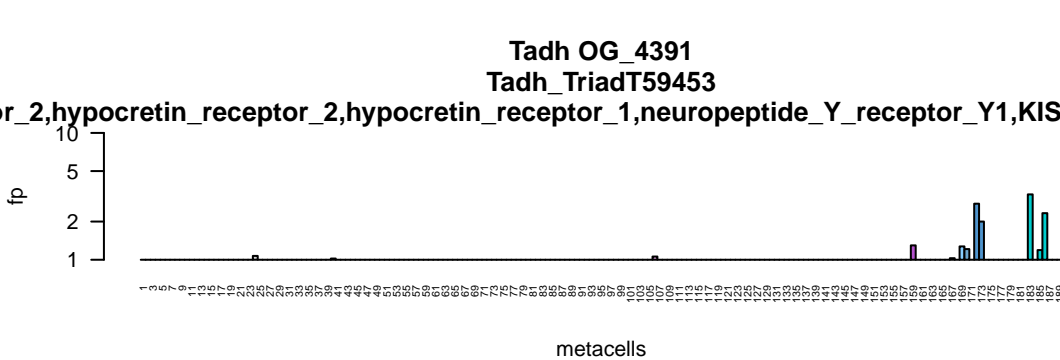
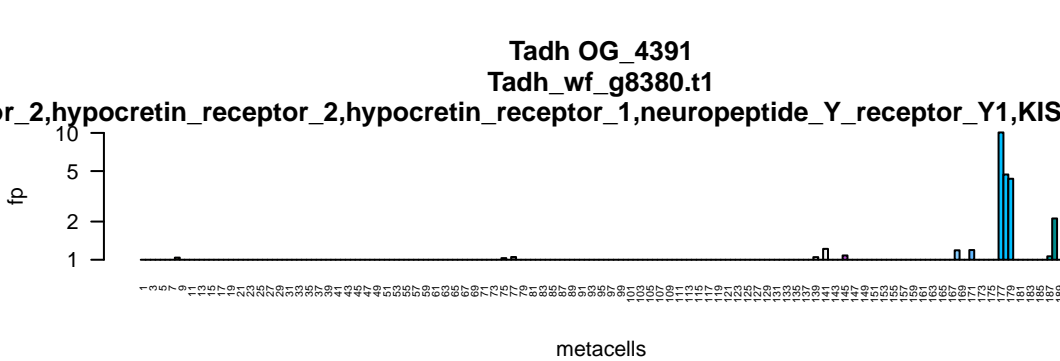


or_2,hypocretin_receptor_2,hypocretin_receptor_1,neuropeptide_Y_receptor_Y1,KIS

metacells	fp
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14	1
15	1
16	1
17	1
18	1
19	1
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21	1
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172	1
173	1
174	1
175	1
176	1
177	2
178	1
179	2
180	1
181	1
182	1
183	1
184	1
185	1
186	1
187	1



r_2,hypocretin_receptor_2,hypocretin_receptor_1,neuropeptide_Y_receptor_Y1,KIS



r_2,hypocretin_receptor_2,hypocretin_receptor_1,neuropeptide_Y_receptor_Y1,KIS

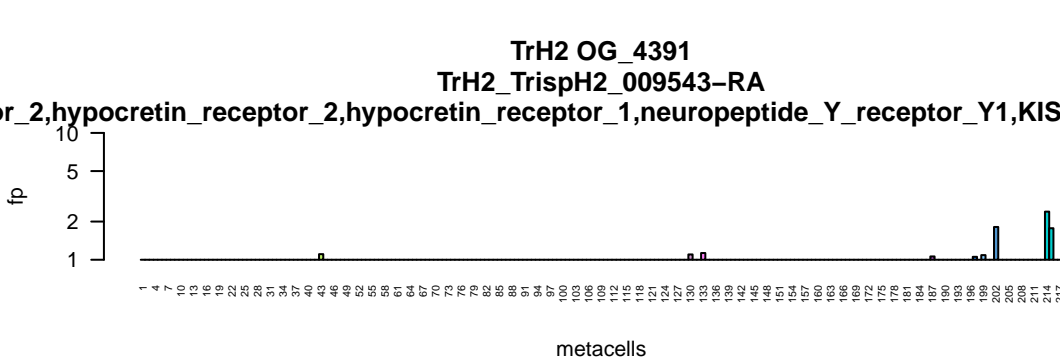
fp

metacells

metacells	fp
r_2	1
hypocretin_receptor_2	1
hypocretin_receptor_1	1
neuropeptide_Y_receptor_Y1	10
KIS	1
...	...
metacells_40	1

or_2,hypocretin_receptor_2,hypocretin_receptor_1,neuropeptide_Y_receptor_Y1,KIS

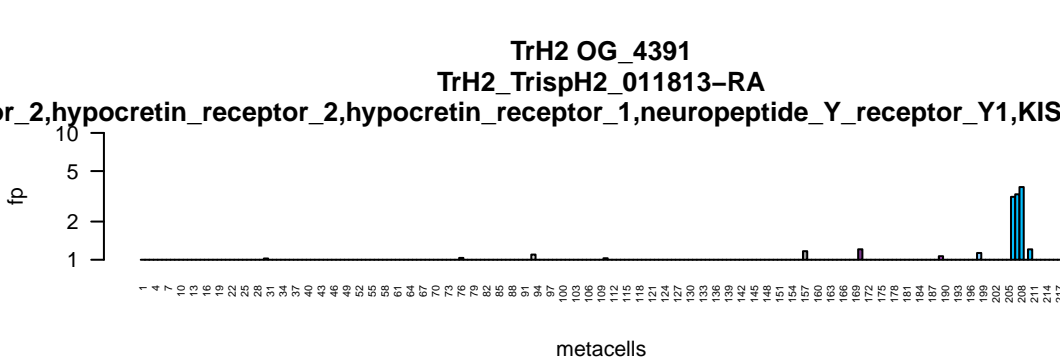
metacells	fp
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137	1
139	1
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147	1
149	1
153	1
157	1
161	1
163	1
167	1
171	1
173	1
177	1
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183	1
187	1



or_2,hypocretin_receptor_2,hypocretin_receptor_1,neuropeptide_Y_receptor_Y1,KIS

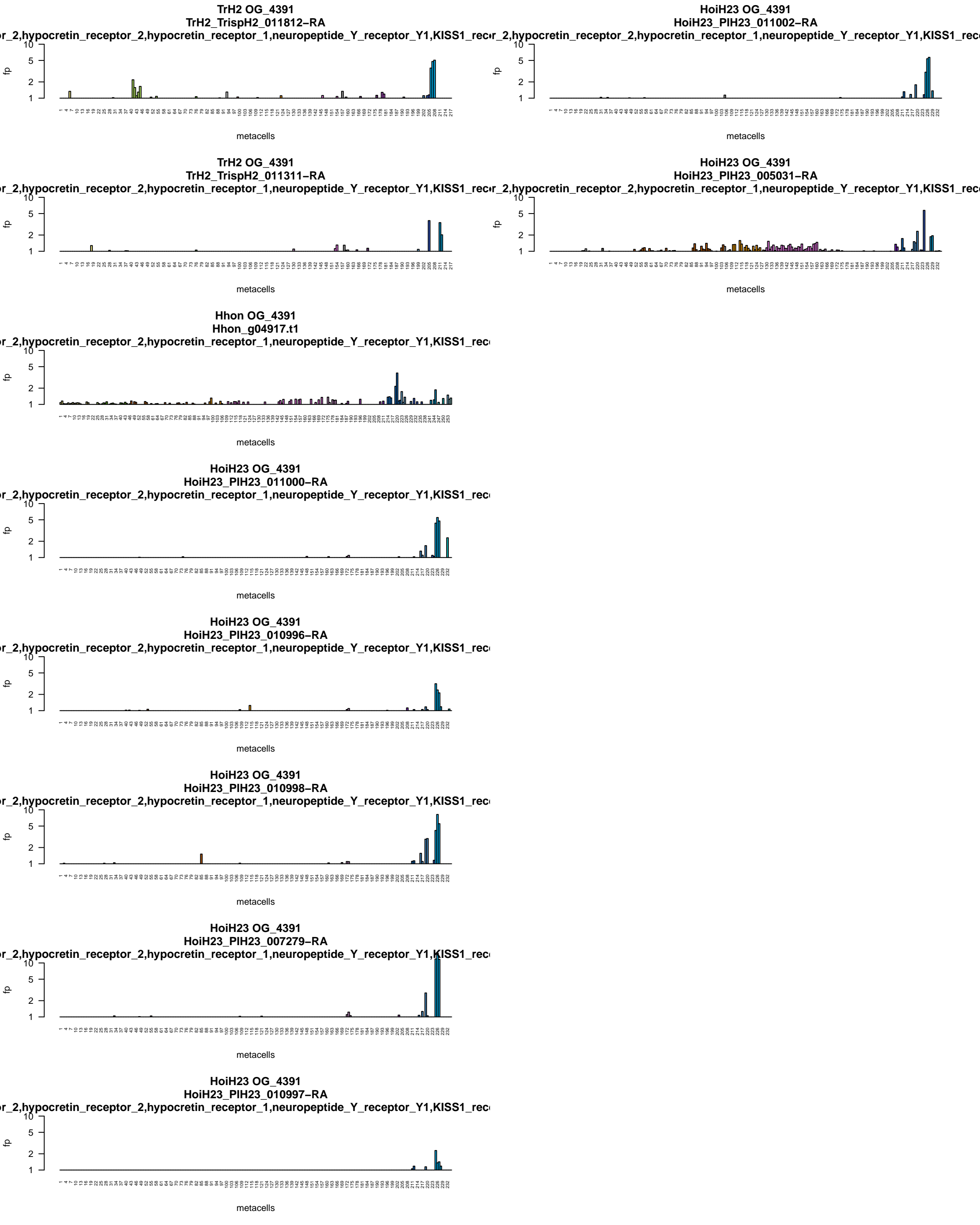
metacell	fp
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172	1
175	1
178	1
181	1
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metacells

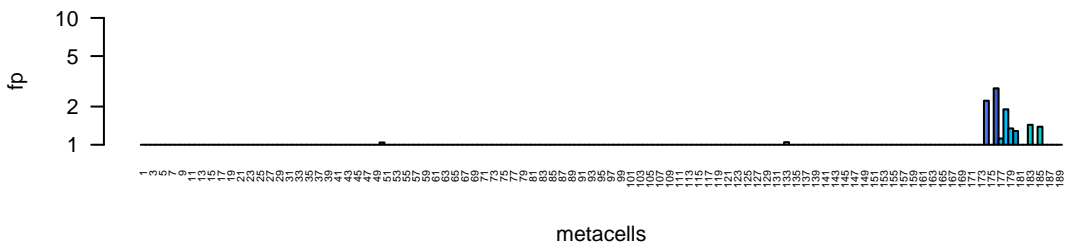


r_2,hypocretin_receptor_2,hypocretin_receptor_1,neuropeptide_Y_receptor_Y1,KIS

metacell	fp
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16	1
19	1
22	1
25	1
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157	1
160	1
163	1
166	1
169	1.5
172	1.5
175	1
178	1
181	1.5
184	1
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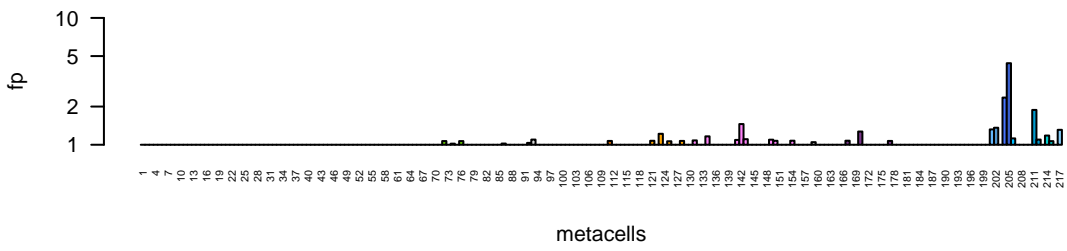


Tadh OG_8094
Tadh_wf_g6497.t1



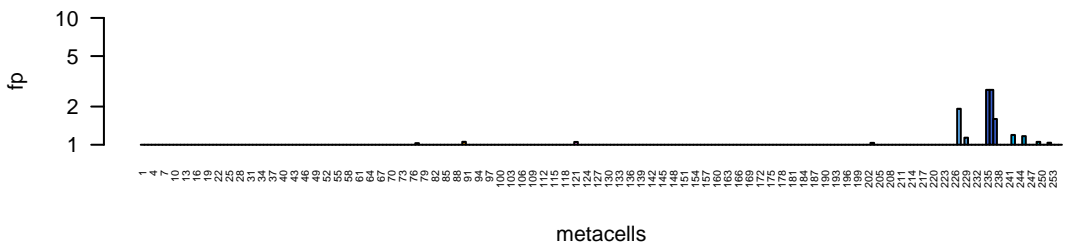
metacells

TrH2 OG_8094
TrH2_TrispH2_011737-RA



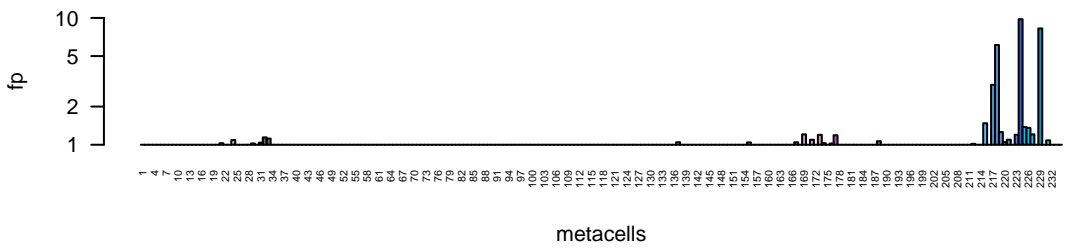
metacells

Hhon OG_8094
Hhon_g08752.t1



metacells

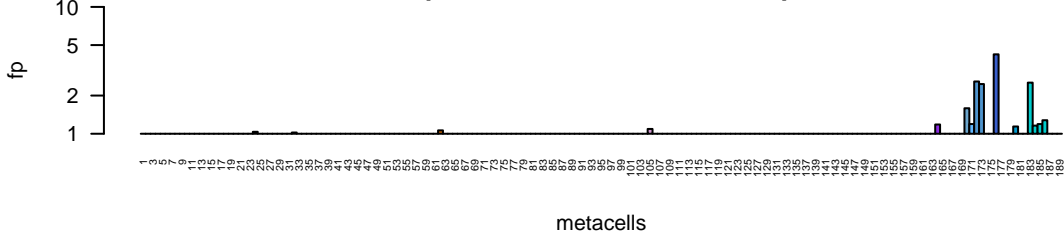
HoiH23 OG_8094
HoiH23_PIH23_007147-RA



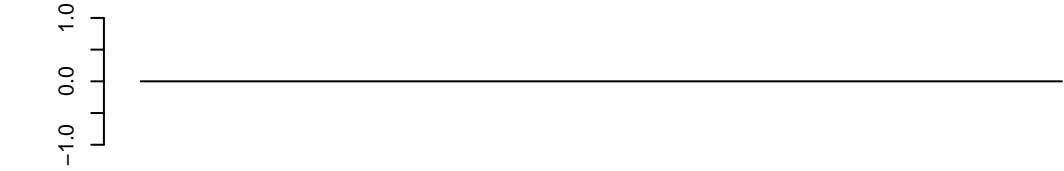
metacells

Tadh OG_4982
Tadh_wf_g7610.t1

trace_amine_associated_receptor_5,adenosine_A2b_receptor,adenosine_A2a_receptor

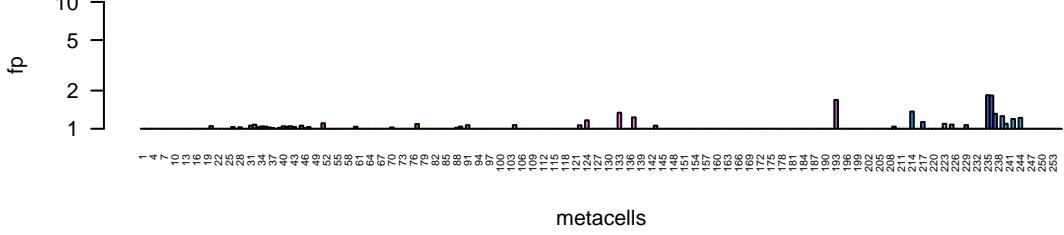


trace_amine_associated_receptor_5,adenosine_A2b_receptor,adenosine_A2a_receptor
TrH2 | no data

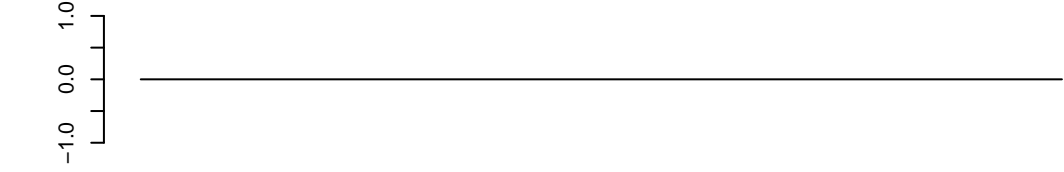


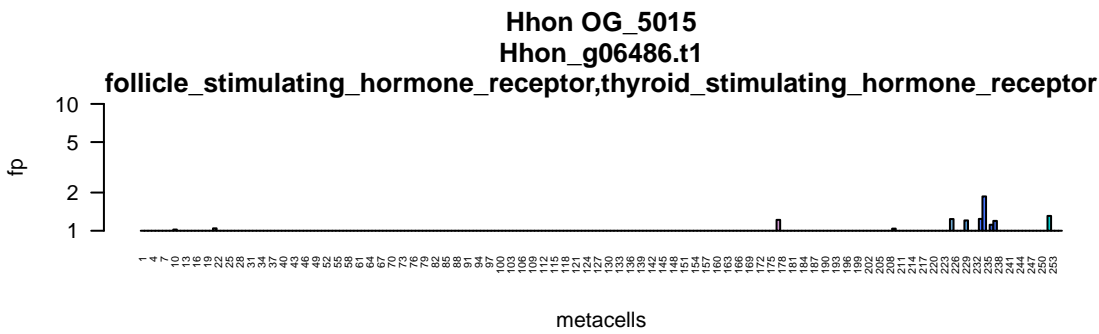
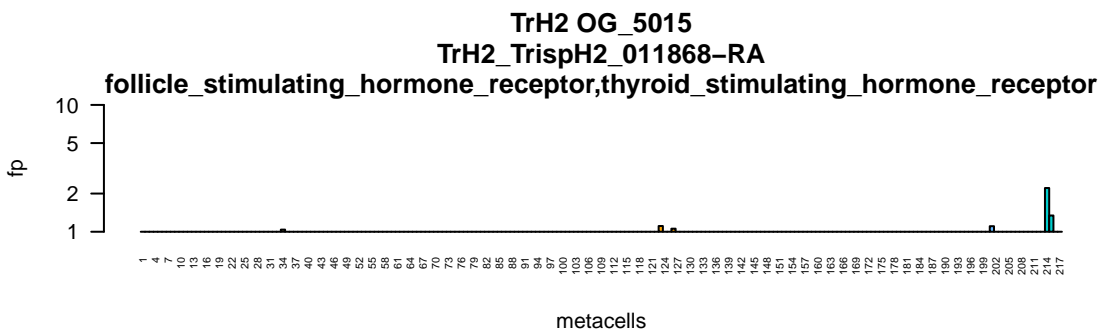
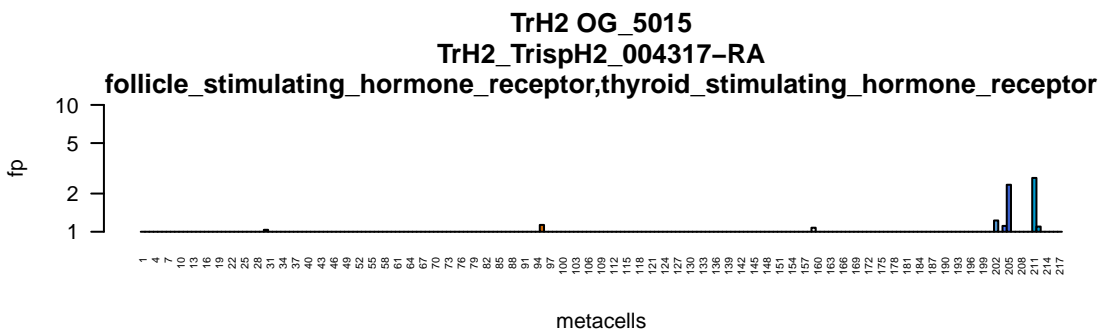
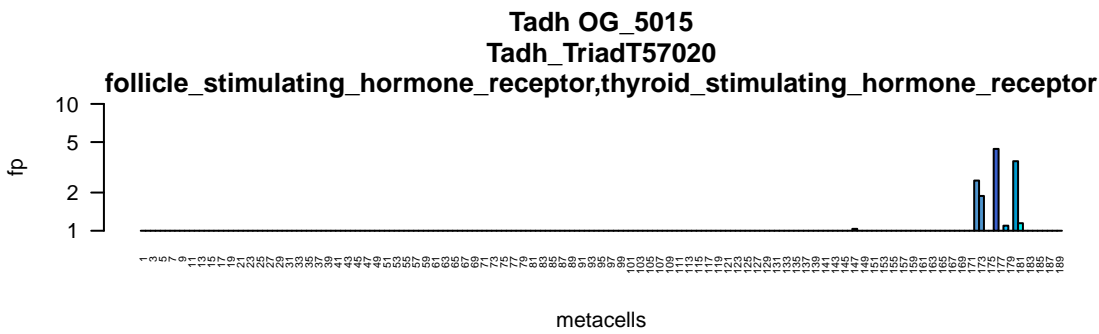
Hhon OG_4982
Hhon_g01340.t1

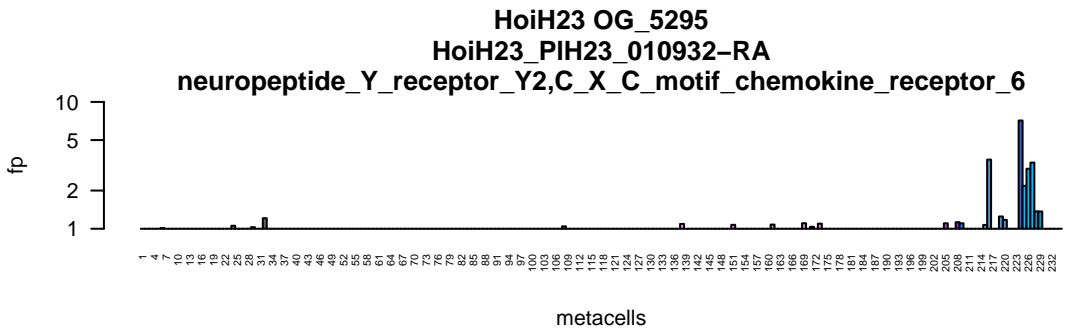
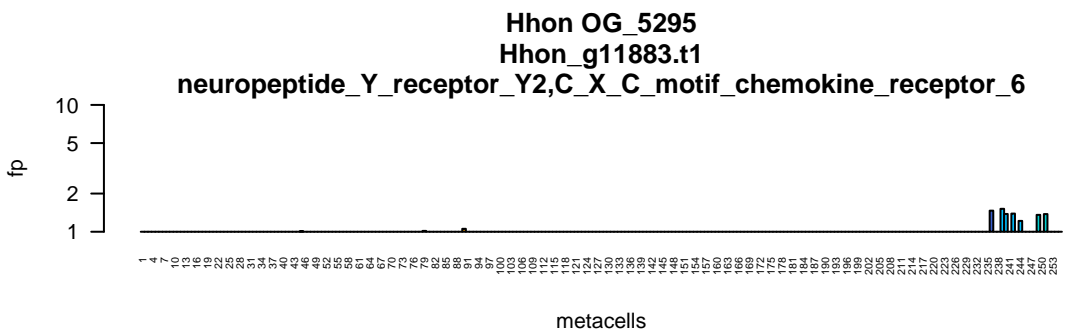
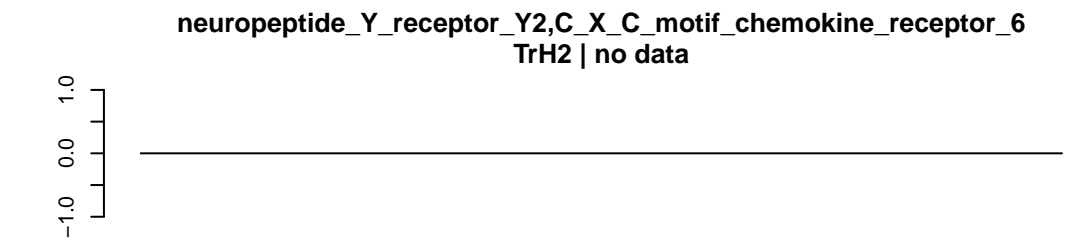
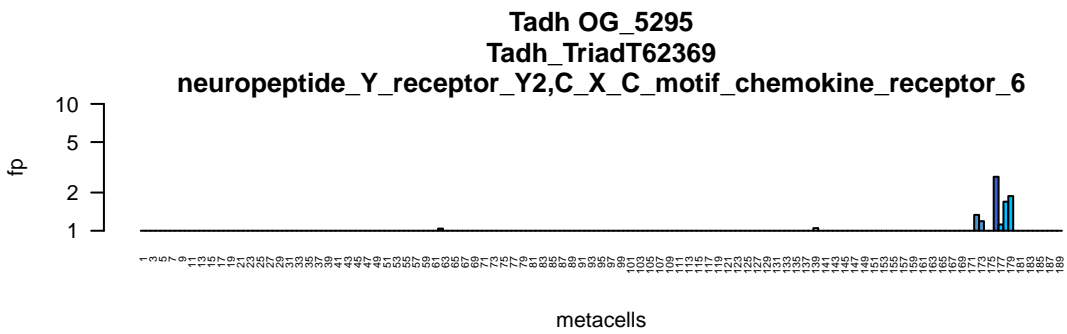
trace_amine_associated_receptor_5,adenosine_A2b_receptor,adenosine_A2a_receptor



trace_amine_associated_receptor_5,adenosine_A2b_receptor,adenosine_A2a_receptor
HoiH23 | no data



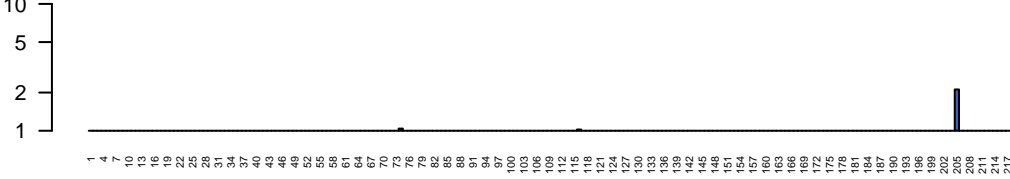




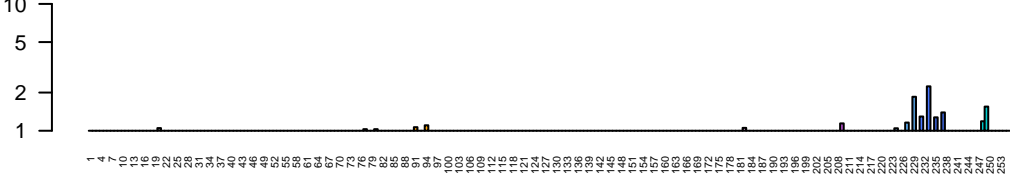
adhesion_G_protein_coupled_receptor_G4,adhesion_G_protein_coupled_receptor_G2
Tadh | no data



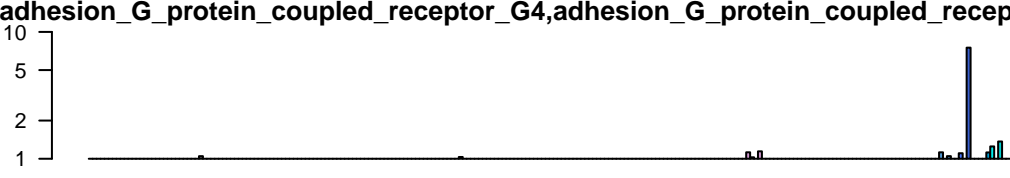
TrH2 OG_8020
TrH2_TrispH2_004376-RA
adhesion_G_protein_coupled_receptor_G4,adhesion_G_protein_coupled_receptor_G2

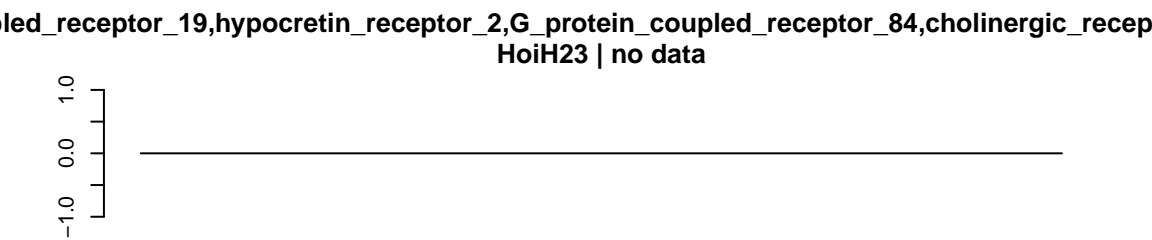
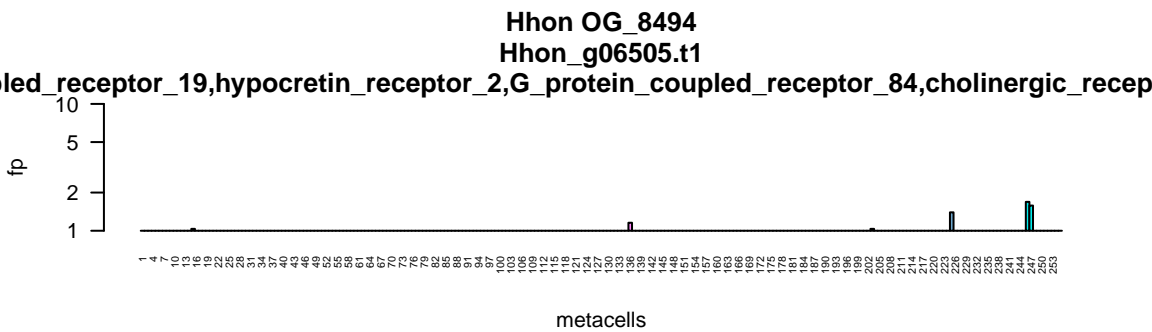
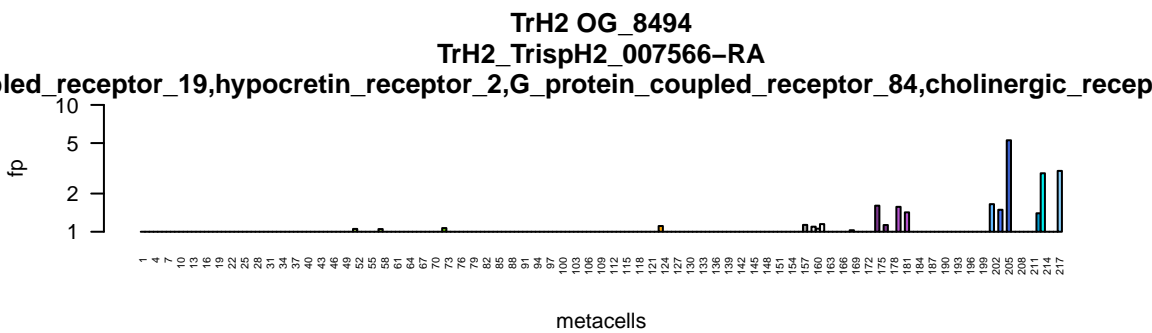
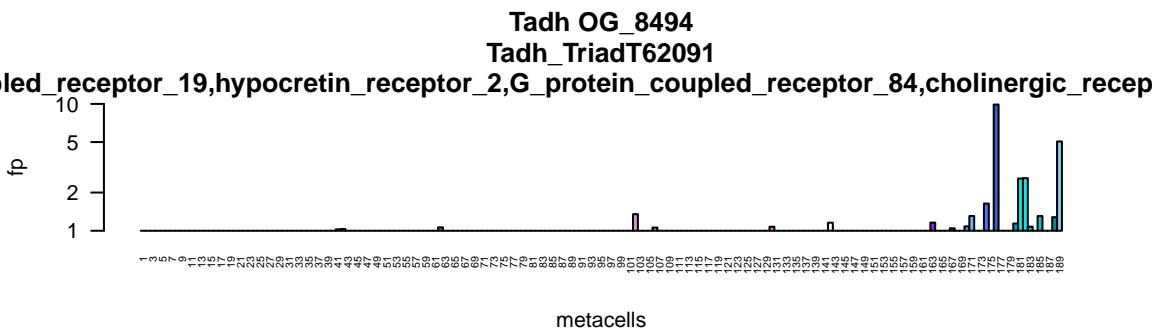


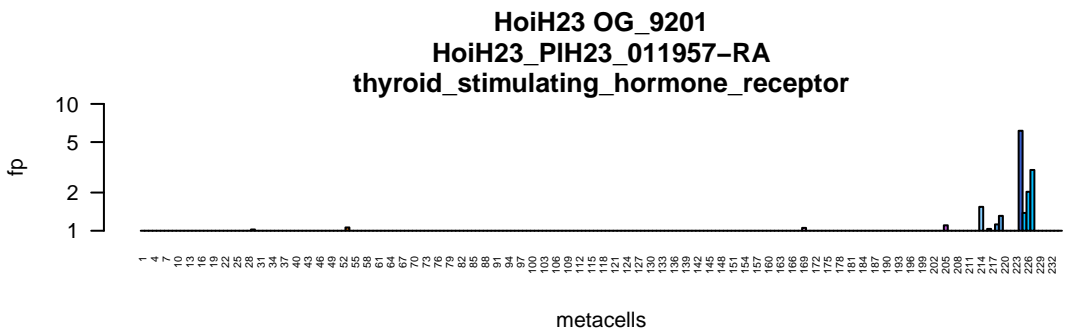
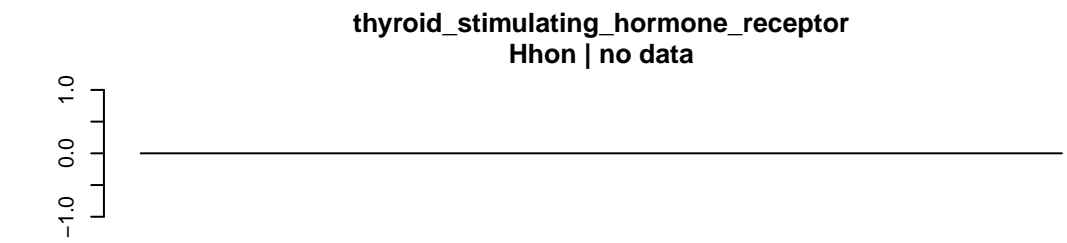
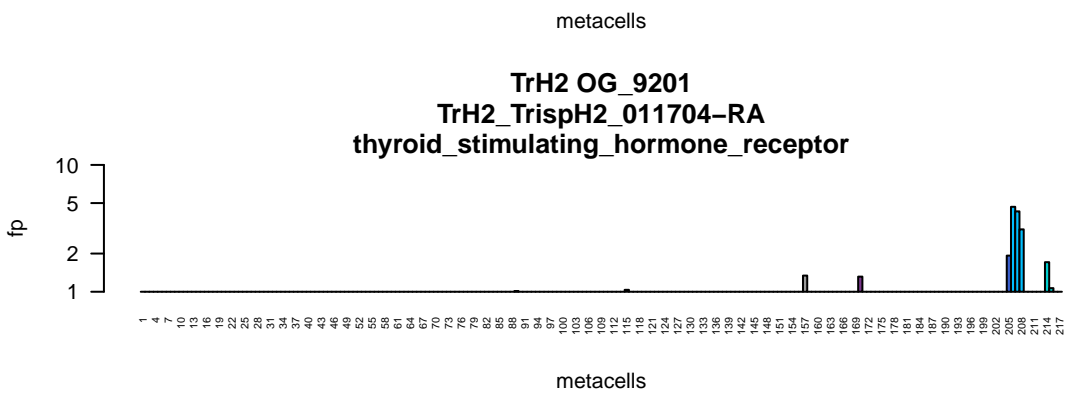
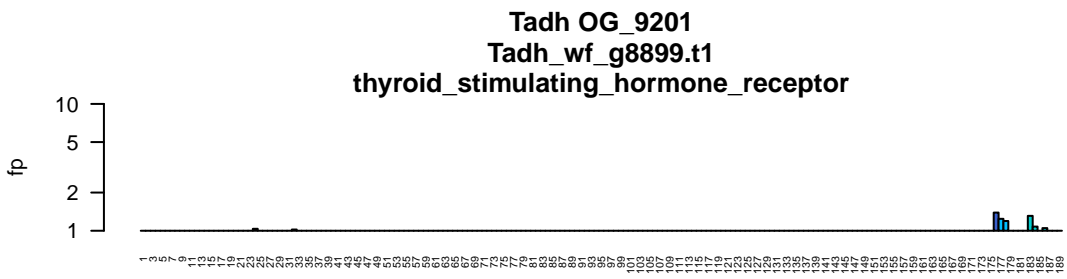
Hhon OG_8020
Hhon_g07984.t1
adhesion_G_protein_coupled_receptor_G4,adhesion_G_protein_coupled_receptor_G2

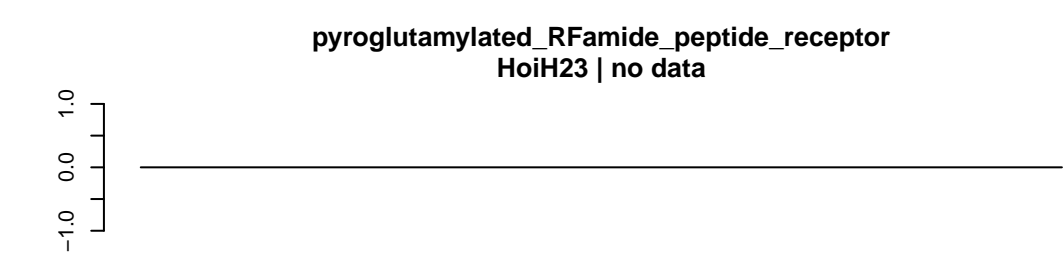
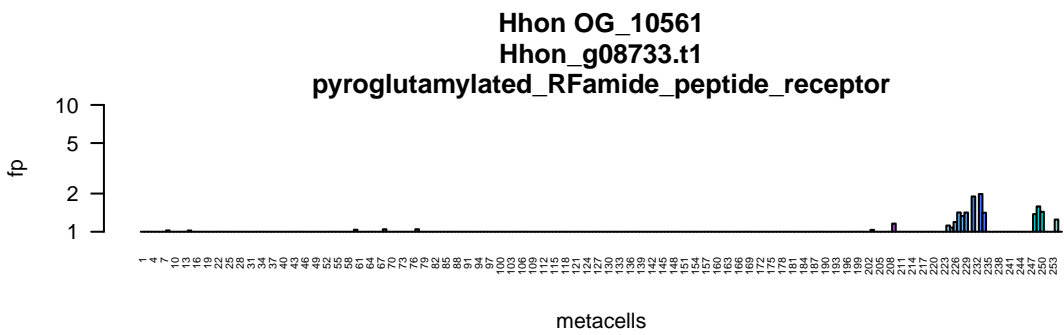
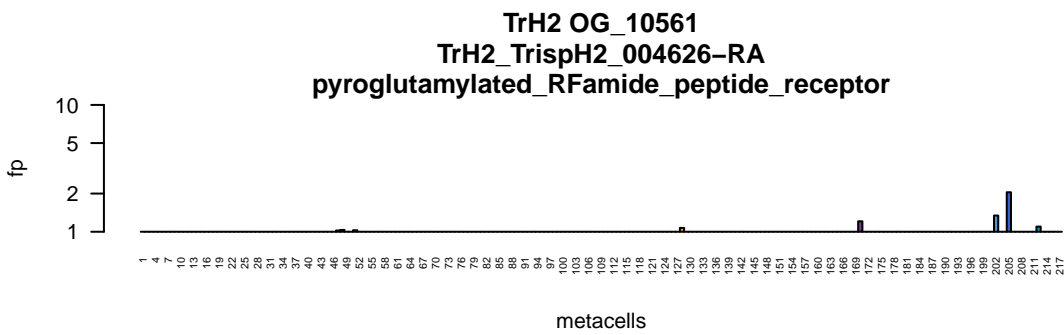
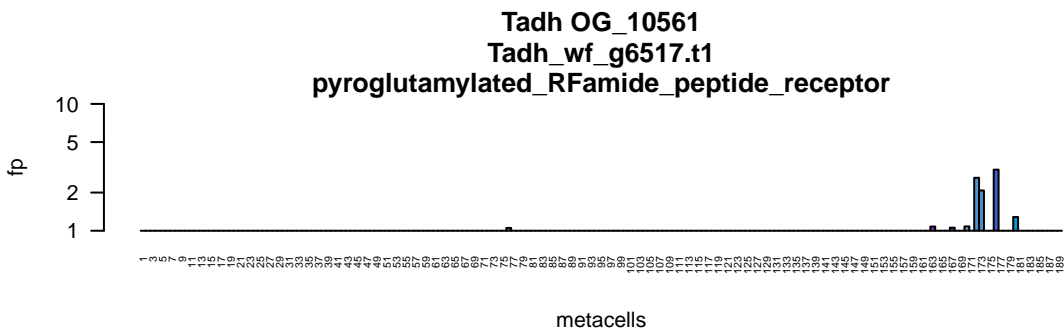


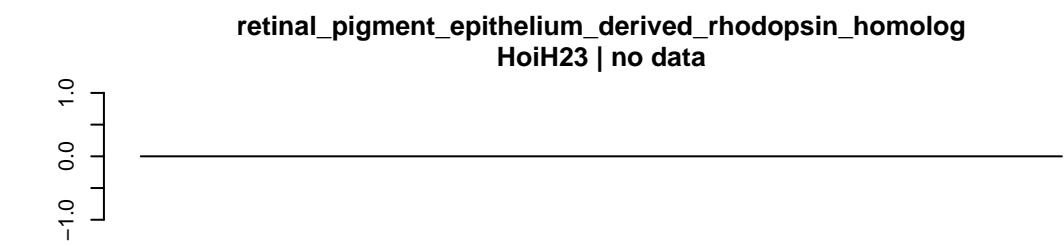
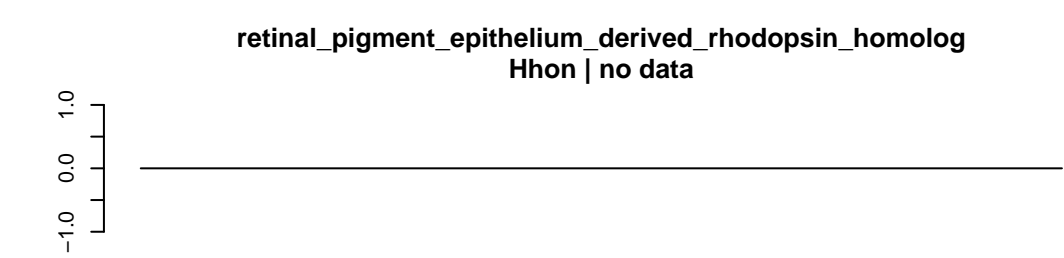
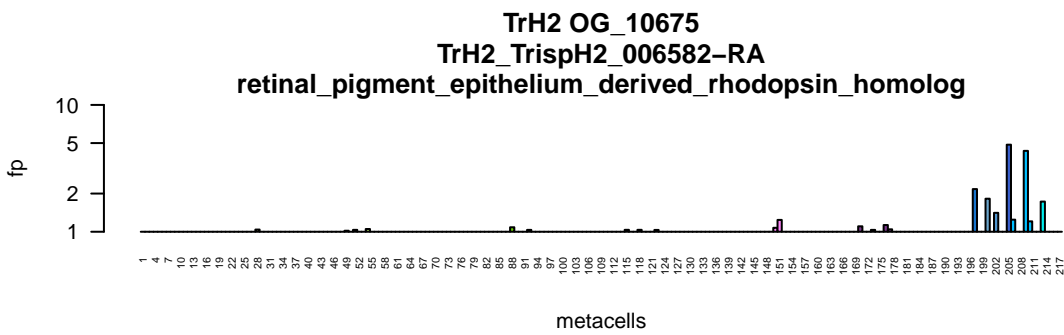
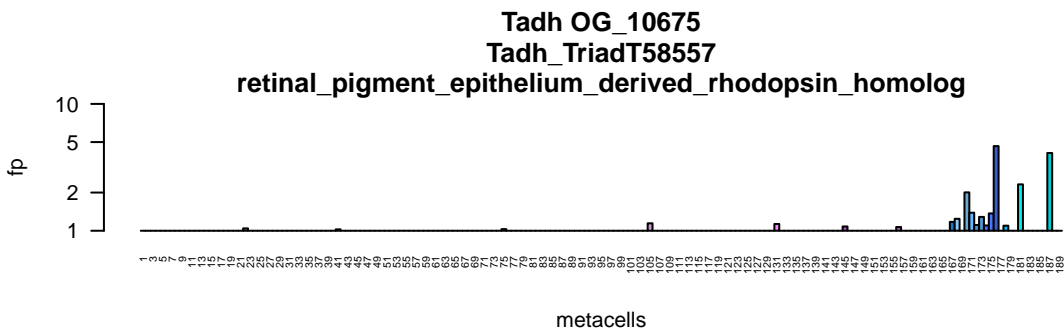
HoiH23 OG_8020
HoiH23_PIH23_006151-RA
adhesion_G_protein_coupled_receptor_G4,adhesion_G_protein_coupled_receptor_G2

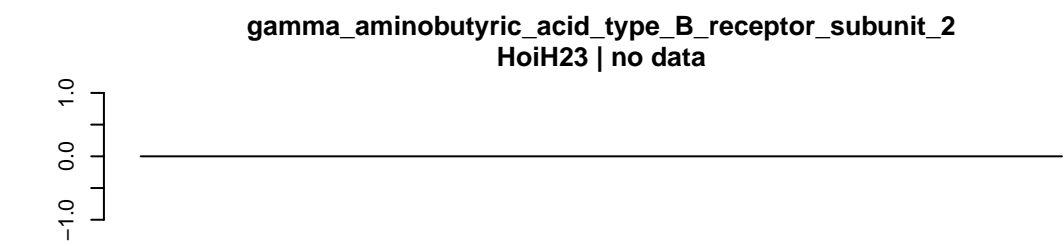
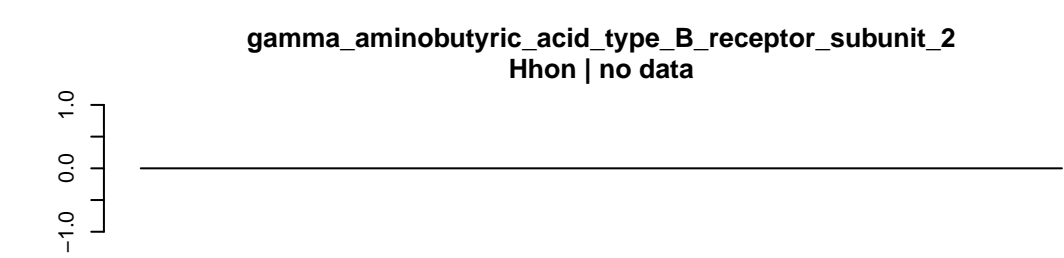
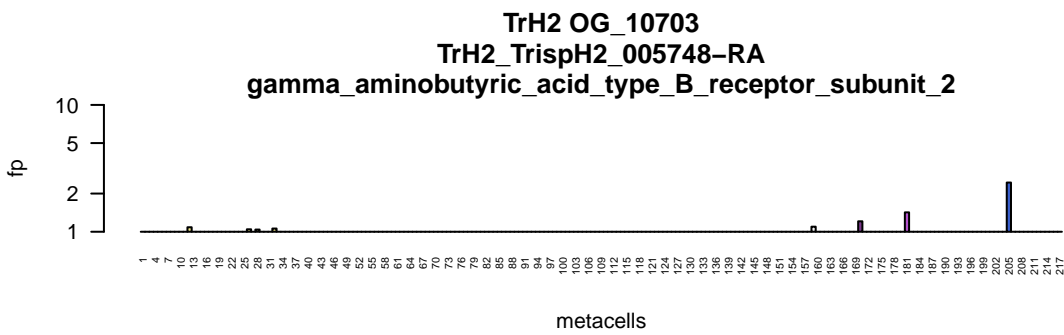
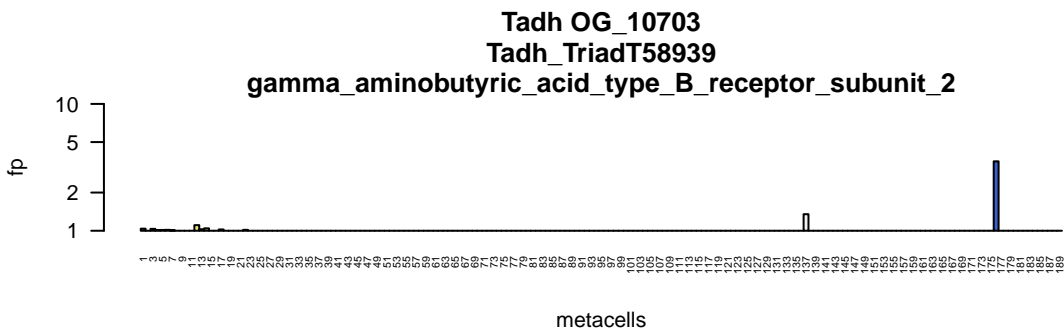


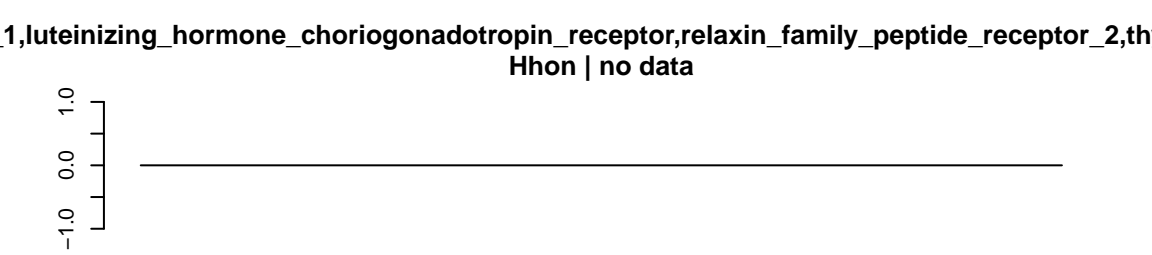
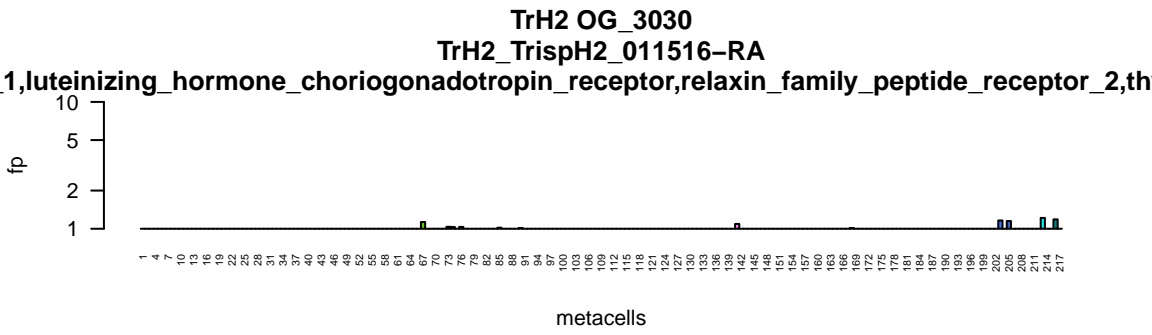
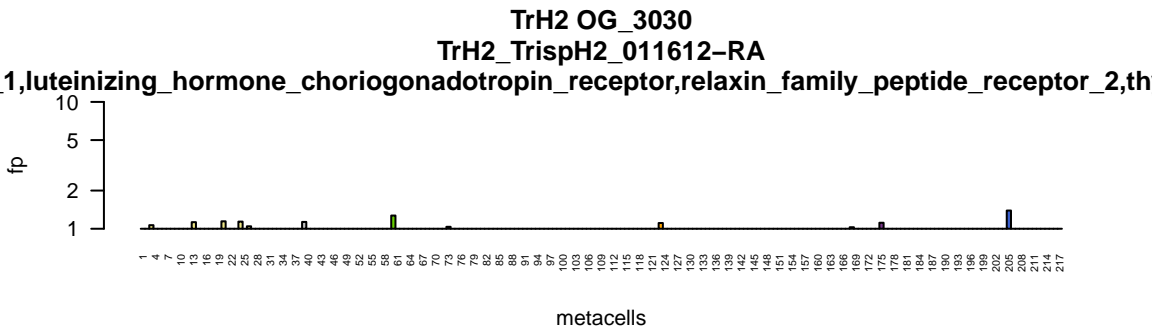
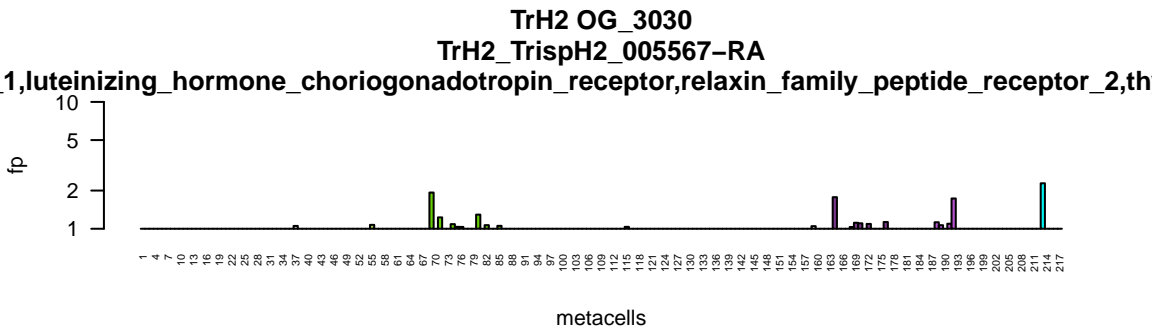
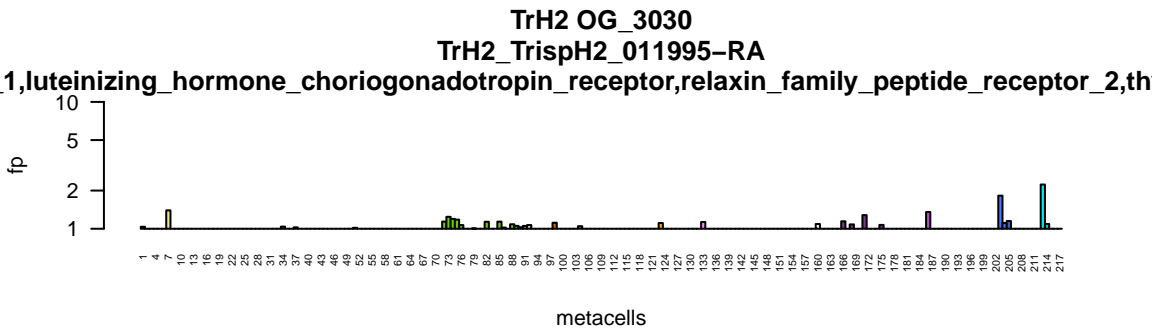
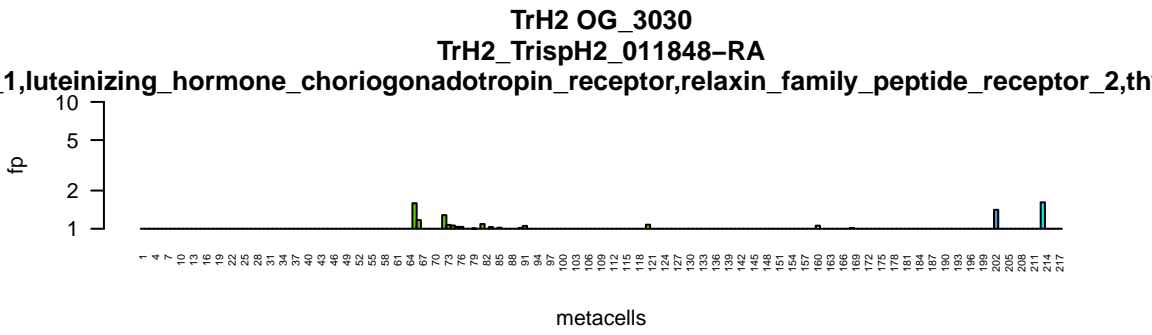
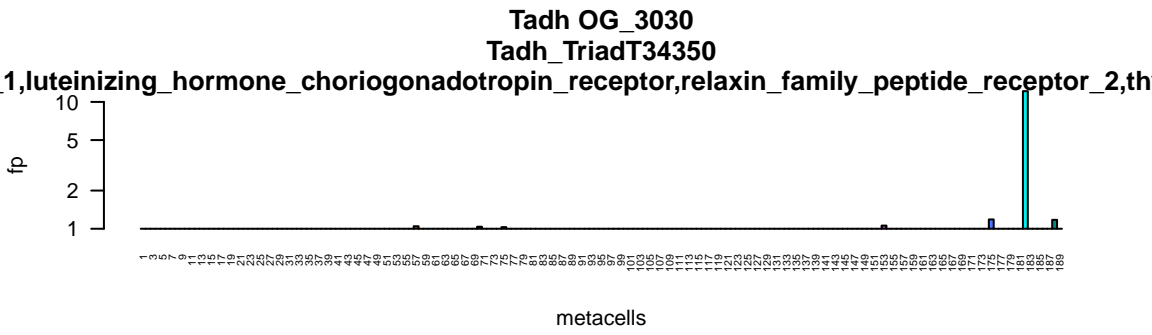
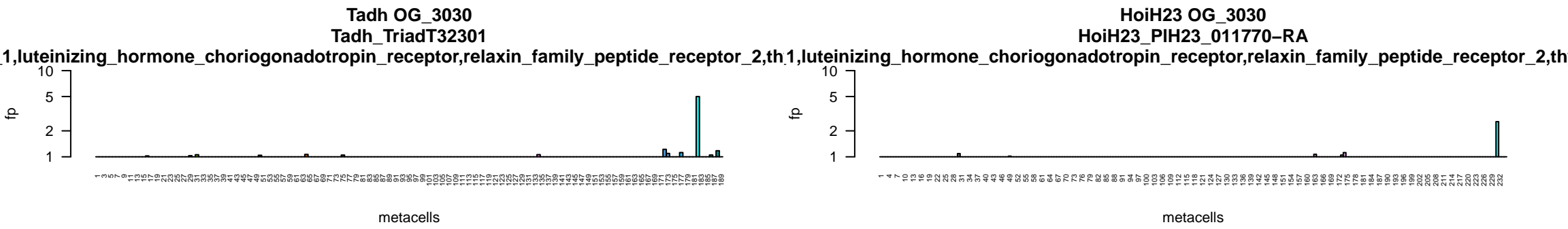


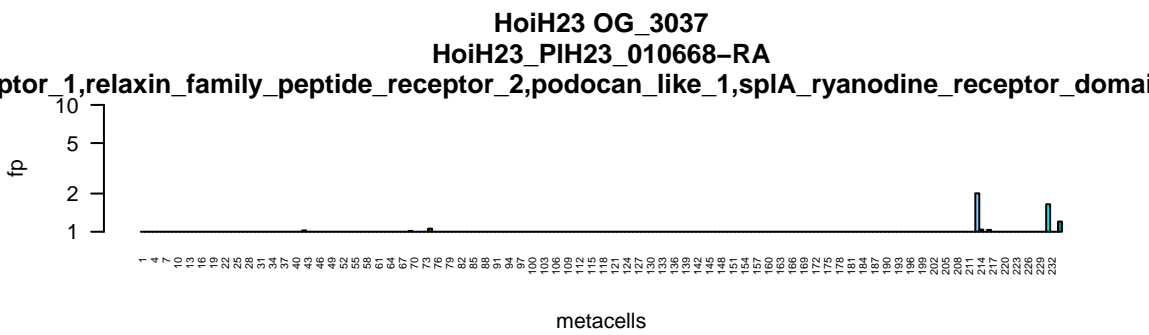
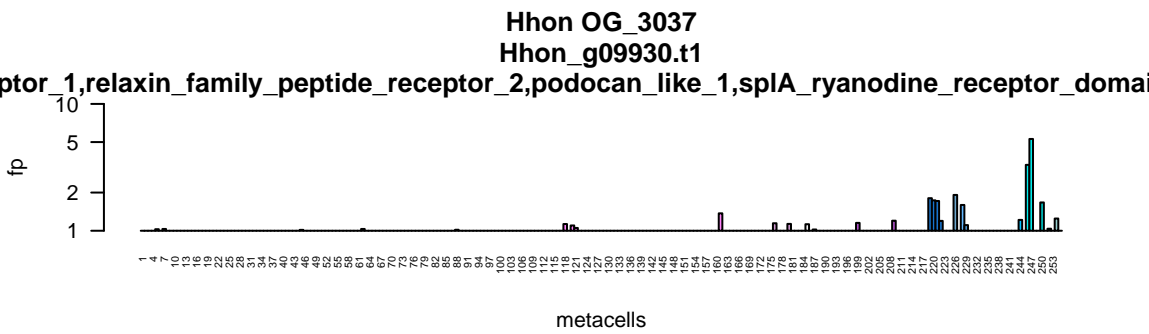
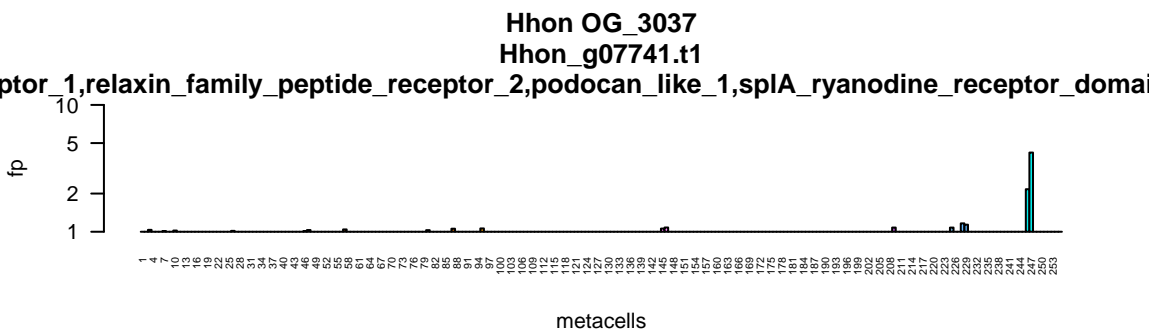
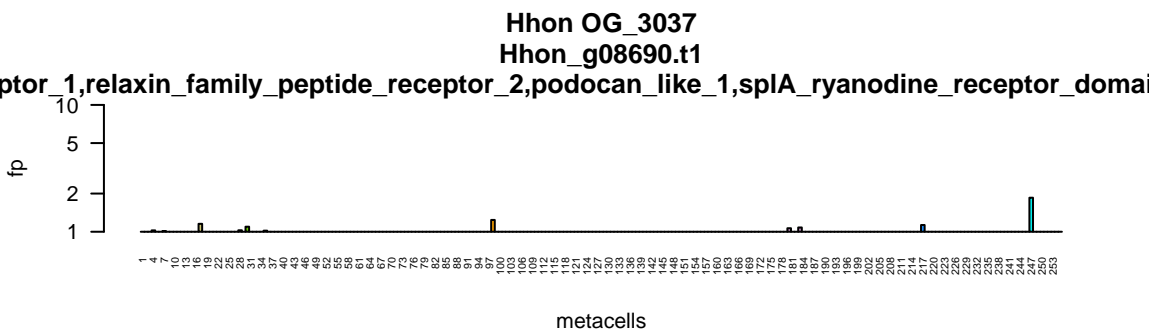
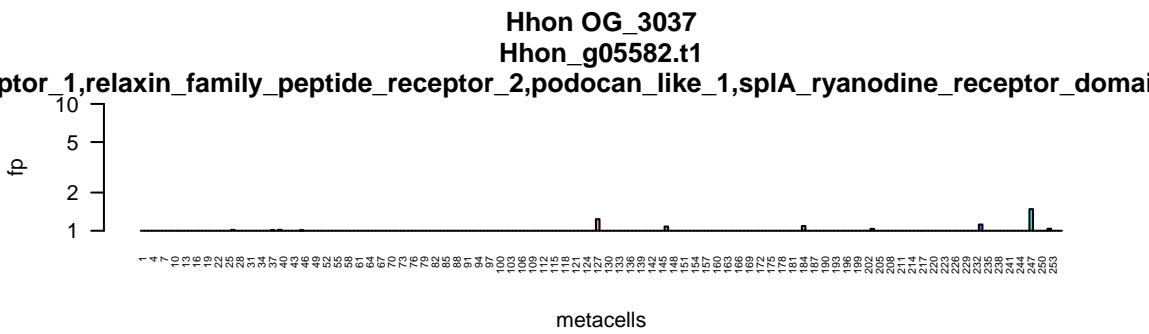
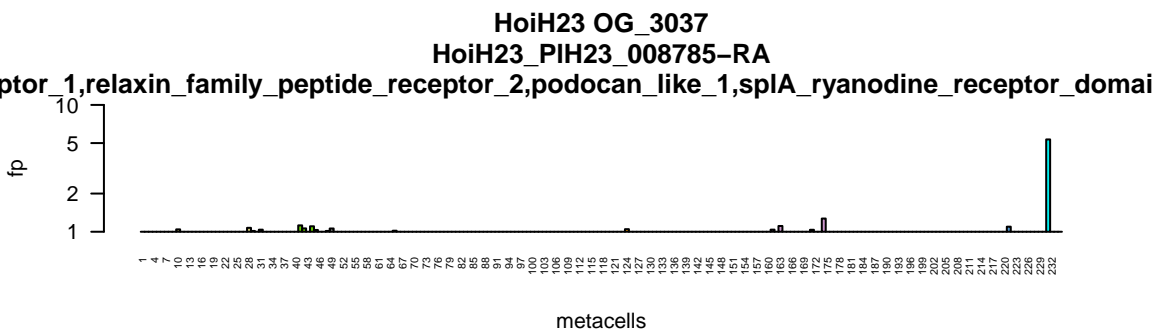
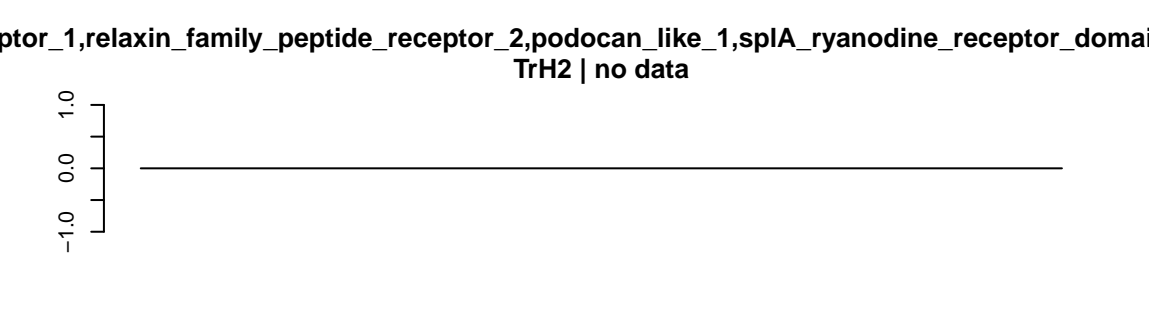
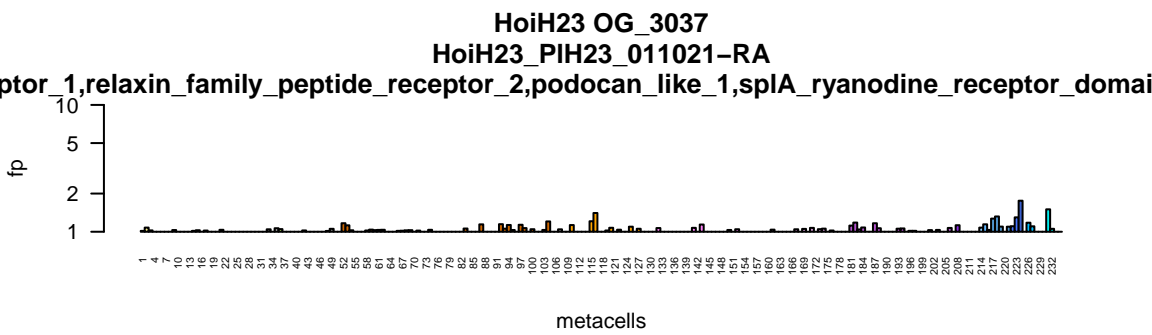
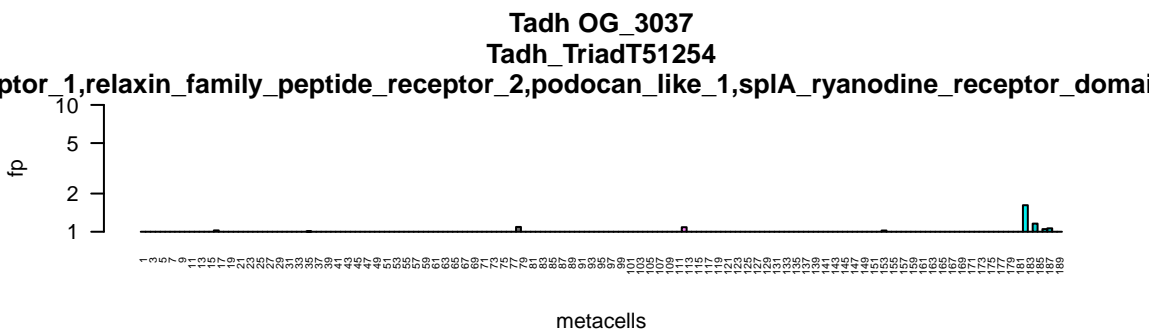
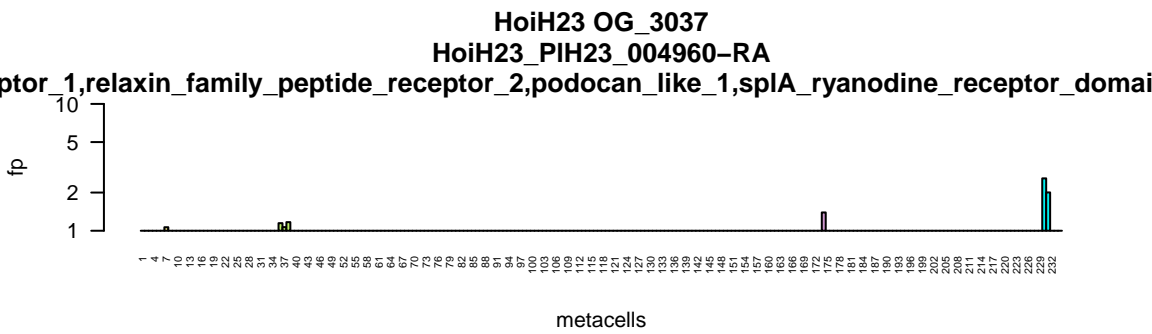
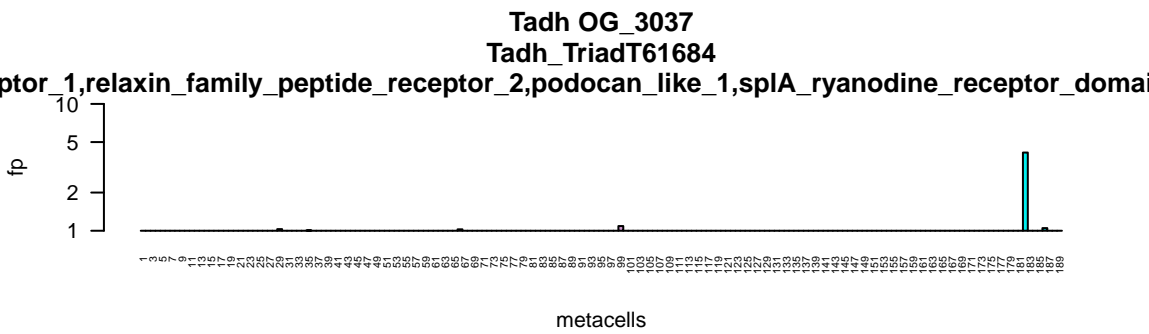


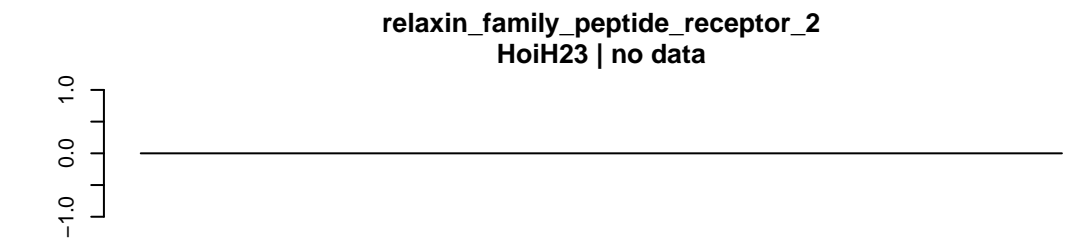
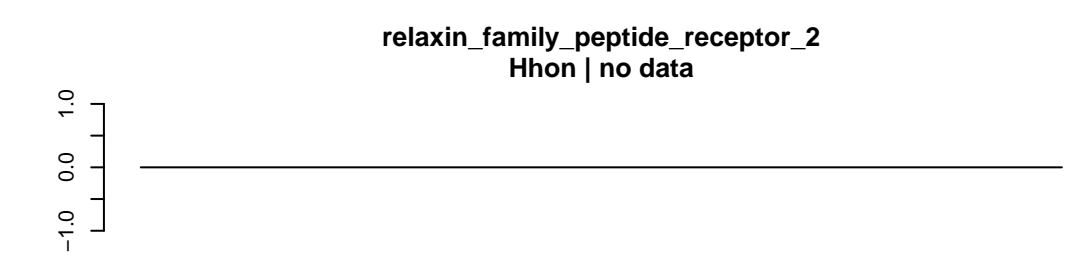
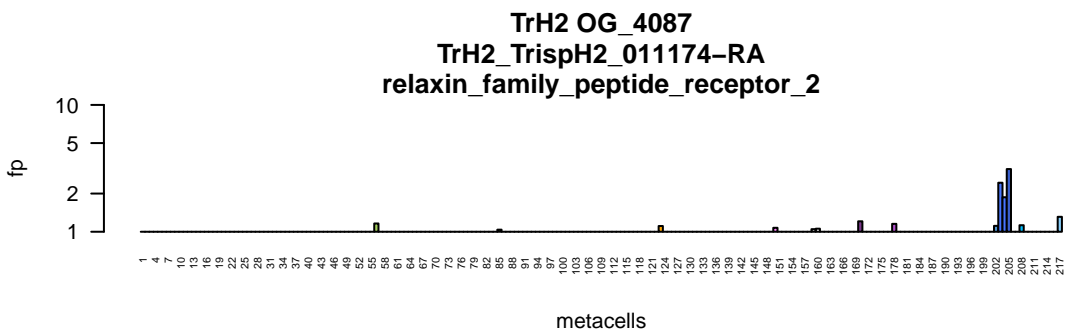
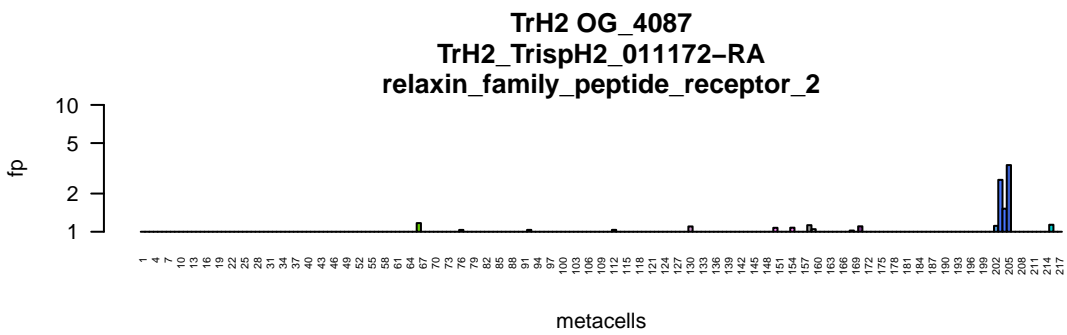
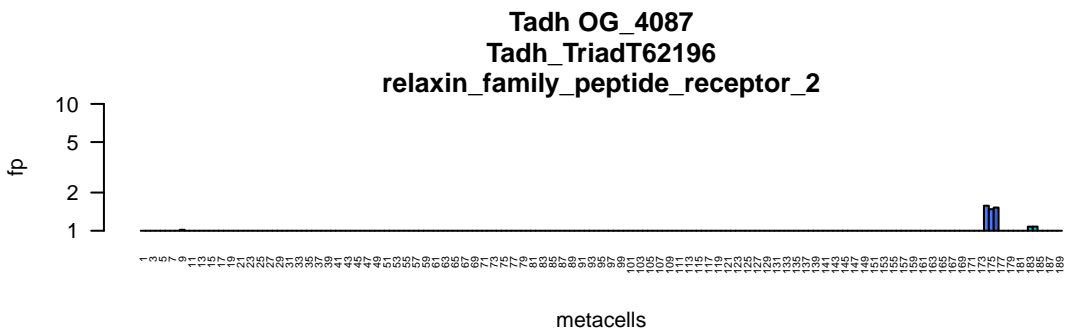


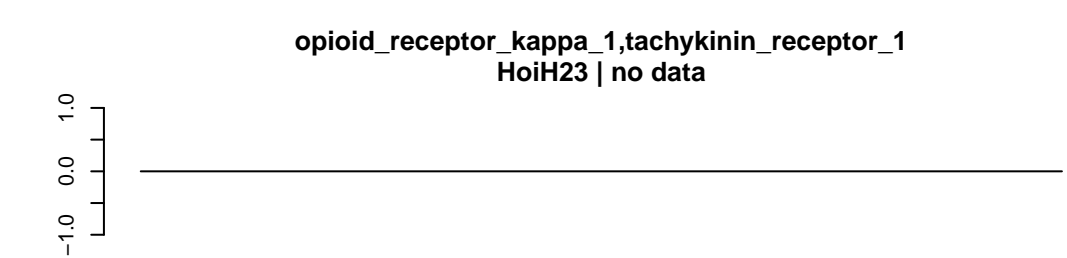
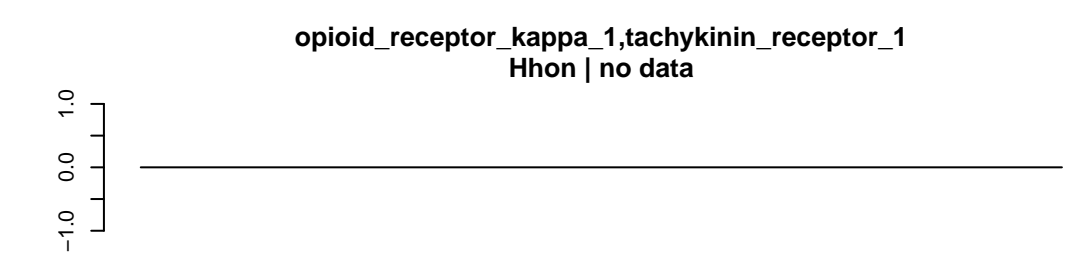
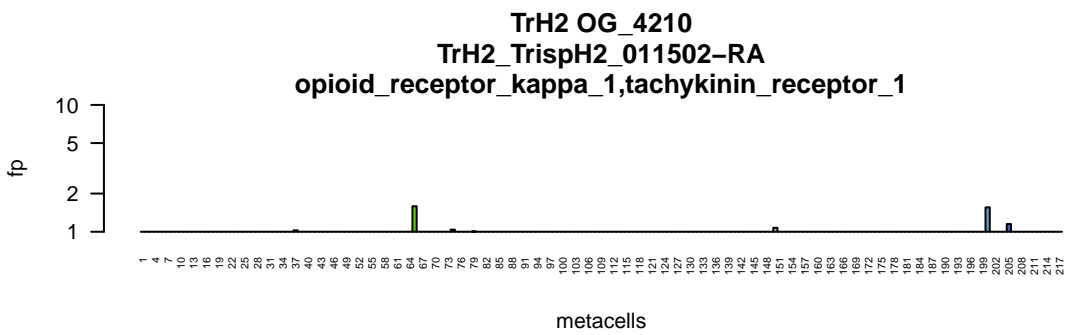
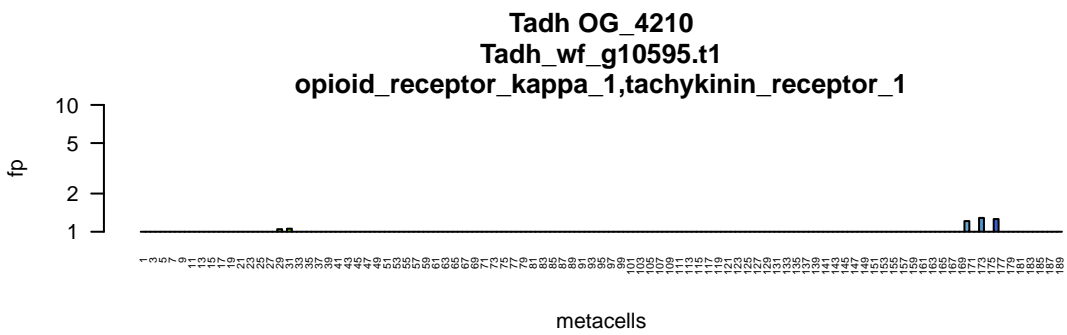
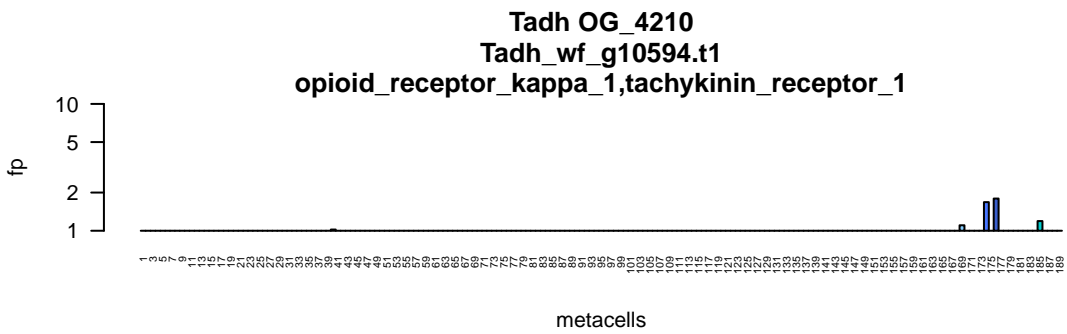


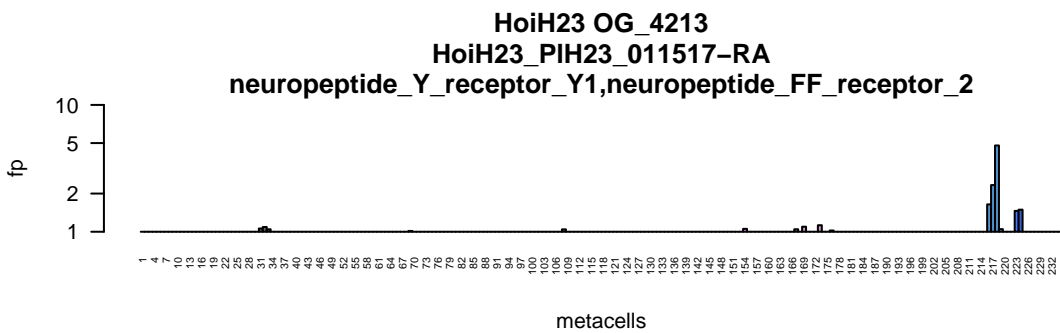
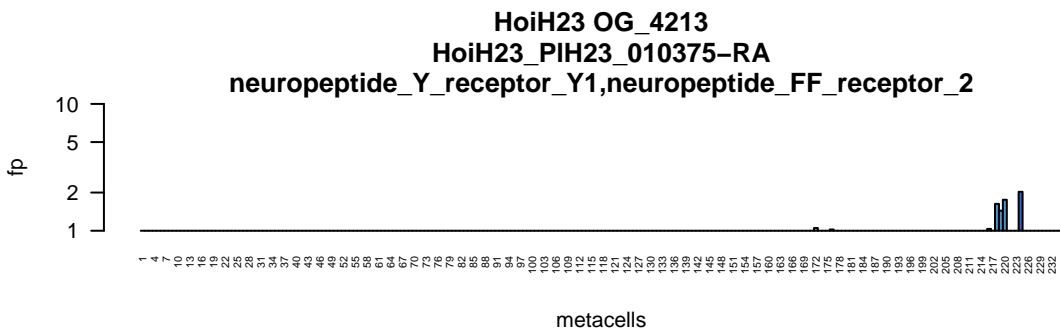
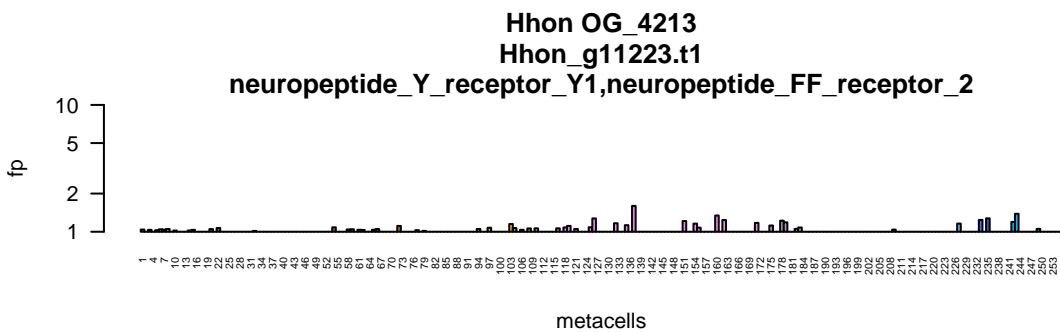
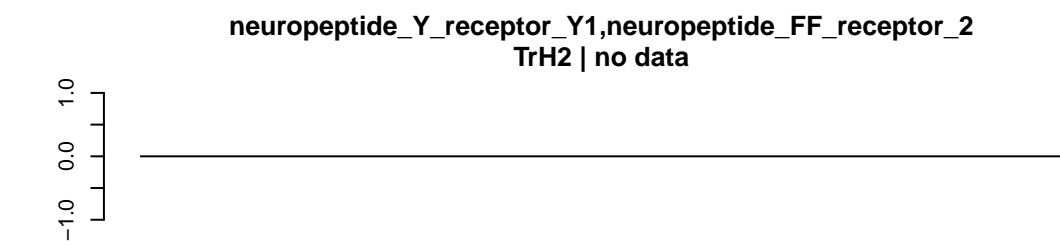
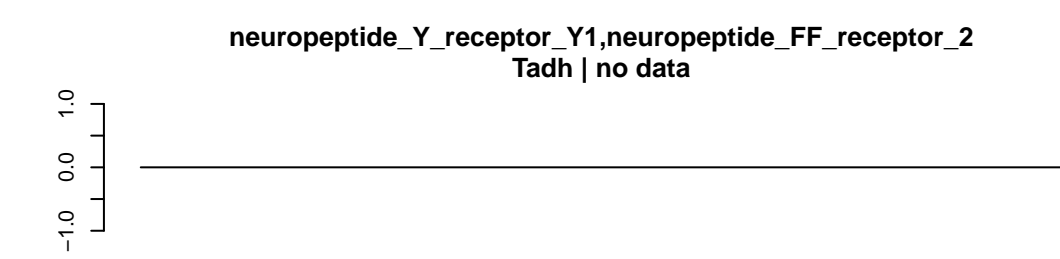












Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists 100 metacells. Most metacells have 0 false positives, but a few have 1, and a small cluster of metacells (173-187) has 2, 3, or 4 false 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
88	0
89	0
90	0
91	0
92	0
93	0
94	0
95	0
96	0
97	0
98	0
99	0
100	0
101	0
102	0
103	0
104	0
105	0
106	0
107	0
108	0
109	0
110	0
111	0
112	0
113	0
114	0
115	0
116	0
117	0
118	0
119	0
120	0
121	0
122	0
123	0
124	0
125	0
126	0
127	0
128	0
129	0
130	0
131	0
132	0
133	0
134	0
135	0
136	0
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 187 metacells. Most metacells have a false positive count of 1, with a few having counts of 2 or 4.

metacell	fp
1	1
2	1
3	1
4	1
5	1
6	1
7	1
8	1
9	1
10	1
11	1
12	1
13	1
14	1
15	1
16	1
17	1
18	1
19	1
20	1
21	1
22	1
23	1
24	1
25	1
26	1
27	1
28	1
29	1
30	1
31	1
32	1
33	1
34	1
35	1
36	1
37	1
38	1
39	1
40	1
41	1
42	1
43	1
44	1
45	1
46	1
47	1
48	1
49	1
50	1
51	1
52	1
53	1
54	1
55	1
56	1
57	1
58	1
59	1
60	1
61	1
62	1
63	1
64	1
65	1
66	1
67	1
68	1
69	1
70	1
71	1
72	1
73	1
74	1
75	1
76	1
77	1
78	1
79	1
80	1
81	1
82	1
83	1
84	1
85	1
86	1
87	1
88	1
89	1
90	1
91	1
92	1
93	1
94	1
95	1
96	1
97	1
98	1
99	1
100	1
101	1
102	1
103	1
104	1
105	1
106	1
107	1
108	1
109	1
110	1
111	1
112	1
113	1
114	1
115	1
116	1
117	1
118	1
119	1
120	1
121	1
122	1
123	1
124	1
125	1
126	1
127	1
128	1
129	1
130	1
131	1
132	1
133	1
134	1
135	1
136	1
137	1
138	1
139	1
140	1
141	1
142	1
143	1
144	1
145	1
146	1
147	1
148	1
149	1
150	1
151	1
152	1
153	1
154	1
155	1
156	1
157	1
158	1
159	1
160	1
161	1
162	1
163	1
164	1
165	1
166	1
167	1
168	1
169	1
170	1
171	1
172	1
173	1
174	1
175	1
176	1
177	1
178	1
179	1
180	1
181	1
182	1
183	1
184	1
185	1
186	1
187	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 40 metacells. Most metacells have a false positive count of 1, with a few having counts of 2 or 3.

metacells	fp
1	1
7	1
10	1
13	1
16	1
19	1
22	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	2
199	1
202	1
205	2
208	3
211	2
214	1

Bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and lists 48 categories. The y-axis is labeled 'fp' and ranges from 1 to 10. Most categories have a frequency of 1, with a few having higher frequencies: 196 (2), 202 (1), 208 (1), 214 (1).

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

metacell	fp
1	1
4	1
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	0
34	0
37	0
40	0
43	0
46	0
49	0
52	0
55	0
58	0
61	0
64	0
67	0
70	0
73	0
76	1
78	1
81	0
84	0
87	0
90	0
93	1
96	0
99	0
102	0
105	0
108	0
111	0
114	0
117	0
120	0
123	0
126	0
129	0
132	0
135	0
138	0
141	0
144	0
147	0
150	0
153	0
156	1
159	1
162	1
165	1
168	1
171	0
174	0
177	0
180	0
183	0
186	0
189	0
192	0
195	0
198	1
201	1
204	1
207	1
210	1
213	1
216	1

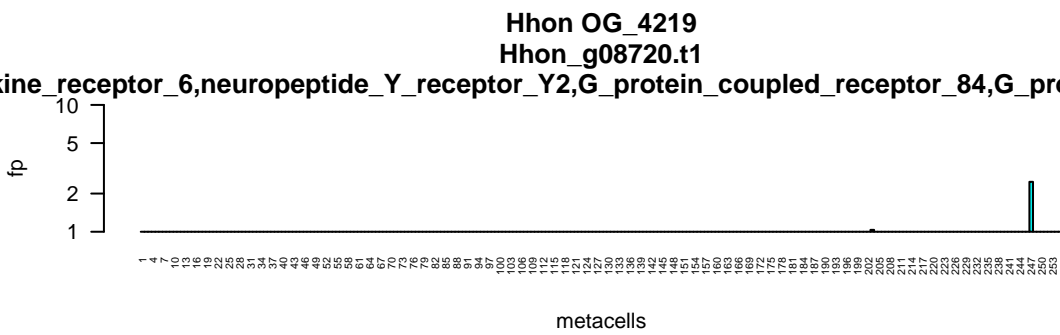
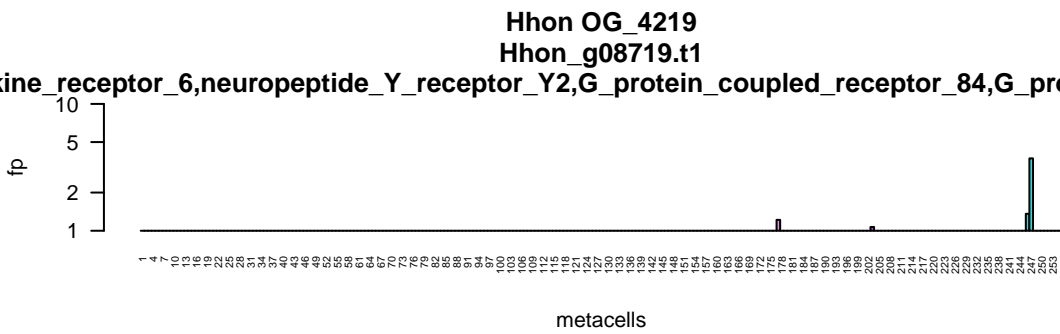
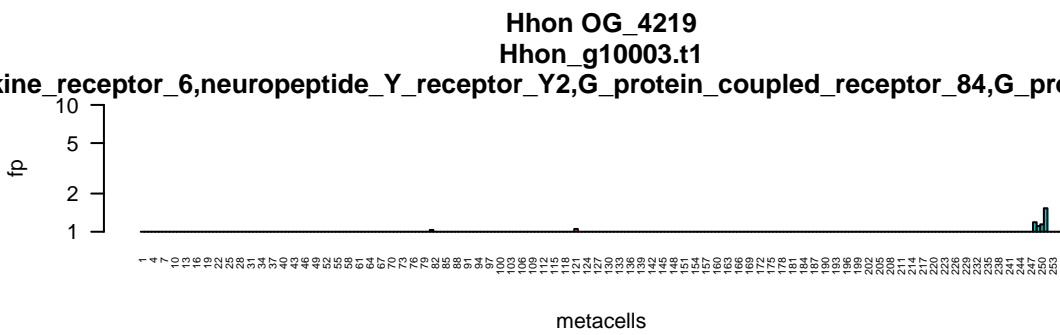
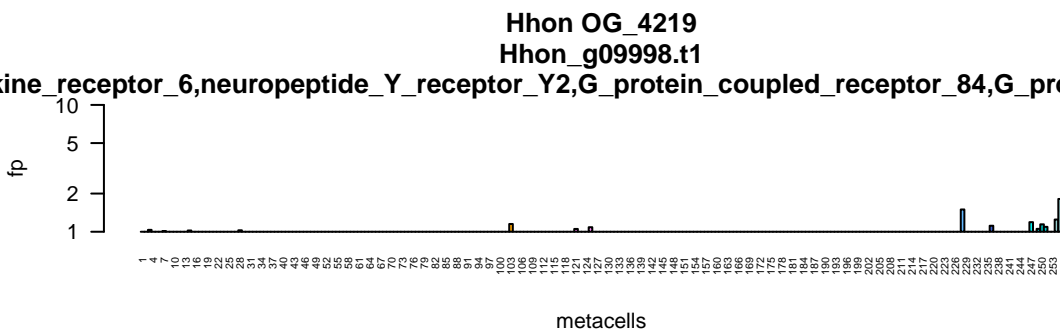
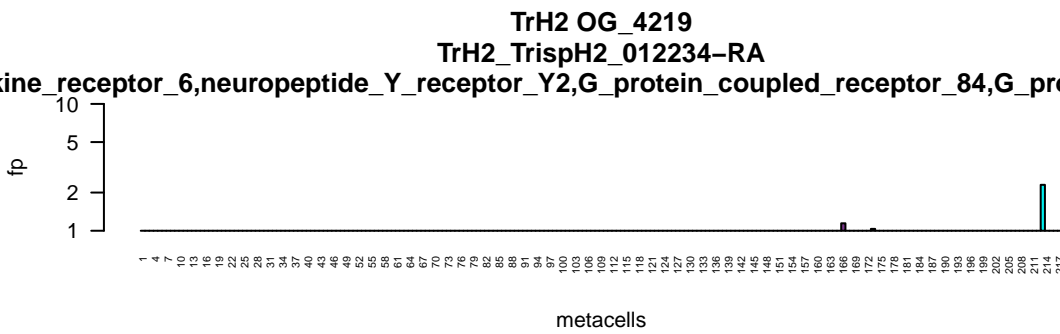
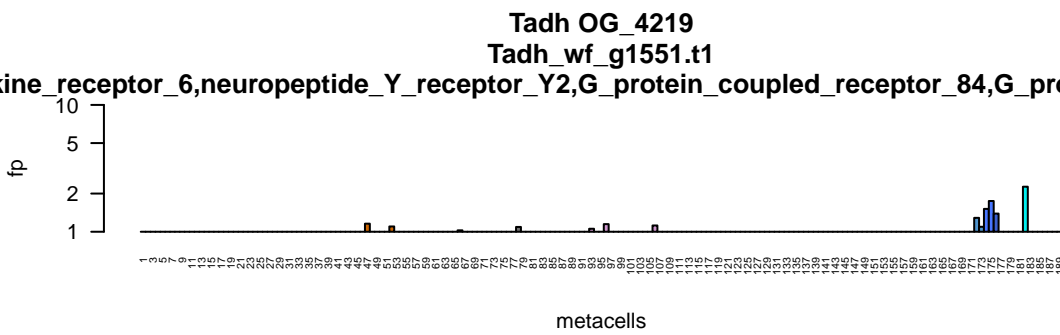
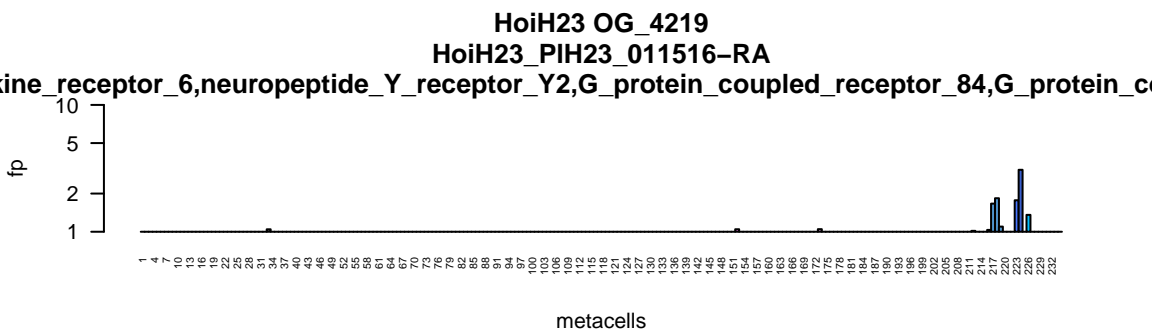
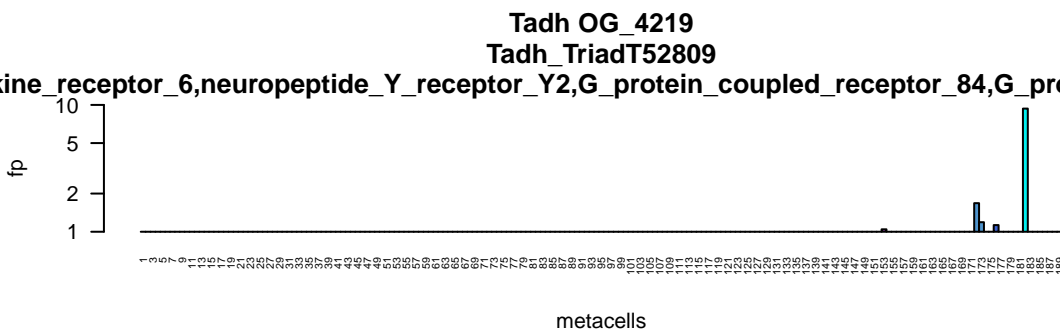
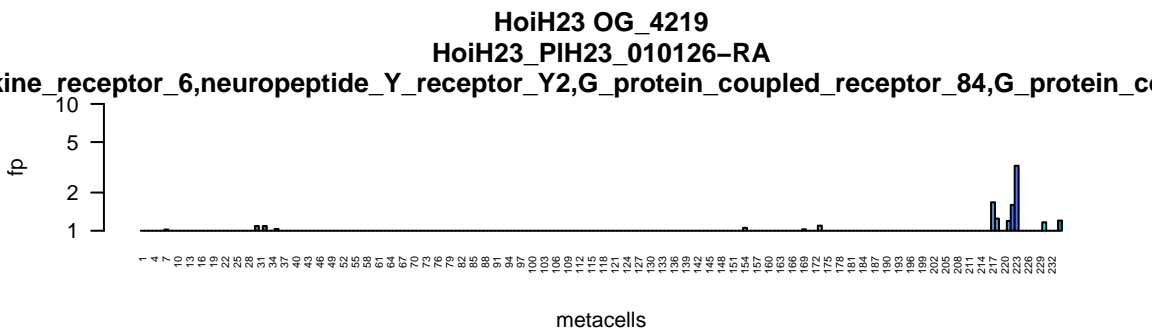
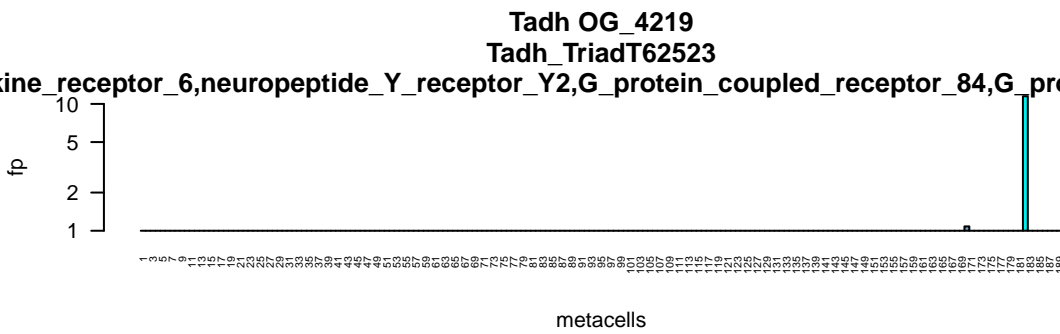
metacell	fp
1	1
2	1
3	1
4	1
5	1
6	1
7	1
8	1
9	1
10	1
11	1
12	1
13	1
14	1
15	1
16	1
17	1
18	1
19	1
20	1
21	1
22	1
23	1
24	1
25	1
26	1
27	1
28	1
29	1
30	1
31	1
32	1
33	1
34	1
35	1
36	1
37	1
38	1
39	1
40	1
41	1
42	1
43	1
44	1
45	1
46	1
47	1
48	1
49	1
50	1
51	1
52	1
53	1
54	1
55	1
56	1
57	1
58	1
59	1
60	1
61	1
62	1
63	1
64	1
65	1
66	1
67	1
68	1
69	1
70	1
71	1
72	1
73	1
74	1
75	1
76	1
77	1
78	1
79	1
80	1
81	1
82	1
83	1
84	1
85	1
86	1
87	1
88	1
89	1
90	1
91	1
92	1
93	1
94	1
95	1
96	1
97	1
98	1
99	1
100	1
101	1
102	1
103	1
104	1
105	1
106	1
107	1
108	1
109	1
110	1
111	1
112	1
113	1
114	1
115	1
116	1
117	1
118	1
119	1
120	1
121	1
122	1
123	1
124	1
125	1
126	1
127	1
128	1
129	1
130	1
131	1
132	1
133	1
134	1
135	1
136	1
137	1
138	1
139	1
140	1
141	1
142	1
143	1
144	1
145	1
146	1
147	1
148	1
149	1
150	1
151	1
152	1
153	1
154	1
155	1
156	1
157	1
158	1
159	1
160	1
161	1
162	1
163	1
164	1
165	1
166	1
167	1
168	1
169	1
170	1
171	1
172	1
173	1
174	1
175	1
176	1
177	1
178	1
179	1
180	1
181	1
182	1
183	1
184	1
185	1
186	1
187	1
188	1
189	1
190	1
191	1
192	1
193	1
194	1
195	1
196	1
197	1
198	1
199	1
200	1
201	1
202	1
203	1
204	1
205	1
206	1
207	1
208	1
209	1
210	1
211	1
212	1
213	1

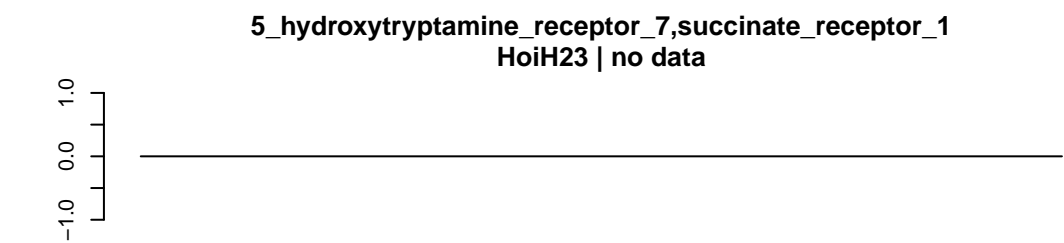
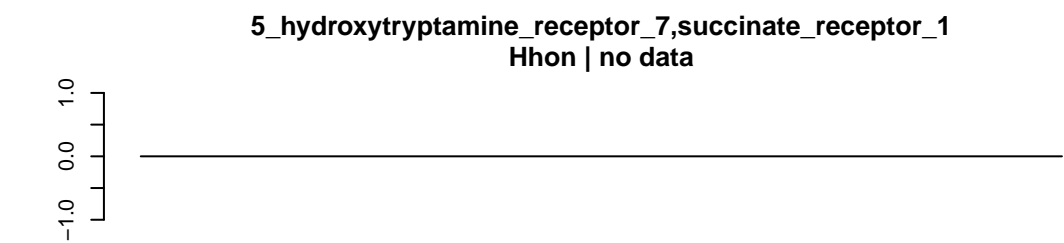
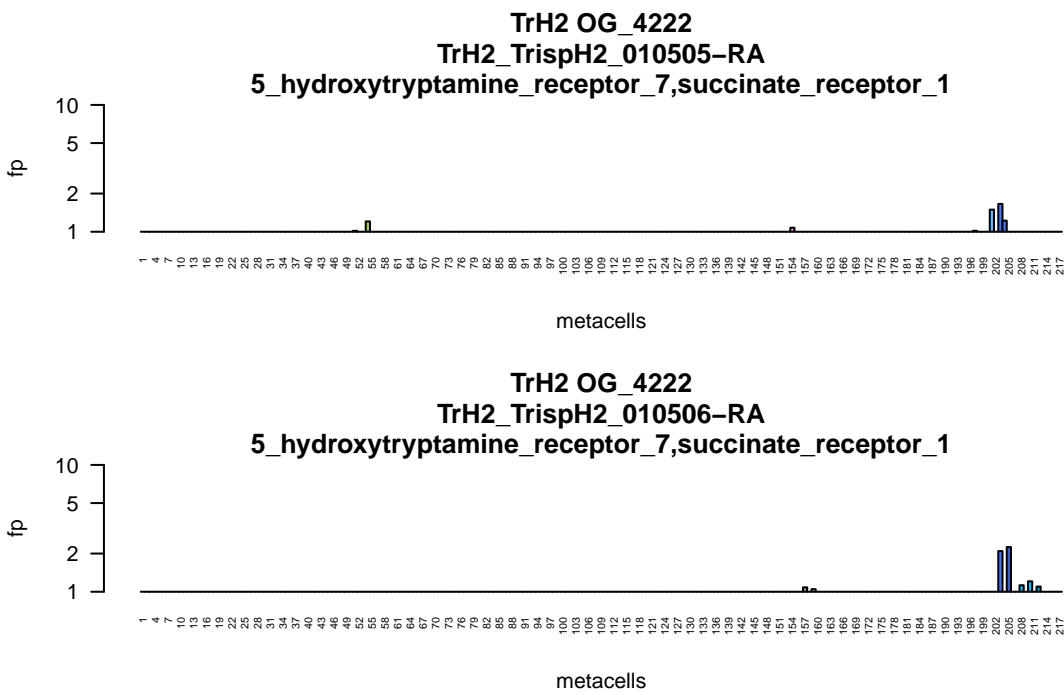
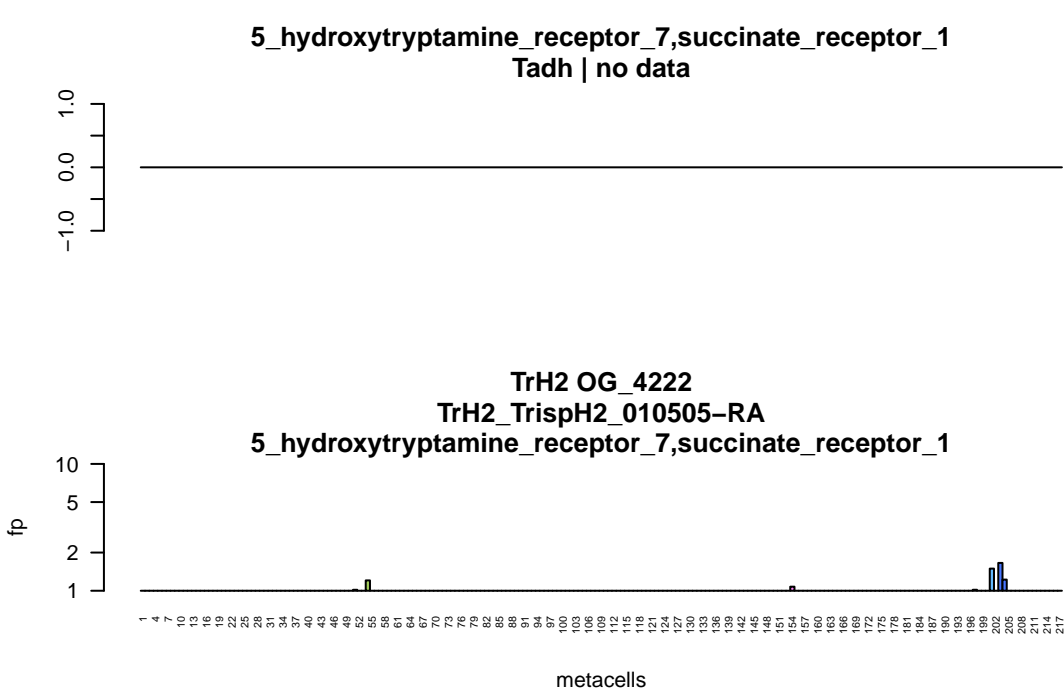
Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists indices from 1 to 232. Most metacells have a false positive count of 1. Metacells 169, 170, 171, 172, 173, 174, 175, 176, 177, 178, 179, 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.

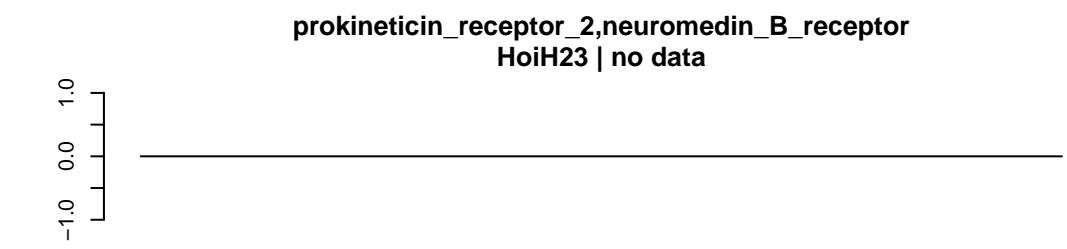
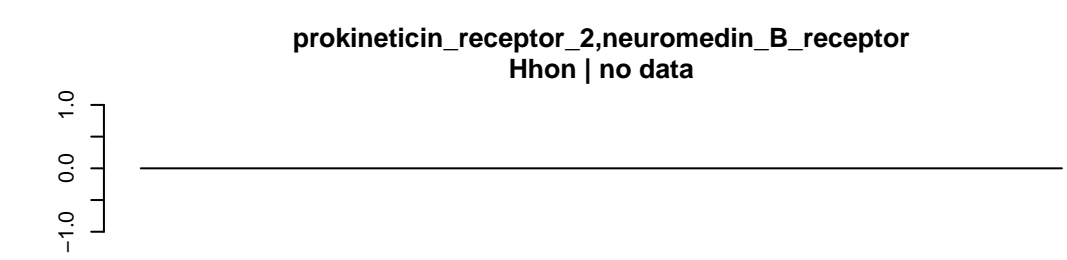
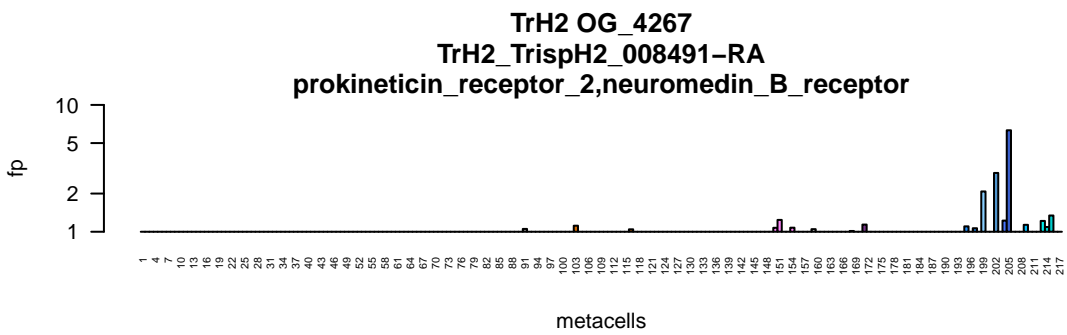
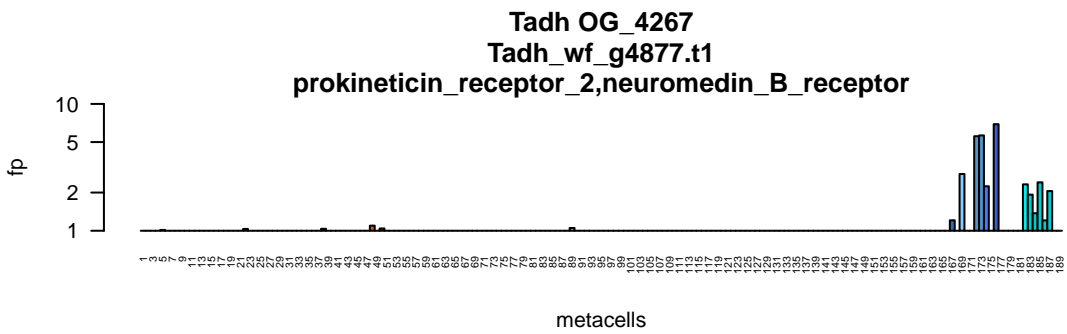
Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 232. Most metacells have 1 false positive, with a small cluster of metacells (211-232) showing higher values up to 2.

metacell	fp
1	0
4	1
10	0
13	0
19	0
22	0
31	0
34	1
40	0
43	0
46	0
52	0
55	0
58	0
60	0
64	0
67	0
73	0
76	1
82	0
85	0
88	0
94	1
97	0
103	0
106	1
109	0
115	0
118	0
124	0
127	1
130	0
136	1
139	0
145	0
151	1
157	0
160	0
166	0
169	0
172	1
178	0
181	0
187	0
190	0
193	0
199	0
202	0
208	0
211	1
214	0
217	3
220	2
221	1
222	6
223	10
229	9
232	1

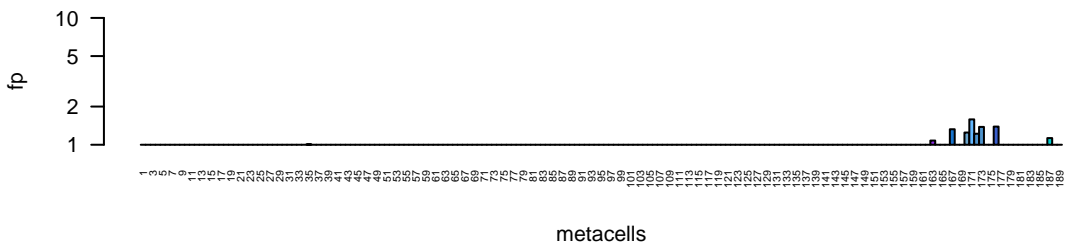
metacell	fp
1	0
10	0
13	0
19	0
22	0
25	0
31	1
34	1
40	0
43	0
46	0
52	1
55	0
58	0
64	0
67	0
73	0
76	0
82	0
85	0
89	0
94	0
97	0
103	0
106	0
109	0
115	0
118	0
124	0
127	0
130	0
136	0
139	0
145	0
148	0
151	1
157	0
160	0
166	0
169	1
172	0
178	0
187	0
190	0
193	0
199	0
202	0
208	0
211	0
214	2
220	2
223	3
229	1
232	2





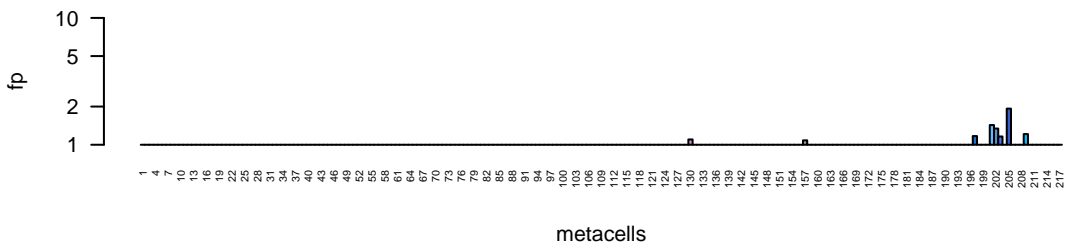


Tadh OG_4270
Tadh_wf_g4875.t1



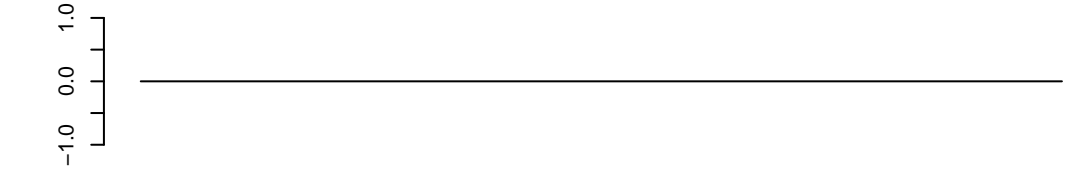
metacells

TrH2 OG_4270
TrH2_TrispH2_008489-RA

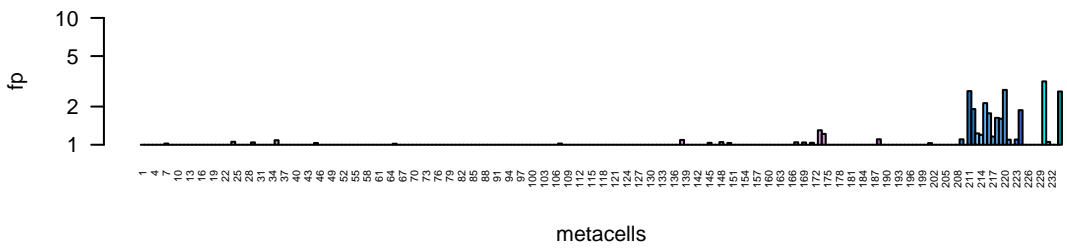


metacells

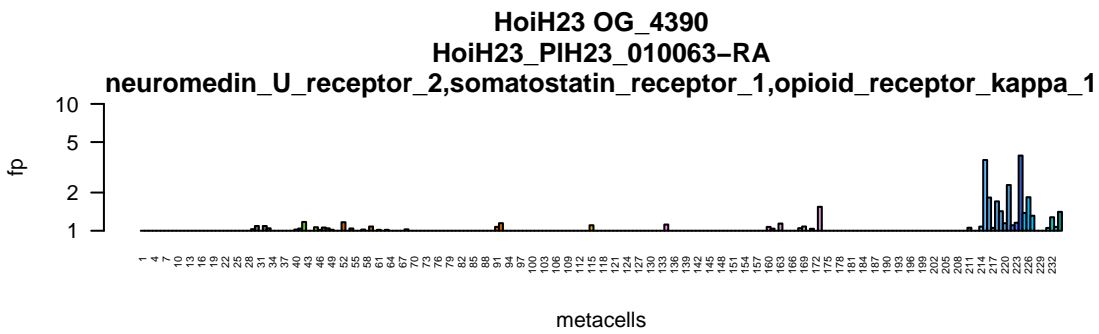
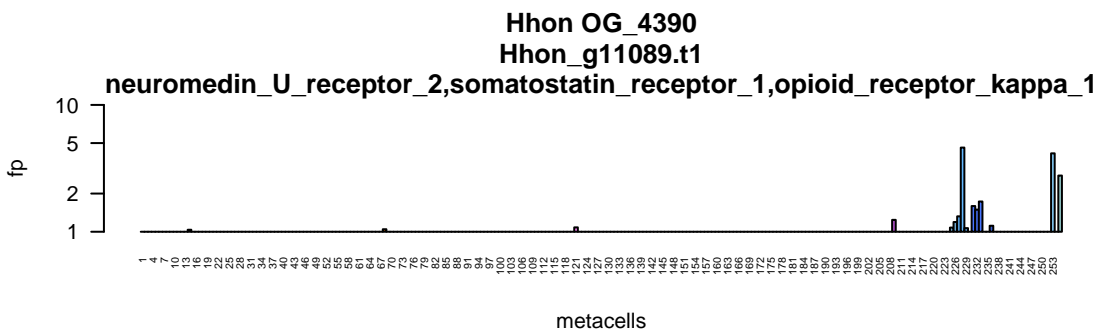
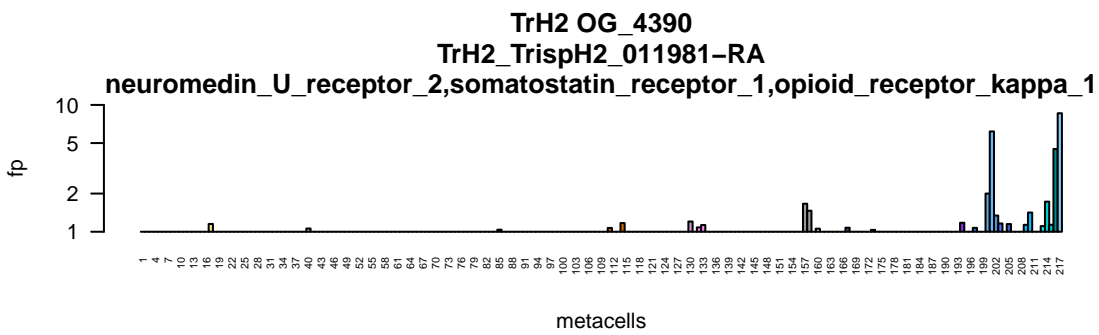
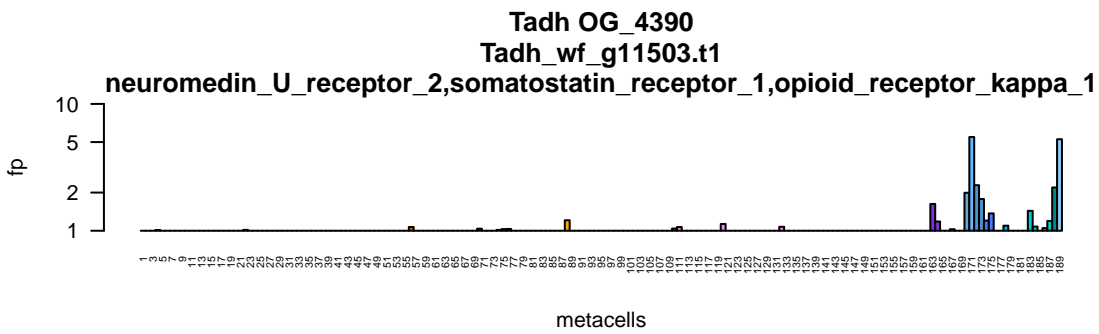
Hhon | no data

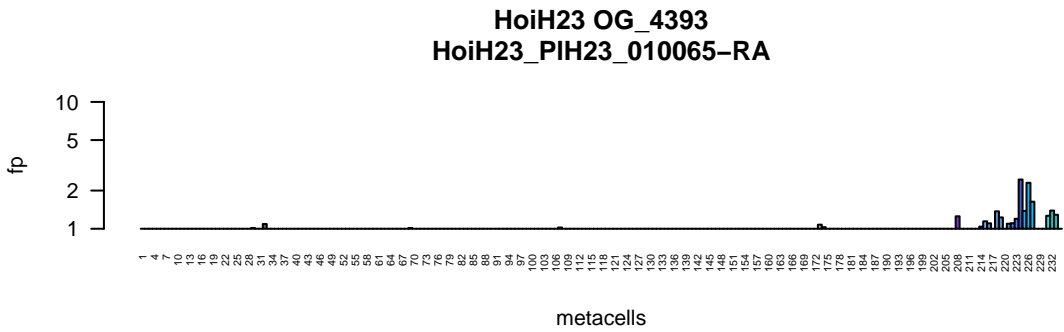
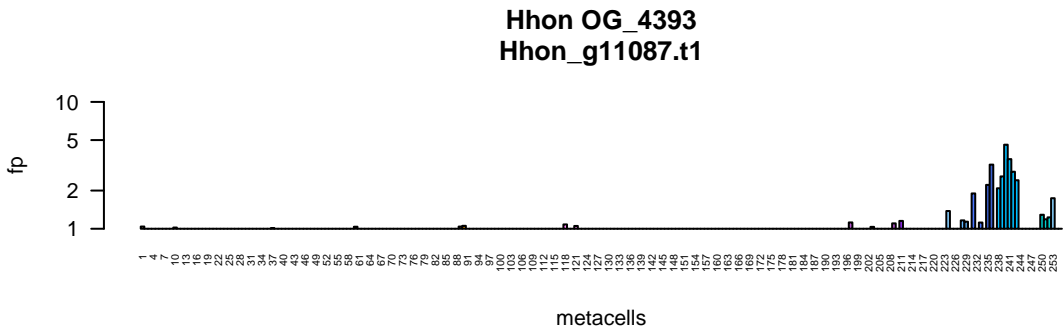
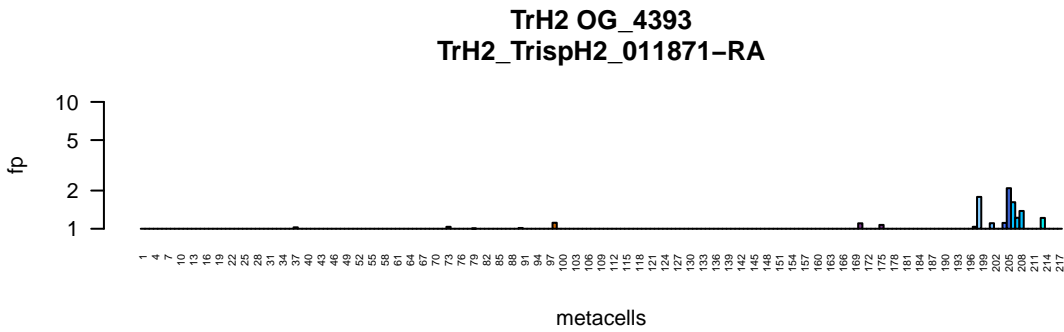
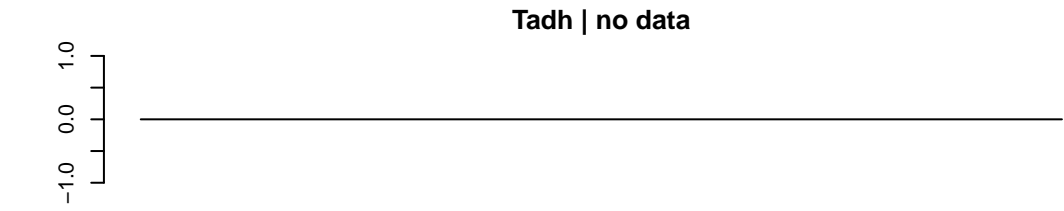


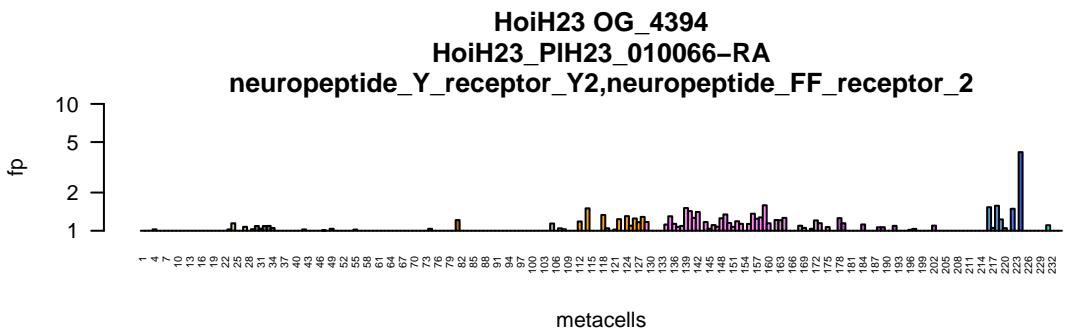
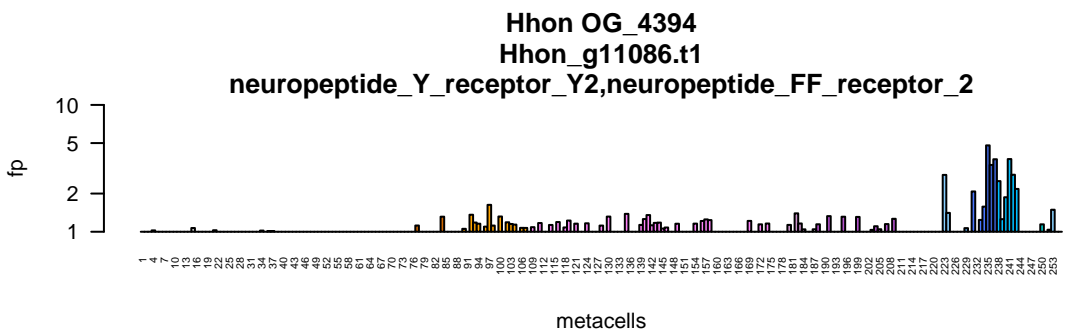
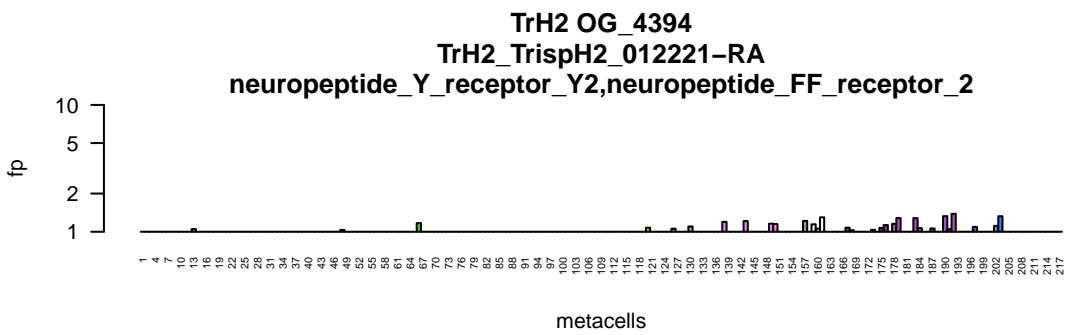
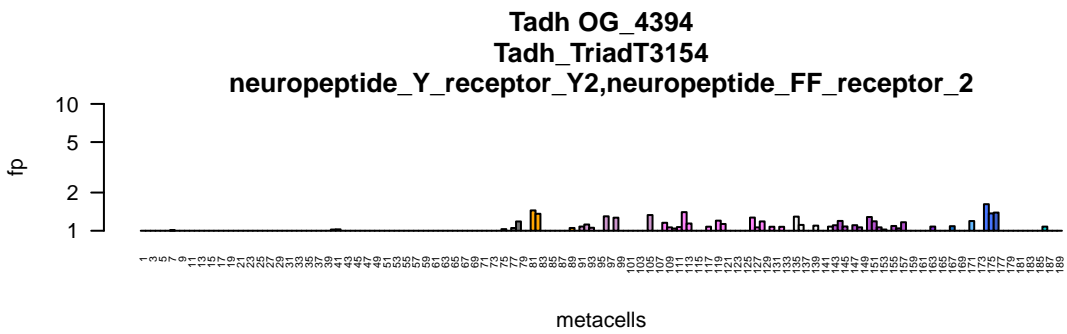
HoiH23 OG_4270
HoiH23_PIH23_009325-RA



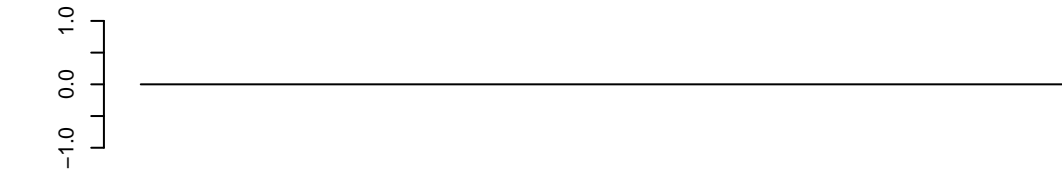
metacells



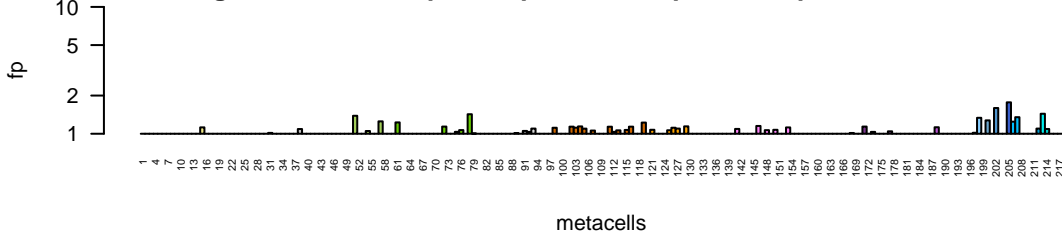




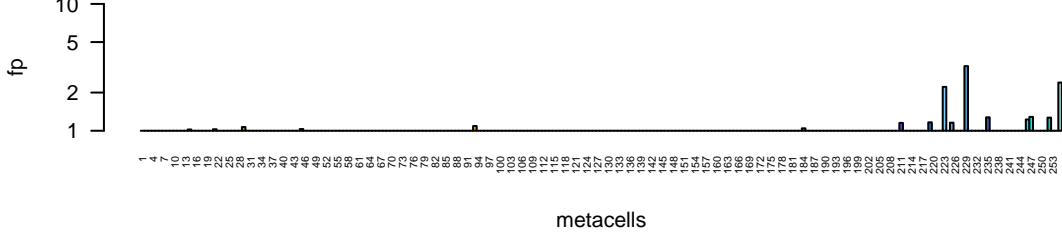
rolactin_releasing_hormone_receptor,G_protein_coupled_receptor_63,somatostatin_recep
Tadh | no data



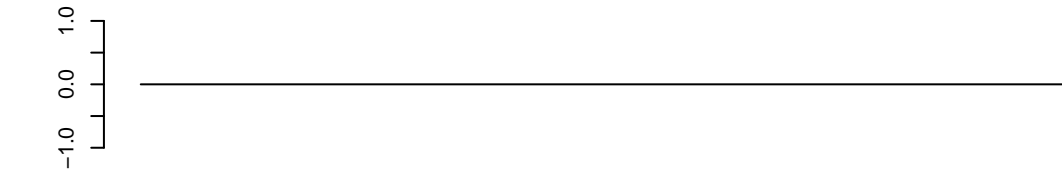
TrH2 OG_4413
TrH2_TrispH2_009693-RA
rolactin_releasing_hormone_receptor,G_protein_coupled_receptor_63,somatostatin_recep

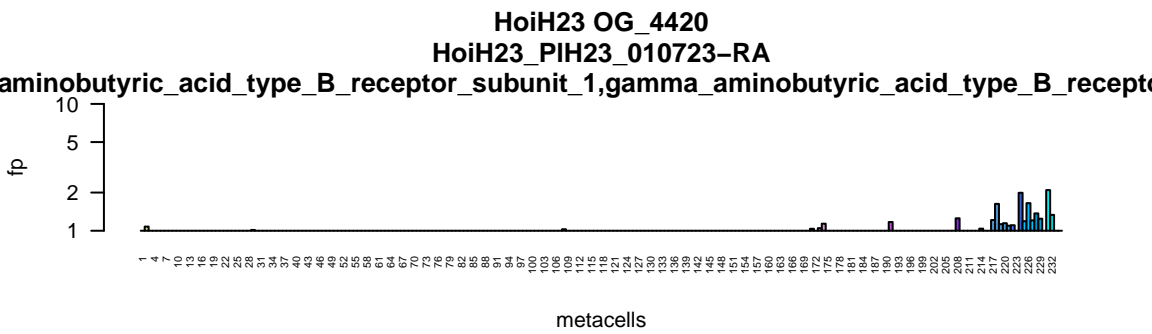
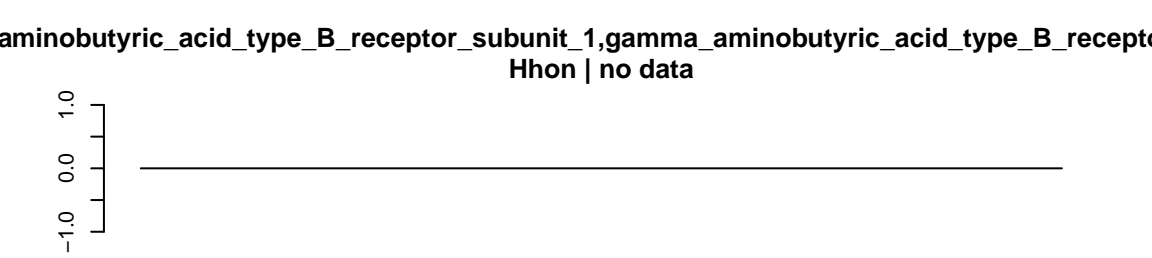
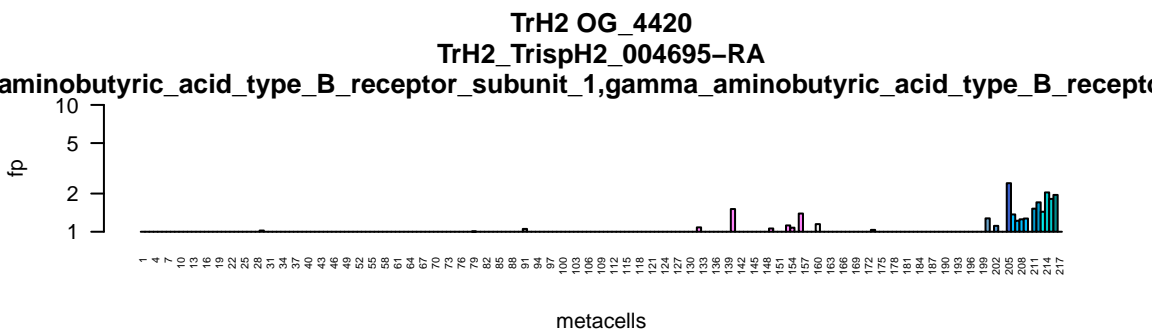
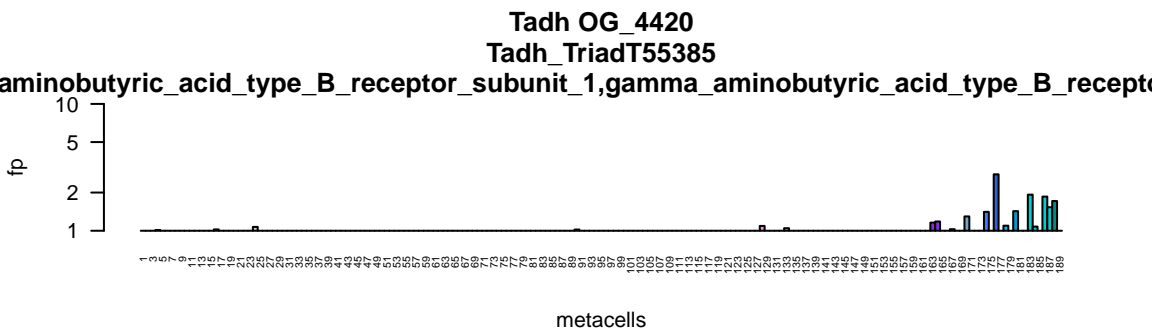


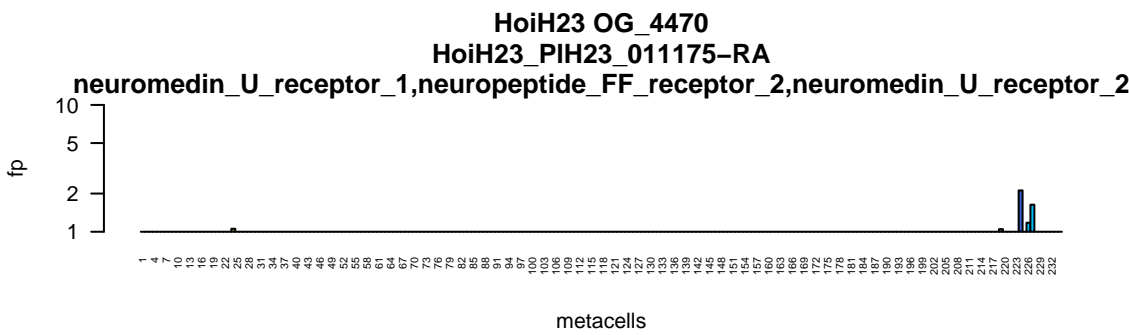
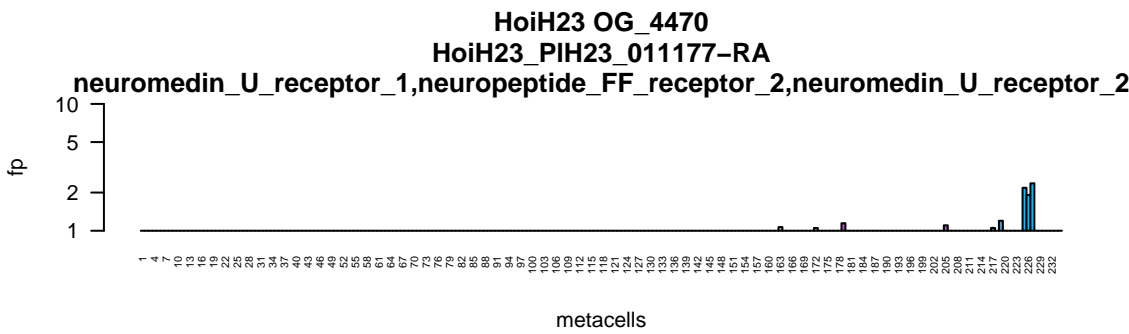
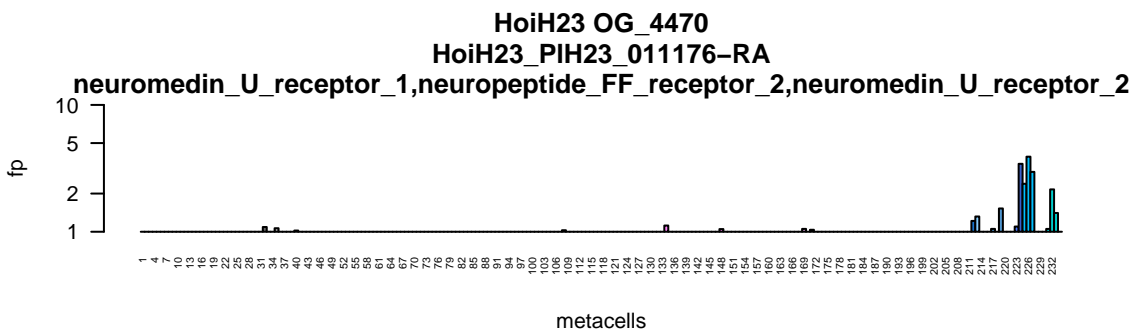
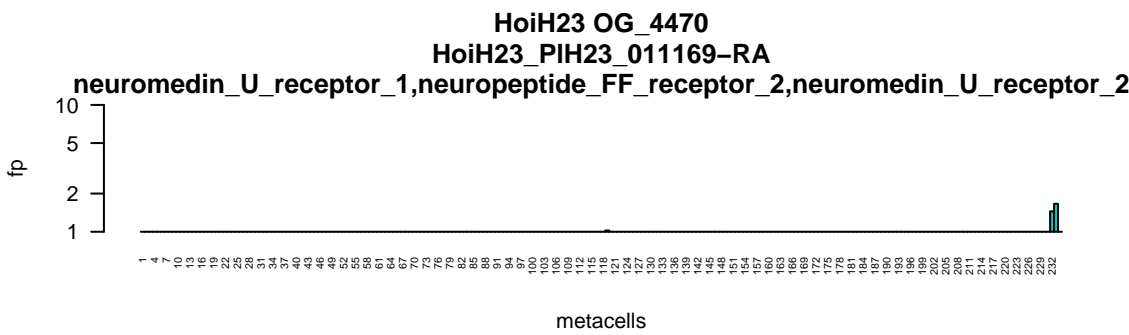
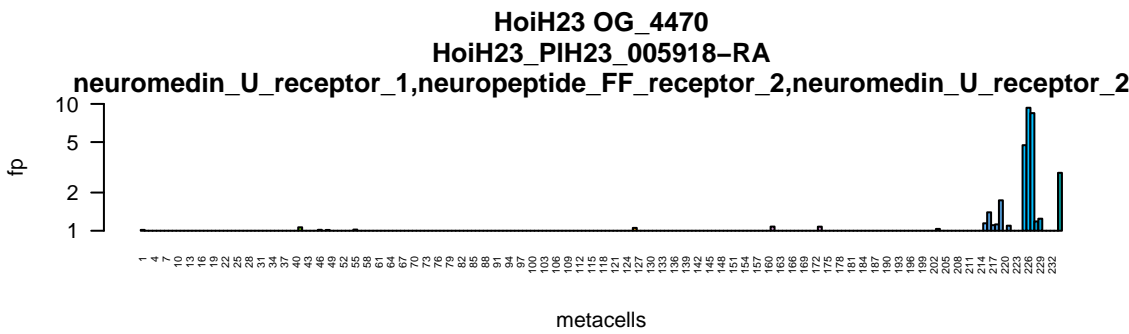
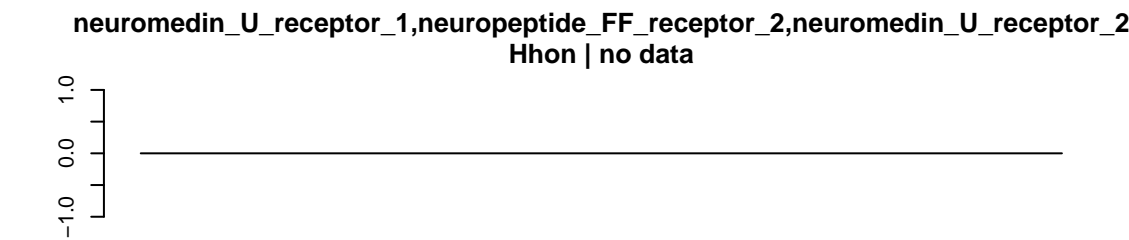
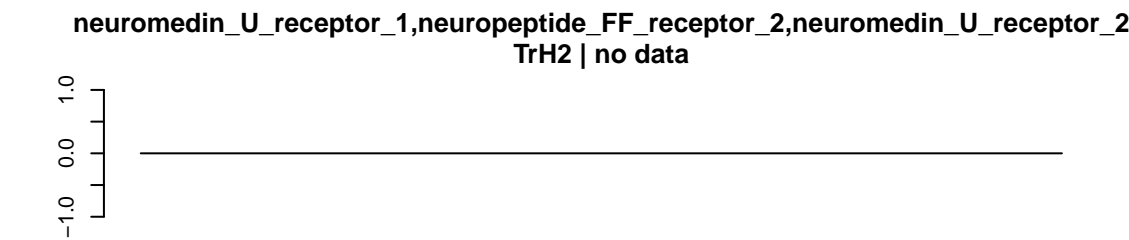
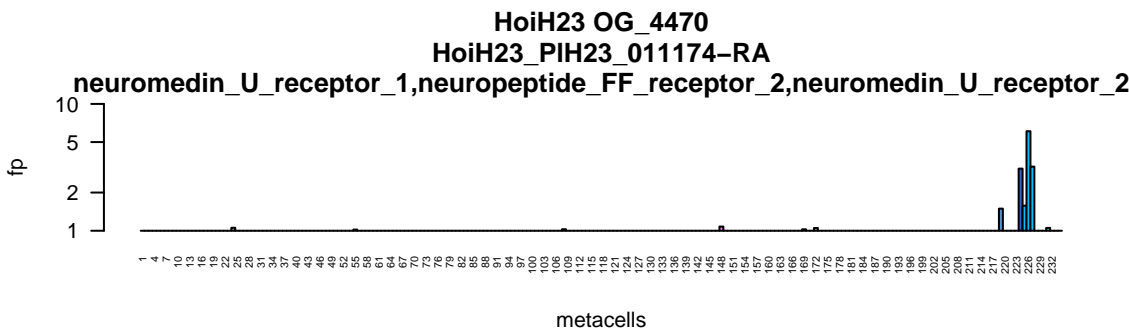
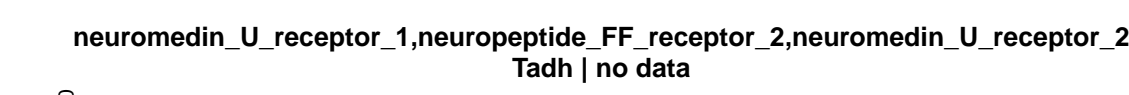
Hhon OG_4413
Hhon_g10372.t1
rolactin_releasing_hormone_receptor,G_protein_coupled_receptor_63,somatostatin_recep

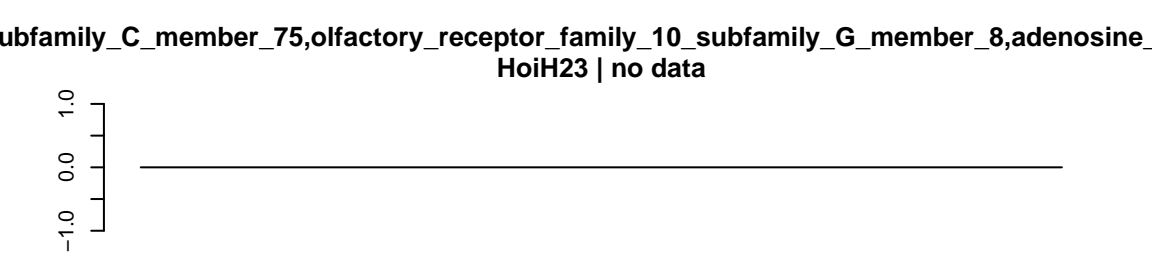
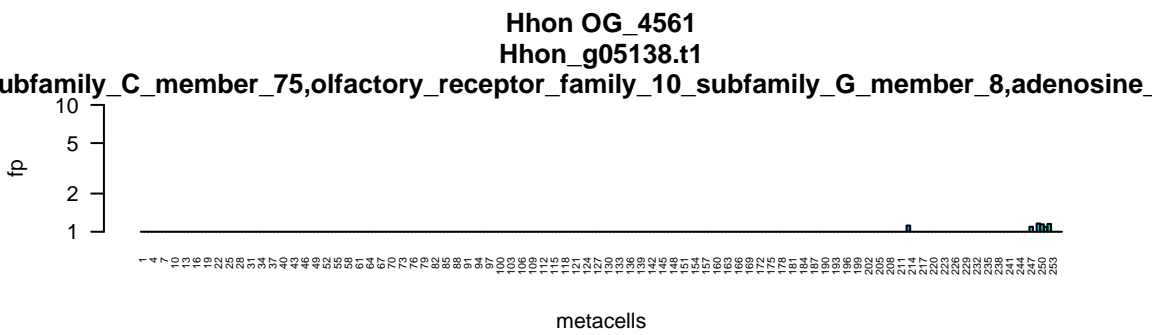
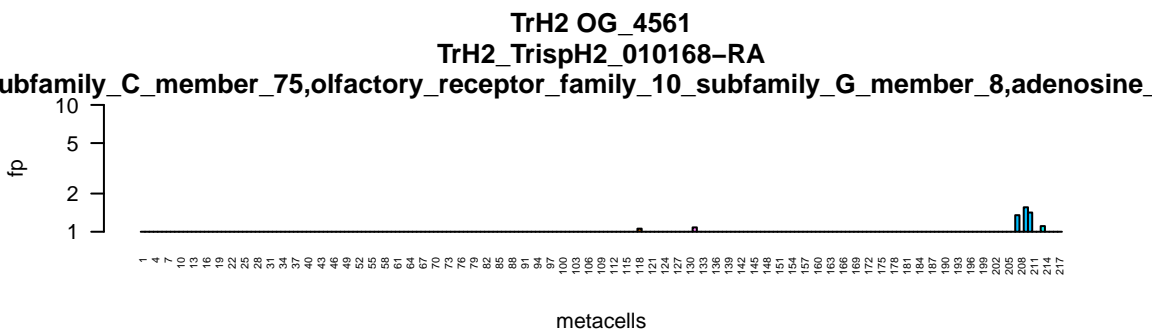
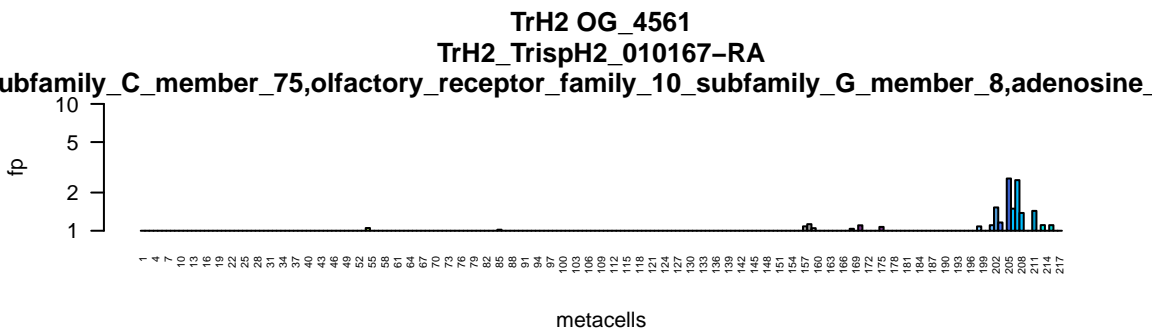
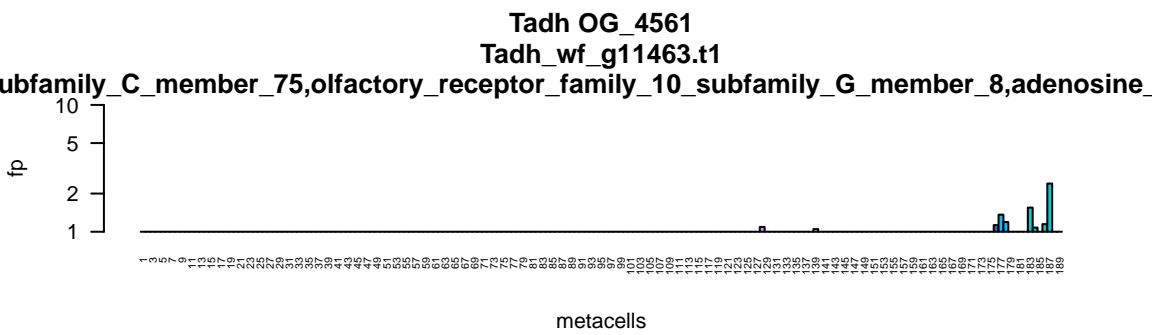
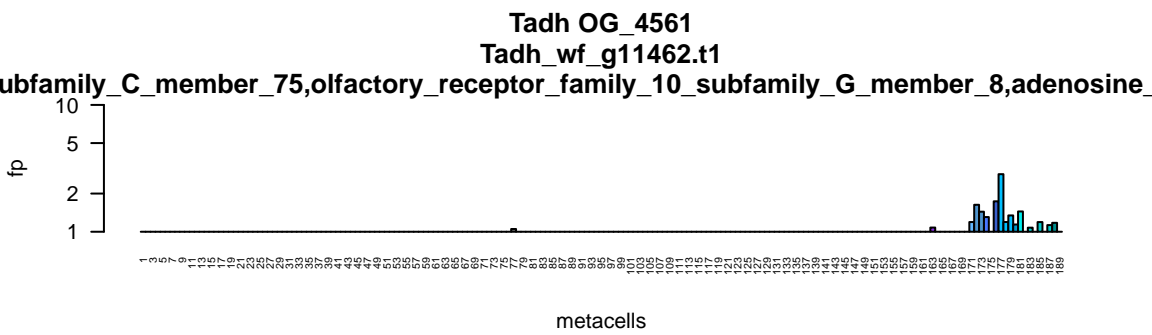
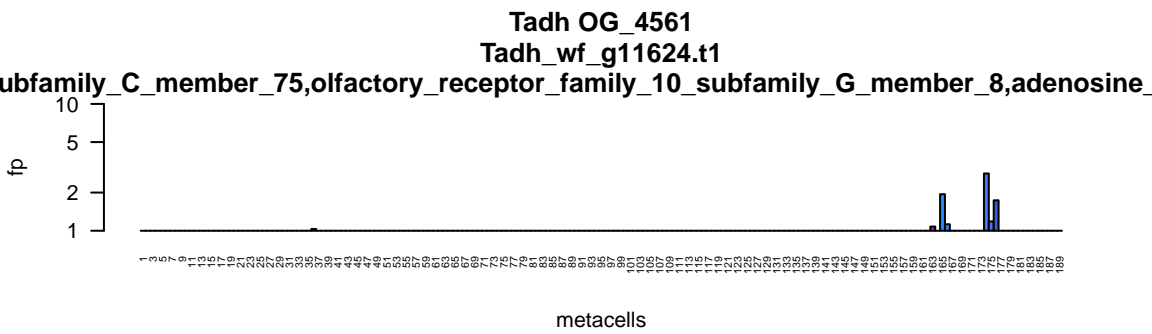


rolactin_releasing_hormone_receptor,G_protein_coupled_receptor_63,somatostatin_recep
HoiH23 | no data









TrH2 OG_4762
TrH2_TrispH2_010632-RA
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

fp

metacells

TrH2 OG_4762
TrH2_TrispH2_010635-RA
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

metacells	fp
1	1
7	1
13	1
16	1
19	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
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76	1
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82	1
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127	1
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133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
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160	1
163	1
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172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	2
202	1
205	1
208	1
211	1
214	2
217	1

TrH2 OG_4762
TrH2_TrispH2_010271-RA
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

fp

metacells

Hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

metacells

Hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

metacells	fp
1	1
4	1
10	1
13	1
19	1
22	1
28	1
31	1
37	1
43	1
46	1
52	1.5
55	1
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79	1
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88	1
91	1
97	1
100	1
106	1
108	1
109	1
115	1
118	1
127	1
133	1
136	1
145	1
151	1
154	1
160	1
163	1
169	1
175	1
178	1
181	1
184	1
187	1
193	1
196	1
202	1
205	1
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253	1

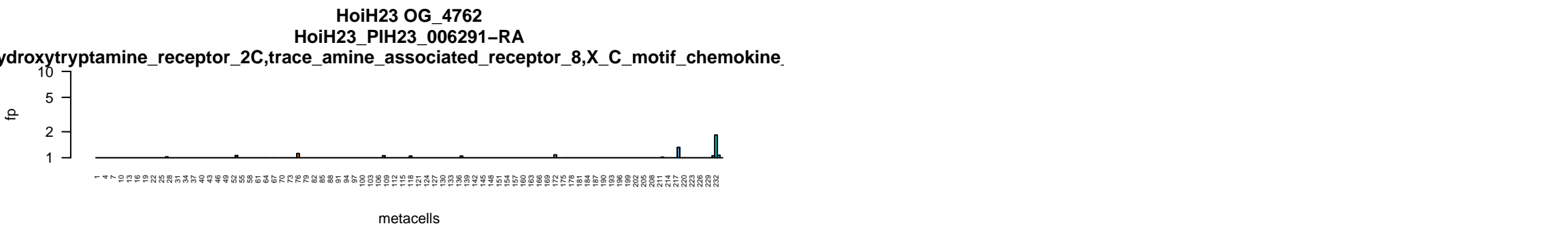
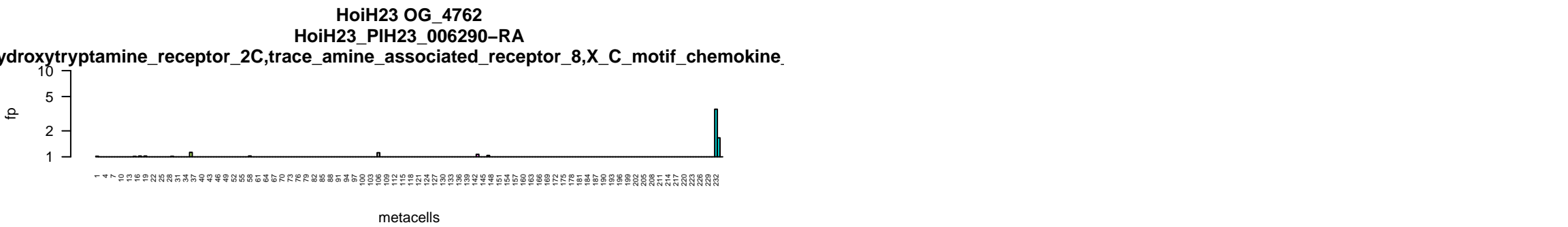
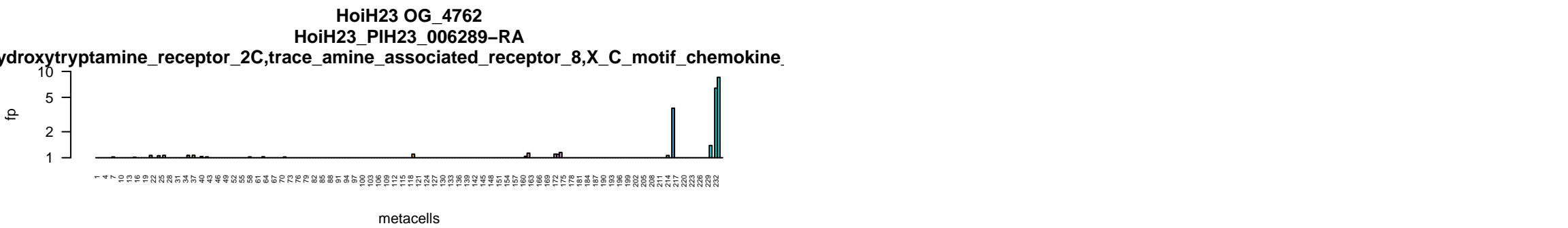
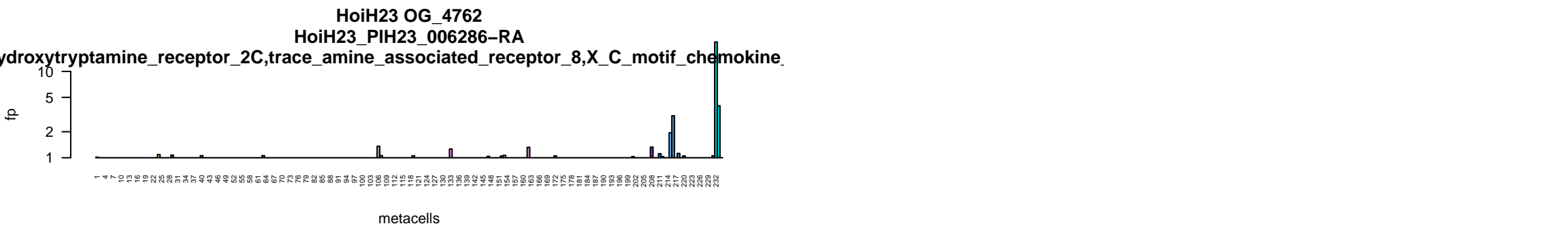
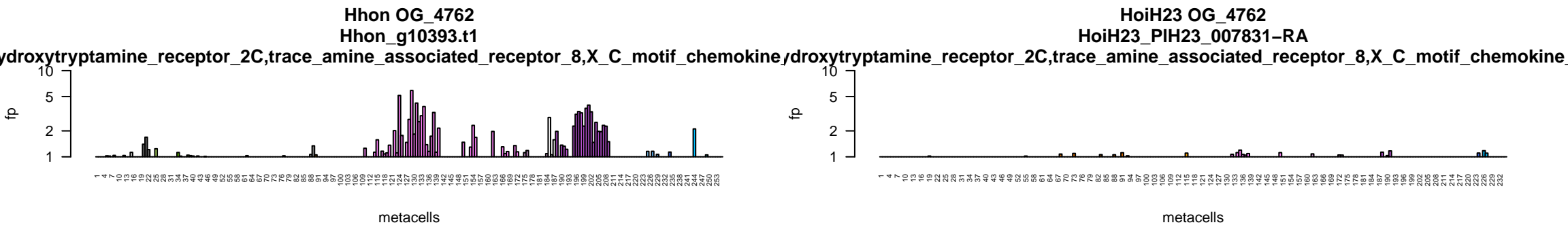
Hhox OG_4762
Hhox_g01703.t1

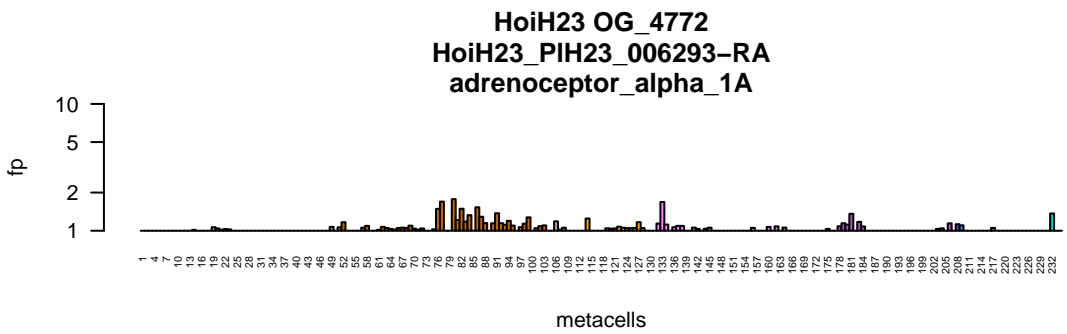
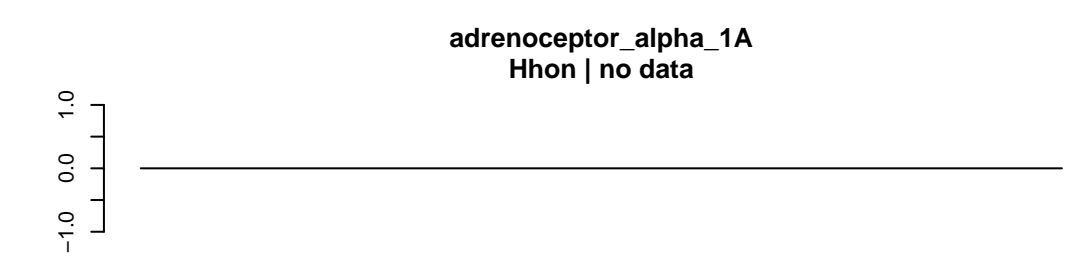
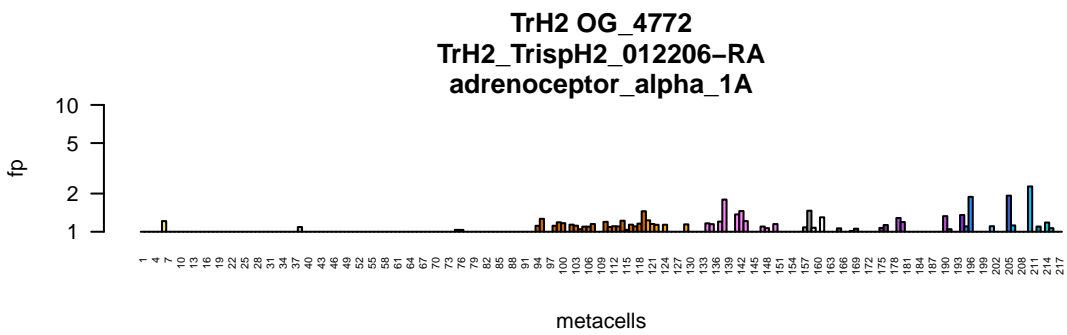
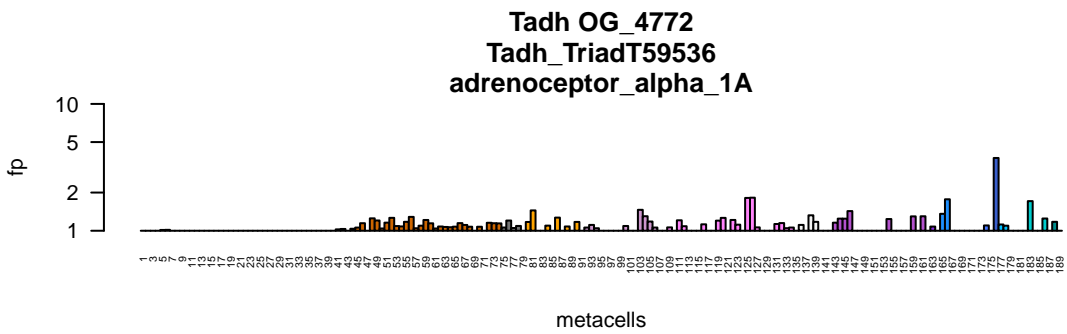
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

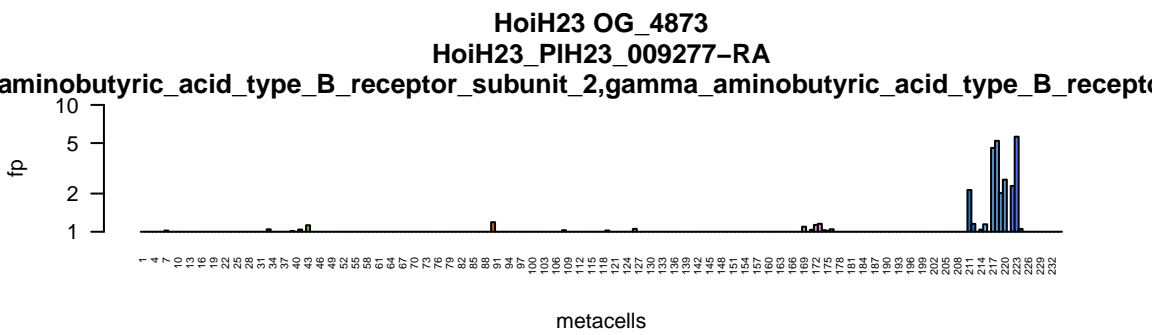
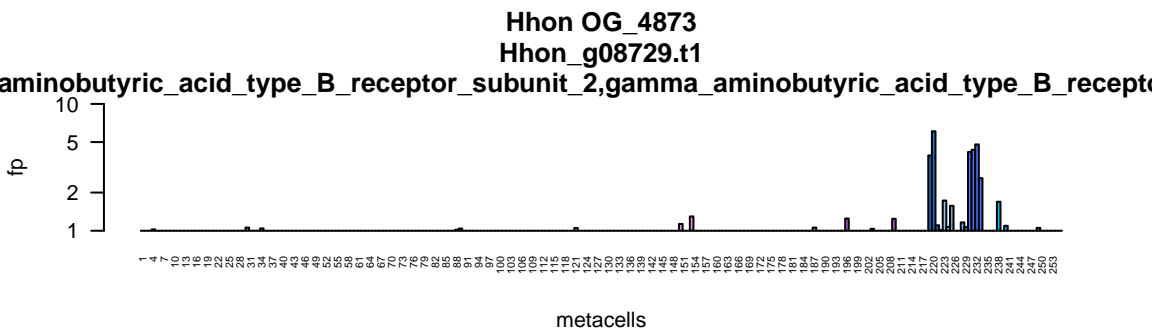
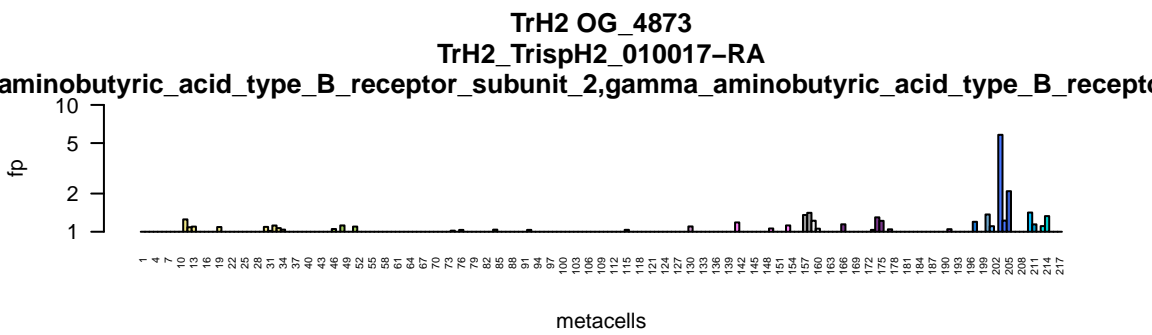
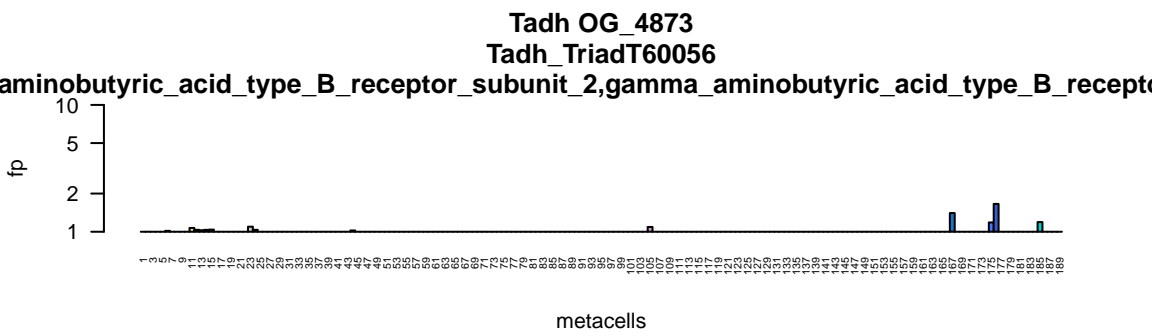
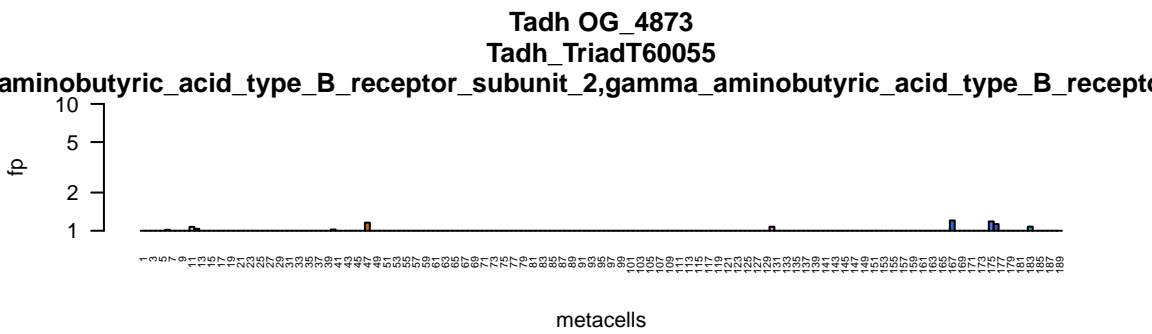
fp

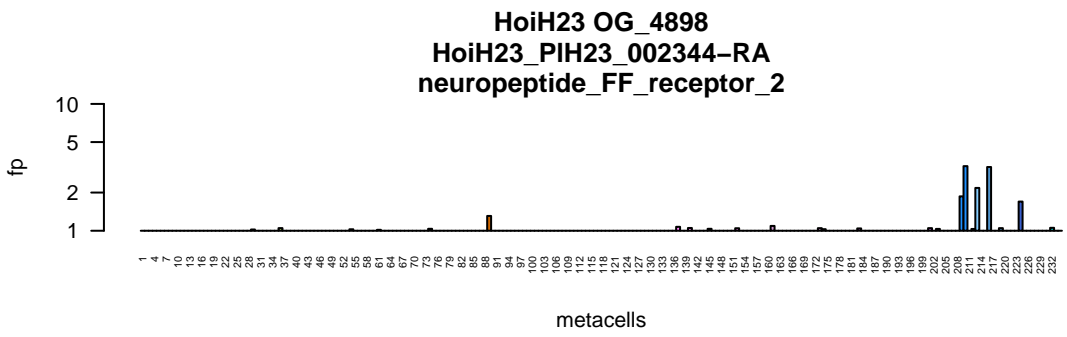
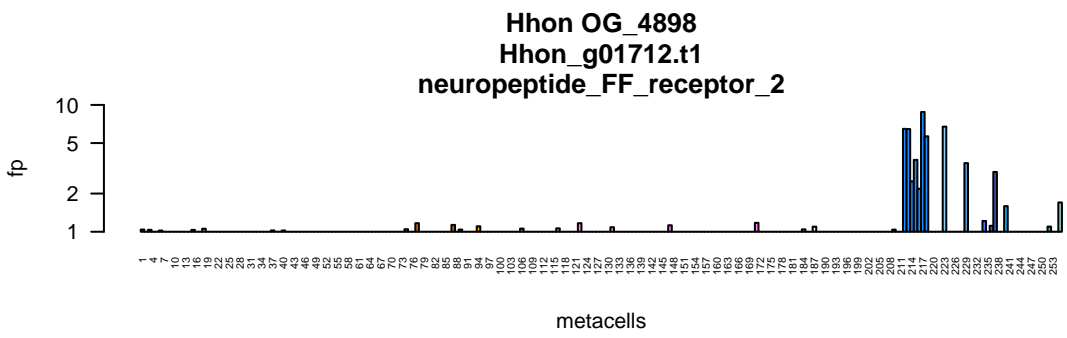
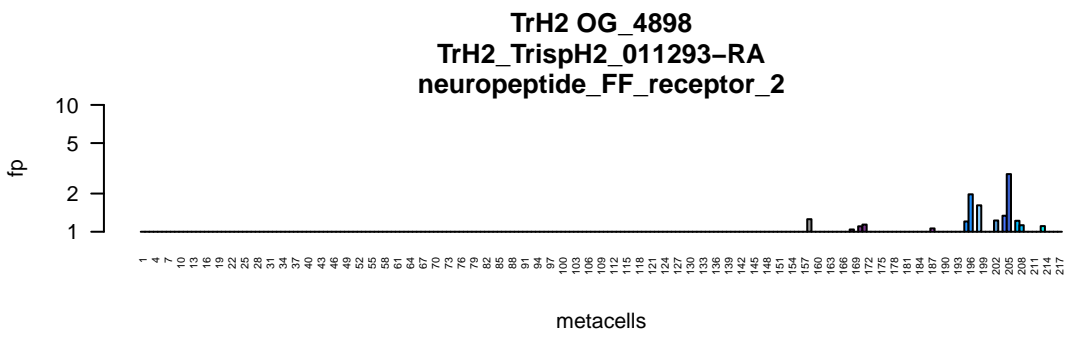
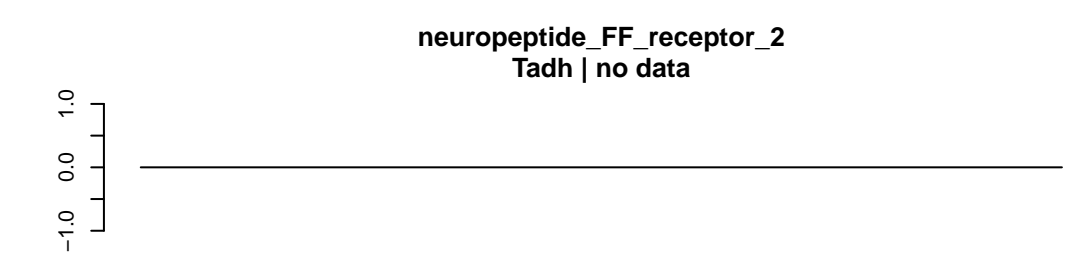
metacells

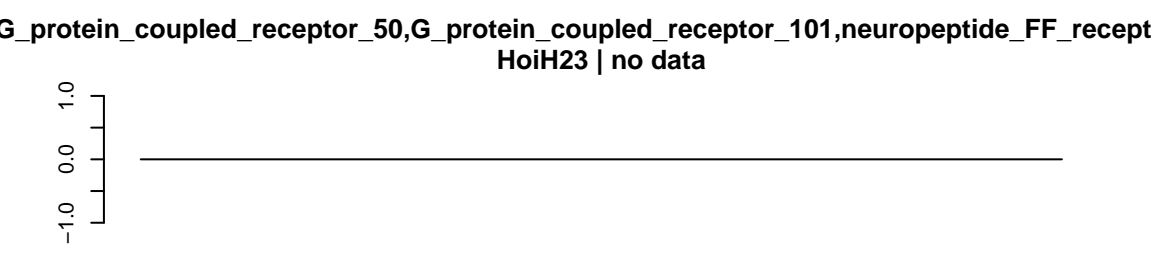
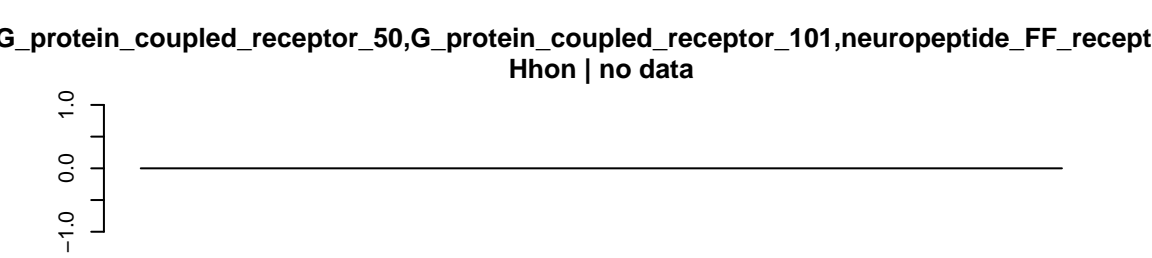
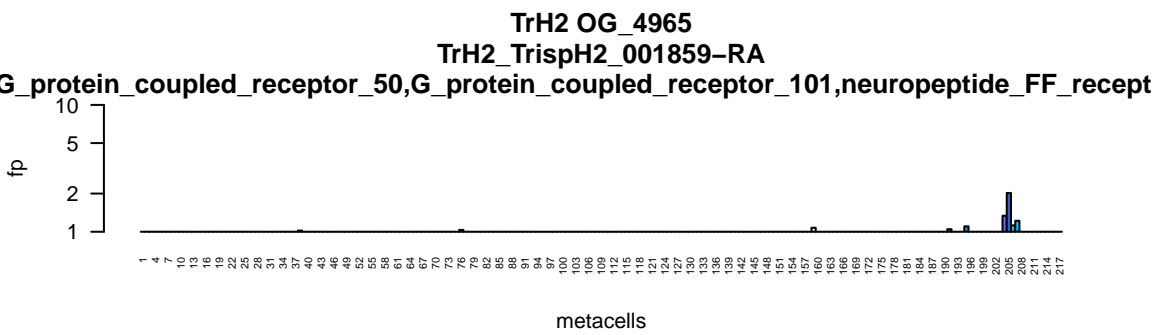
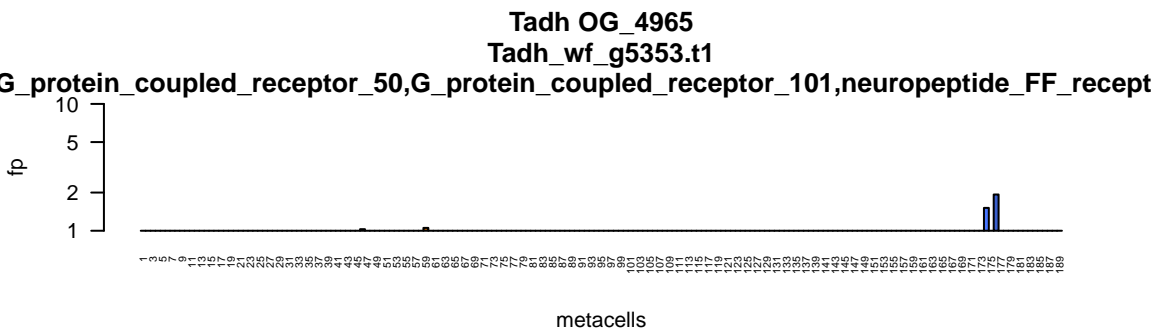
metacell	fp
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193	1
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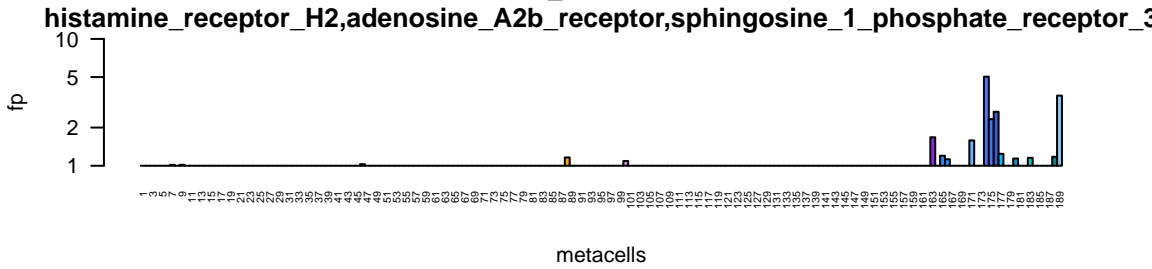




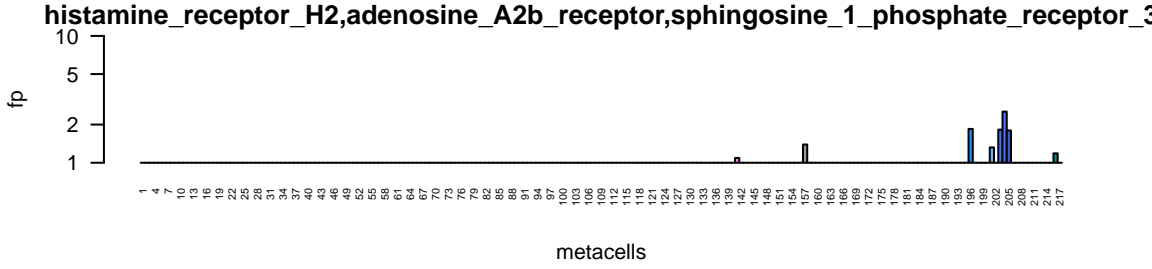




Tadh OG_4983
Tadh_TriadT58722



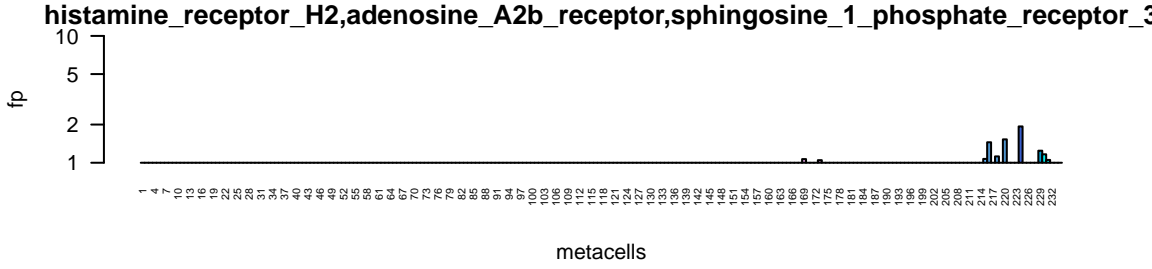
TrH2 OG_4983
TrH2_TrispH2_006523-RA



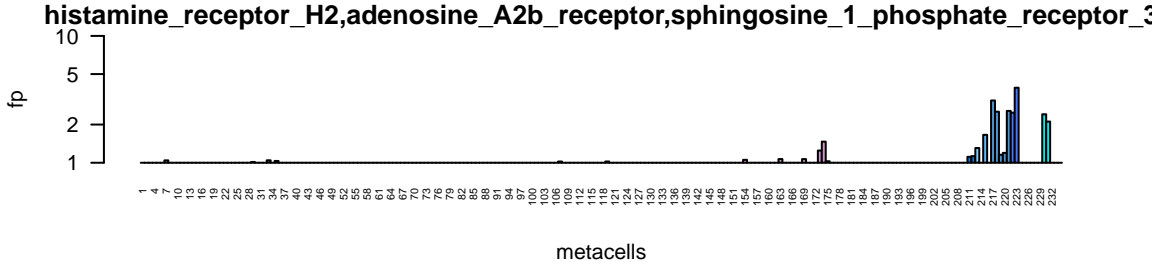
Hhon OG_4983
Hhon_g01339.t1

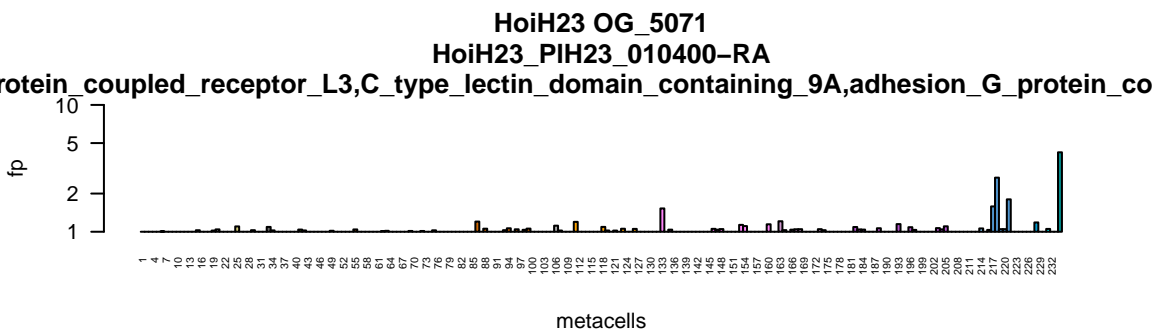
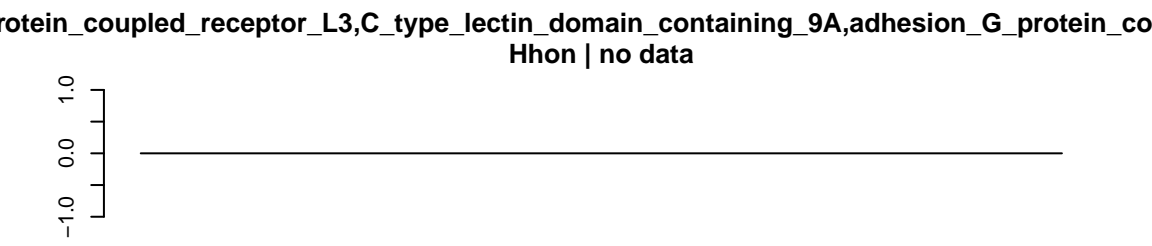
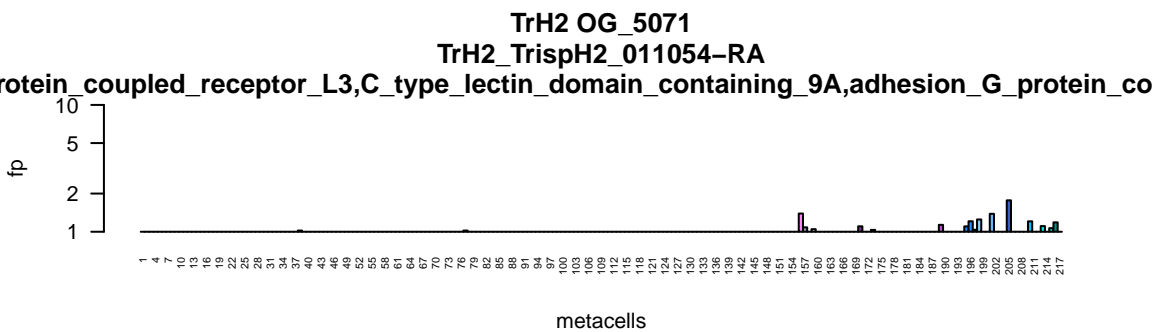
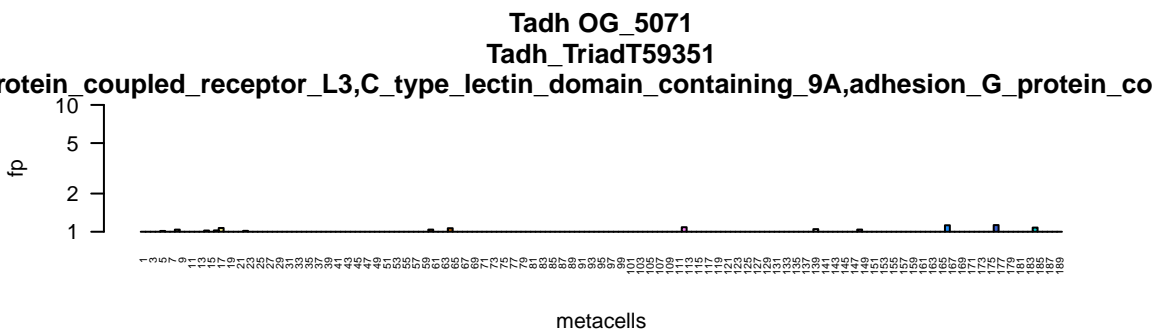
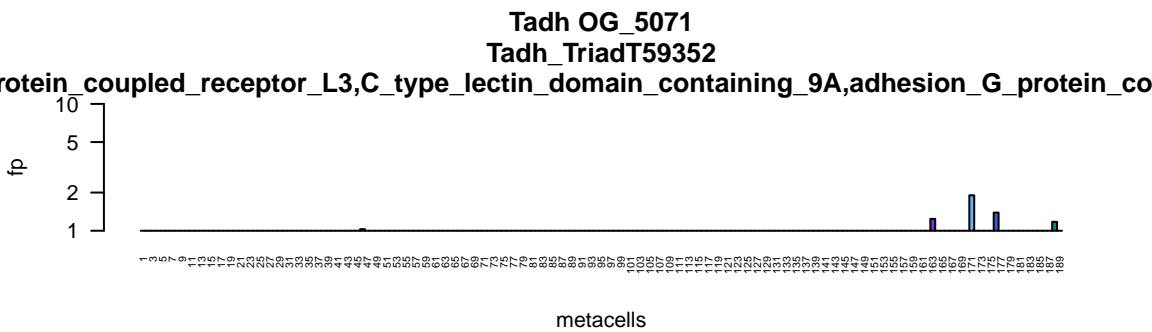


HoiH23 OG_4983
HoiH23_PIH23_009203-RA

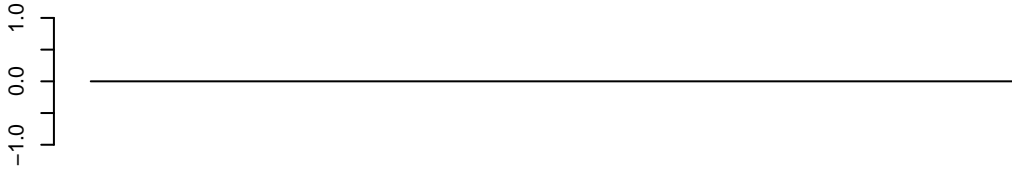


HoiH23 OG_4983
HoiH23_PIH23_009204-RA

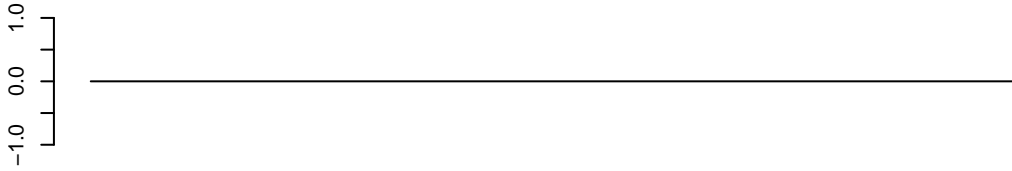




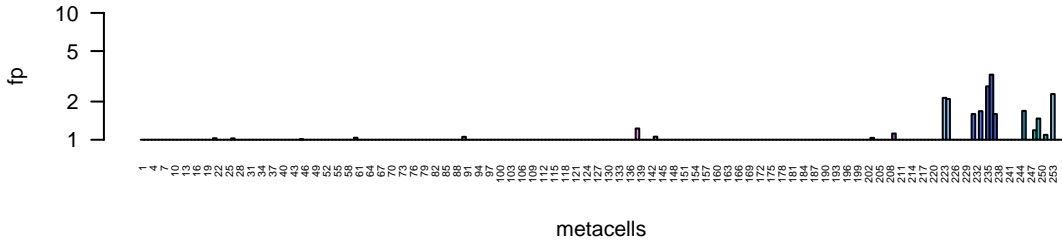
Tadh | no data



TrH2 | no data

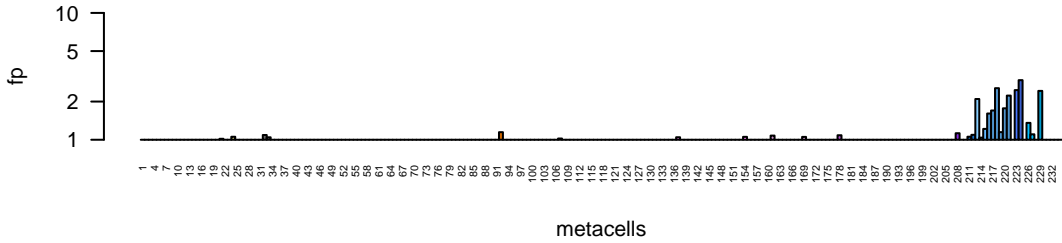


Hhon OG_5311
Hhon_g07916.t1

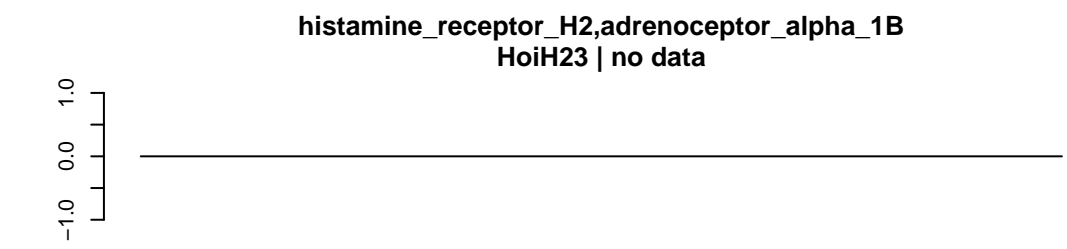
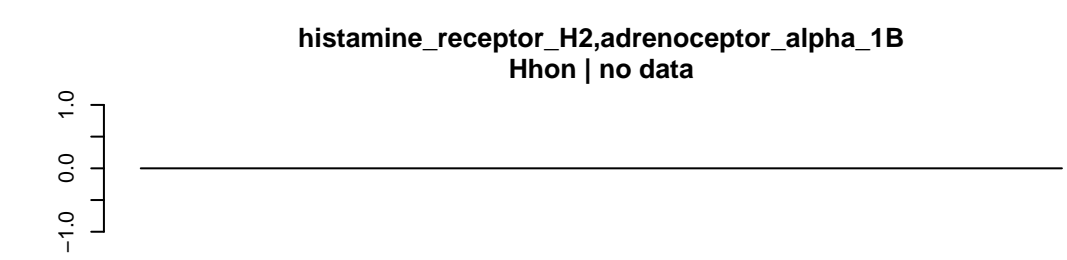
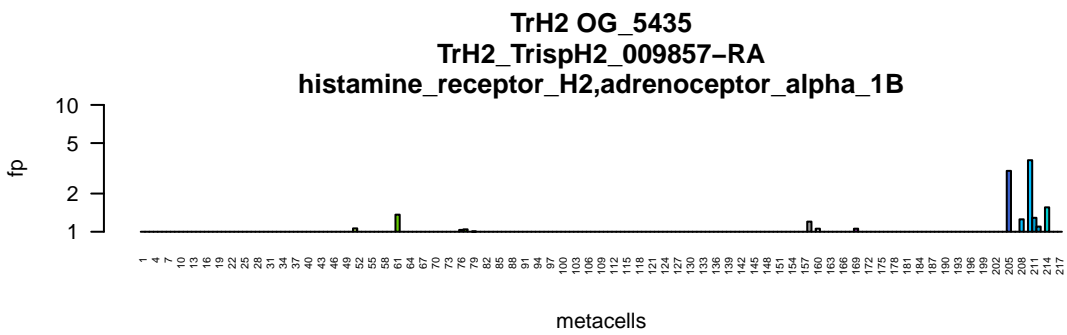
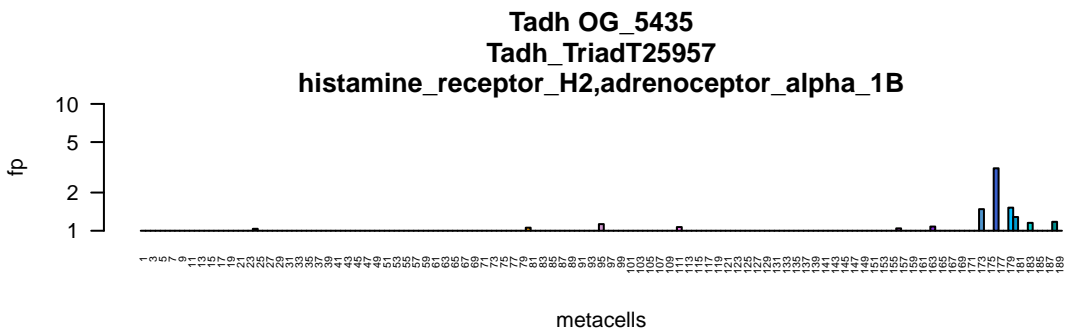


metacells

HoiH23 OG_5311
HoiH23_PIH23_006433-RA



metacells



Bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis lists metacells from 1 to 189. The y-axis ranges from 1 to 10. The chart shows a distribution of feature counts across metacells, with most metacells having 1 or 2 features, and a few having up to 6 features.

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 189 metacells. Most metacells have a false positive count of 1, with a few having counts of 2 or 3. Metacells 175, 176, and 179 have the highest counts, all at 4.

metacell	fp
1	1
2	1
3	1
4	1
5	1
6	1
7	1
8	1
9	1
10	1
11	1
12	1
13	1
14	1
15	1
16	1
17	1
18	1
19	1
20	1
21	1
22	1
23	1
24	1
25	1
26	1
27	1
28	1
29	1
30	1
31	1
32	1
33	1
34	1
35	1
36	1
37	1
38	1
39	1
40	1
41	1
42	1
43	1
44	1
45	1
46	1
47	1
48	1
49	1
50	1
51	1
52	1
53	1
54	1
55	1
56	1
57	1
58	1
59	1
60	1
61	1
62	1
63	1
64	1
65	1
66	1
67	1
68	1
69	1
70	1
71	1
72	1
73	1
74	1
75	1
76	1
77	1
78	1
79	1
80	1
81	1
82	1
83	1
84	1
85	1
86	1
87	1
88	1
89	1
90	1
91	1
92	1
93	1
94	1
95	1
96	1
97	1
98	1
99	1
100	1
101	1
102	1
103	1
104	1
105	1
106	1
107	1
108	1
109	1
110	1
111	1
112	1
113	1
114	1
115	1
116	1
117	1
118	1
119	1
120	1
121	1
122	1
123	1
124	1
125	1
126	1
127	1
128	1
129	1
130	1
131	1
132	1
133	1
134	1
135	1
136	1
137	1
138	1
139	1
140	1
141	1
142	1
143	1
144	1
145	1
146	1
147	1
148	1
149	1
150	1
151	1
152	1
153	1
154	1
155	1
156	1
157	1
158	1
159	1
160	1
161	1
162	1
163	1
164	1
165	1
166	1
167	1
168	1
169	1
170	1
171	1
172	1
173	1
174	1
175	4
176	4
177	1
178	1
179	4
180	1
181	1
182	1
183	1
184	1
185	1
186	1
187	1
188	1
189	1

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 180 metacells. Most metacells have 0 false positives, but some have 1, 2, or more. The bars are colored in a repeating pattern of black, purple, and green.

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 189 metacells. The bars are colored in a gradient from blue to red. Most metacells have a false positive count of 1, with a few having counts of 2 or 3.

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 189 metacells. Most metacells have 1 false positive, with a few having 2 or 4. Metacells 175, 176, 177, 178, and 179 have 4 false positives each.

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 metacells from 1 to 217. The bars are colored in a repeating pattern of light blue, light green, and light red. Most metacells have a false positive count of 1, with some having 2 or 3. Metacells 178 and 181 have the highest false positive counts, at 5 and 4 respectively.

Bar chart showing the frequency of metacells (x-axis) versus the frequency of the top 1000 most frequent metacells (y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows a distribution of frequencies across the metacells, with a peak around metacell 145.

A bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows 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
23	1
25	1
28	1
31	1
32	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
202	1
205	1
208	1
211	1
214	1
217	1

A bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a few outliers reaching up to 7.

metacells	fp
1	1
4	1
10	1
13	1
16	1.5
22	1
25	1
28	1
31	1
32	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1.2
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1.5
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
202	2.5
205	1.5
208	4.5
211	4
214	1
217	1

A bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a few outliers reaching up to 2.

metacells	fp
1	1
4	1
10	1
13	1
16	1
22	1
23	1
25	1
28	1
31	1
32	1
37	1
40	1
43	2
44	1
48	1
49	1
52	1
55	1
59	1
61	1
64	1
67	1
73	1
76	1
79	1
82	1
83	1
88	1
91	1
94	1
97	1
100	1
103	1
108	1
110	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	2
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
200	1
202	2
205	1
208	1
211	1
214	1
217	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 217. Most metacells have a false positive count of 1, with a few having counts of 2 or 3. Metacells 205 and 206 show the highest counts, around 4.

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 metacells from 1 to 217. Most metacells have 0 false positives, but some have 1 or 2. Metacells 181 and 202 have 2 false positives each.

metacell	fp
1	0
4	0
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	0
34	0
37	1
40	1
43	0
46	0
49	0
52	0
55	0
58	0
61	0
64	1
67	0
70	0
73	0
76	0
79	0
82	0
85	0
88	0
91	0
94	1
97	1
100	0
103	0
106	0
109	0
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	0
136	0
139	0
142	1
145	1
148	1
151	1
154	0
157	0
160	0
163	0
166	0
169	0
172	0
175	0
178	1
181	2
184	0
187	0
190	0
193	0
196	1
199	1
202	2
205	0
208	0
211	0
214	0
217	1

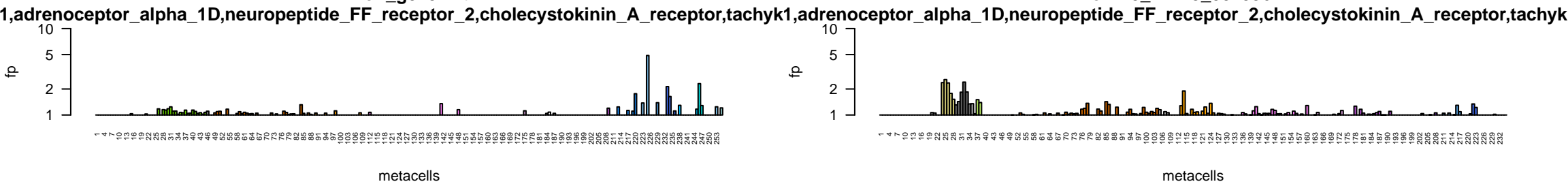
A bar chart showing the number of false positives (fp) for each metacell. The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 0 to 10. The chart shows a distribution of false positives across metacells, with most having 0 or 1 false positive, and a few having up to 4.

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

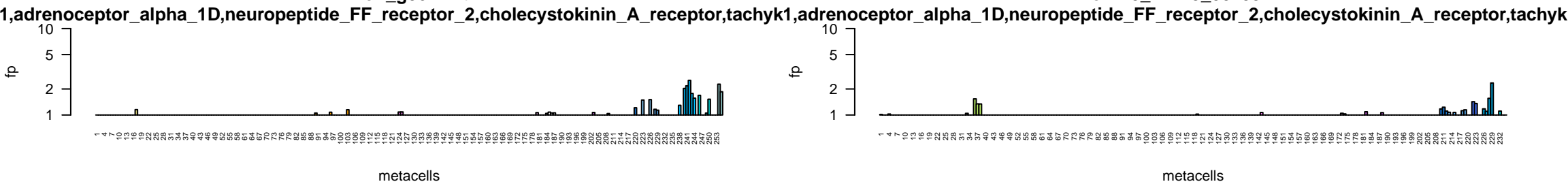
A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis ranges from 1 to 250, and the y-axis ranges from 1 to 10. Most metacells have a frequency of 1, with a small cluster of higher frequencies around metacell 240.

metacells	fp
1	1
7	1
10	1
16	1
18	1
19	1
25	1
26	1
28	1
34	1
37	1
43	1
46	1
55	1
61	1
64	1
70	1
76	1
79	1
85	1
88	1
94	1
103	1
112	1
115	1
121	1
122	1
127	1
130	1
136	1
145	1
148	1
154	1
157	1
163	1
166	1
172	1
175	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
205	1
208	1
214	1
217	1
223	1
228	1
232	1
235	1
236	1
241	1
242	1
244	1
247	1
250	1
253	1

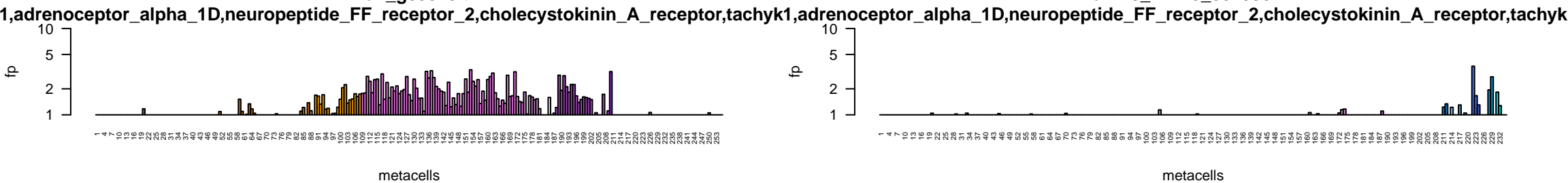
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Hhon_g04574.t1



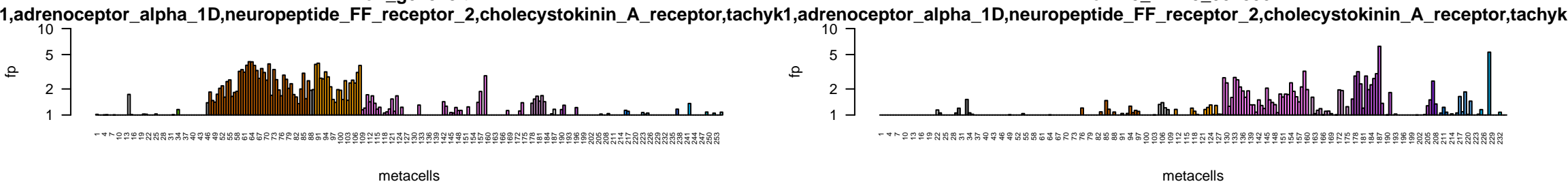
Hhon OG_5725
Hhon_g06114.t1



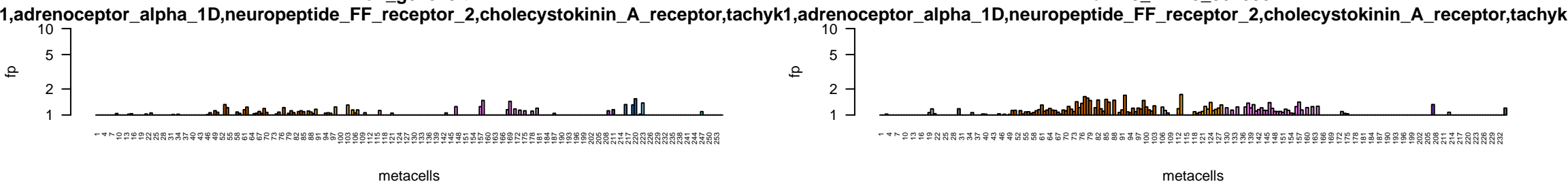
Hhon OG_5725
Hhon_g09573.t1



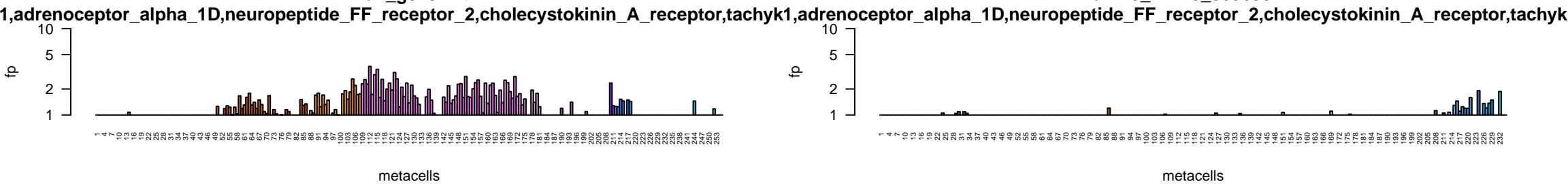
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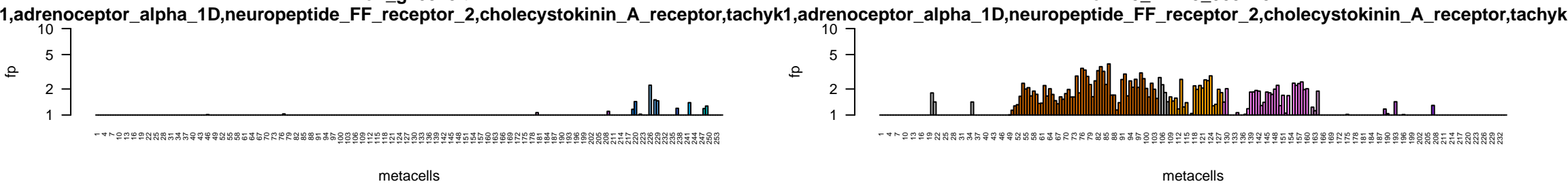
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Hhon_g07529.t1



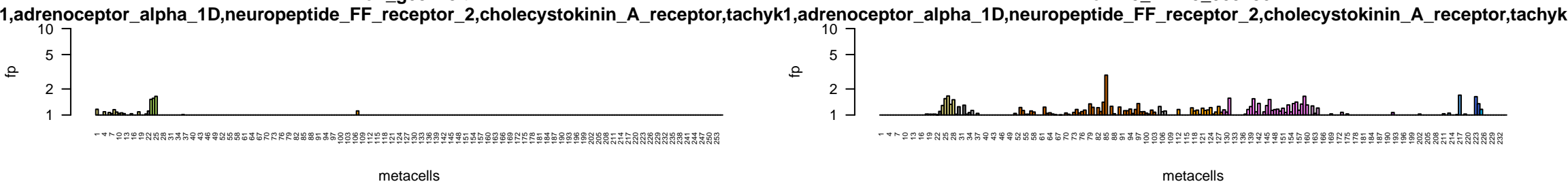
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Hhon_g07527.t1

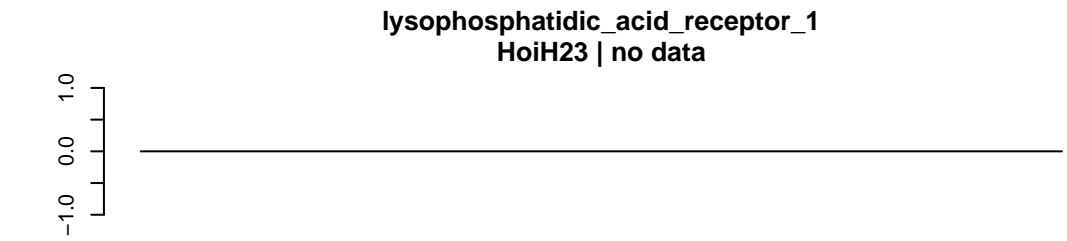
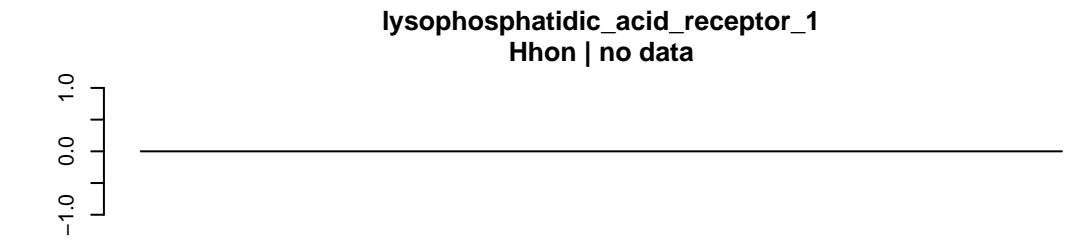
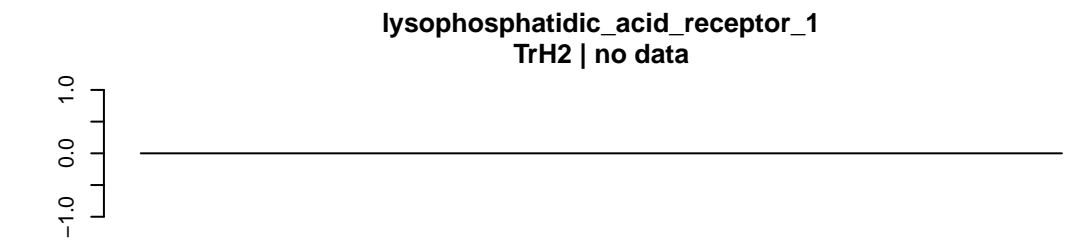
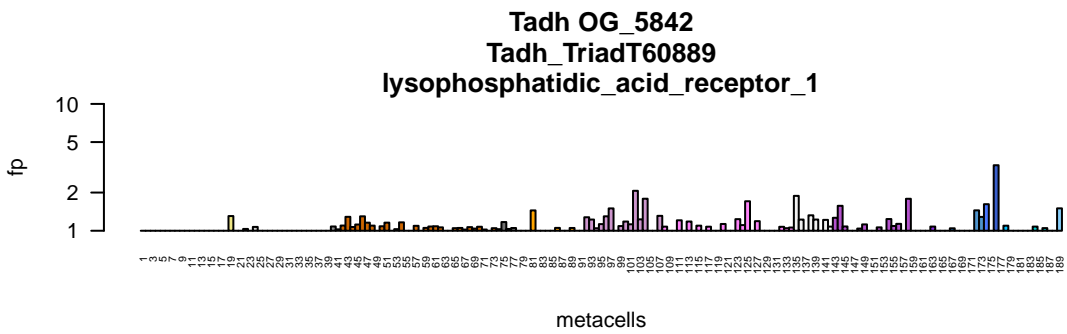


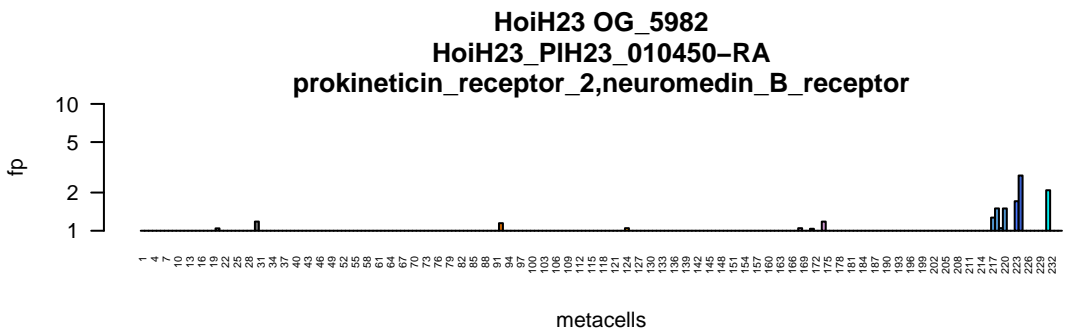
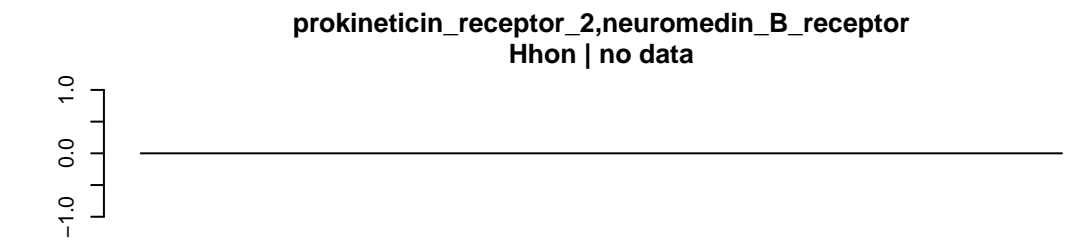
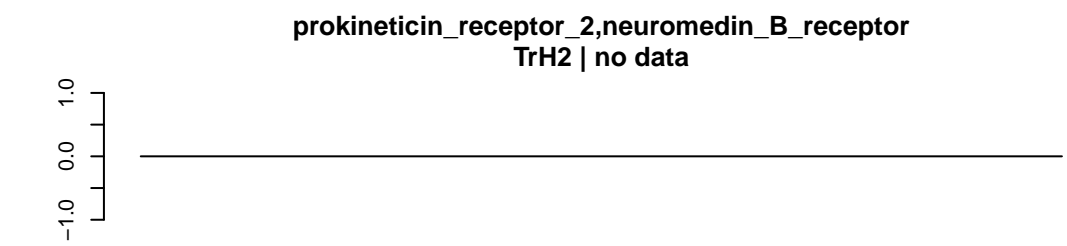
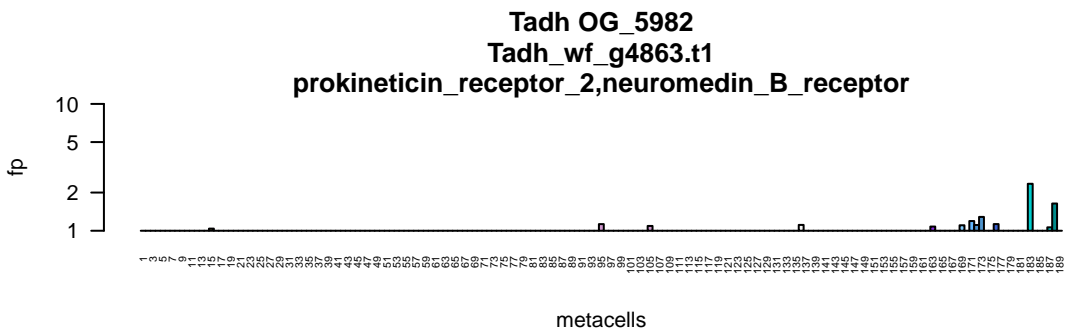
Hhon OG_5725
Hhon_g10923.t1

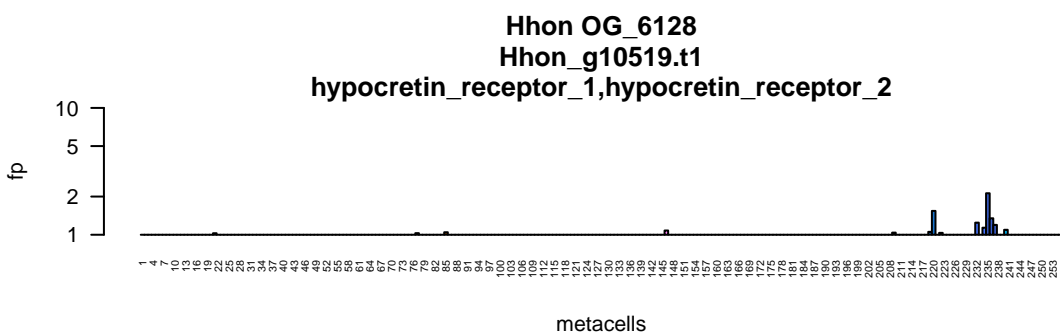
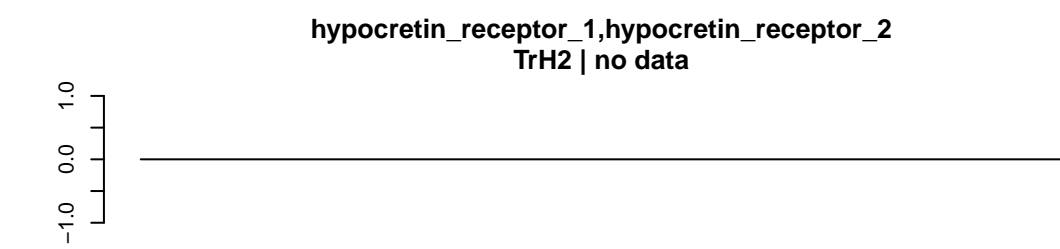
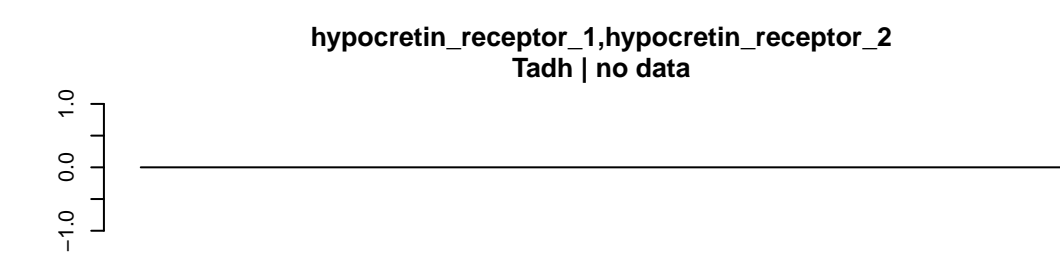


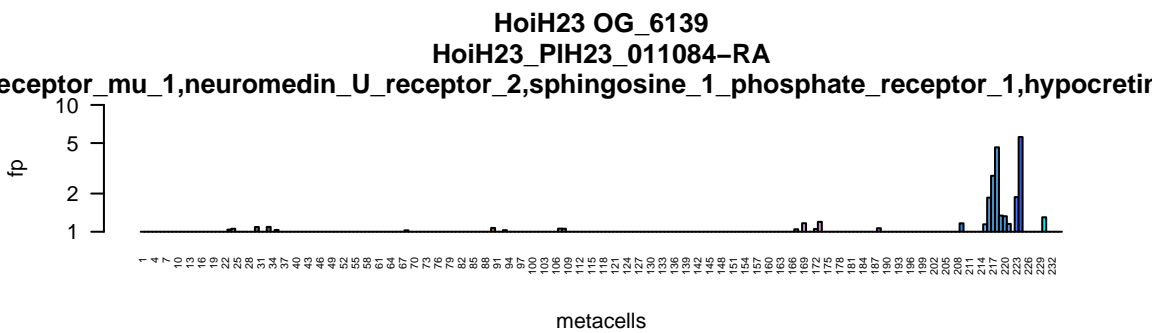
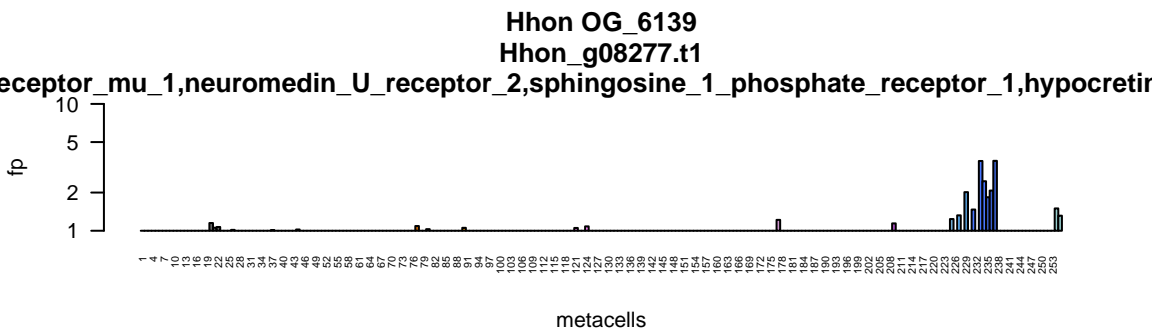
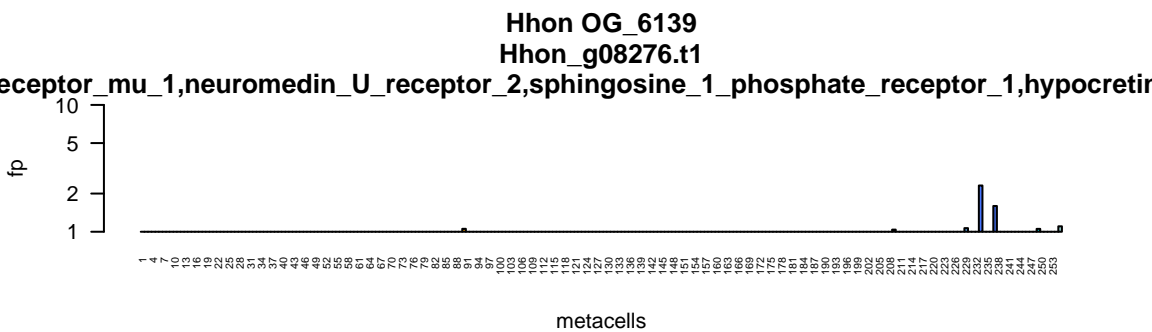
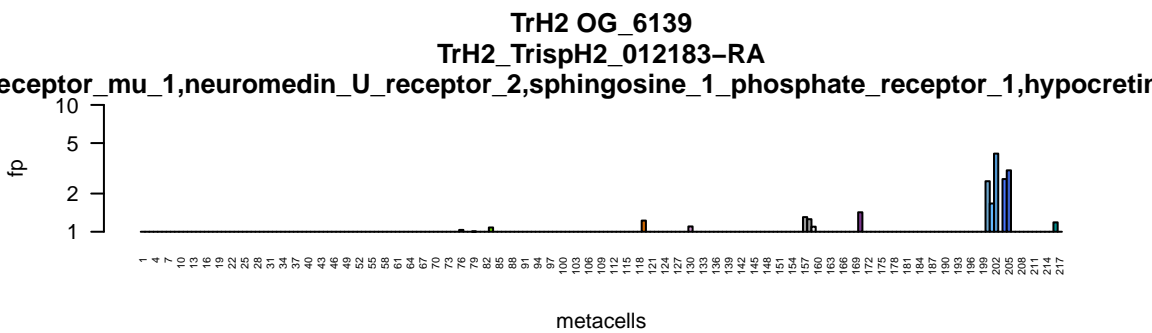
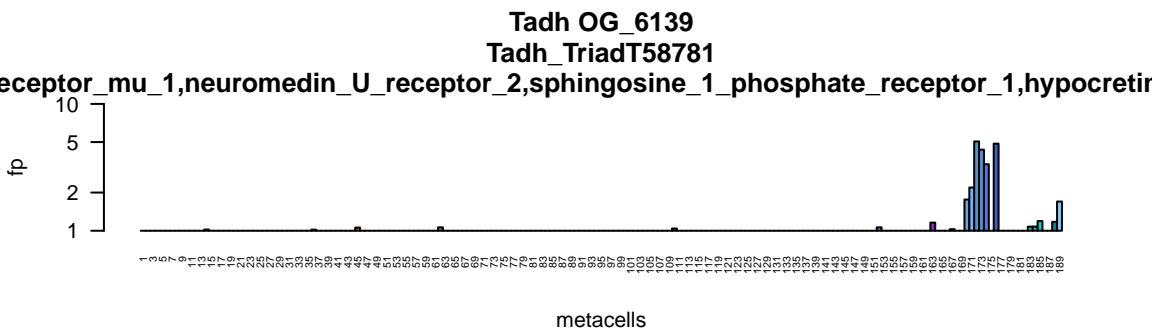
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Hhon_g08418.t1

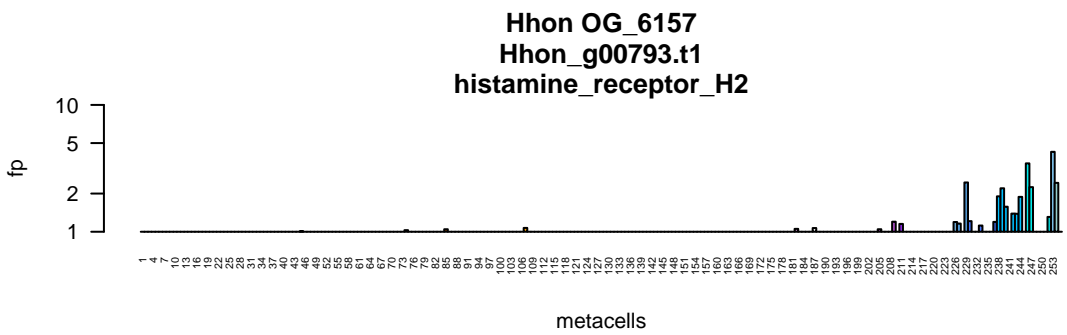
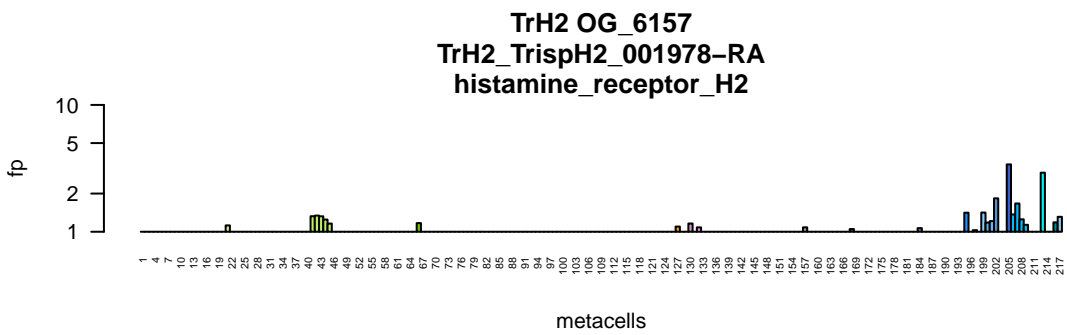
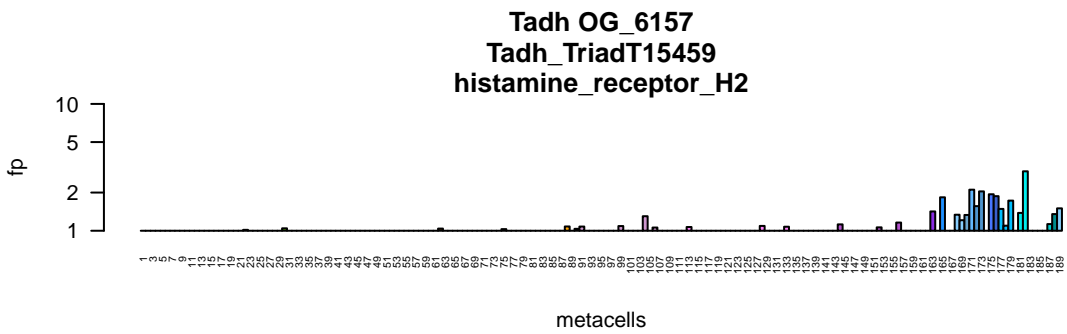












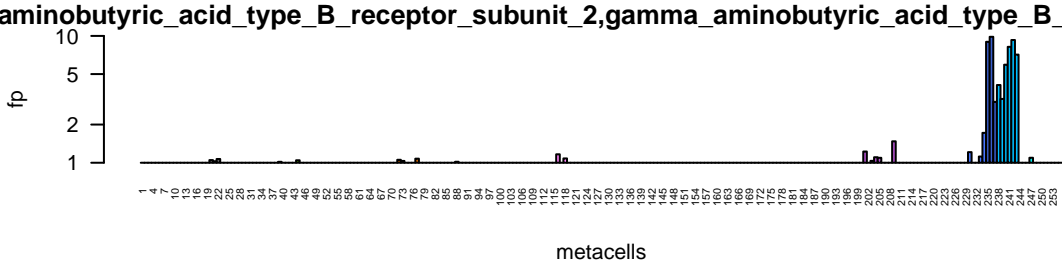
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Tadh | no data



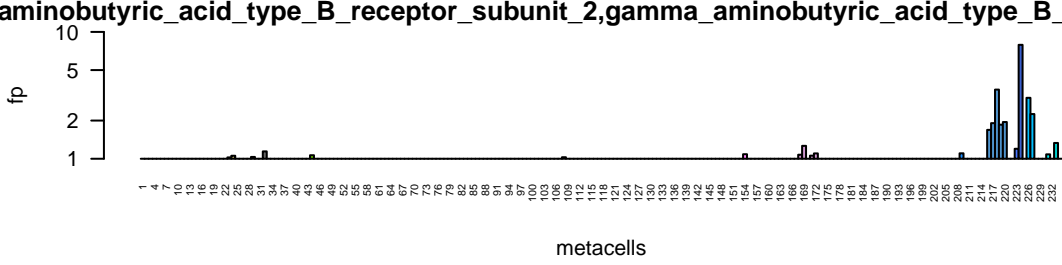
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recept
TrH2 | no data



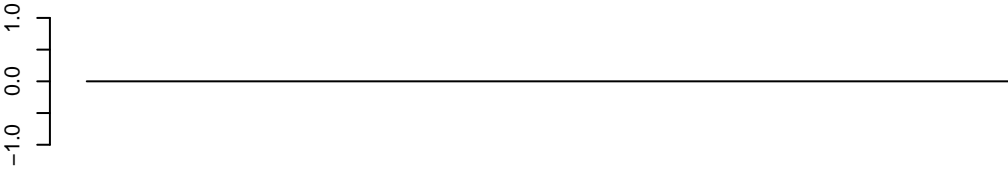
Hhon OG_6183
Hhon_g07966.t1
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recept



HoiH23 OG_6183
HoiH23_PIH23_003840-RA
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recept



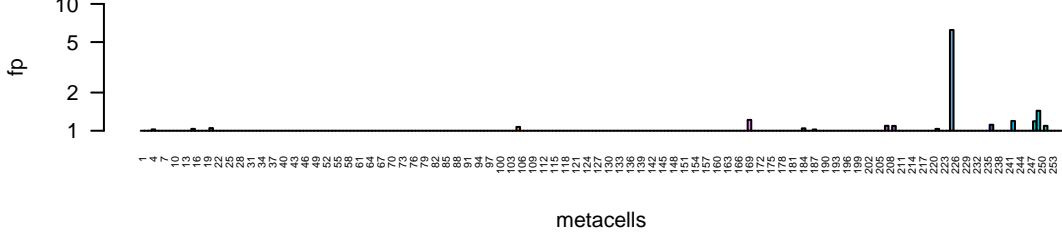
gamma_aminobutyric_acid_type_B_receptor_subunit_2
Tadh | no data



gamma_aminobutyric_acid_type_B_receptor_subunit_2
TrH2 | no data

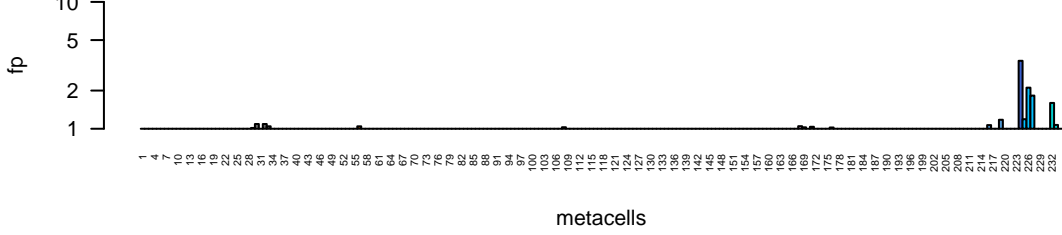


Hhon OG_6184
Hhon_g07965.t1
gamma_aminobutyric_acid_type_B_receptor_subunit_2

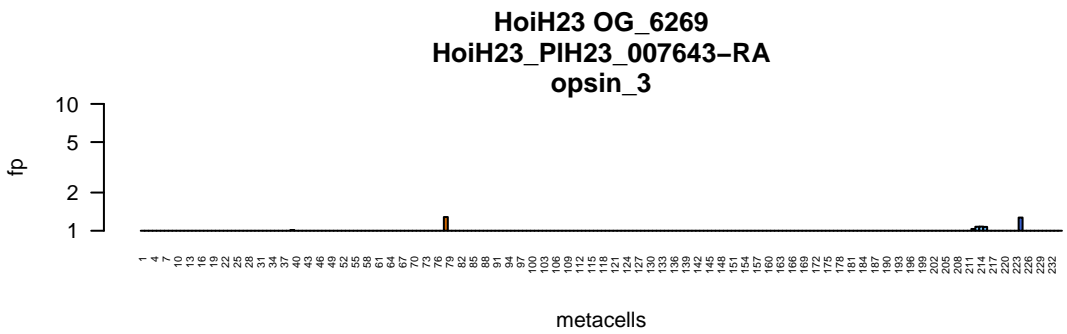
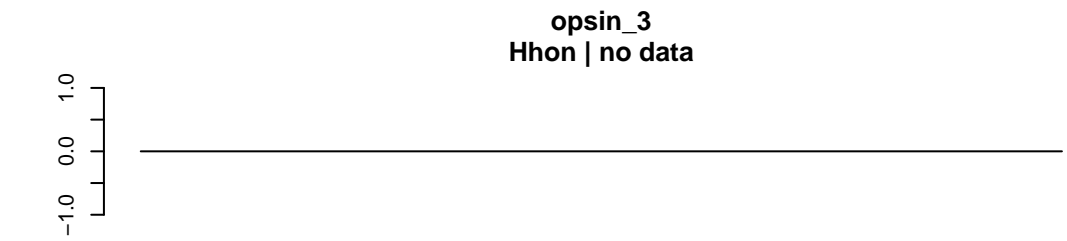
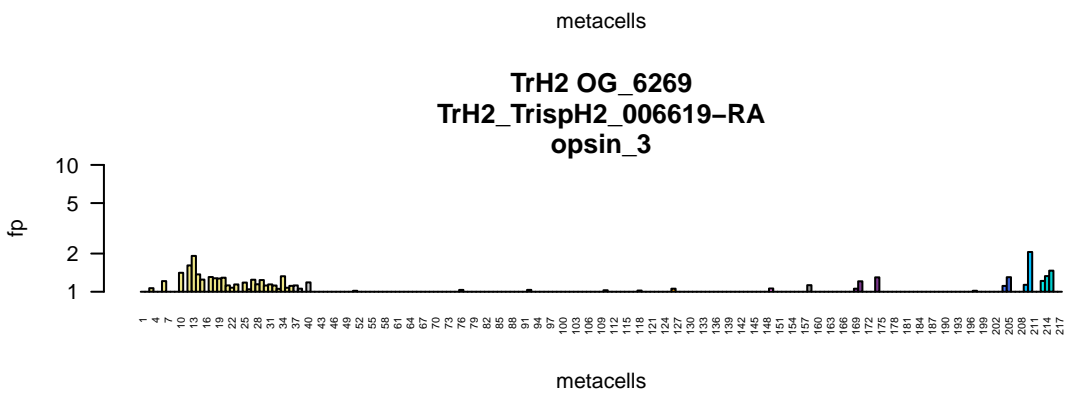
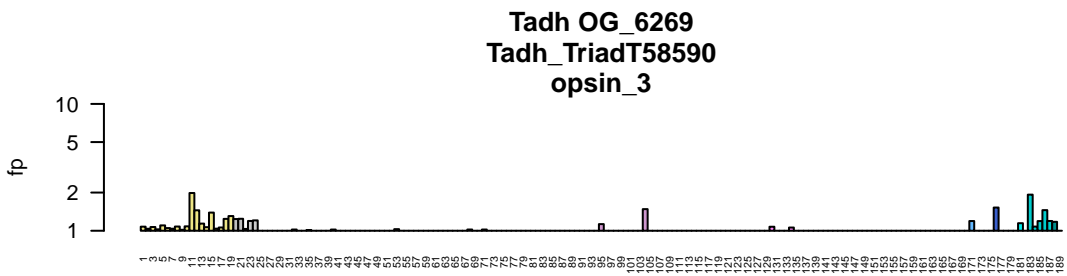


metacells

HoiH23 OG_6184
HoiH23_PIH23_003841-RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2



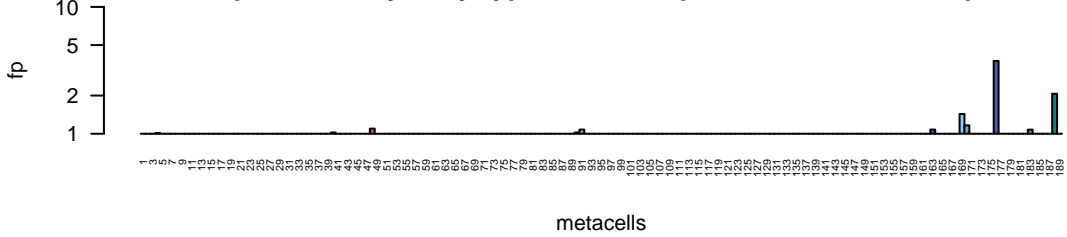
metacells



Tadh OG_6359

Tadh_wf_g9040.t1

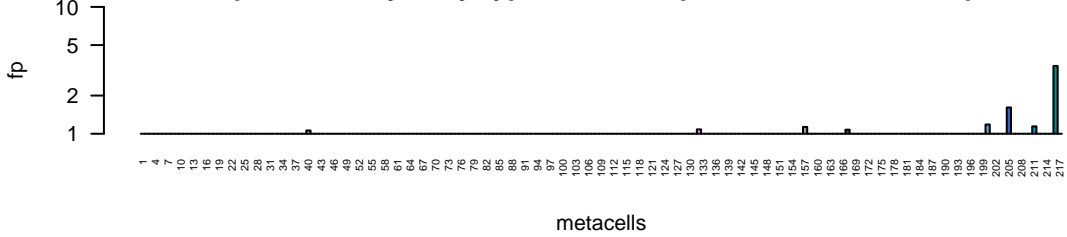
histamine_receptor_H4,5_hydroxytryptamine_receptor_7,bombesin_receptor_subtype_:



metacells

TrH2 OG_6359
TrH2_TrispH2_010326-RA

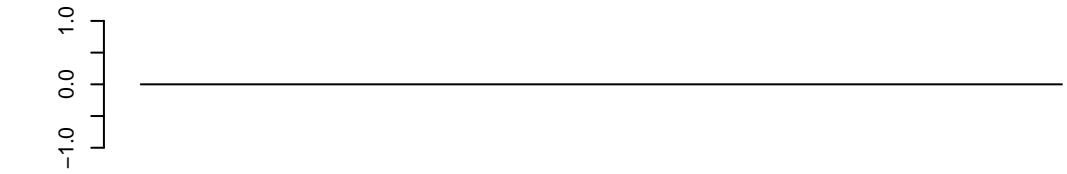
histamine_receptor_H4,5_hydroxytryptamine_receptor_7,bombesin_receptor_subtype_:



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histamine_receptor_H4,5_hydroxytryptamine_receptor_7,bombesin_receptor_subtype_:

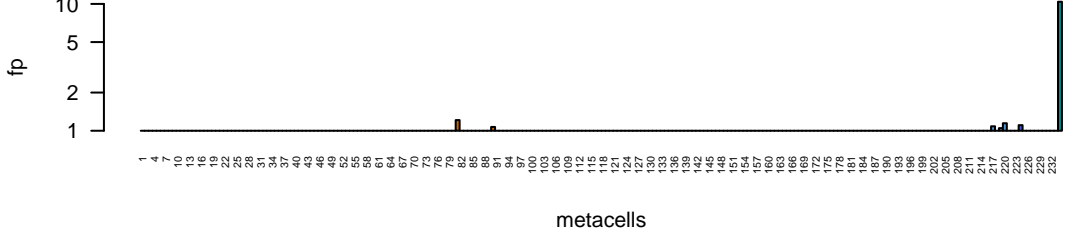
Hhon | no data



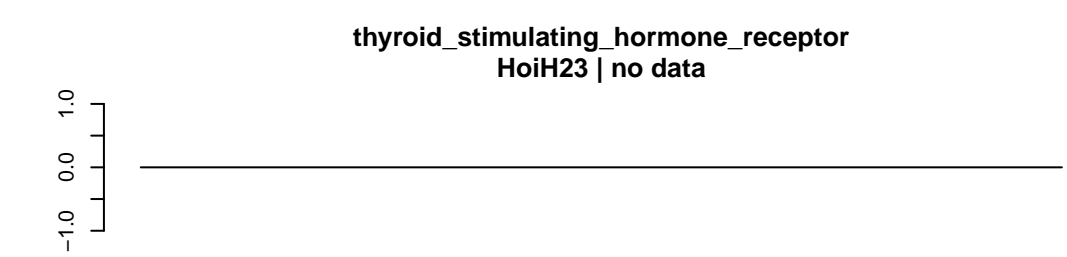
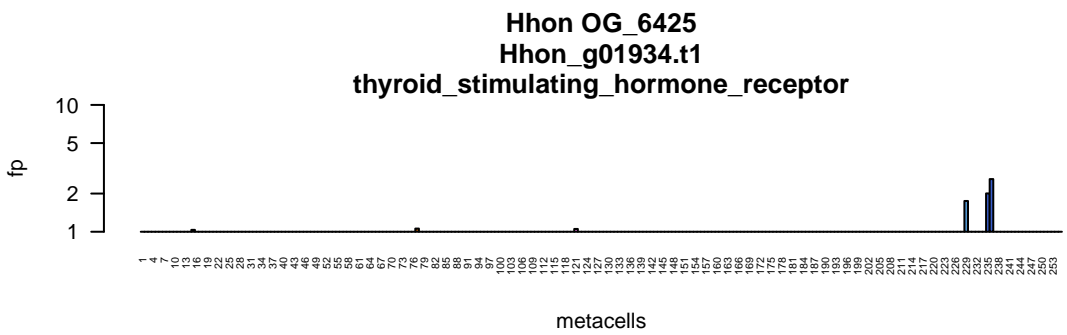
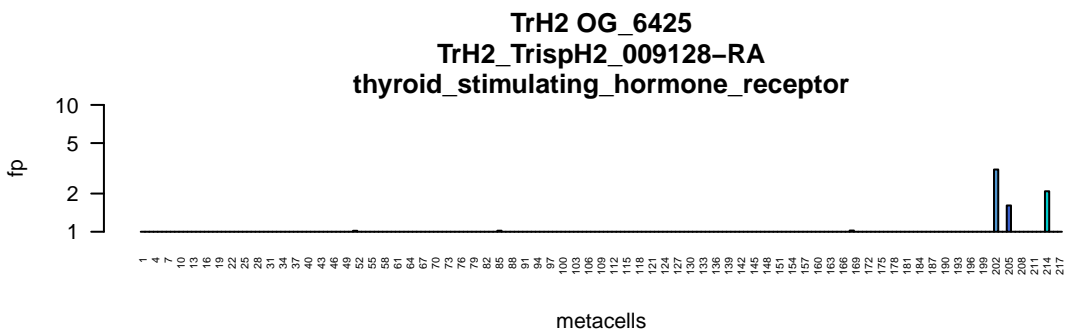
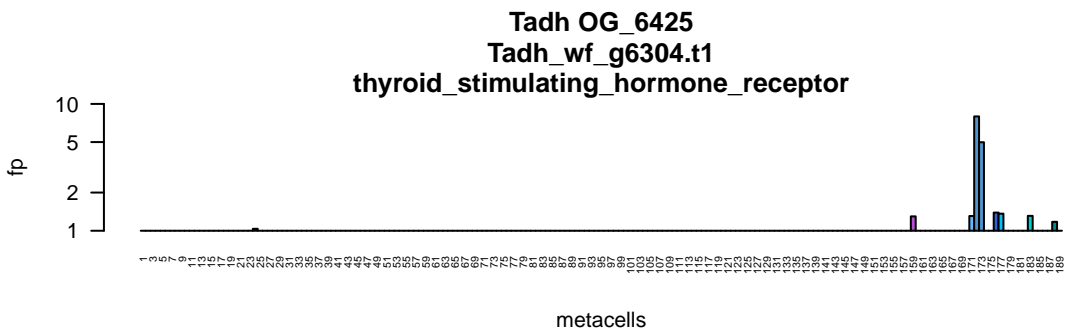
HoiH23 OG_6359

HoiH23_PIH23_011128-RA

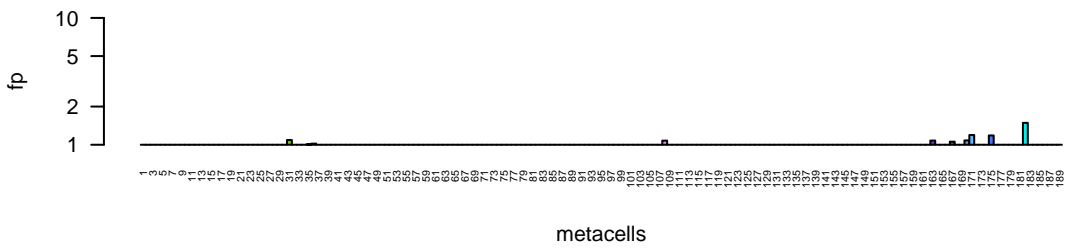
histamine_receptor_H4,5_hydroxytryptamine_receptor_7,bombesin_receptor_subtype_:



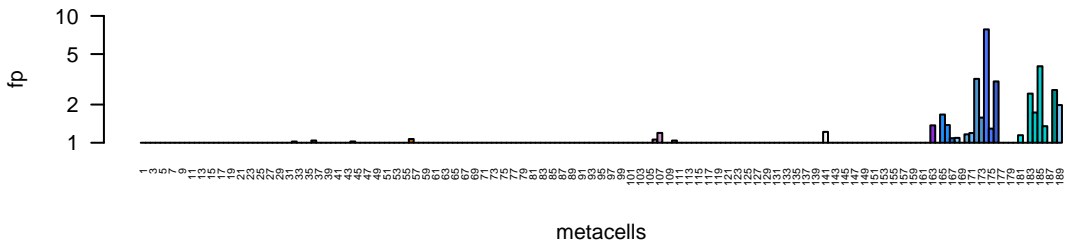
metacells



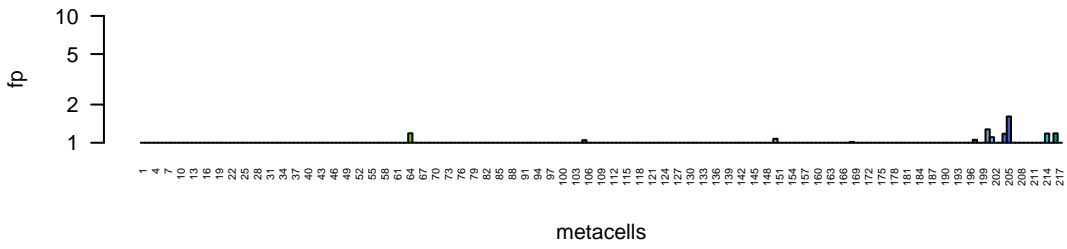
Tadh OG_6482
Tadh_wf_g9560.t1



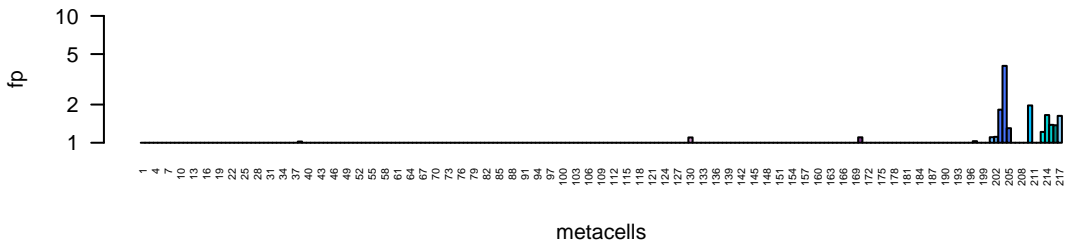
Tadh OG_6482
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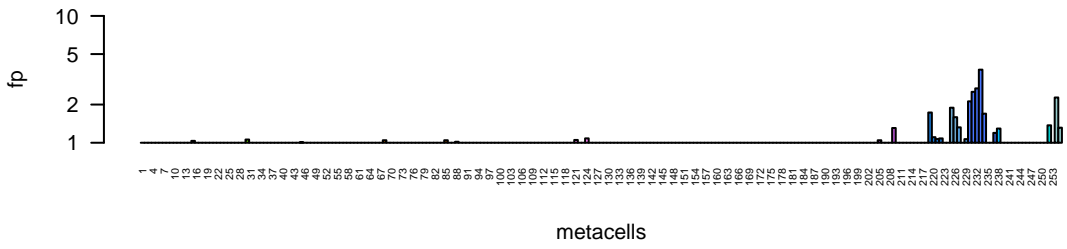
TrH2 OG_6482
TrH2_TrispH2_009482-RA



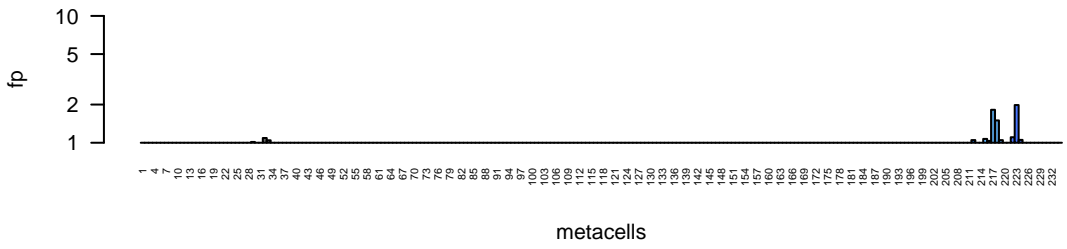
TrH2 OG_6482
TrH2_TrispH2_011478-RA

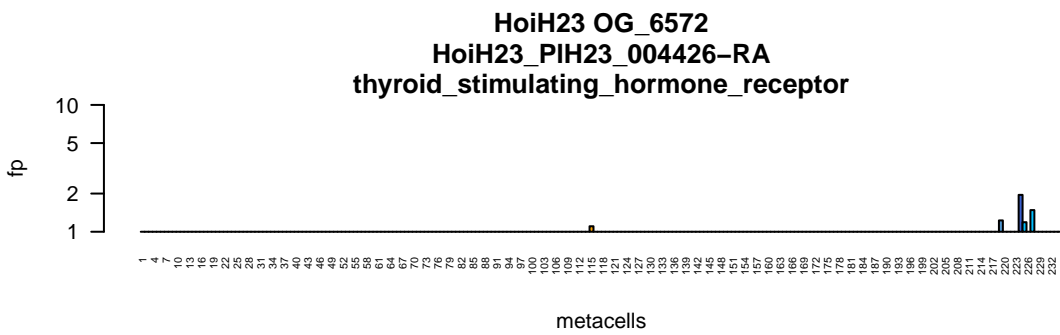
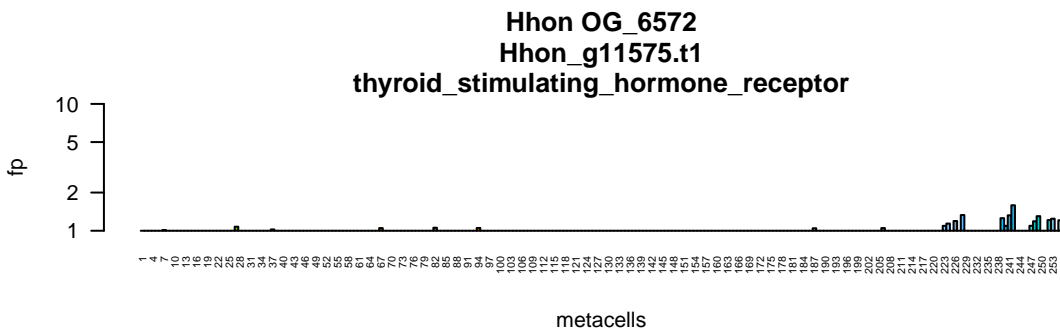
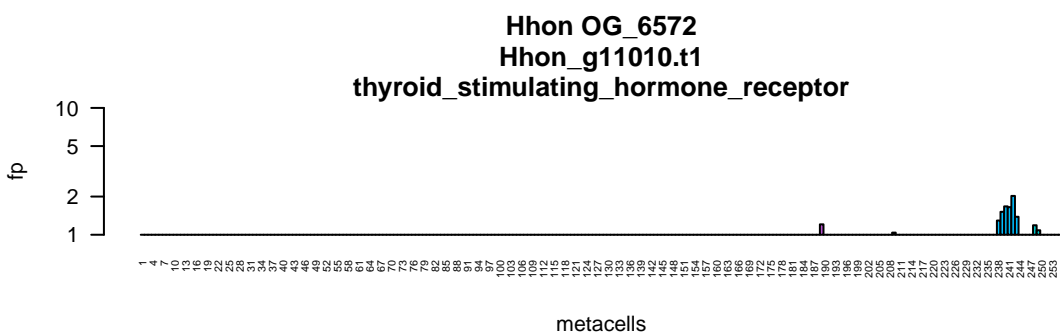
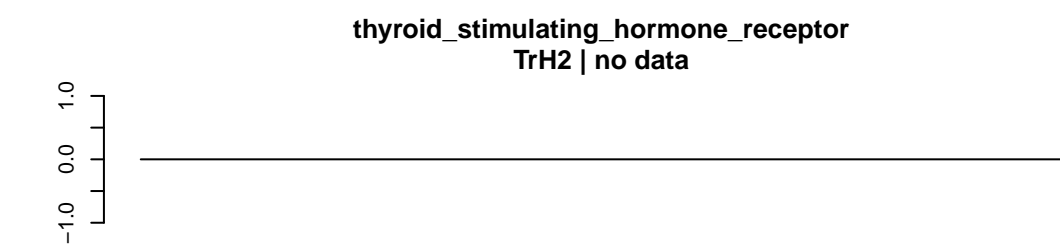
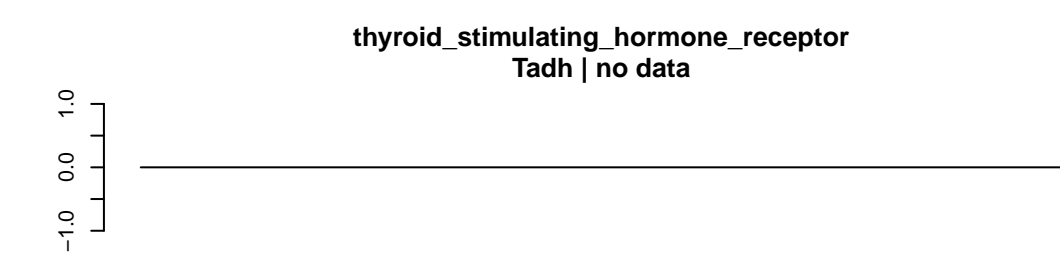


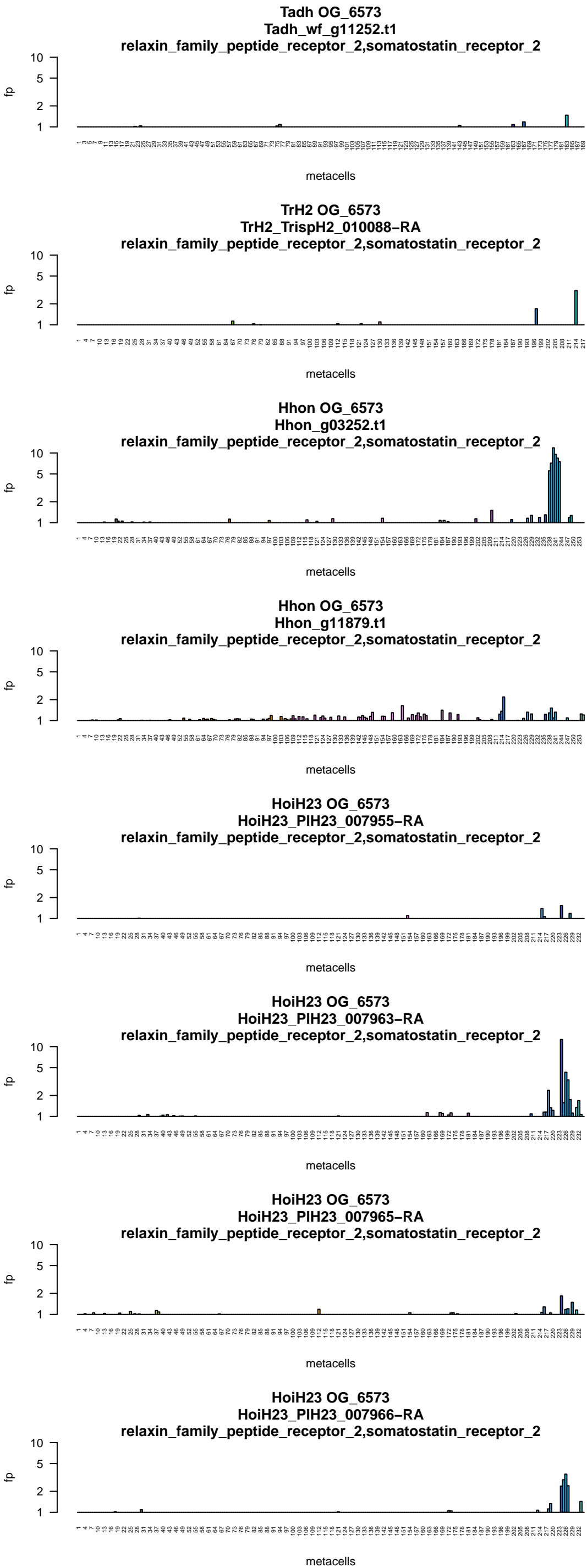
Hhon OG_6482
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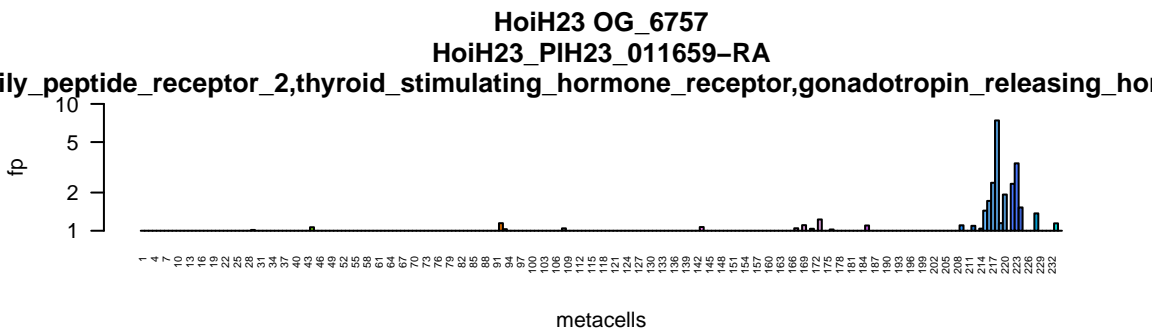
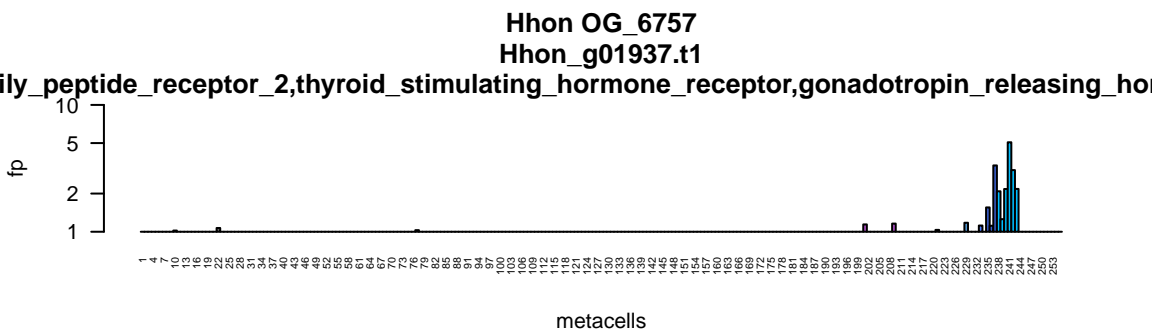
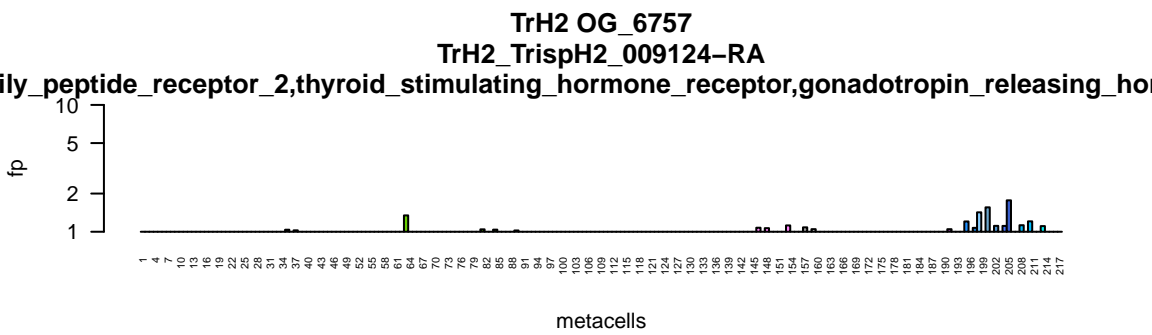
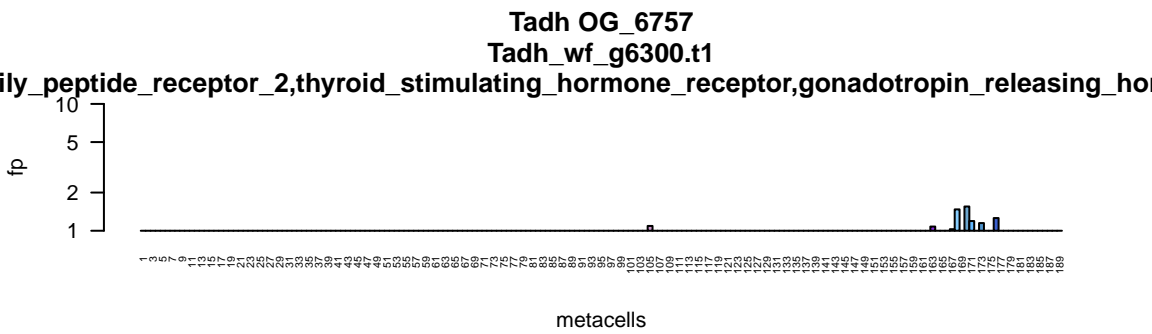


HoiH23 OG_6482
HoiH23_PIH23_011006-RA





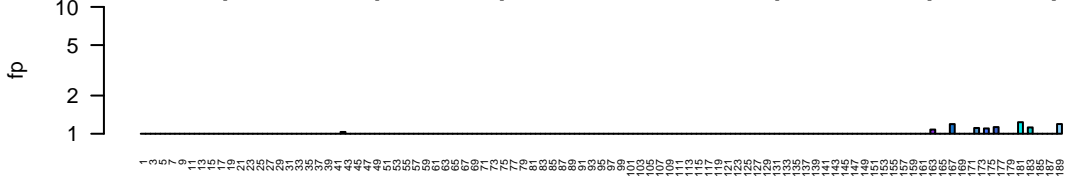




Tadh OG_7054

Tadh_TriadT59349

adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3

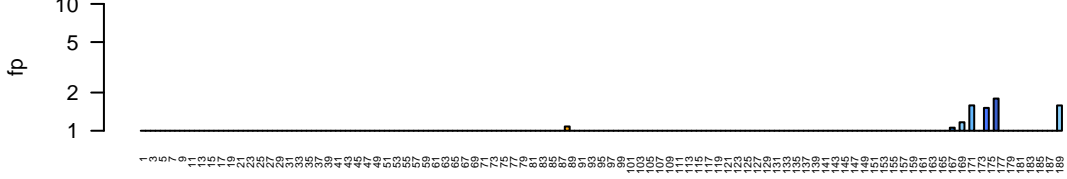


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Tadh OG_7054

Tadh_TriadT59345

adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3

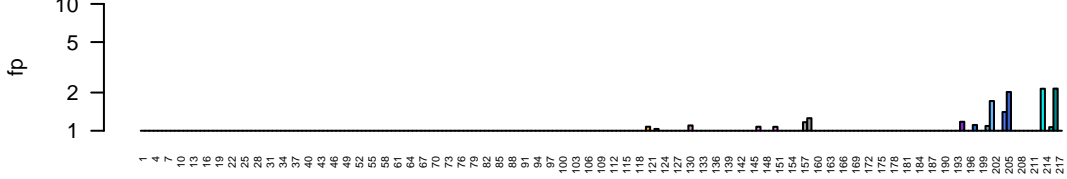


metacells

TrH2 OG_7054

TrH2_TrispH2_011841-RA

adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3

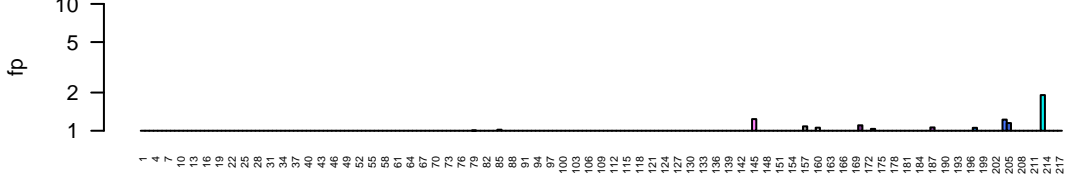


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TrH2 OG_7054

TrH2_TrispH2_010220-RA

adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3

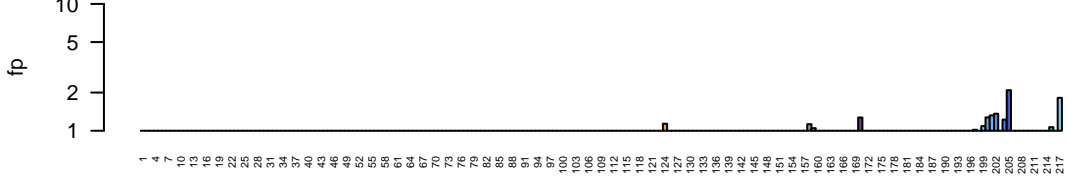


metacells

TrH2 OG_7054

TrH2_TrispH2_010222-RA

adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3

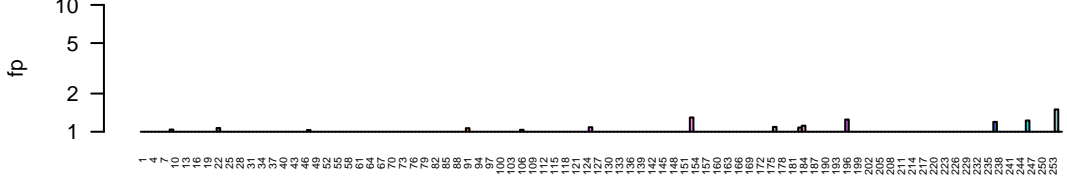


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Hhon OG_7054

Hhon_g10892.t1

adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3

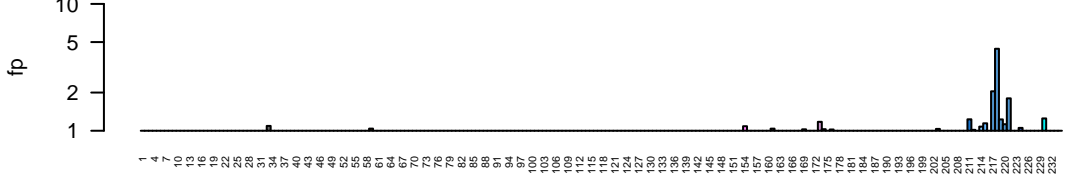


metacells

HoiH23 OG_7054

HoiH23_PIH23_010401-RA

adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3

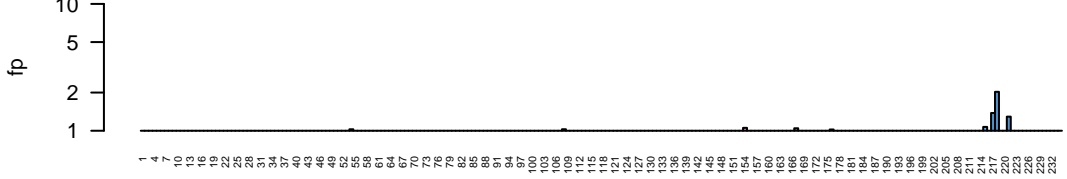


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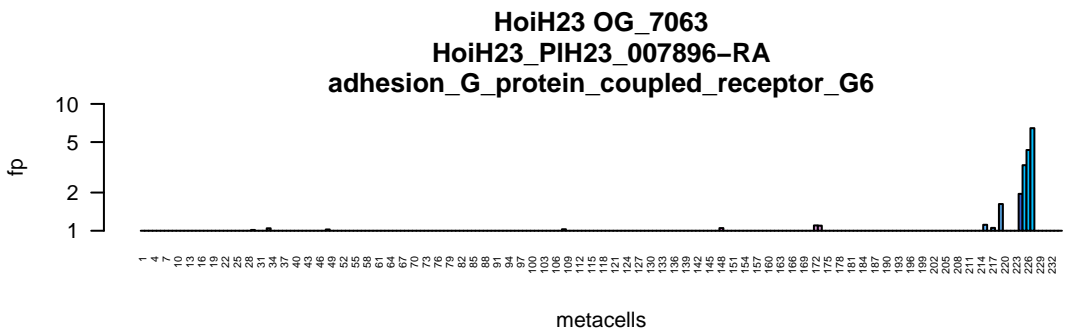
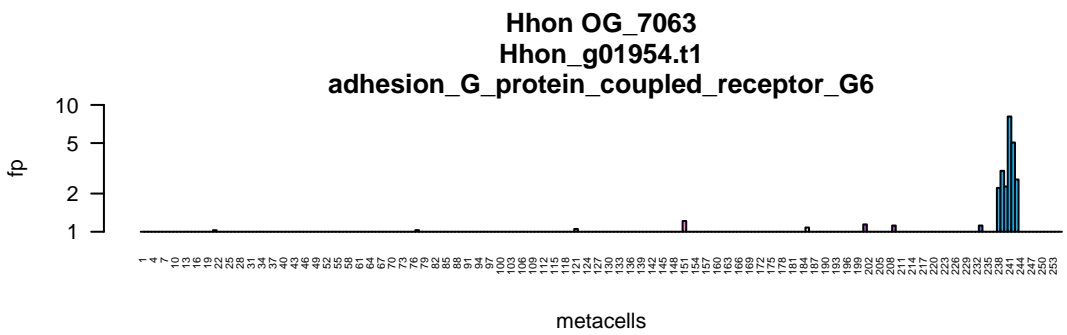
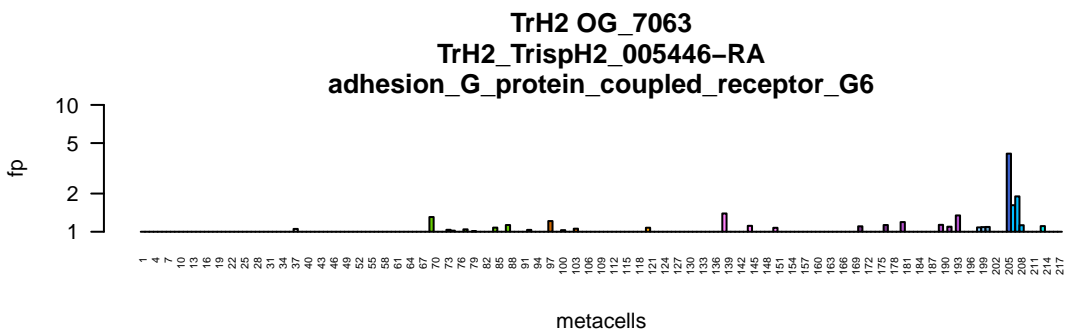
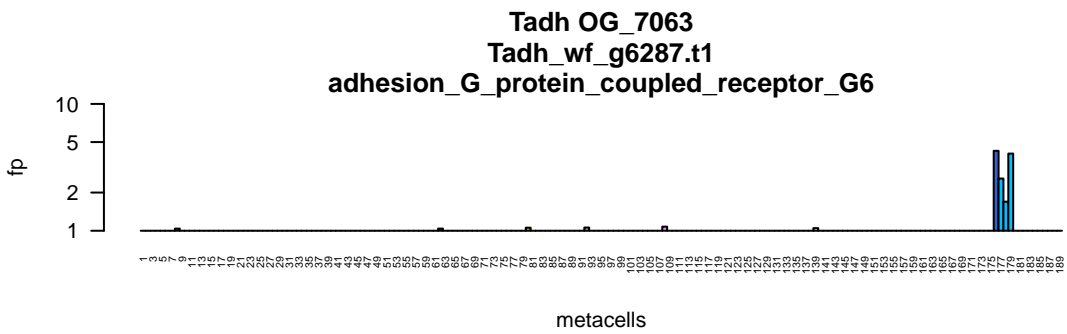
HoiH23 OG_7054

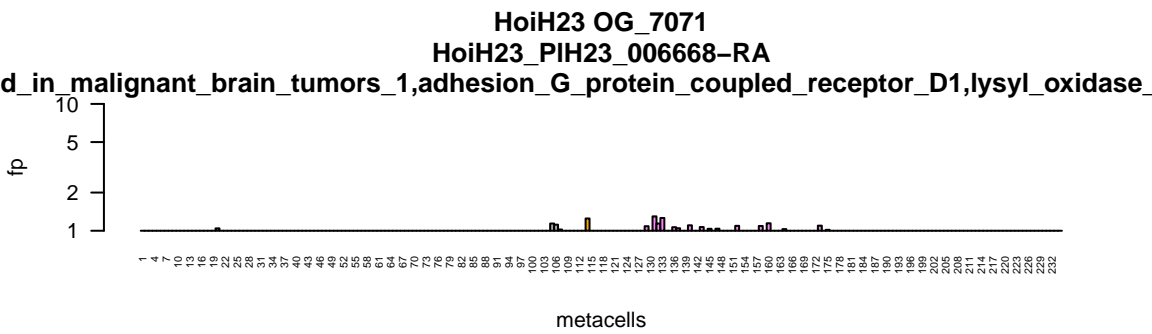
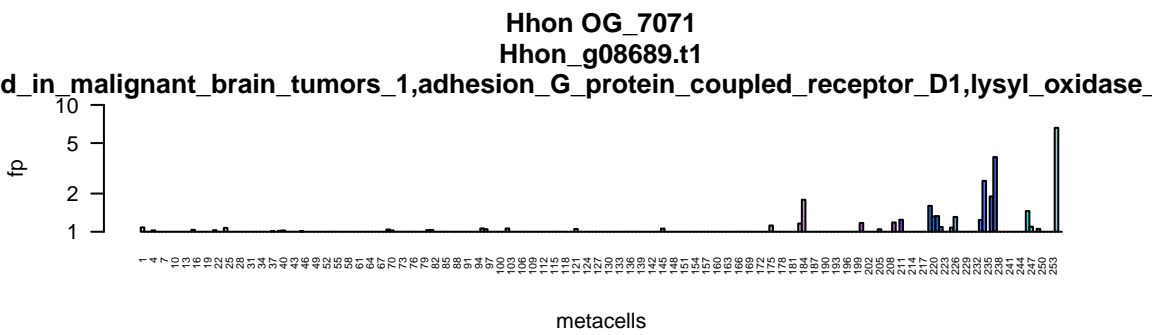
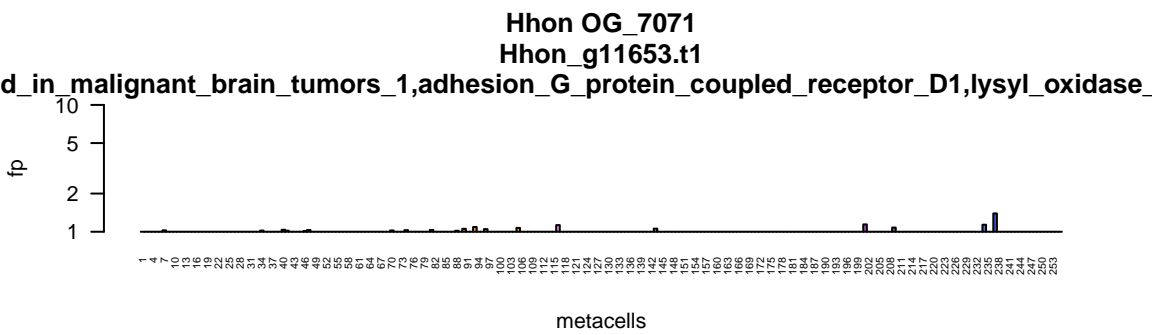
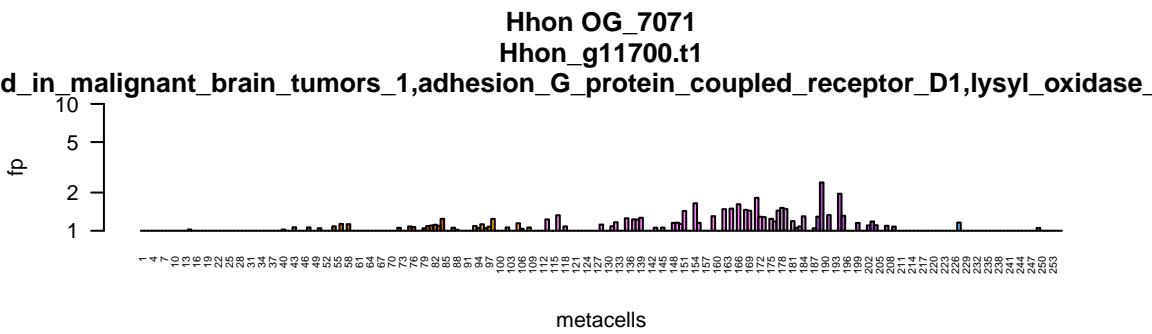
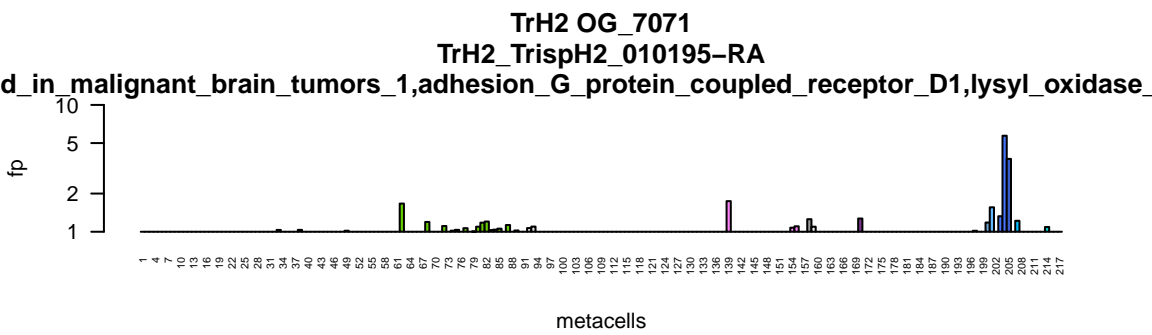
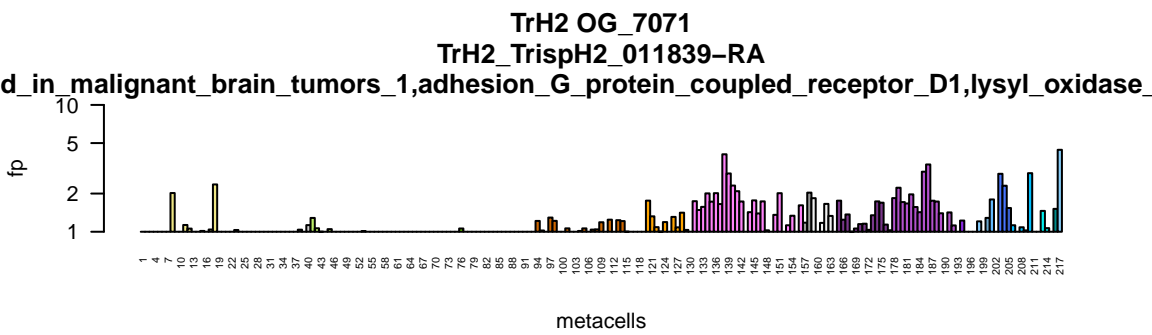
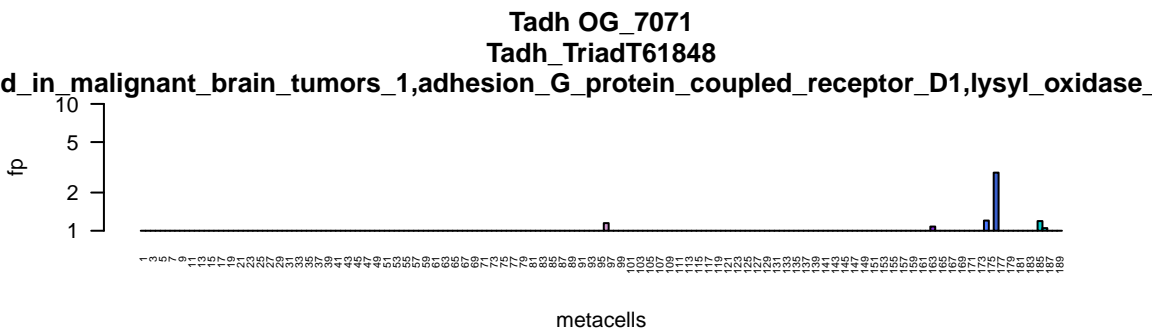
HoiH23_PIH23_006966-RA

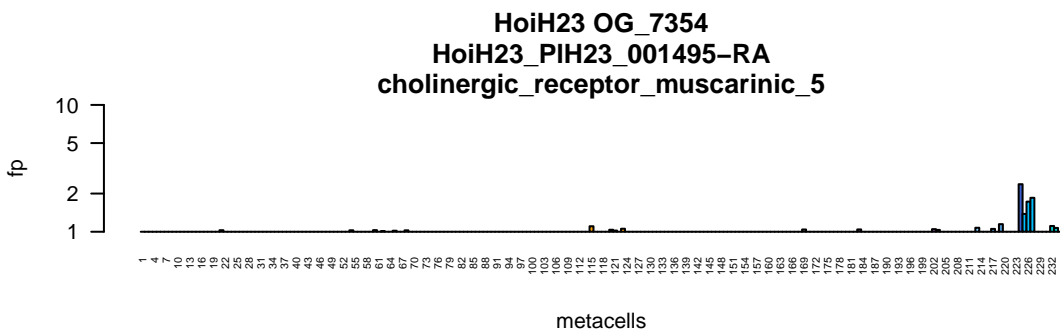
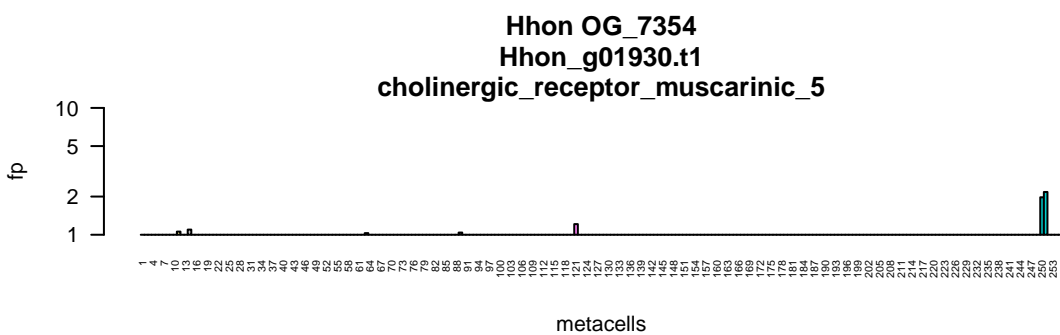
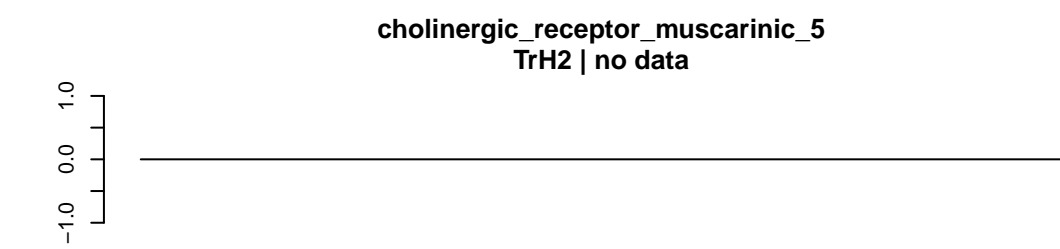
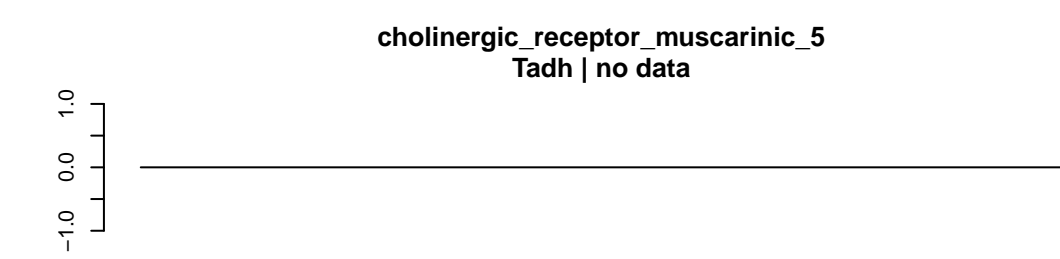
adhesion_G_protein_coupled_receptor_D1,adhesion_G_protein_coupled_receptor_L3

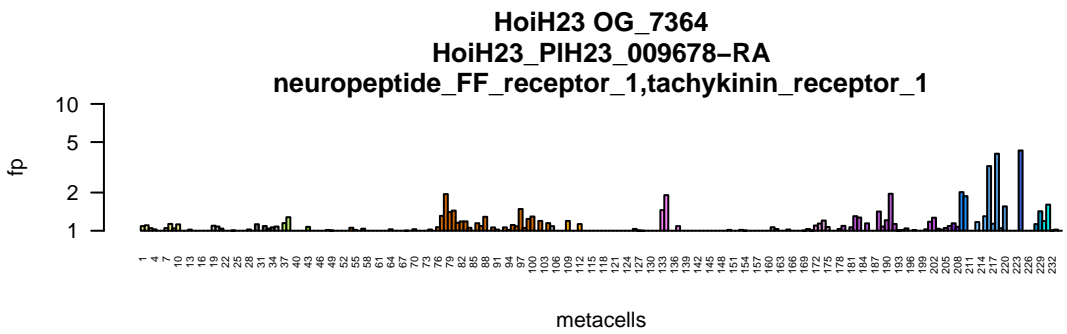
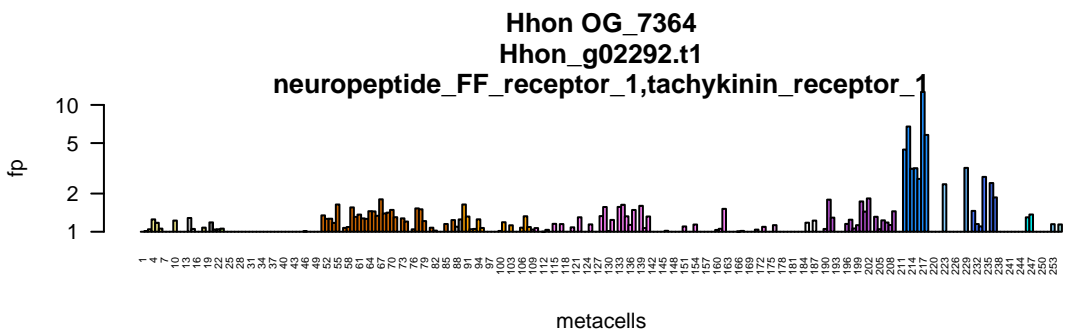
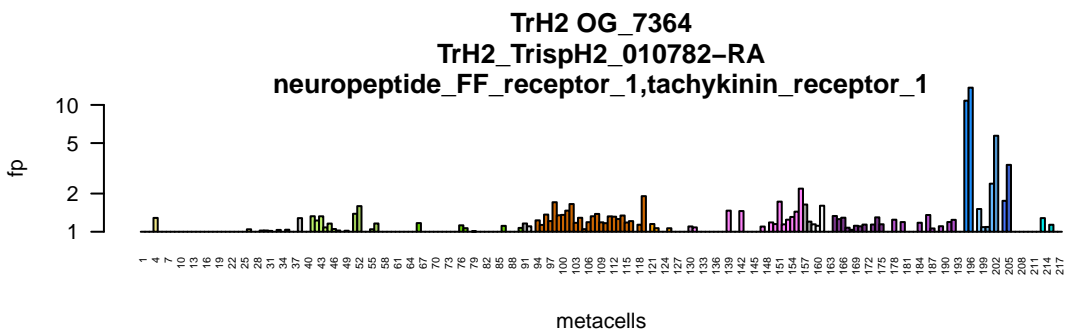
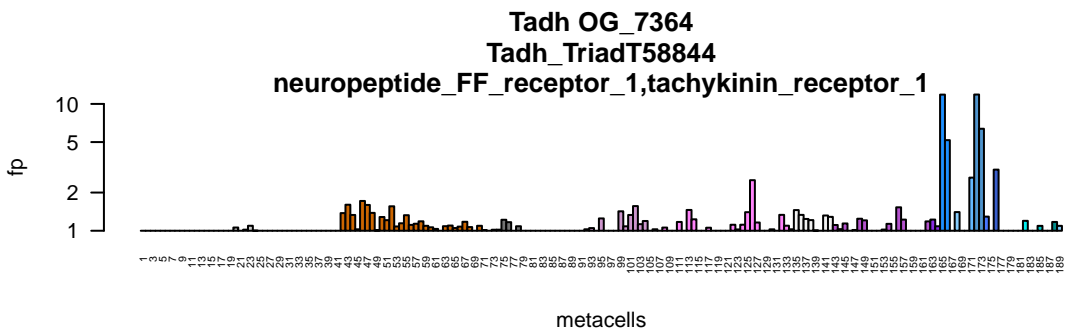


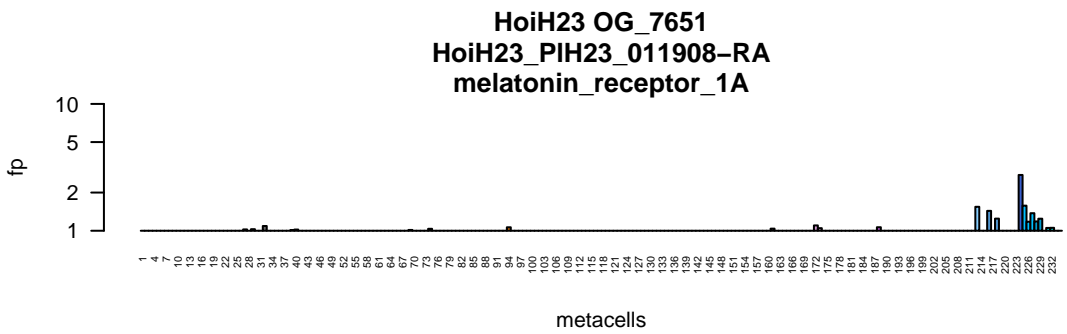
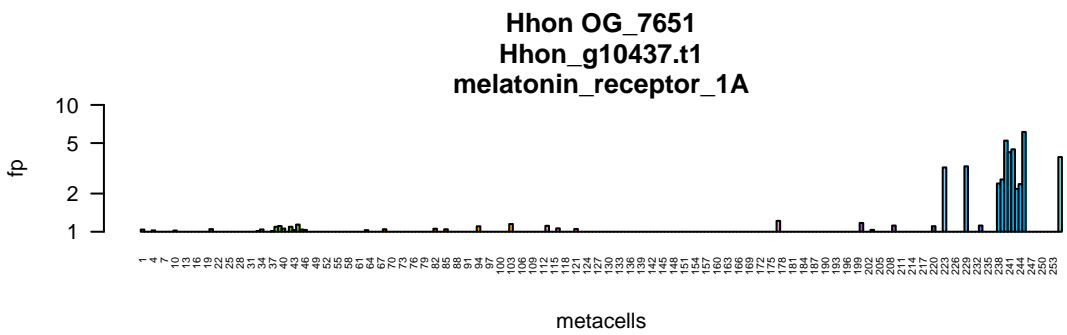
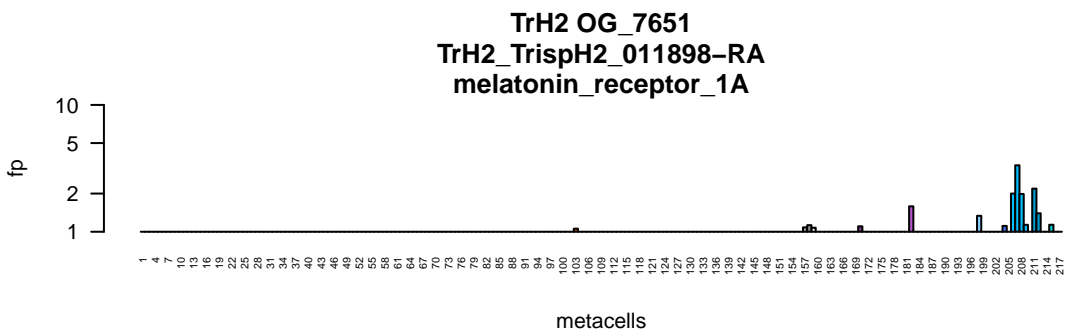
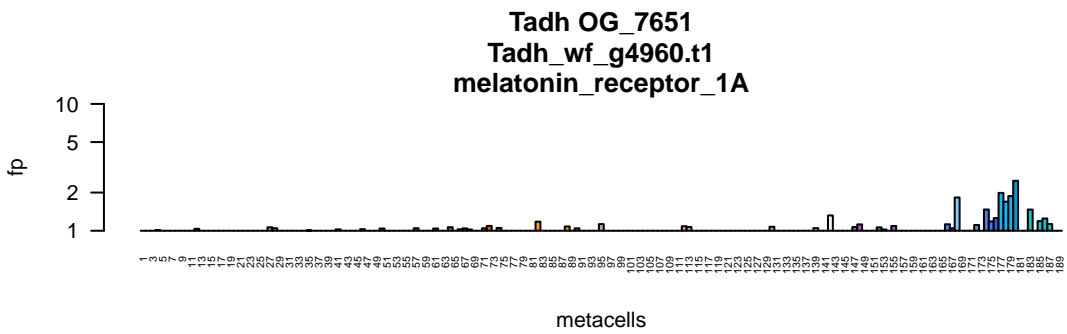
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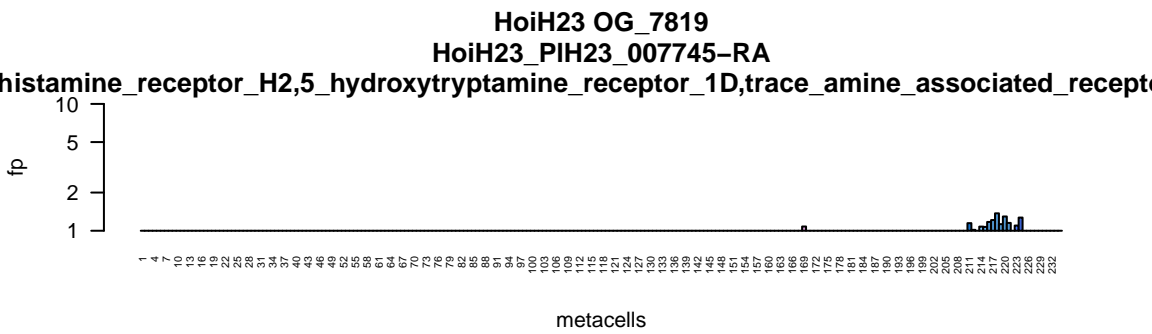
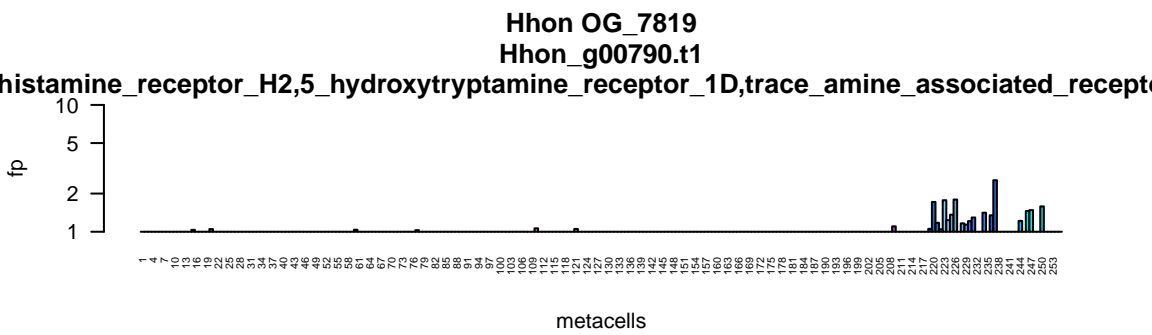
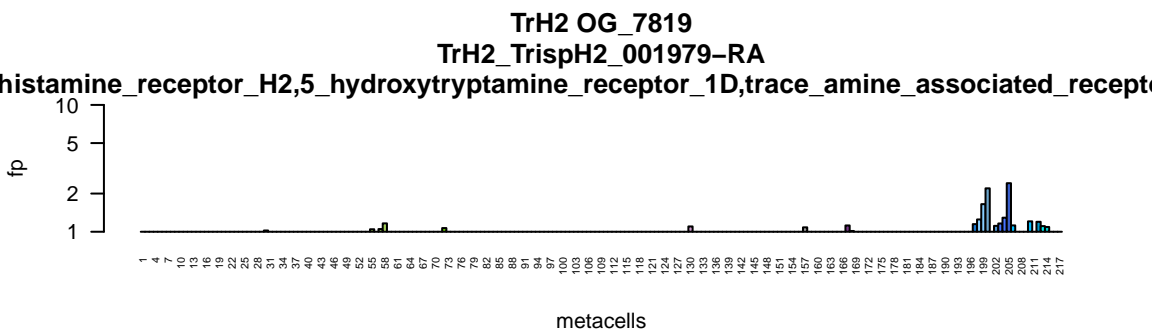
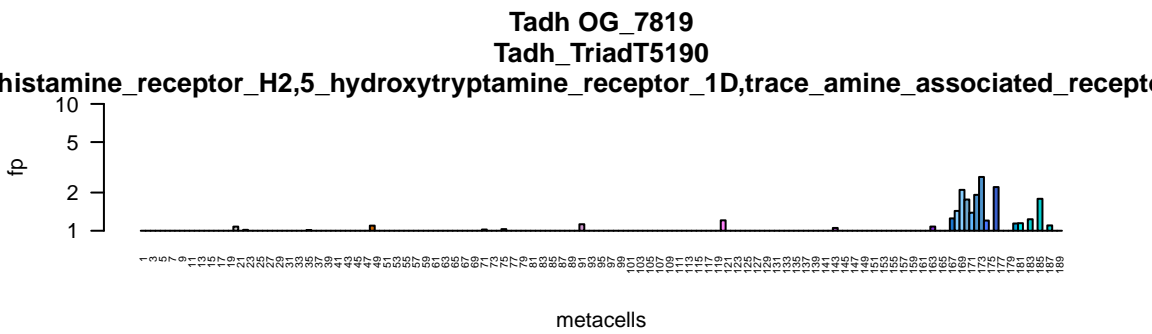


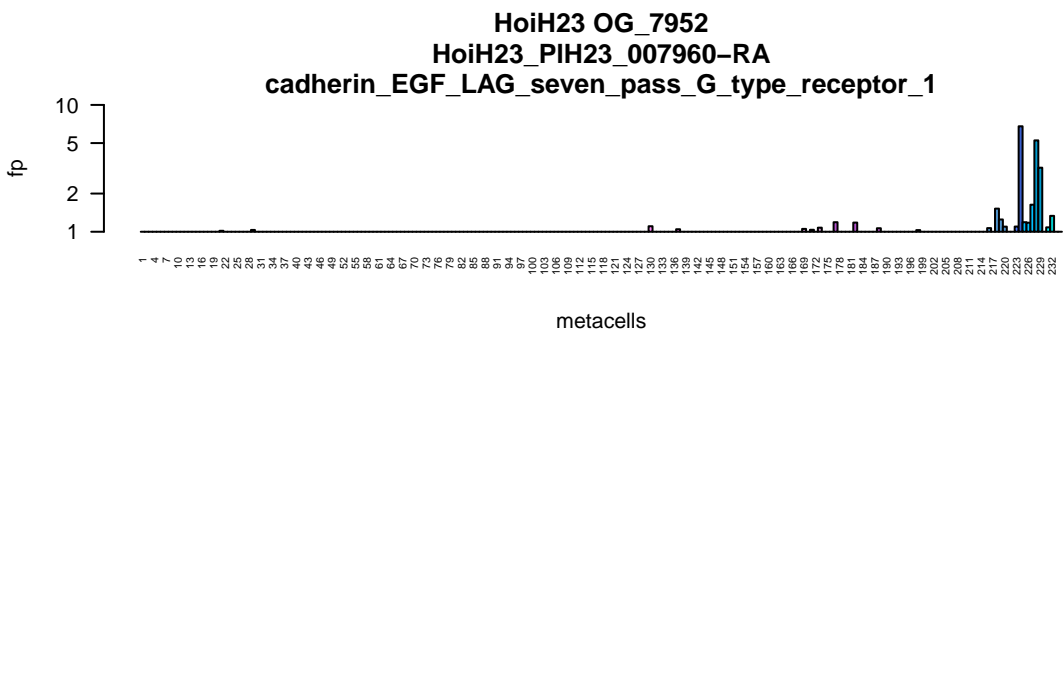
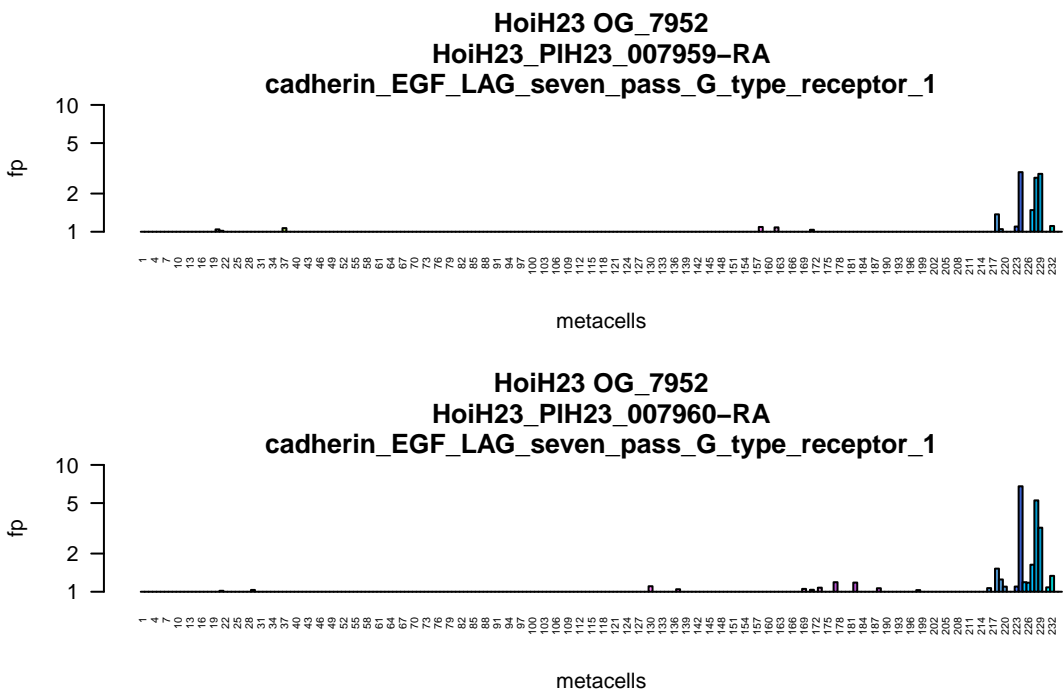
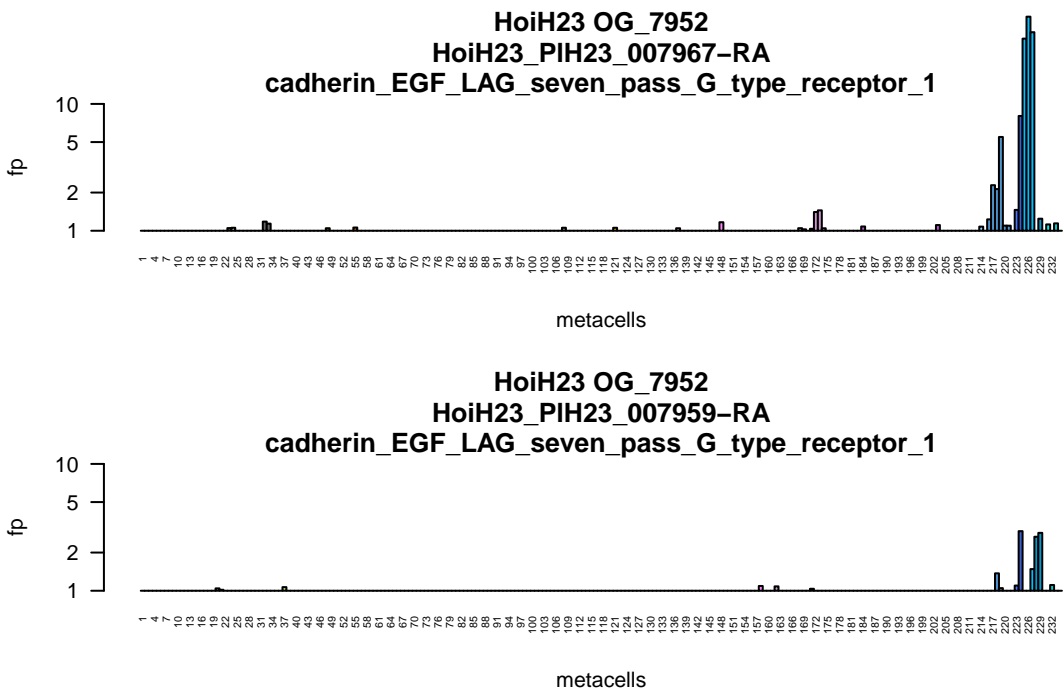
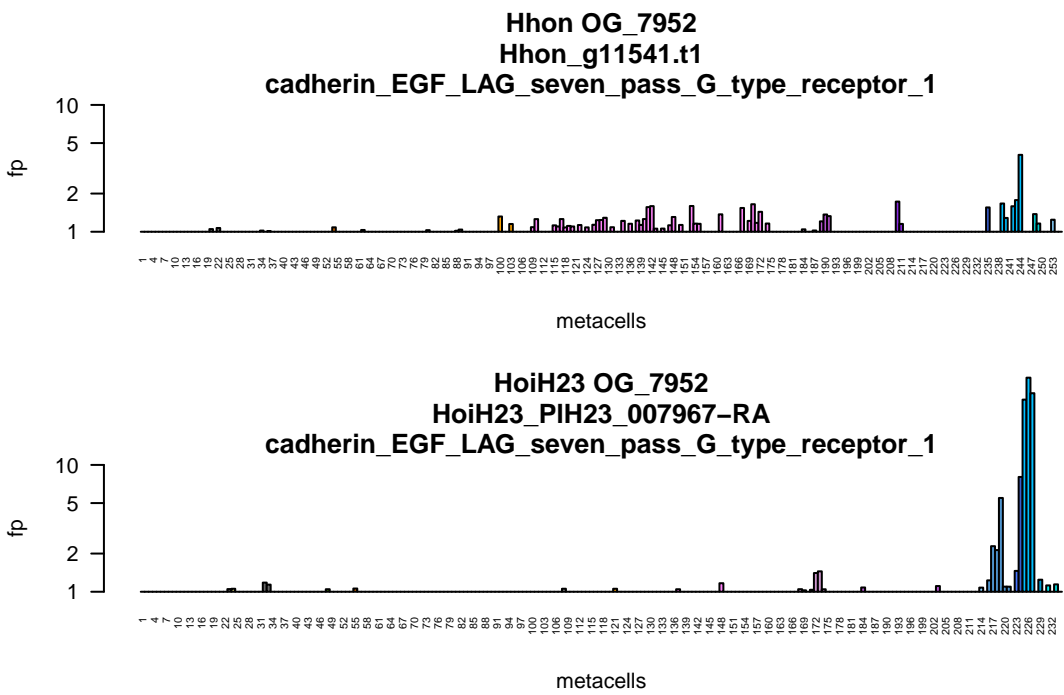
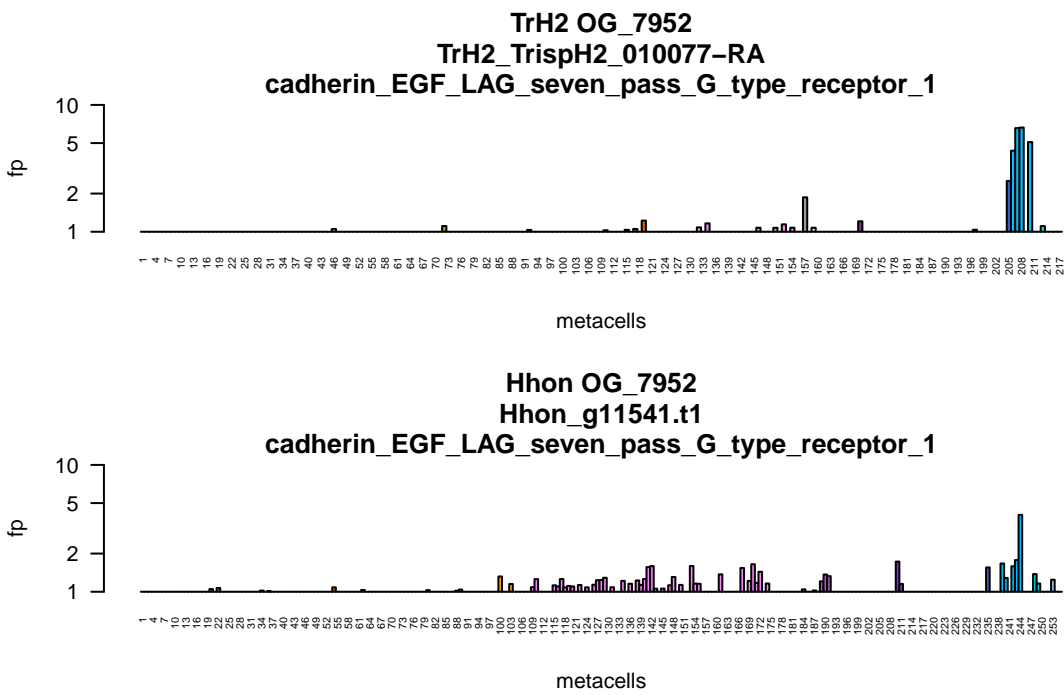
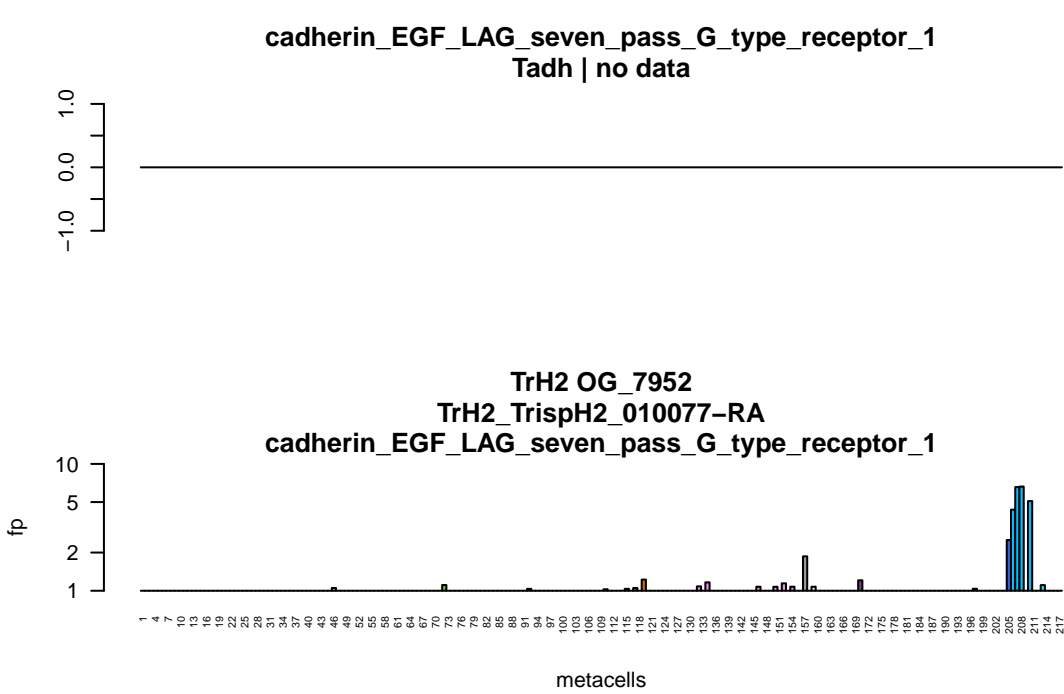


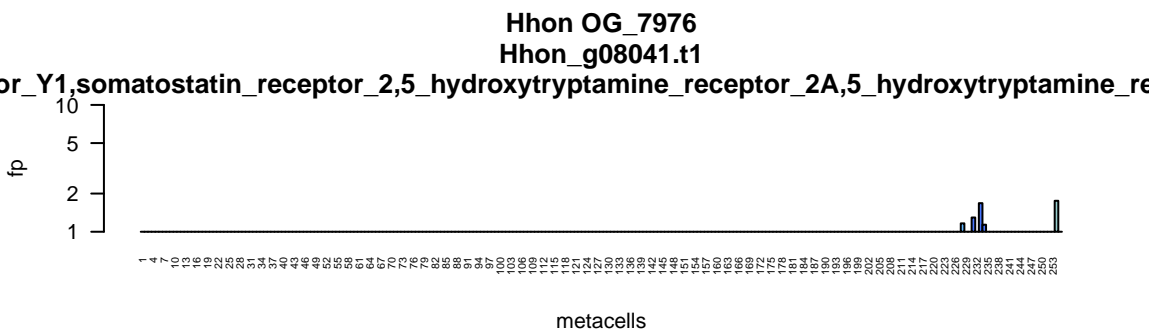
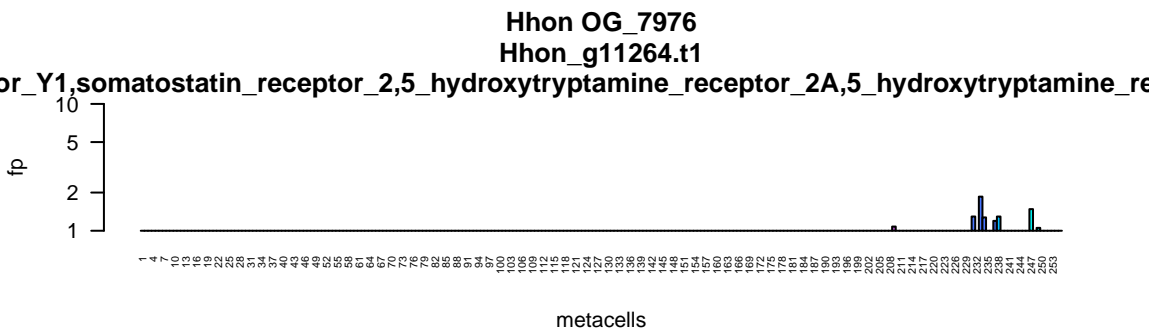
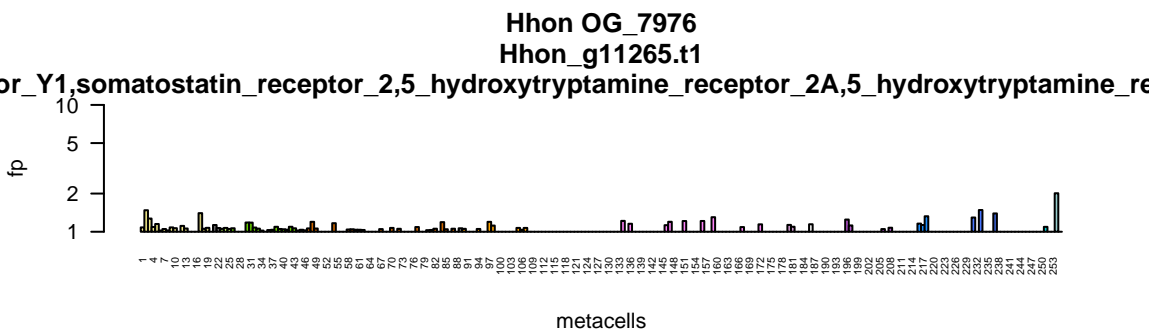
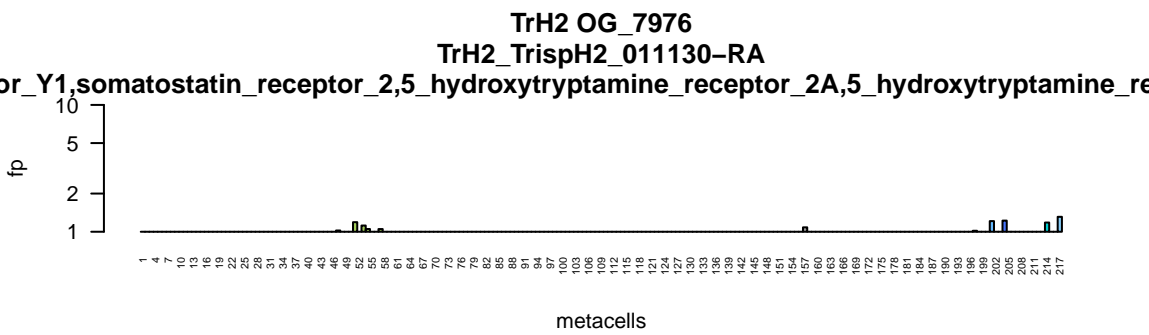
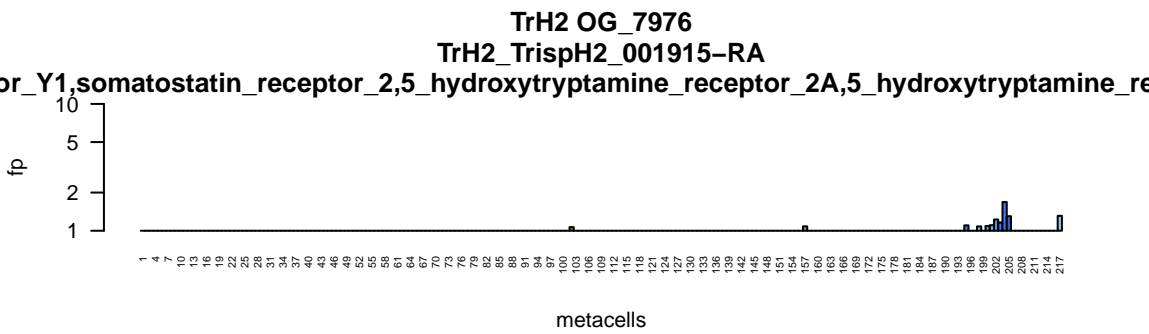
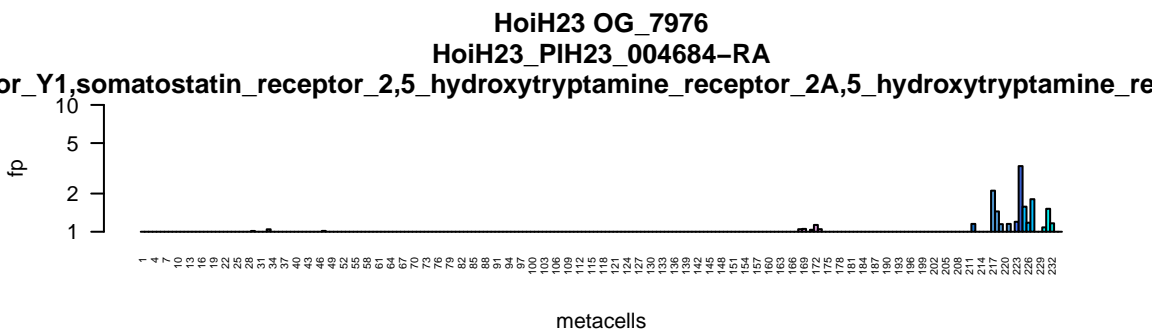
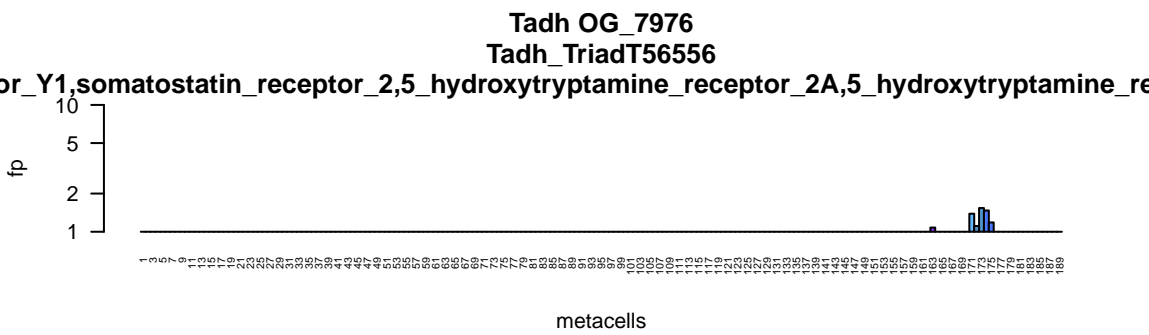
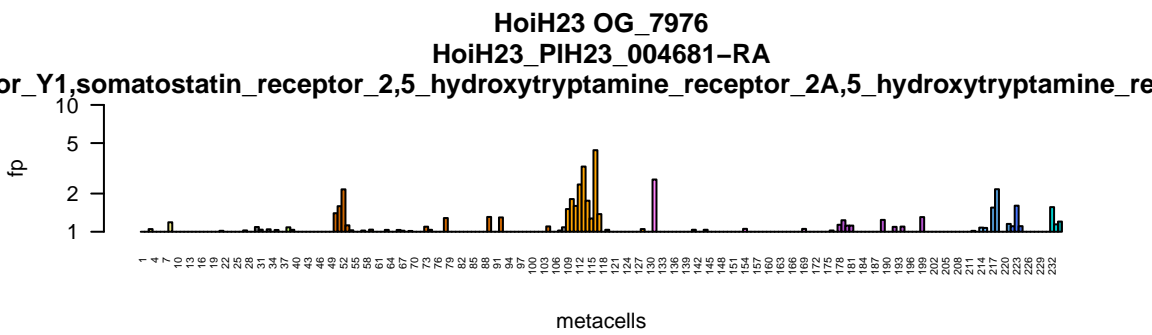
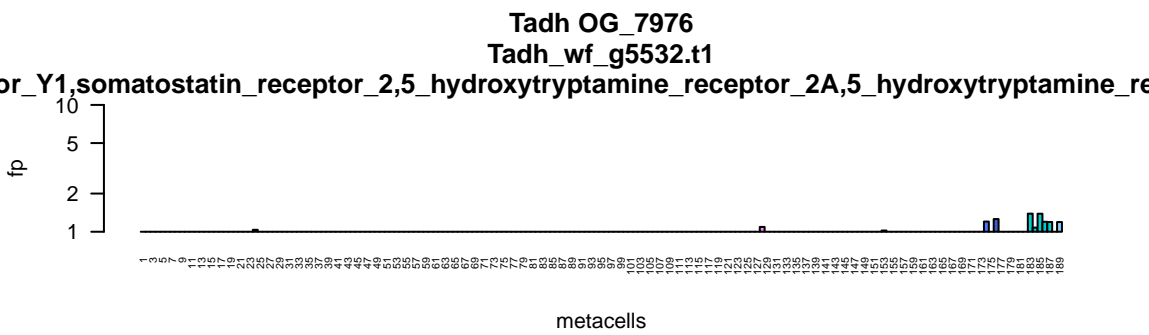
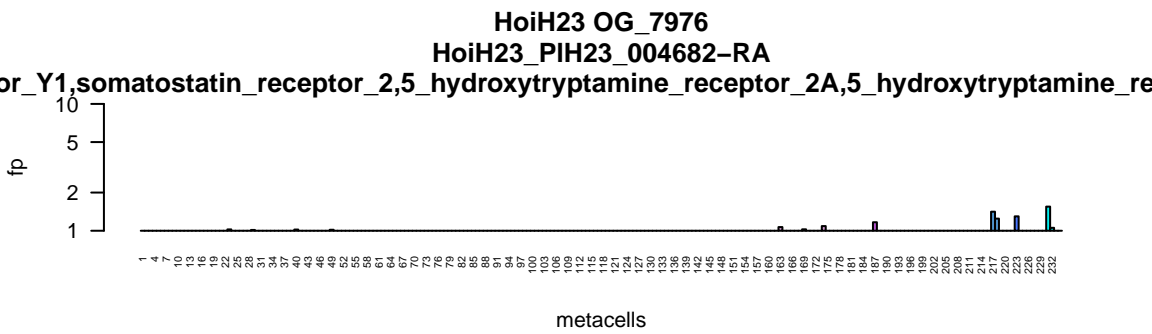
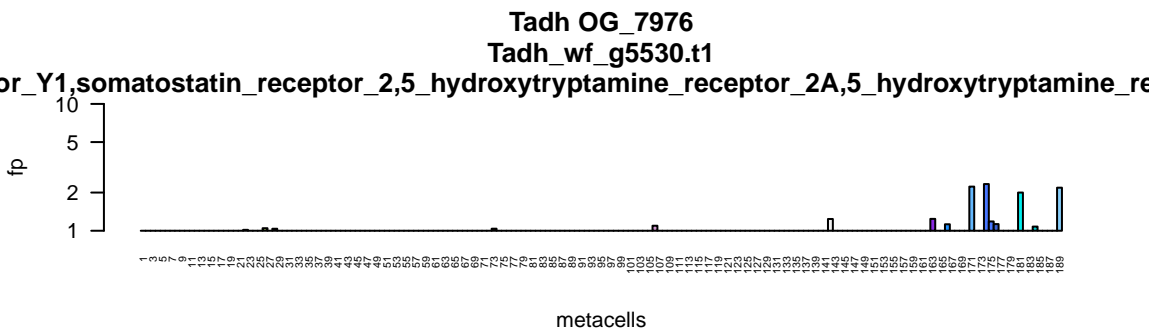


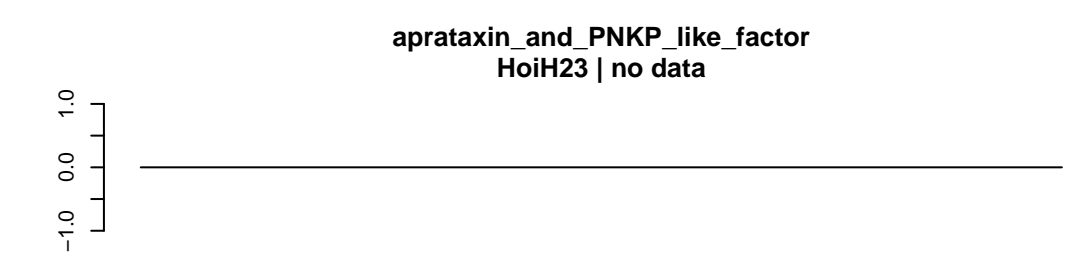
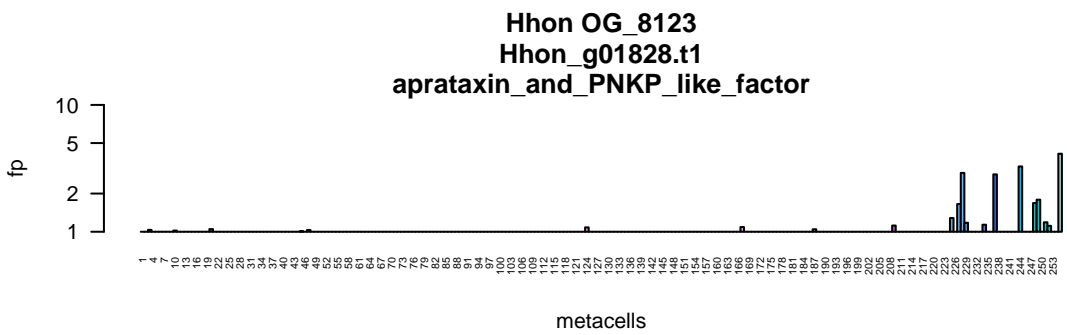
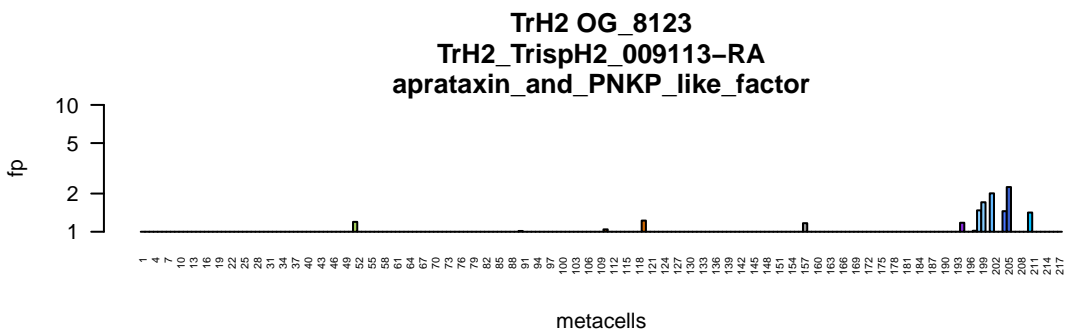
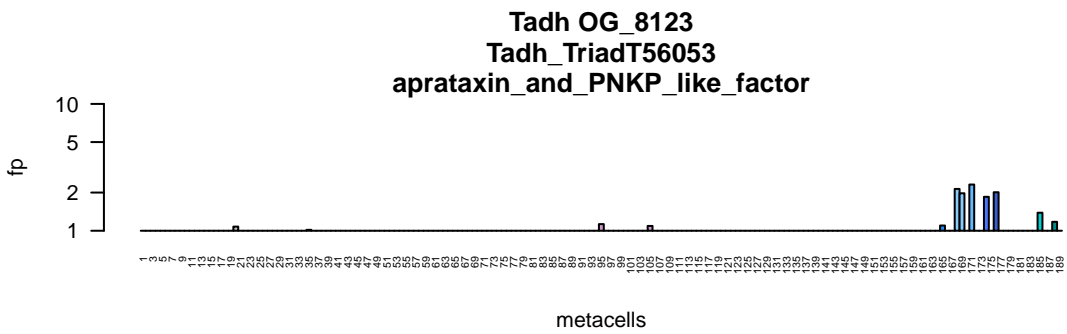


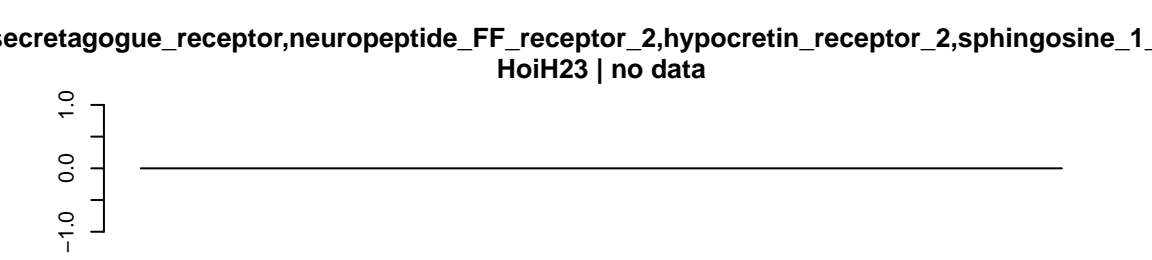
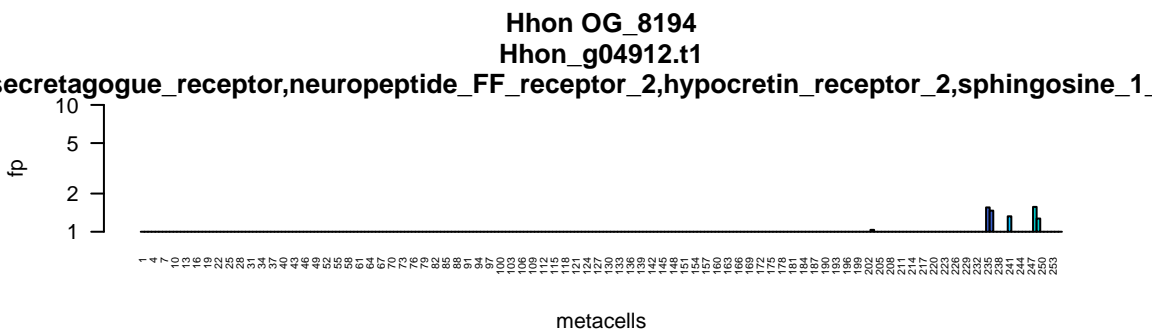
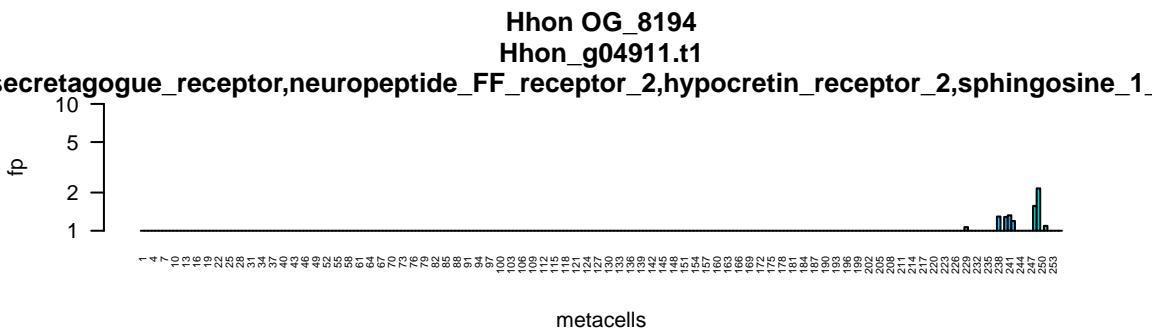
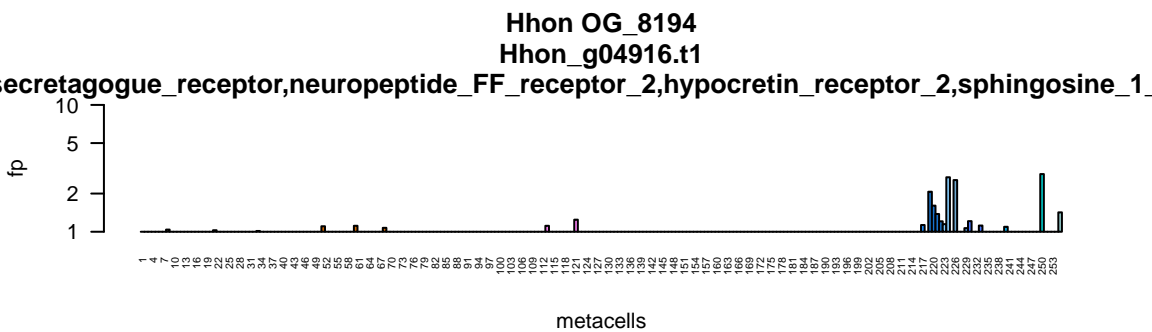
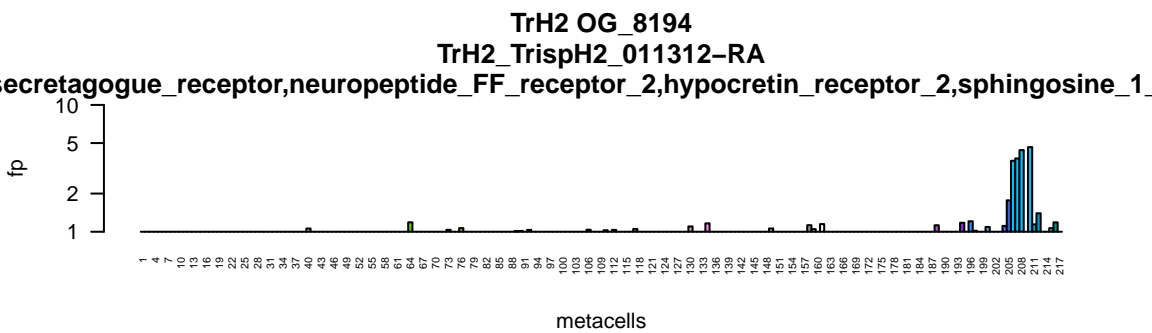
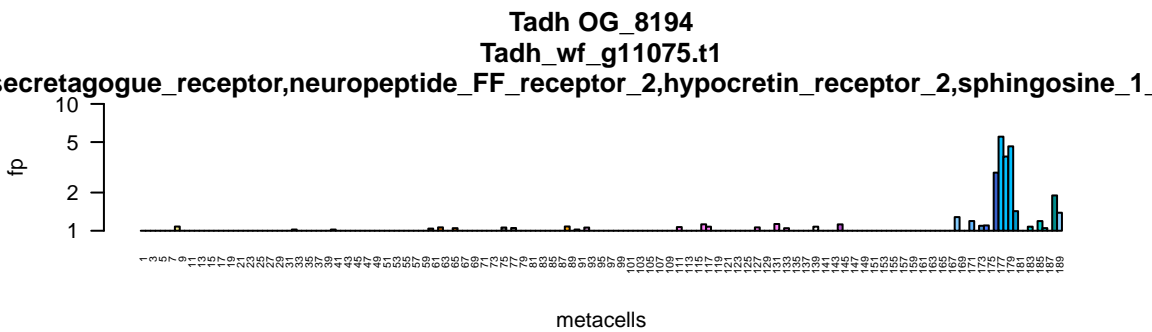


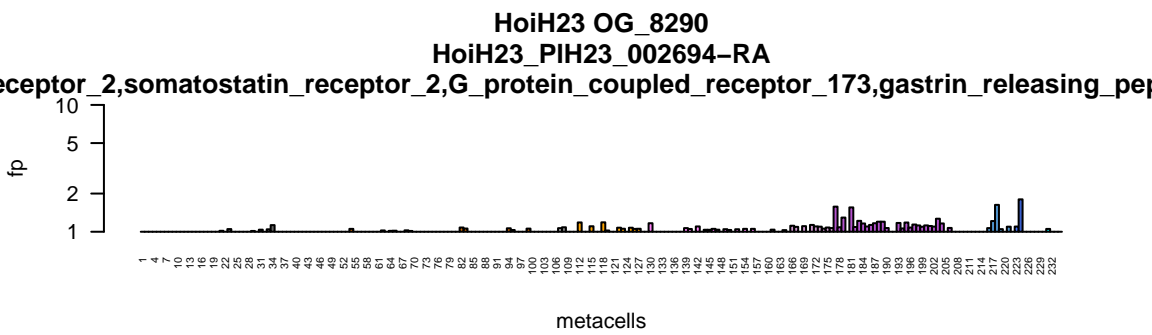
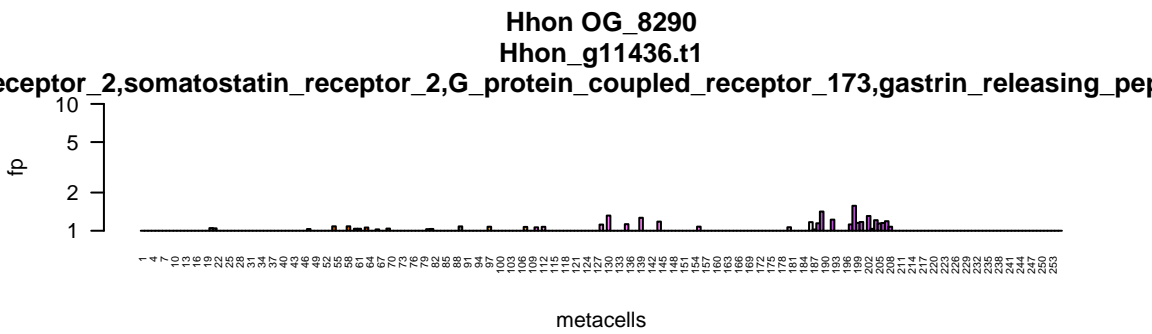
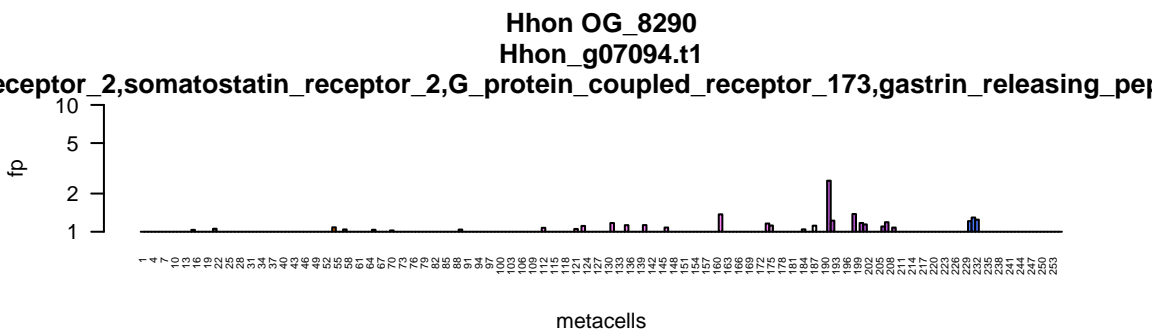
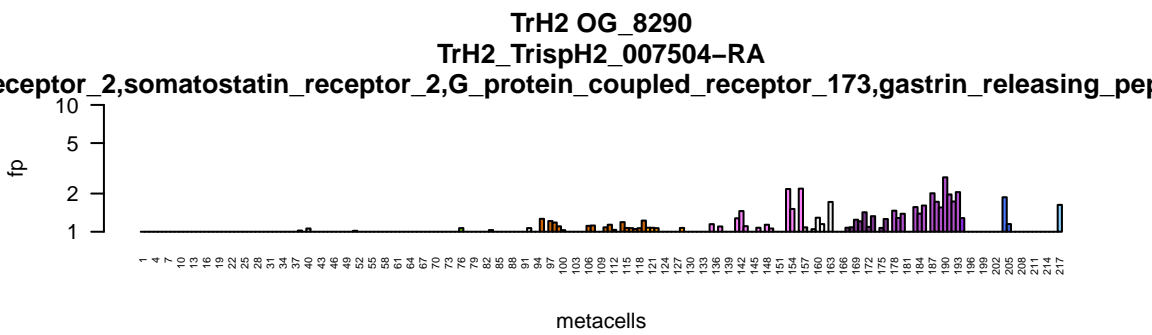
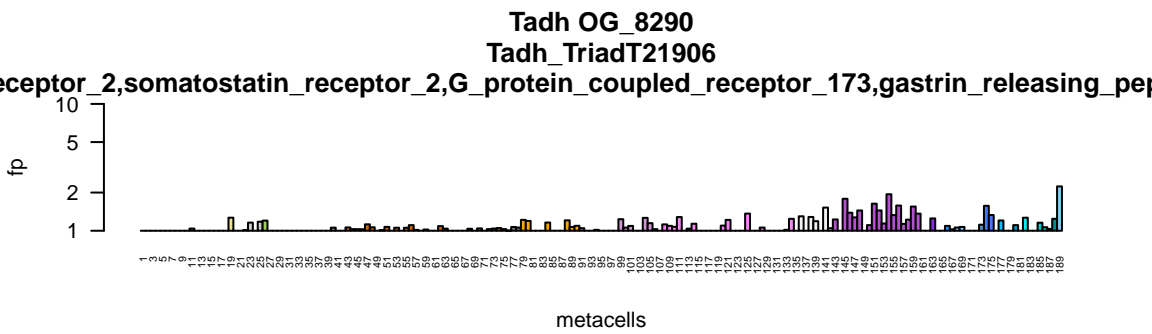








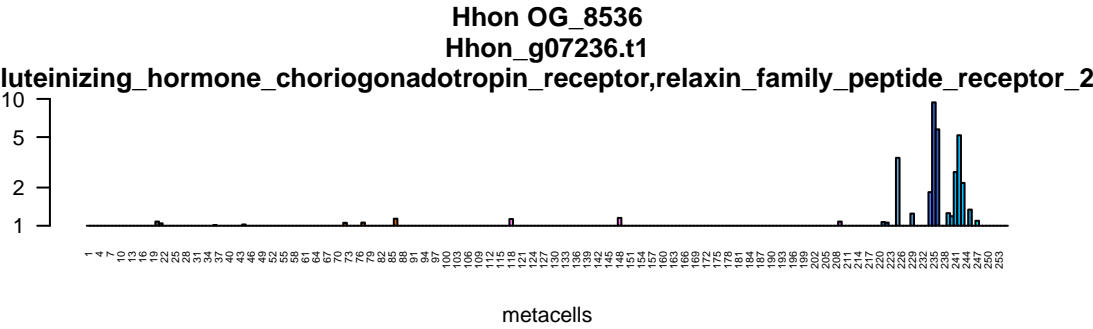




luteinizing_hormone_choriogonadotropin_receptor,relaxin_family_peptide_receptor_2
Tadh | no data

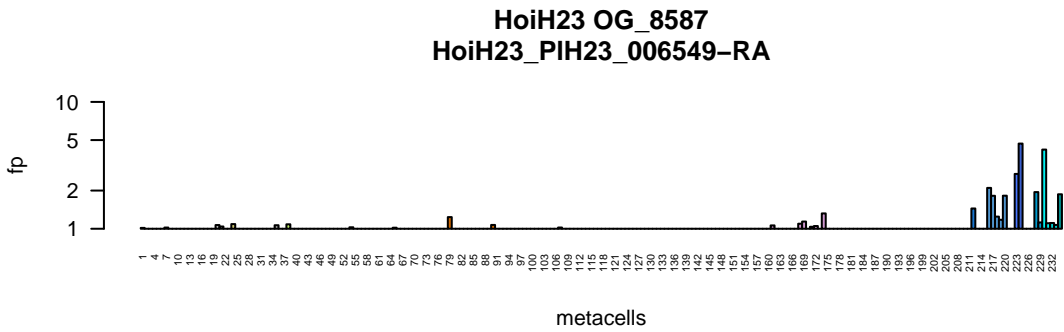
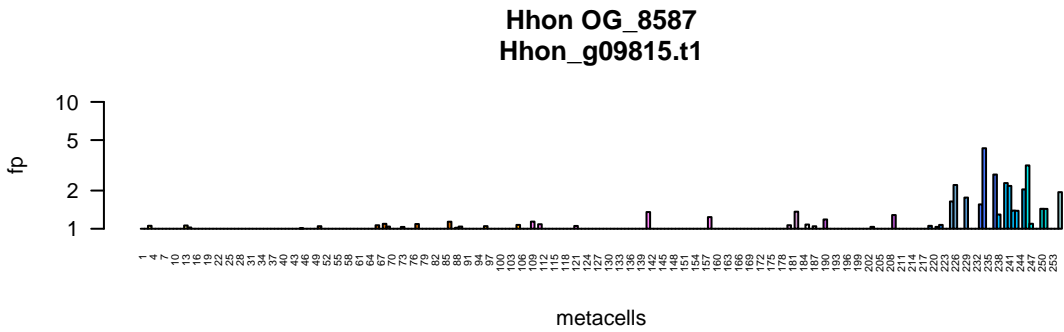
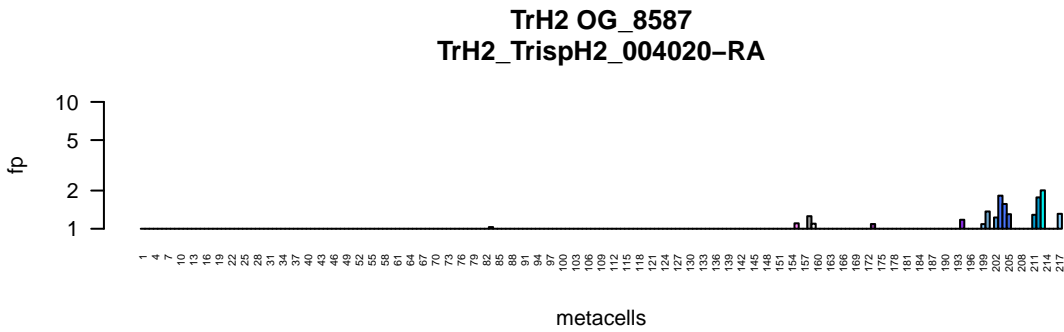
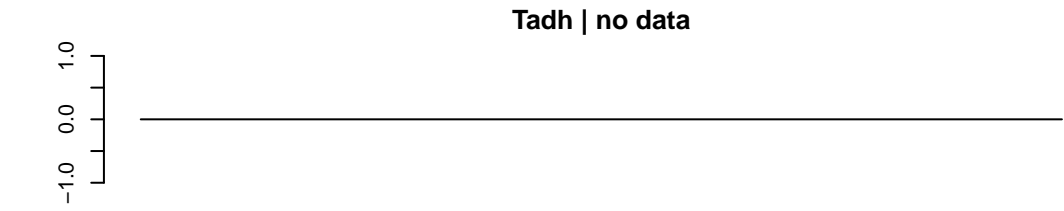


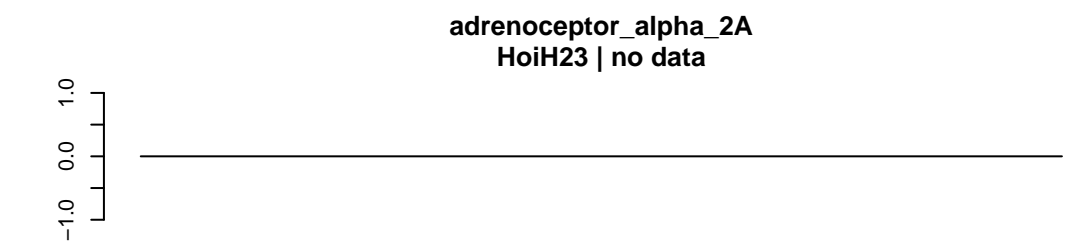
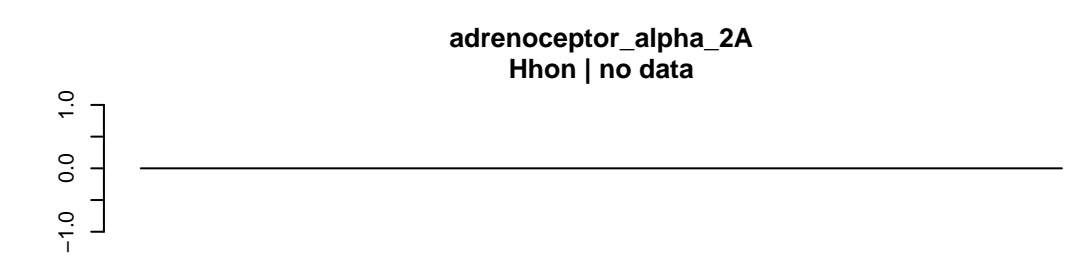
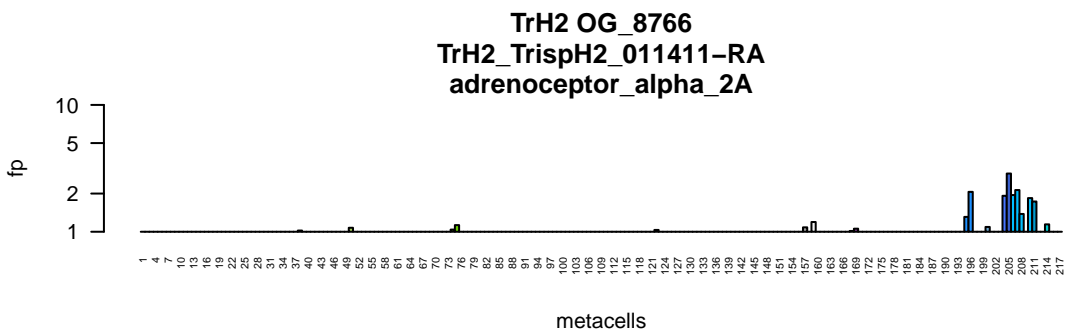
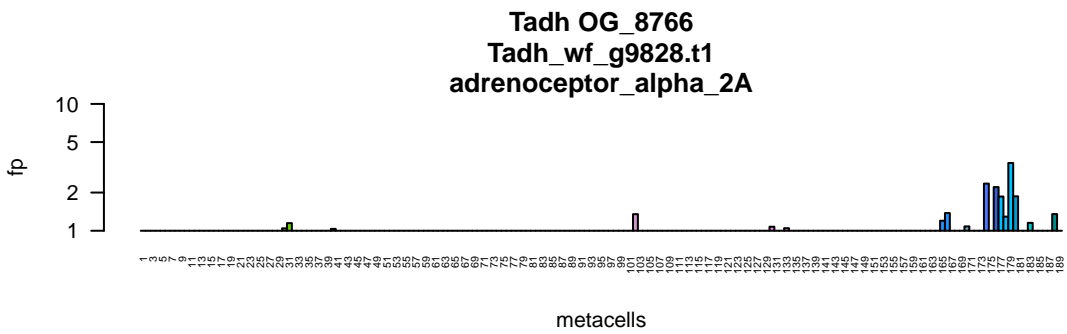
luteinizing_hormone_choriogonadotropin_receptor,relaxin_family_peptide_receptor_2
TrH2 | no data



luteinizing_hormone_choriogonadotropin_receptor,relaxin_family_peptide_receptor_2
HoiH23 | no data



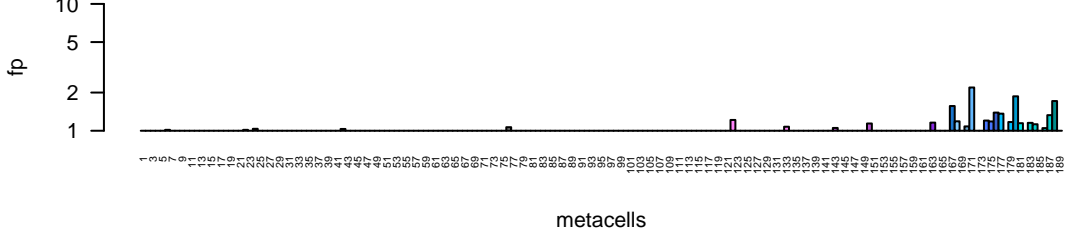




Tadh OG_9223

Tadh_TriadT51796

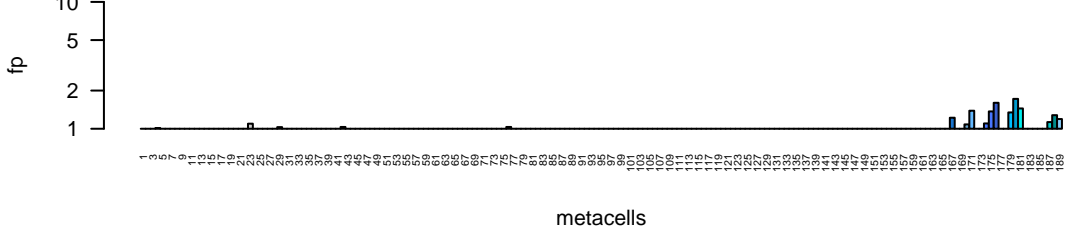
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Tadh OG_9223

Tadh_TriadT51797

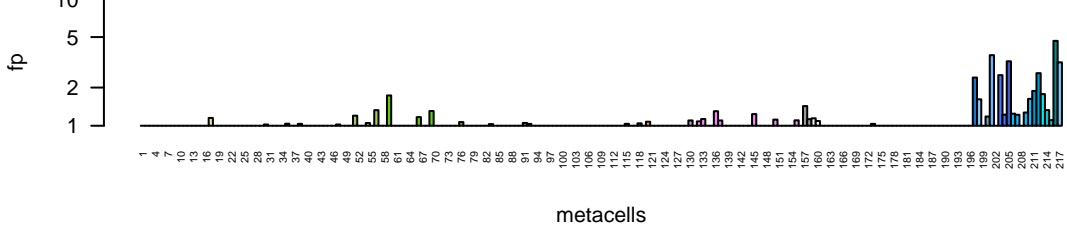
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



TrH2 OG_9223

TrH2_TrispH2_008387-RA

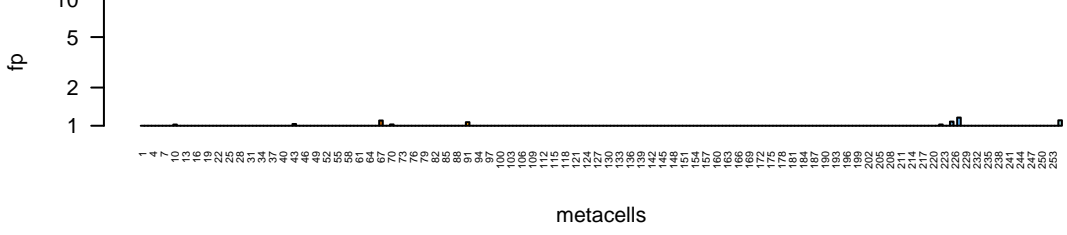
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Hhon OG_9223

Hhon_g11380.t1

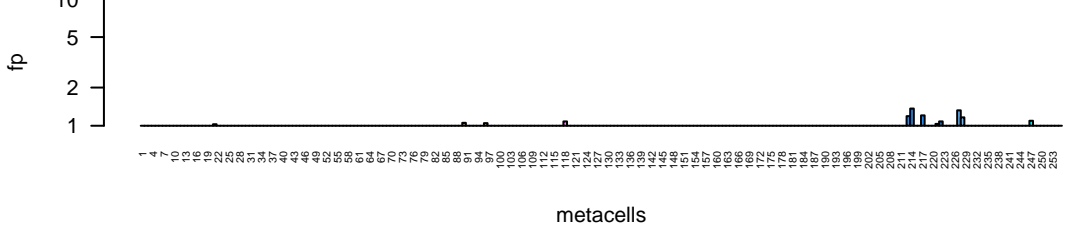
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Hhon OG_9223

Hhon_g01959.t1

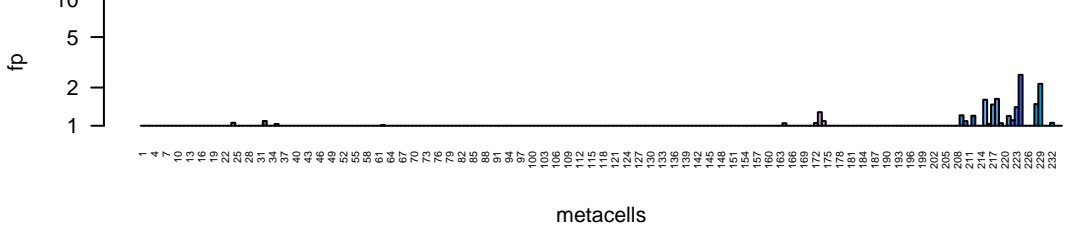
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto

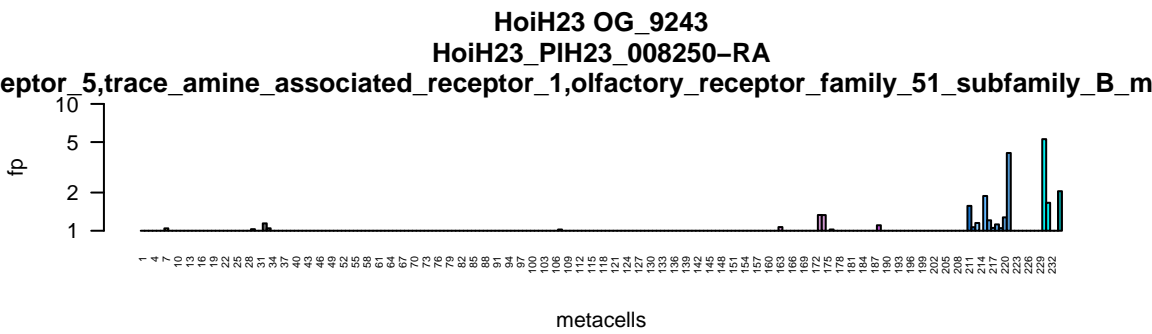
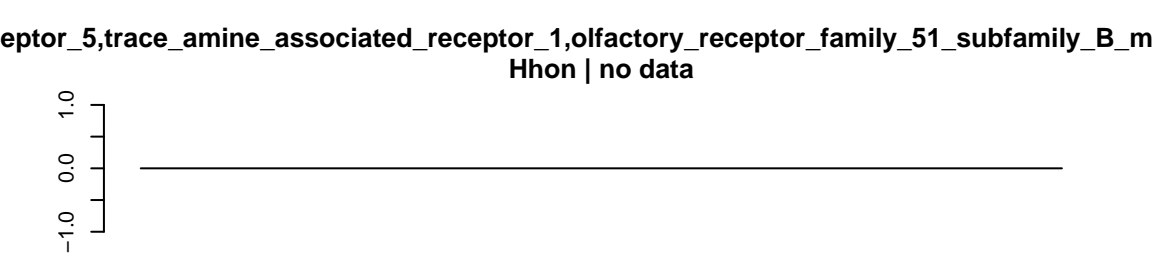
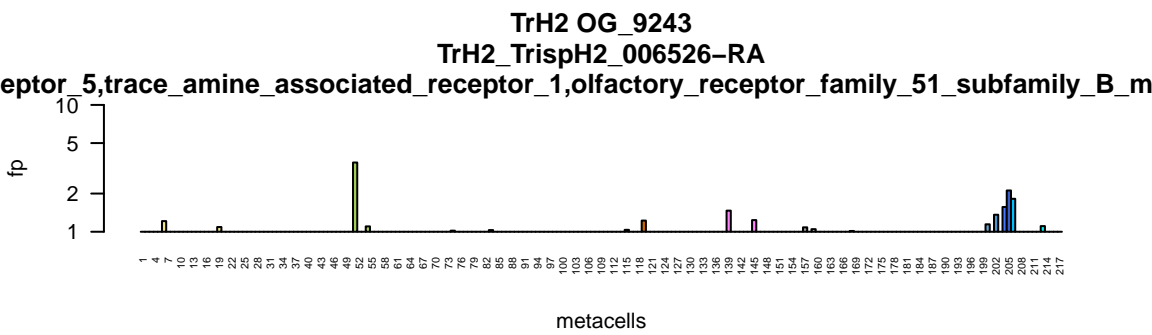
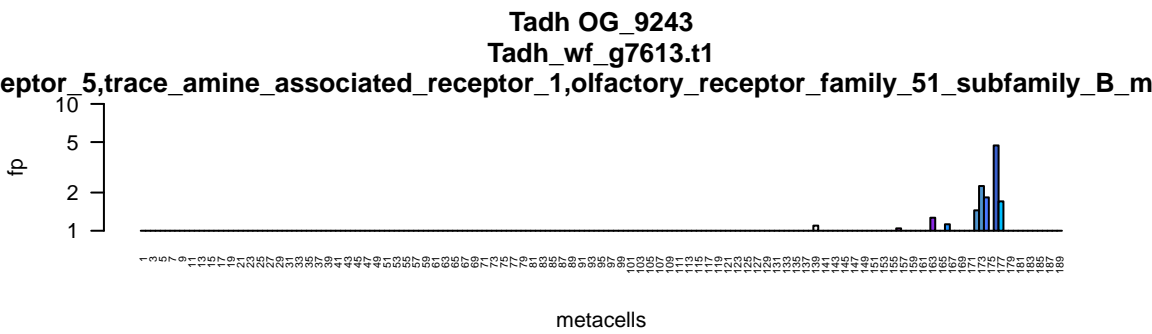


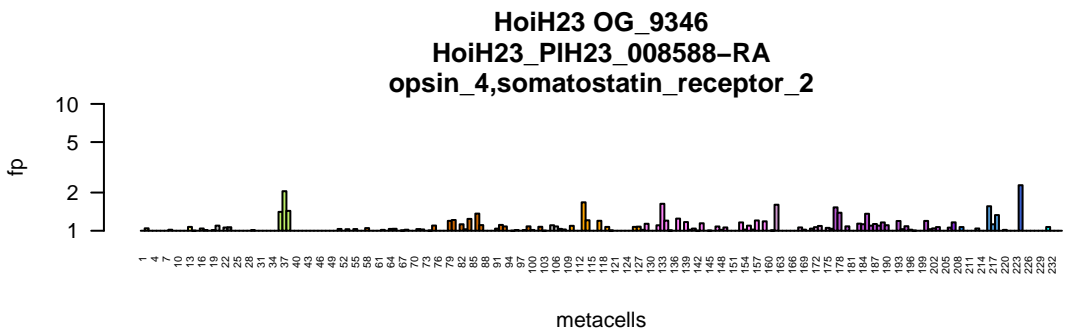
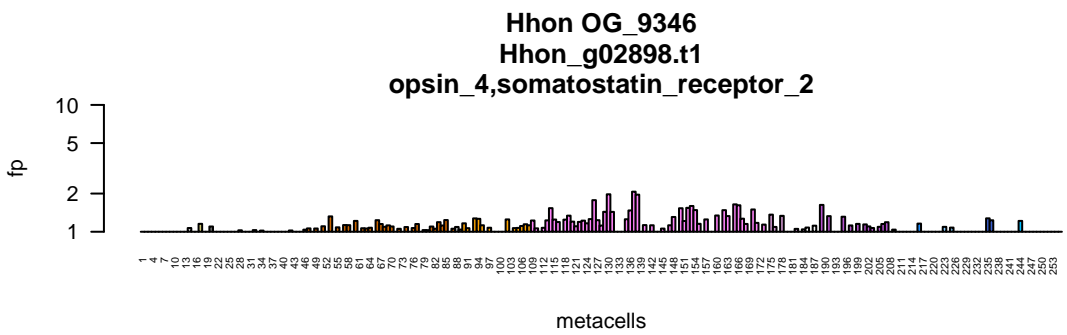
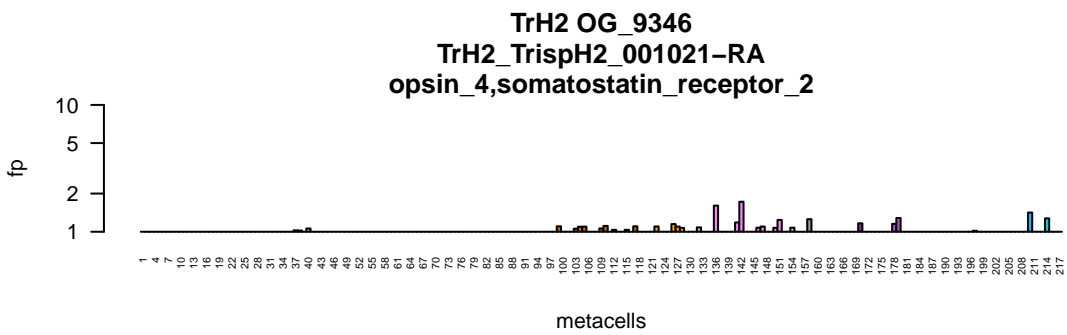
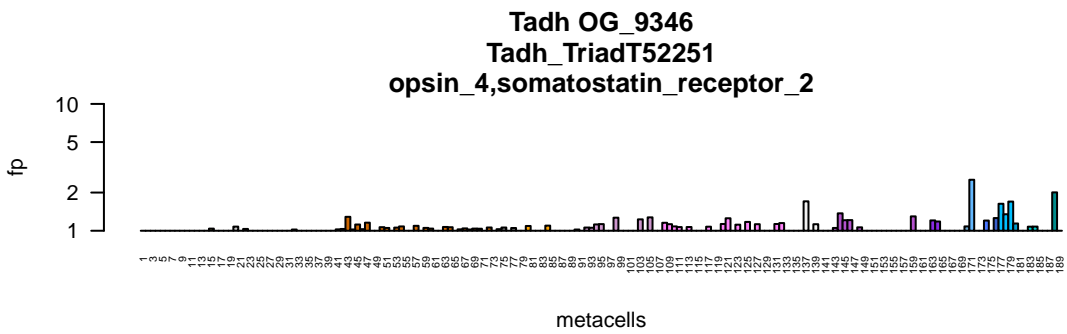
HoiH23 OG_9223

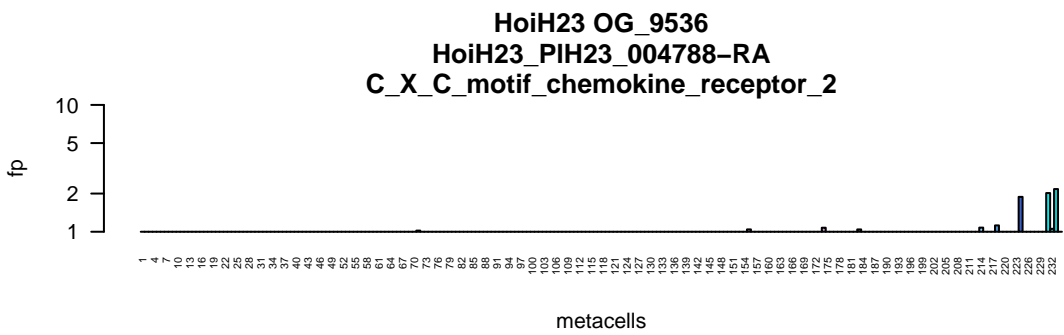
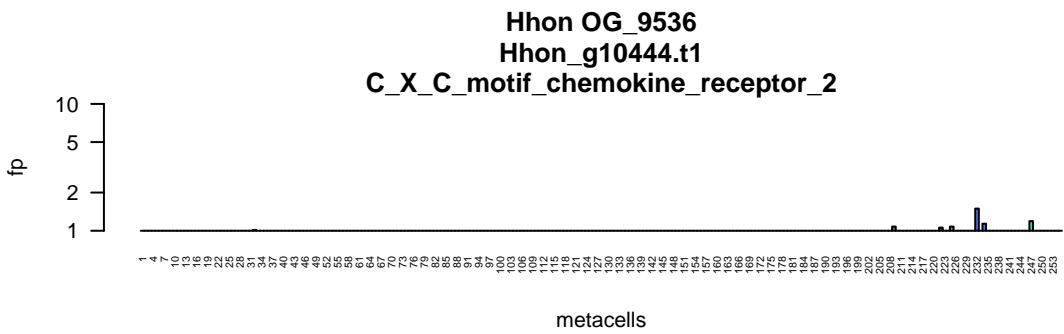
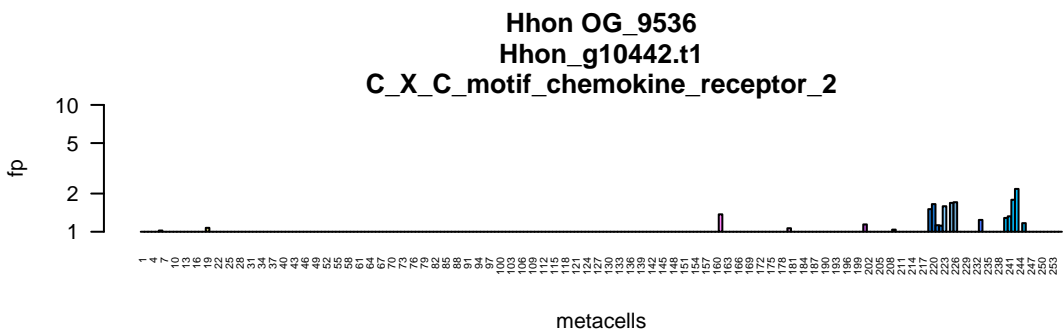
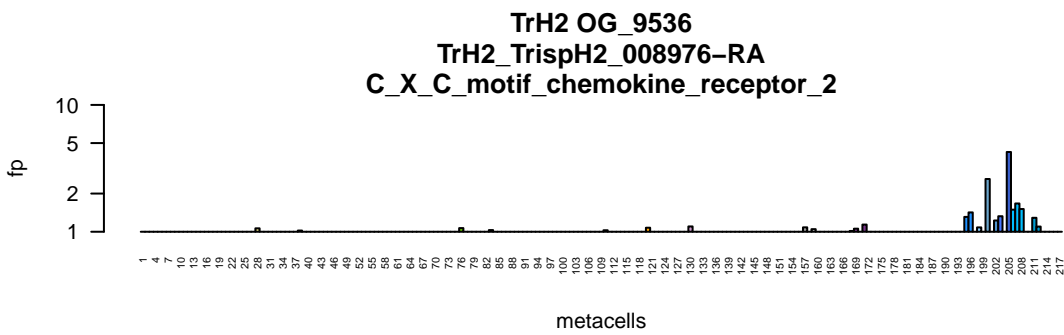
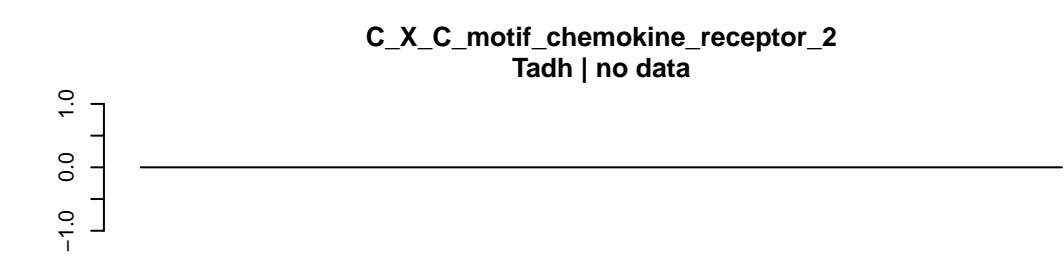
HoiH23_PIH23_004873-RA

relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto

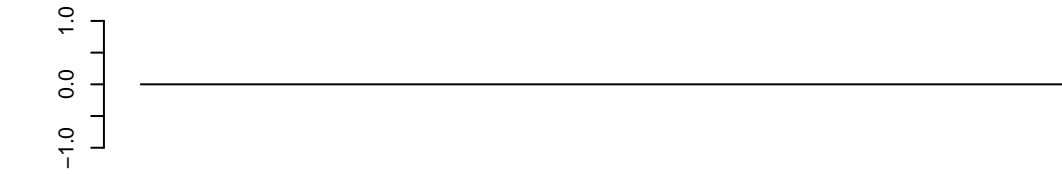




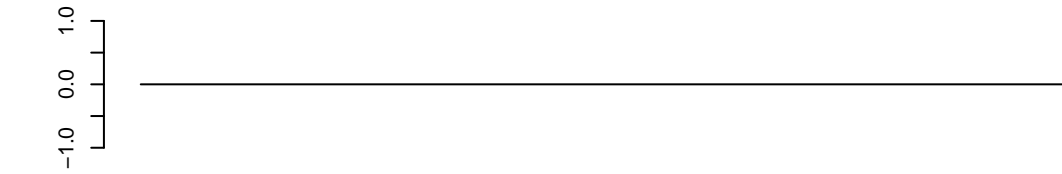




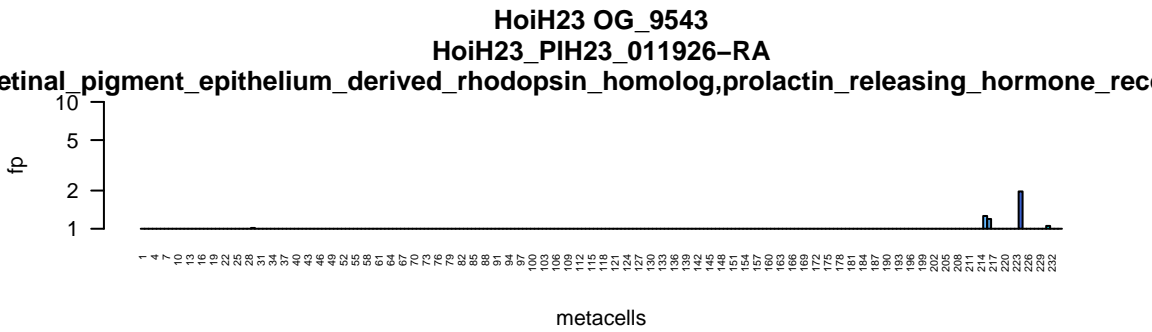
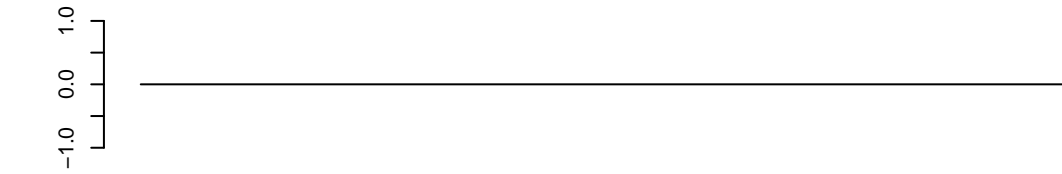
etinal_pigment_epithelium_derived_rhodopsin_homolog,prolactin_releasing_hormone_rec
Tadh | no data

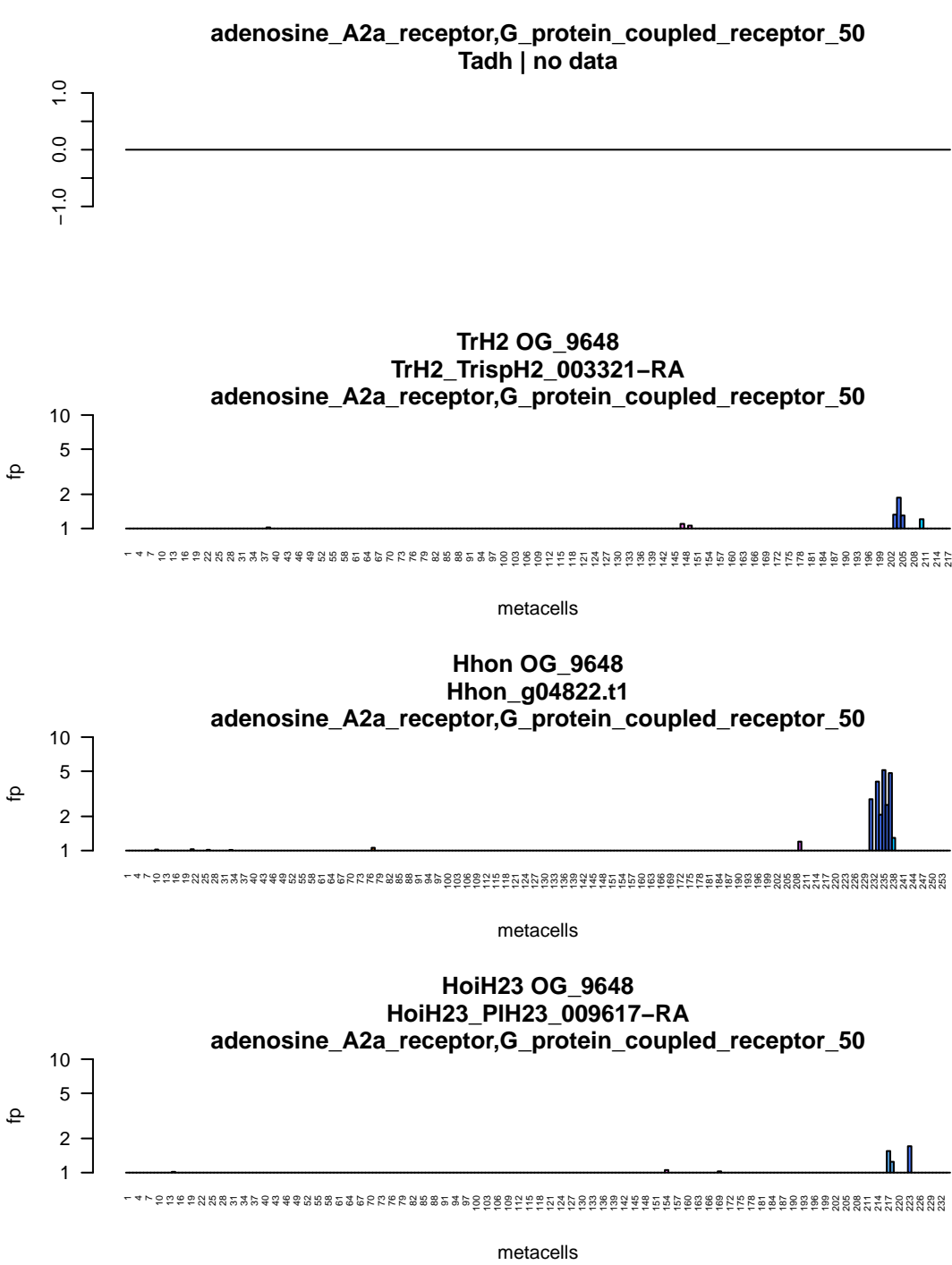


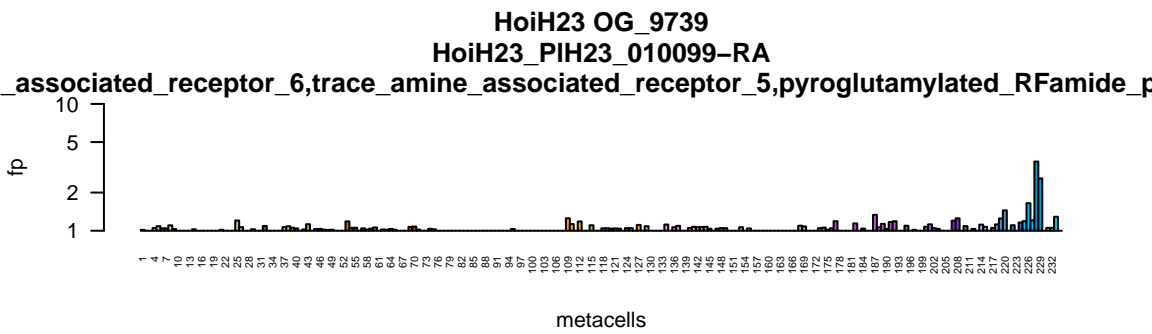
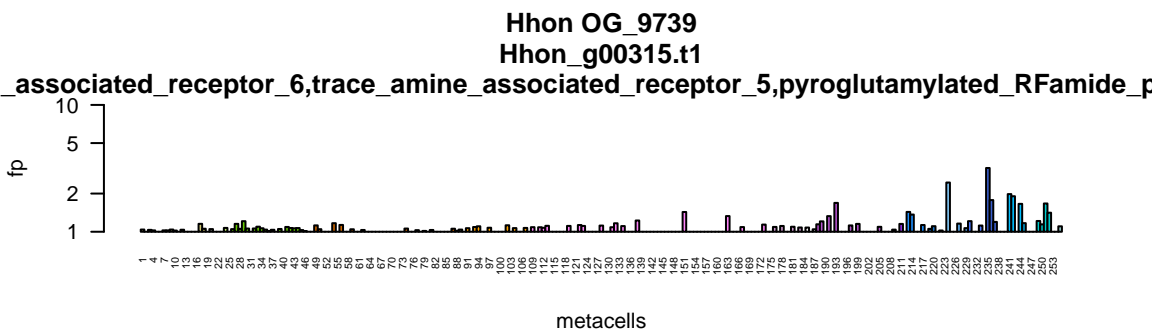
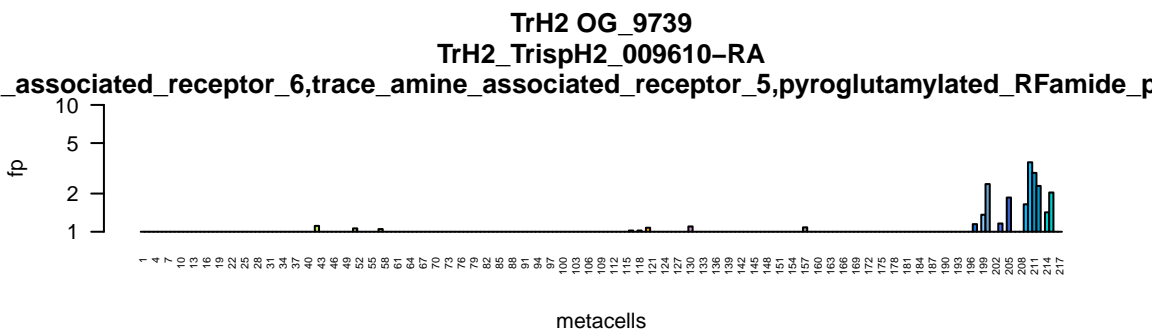
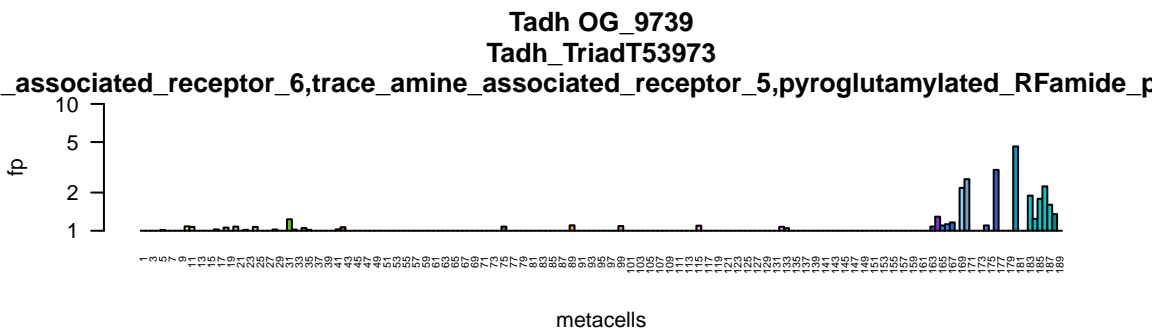
etinal_pigment_epithelium_derived_rhodopsin_homolog,prolactin_releasing_hormone_rec
TrH2 | no data



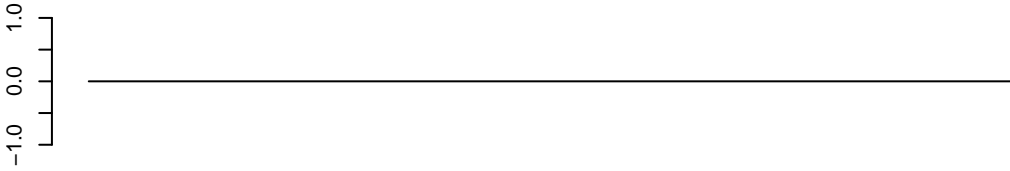
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Hhon | no data



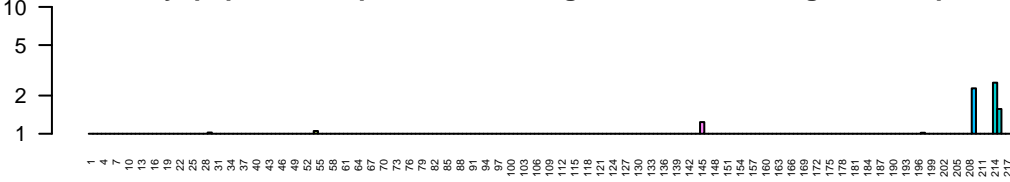




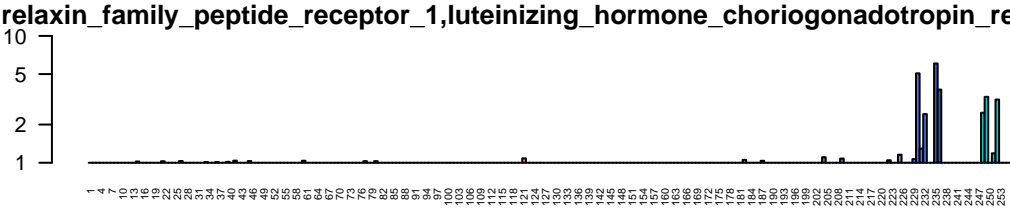
relaxin_family_peptide_receptor_1,luteinizing_hormone_choriogonadotropin_receptor
Tadh | no data



TrH2 OG_9876
TrH2_TrispH2_009947-RA
relaxin_family_peptide_receptor_1,luteinizing_hormone_choriogonadotropin_receptor



Hhon OG_9876
Hhon_g04760.t1
relaxin_family_peptide_receptor_1,luteinizing_hormone_choriogonadotropin_receptor



HoiH23 OG_9876
HoiH23_PIH23_006596-RA
relaxin_family_peptide_receptor_1,luteinizing_hormone_choriogonadotropin_receptor

