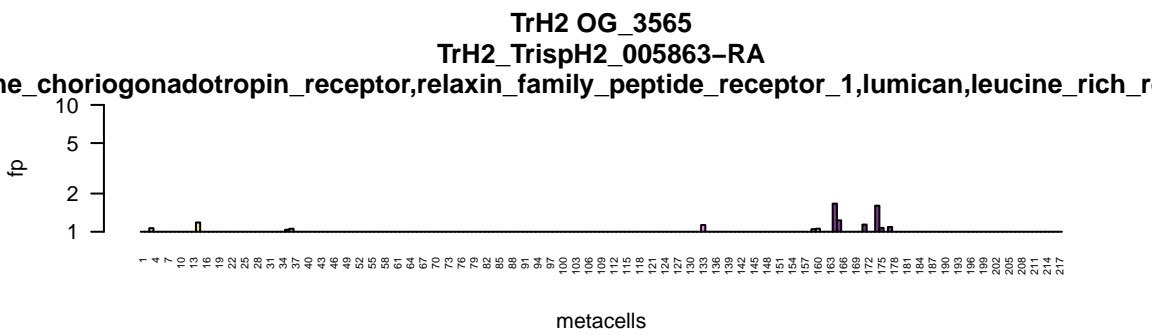
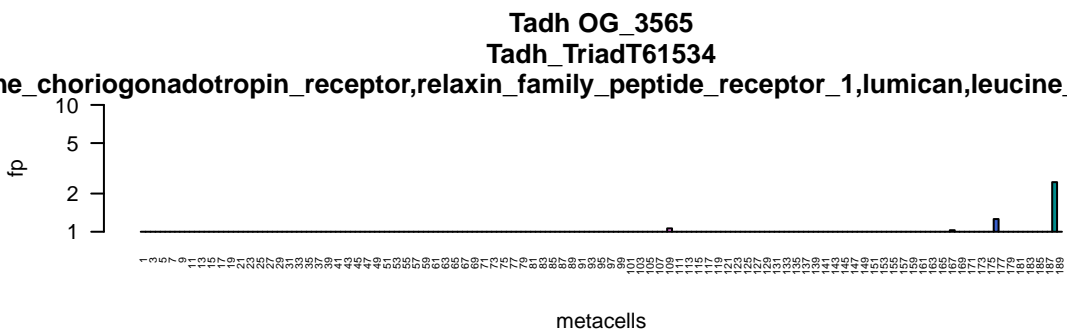
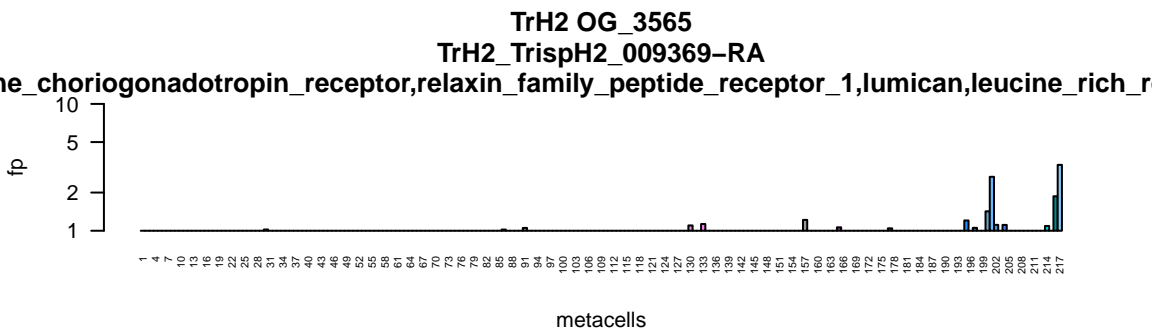
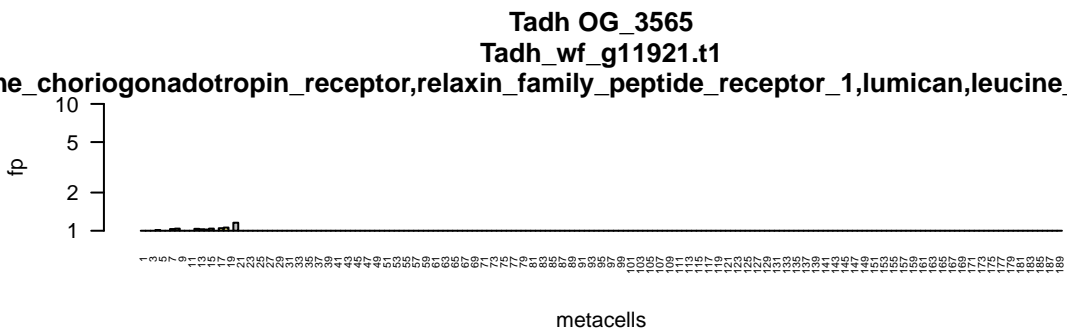
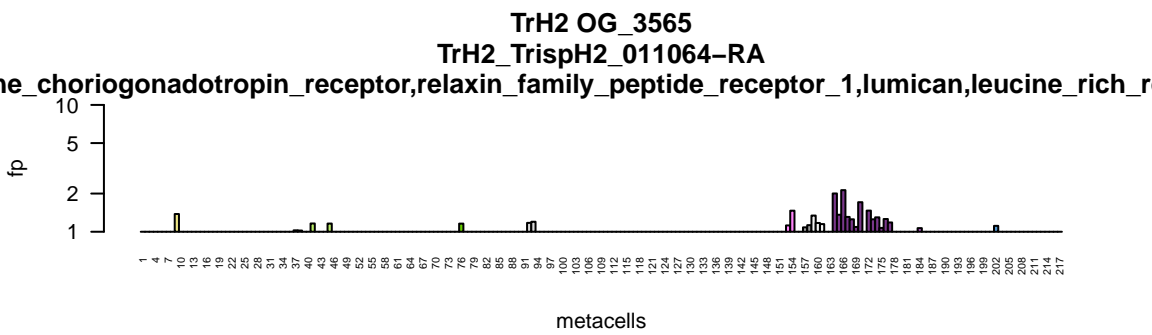
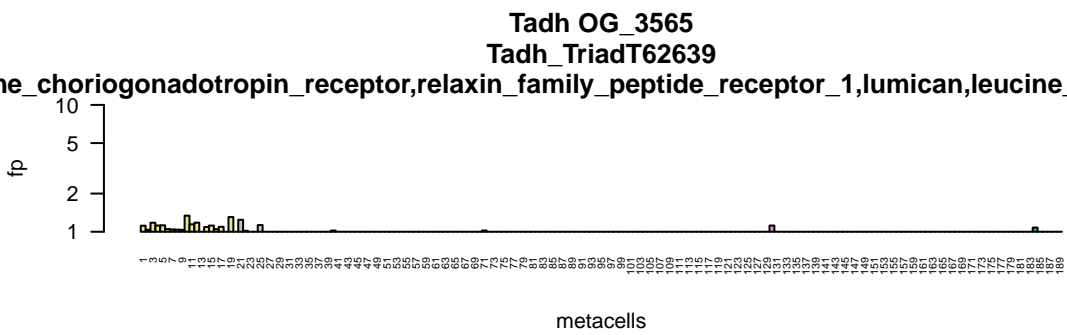
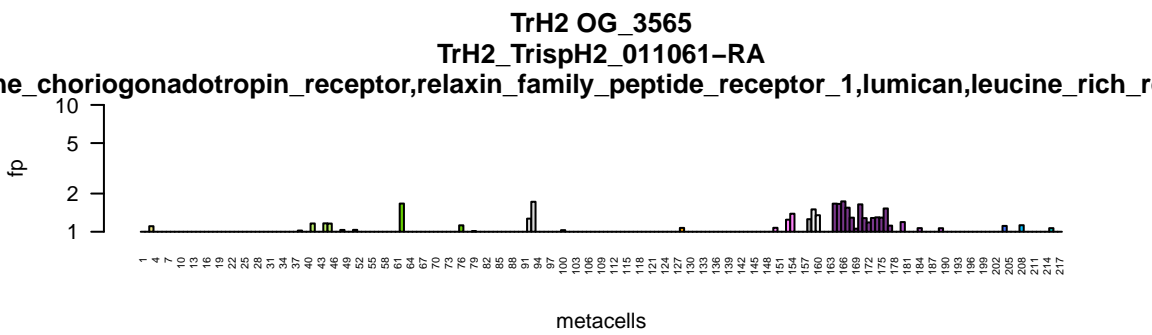
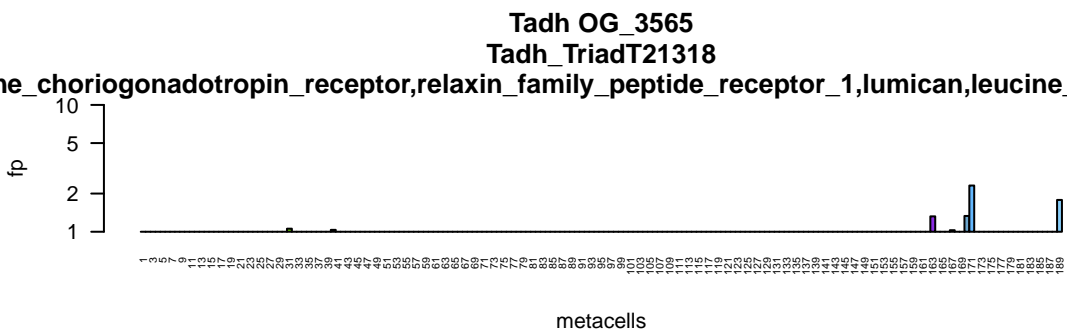
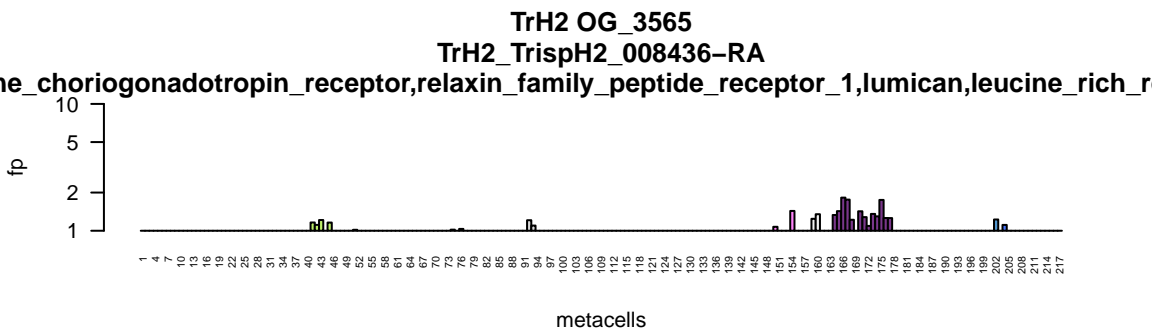
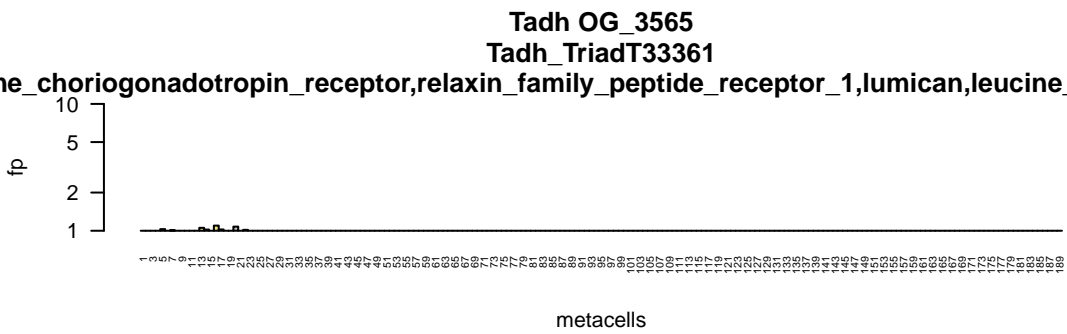
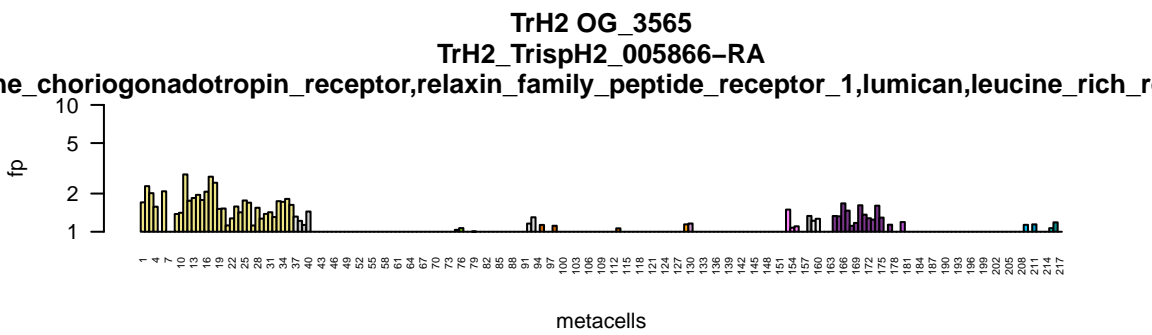
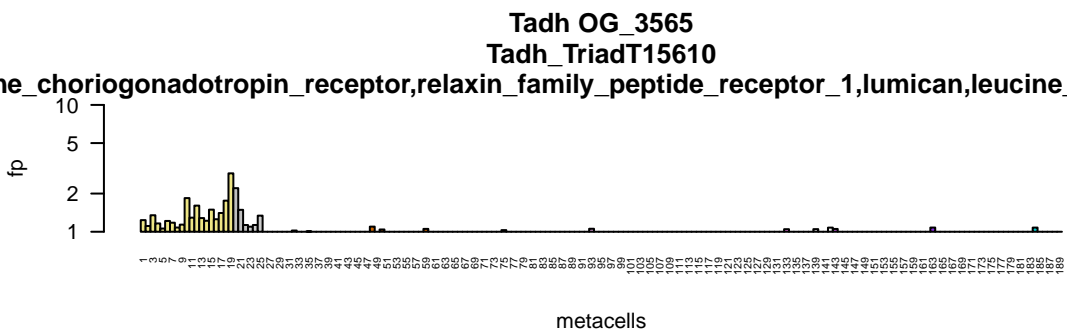
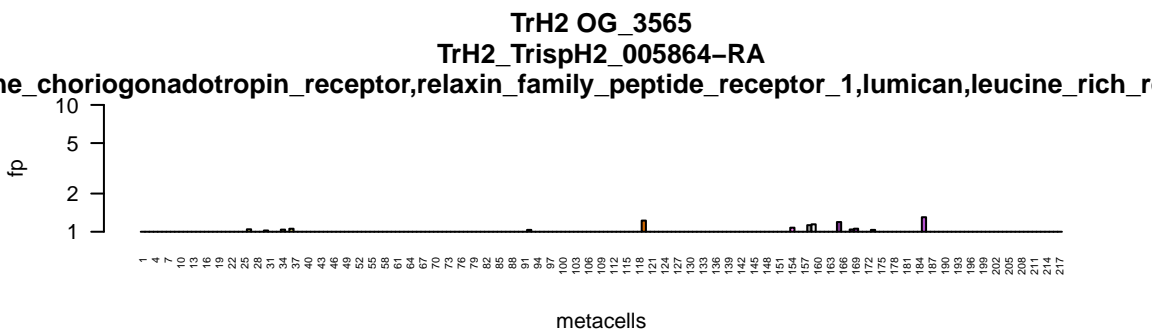
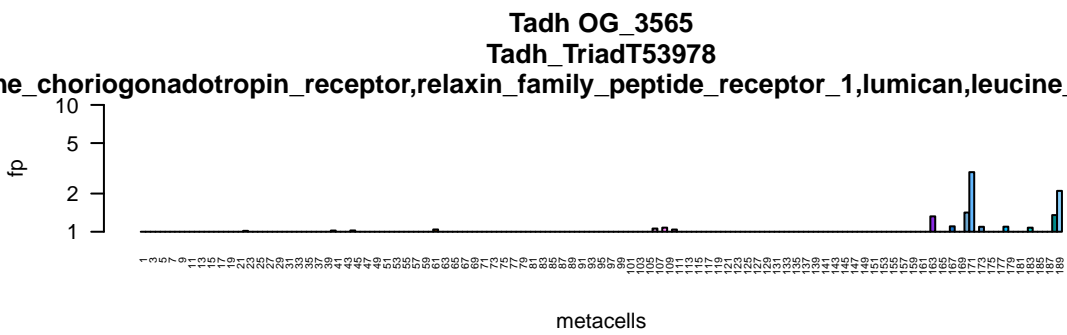
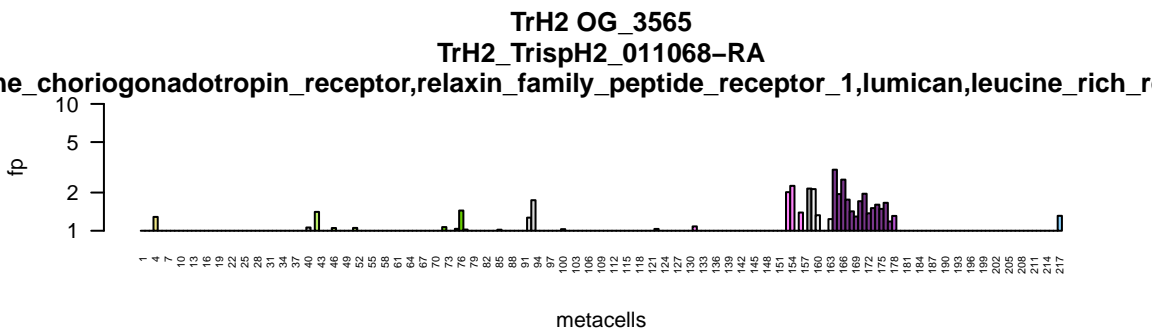
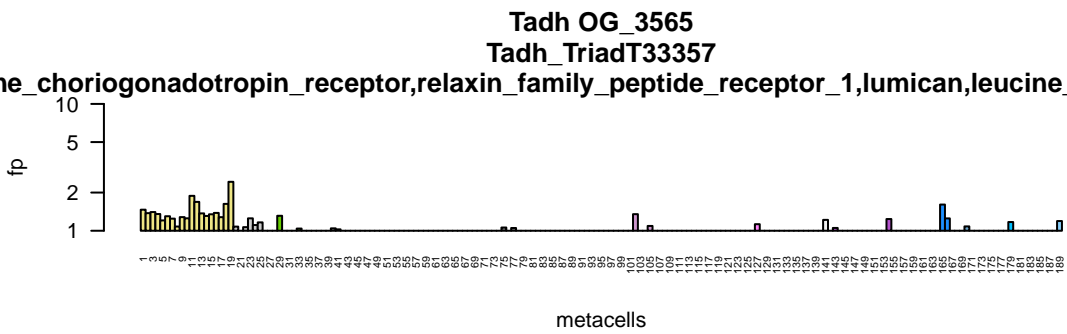


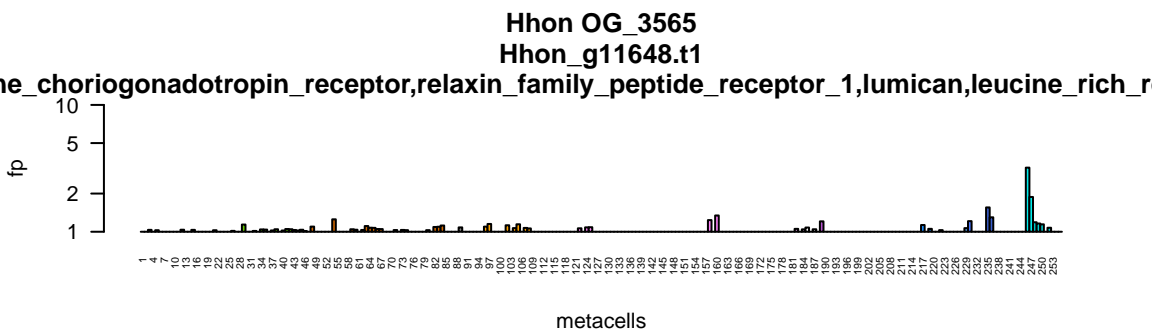
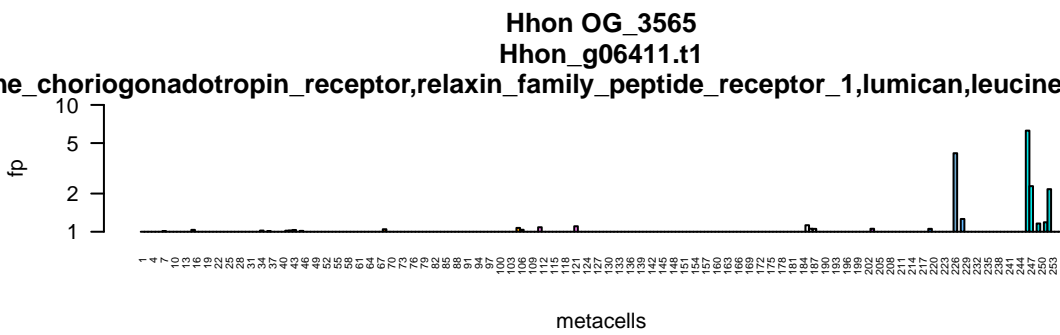
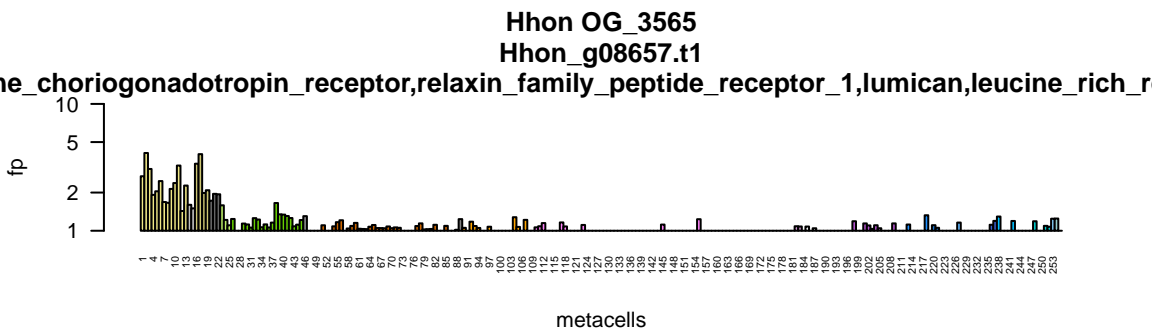
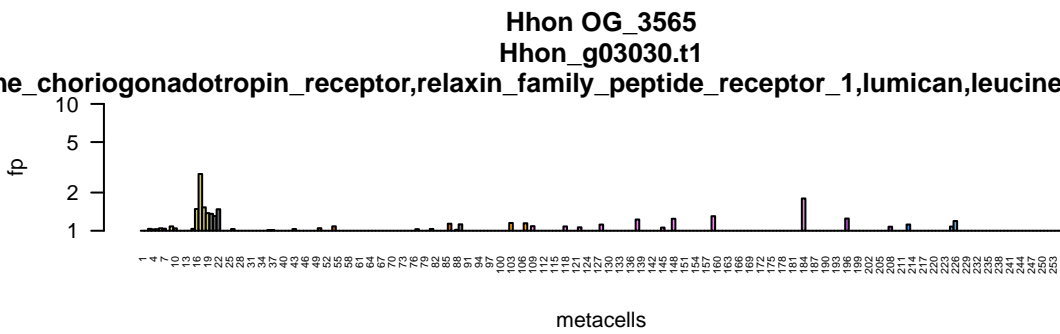
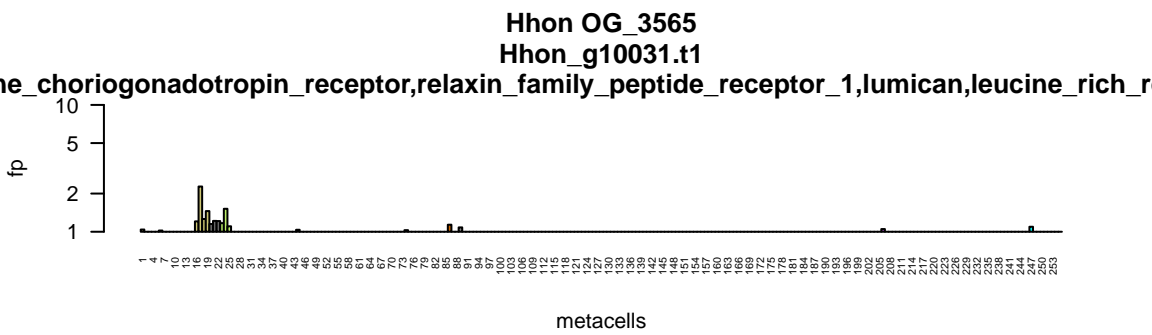
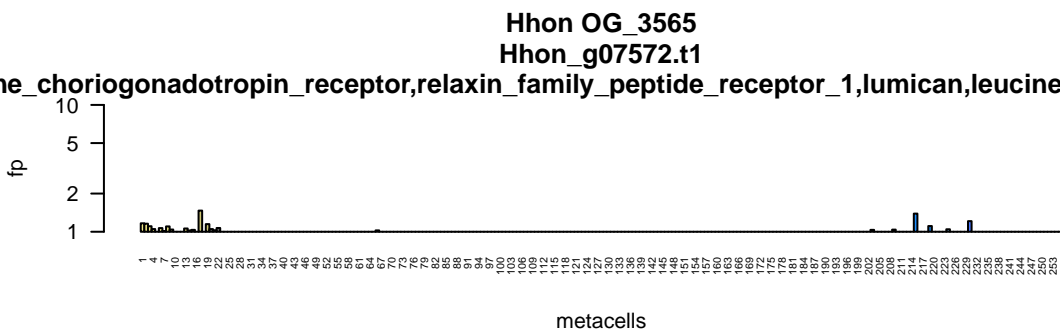
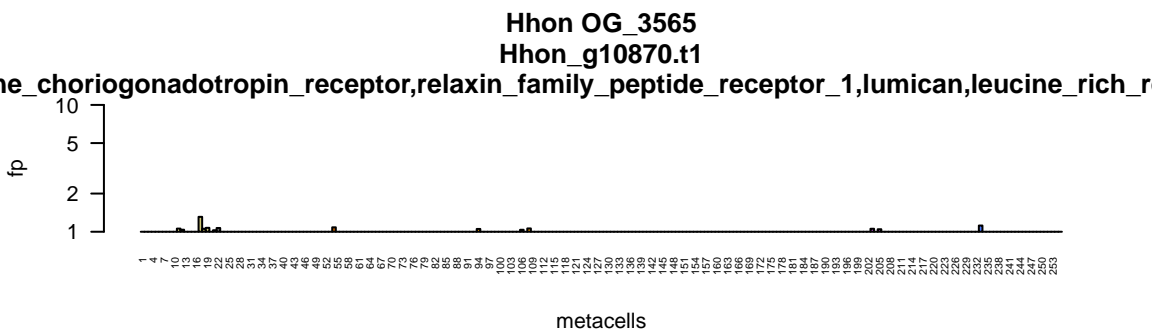
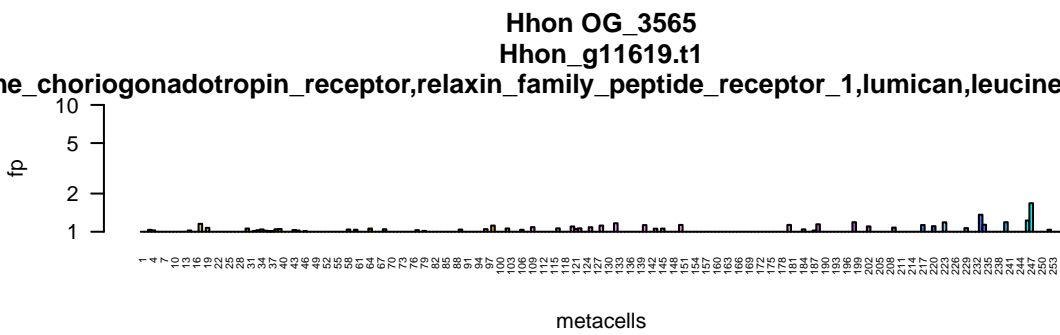
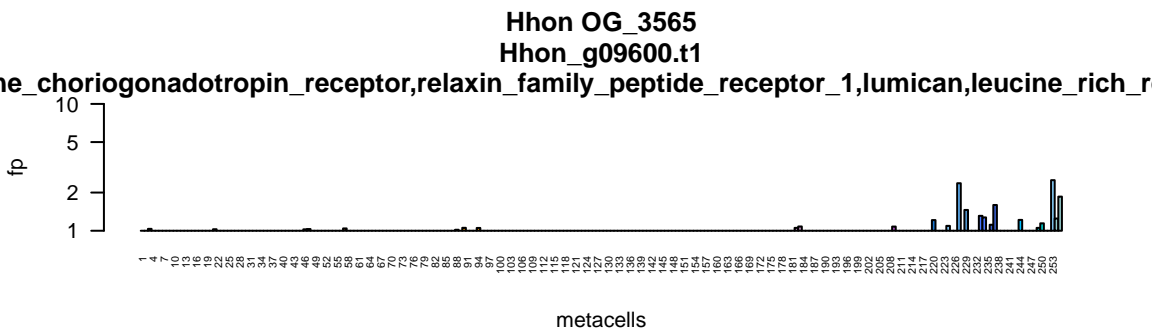
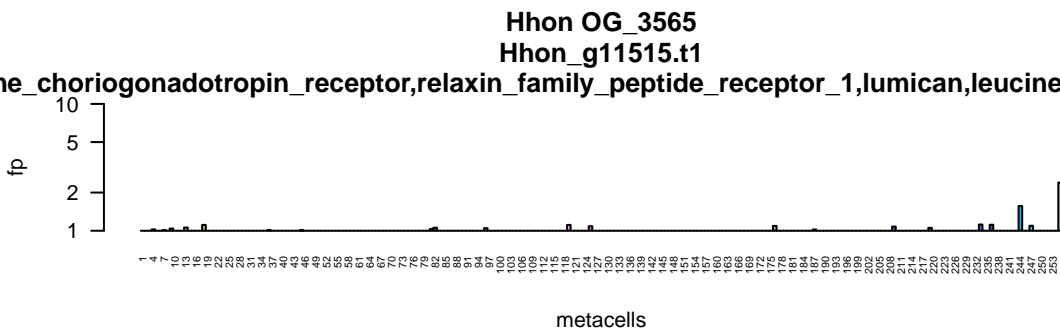
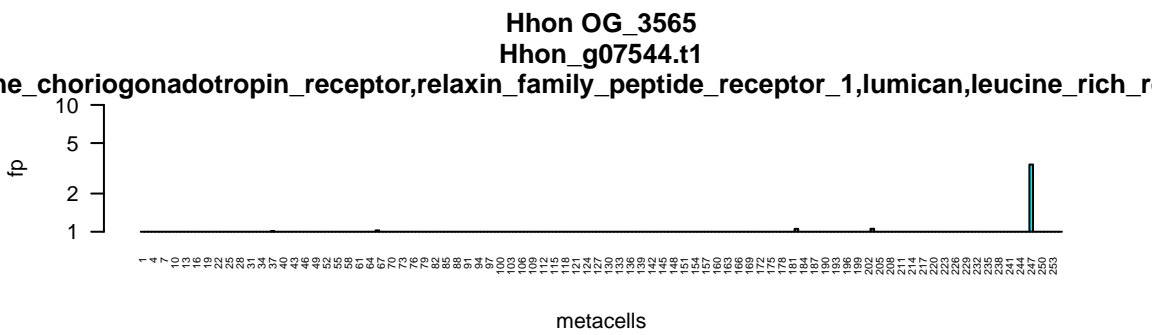
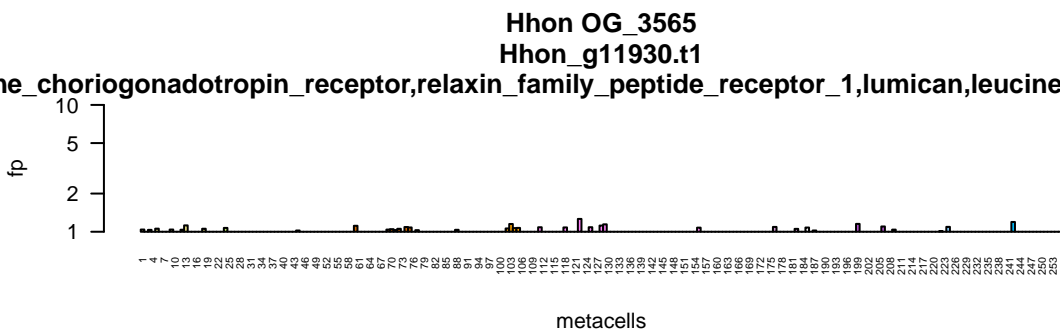
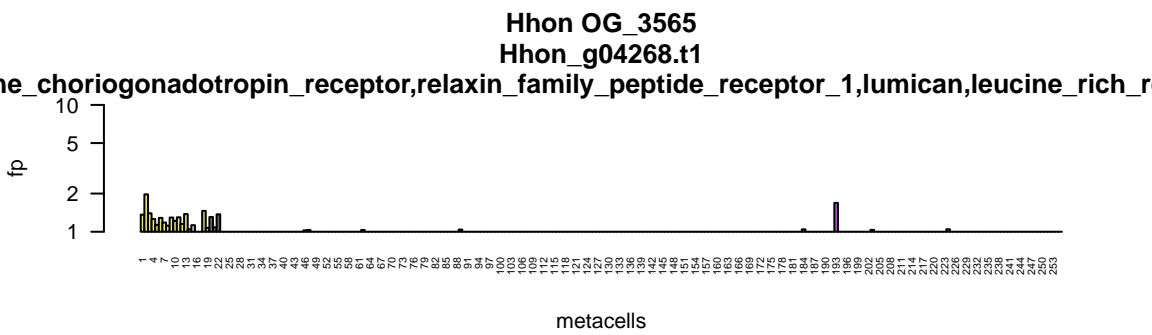
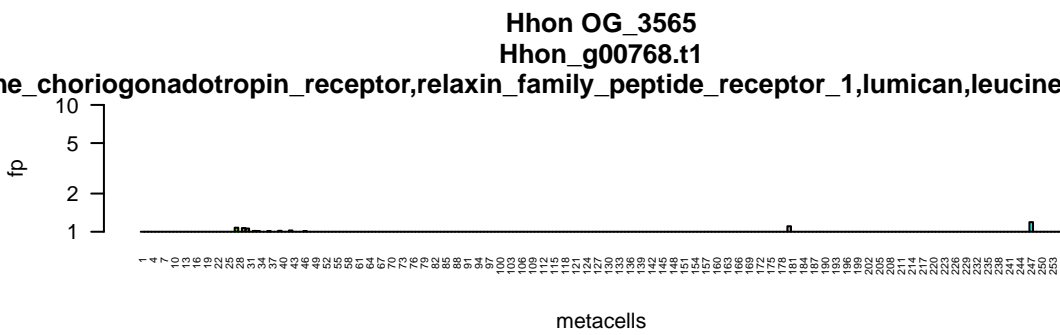
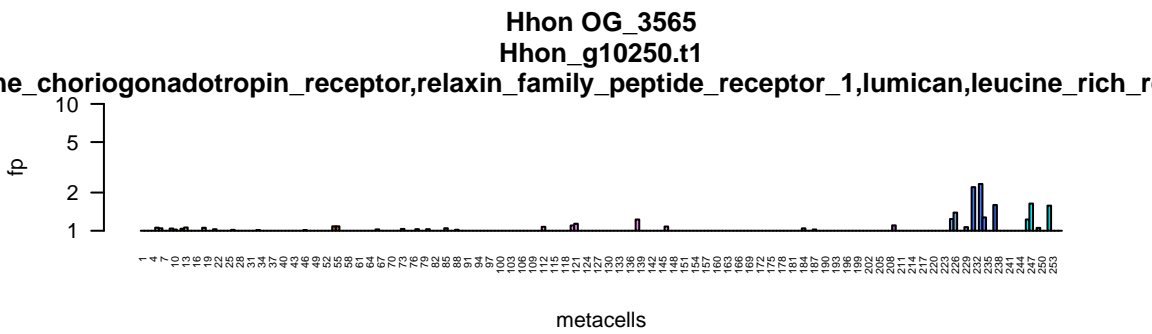
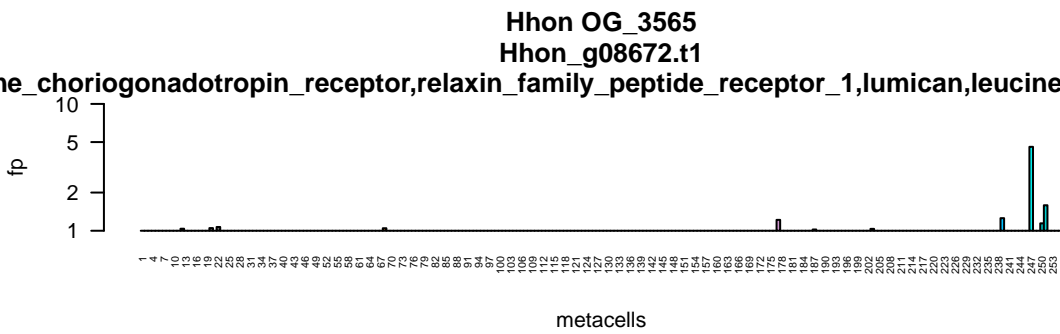


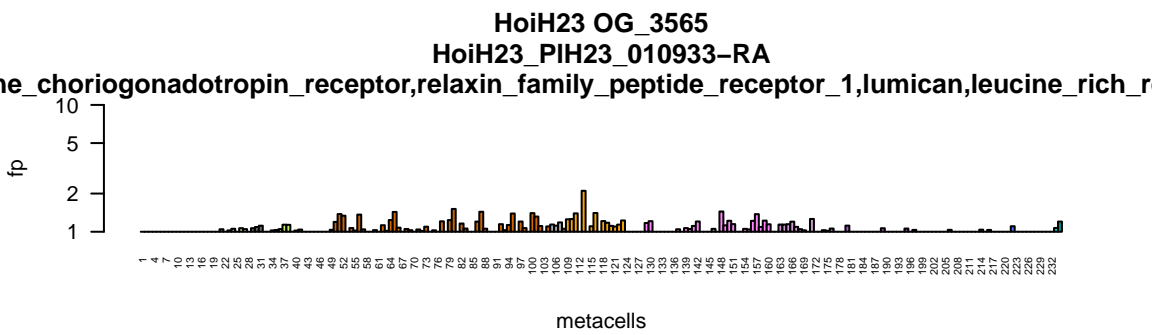
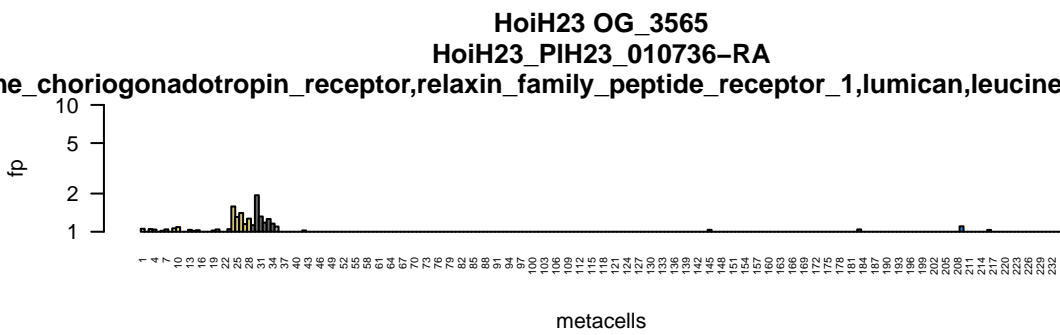
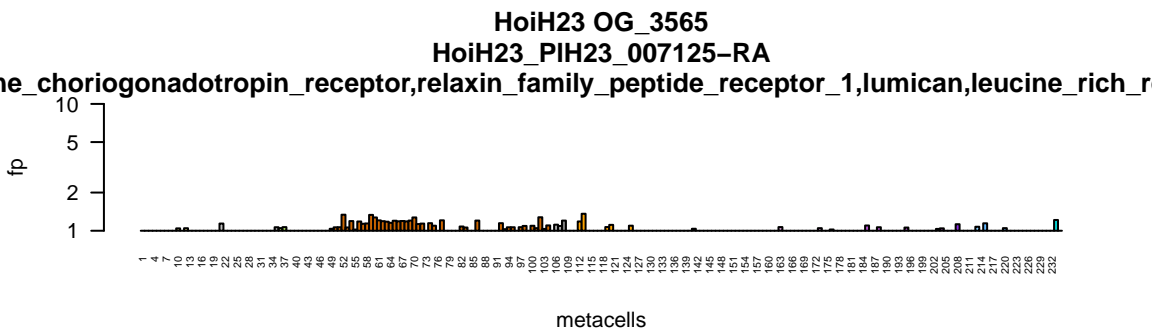
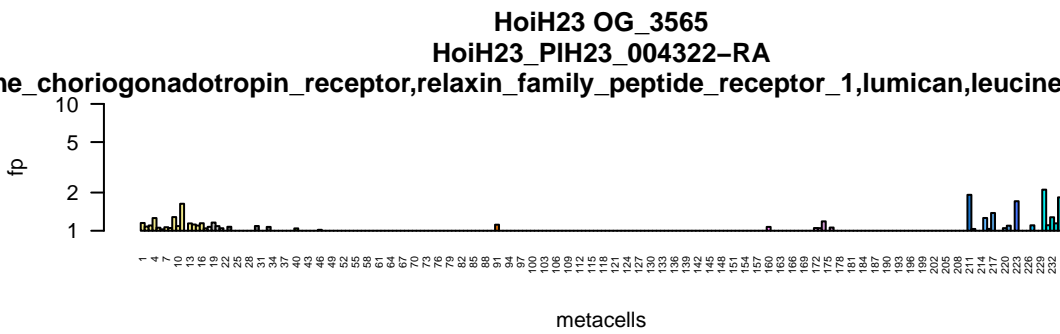
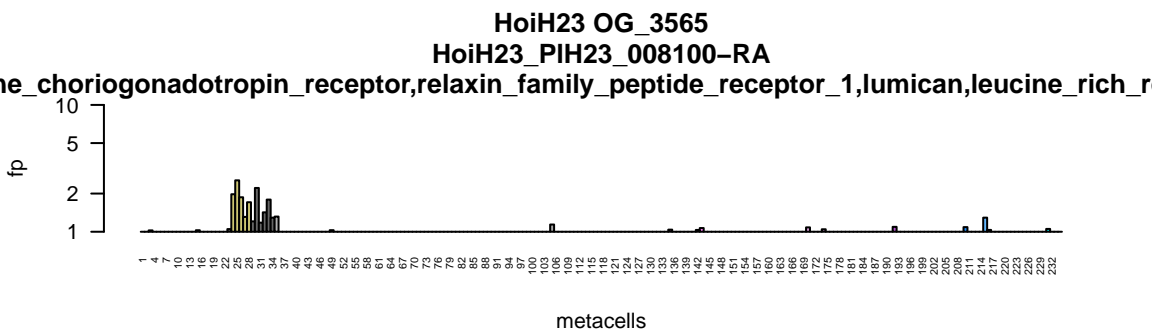
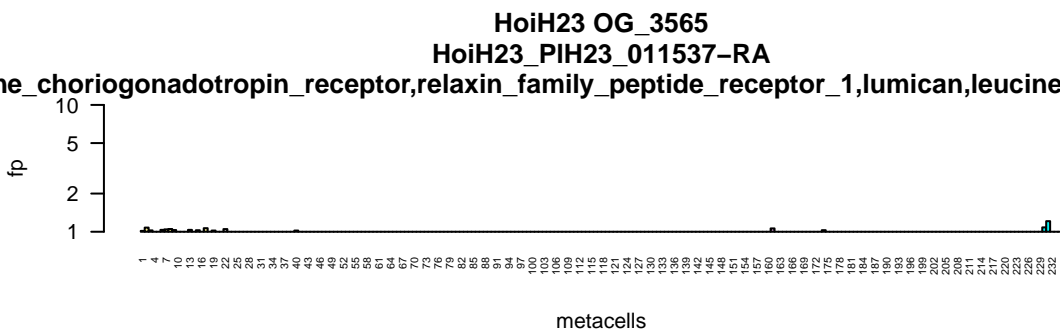
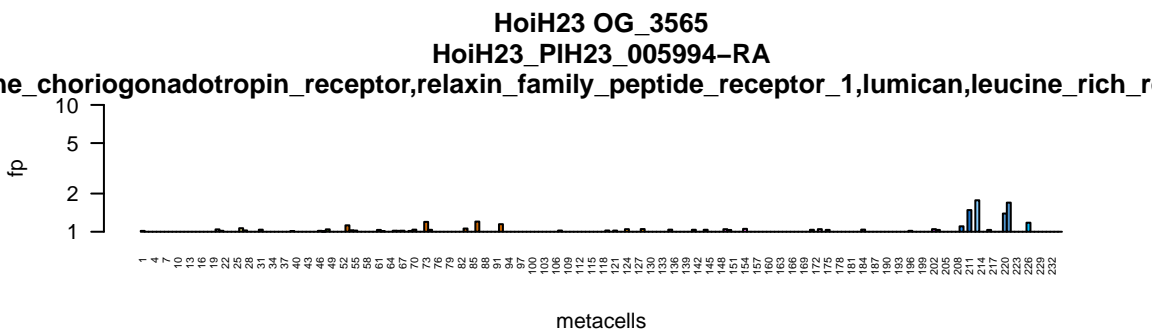
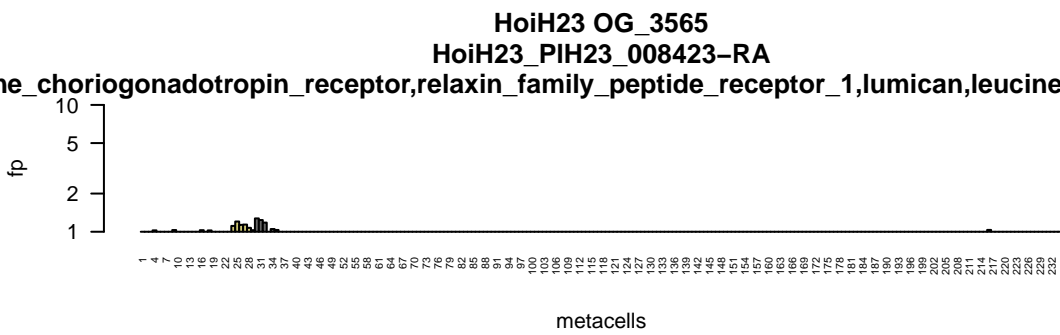
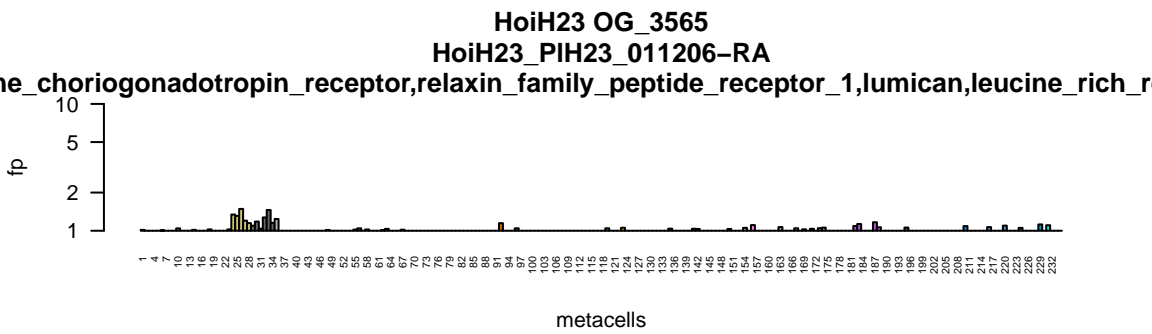
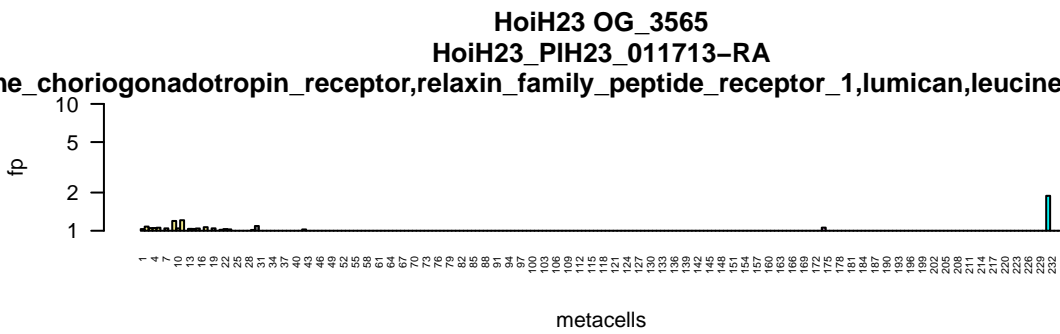
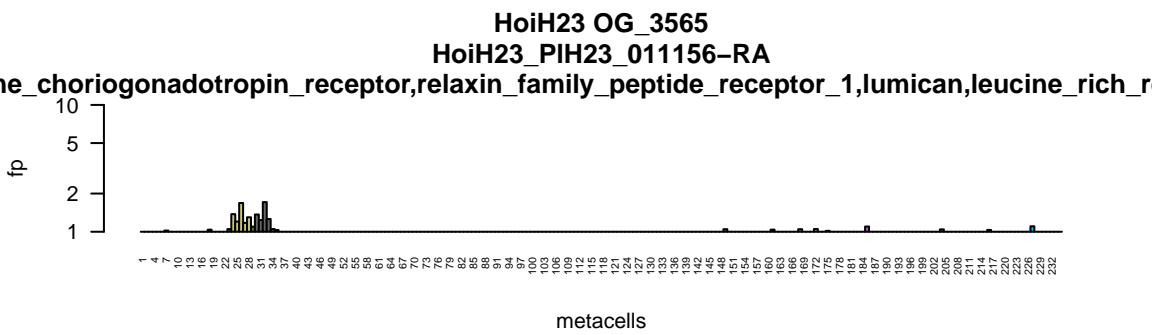
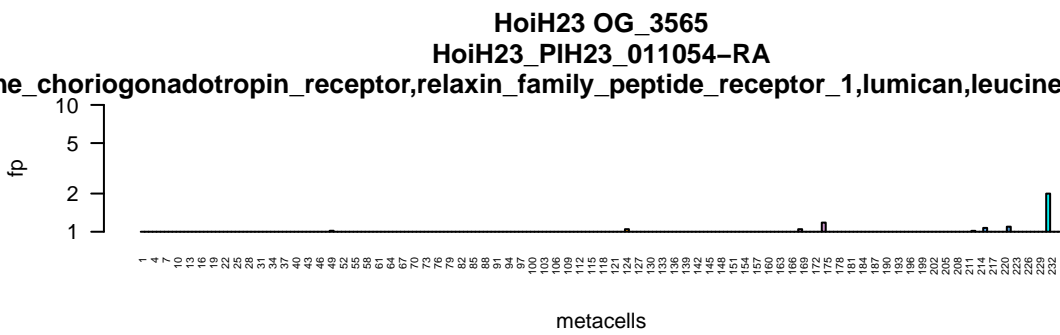
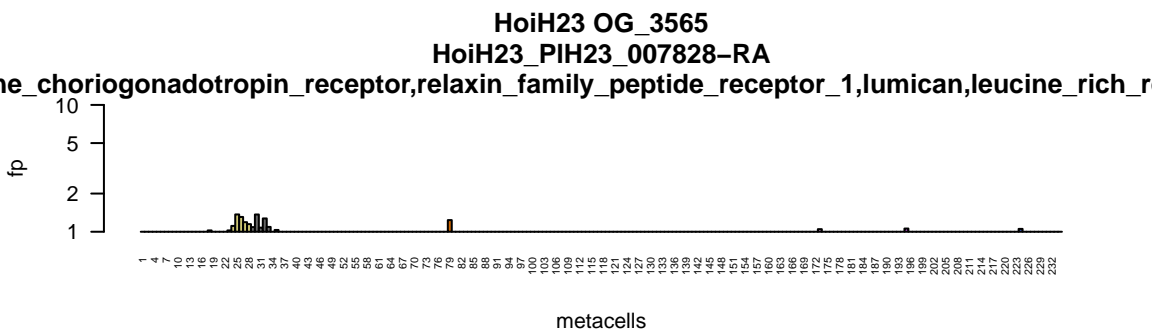
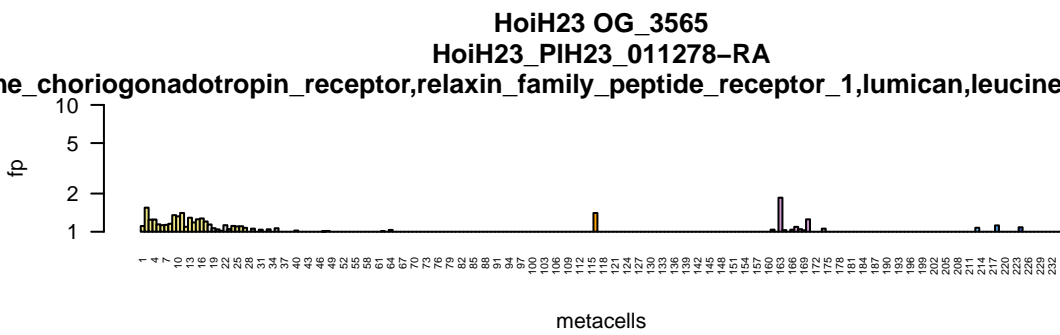
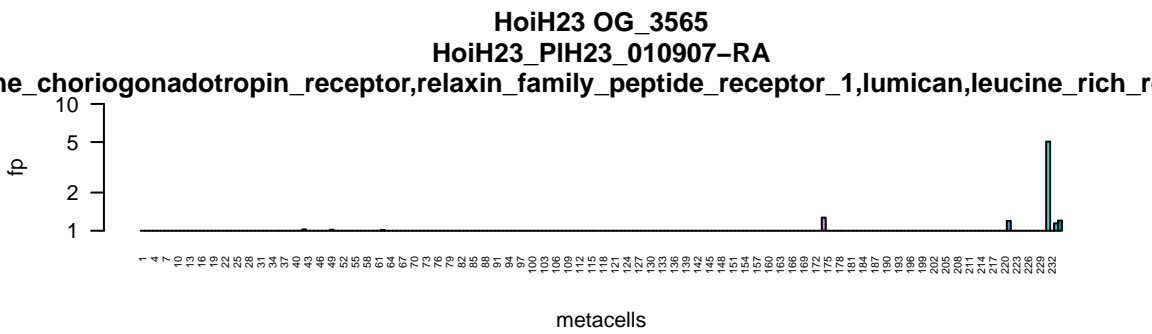
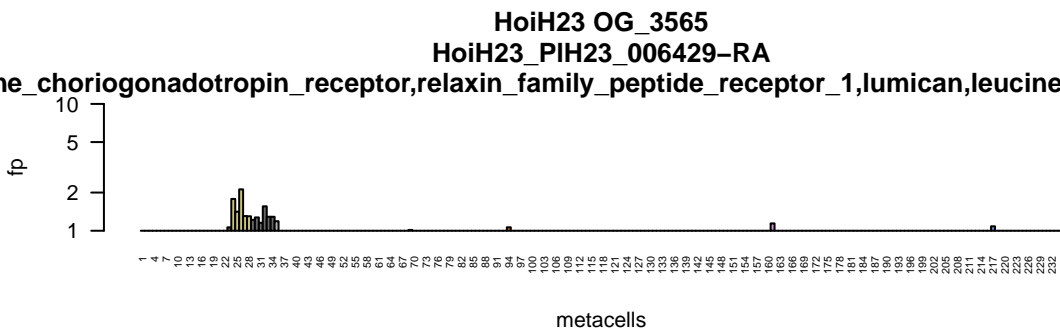


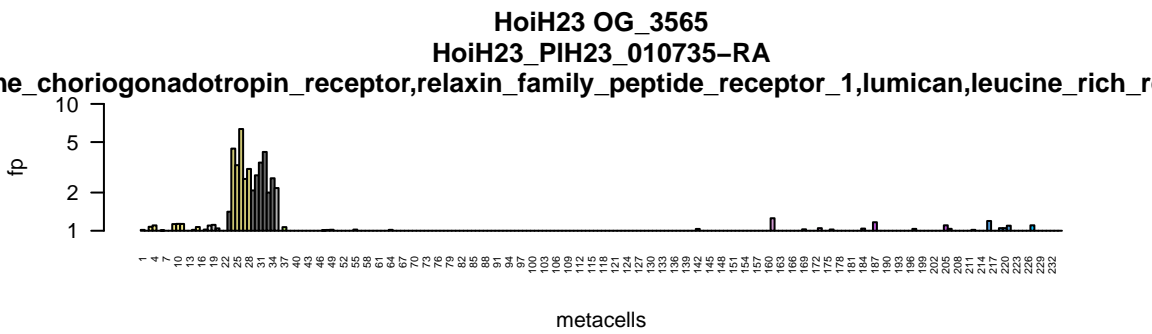
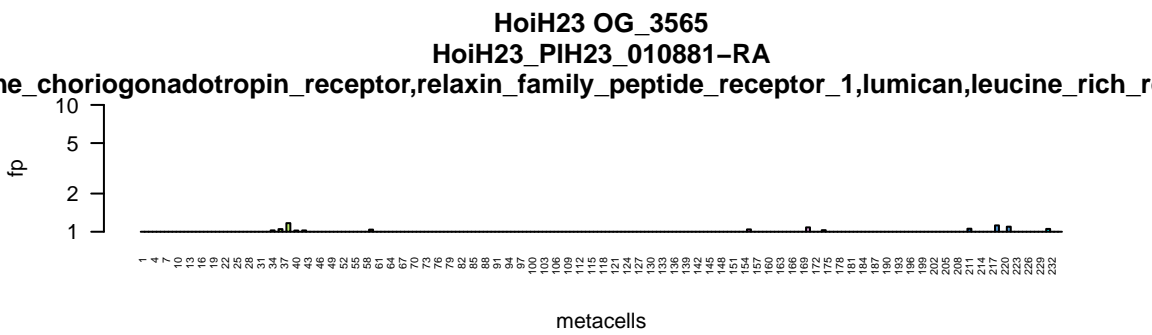
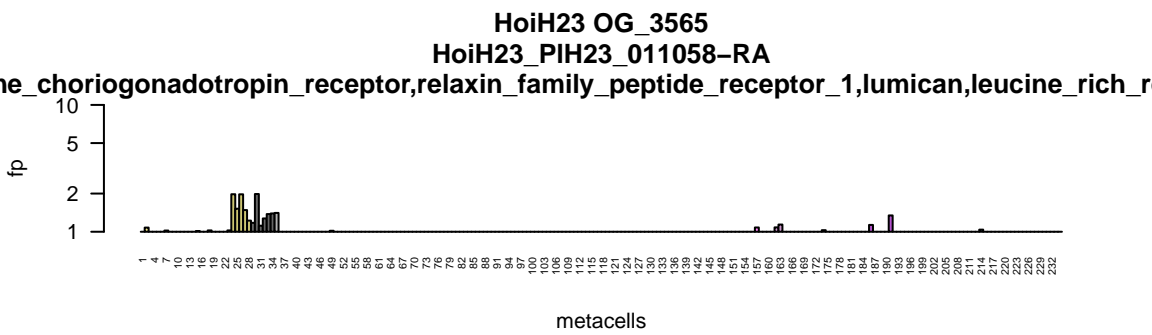
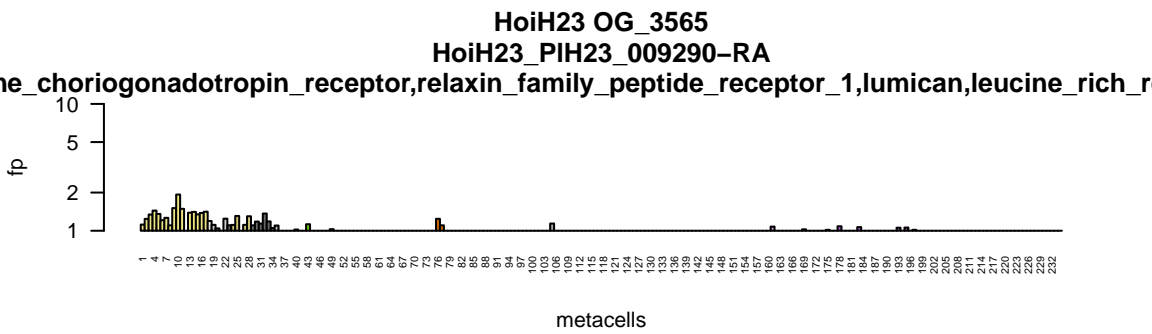


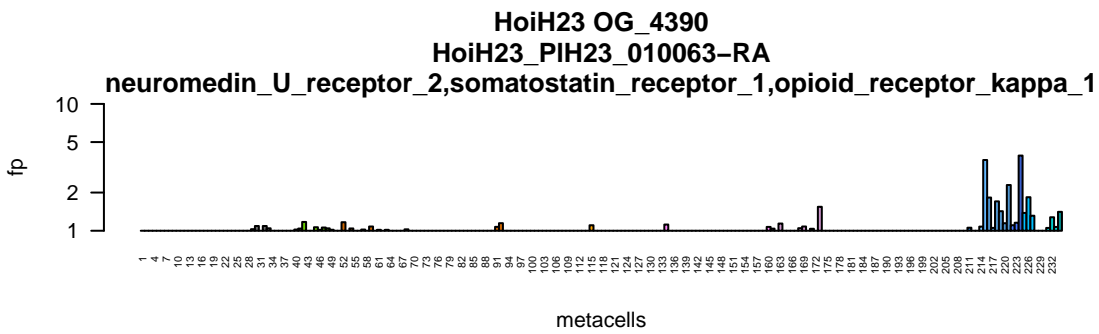
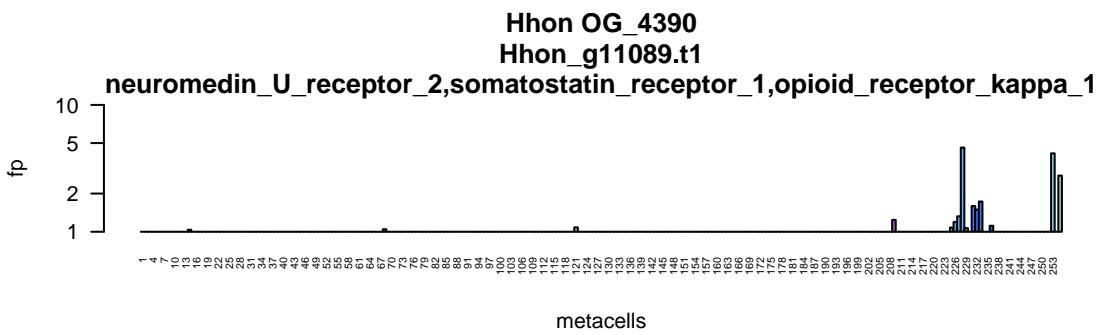
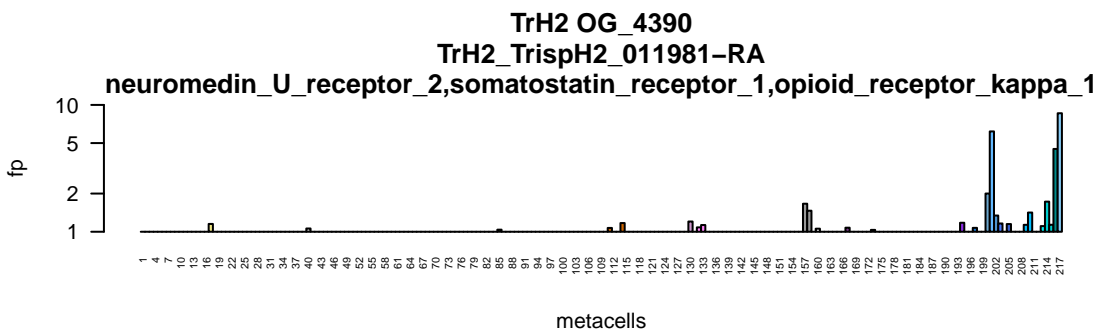
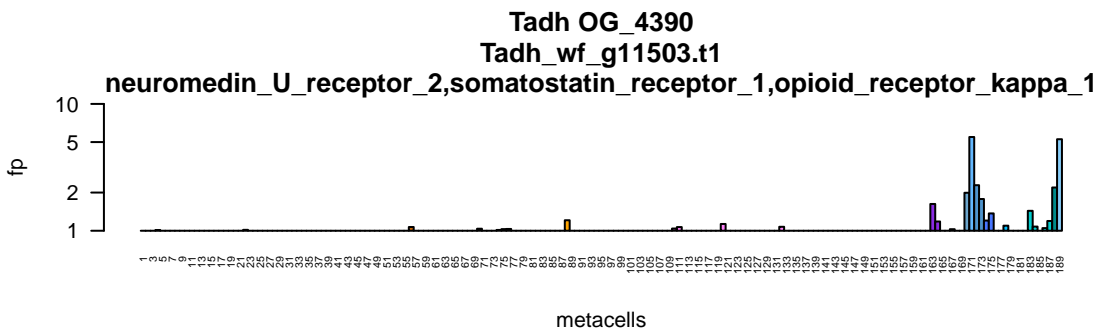




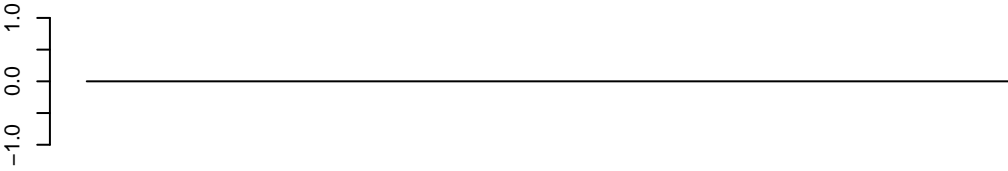




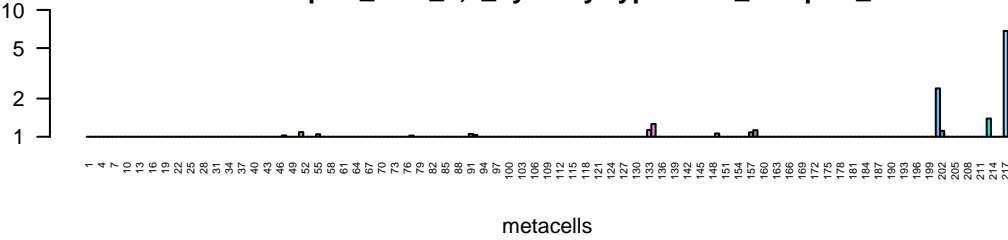




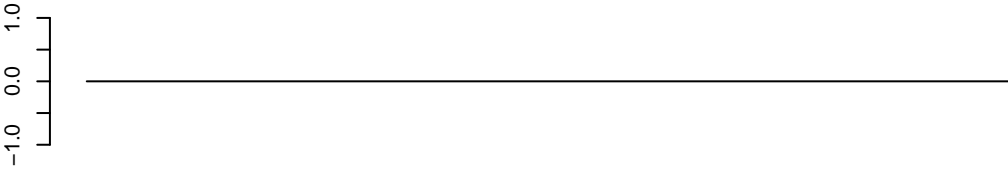
adrenoceptor_beta_2,5_hydroxytryptamine_receptor_2A
Tadh | no data



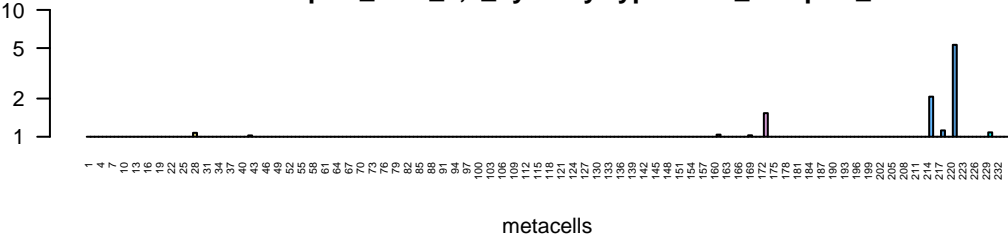
TrH2 OG_4563
TrH2_TrispH2_003322-RA
adrenoceptor_beta_2,5_hydroxytryptamine_receptor_2A

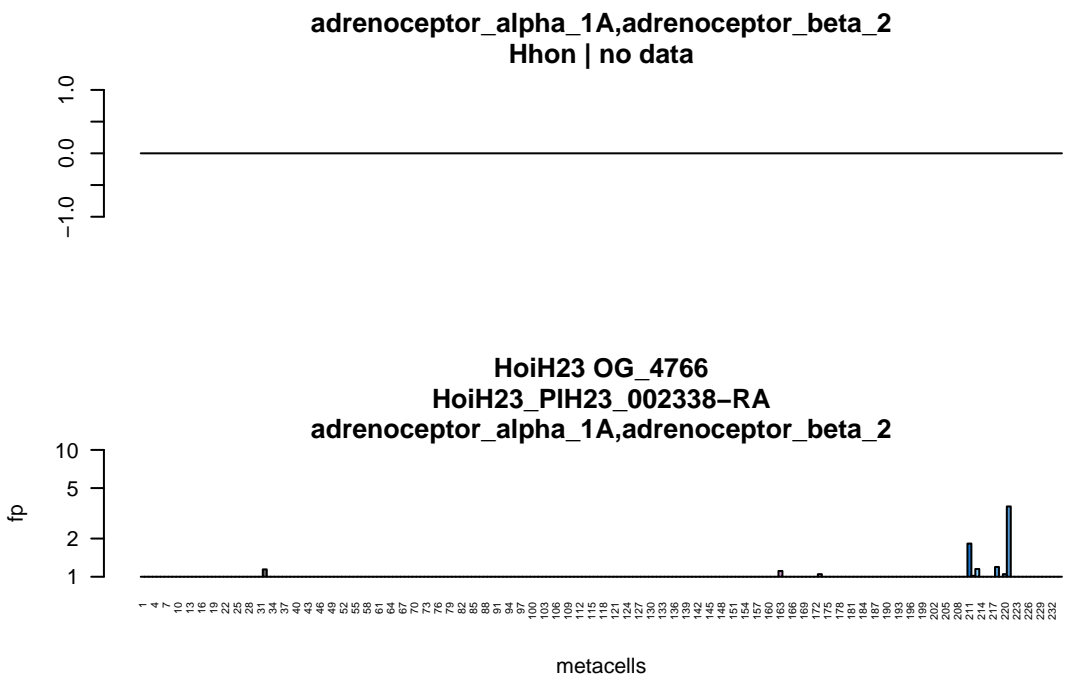
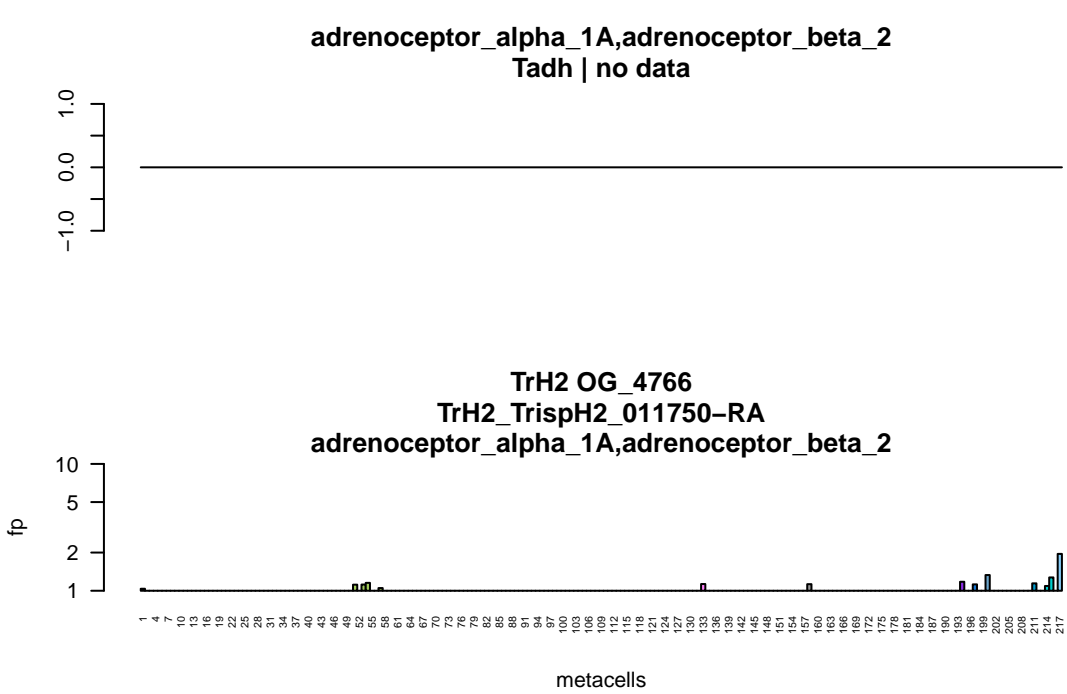


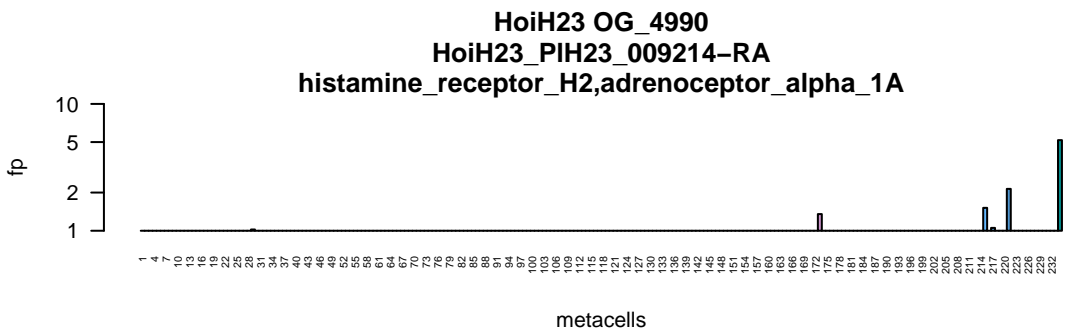
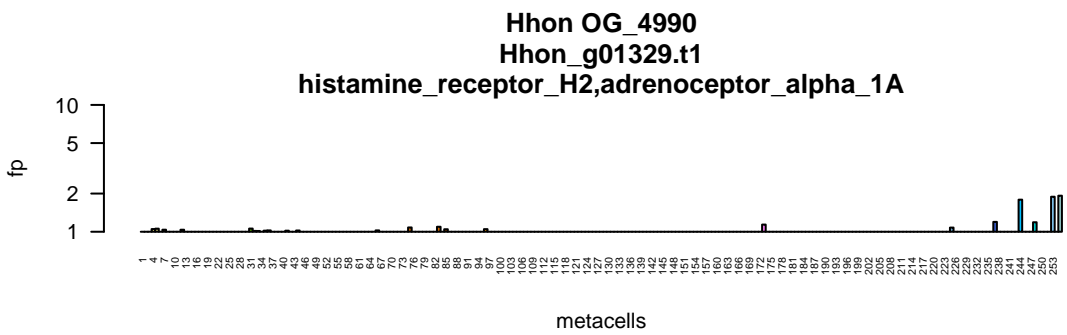
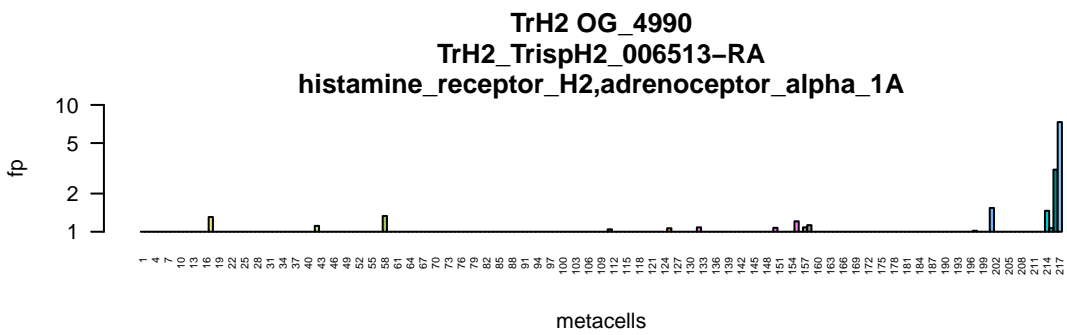
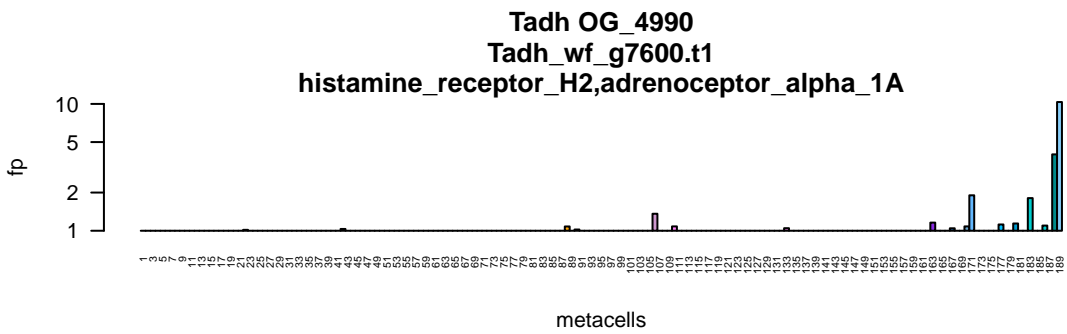
adrenoceptor_beta_2,5_hydroxytryptamine_receptor_2A
Hhon | no data

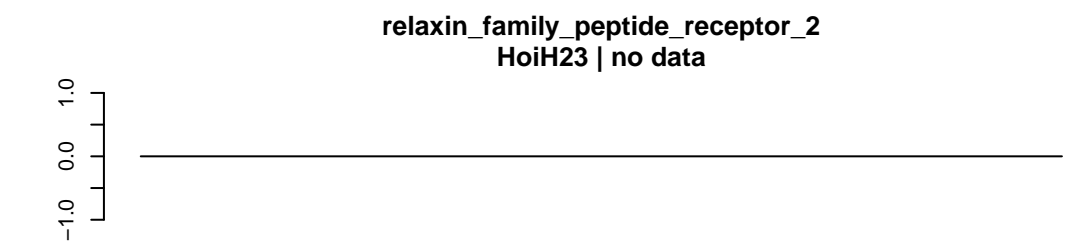
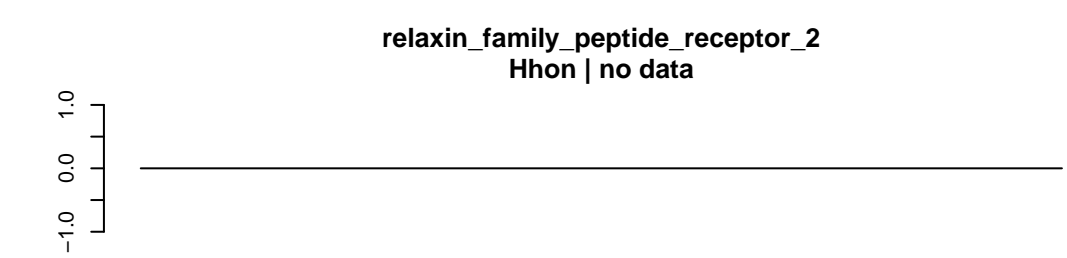
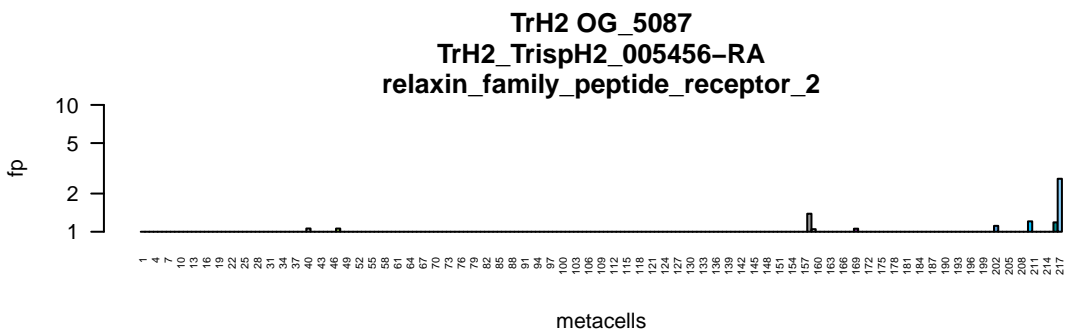
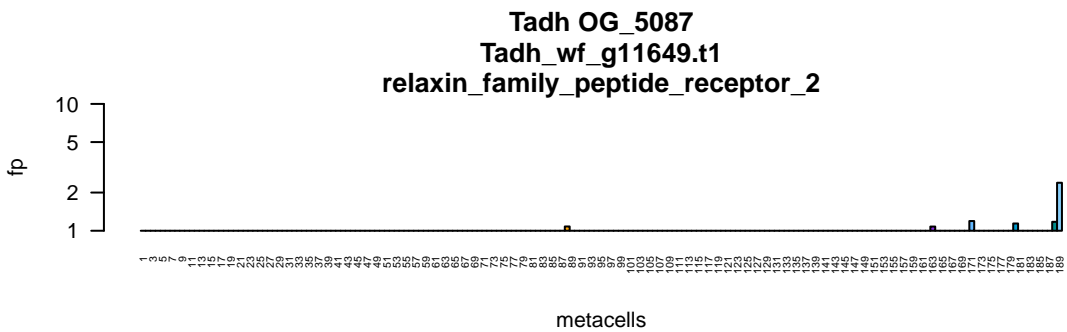


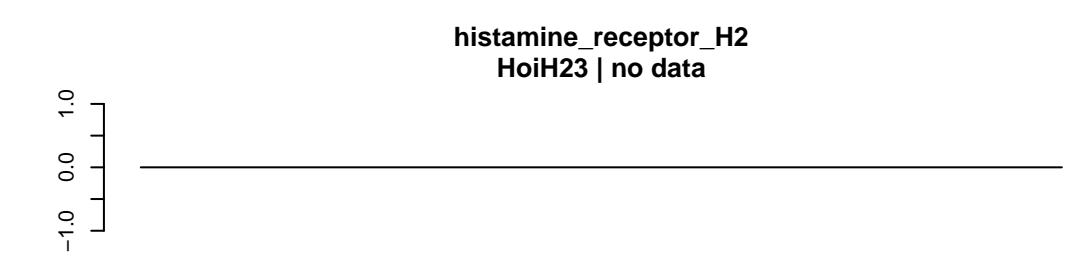
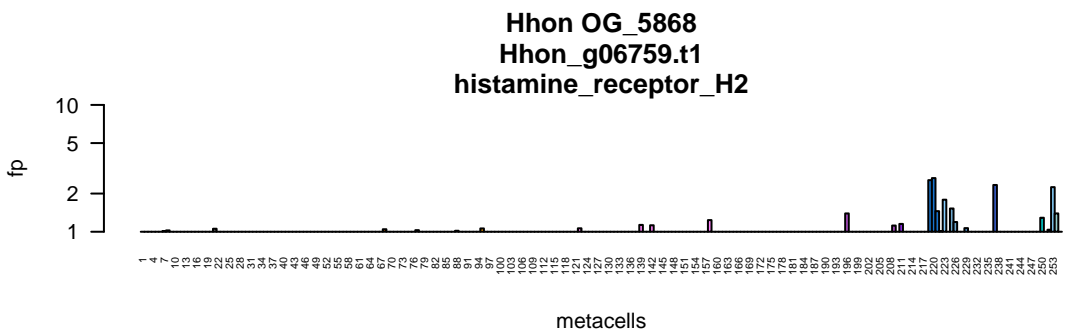
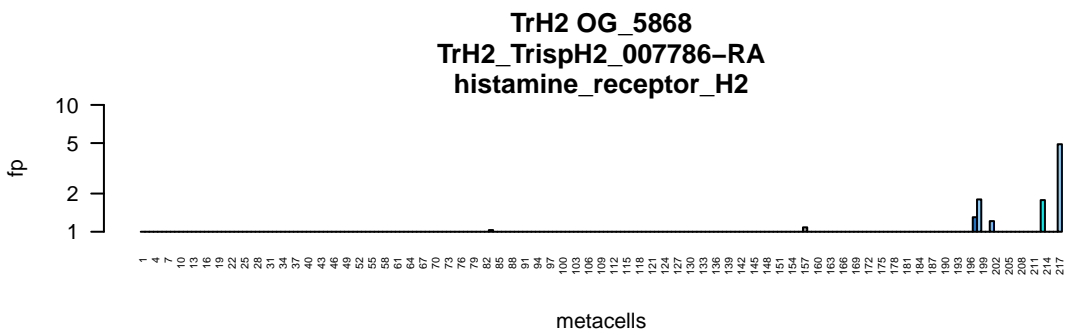
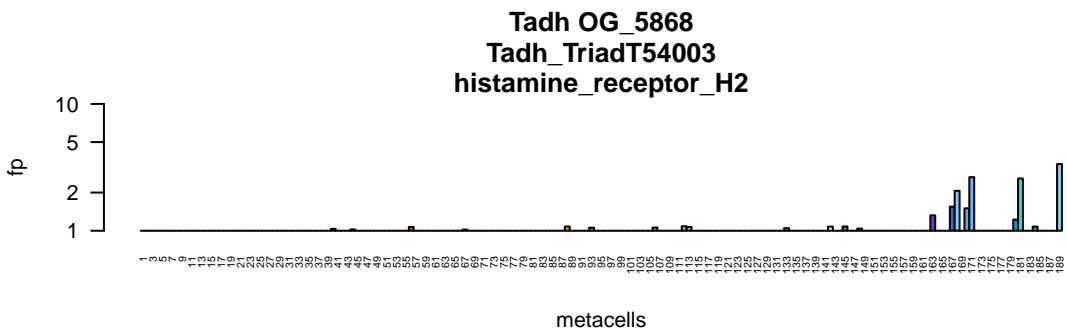
HoiH23 OG_4563
HoiH23_PIH23_009618-RA
adrenoceptor_beta_2,5_hydroxytryptamine_receptor_2A



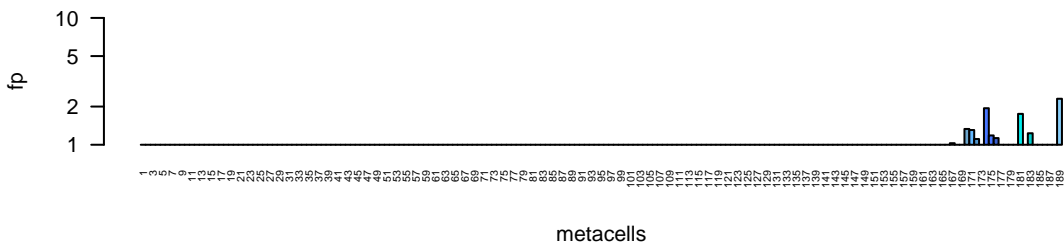




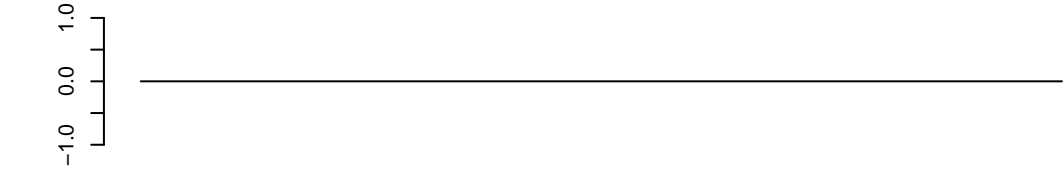




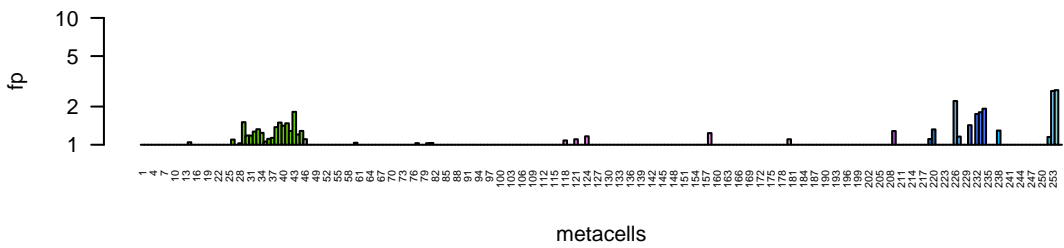
Tadh OG_6481
Tadh_wf_g9558.t1



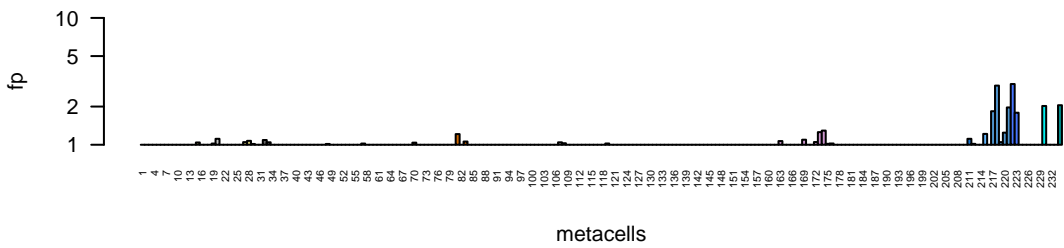
TrH2 | no data

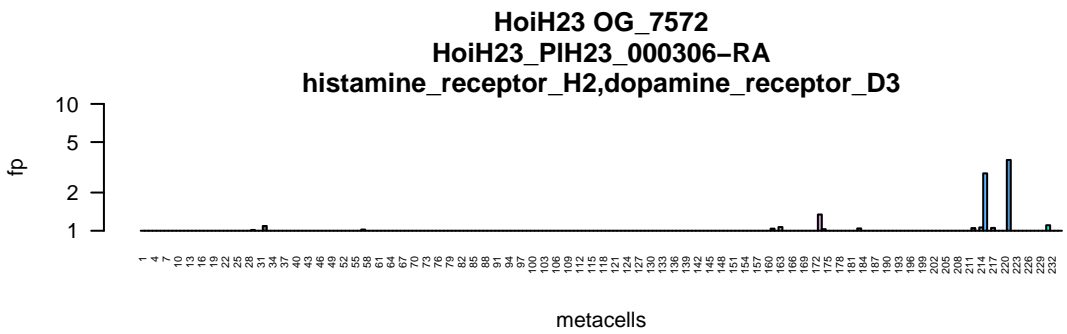
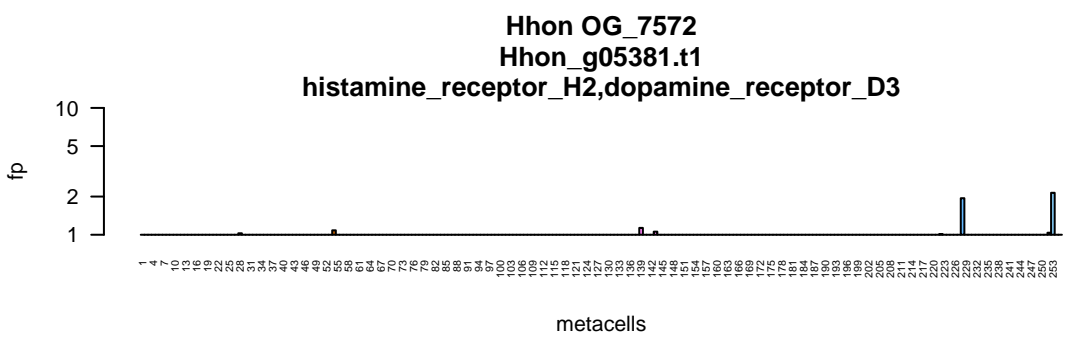
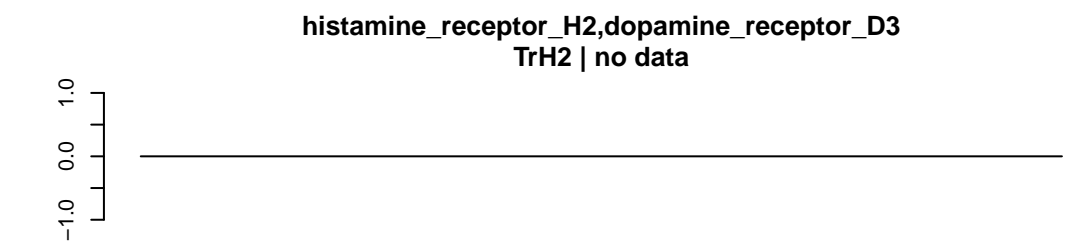
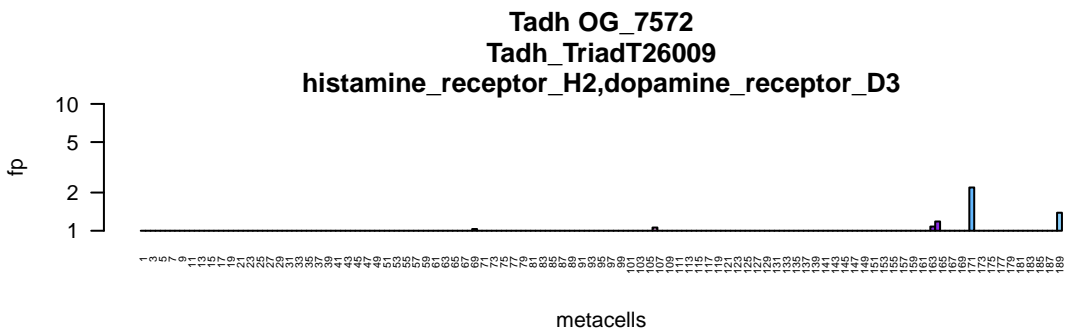


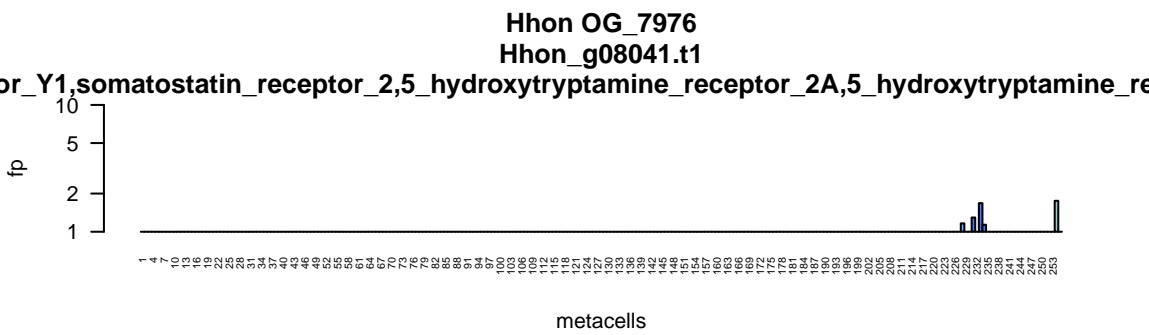
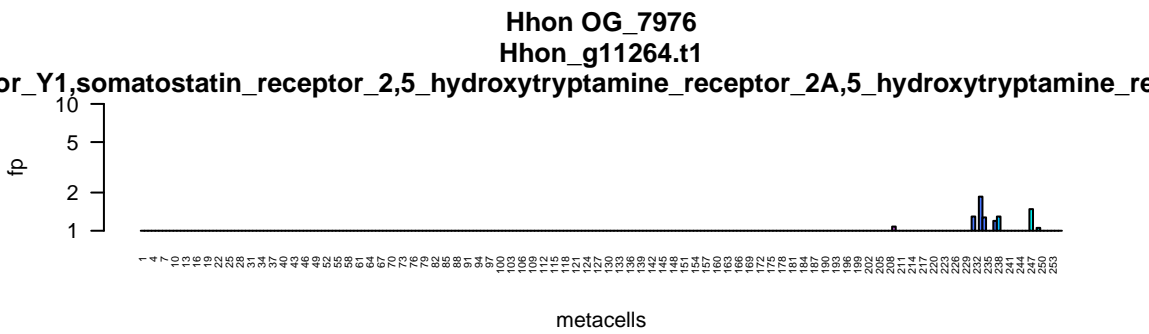
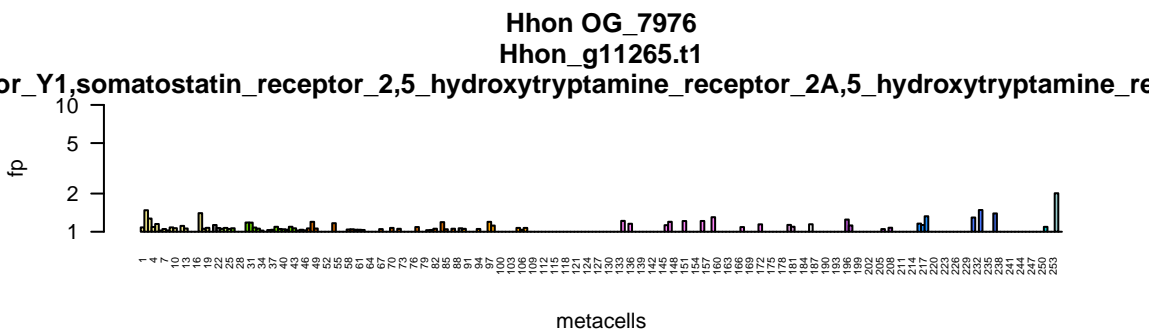
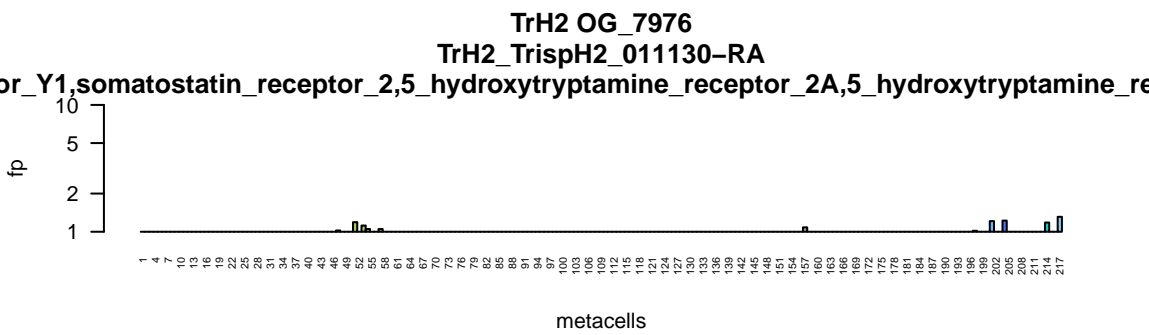
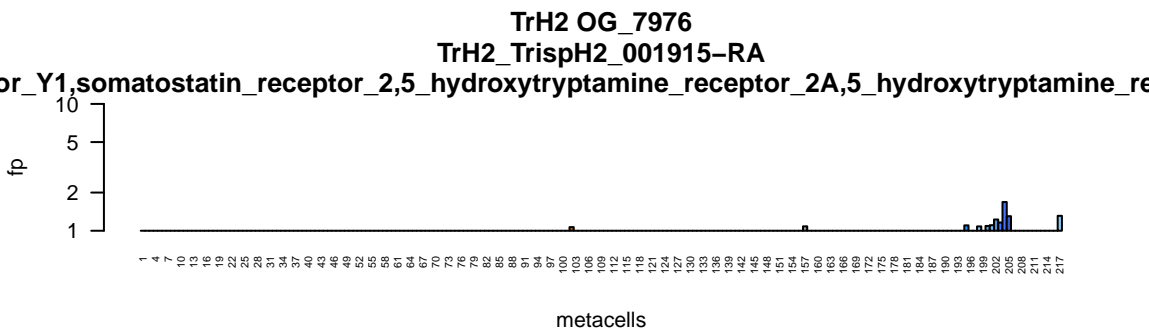
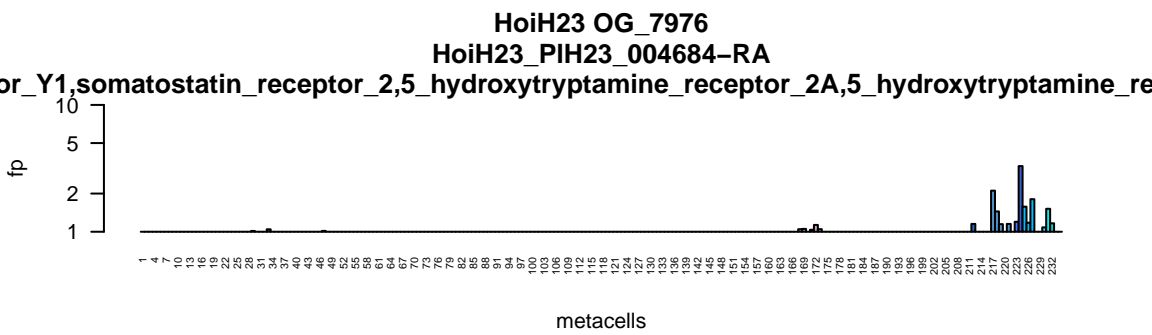
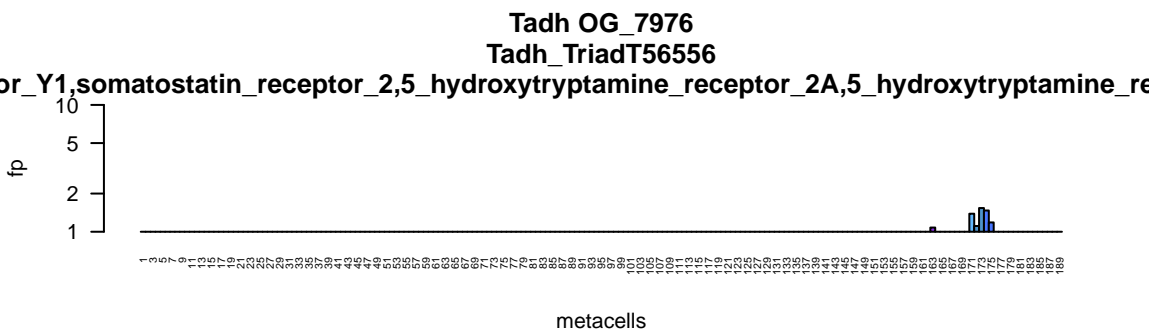
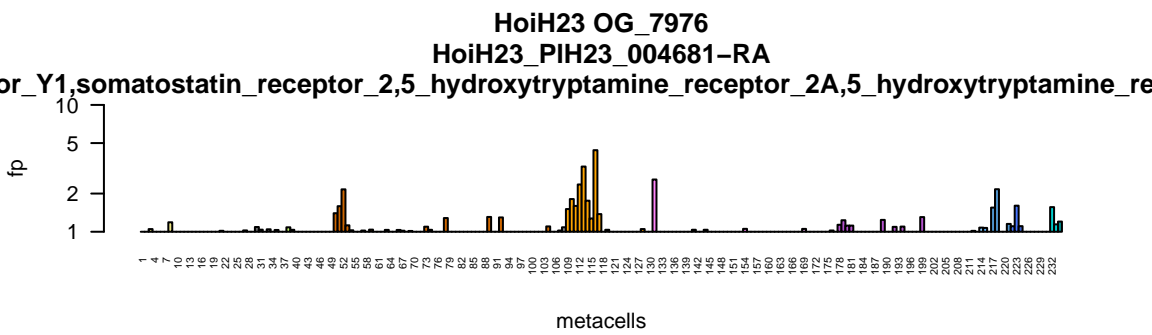
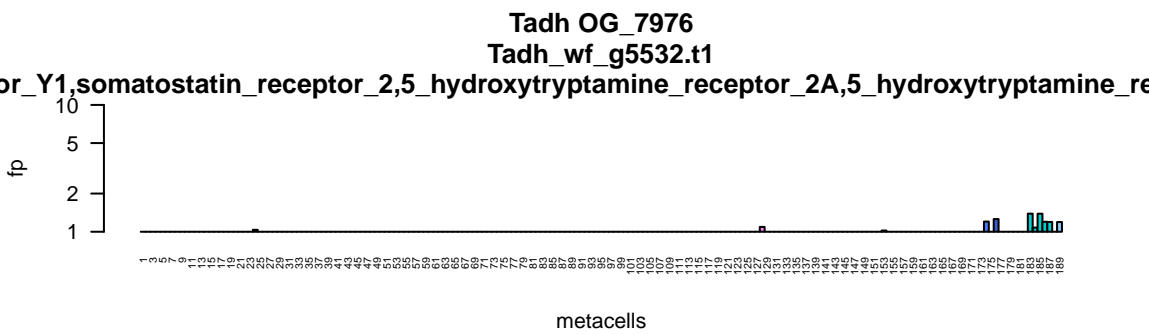
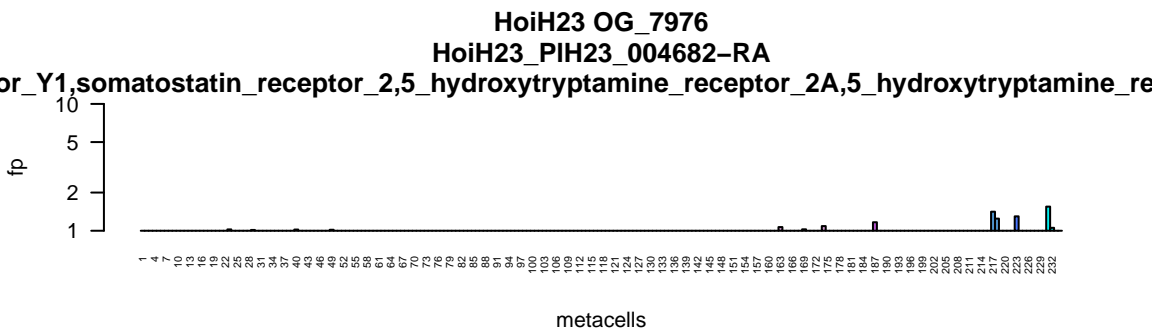
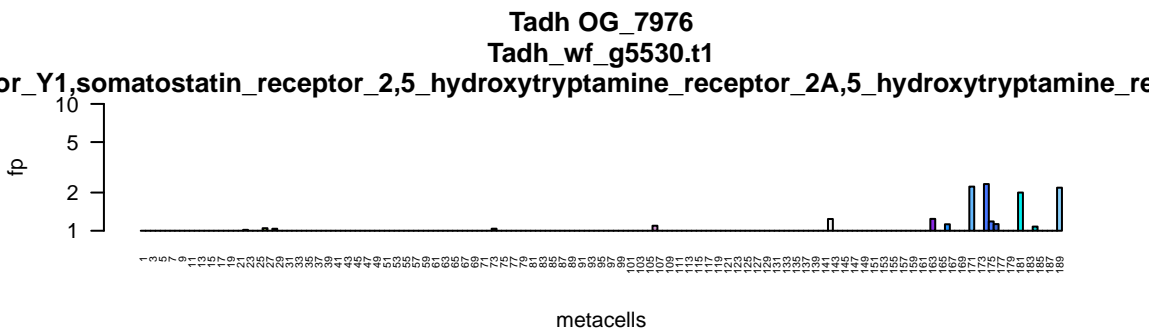
Hhon OG_6481
Hhon_g10968.t1

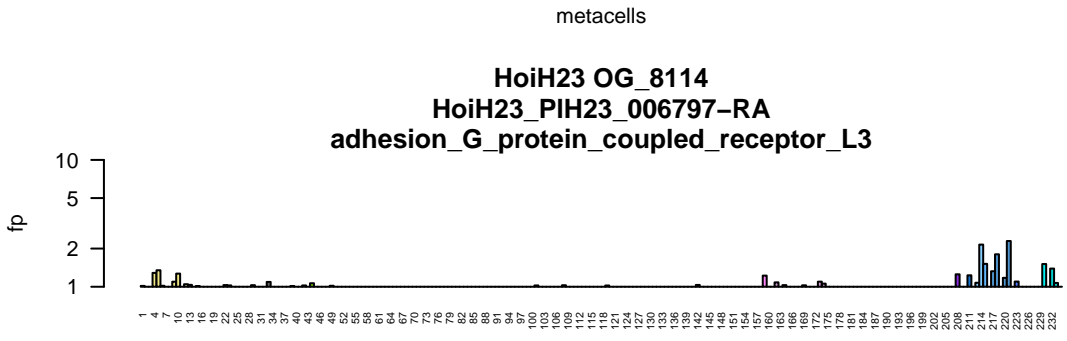
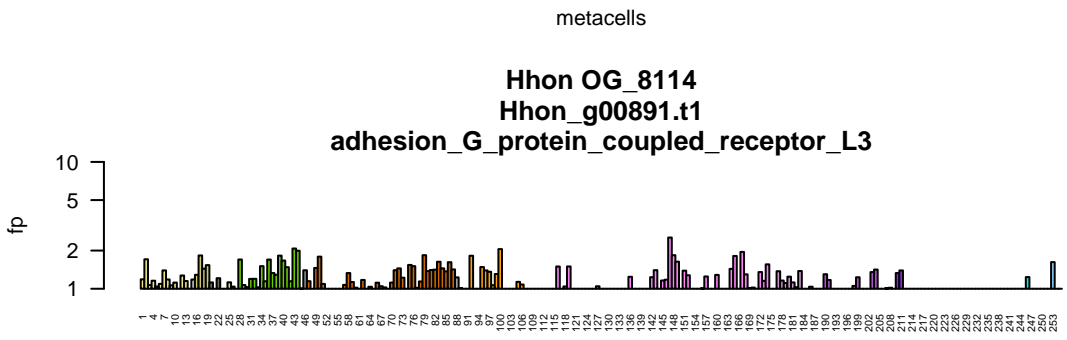
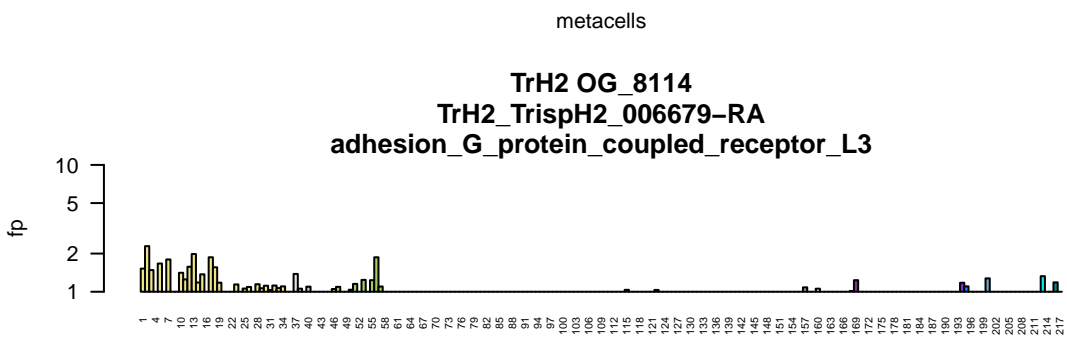
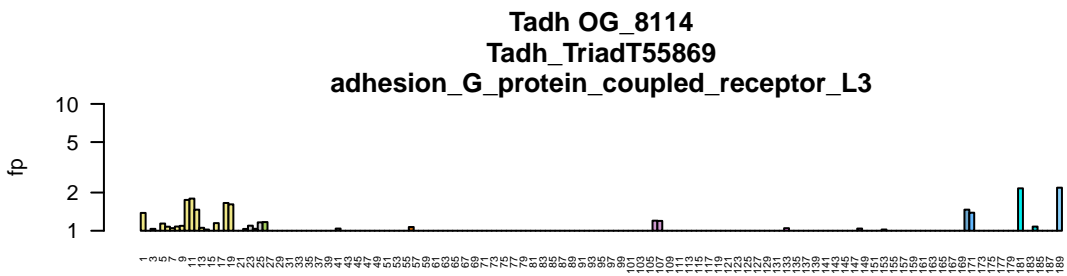


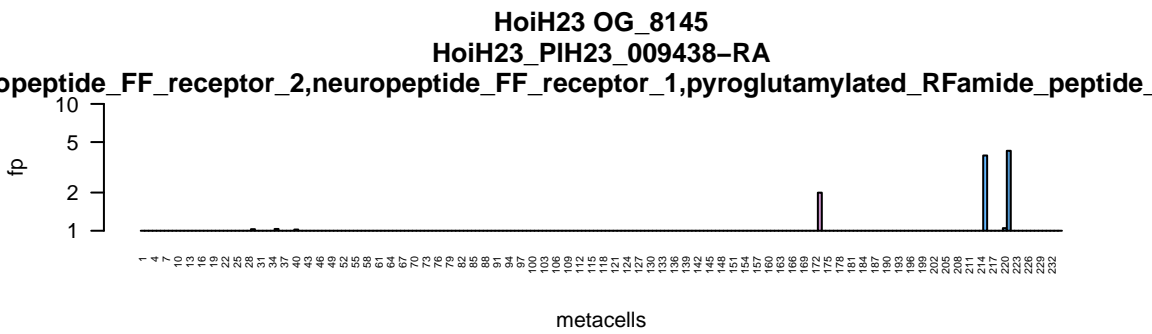
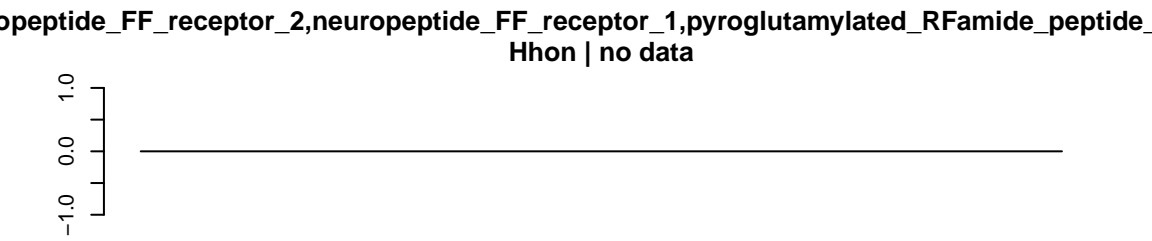
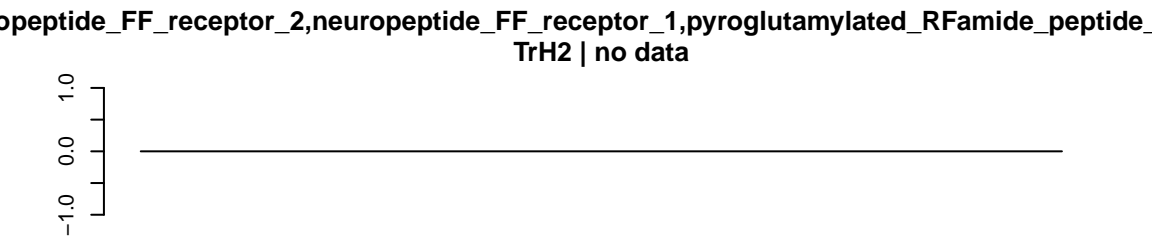
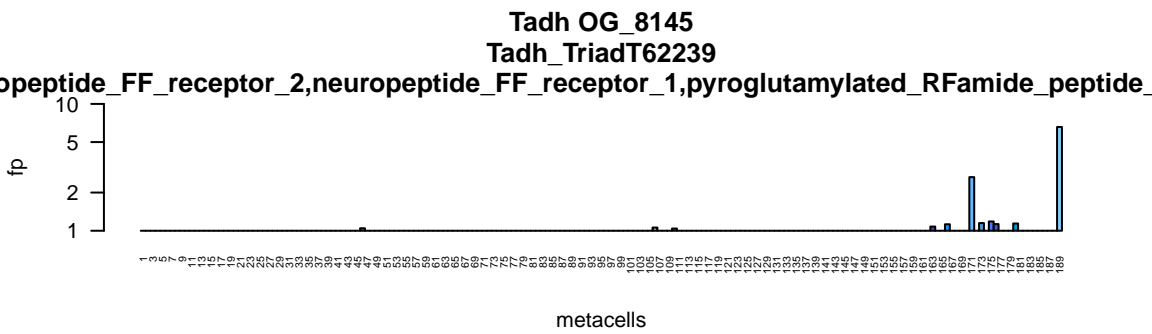
HoiH23 OG_6481
HoiH23_PIH23_011005-RA

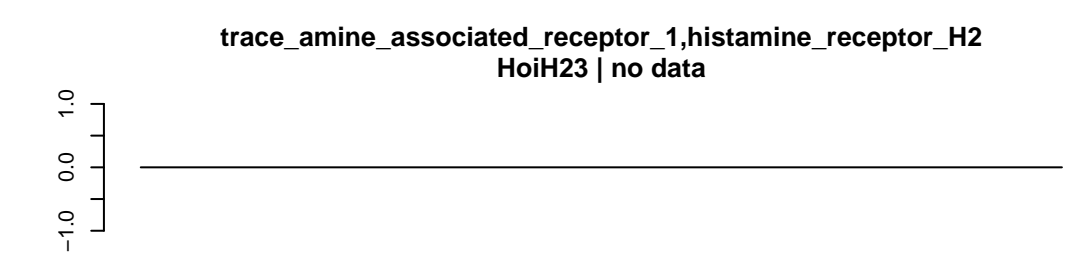
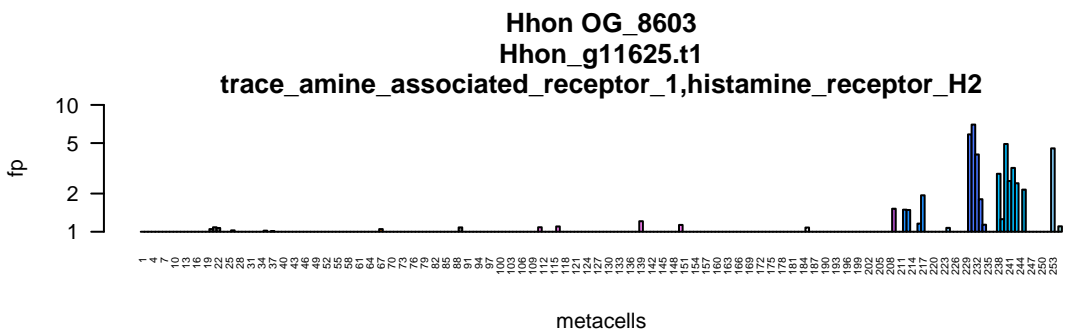
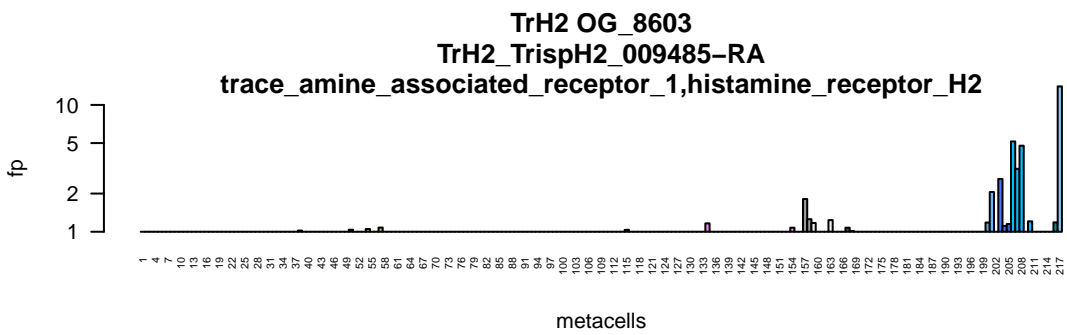
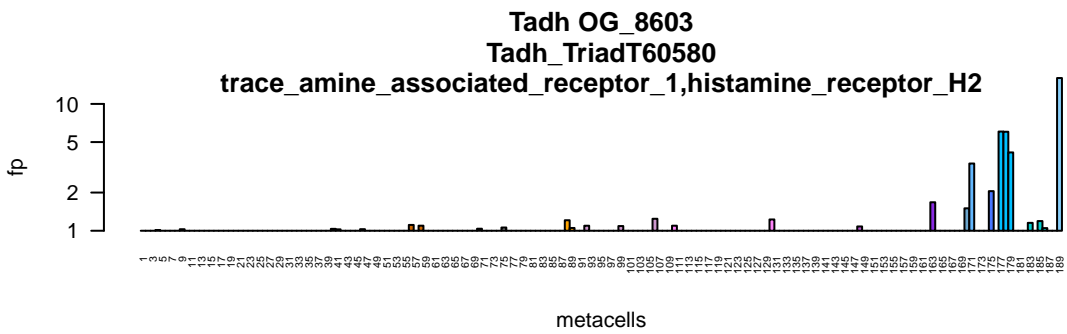


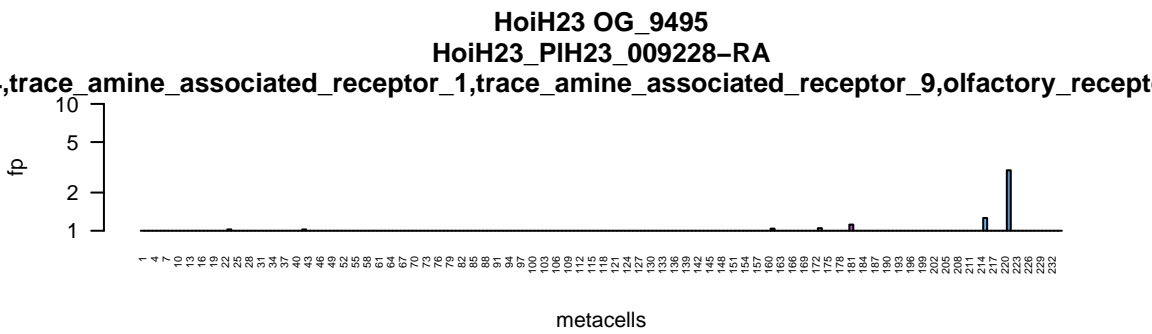
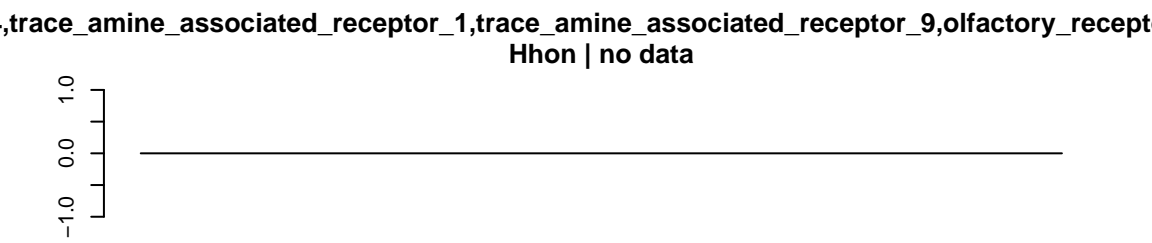
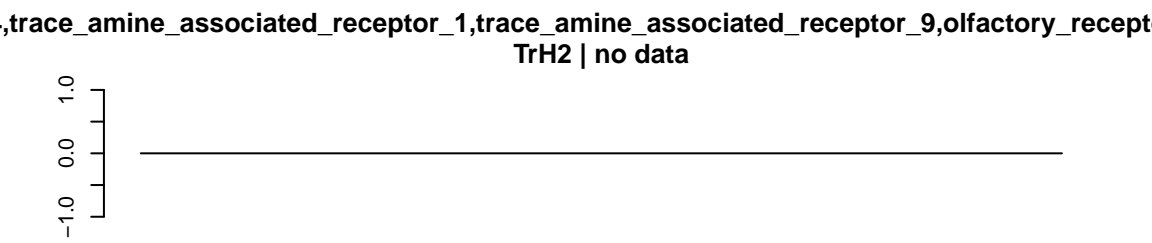
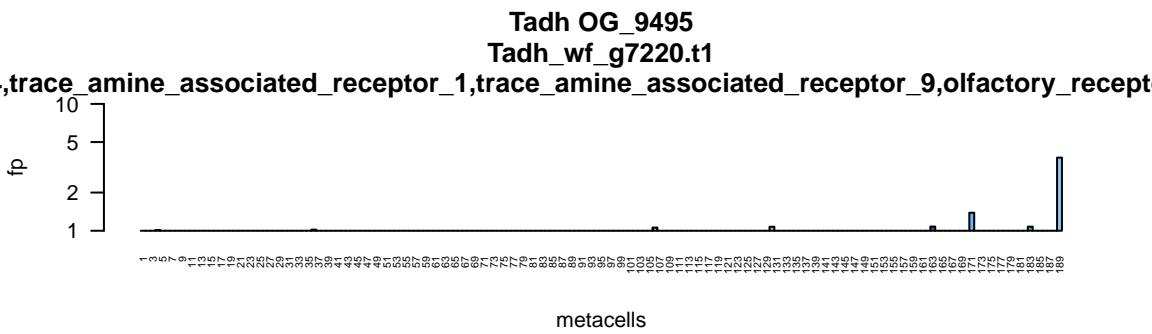


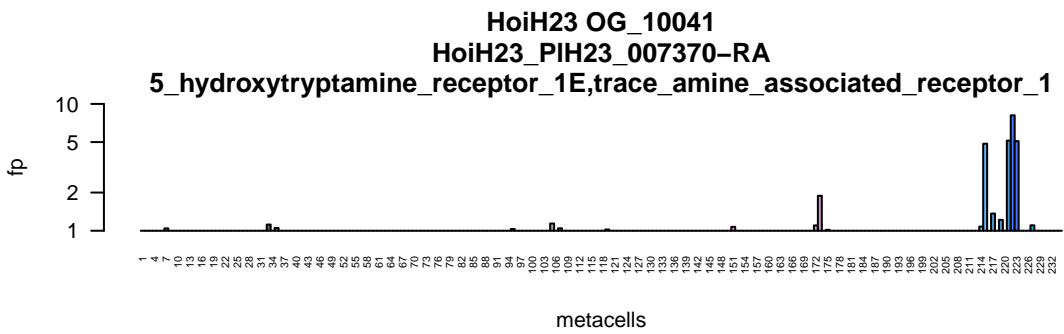
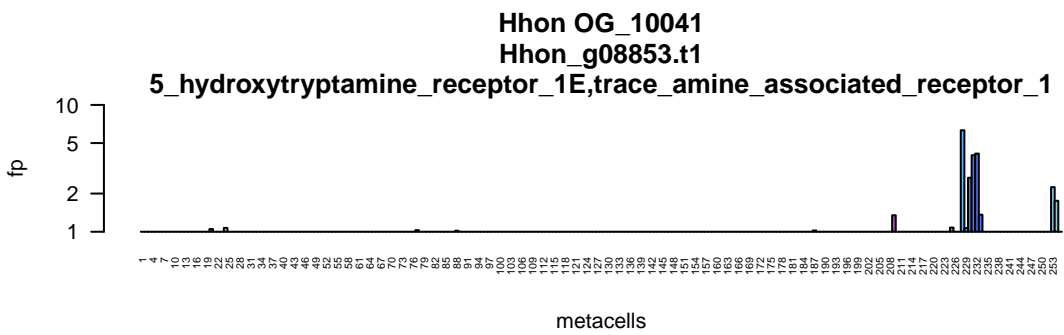
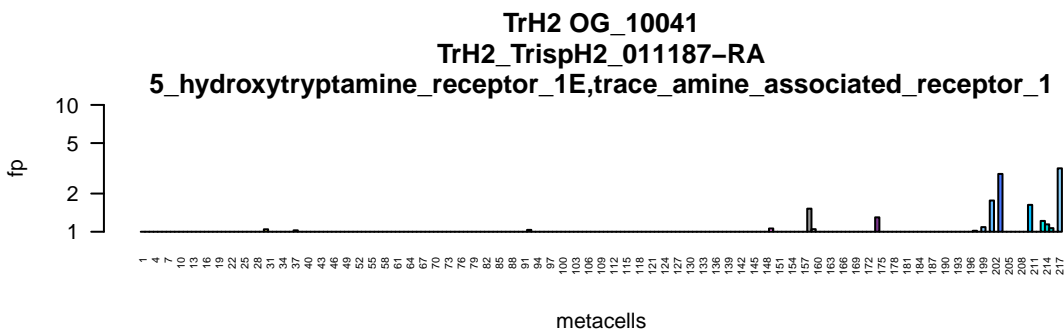
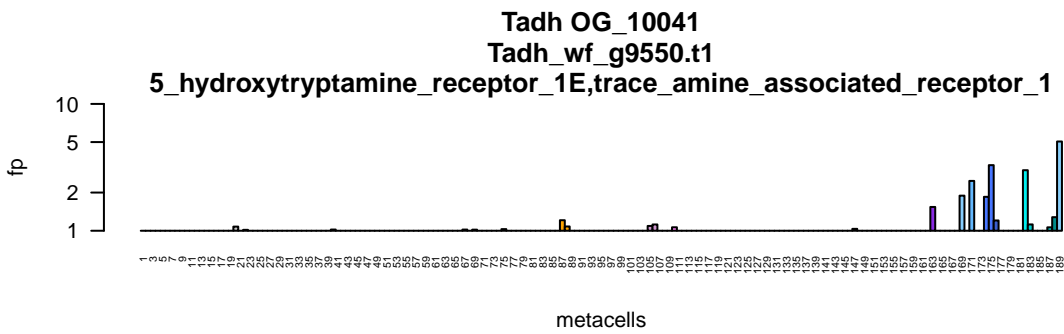




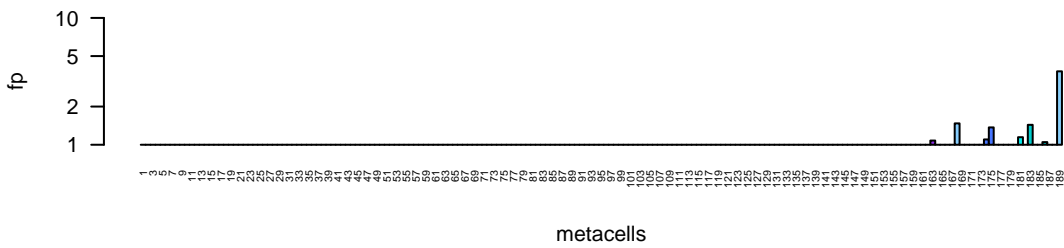




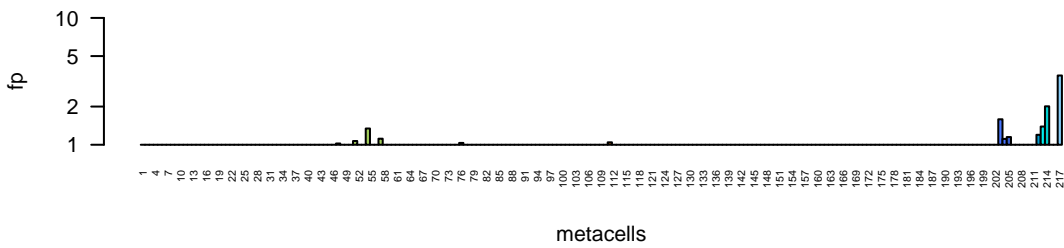




Tadh OG_10104
Tadh_TriadT60048



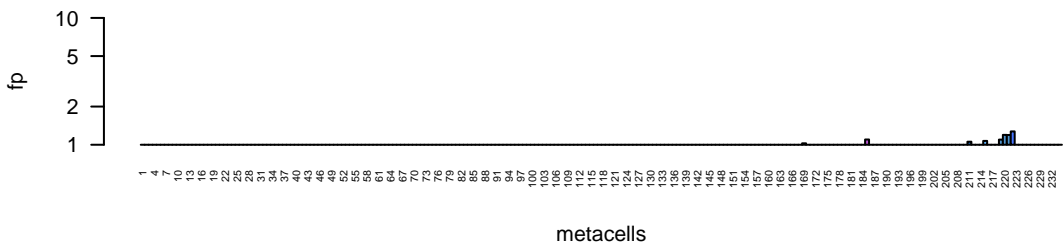
TrH2 OG_10104
TrH2_TrispH2_011300-RA

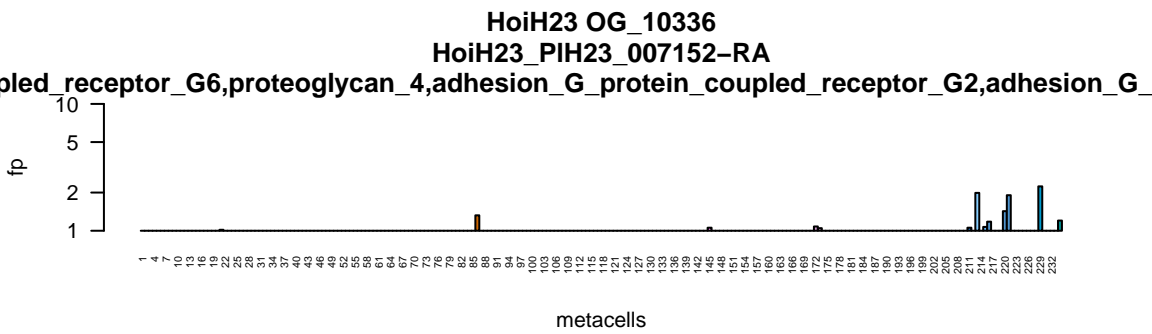
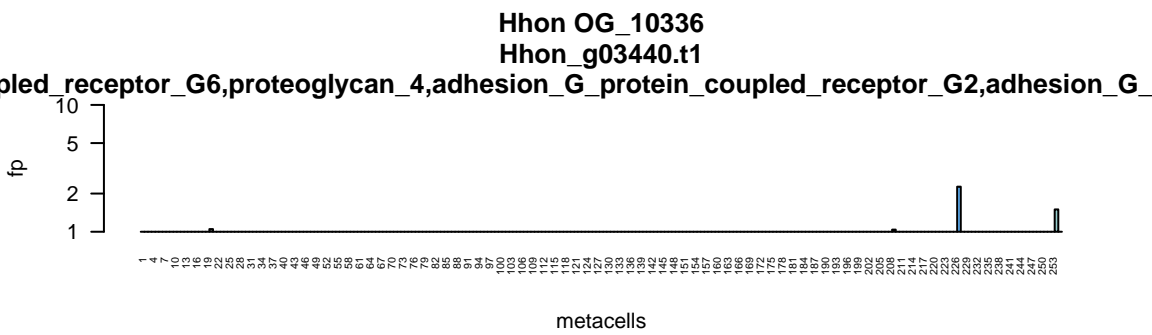
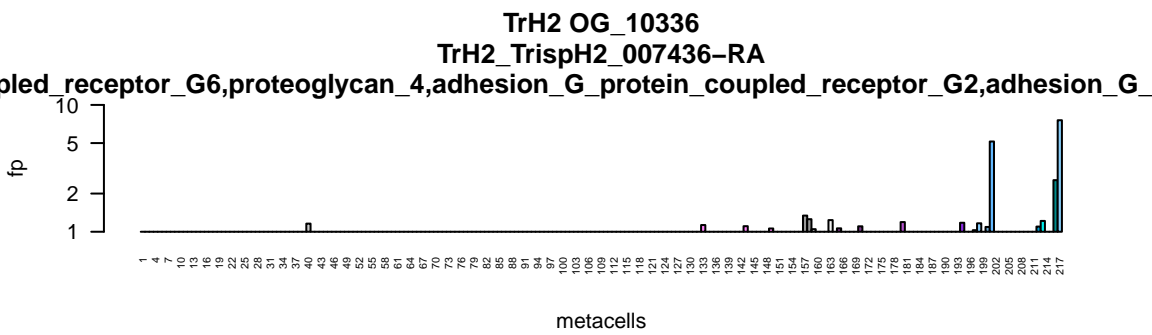
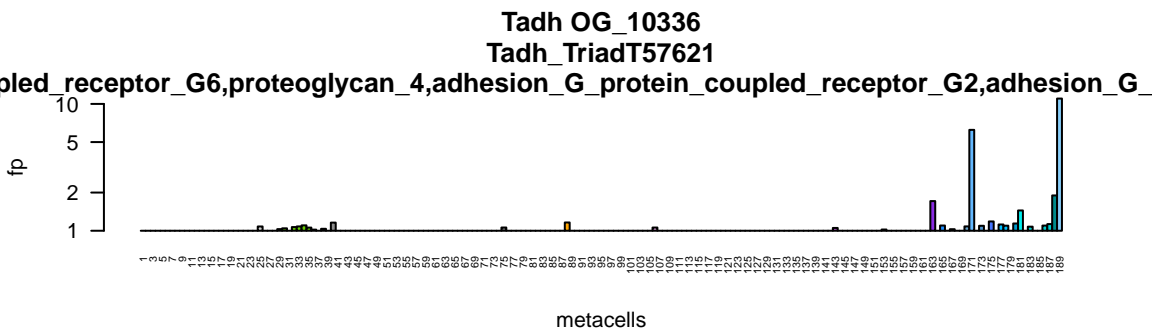


Hhon | no data



HoiH23 OG_10104
HoiH23_PIH23_006092-RA





Tadh_TriadT57663

[illegible]

12 OG 3871

TrH2_TrispH2_004645-RA

on OG 3871

Hhon_g01812.t1

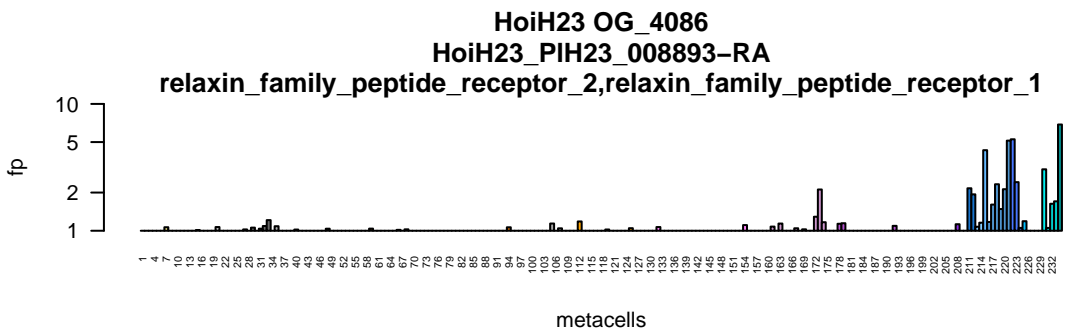
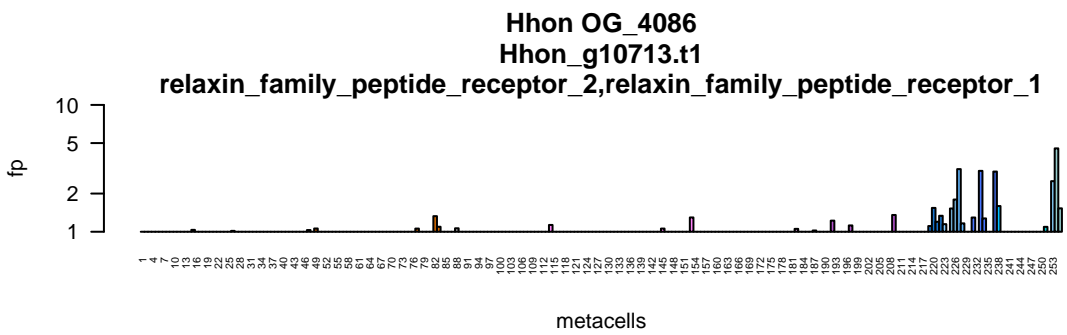
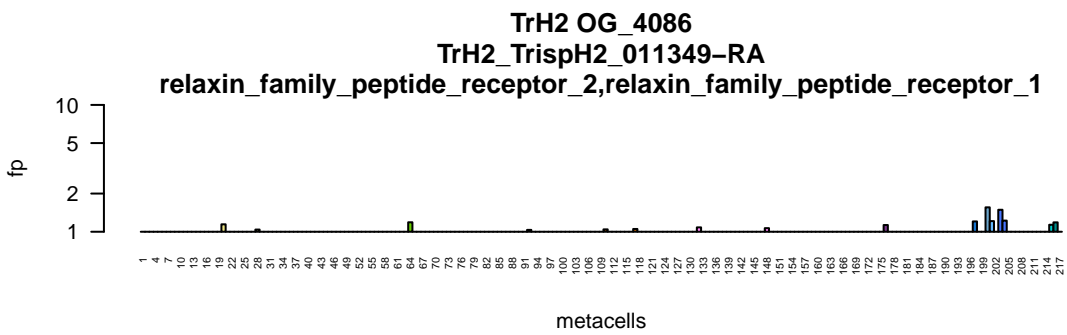
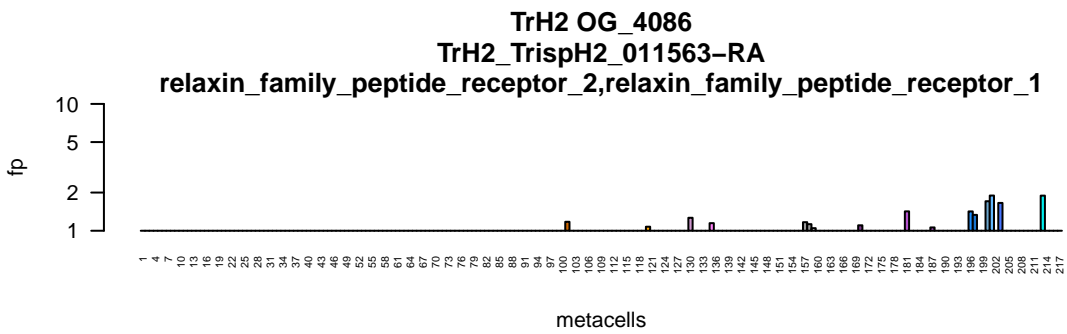
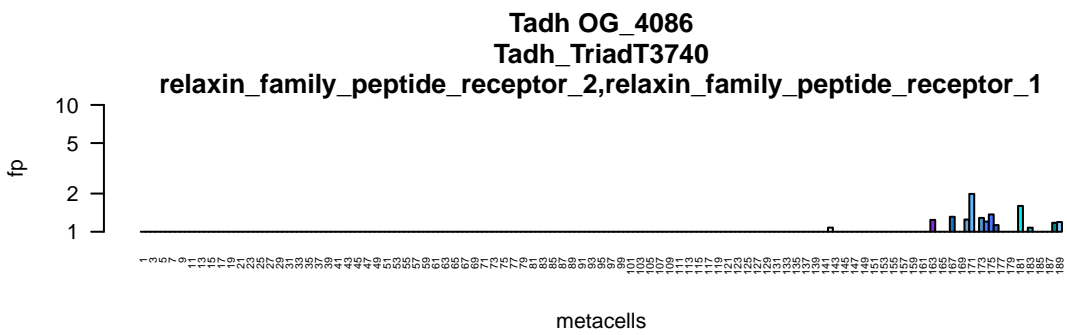
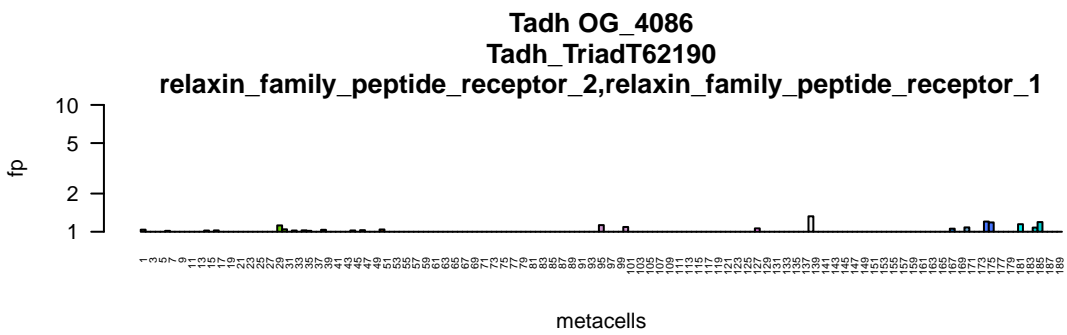
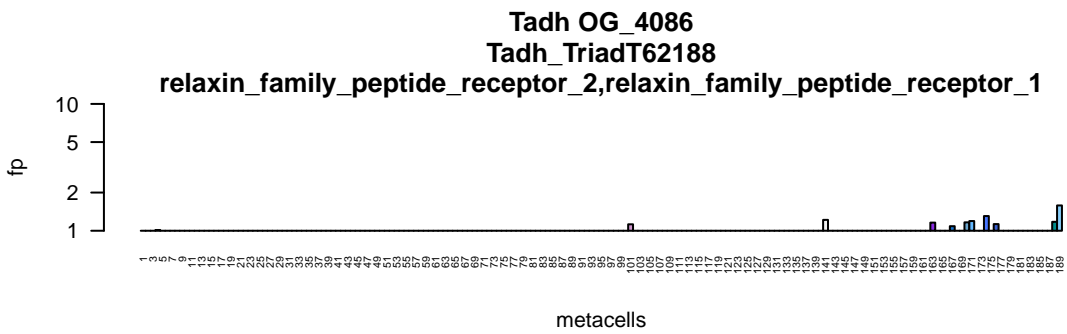
Bar chart showing the frequency of the number of factors (fp) for each variable. The y-axis is labeled fp and ranges from 1 to 10. The x-axis lists 63 variables. Most variables have a frequency of 1, with a few having frequencies of 2 or 3. The variable x_{47} has the highest frequency of 4.

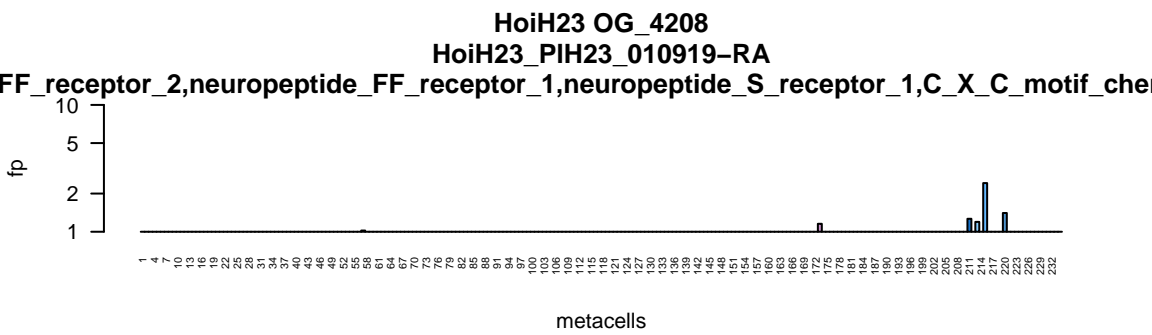
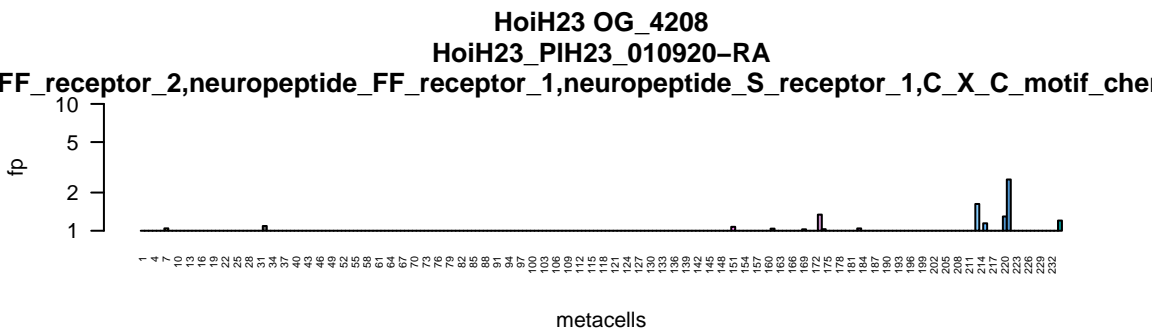
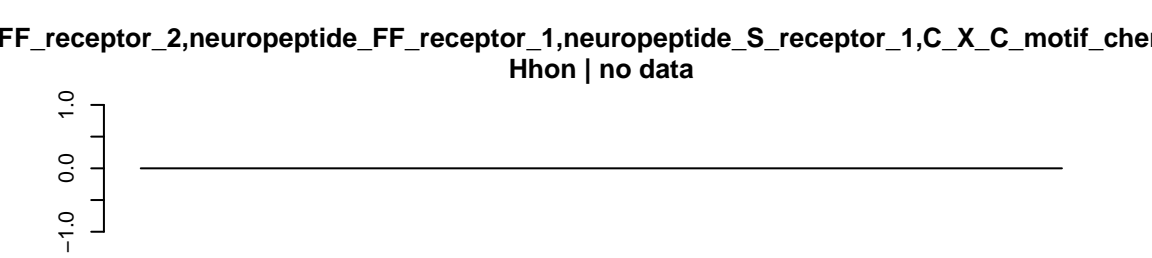
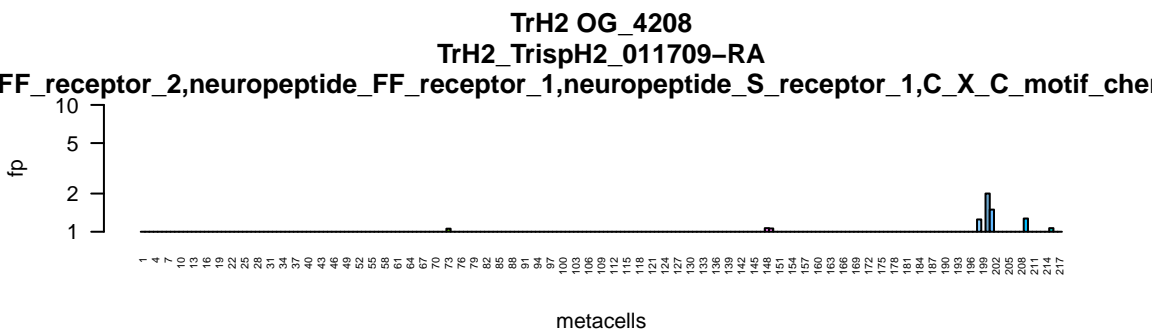
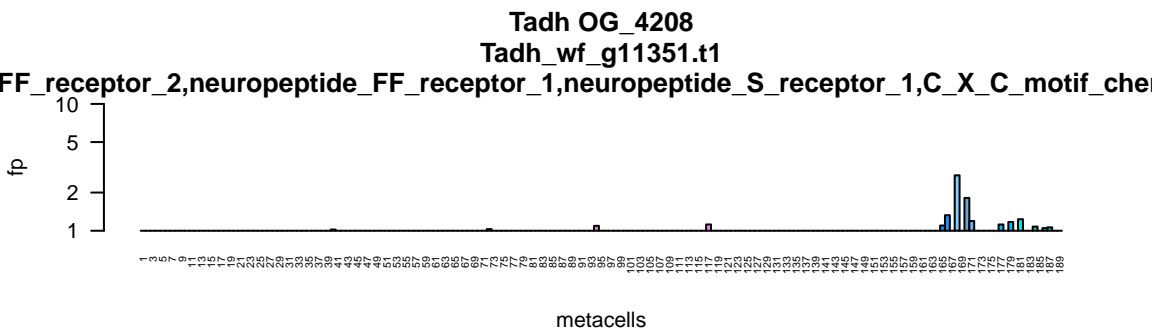
HoiH23 OG 3871

HoiH23_PIH23_009268-RA

Bar chart showing the frequency of the number of features (fp) for each variable. The y-axis is labeled fp and ranges from 1 to 10. The x-axis lists 63 variables. Most variables have a frequency of 1. Variables 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63 have frequencies greater than 1, with some reaching up to 10.

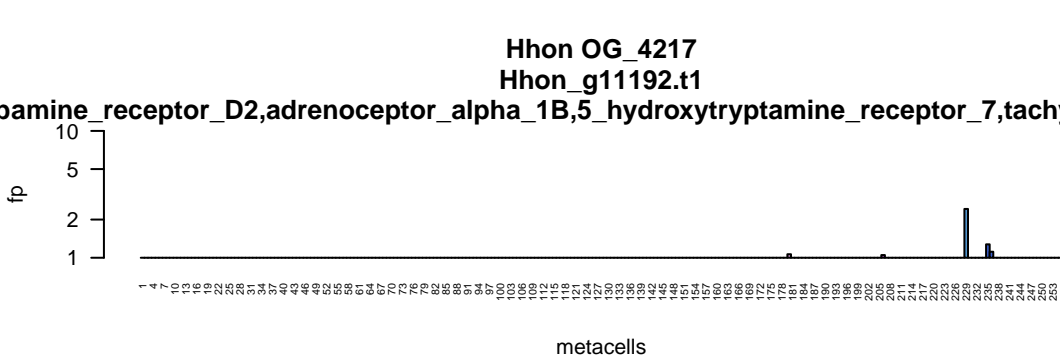
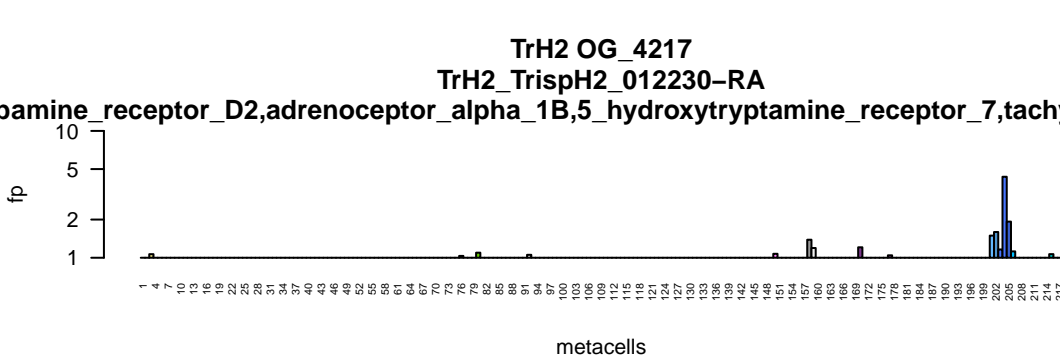
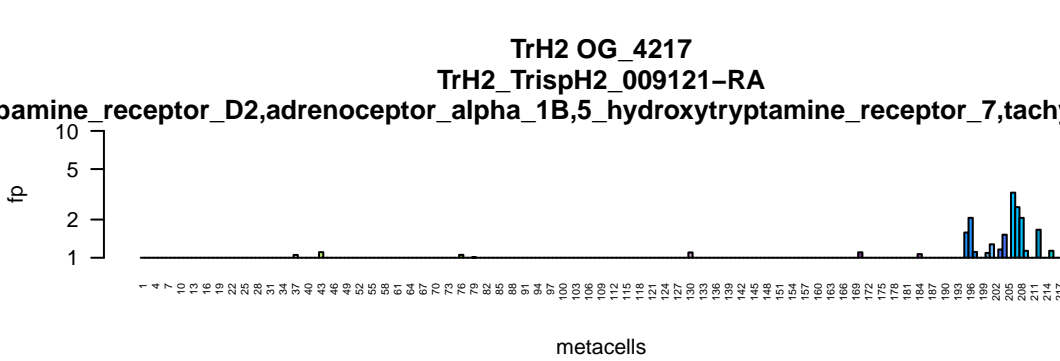
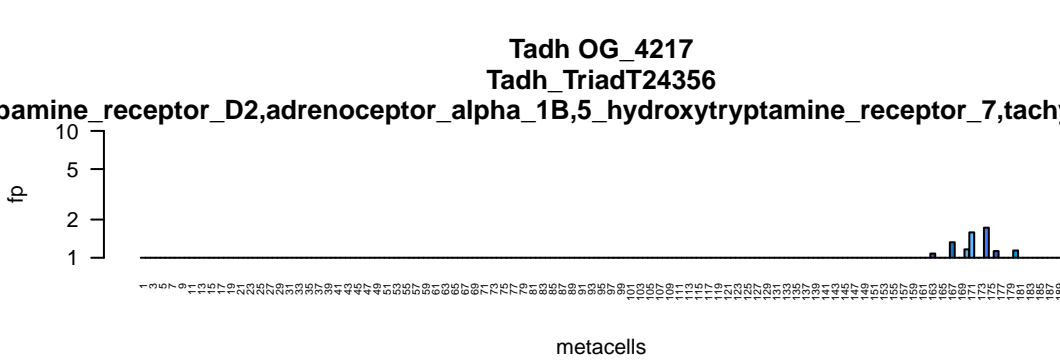
metacells



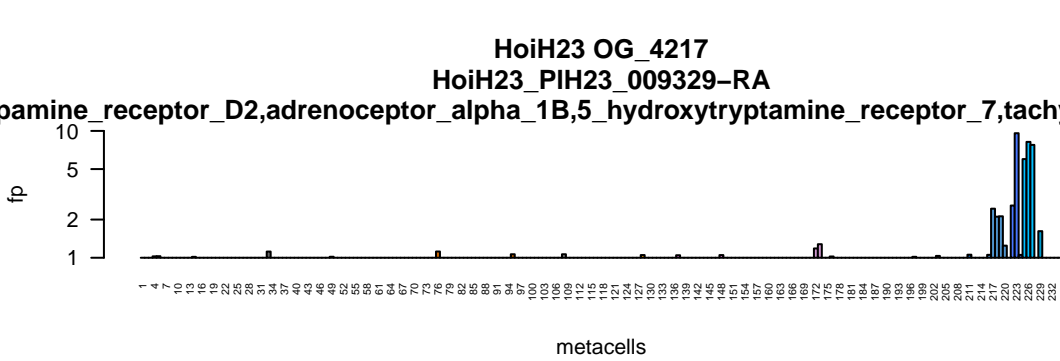


Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists 100 metacells. Most metacells have 0 false positives, but a few have 1, and a small cluster of metacells (173-187) has 2, 3, or 4 false positives.

metacell	fp
1	0
2	0
3	0
4	0
5	0
6	0
7	0
8	0
9	0
10	0
11	0
12	0
13	0
14	0
15	0
16	0
17	0
18	0
19	0
20	0
21	0
22	0
23	0
24	0
25	0
26	0
27	0
28	0
29	0
30	0
31	0
32	0
33	0
34	0
35	0
36	0
37	0
38	0
39	0
40	0
41	0
42	0
43	0
44	0
45	0
46	0
47	0
48	0
49	0
50	0
51	0
52	0
53	0
54	0
55	0
56	0
57	0
58	0
59	0
60	0
61	0
62	0
63	0
64	0
65	0
66	0
67	0
68	0
69	0
70	0
71	0
72	0
73	0
74	0
75	0
76	0
77	0
78	0
79	0
80	0
81	0
82	0
83	0
84	0
85	0
86	0
87	0
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119	0
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125	0
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130	0
131	0
132	0
133	0
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137	0
138	0
139	0
140	0
141	0
142	0
143	0
144	0
145	0
146	0
147	0
148	0
149	0
150	0
151	0
152	0
153	0
154	0
155	0
156	0
157	0
158	0
159	0
160	0
161	0
162	0
163	0
164	0
165	0
166	0
167	0
168	0
169	0
170	0
171	0
172	0
173	1
174	2
175	2
176	3
177	4
178	3
179	2
180	1
181	1
182	1
183	1
184	1
185	1
186	1
187	1
188	1
189	1
190	1



Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 232. Most metacells have a false positive count of 1. Metacells 169, 170, 171, 172, 173, 174, 175, 176, 177, 178, 179, 180, 181, 182, 183, 184, 185, 186, 187, 188, 189, 190, 191, 192, 193, 194, 195, 196, 197, 198, 199, 200, 201, 202, 203, 204, 205, 206, 207, 208, 209, 210, 211, 212, 213, 214, 215, 216, 217, 218, 219, 220, 221, 222, 223, 224, 225, 226, 227, 228, 229, 230, 231, 232 have higher false positive counts, with some reaching up to 10.



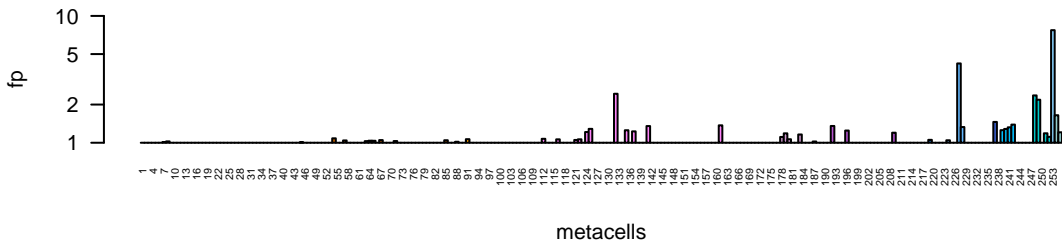
Tadh | no data



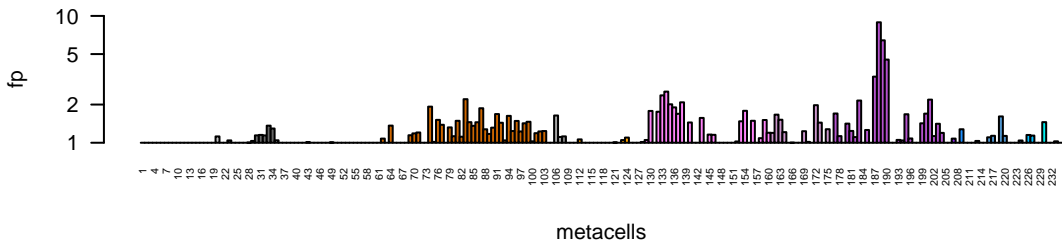
TrH2 | no data



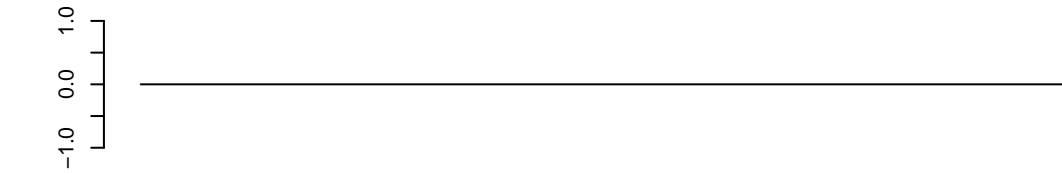
Hhon OG_4260
Hhon_g04196.t1



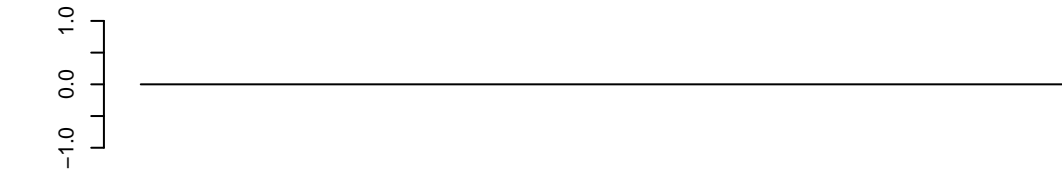
HoiH23 OG_4260
HoiH23_PIH23_009314-RA



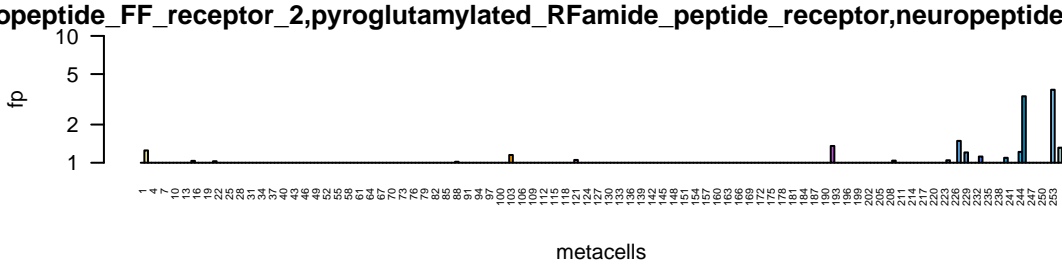
peptide_FF_receptor_2,pyroglutamylated_RFamide_peptide_receptor,neuropeptide_Y_rec
Tadh | no data



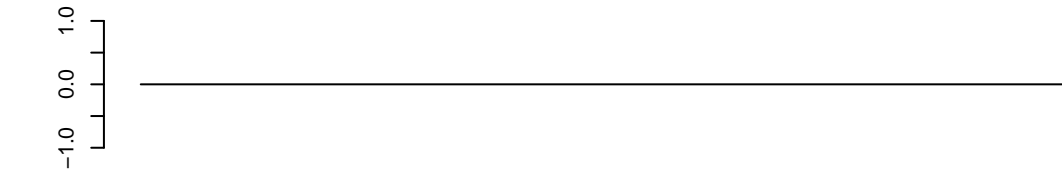
peptide_FF_receptor_2,pyroglutamylated_RFamide_peptide_receptor,neuropeptide_Y_rec
TrH2 | no data

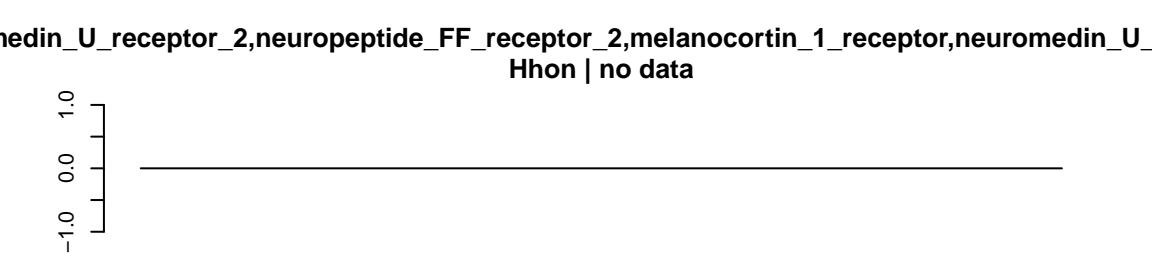
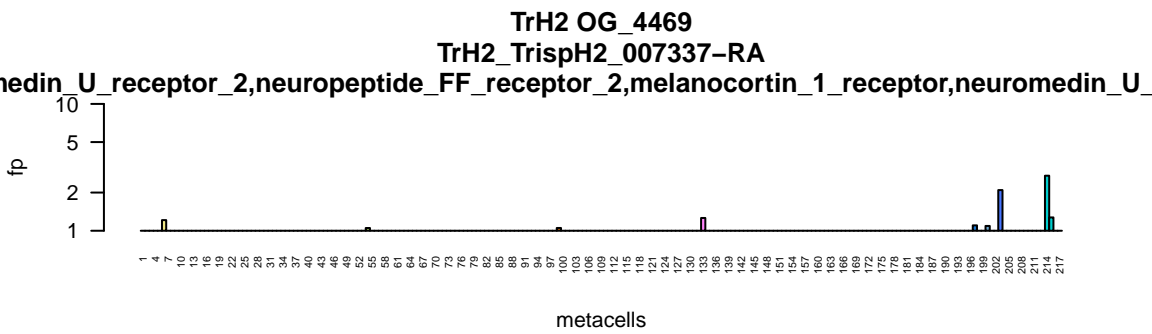
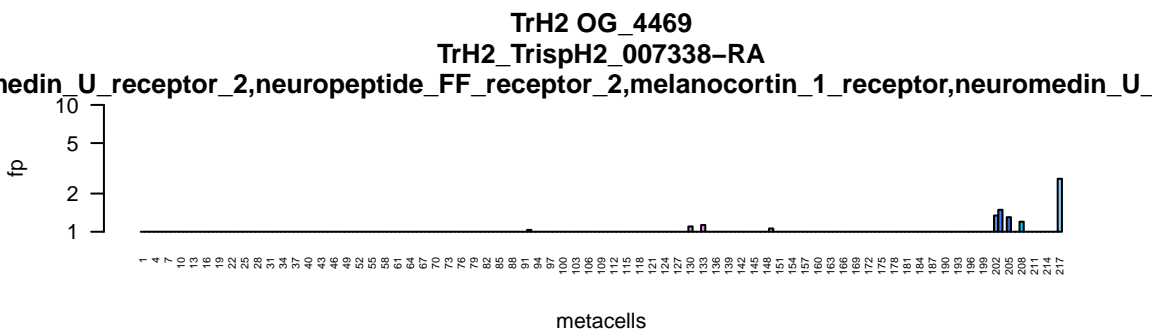
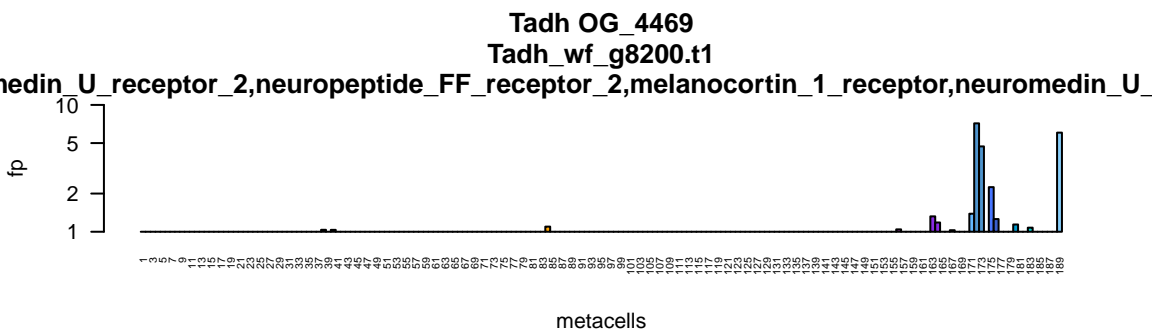
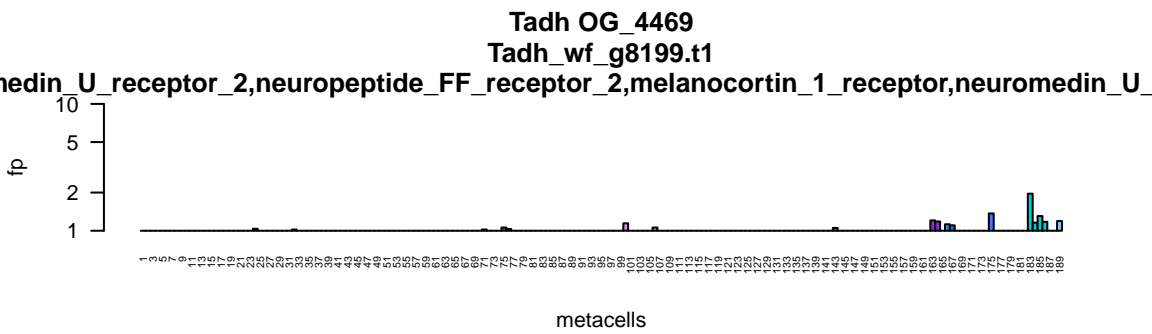


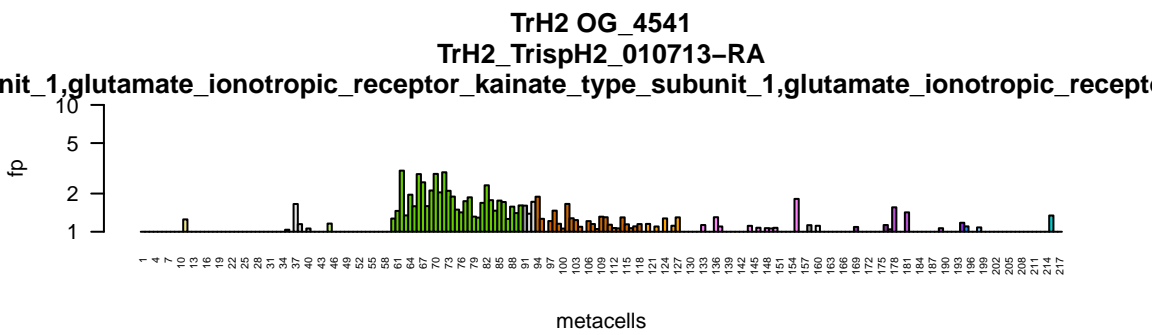
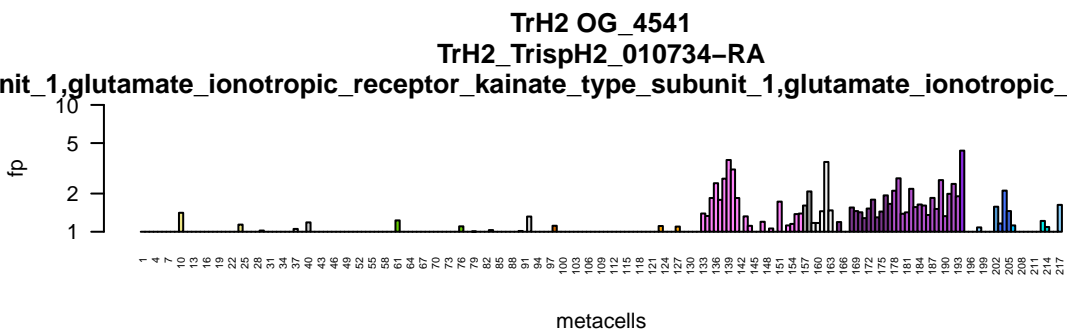
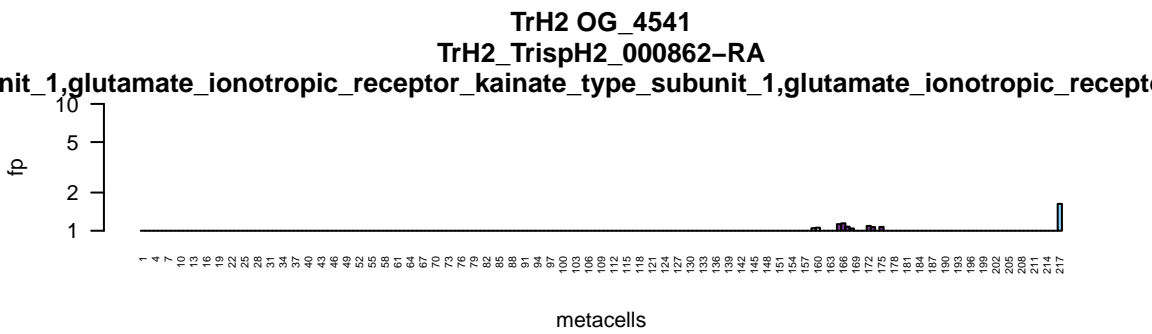
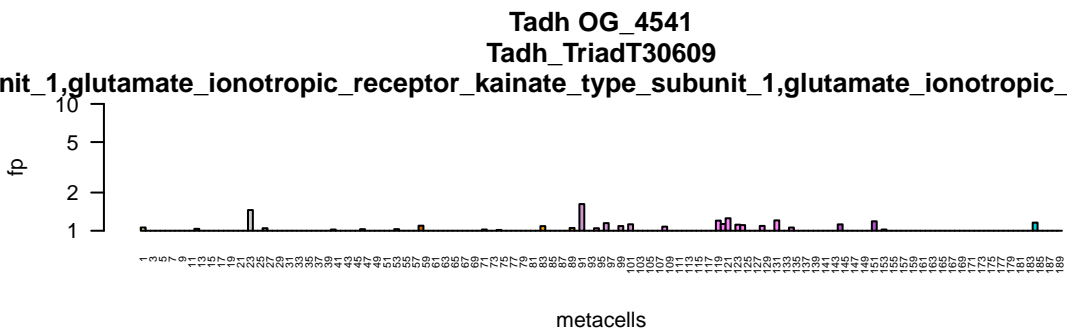
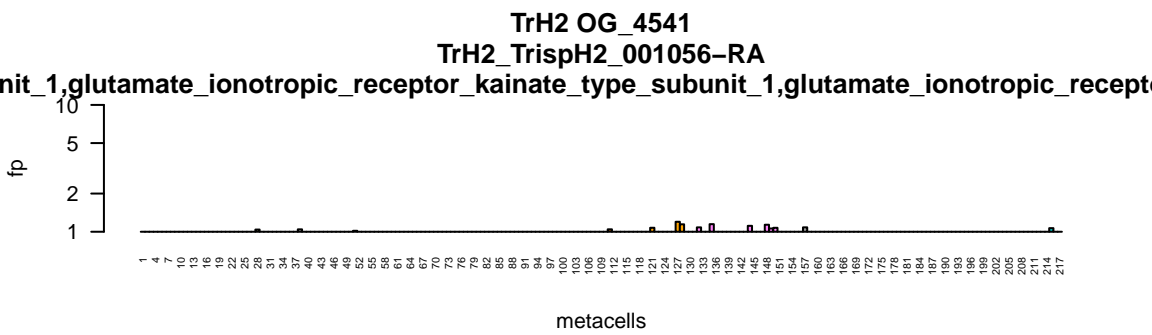
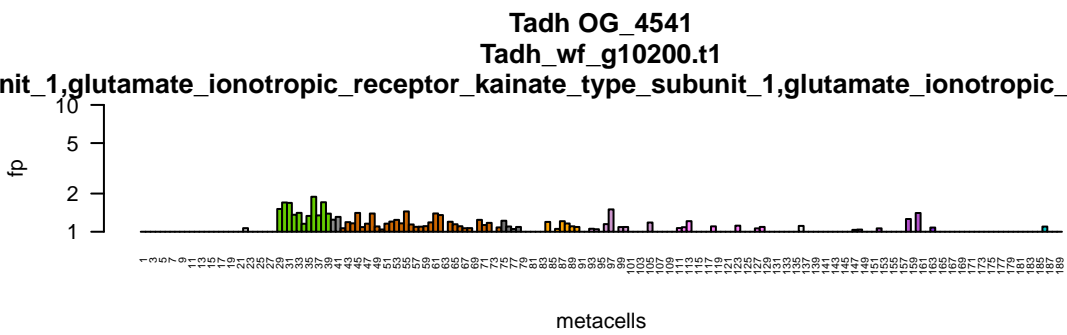
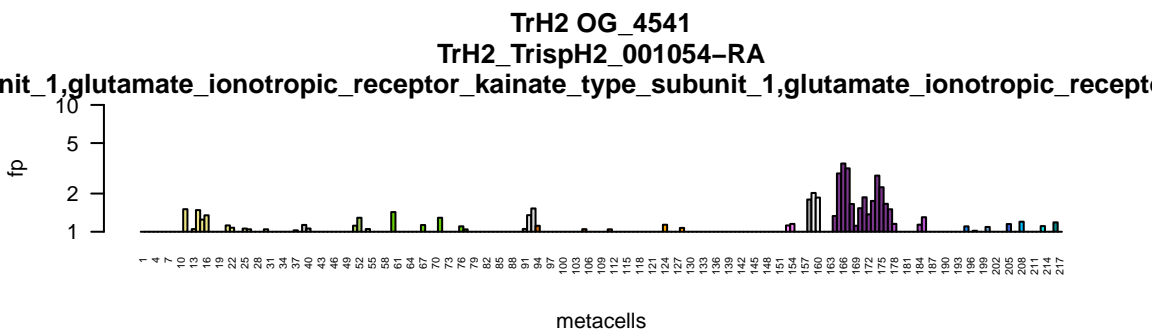
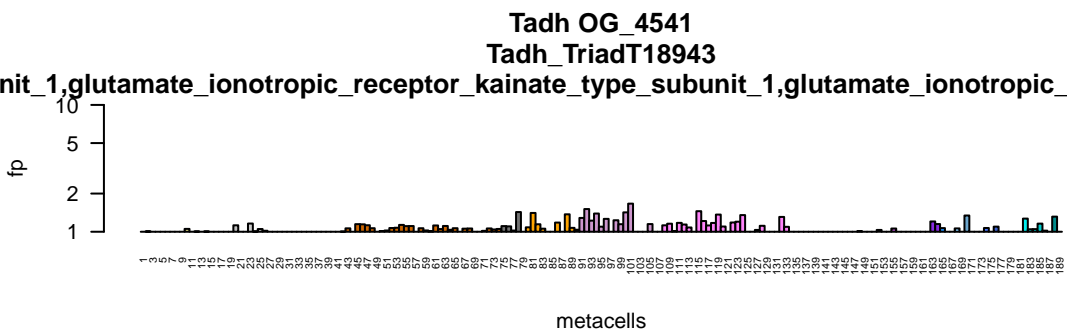
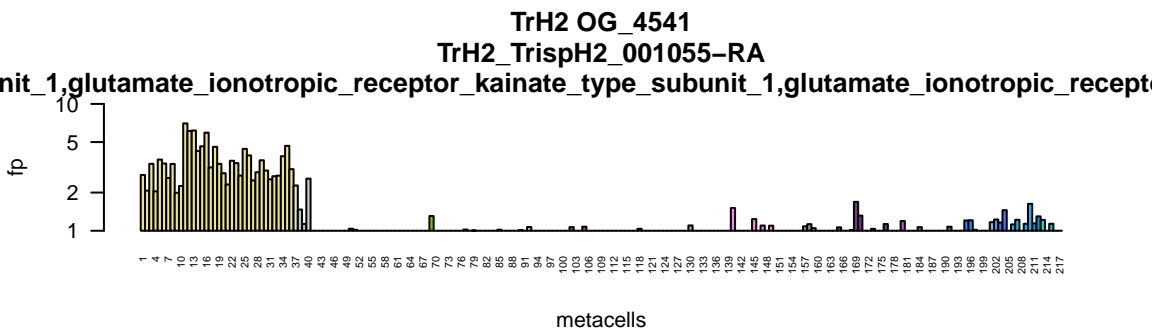
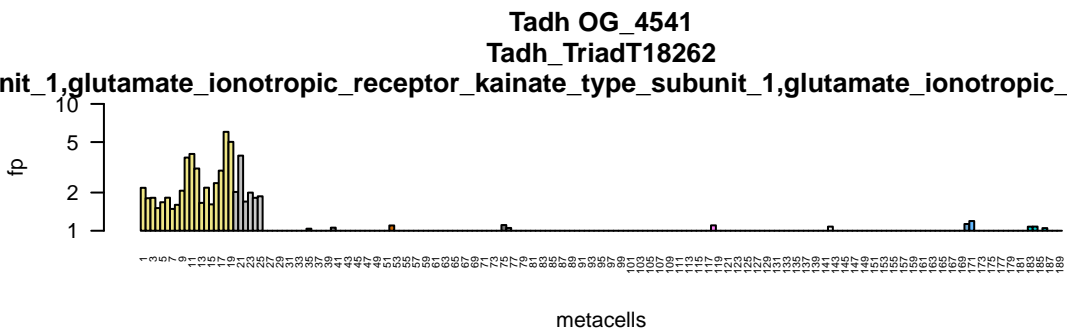
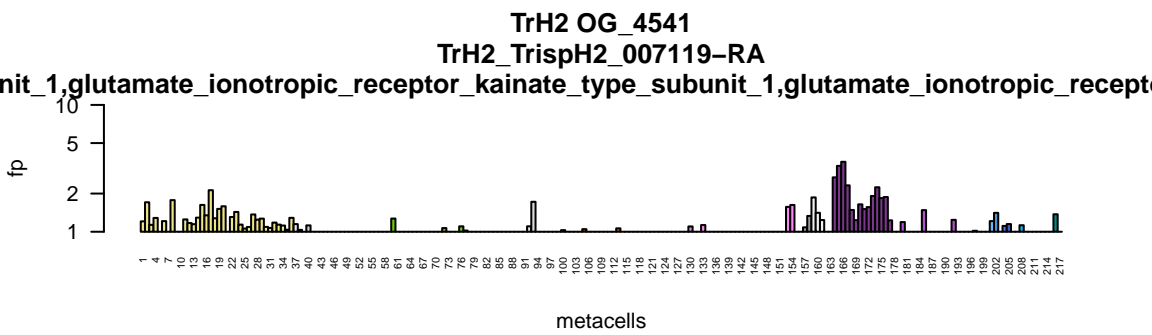
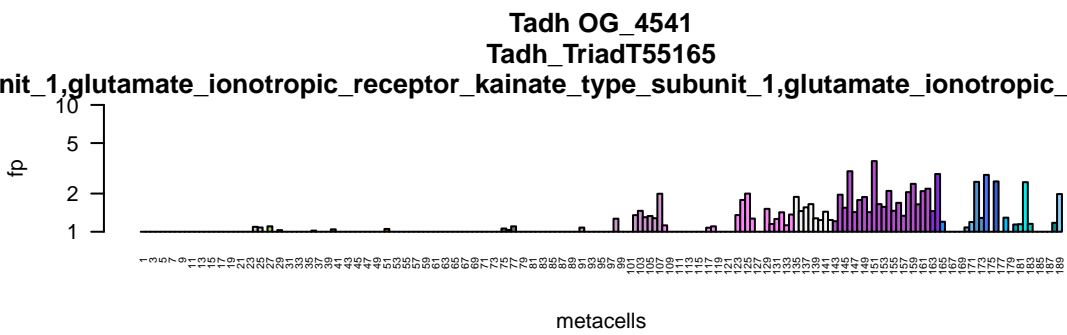
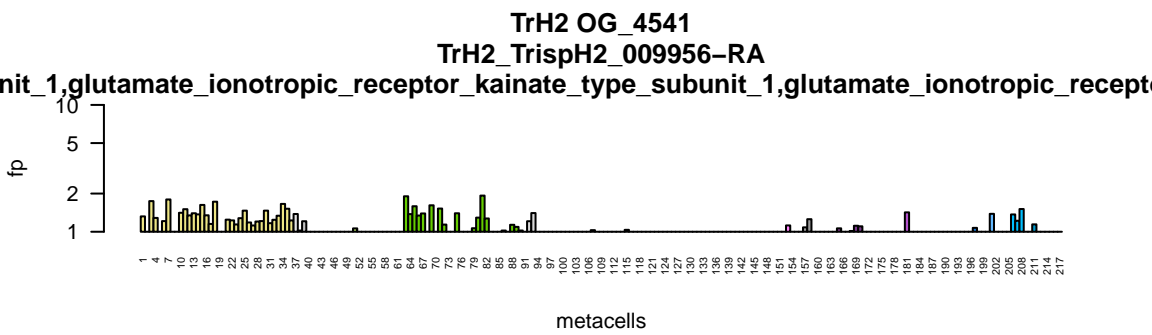
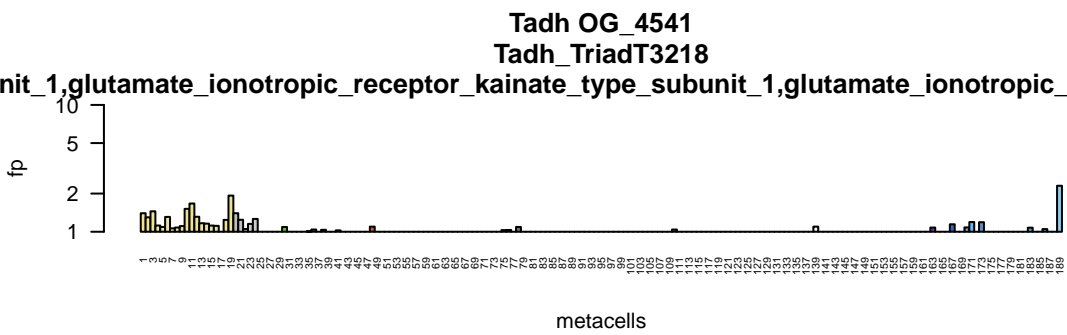
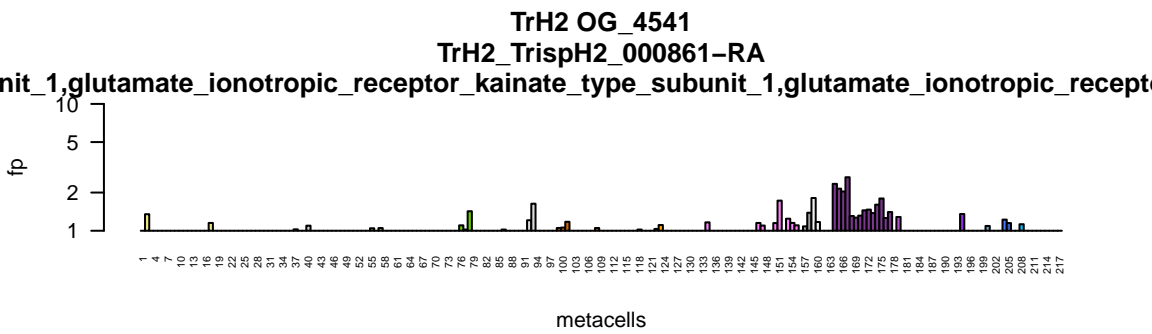
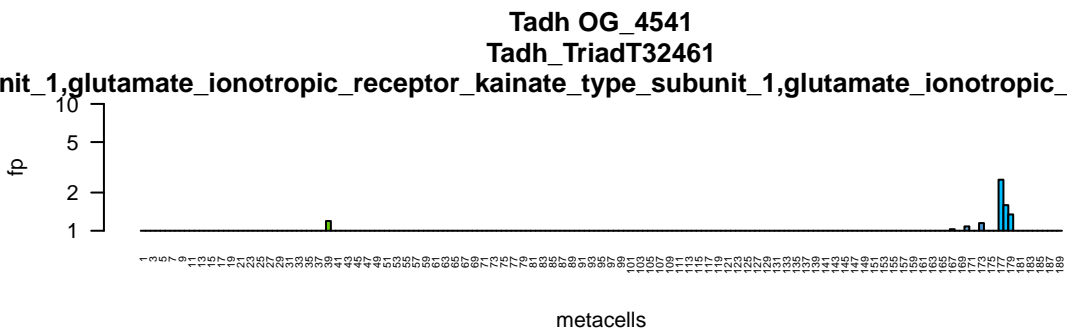
Hhon OG_4269
Hhon_g06774.t1
peptide_FF_receptor_2,pyroglutamylated_RFamide_peptide_receptor,neuropeptide_Y_rec

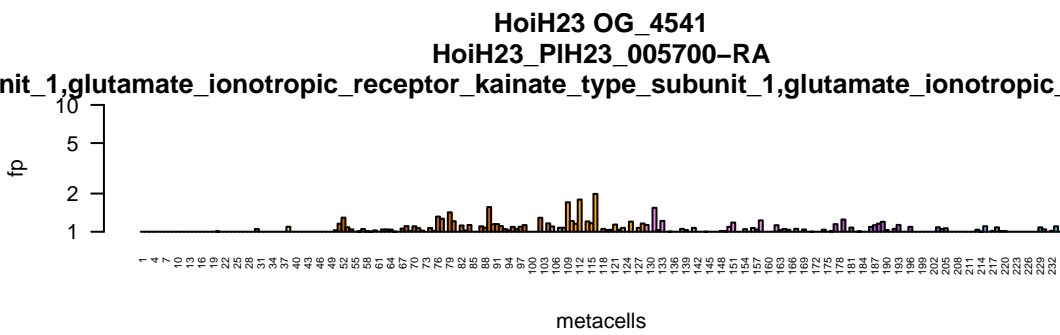
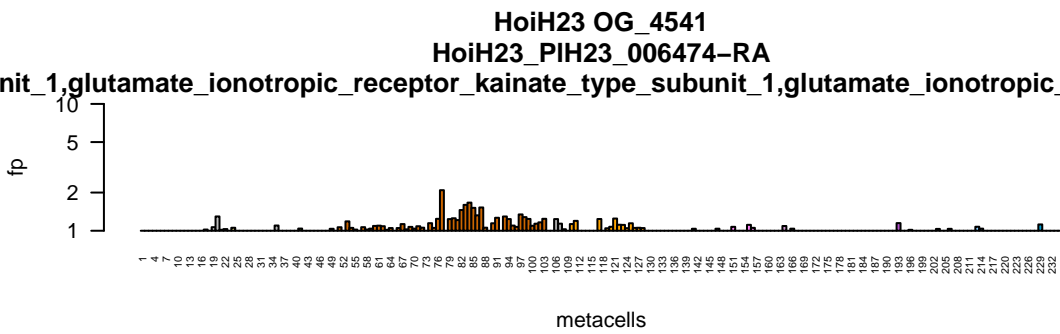
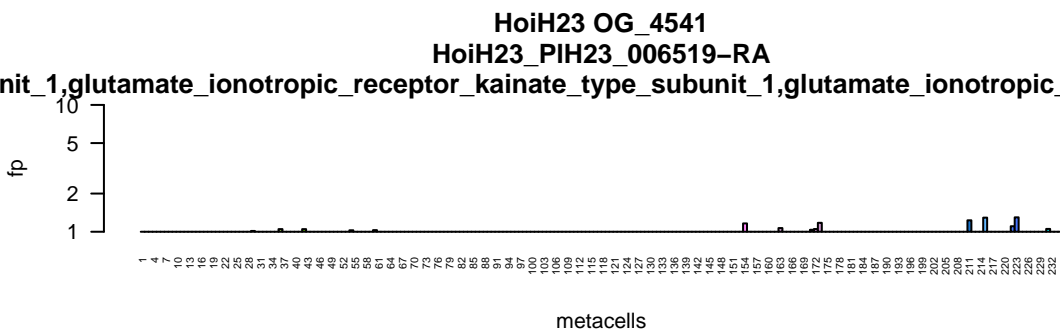
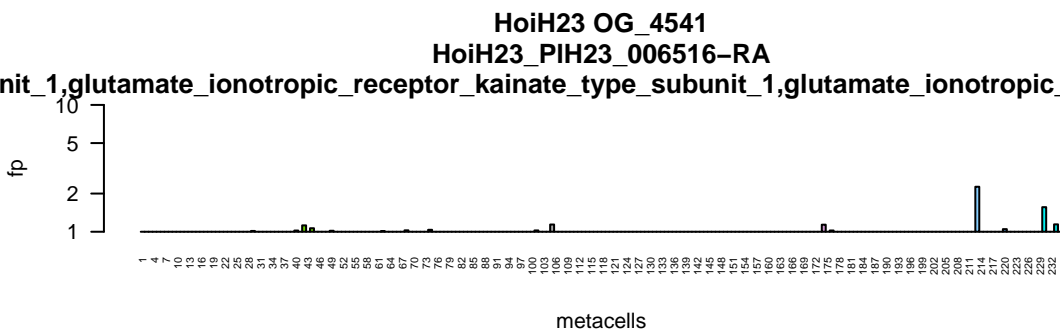
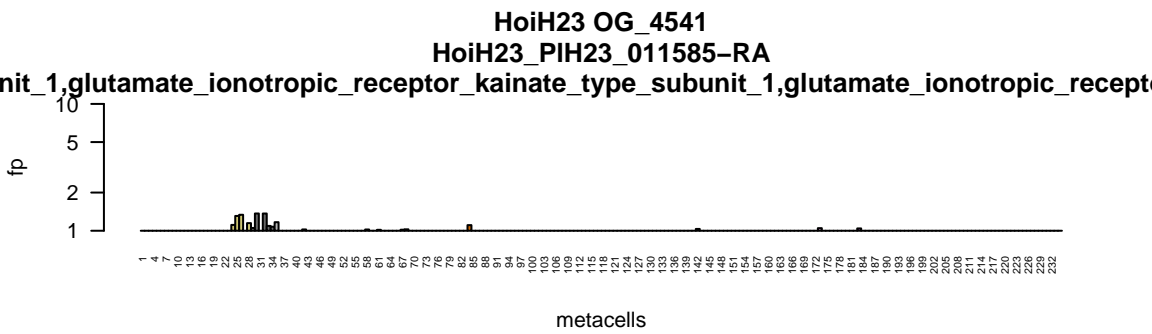
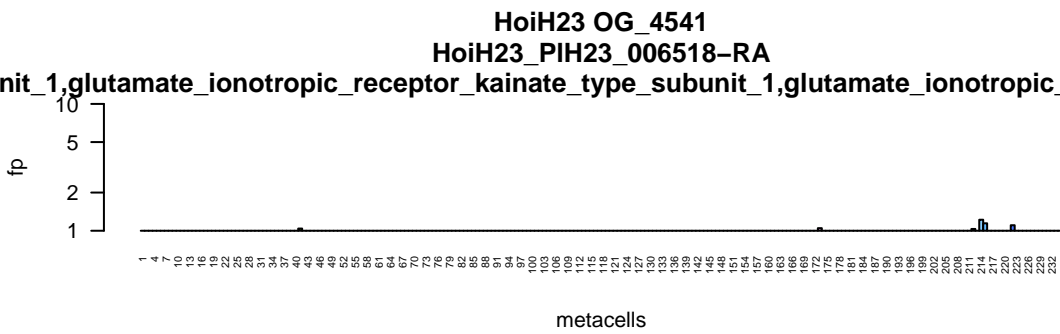
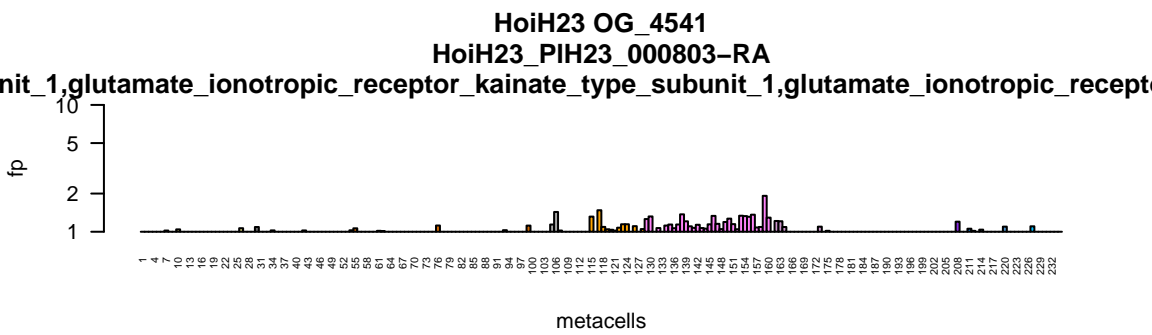
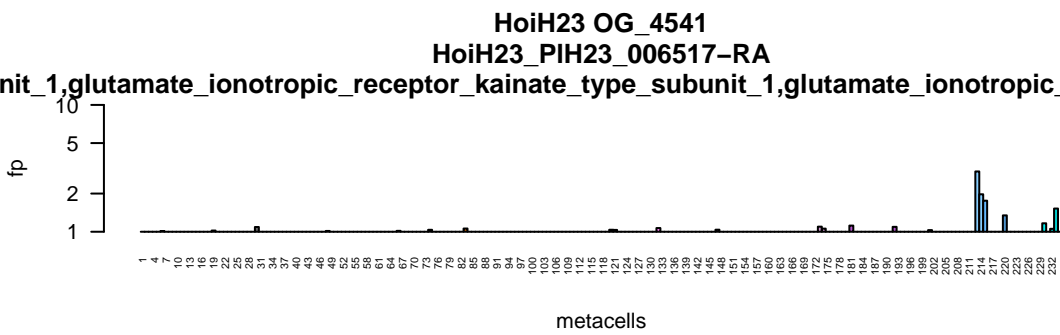
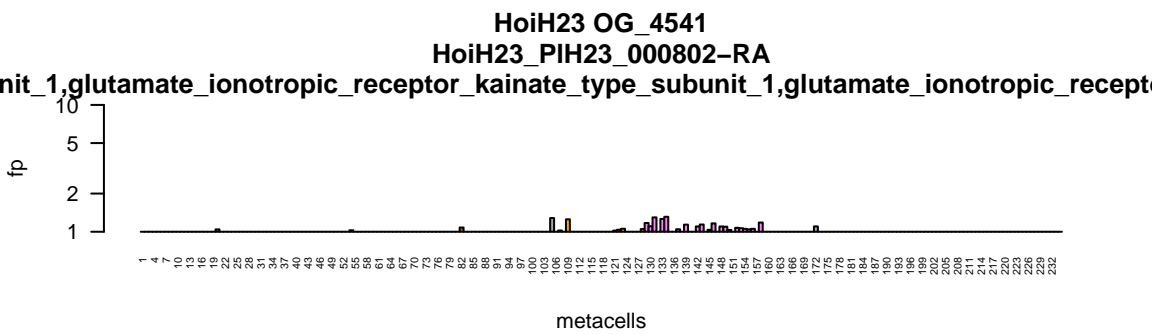
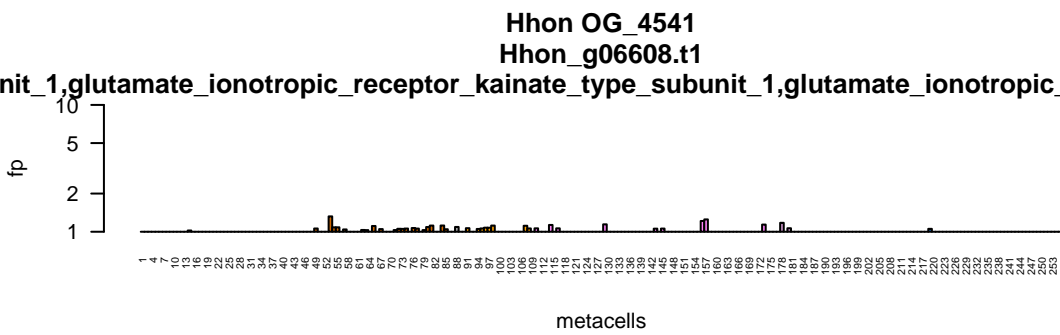
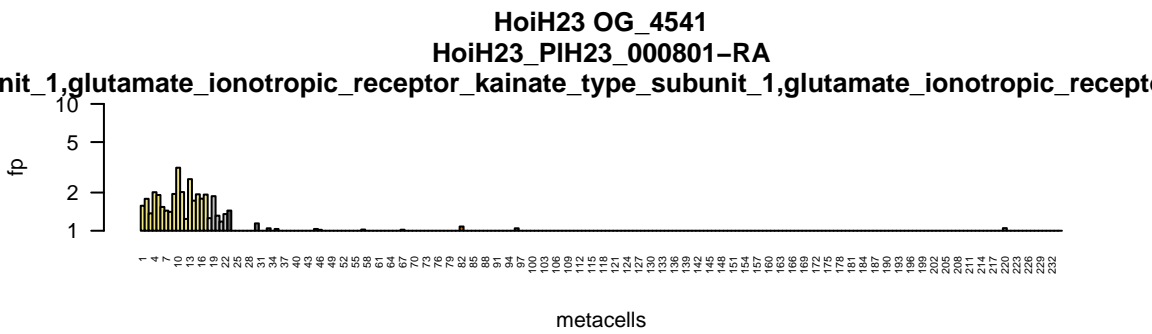
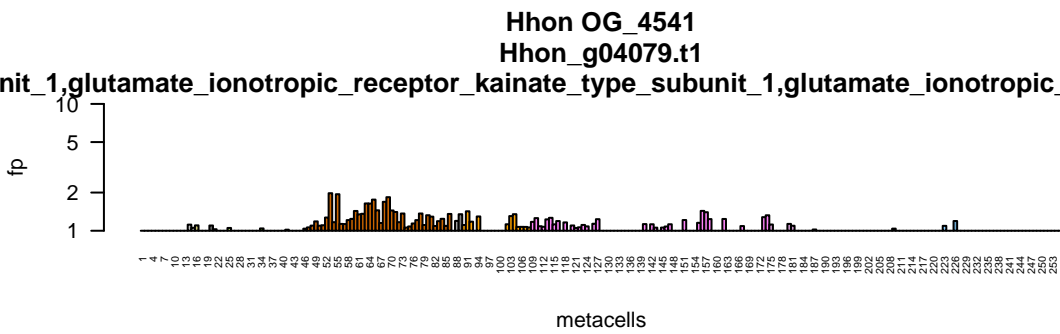


peptide_FF_receptor_2,pyroglutamylated_RFamide_peptide_receptor,neuropeptide_Y_rec
HoiH23 | no data

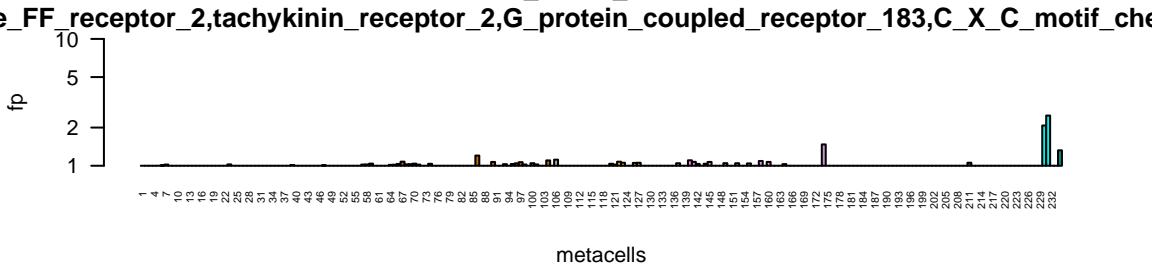


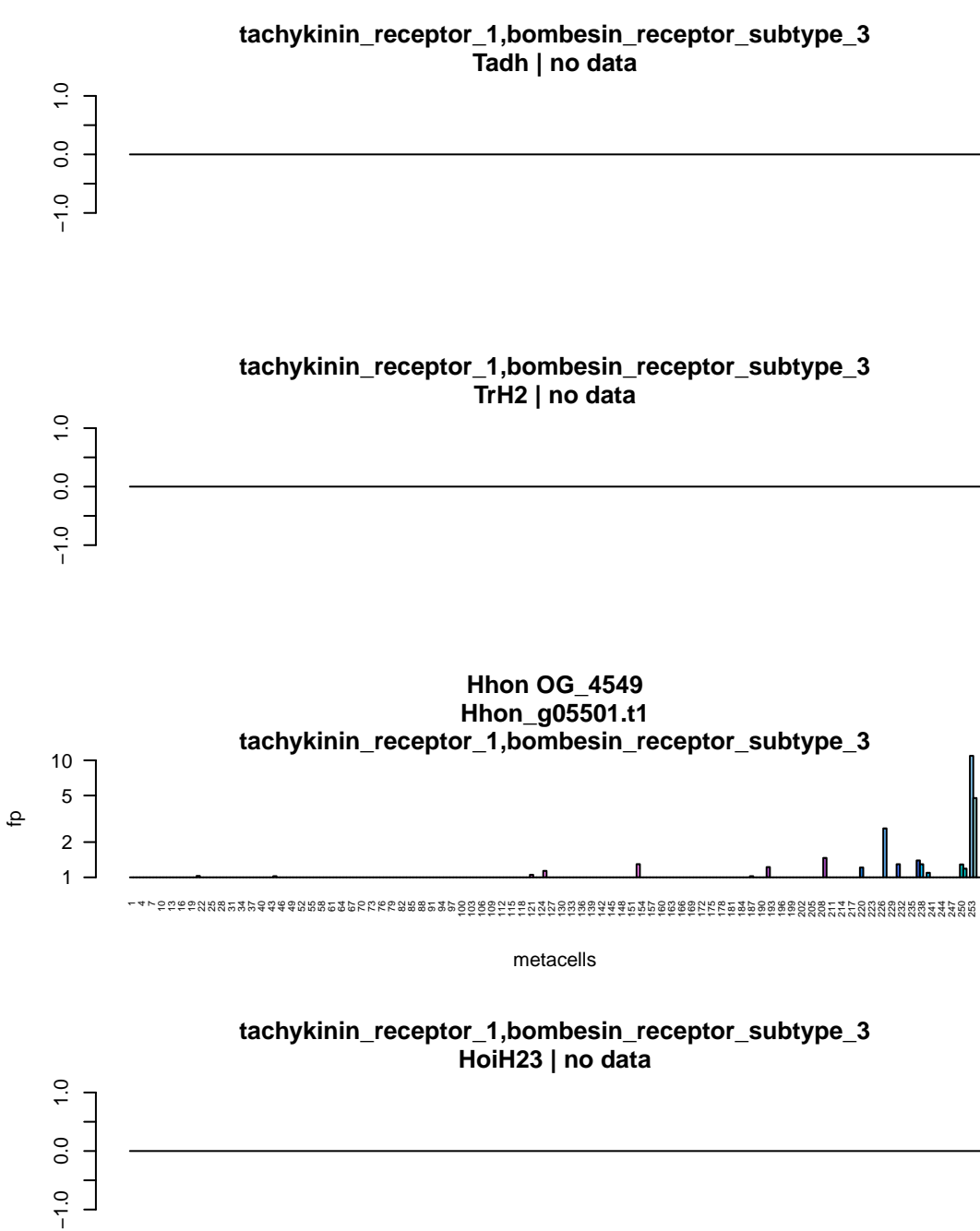


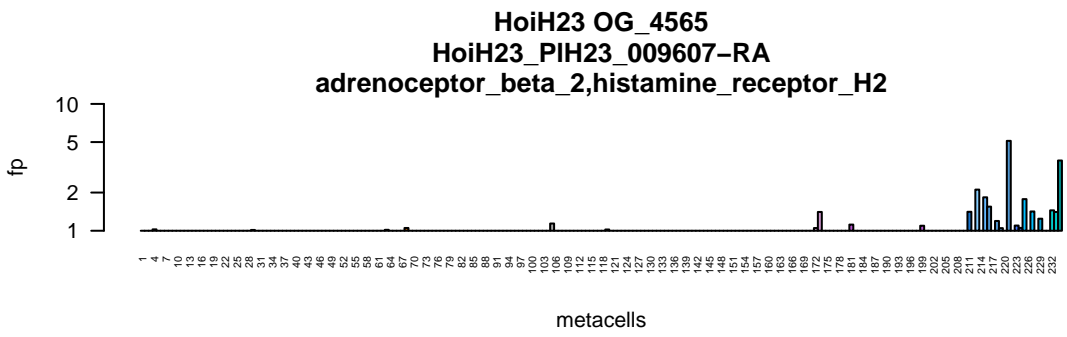
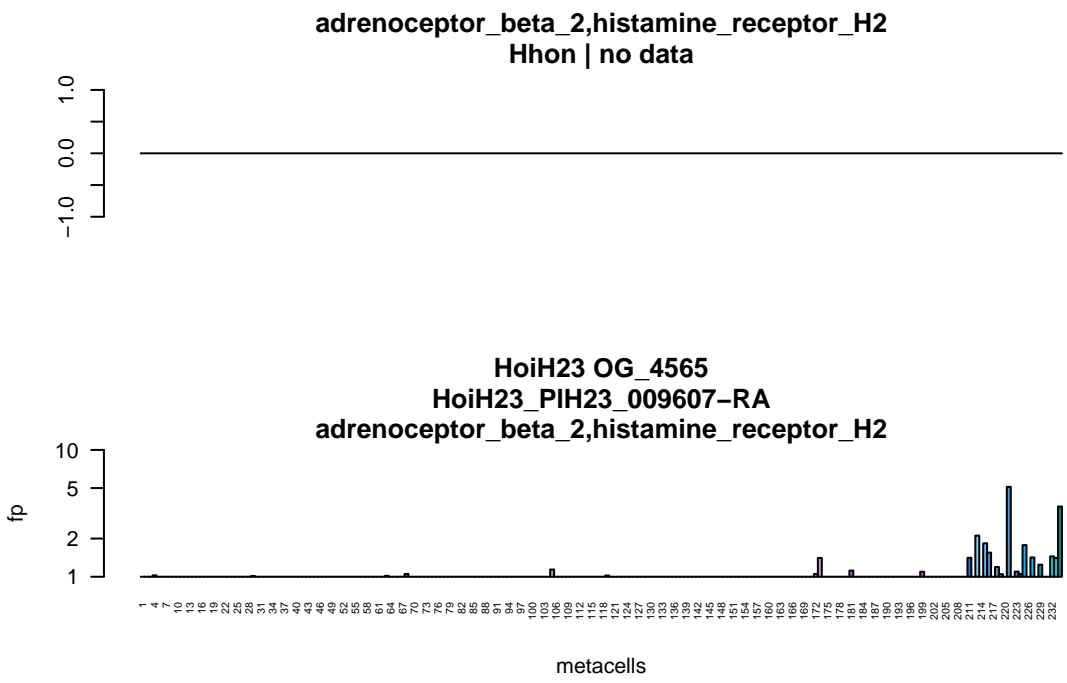
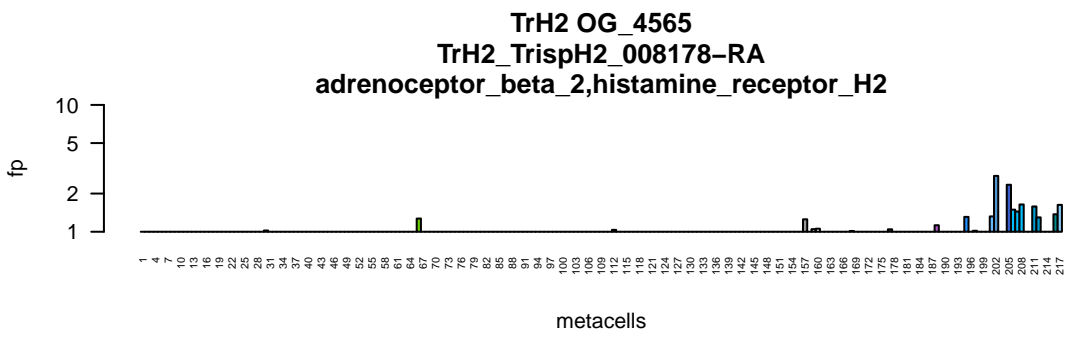
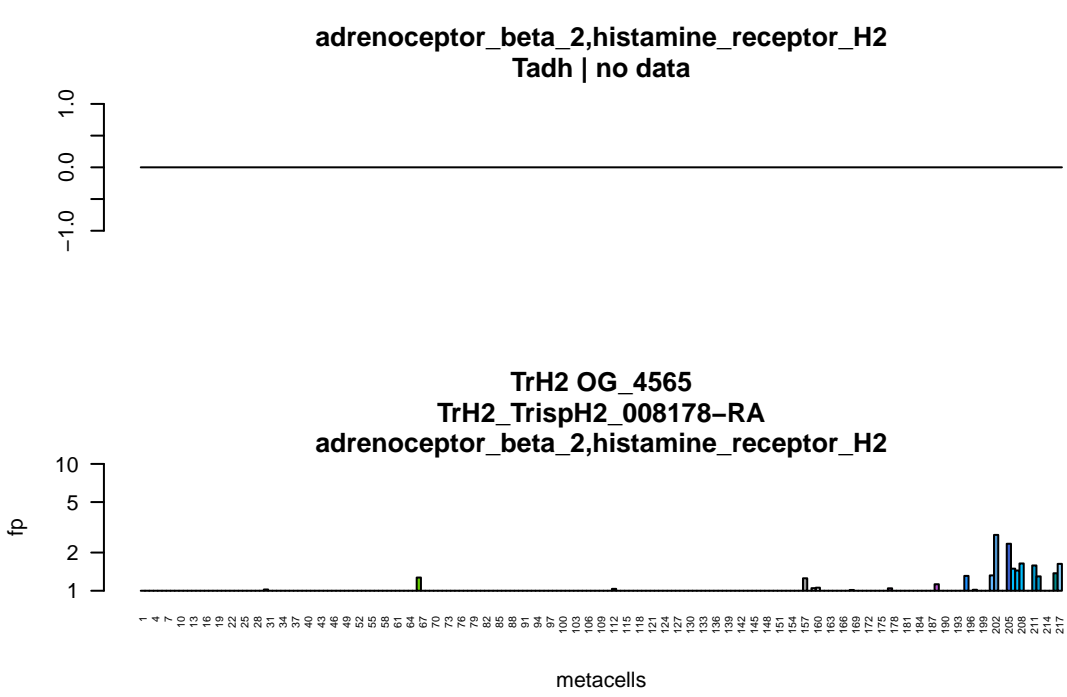


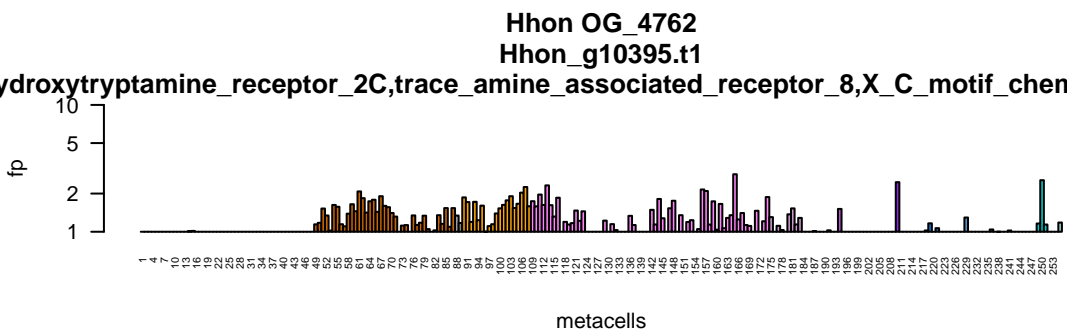
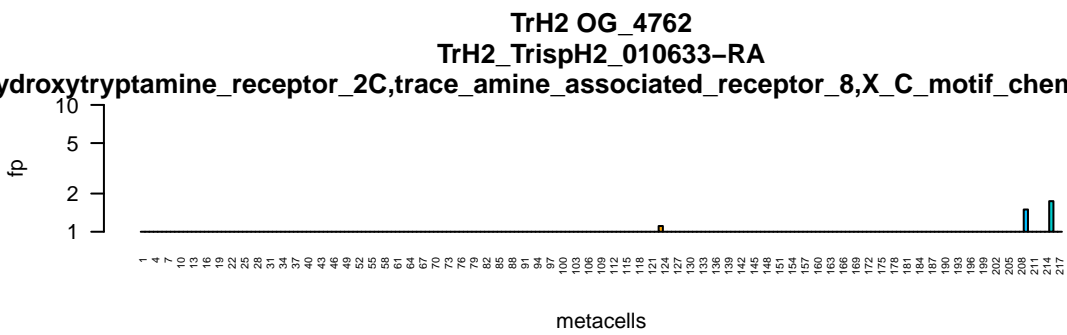
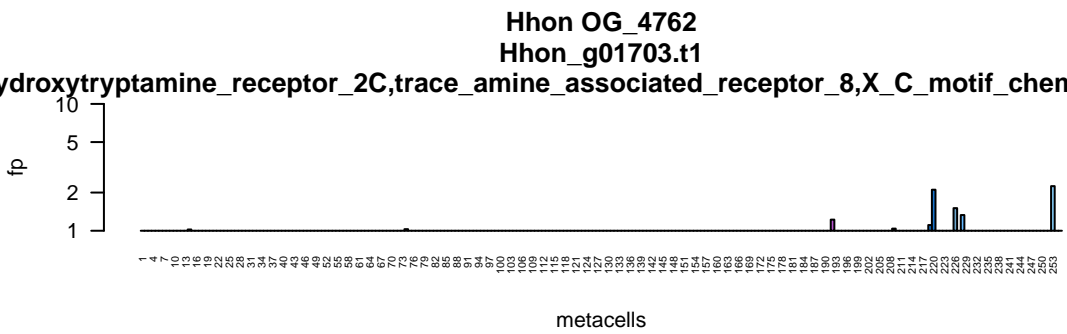
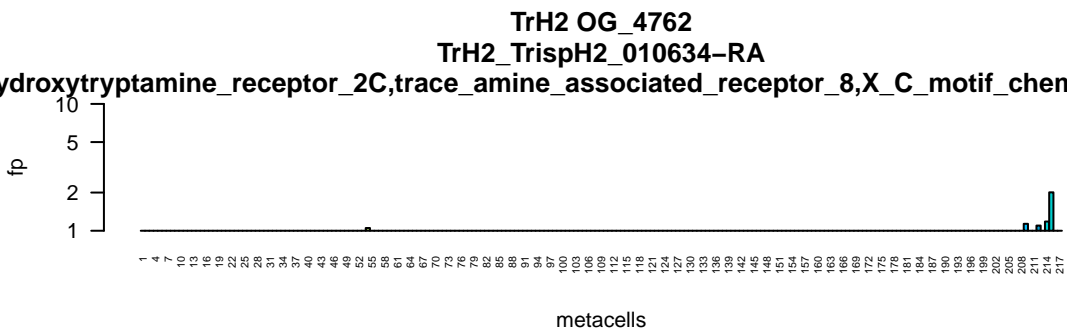
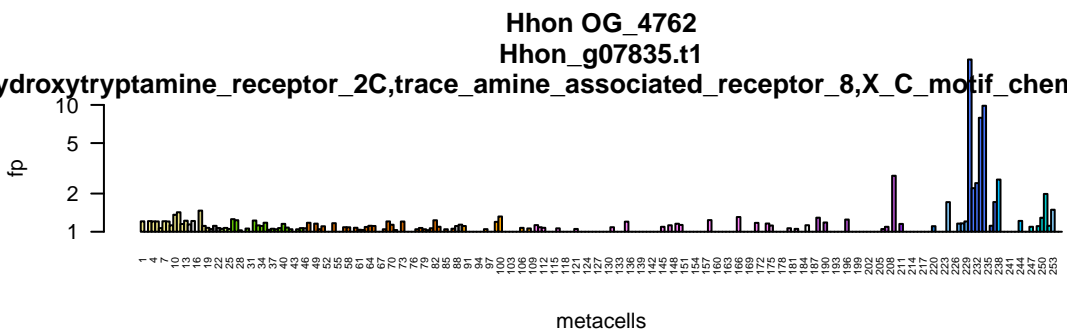
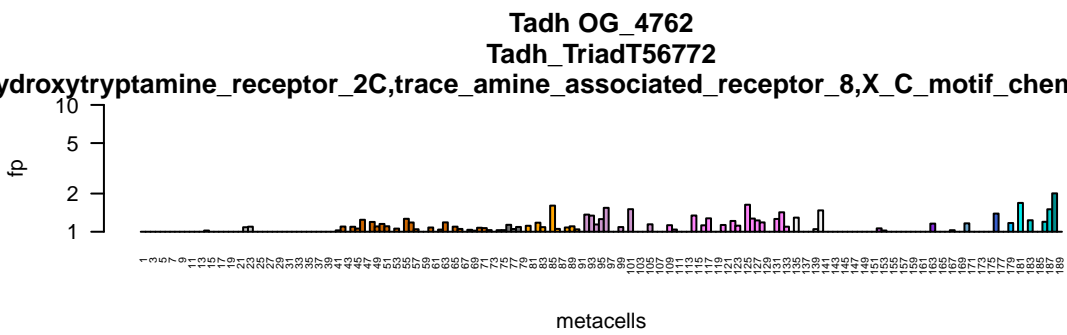
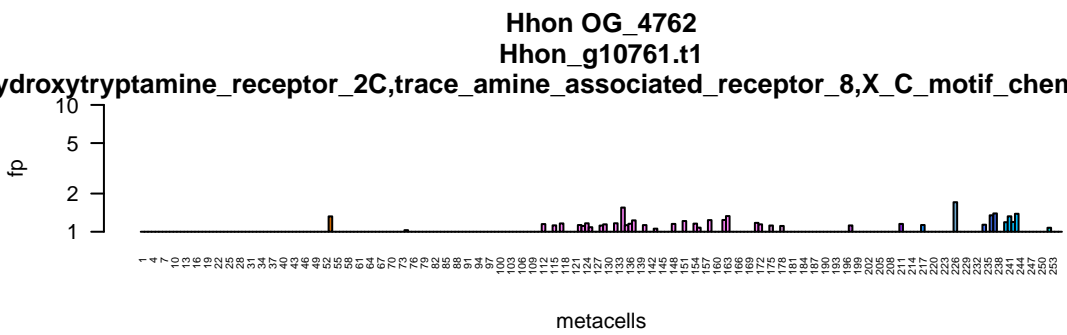
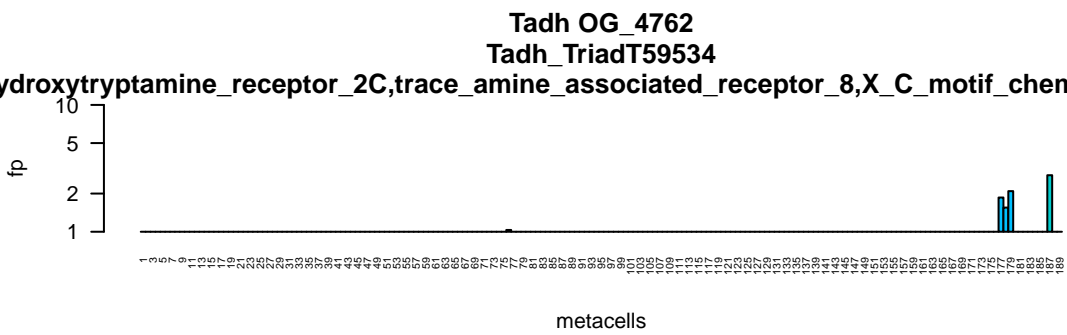
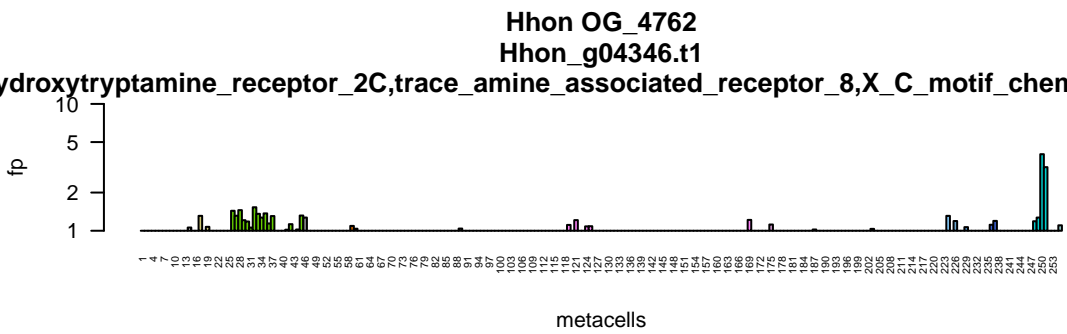
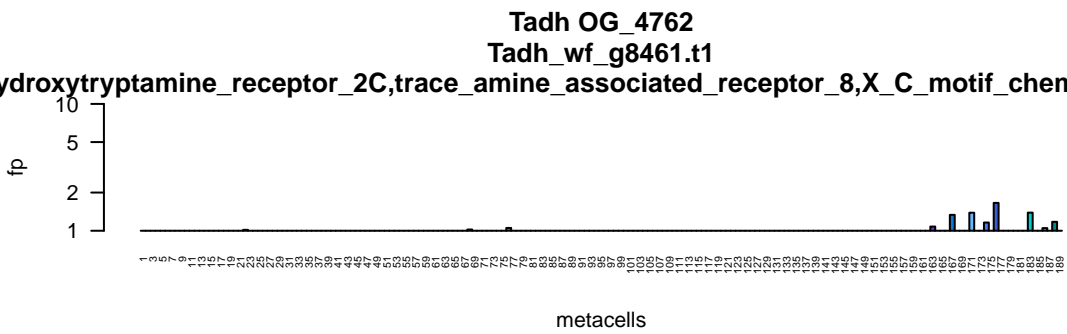
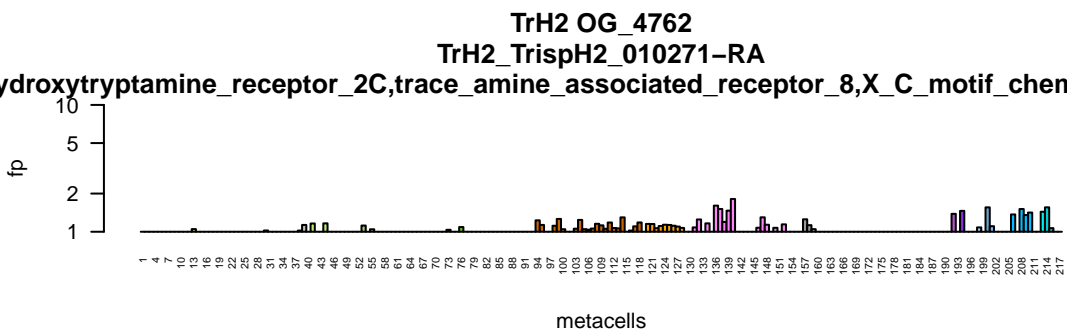
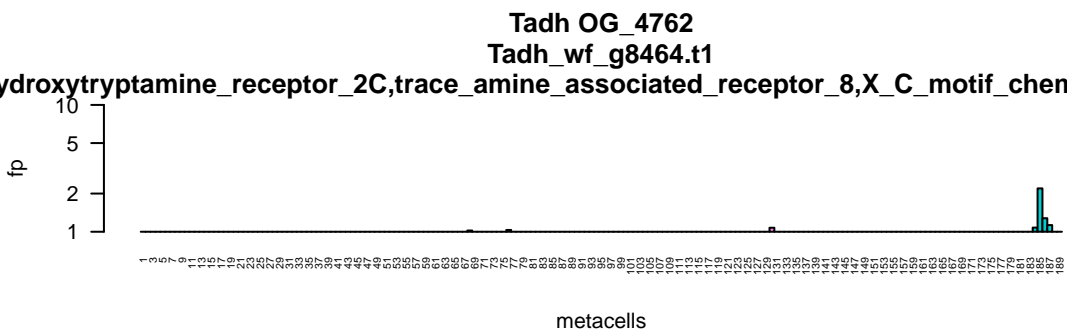
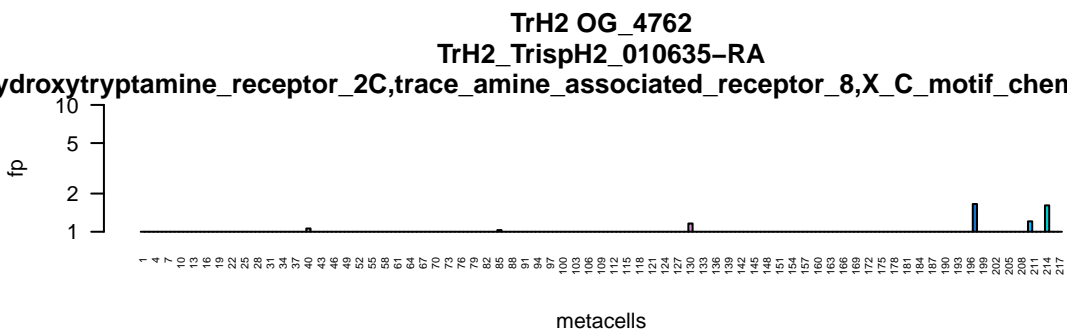
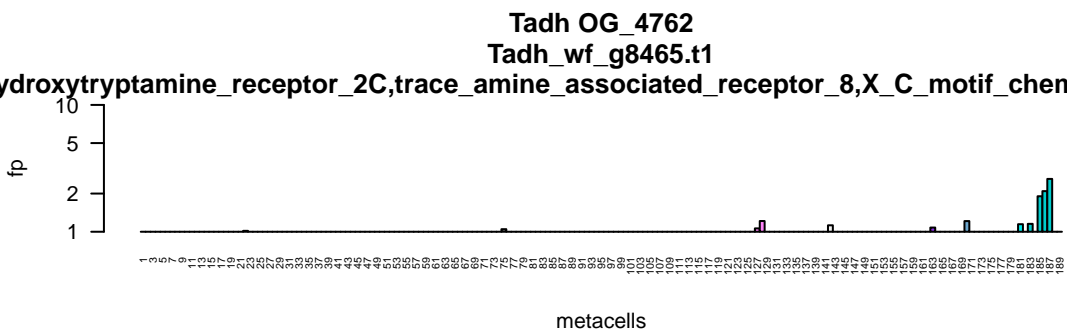
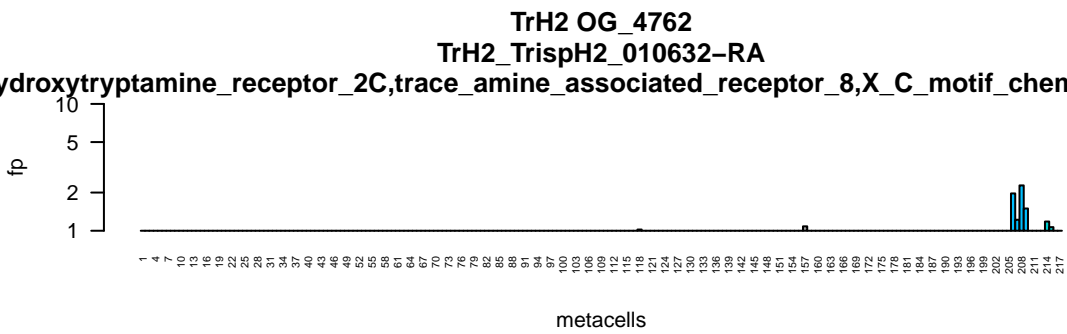
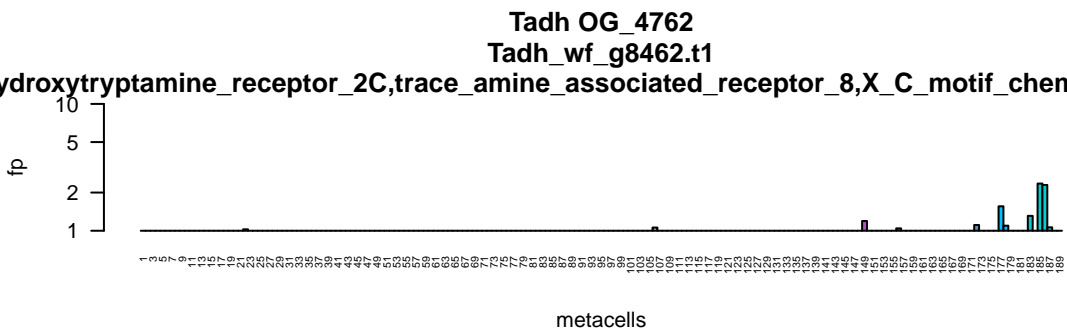


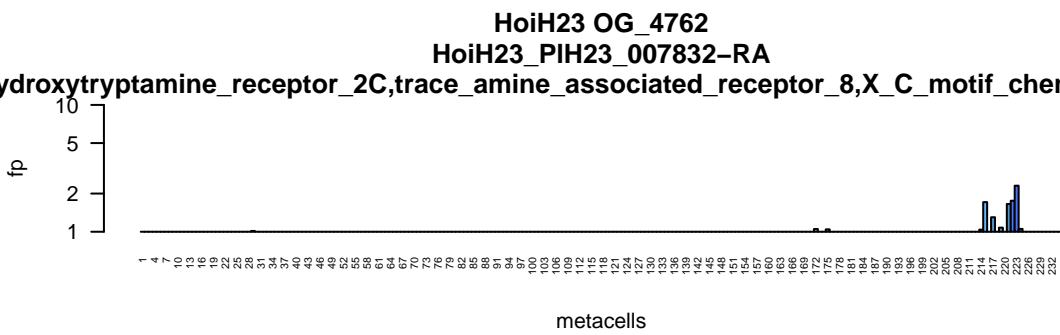
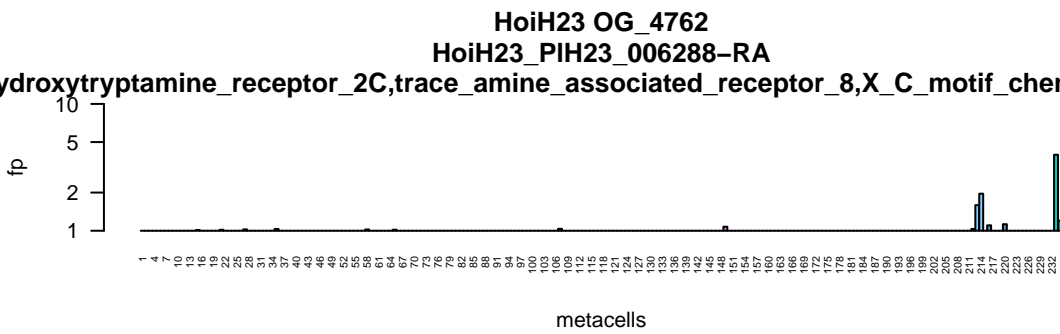
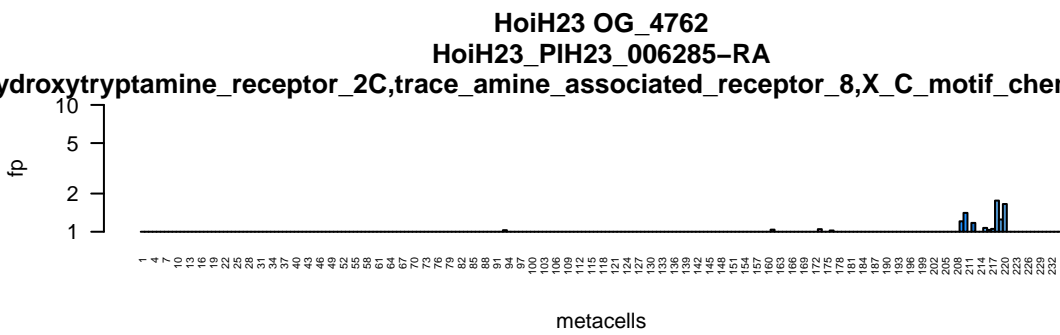
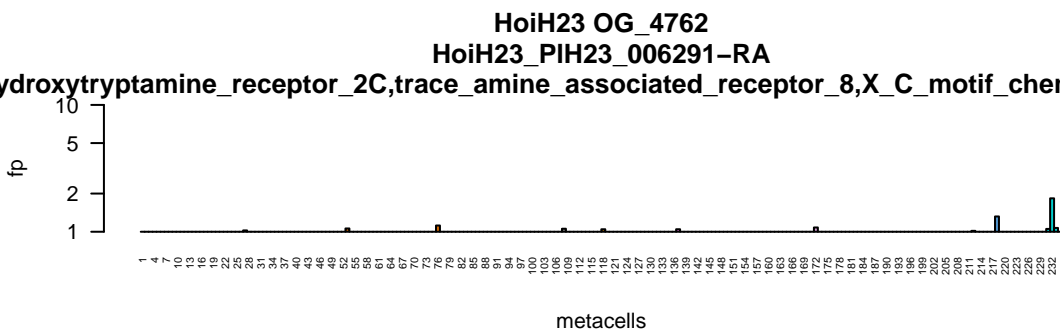
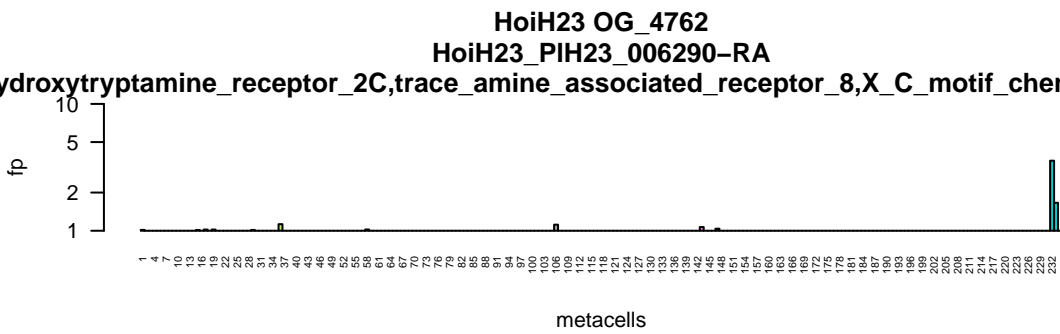
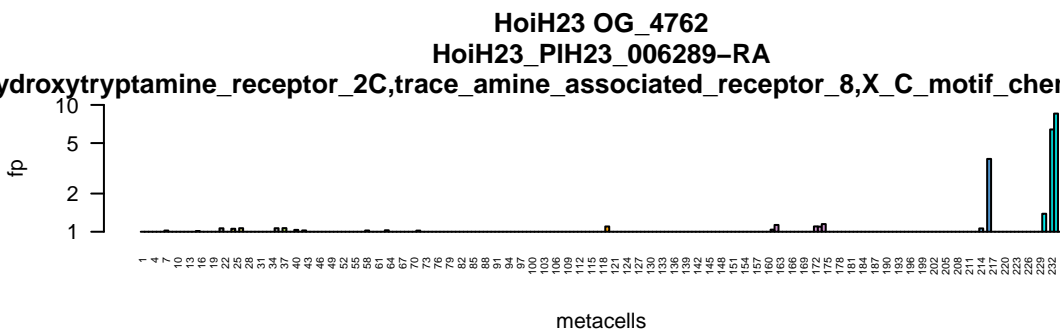
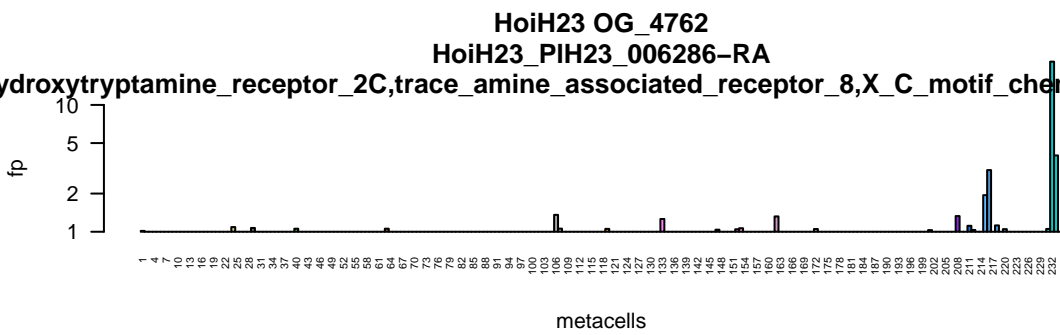
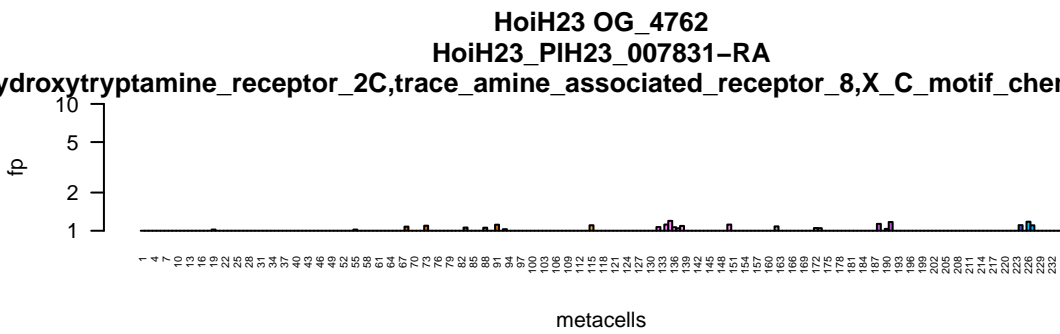
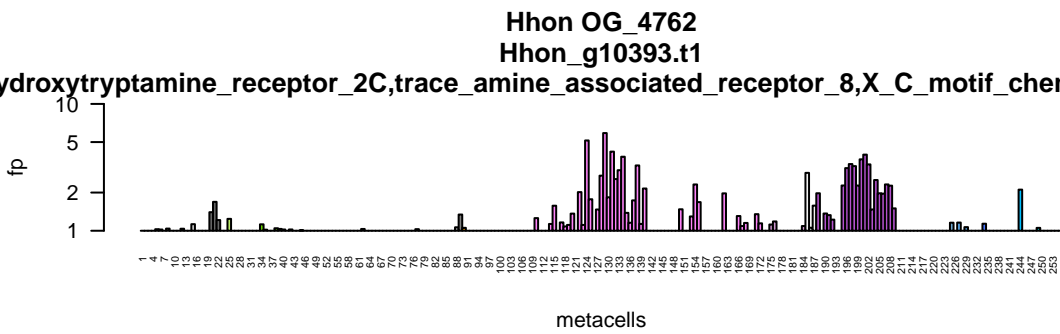
HoiH23 OG_4547
HoiH23_PIH23_002323-RA

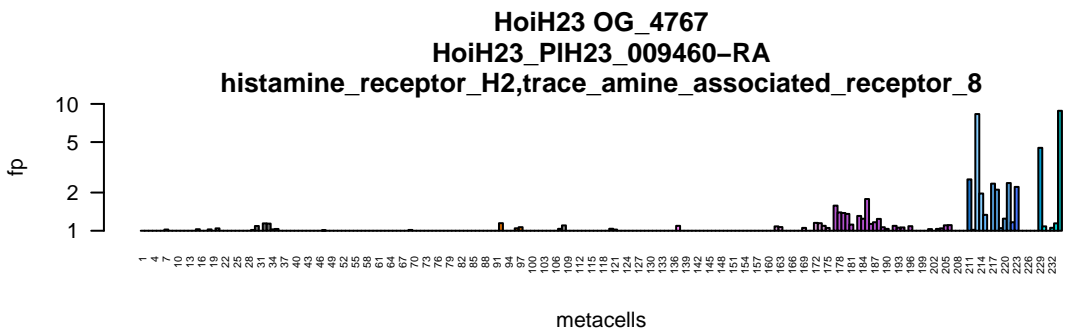
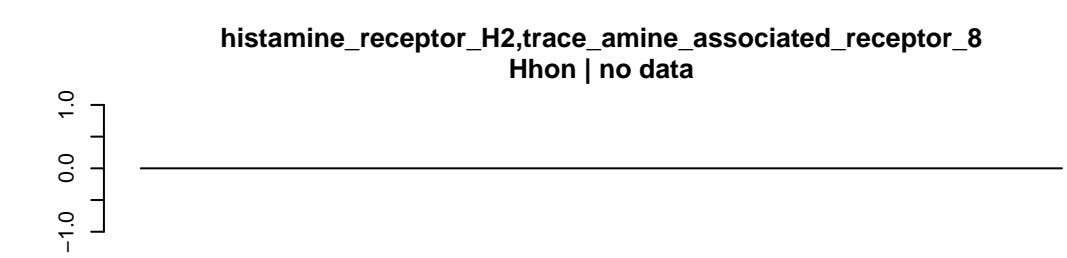
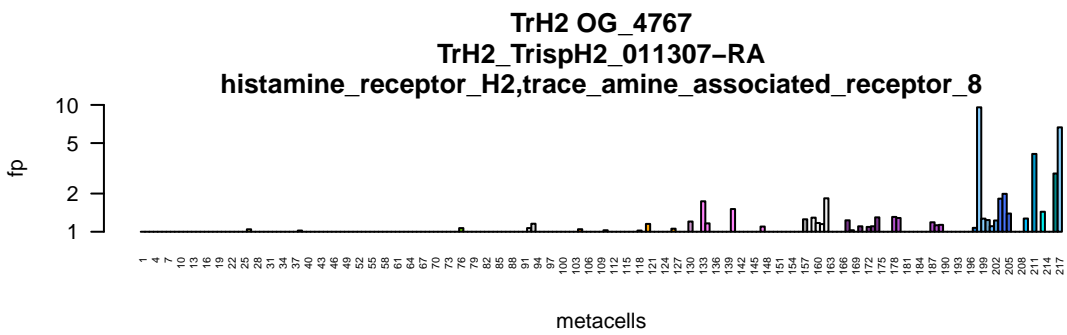
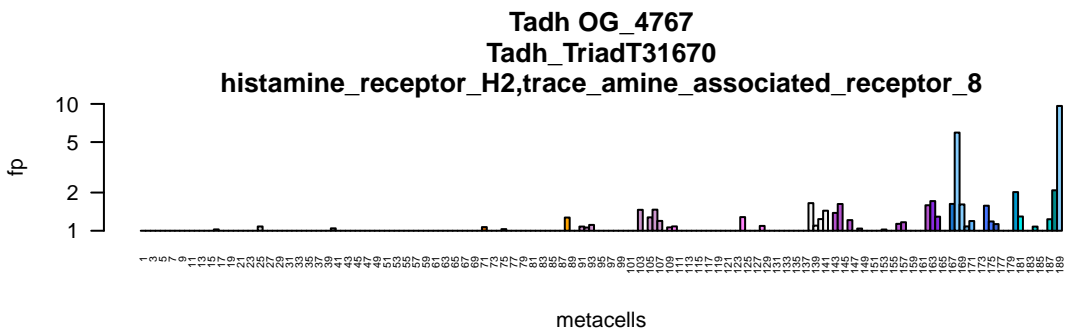




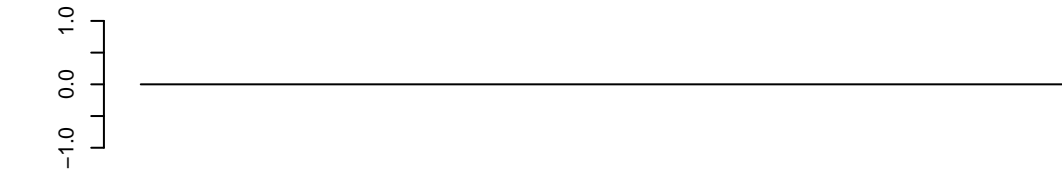




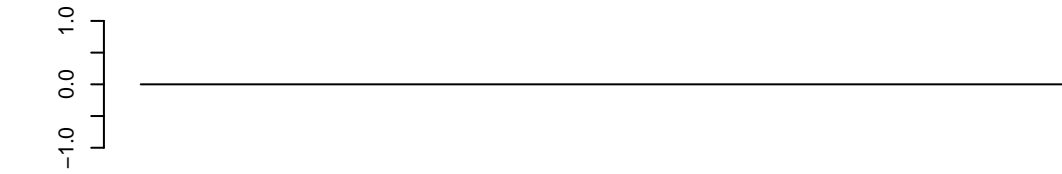




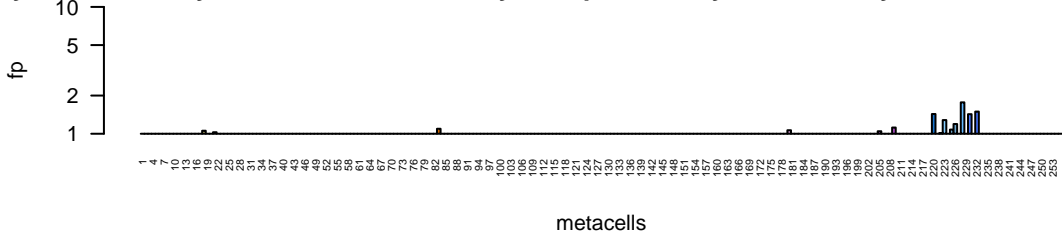
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi
Tadh | no data



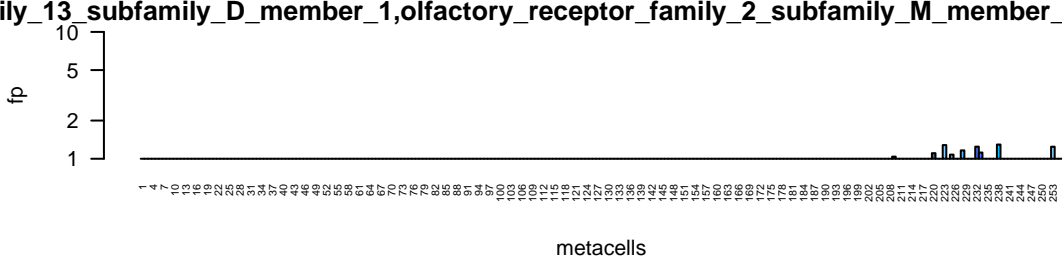
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TrH2 | no data



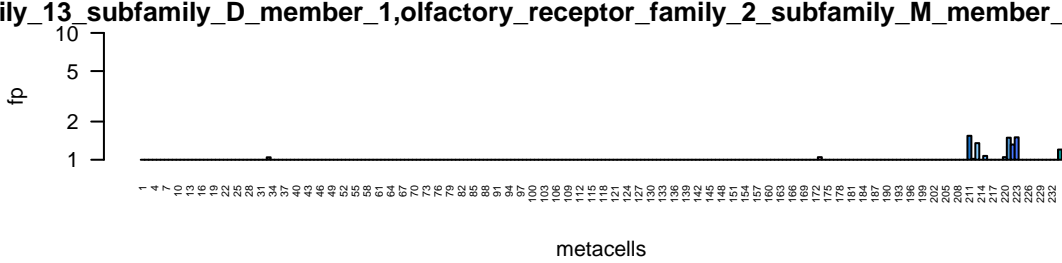
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Hhon OG_4788
Hhon_g08016.t1



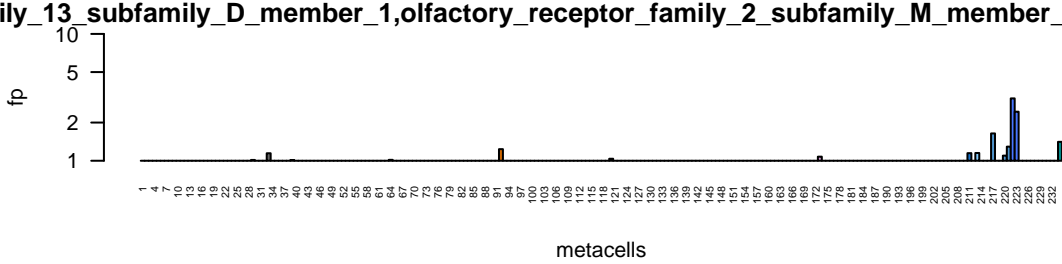
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi
Hhon OG_4788
Hhon_g08015.t1

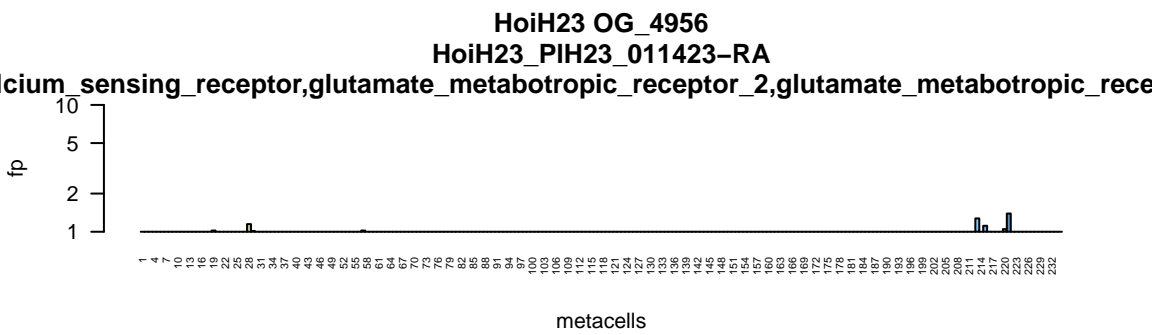
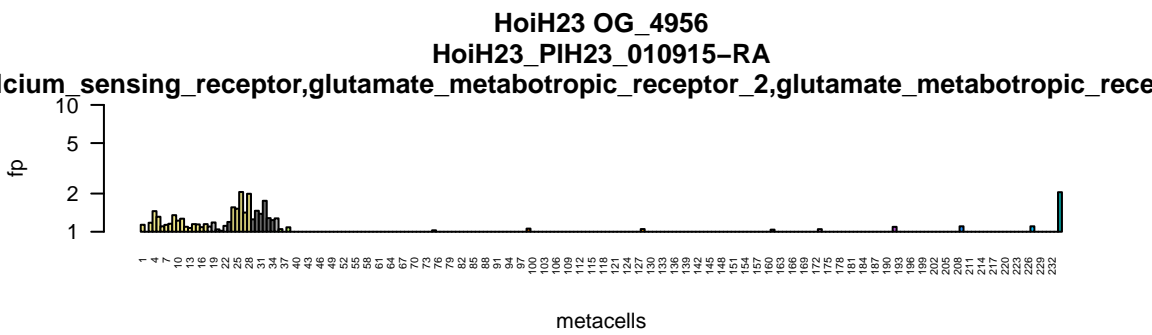
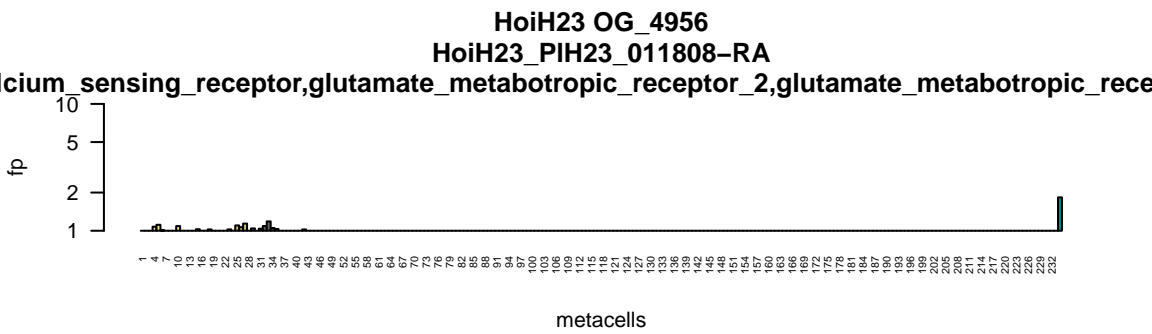
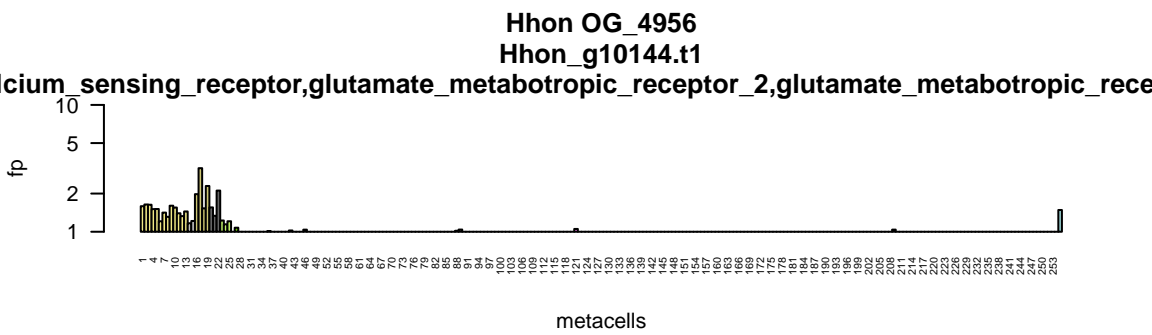
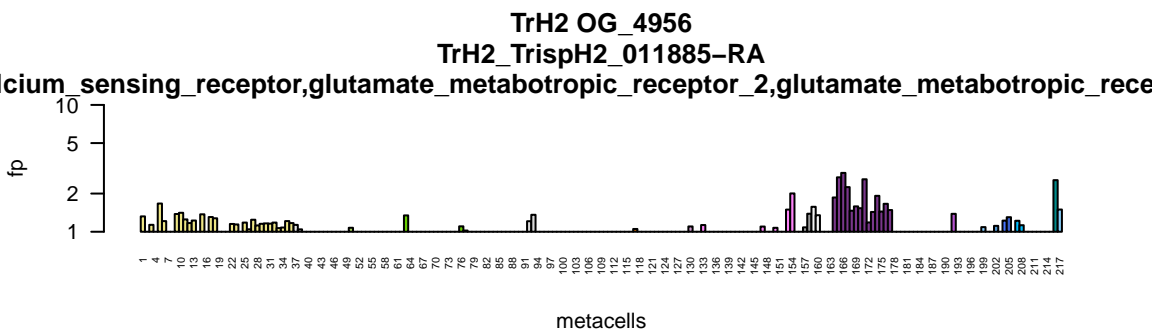
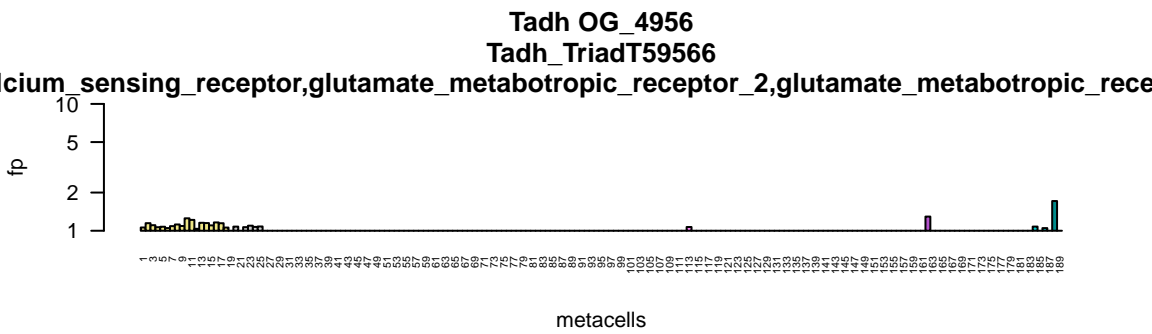


ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi
HoiH23 OG_4788
HoiH23_PIH23_005444-RA

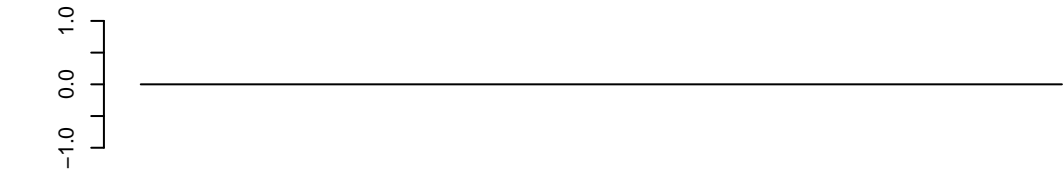


ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi
HoiH23 OG_4788
HoiH23_PIH23_005445-RA

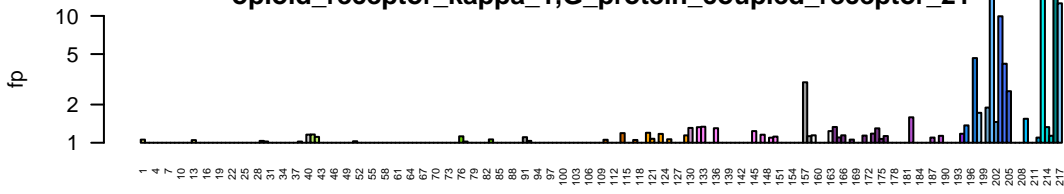




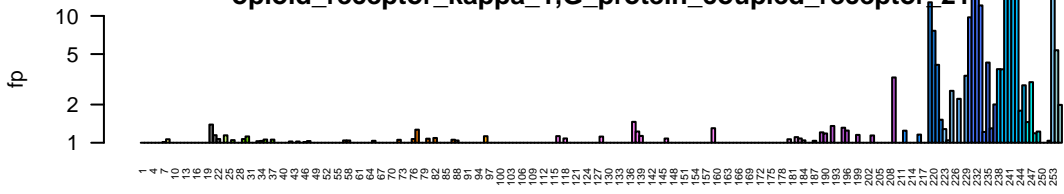
opioid_receptor_kappa_1,G_protein_coupled_receptor_21
Tadh | no data



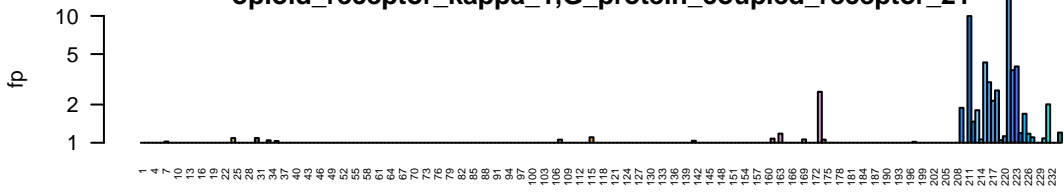
TrH2 OG_5203
TrH2_TrispH2_004138-RA
opioid_receptor_kappa_1,G_protein_coupled_receptor_21



Hhon OG_5203
Hhon_g01680.t1
opioid_receptor_kappa_1,G_protein_coupled_receptor_21



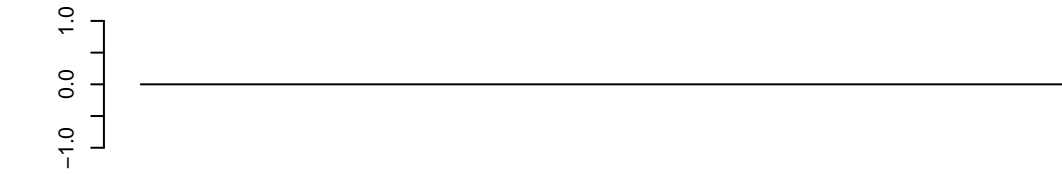
HoiH23 OG_5203
HoiH23_PIH23_003635-RA
opioid_receptor_kappa_1,G_protein_coupled_receptor_21



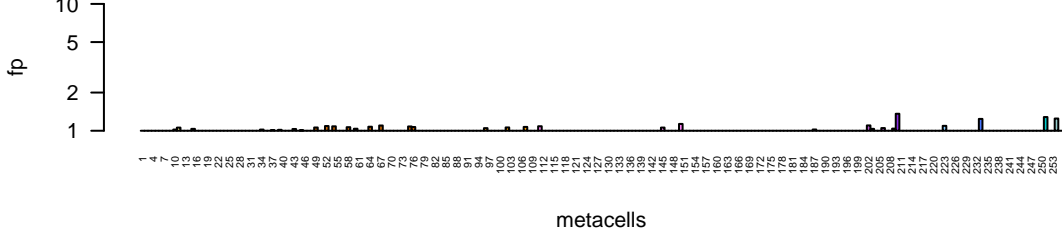
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recept
Tadh | no data



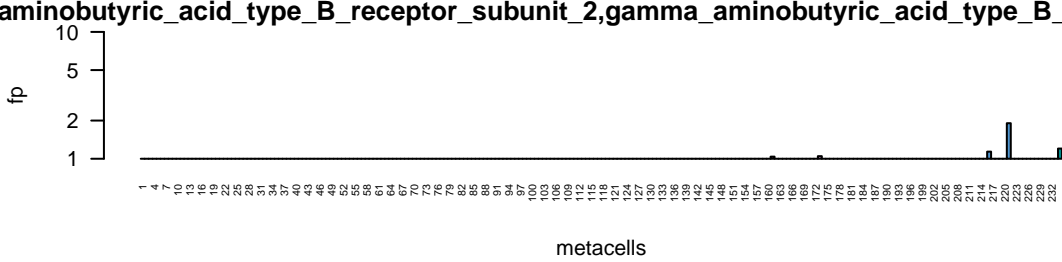
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recept
TrH2 | no data

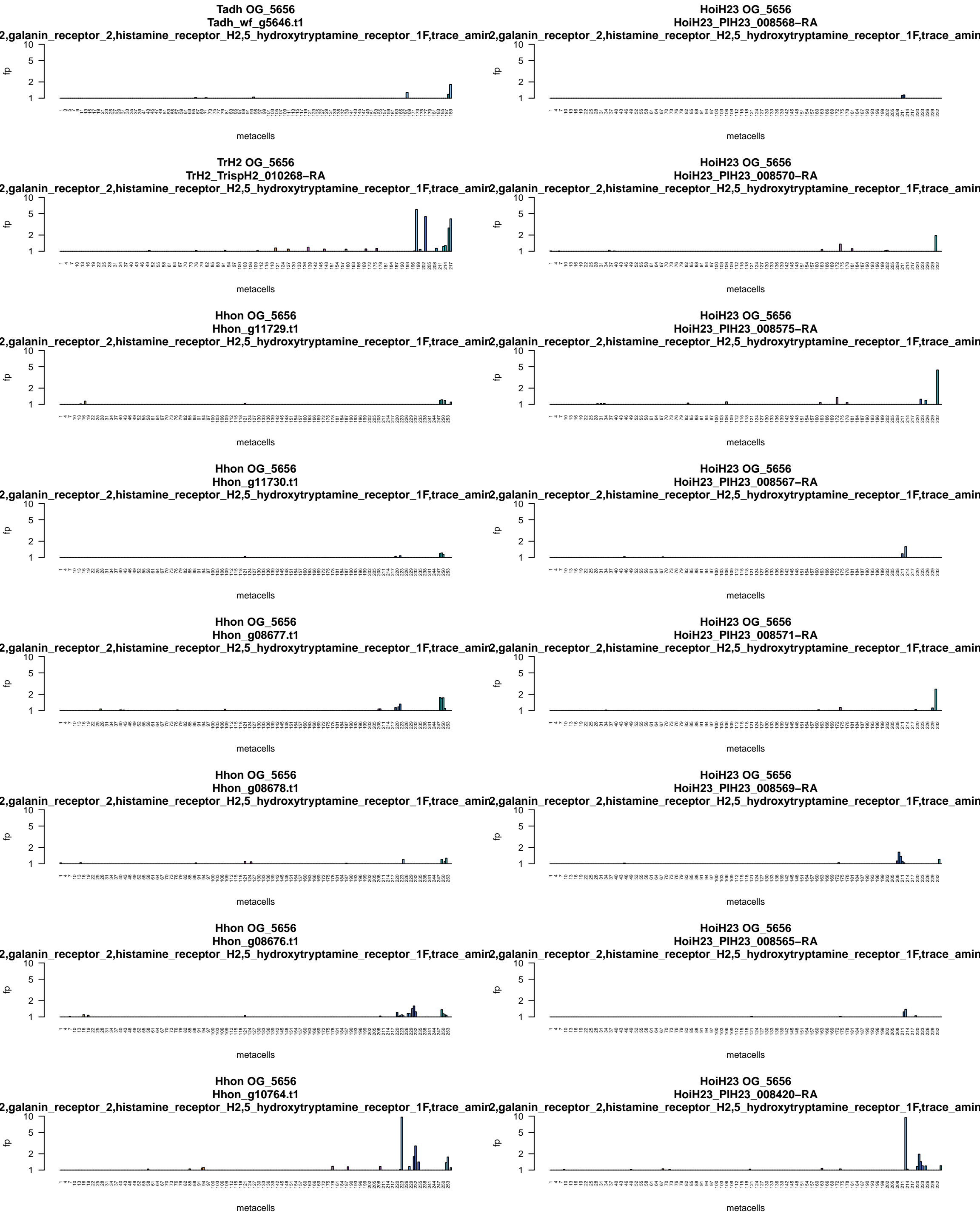


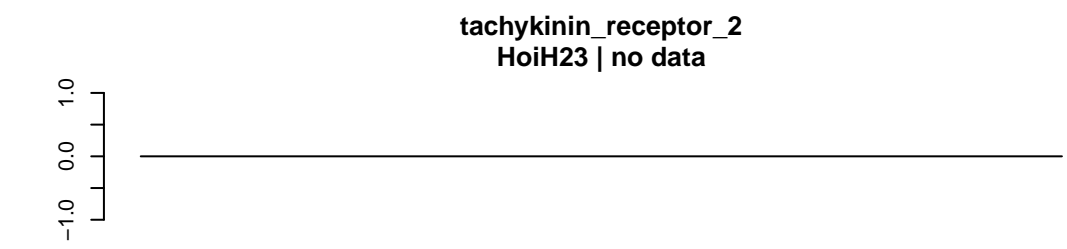
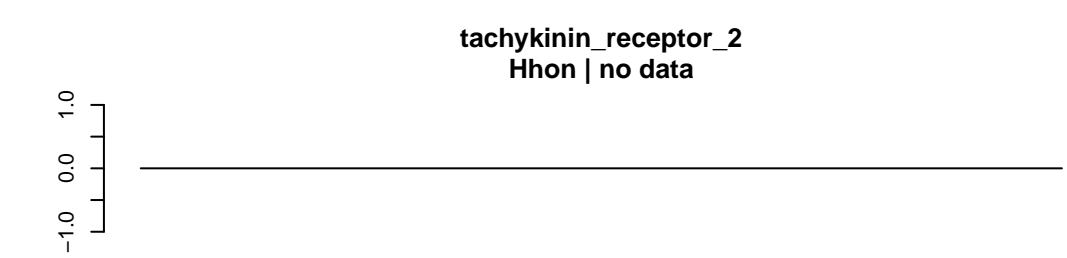
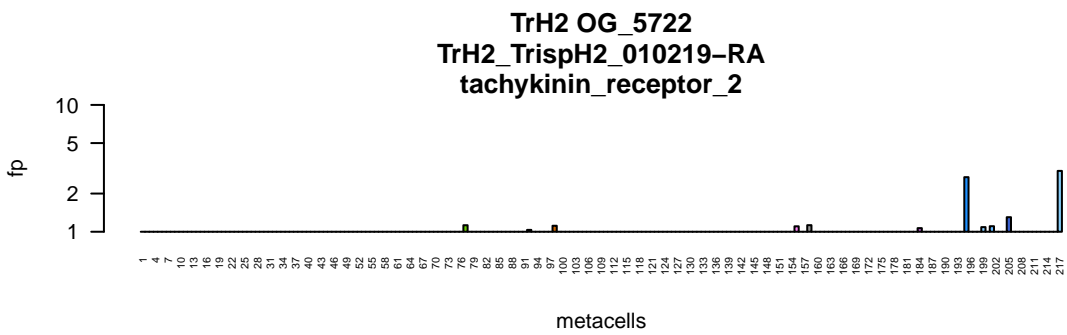
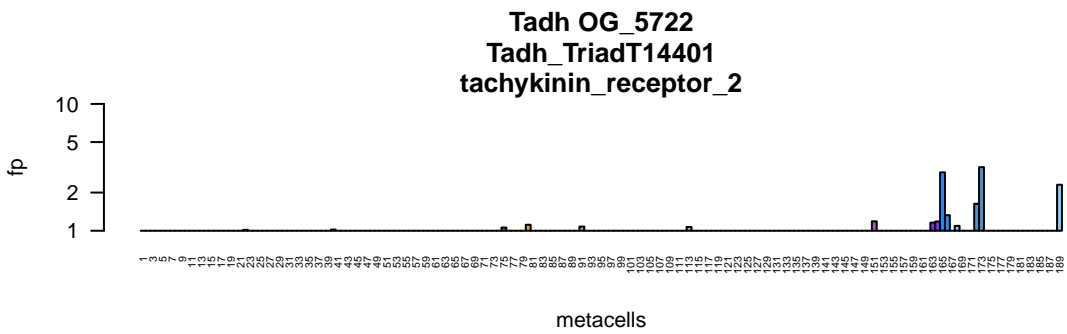
Hhon OG_5437
Hhon_g11390.t1
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recept

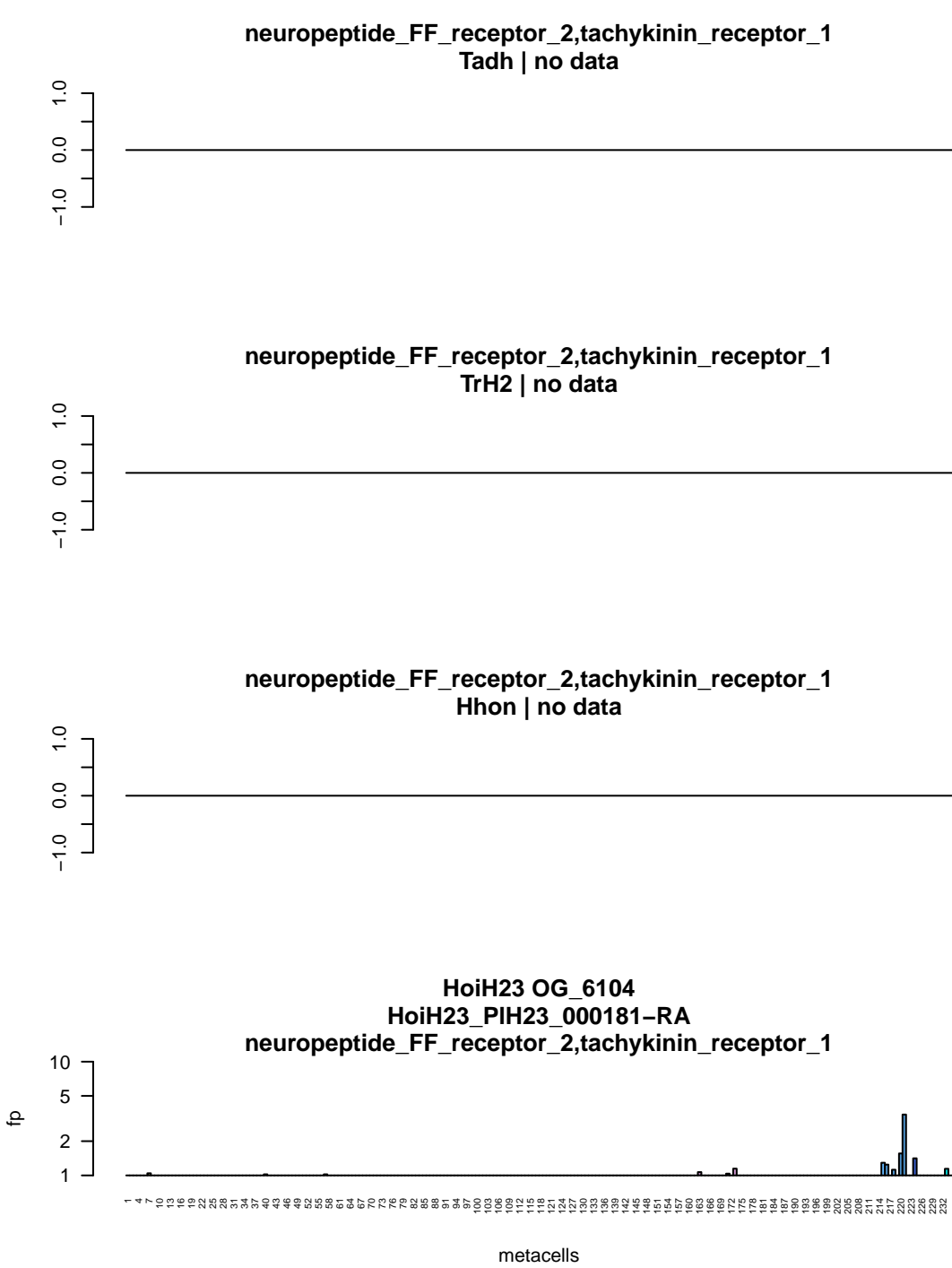


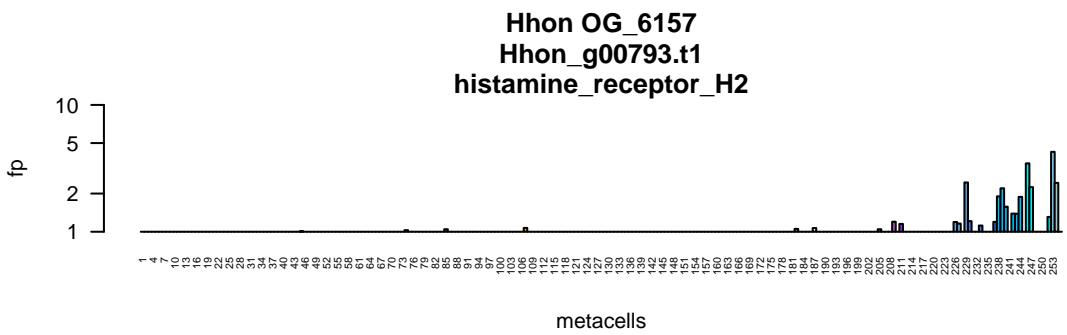
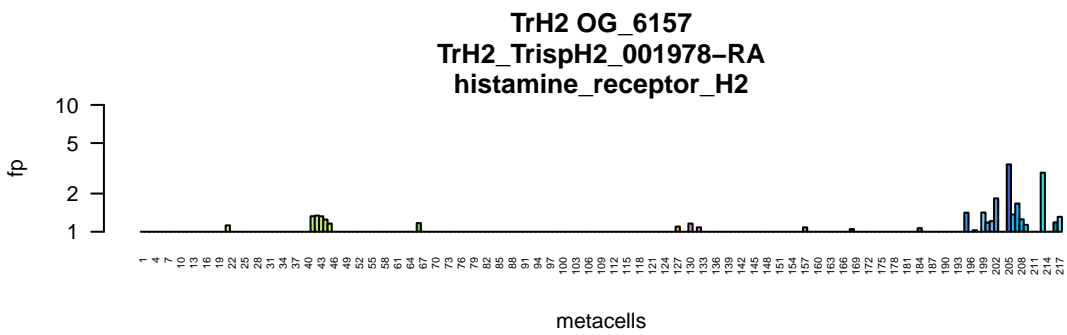
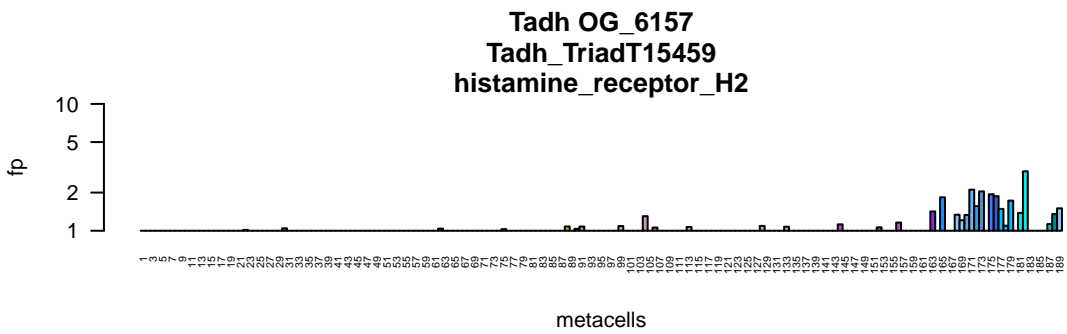
HoiH23 OG_5437
HoiH23_PIH23_011579-RA
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recept

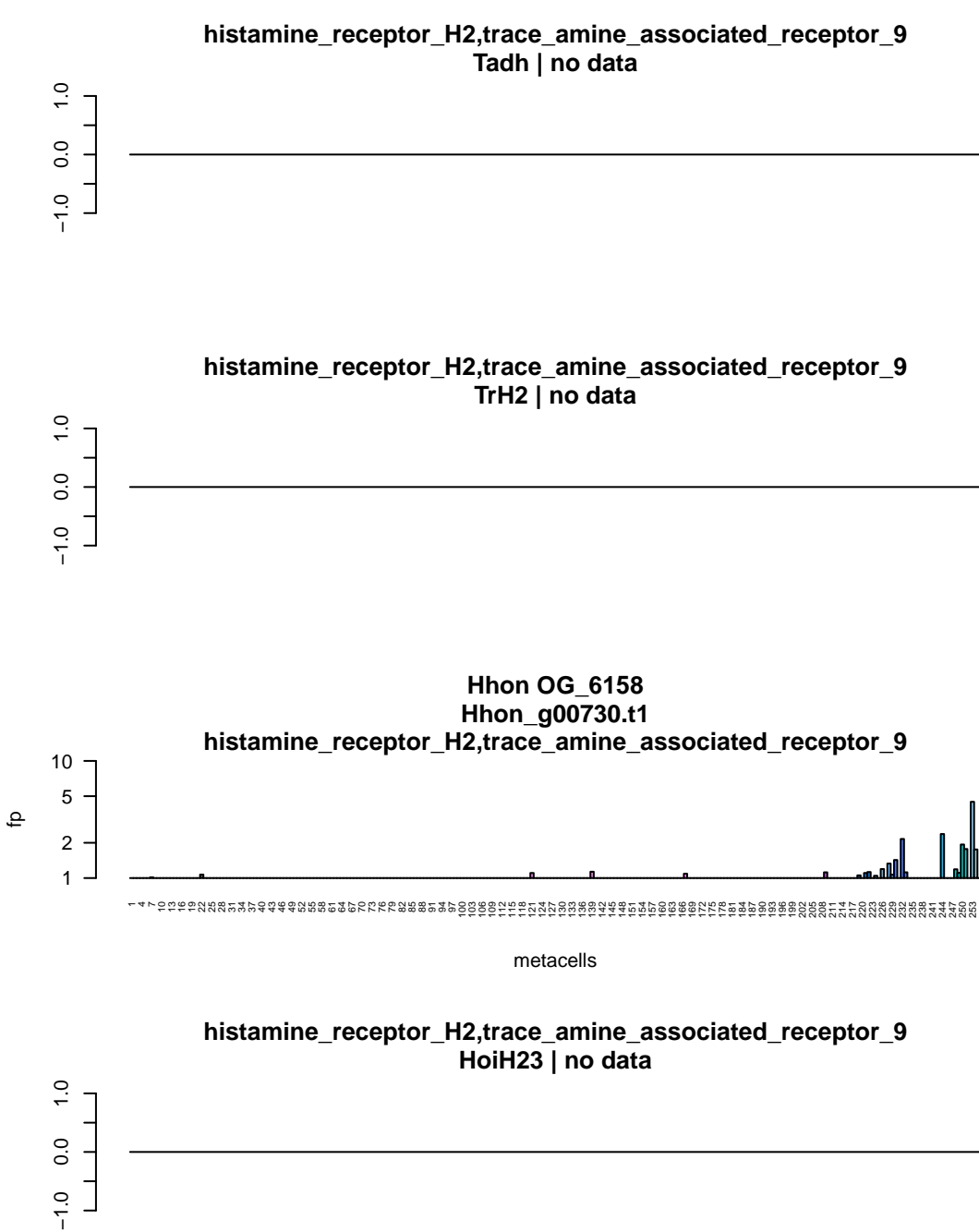




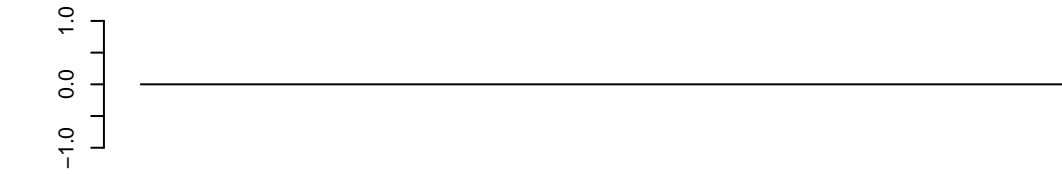




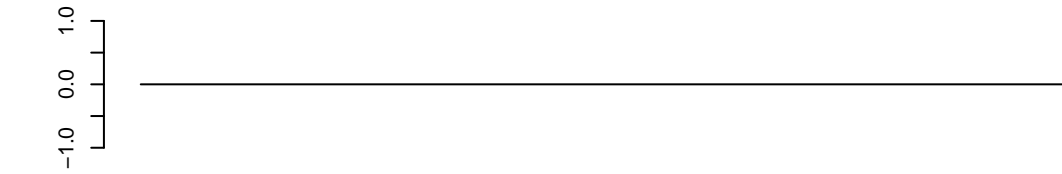




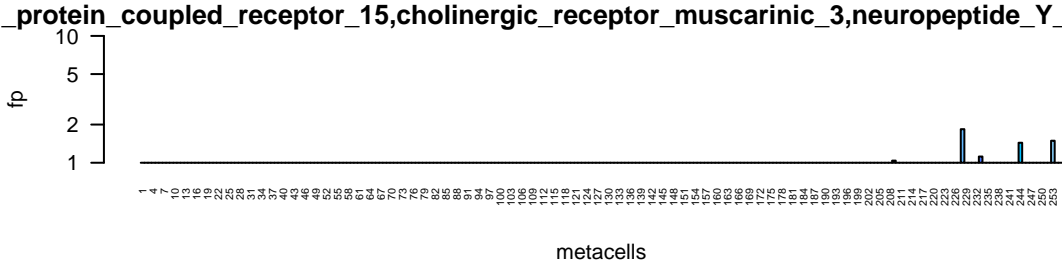
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept
Tadh | no data



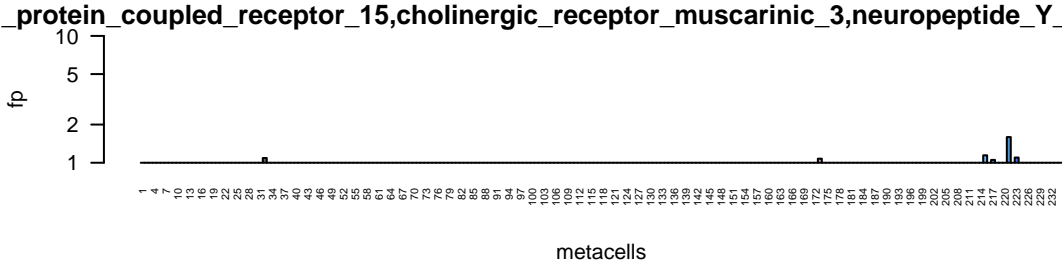
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept
TrH2 | no data

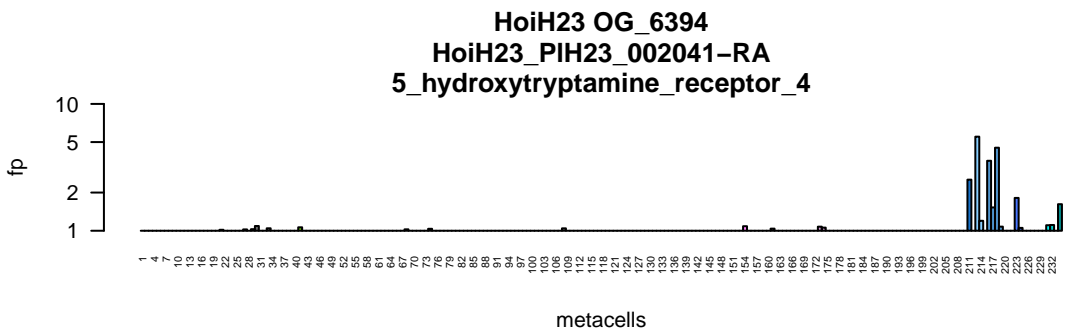
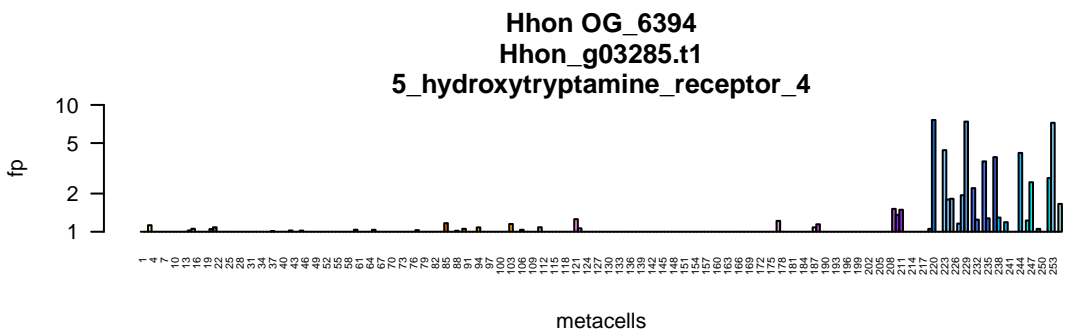
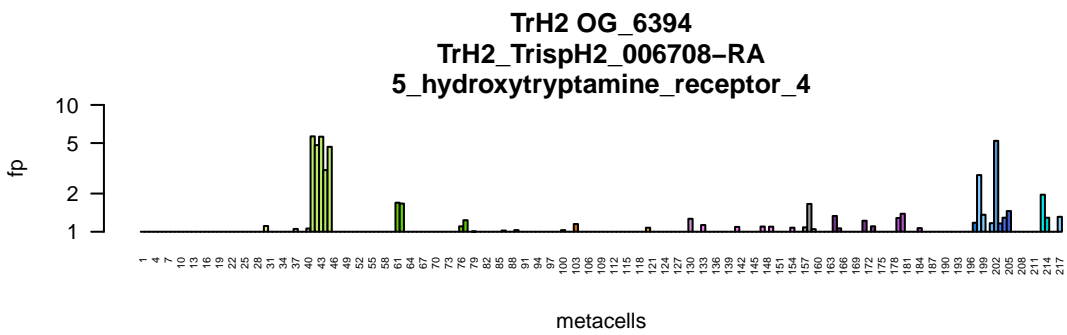
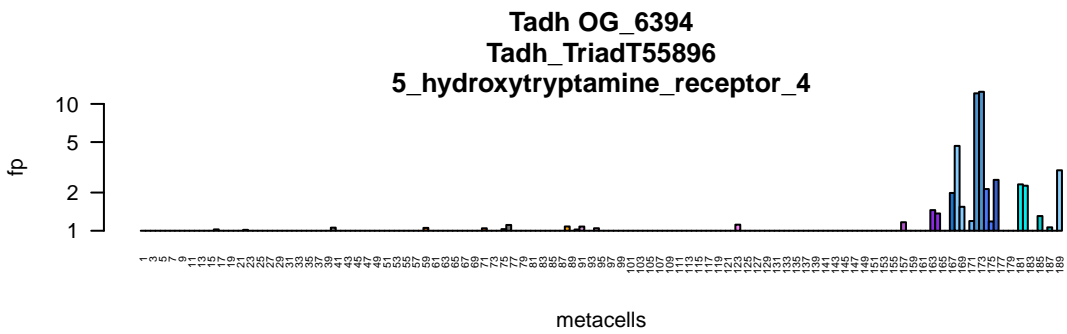


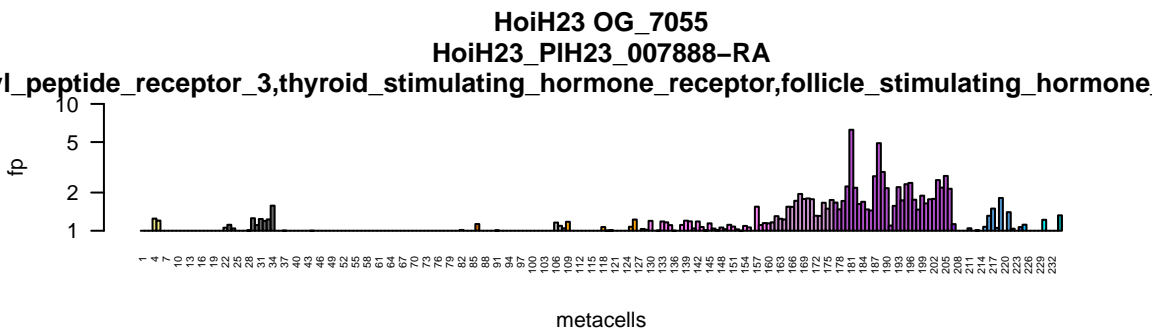
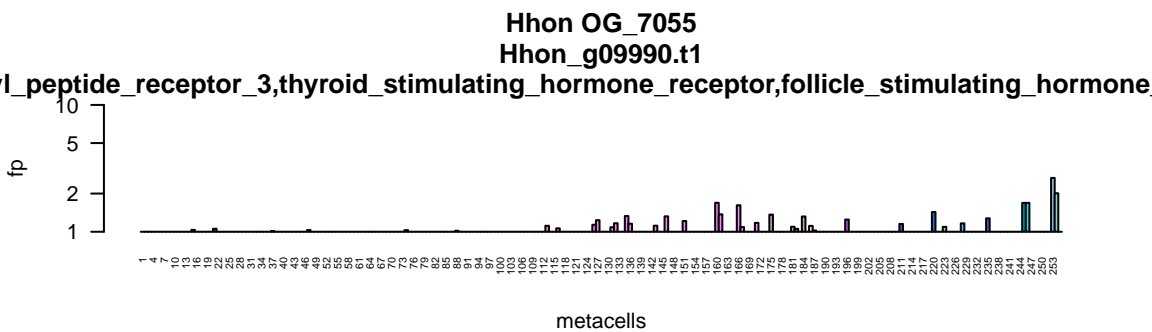
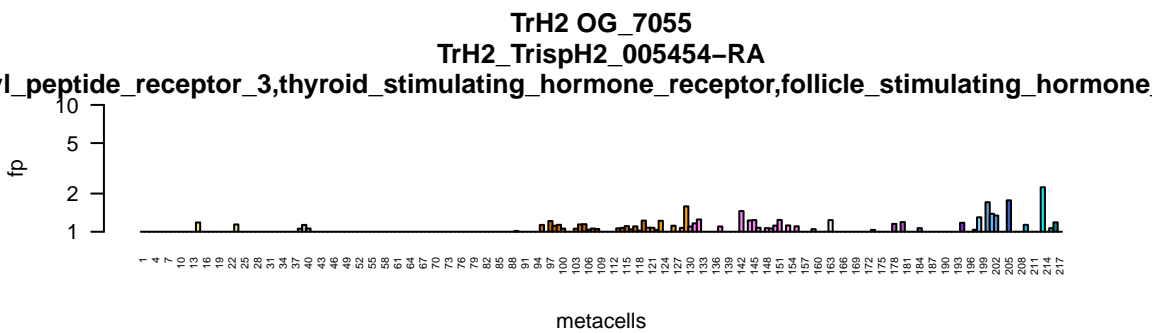
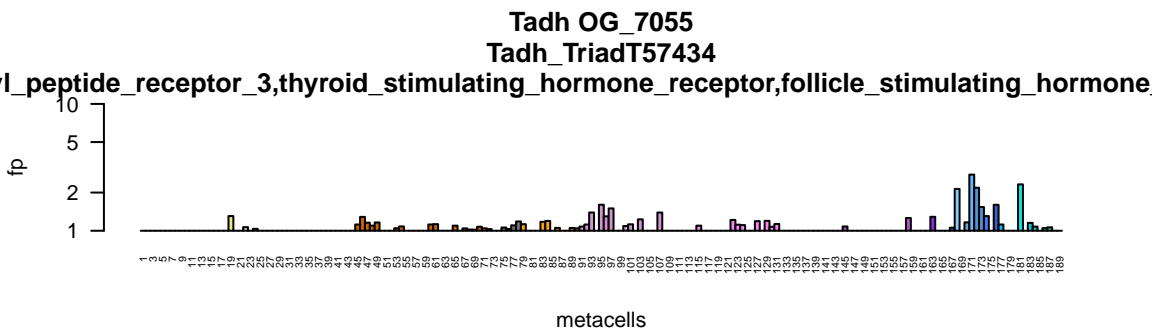
Hhon OG_6355
Hhon_g11199.t1
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept

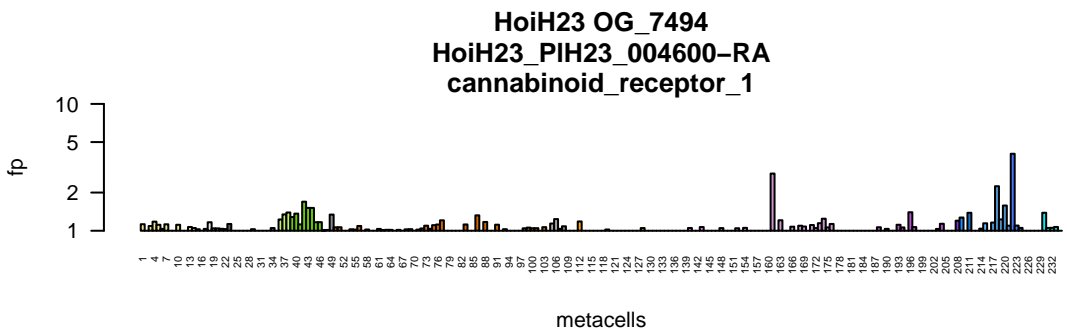
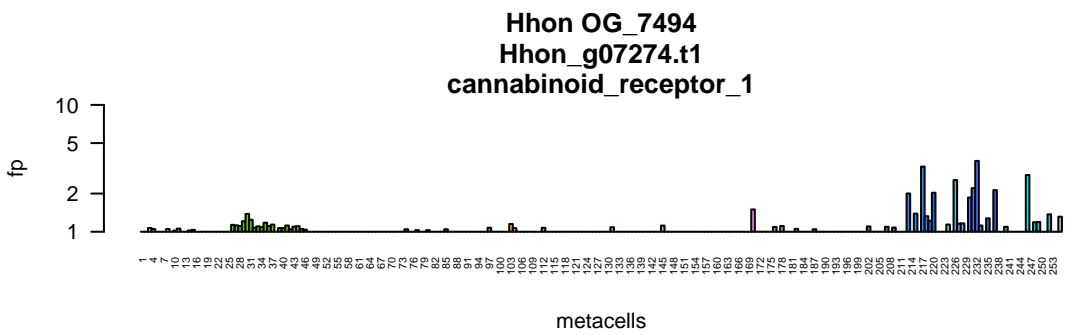
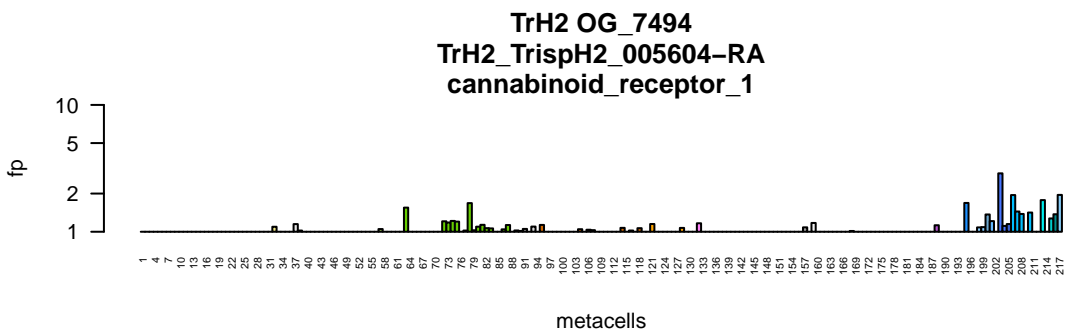
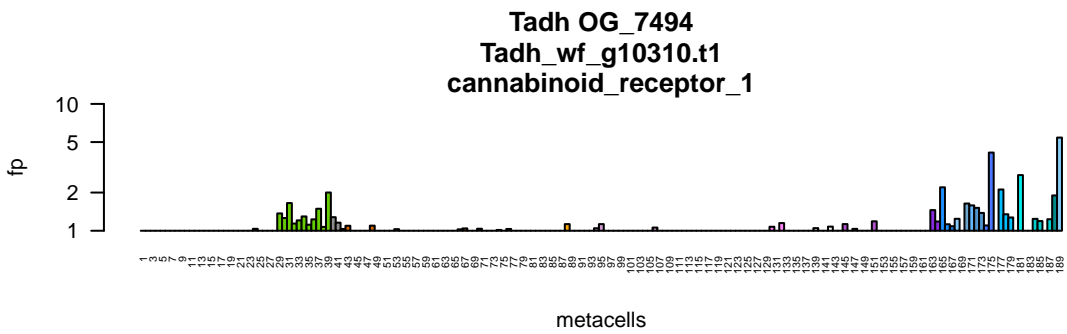


HoiH23 OG_6355
HoiH23_PIH23_012003-RA
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept

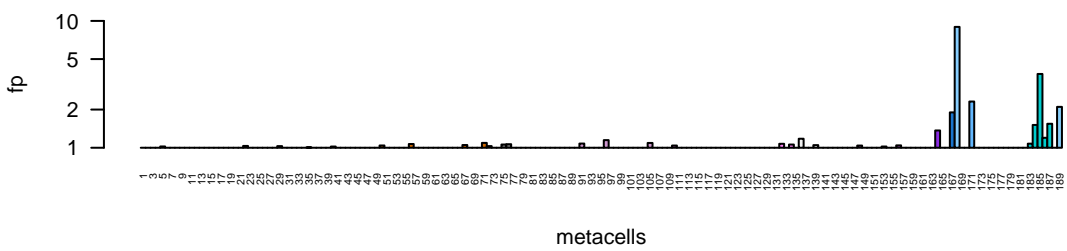






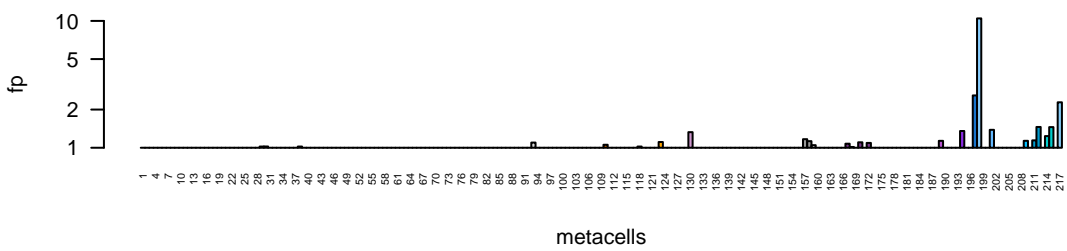


Tadh OG_7537
Tadh_wf_g5232.t1



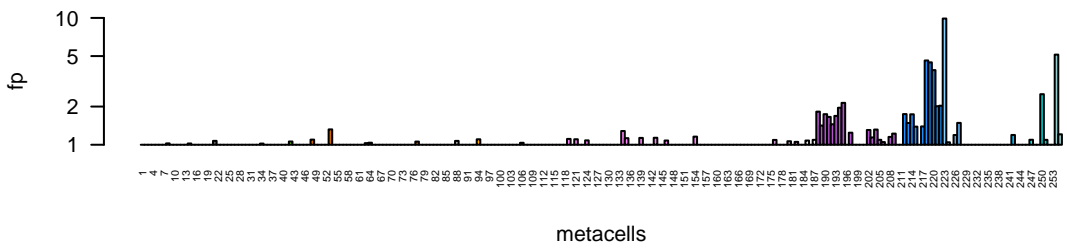
metacells

TrH2 OG_7537
TrH2_TrispH2_001160-RA



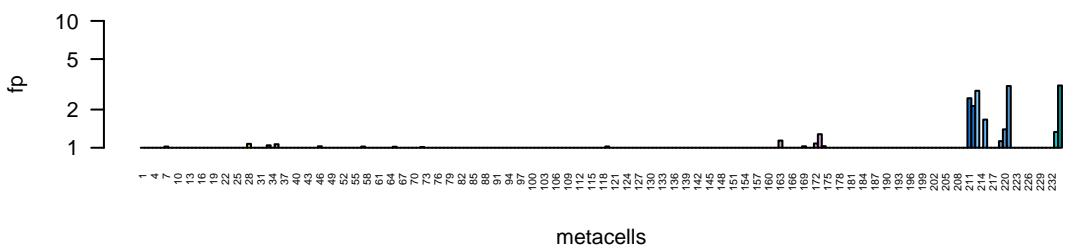
metacells

Hhon OG_7537
Hhon_g10209.t1

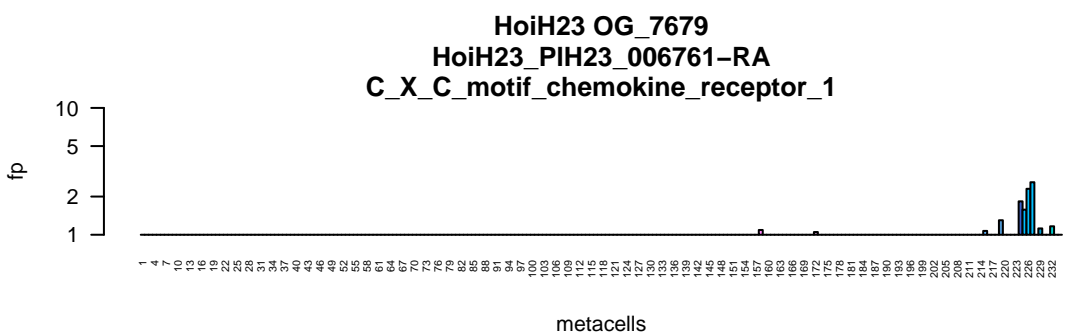
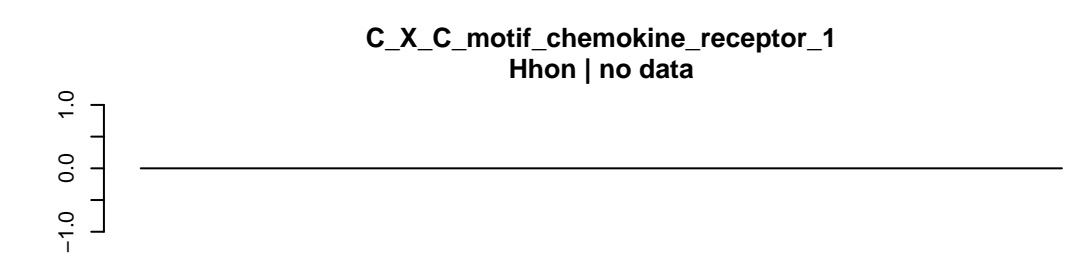
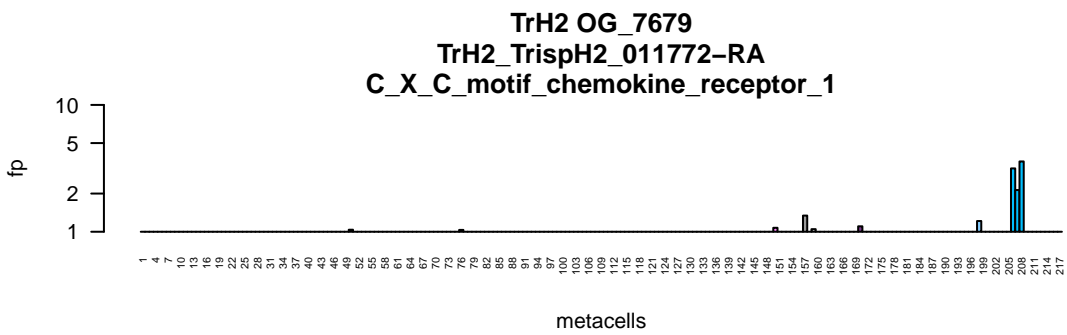
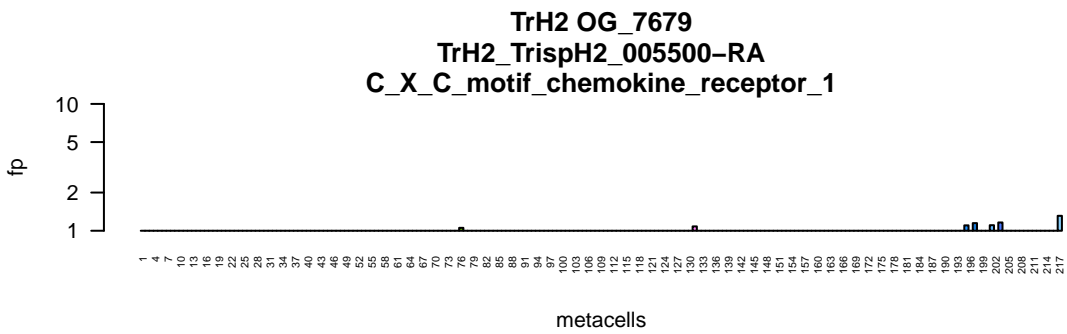
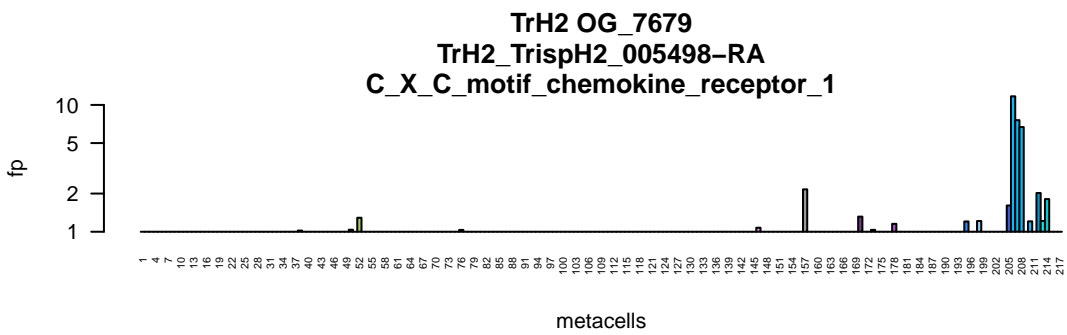
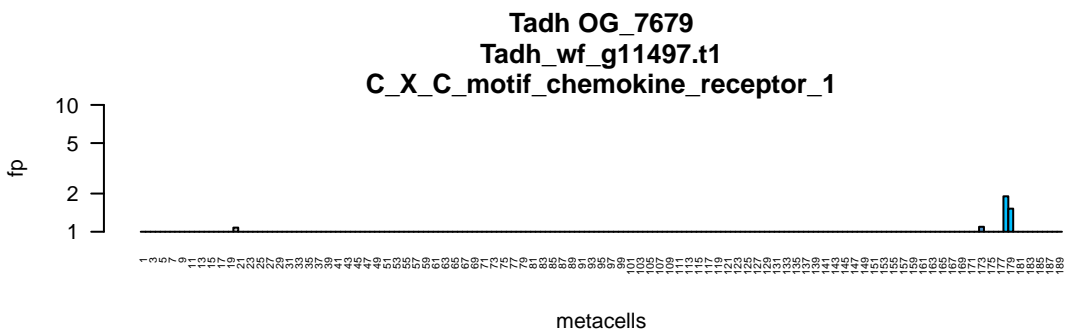
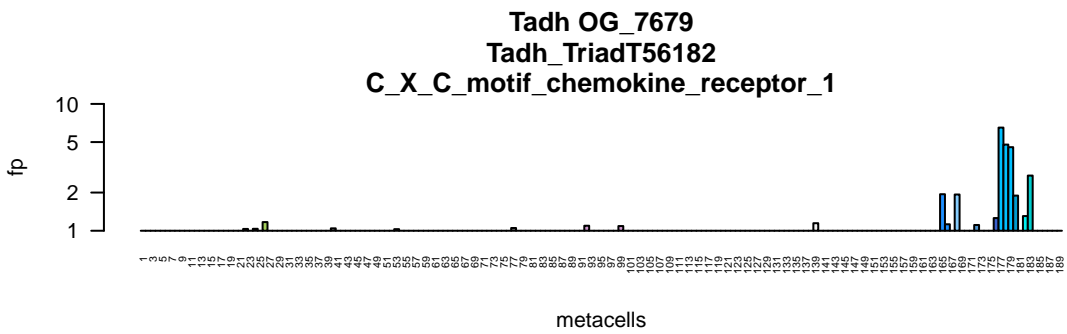


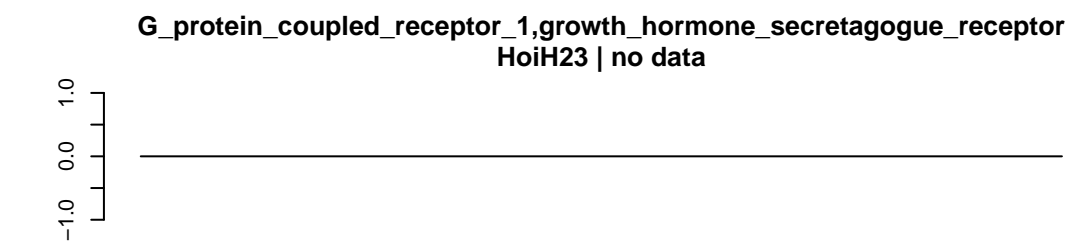
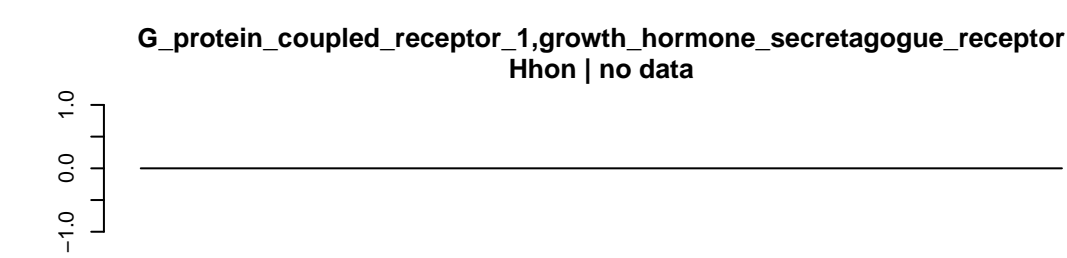
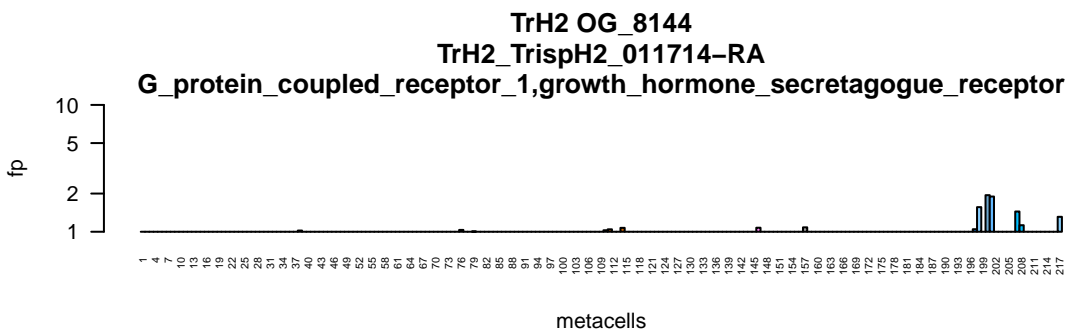
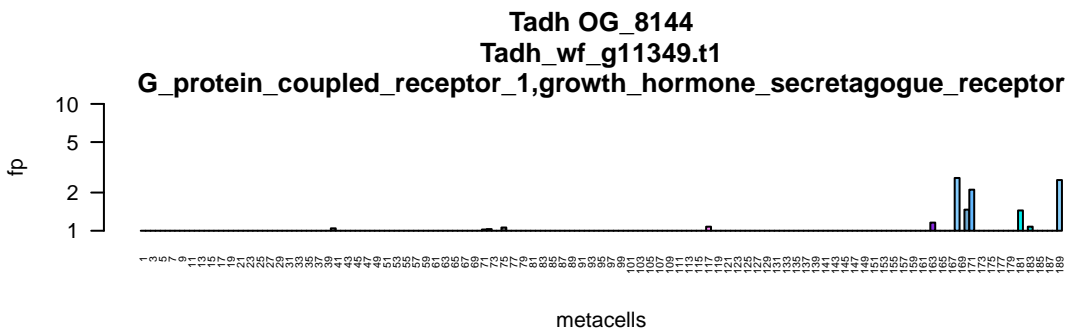
metacells

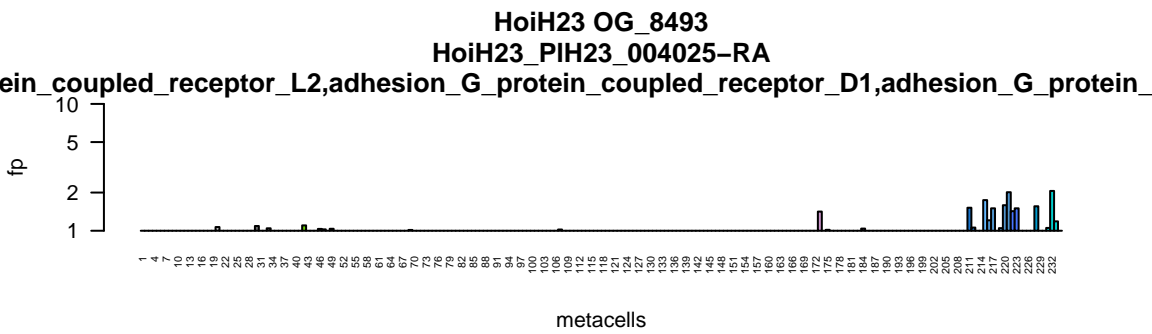
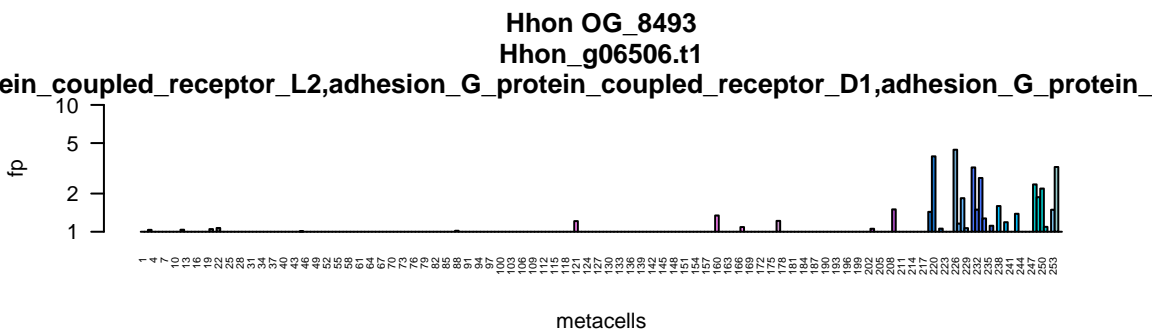
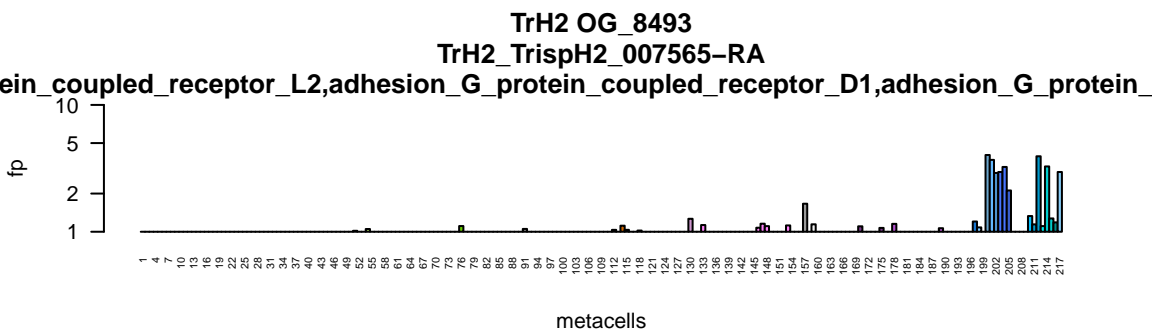
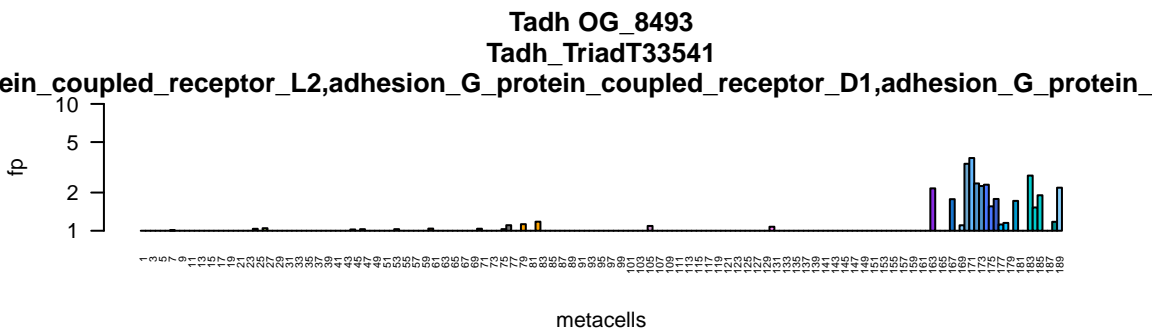
HoiH23 OG_7537
HoiH23_PIH23_000241-RA



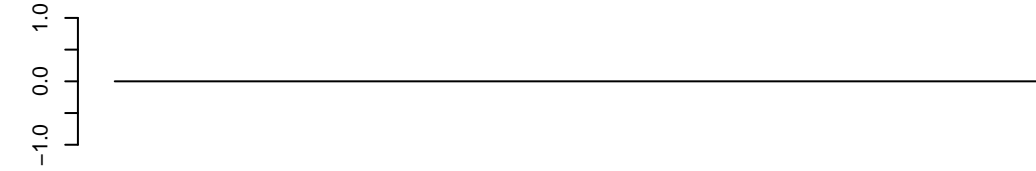
metacells



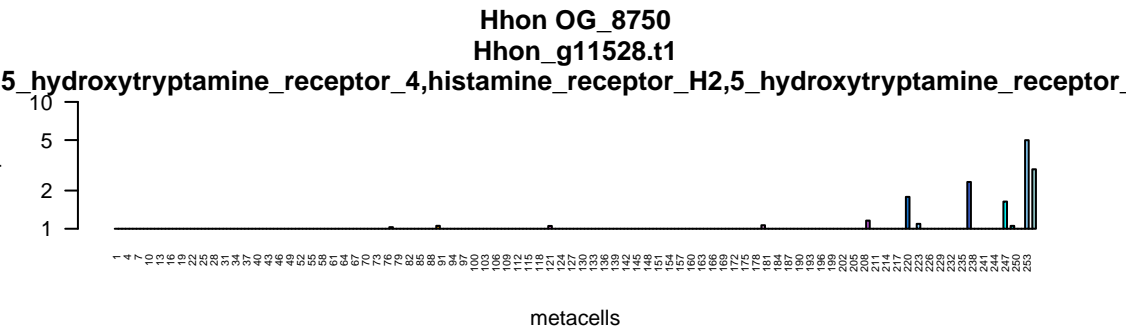




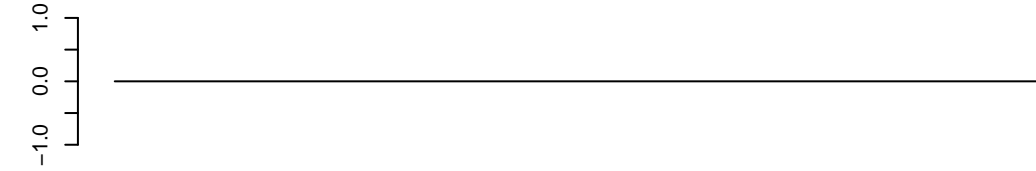
5_hydroxytryptamine_receptor_4,histamine_receptor_H2,5_hydroxytryptamine_receptor_
Tadh | no data



5_hydroxytryptamine_receptor_4,histamine_receptor_H2,5_hydroxytryptamine_receptor_
TrH2 | no data



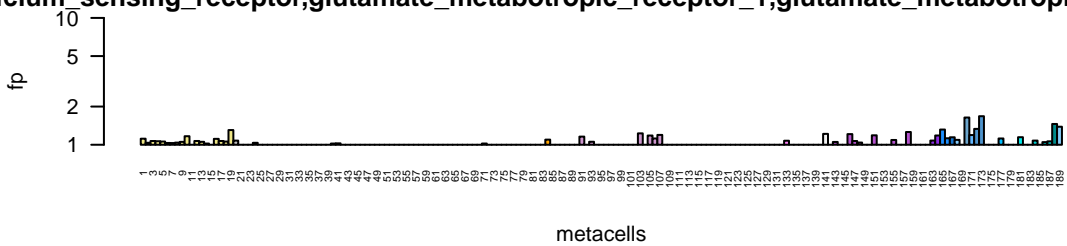
5_hydroxytryptamine_receptor_4,histamine_receptor_H2,5_hydroxytryptamine_receptor_
HoiH23 | no data



Tadh OG 8756

Tadh_wf_g965.t1

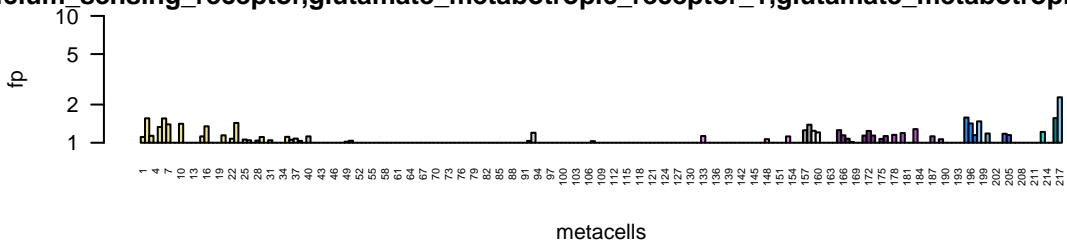
calcium_sensing_receptor,glutamate_metabotropic_receptor_1,glutamate_metabotropic_rece



TrH2 OG_8756

TrH2_TrispH2_000992-RA

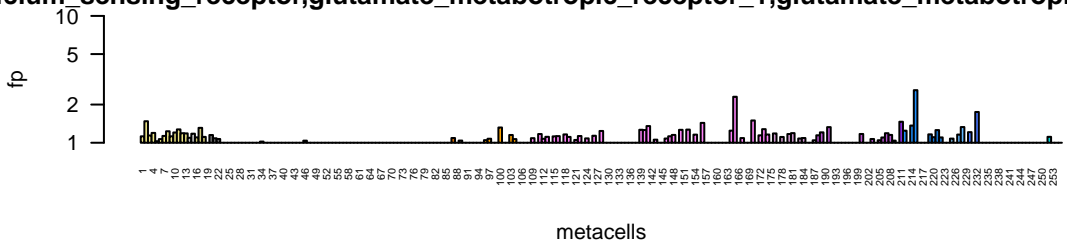
calcium_sensing_receptor,glutamate_metabotropic_receptor_1,glutamate_metabotropic_rece



Hhon OG_8756

Hhon_g02927.t1

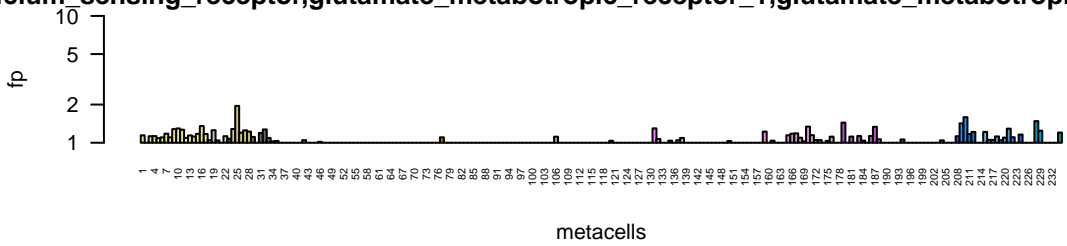
calcium_sensing_receptor,glutamate_metabotropic_receptor_1,glutamate_metabotropic_rece

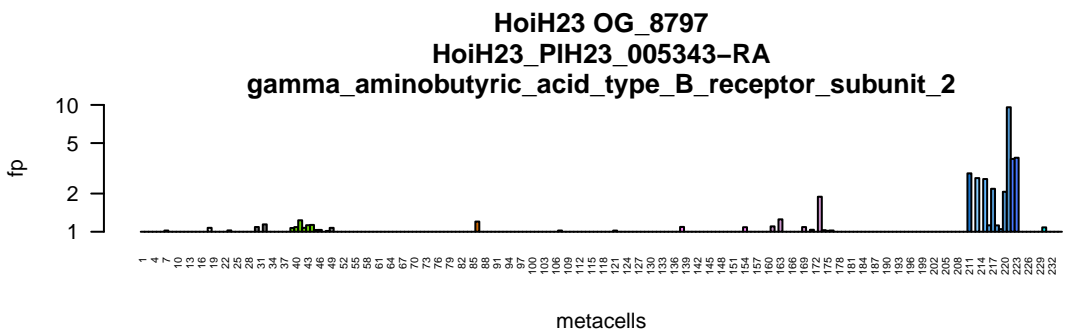
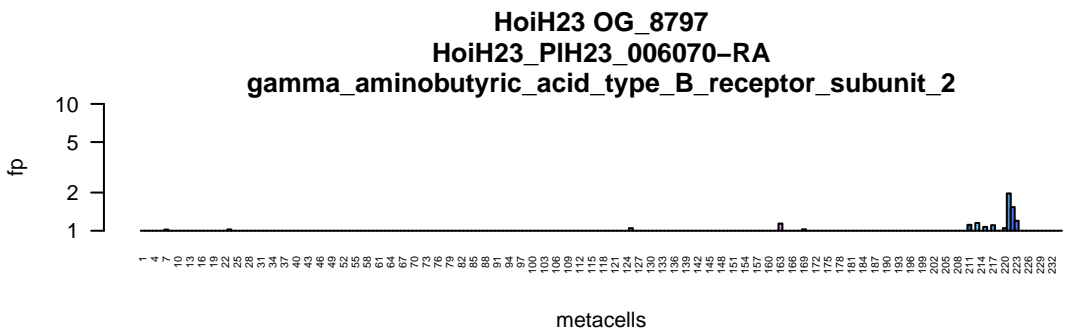
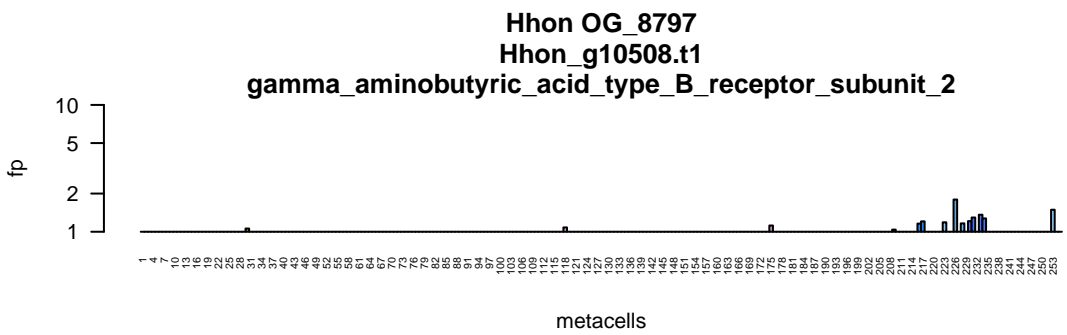
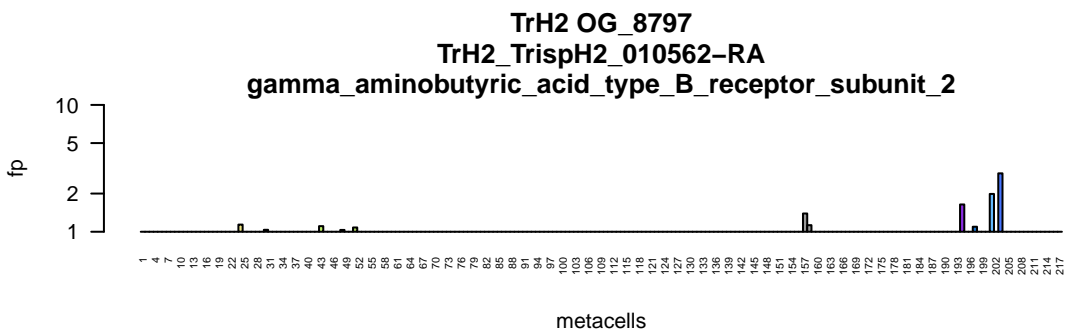
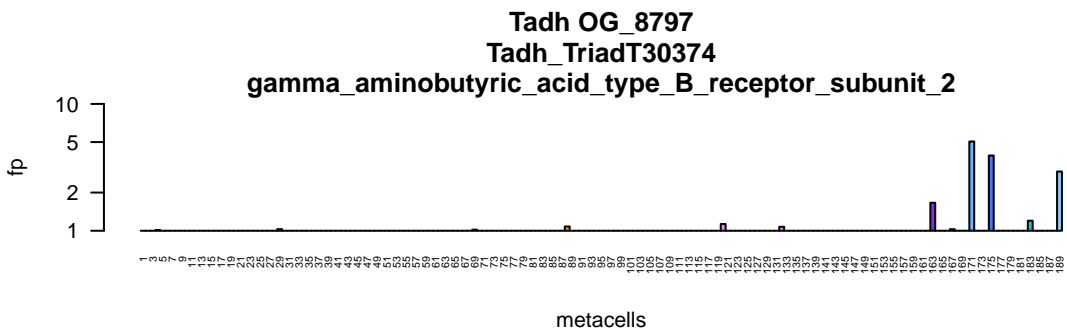


HoiH23 OG_8756

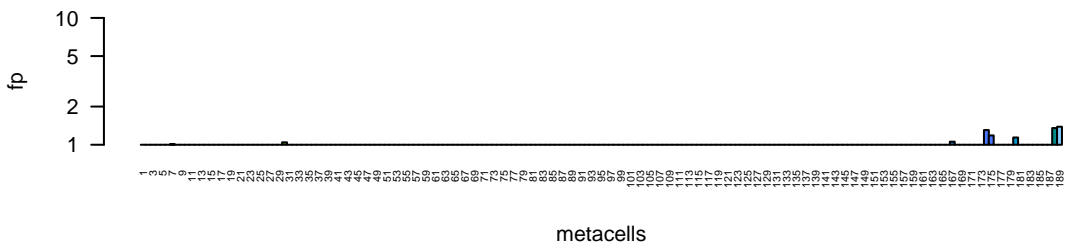
HoiH23_PIH23_010019-RA

calcium_sensing_receptor,glutamate_metabotropic_receptor_1,glutamate_metabotropic_rece





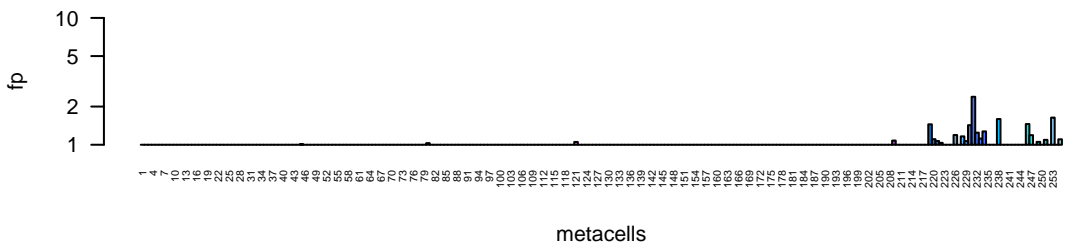
Tadh OG_9610
Tadh_TriadT60281



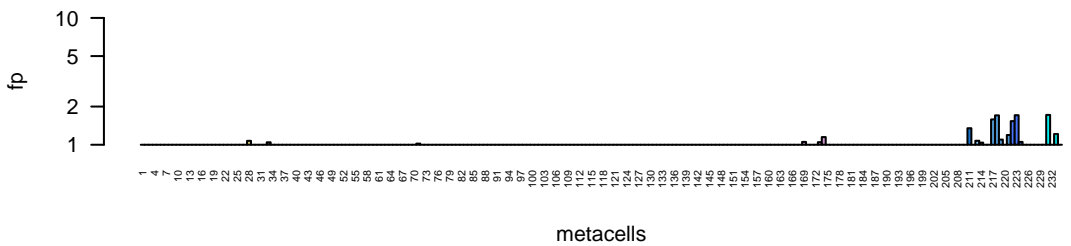
TrH2 | no data

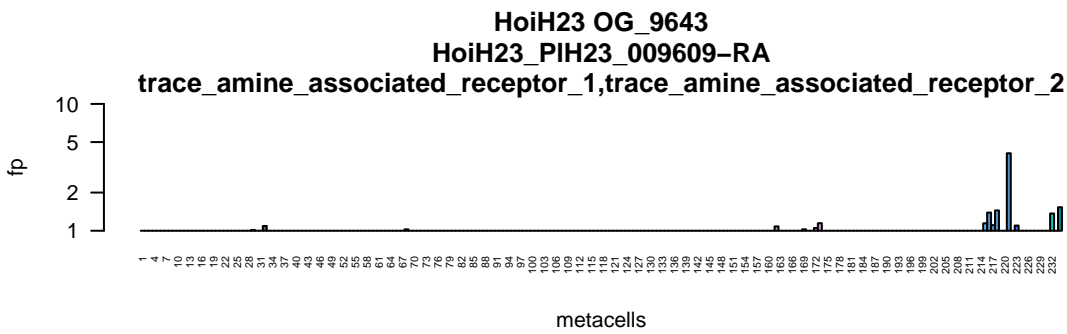
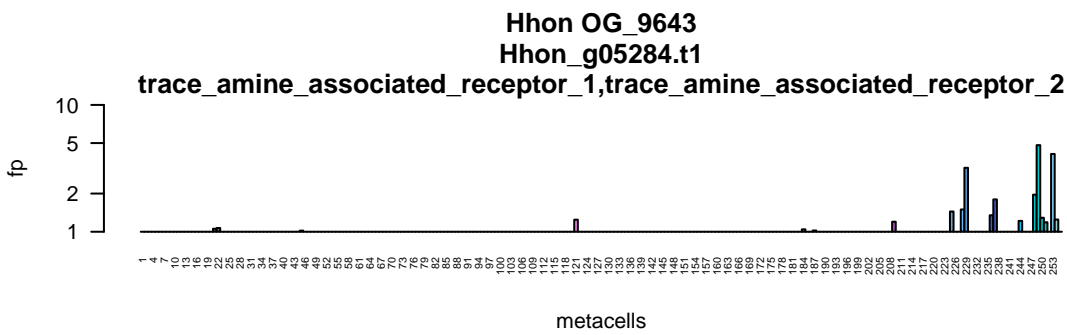
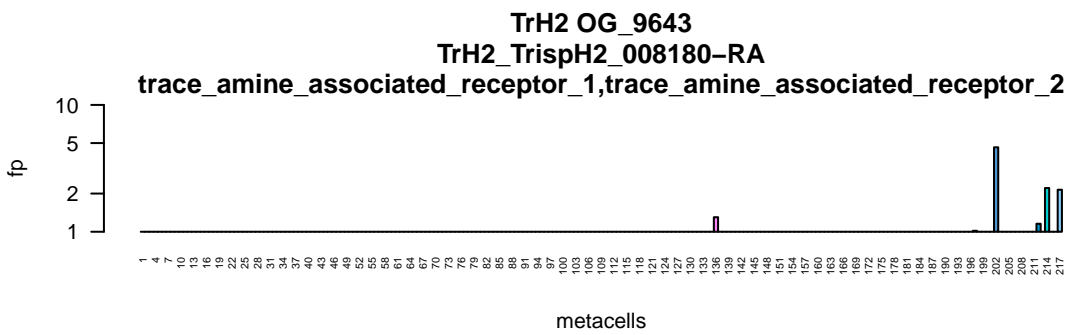
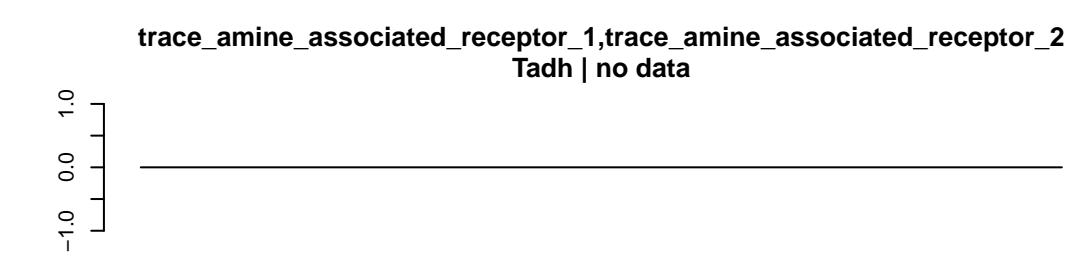


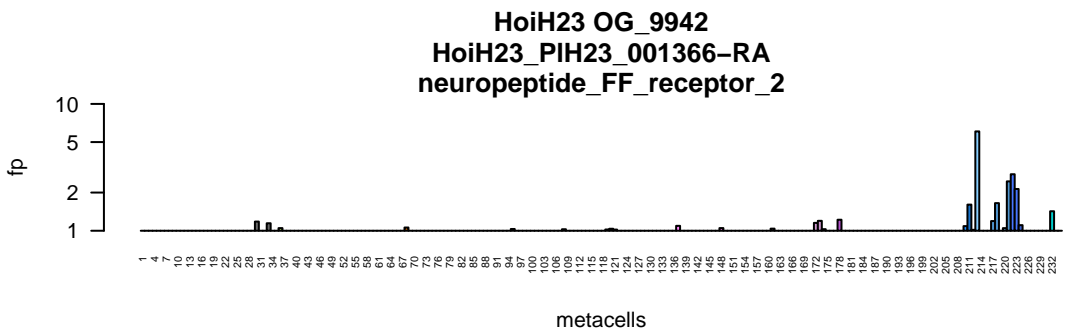
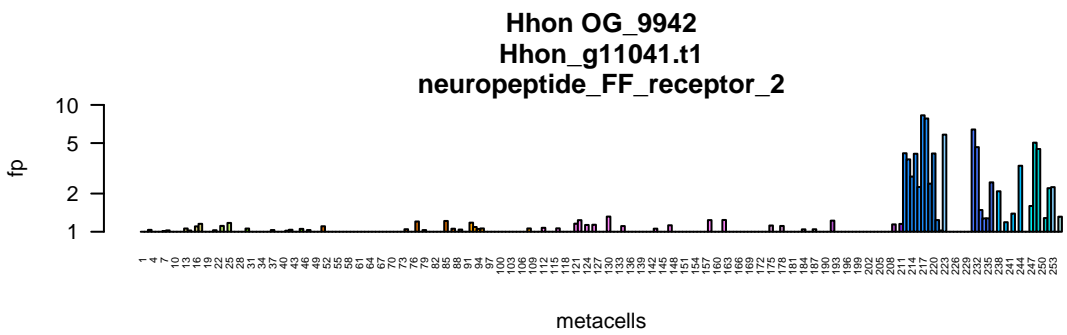
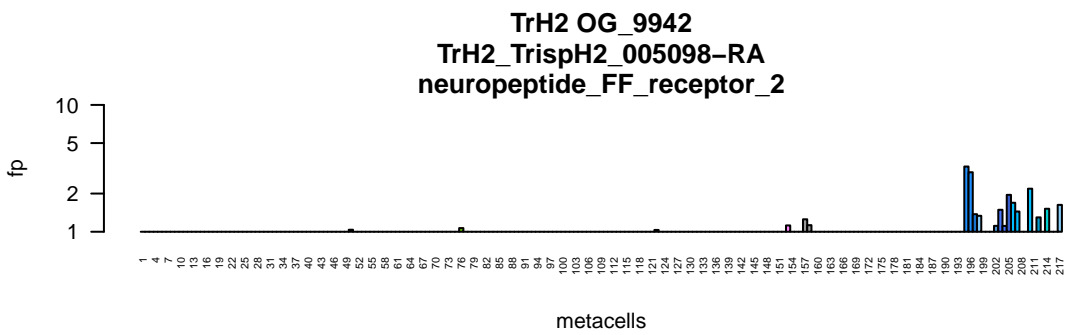
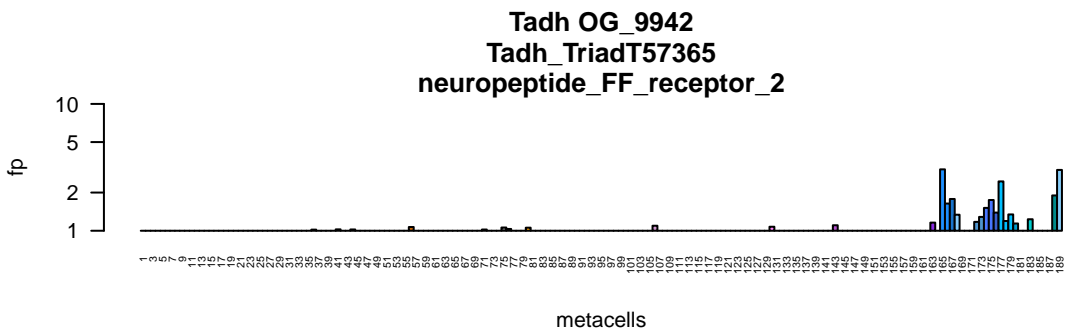
Hhon OG_9610
Hhon_g08104.t1

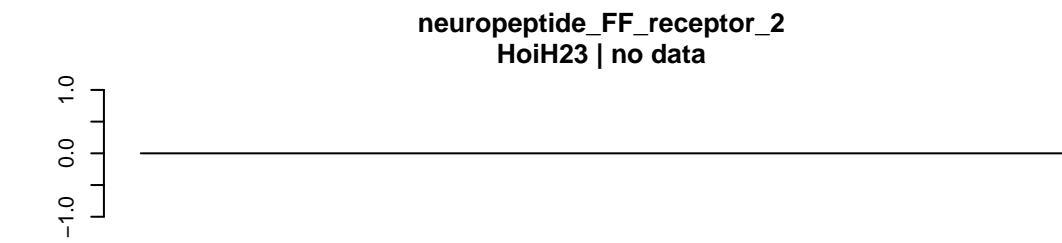
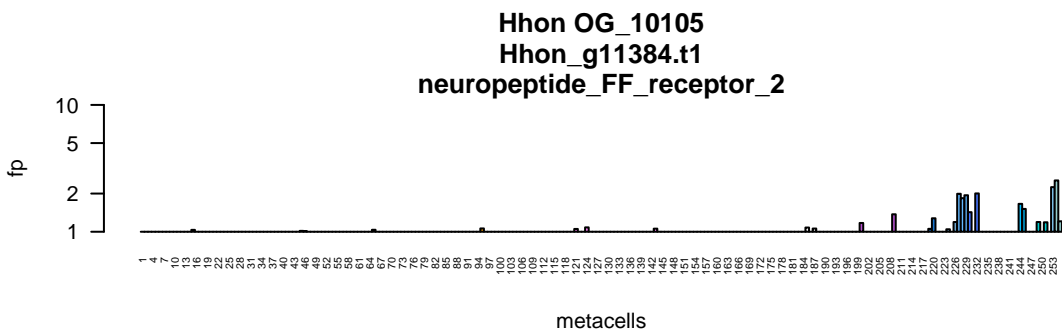
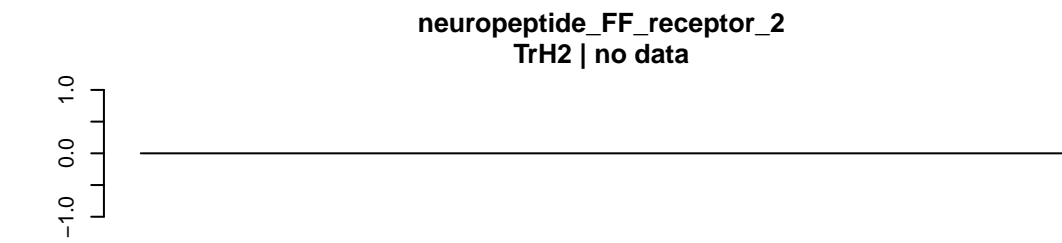
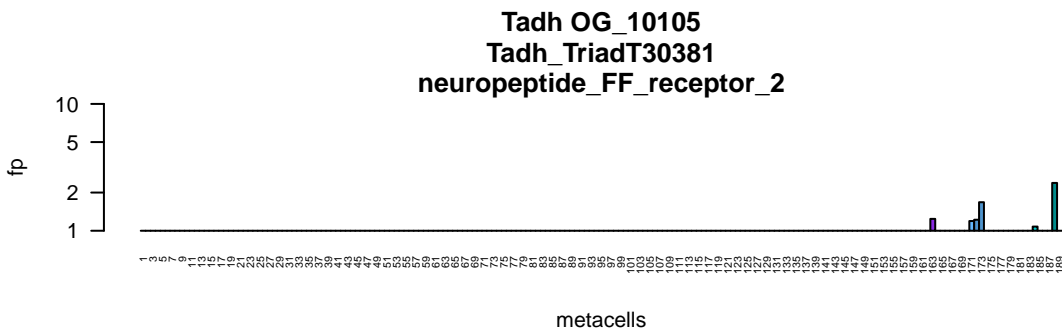


HoiH23 OG_9610
HoiH23_PIH23_005684-RA

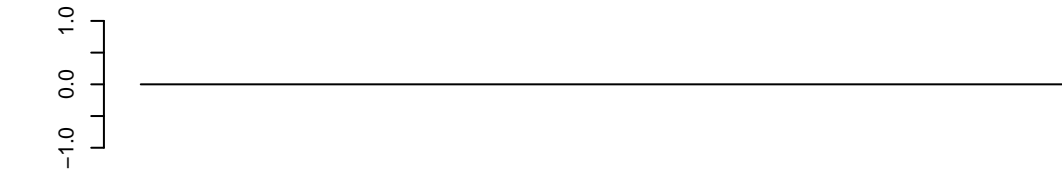




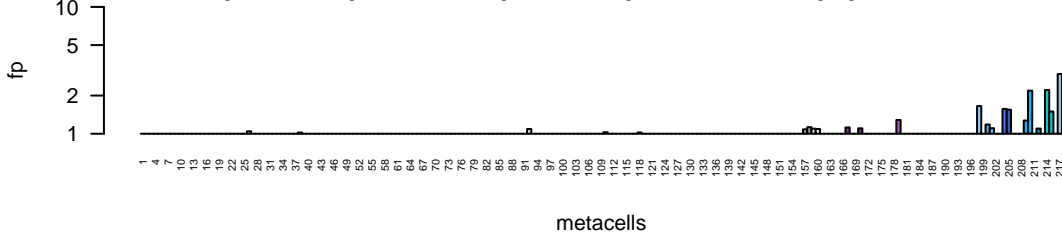




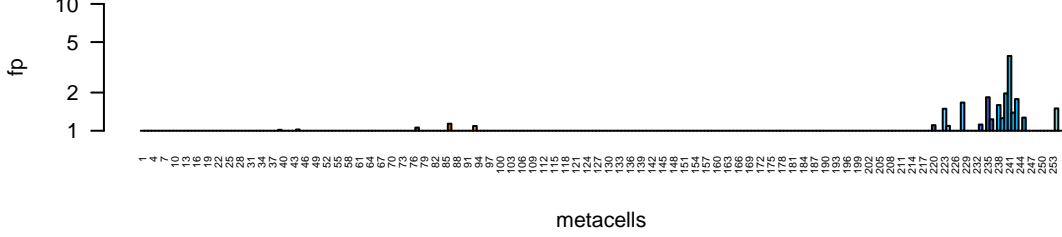
omatostatin_receptor_2,G_protein_coupled_receptor_183,neuropeptides_B_and_W_recept
Tadh | no data



omatostatin_receptor_2,G_protein_coupled_receptor_183,neuropeptides_B_and_W_recept
TrH2 OG_10233
TrH2_TrispH2_006452-RA



omatostatin_receptor_2,G_protein_coupled_receptor_183,neuropeptides_B_and_W_recept
Hhon OG_10233
Hhon_g05148.t1



omatostatin_receptor_2,G_protein_coupled_receptor_183,neuropeptides_B_and_W_recept
HoiH23 OG_10233
HoiH23_PIH23_001231-RA

