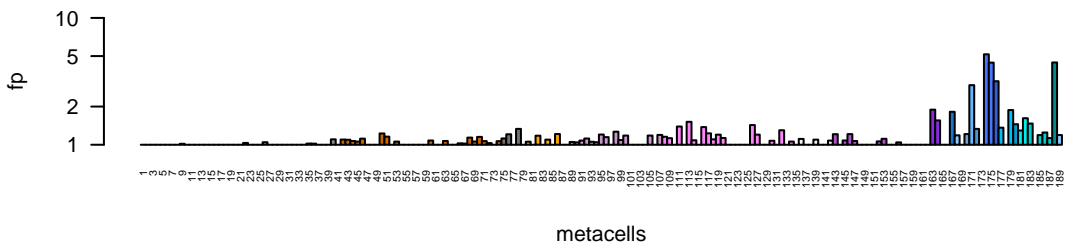
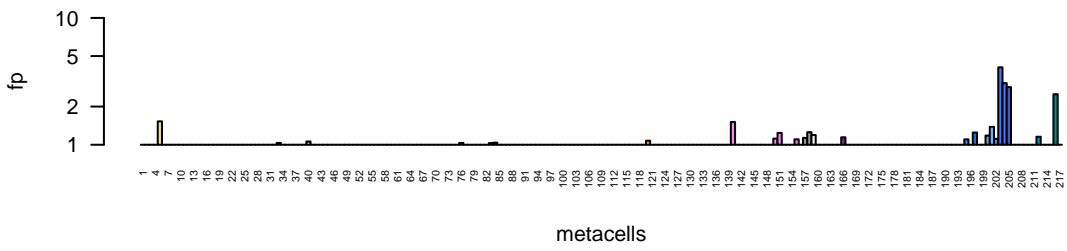


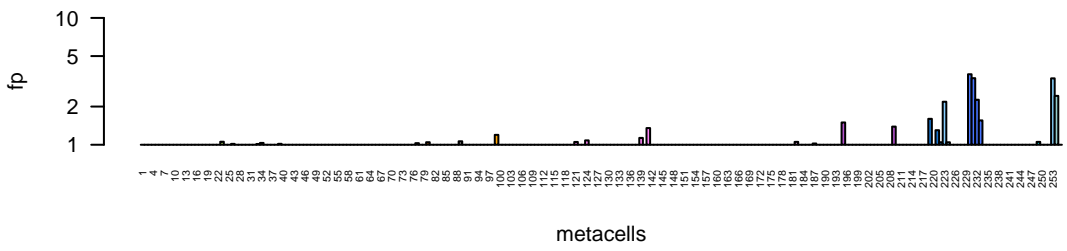
Tadh OG_4774
Tadh_wf_g8472.t1



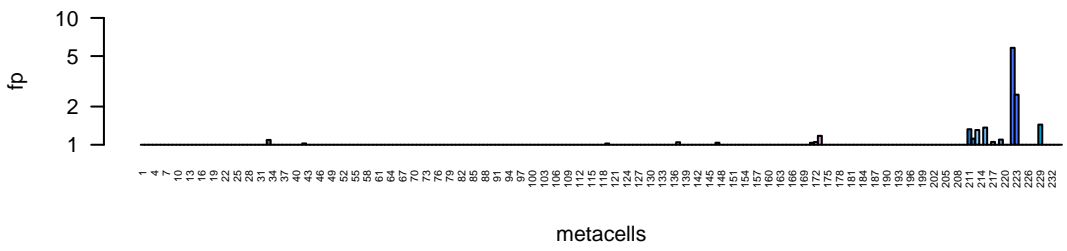
TrH2 OG_4774
TrH2_TrispH2_010103-RA



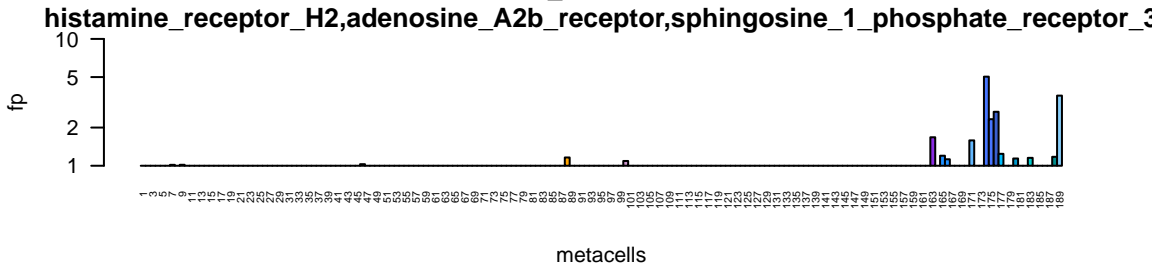
Hhon OG_4774
Hhon_g10392.t1



HoiH23 OG_4774
HoiH23_PIH23_006299-RA



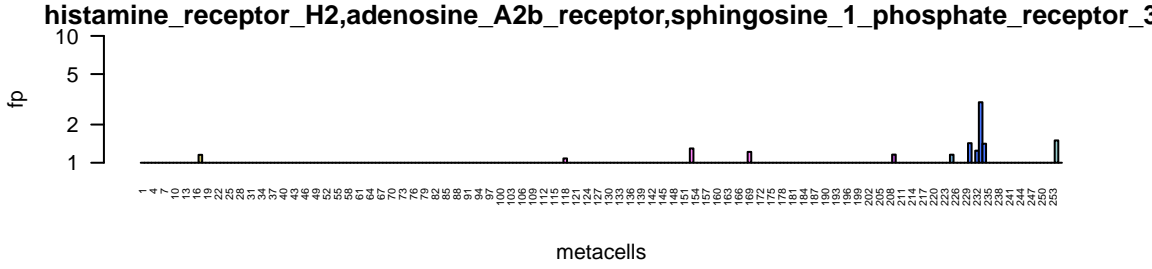
Tadh OG_4983
Tadh_TriadT58722



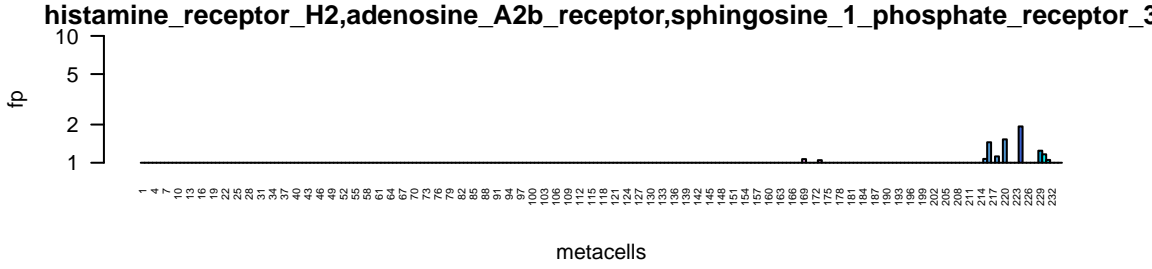
TrH2 OG_4983
TrH2_TrispH2_006523-RA



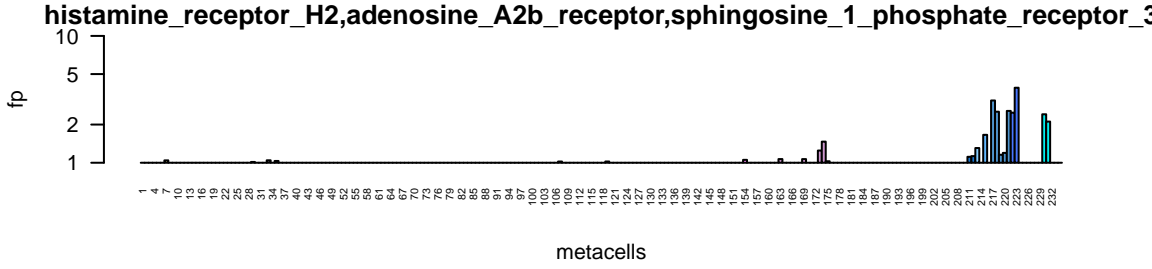
Hhon OG_4983
Hhon_g01339.t1

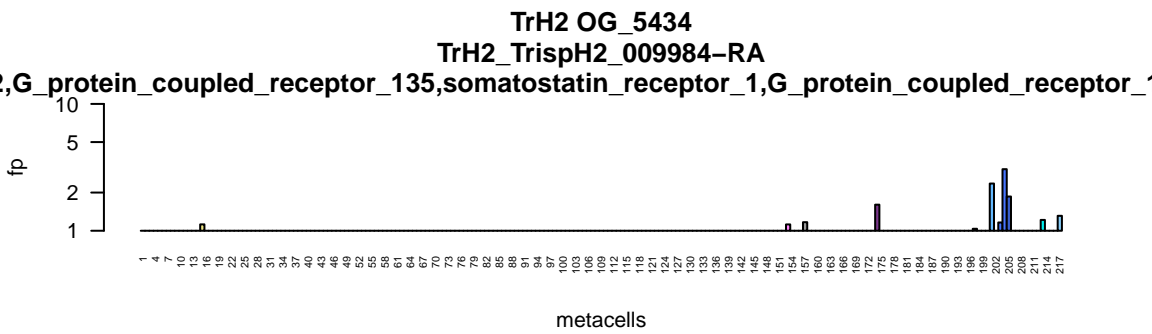
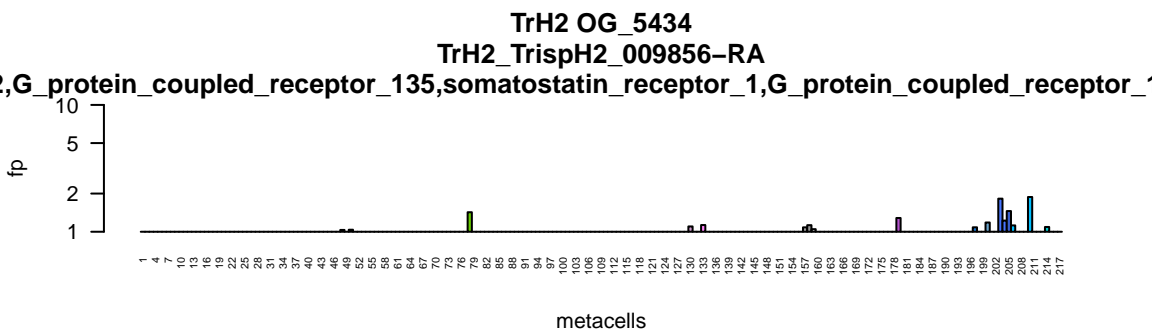
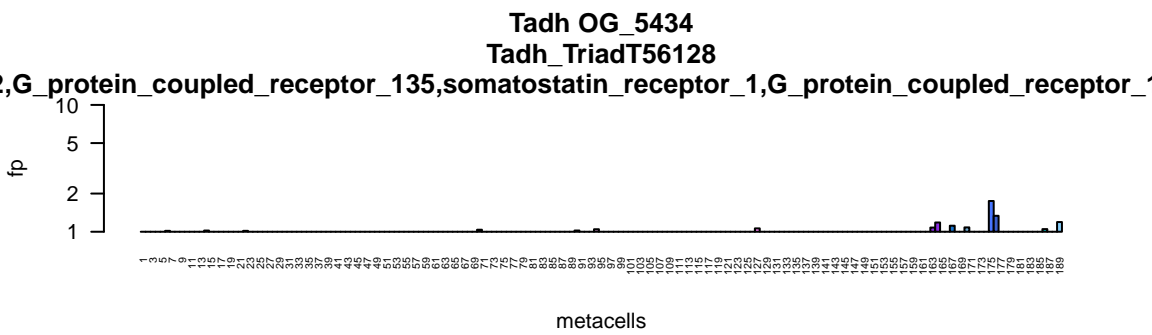
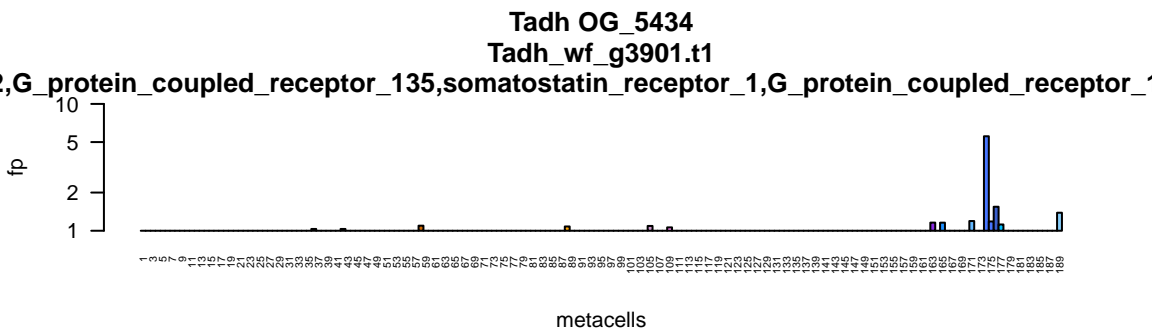


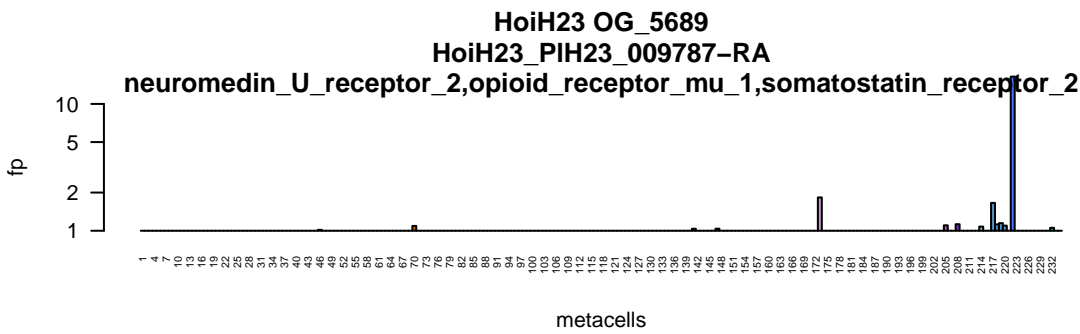
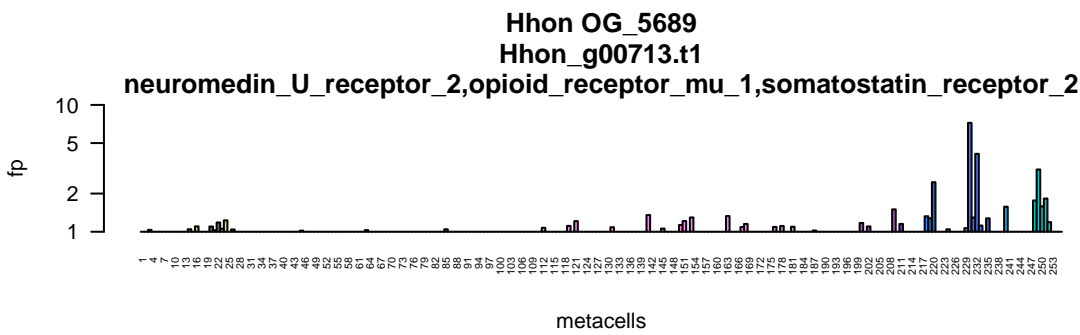
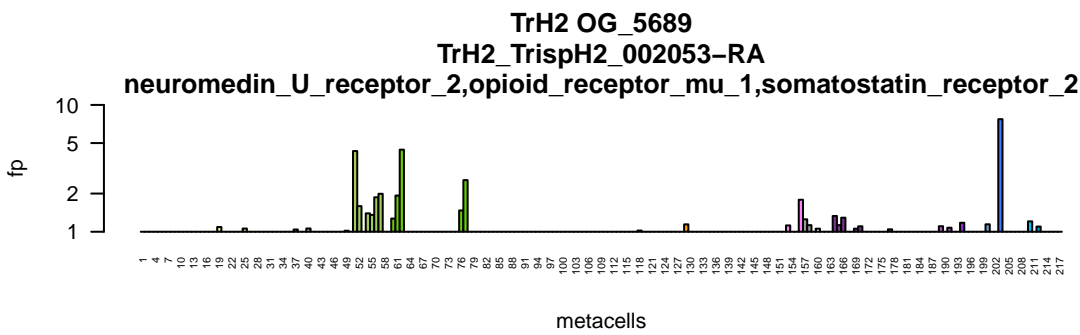
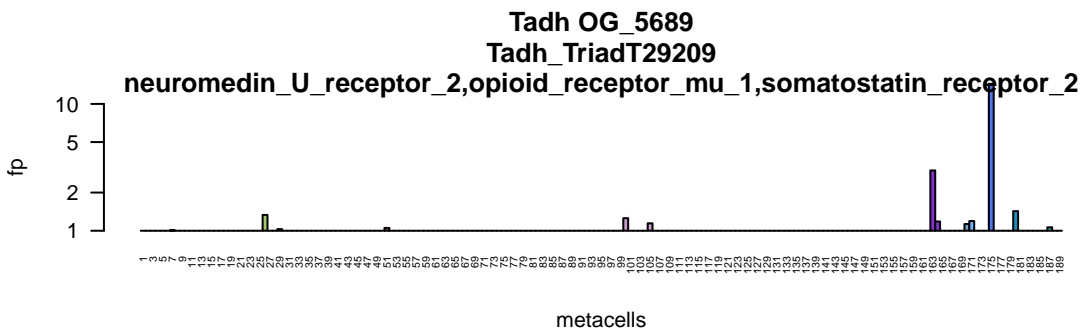
HoiH23 OG_4983
HoiH23_PIH23_009203-RA

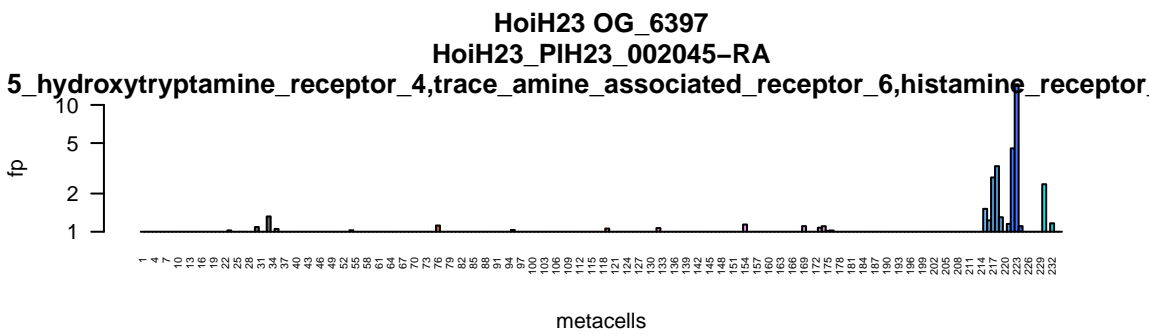
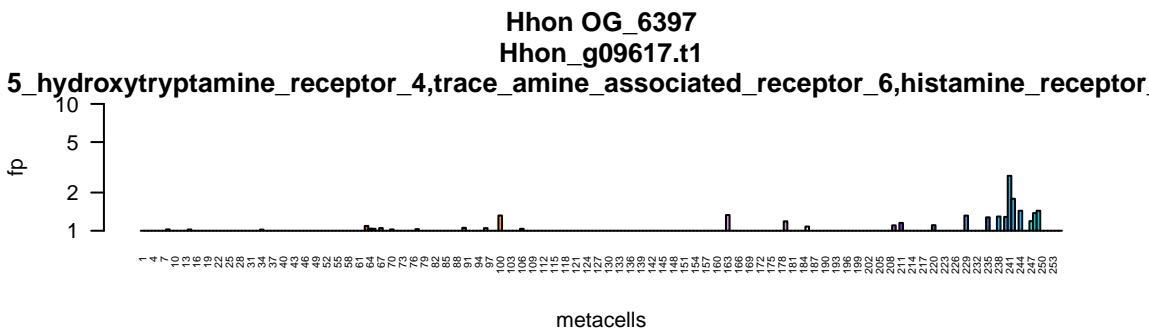
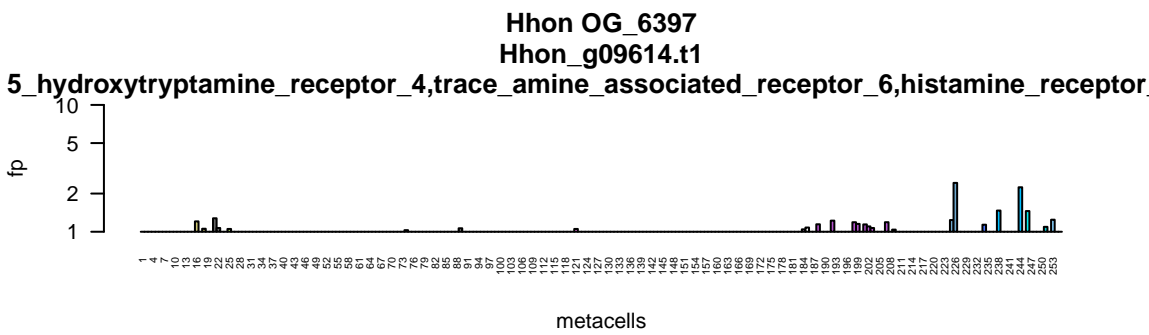
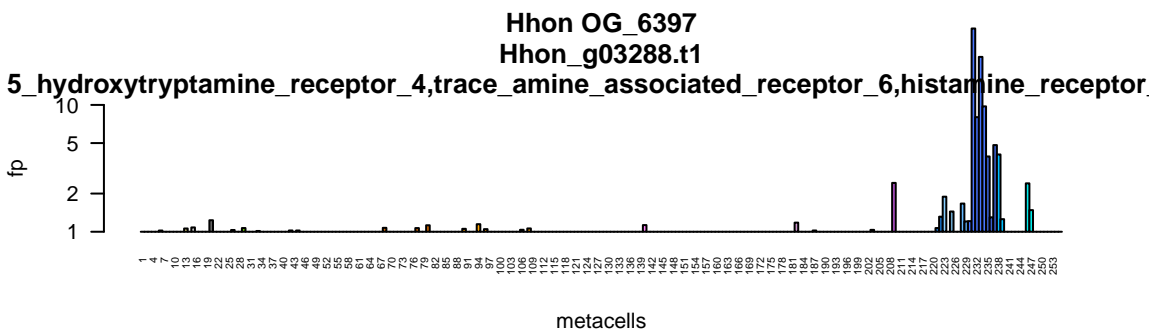
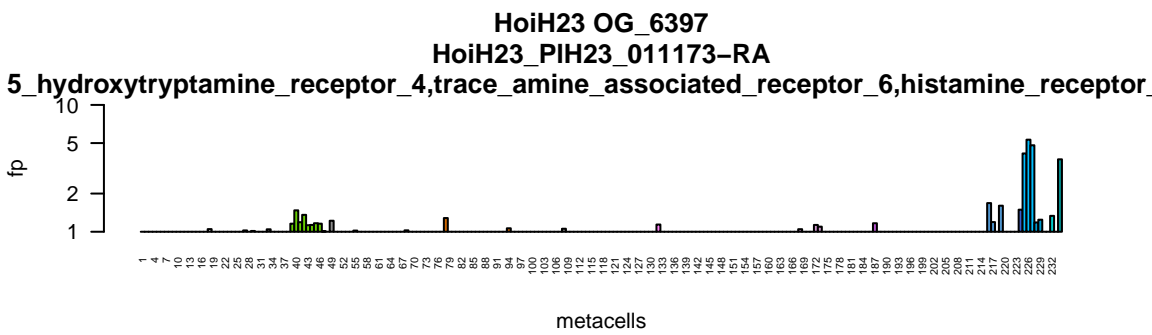
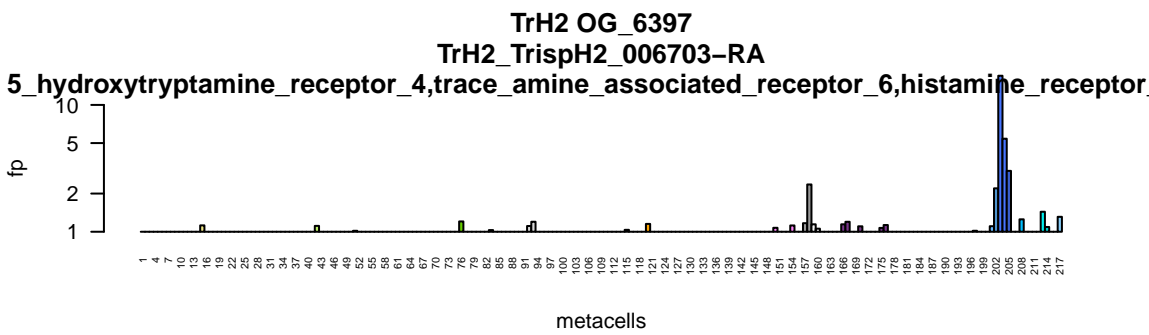
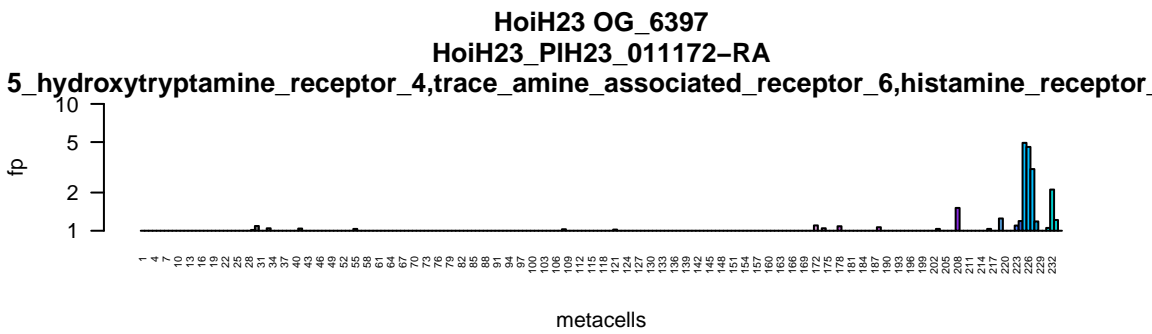
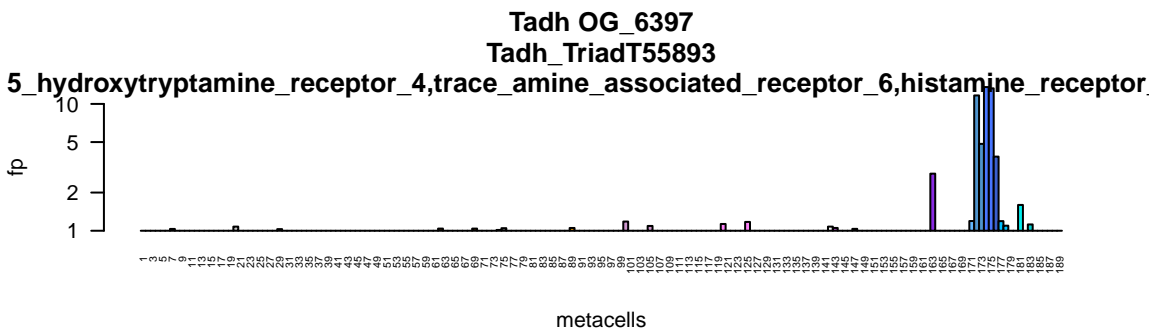


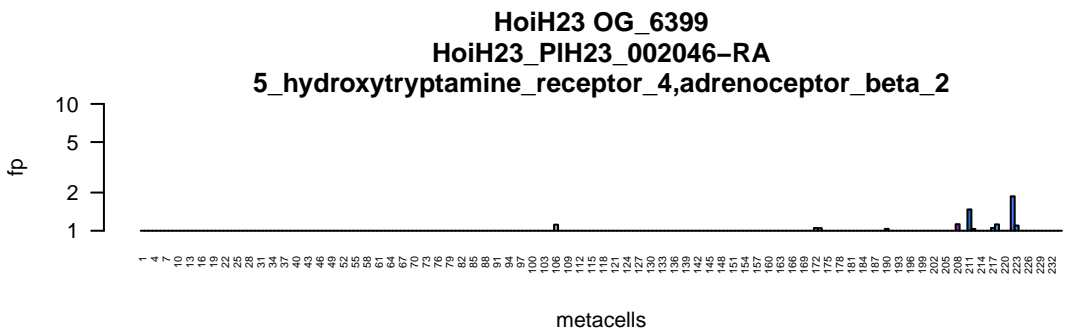
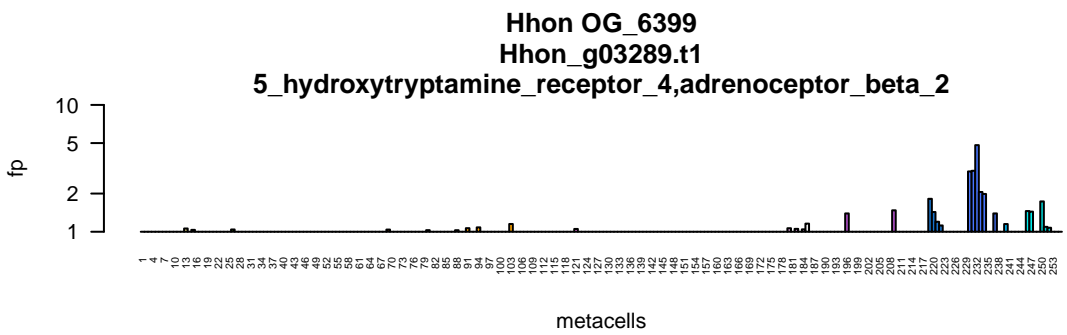
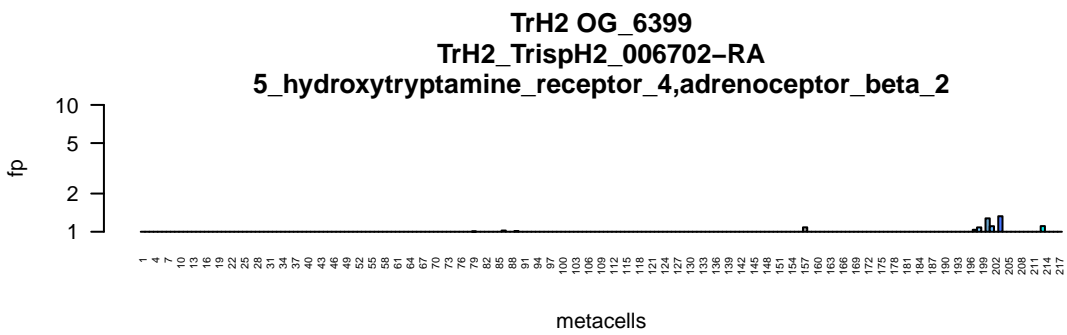
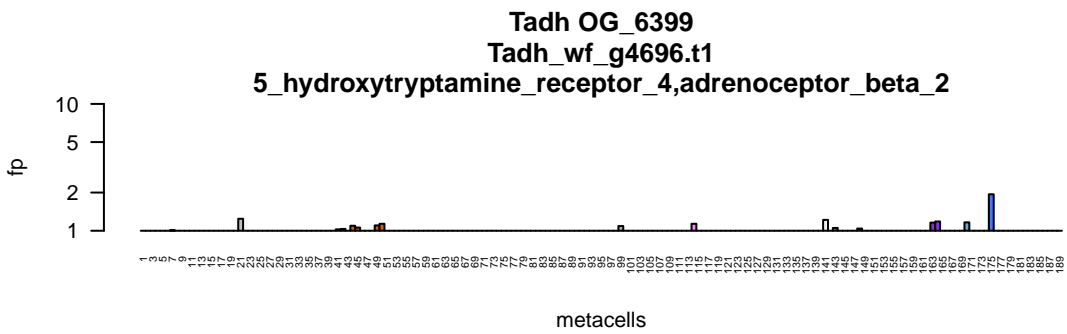
HoiH23 OG_4983
HoiH23_PIH23_009204-RA



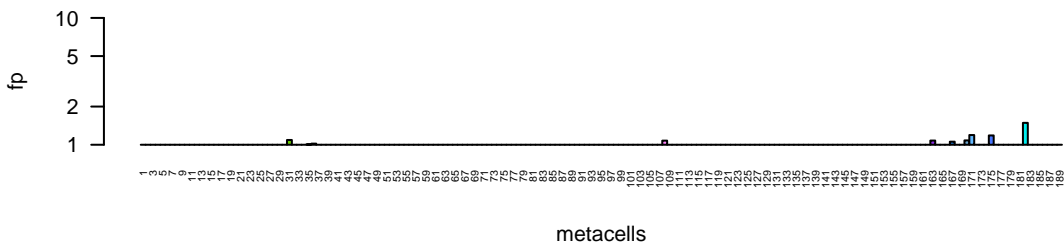




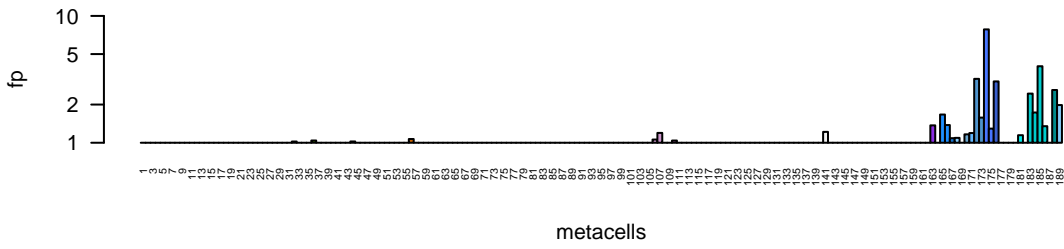




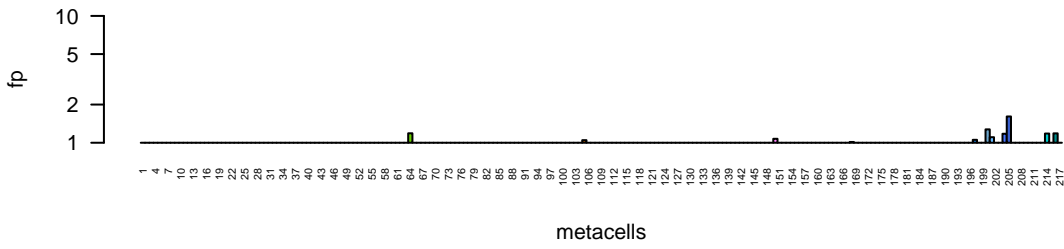
Tadh OG_6482
Tadh_wf_g9560.t1



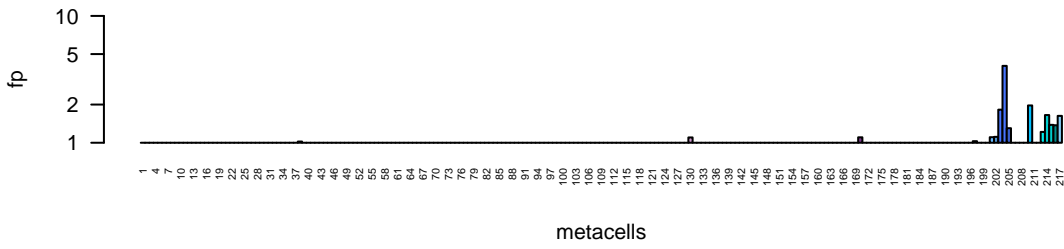
Tadh OG_6482
Tadh_wf_g9559.t1



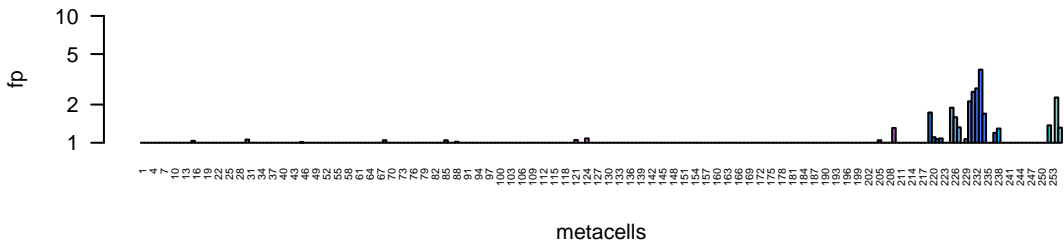
TrH2 OG_6482
TrH2_TrispH2_009482-RA



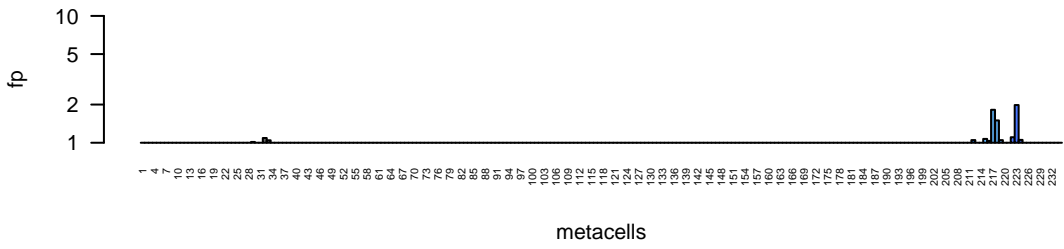
TrH2 OG_6482
TrH2_TrispH2_011478-RA

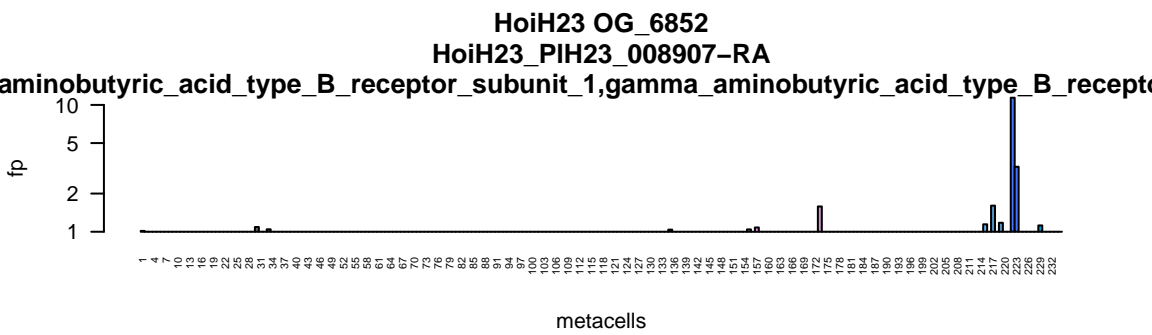
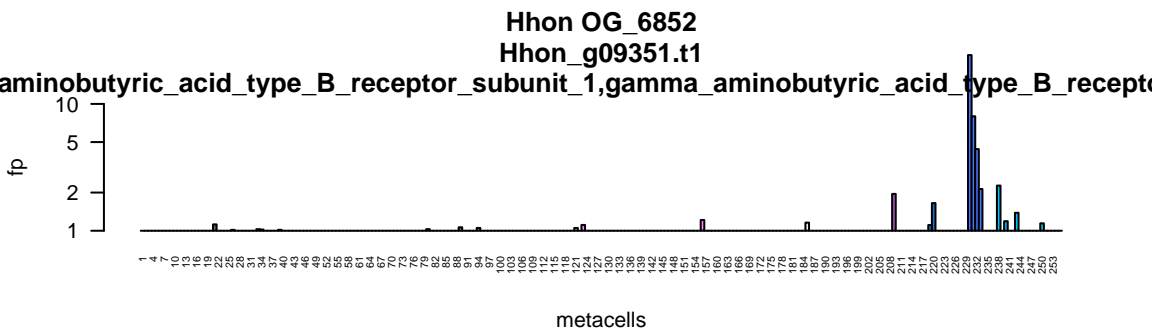
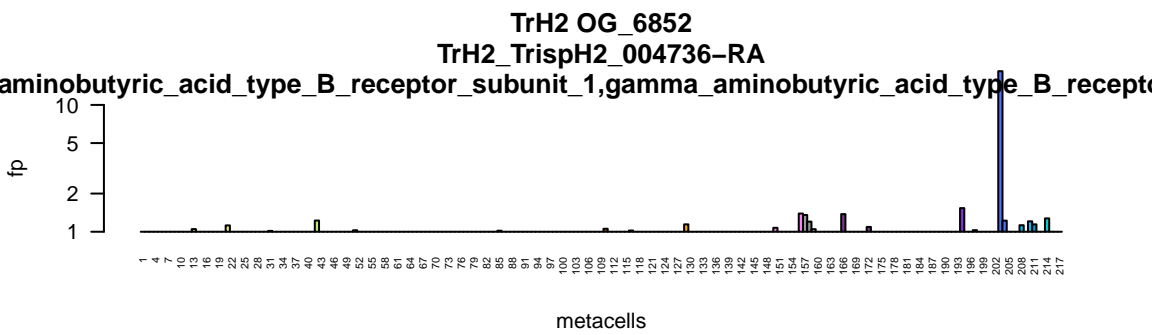
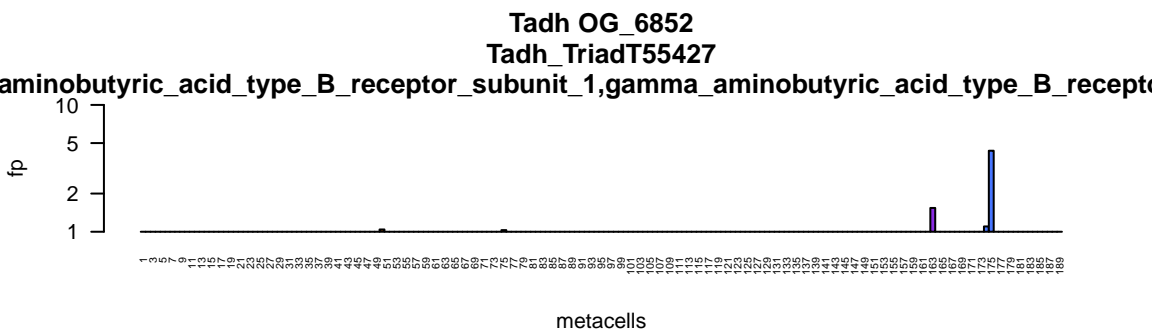
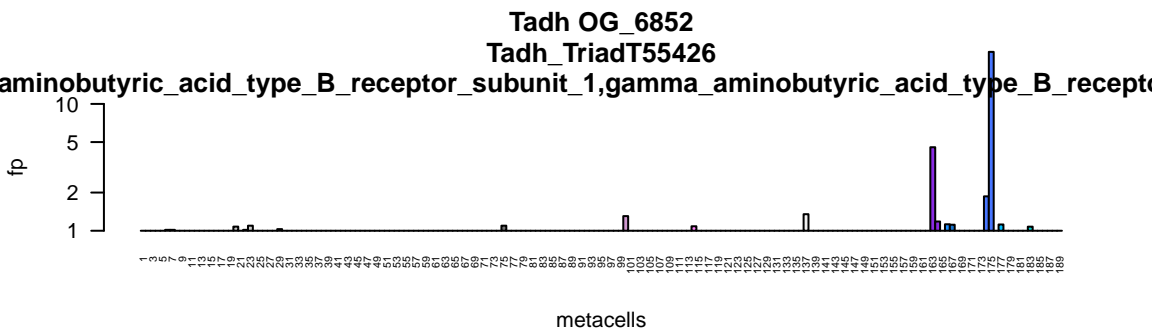


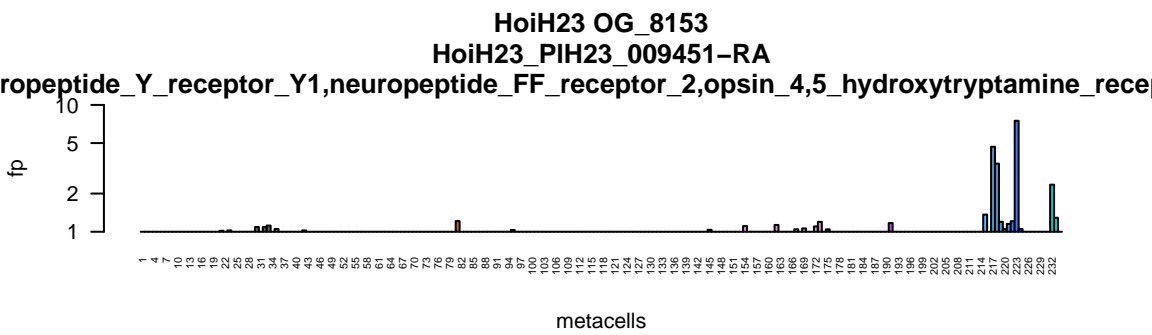
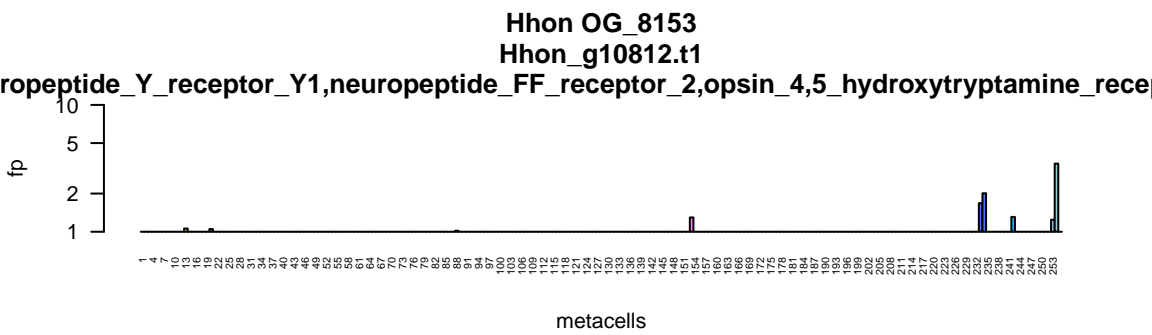
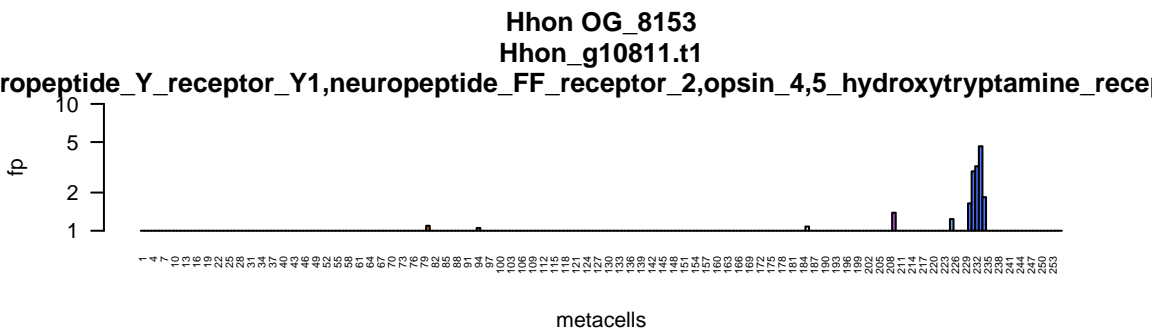
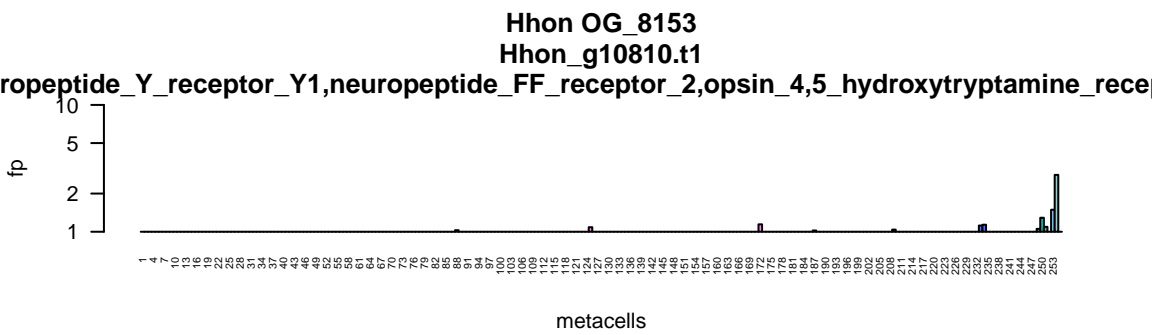
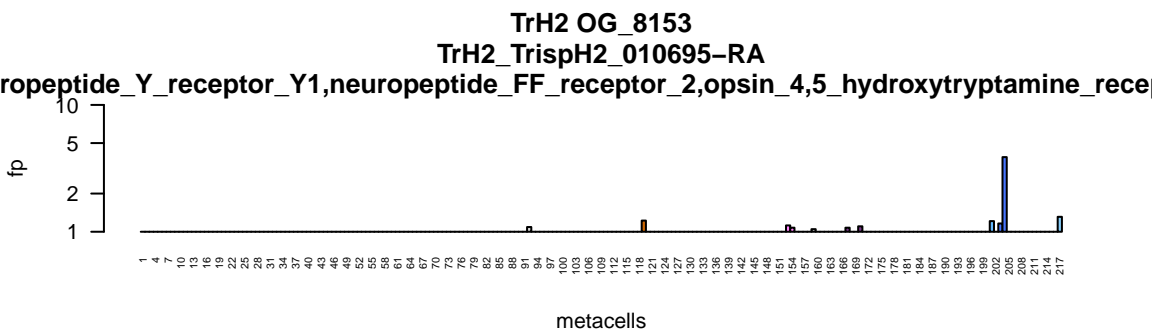
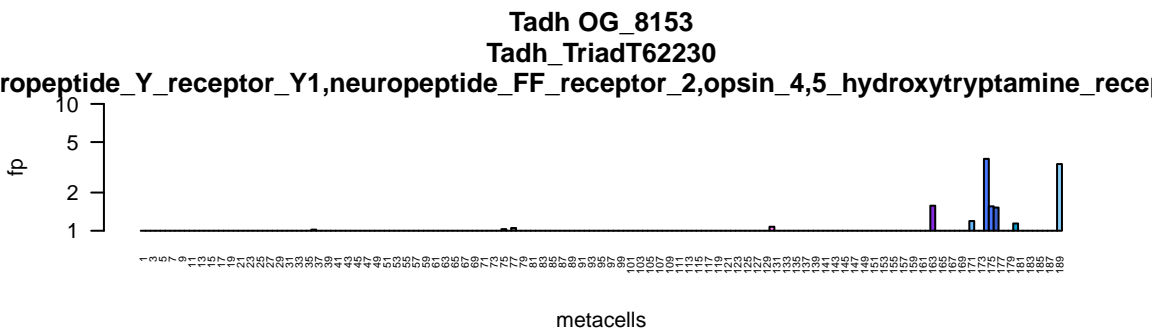
Hhon OG_6482
Hhon_g10969.t1



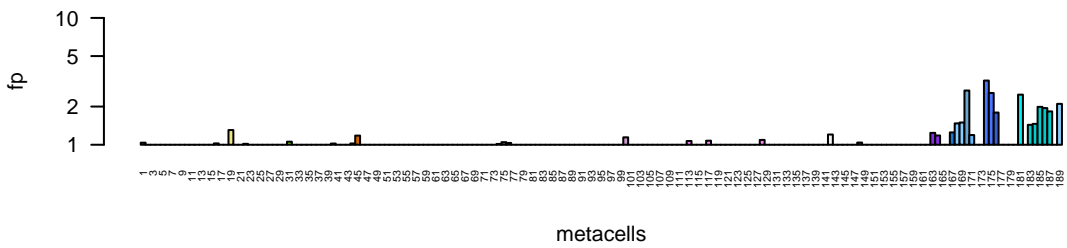
HoiH23 OG_6482
HoiH23_PIH23_011006-RA



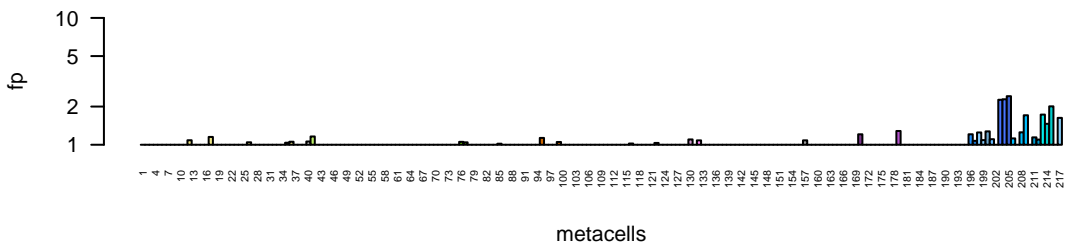




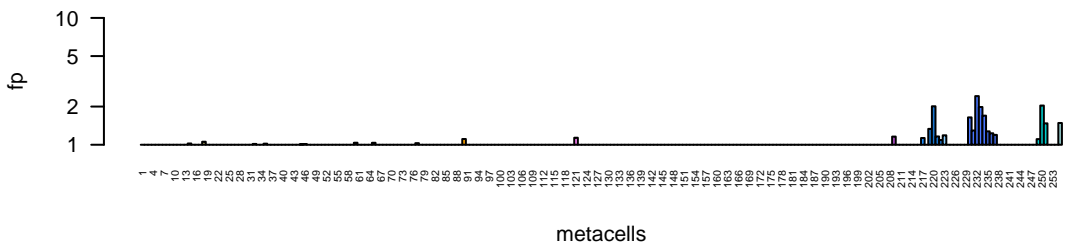
Tadh OG_1238
Tadh_wf_g1259.t1



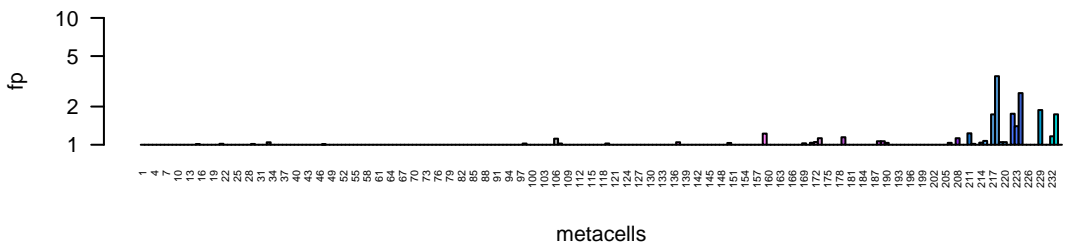
TrH2 OG_1238
TrH2_TrispH2_000180-RA

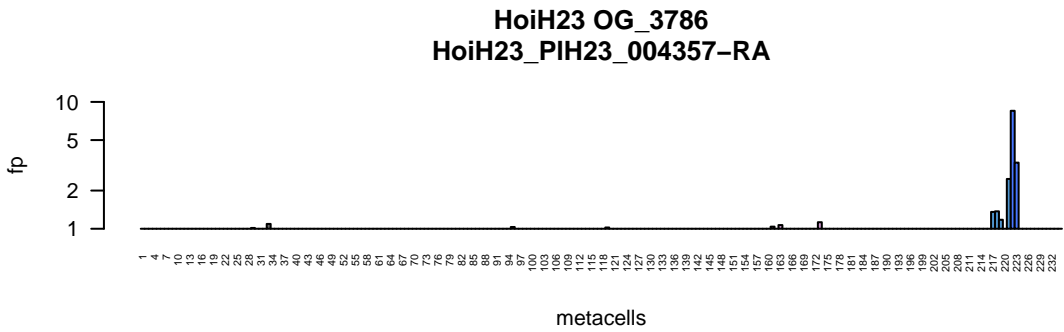
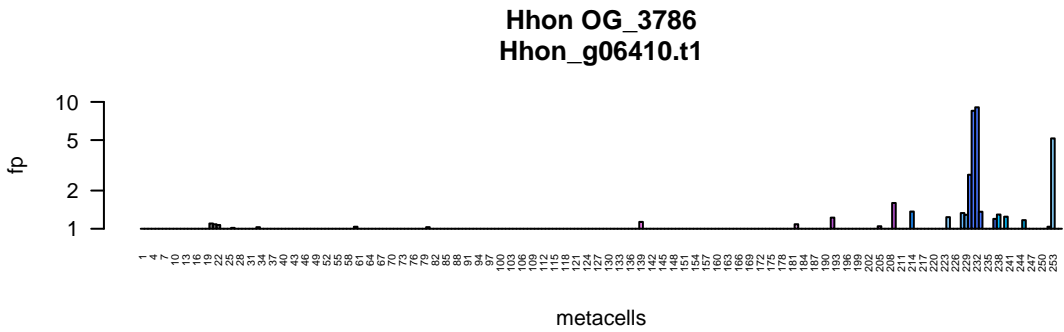
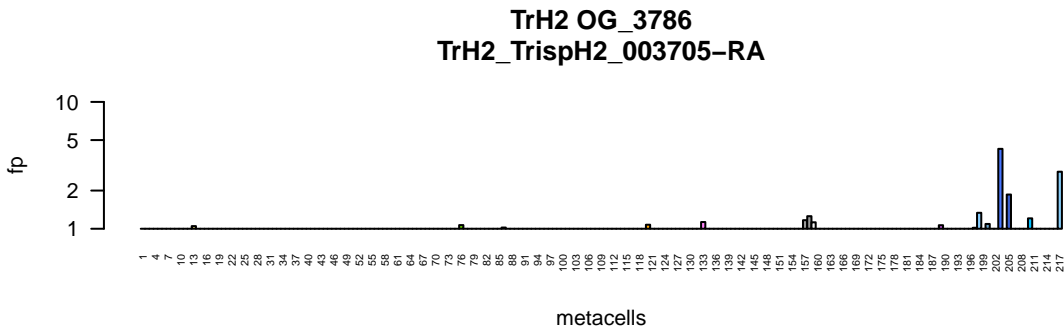
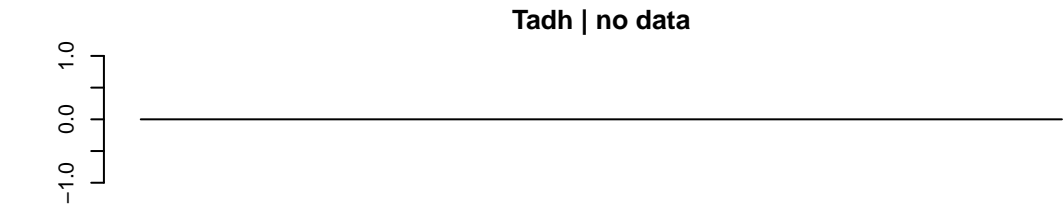


Hhon OG_1238
Hhon_g04472.t1



HoiH23 OG_1238
HoiH23_PIH23_000771-RA





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	1
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 frequency of metacells. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists metacell IDs from 1 to 187. Most metacells have a frequency of 1, with a few having higher frequencies: metacell 167 has a frequency of 2, metacell 171 has a frequency of 3, metacell 172 has a frequency of 4, and metacell 173 has a frequency of 3.

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 48 metacells. Most metacells have a false positive count of 1, with a few having 2 or 3. Metacells 199, 202, 203, 208, and 214 show higher counts, up to 4.

metacell	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	2
202	1
203	4
206	2
208	3
211	2
214	1

metacells	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	2
20	1
21	2
22	1
23	2
24	1
25	2
26	1
27	2
28	1
29	2
30	1
31	2
32	1
33	2
34	1
35	2
36	1
37	2
38	1
39	2
40	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
79	1
82	1
85	0
88	0
91	1
94	1
97	0
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	0
130	0
133	0
136	0
139	0
142	0
145	1
148	1
151	0
154	1
157	1
160	1
163	1
166	0
169	1
172	1
175	1
178	0
181	1
184	1
187	1
190	0
193	1
196	1
199	1
202	4
203	5
208	1
211	1
214	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
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26	1
27	1
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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 metacells from 1 to 232. Most metacells have a false positive count of 1. Metacells 169, 170, 171, 172, 173, 174, 175, 176, 177, 178, 179, 180, 181, 182, 183, 184, 185, 186, 187, 188, 189, 190, 191, 192, 193, 194, 195, 196, 197, 198, 199, 200, 201, 202, 203, 204, 205, 206, 207, 208, 209, 210, 211, 212, 213, 214, 215, 216, 217, 218, 219, 220, 221, 222, 223, 224, 225, 226, 227, 228, 229, 230, 231, 232 have higher false positive counts, with some reaching up to 10.

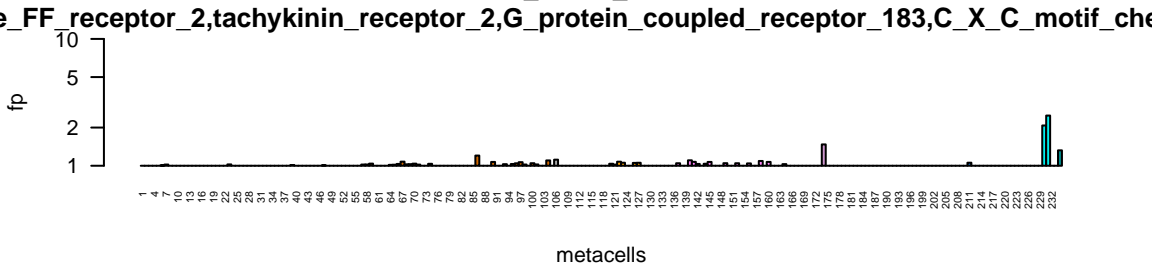
Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 232. Most metacells have 1 false positive, with a small cluster of metacells (211-232) showing higher values up to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

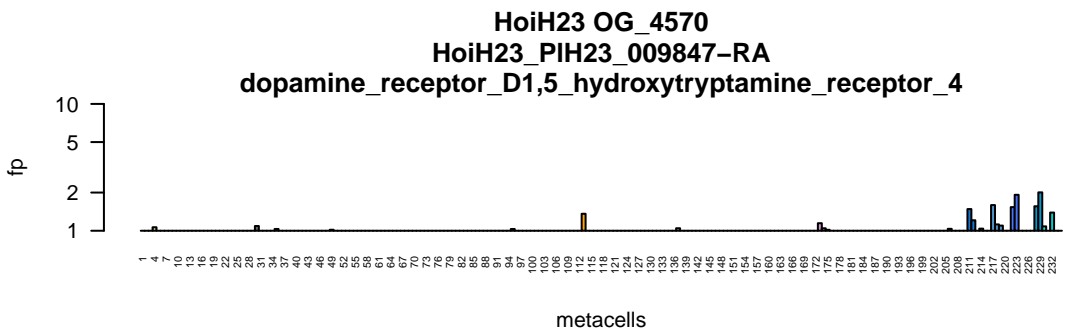
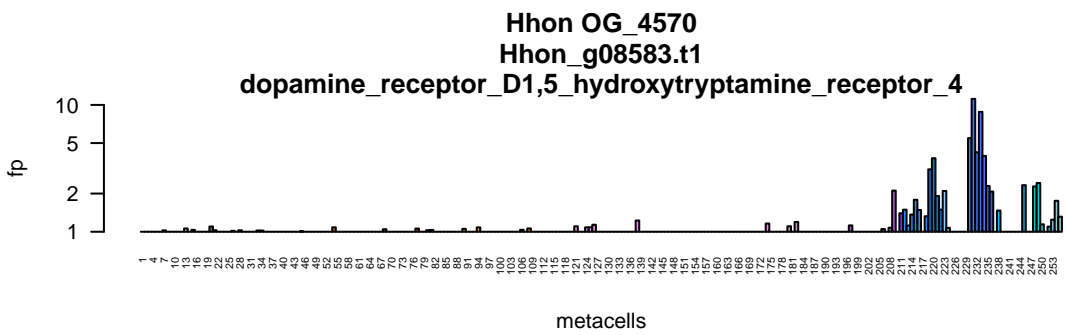
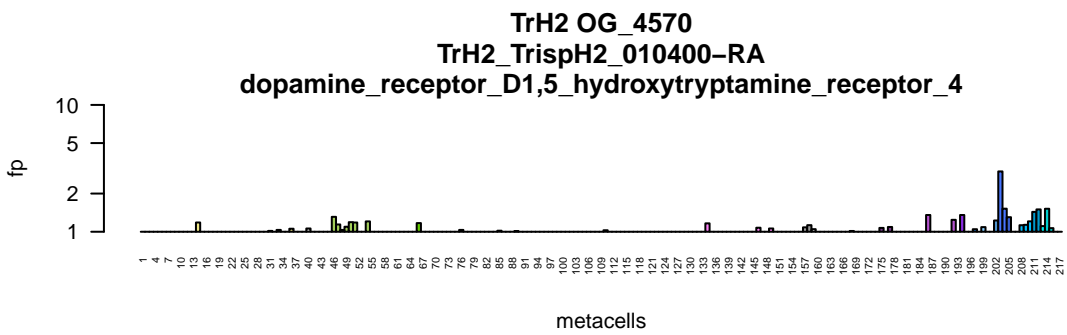
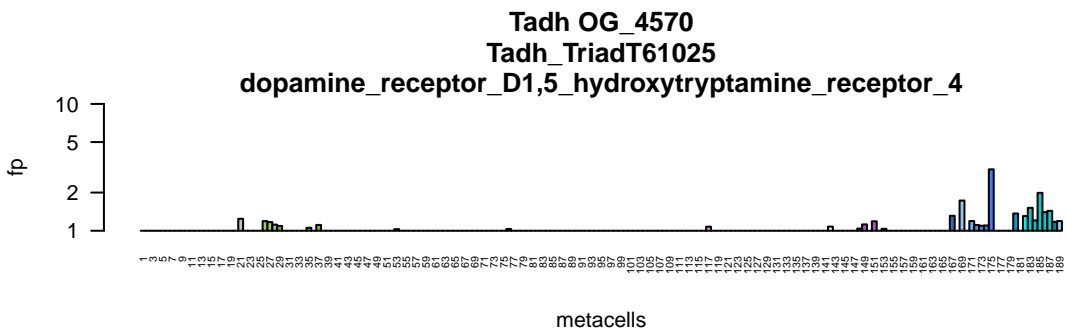
Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 232. Most metacells have 0 false positives. Metacells 220, 221, 222, and 223 show higher counts, with 223 having the highest count at 10.

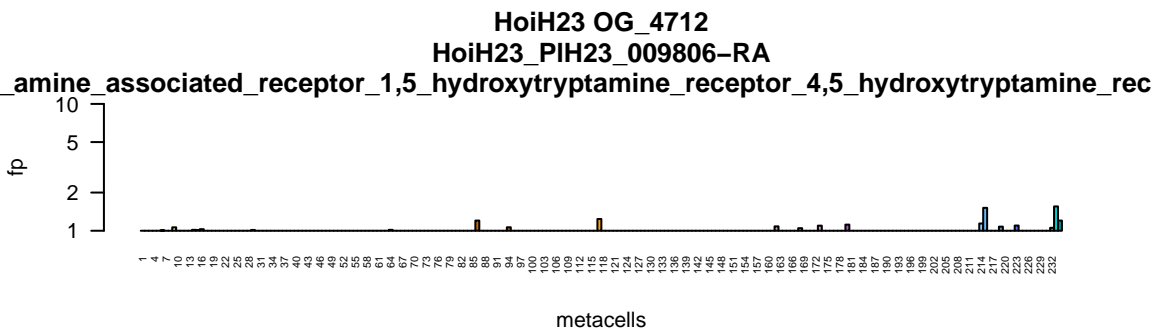
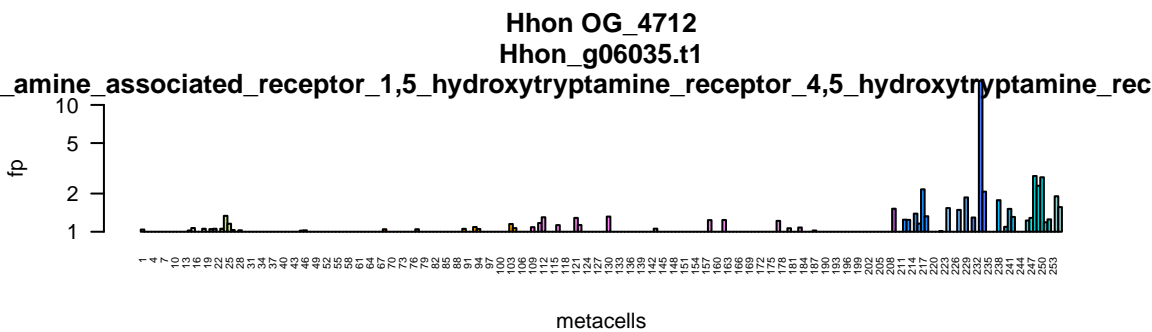
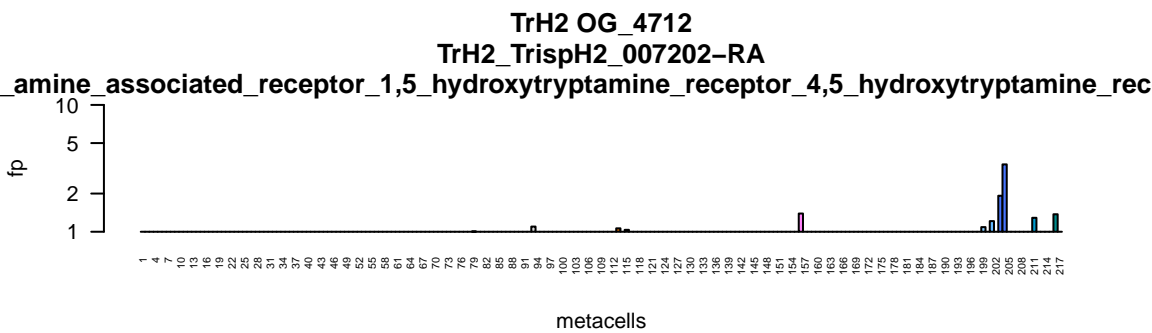
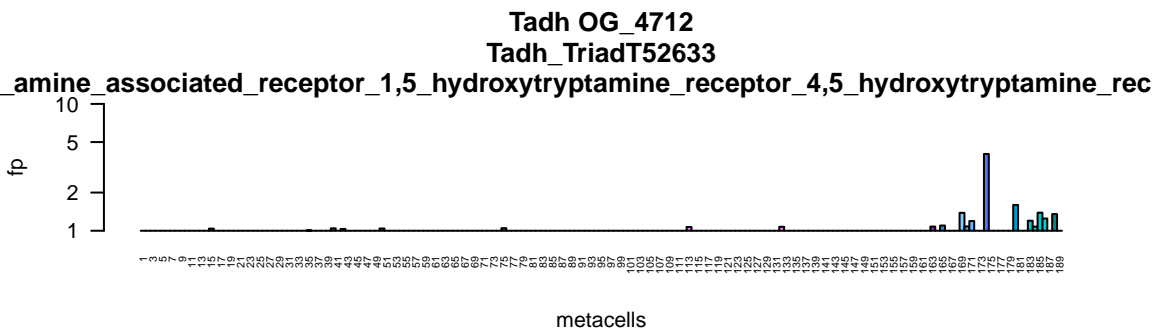
metacell	fp
1	0
4	1
10	0
13	0
19	0
22	0
25	0
31	0
34	1
40	0
43	0
46	0
52	0
55	0
58	0
60	0
64	0
67	0
73	0
76	1
82	0
85	0
88	0
94	1
97	0
103	0
106	1
109	0
115	0
118	0
124	0
127	1
130	0
136	1
139	0
145	0
151	1
157	0
160	0
166	0
169	0
172	1
178	0
181	0
187	0
190	0
193	0
199	0
202	0
208	0
211	1
214	0
217	3
220	2
221	1
222	6
223	10
229	9
232	1

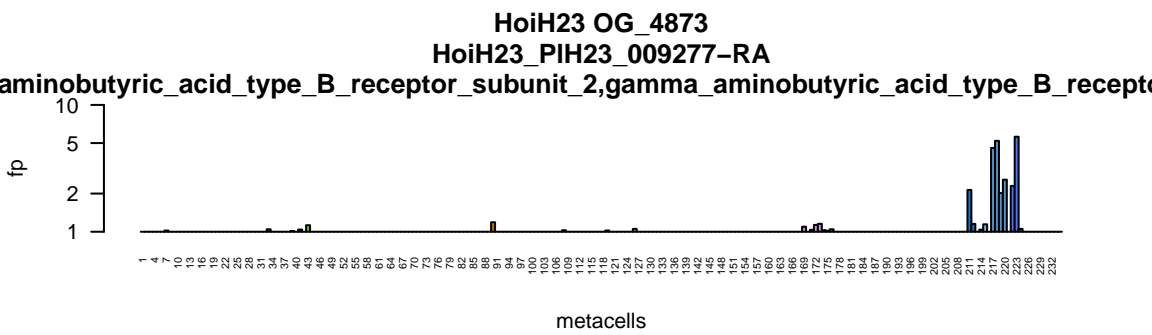
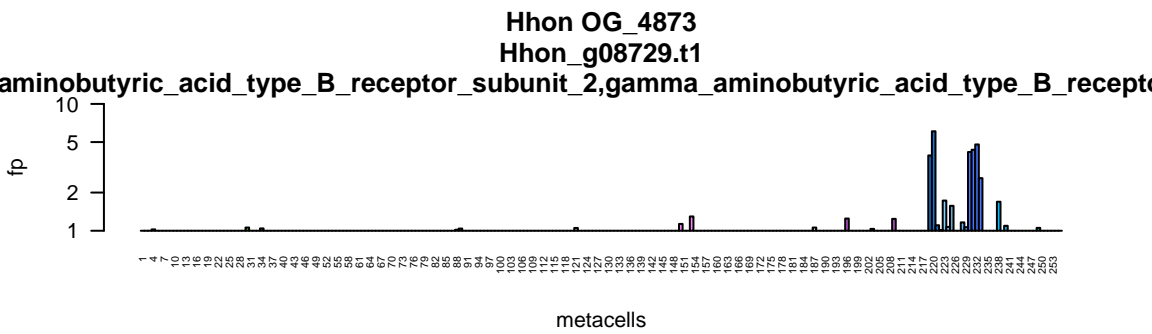
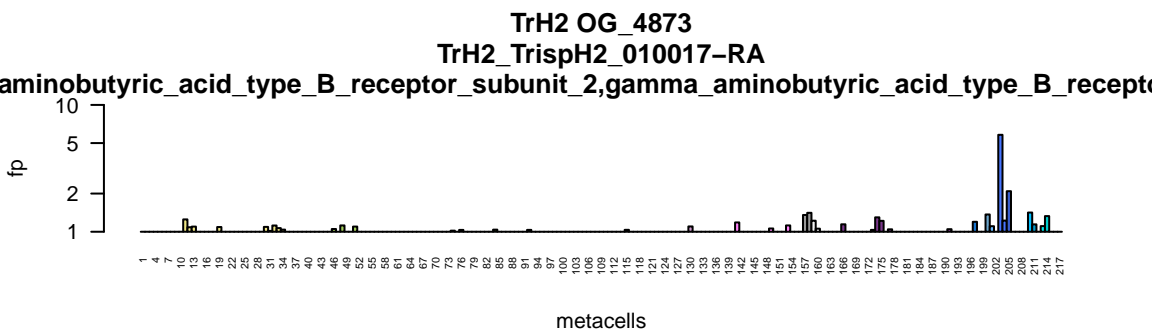
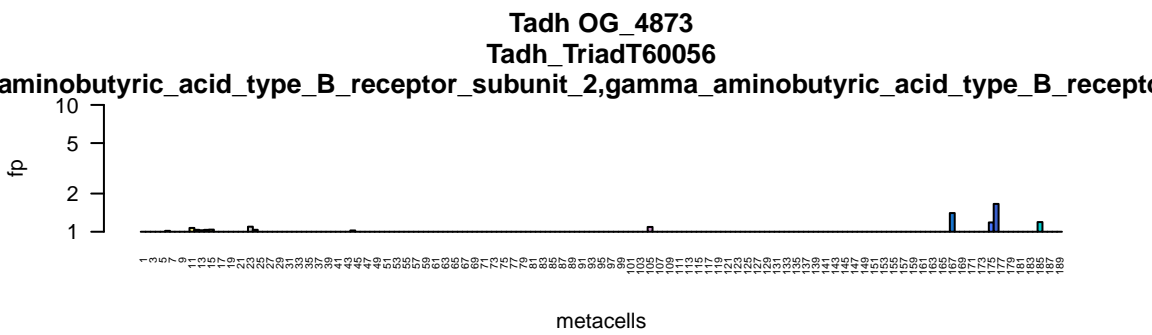
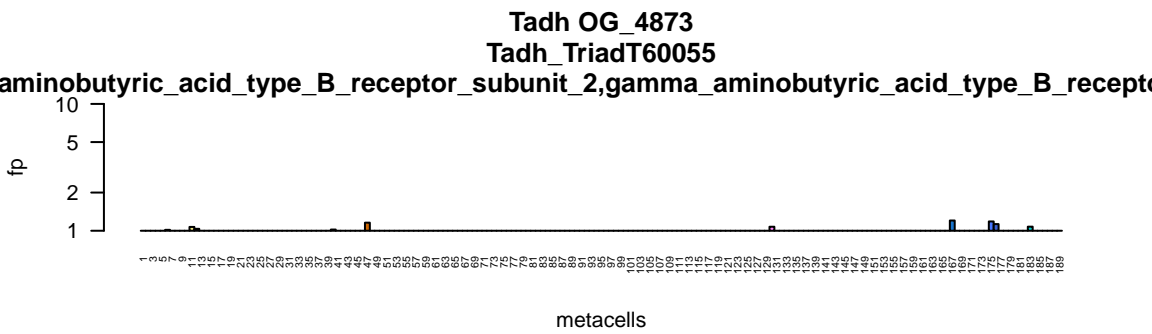
metacell	fp
1	0
4	0
10	0
13	0
19	0
22	0
25	0
31	1
34	1
40	0
43	0
46	0
52	1
55	0
58	0
64	0
67	0
73	0
76	0
82	0
85	0
89	0
94	0
97	0
103	0
106	0
109	0
115	0
118	0
124	0
127	0
130	0
136	0
139	0
145	0
148	0
151	1
157	0
160	0
166	0
169	1
172	0
178	0
187	0
190	0
193	0
199	0
202	0
208	0
211	0
214	2
220	2
223	3
229	1
232	2

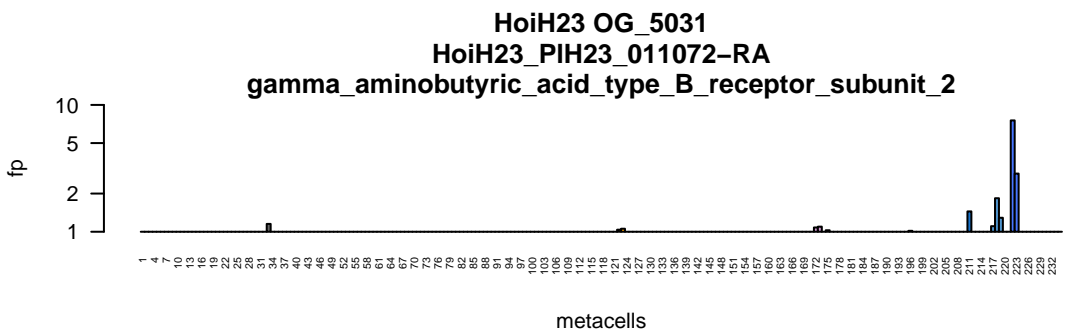
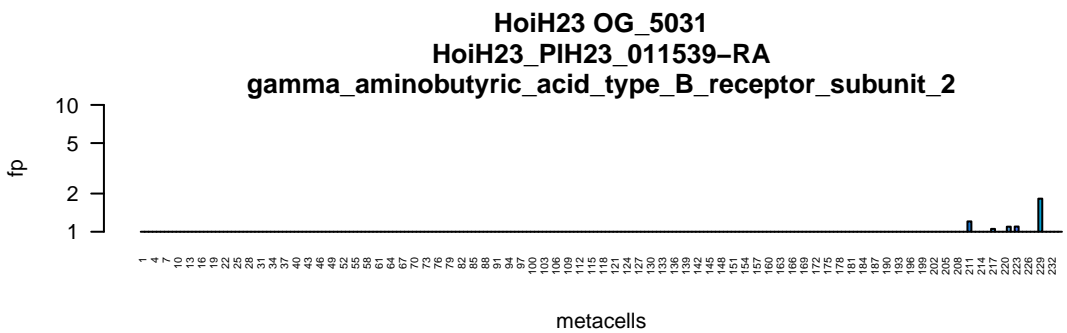
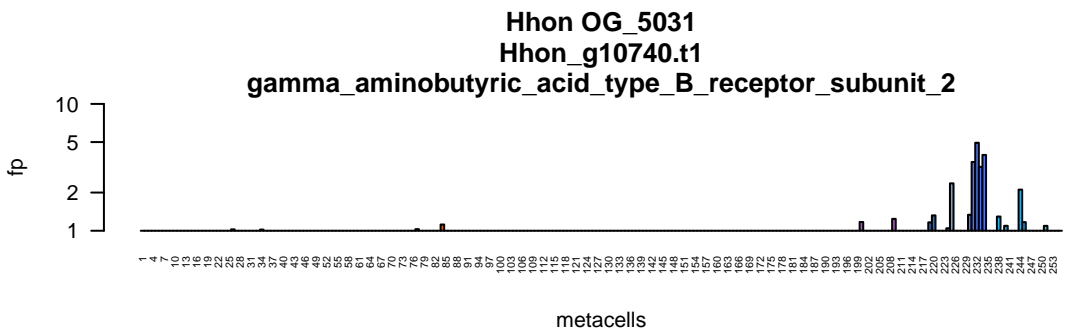
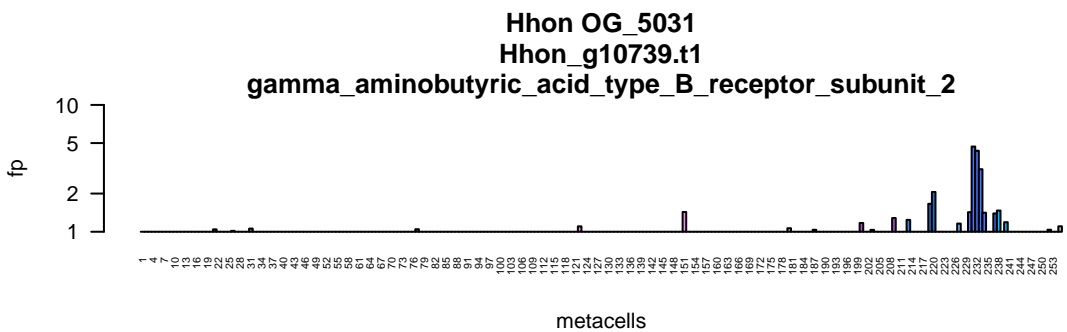
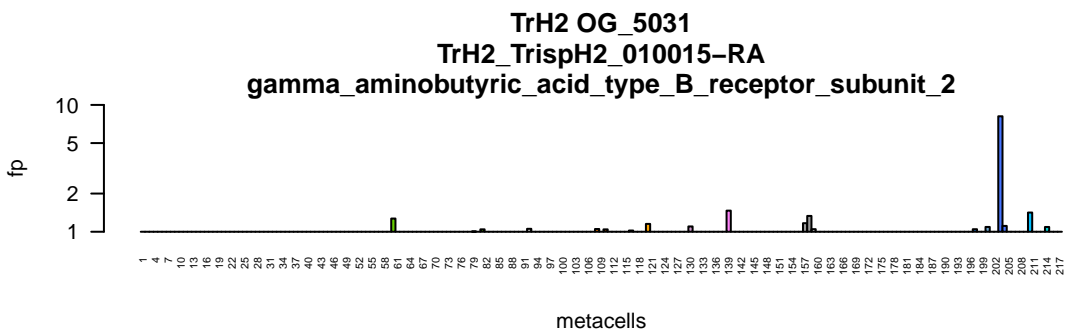
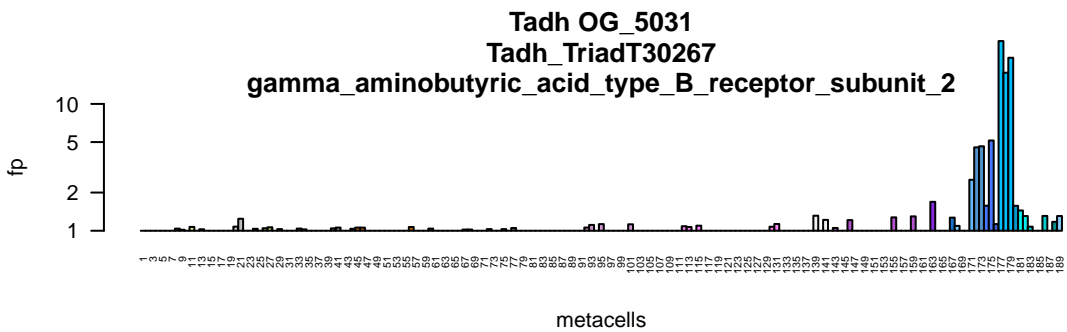
HoiH23 OG_4547
HoiH23_PIH23_002323-RA







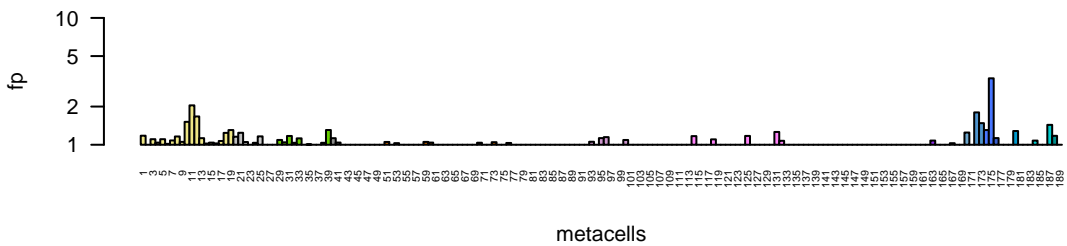




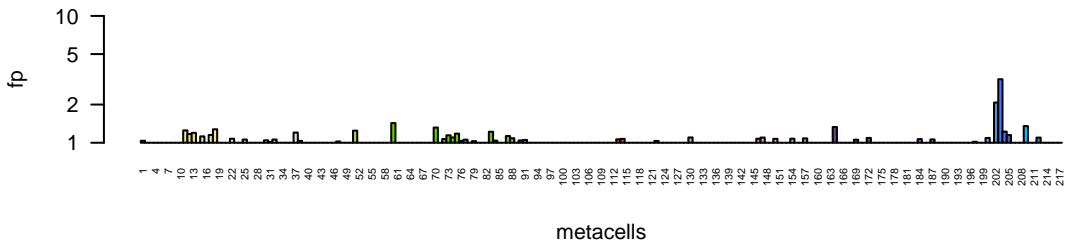


Tadh OG_5119

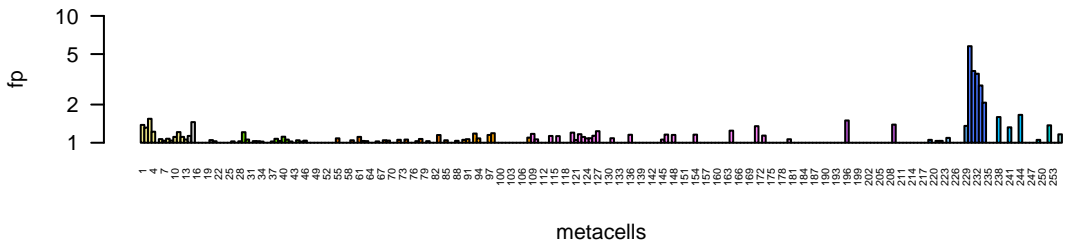
Tadh_TriadT61436



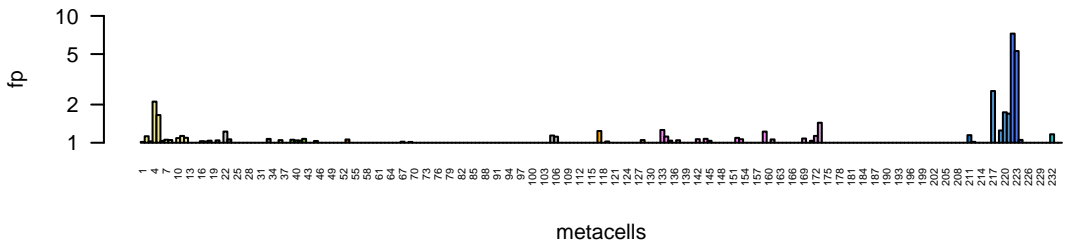
TrH2 OG_5119
TrH2_TrispH2_003533-RA



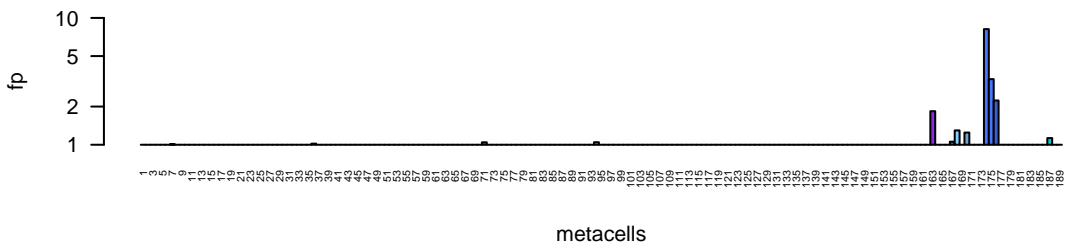
Hhon OG_5119
Hhon_g07857.t1



HoiH23 OG_5119
HoiH23_PIH23_007705-RA

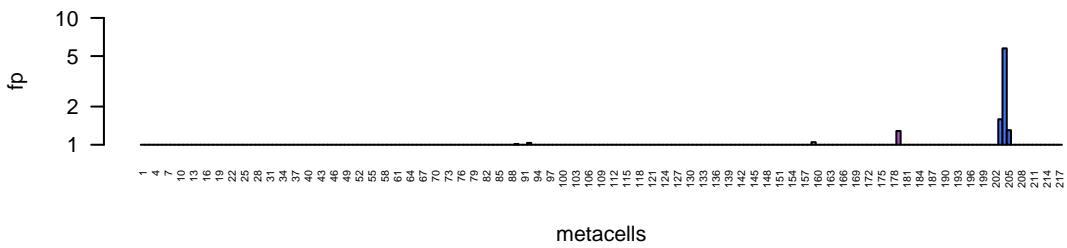


Tadh OG_5432
Tadh_wf_g4961.t1



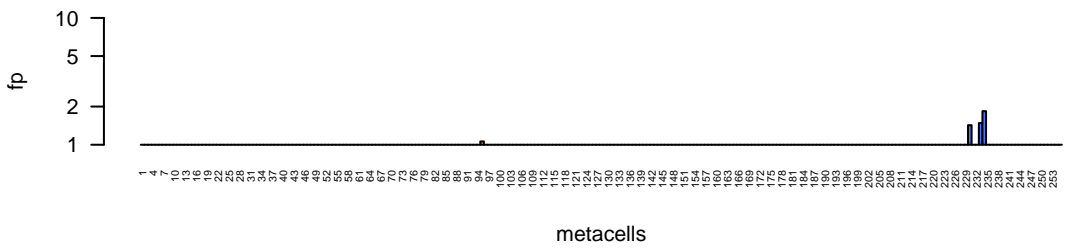
metacells

TrH2 OG_5432
TrH2_TrispH2_009854-RA



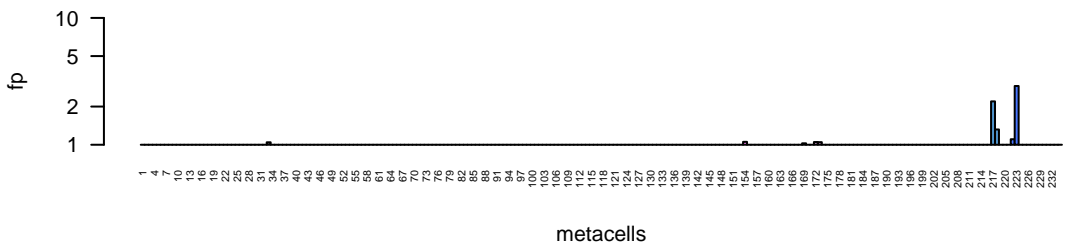
metacells

Hhon OG_5432
Hhon_g10436.t1

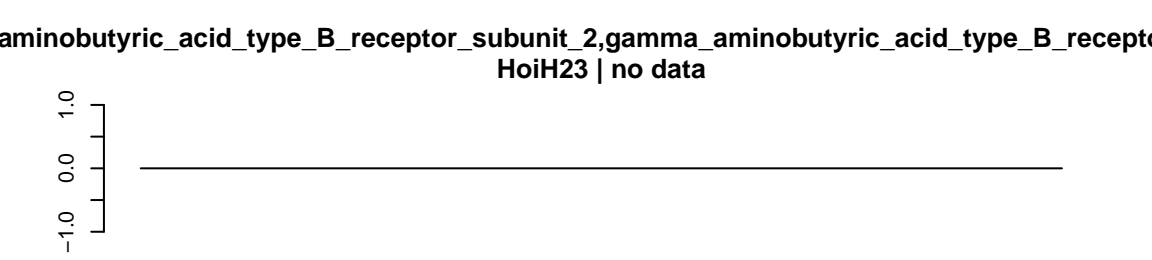
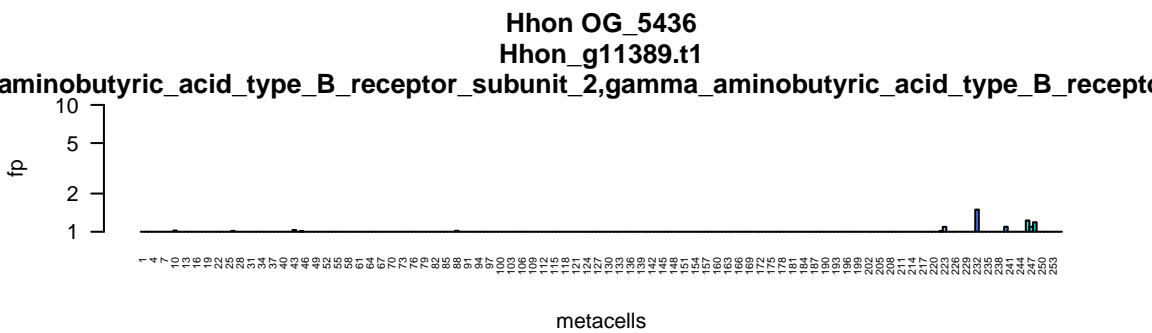
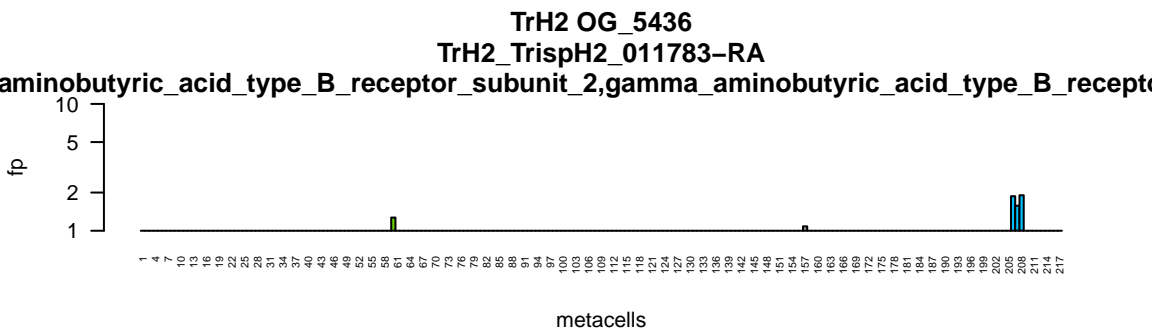
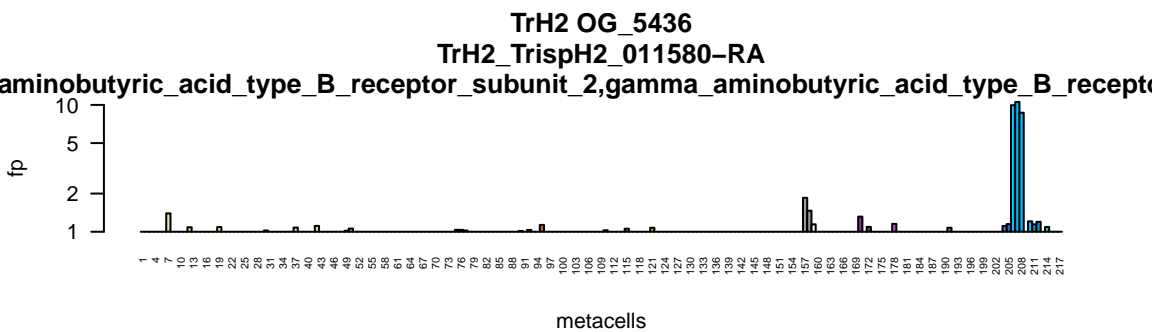
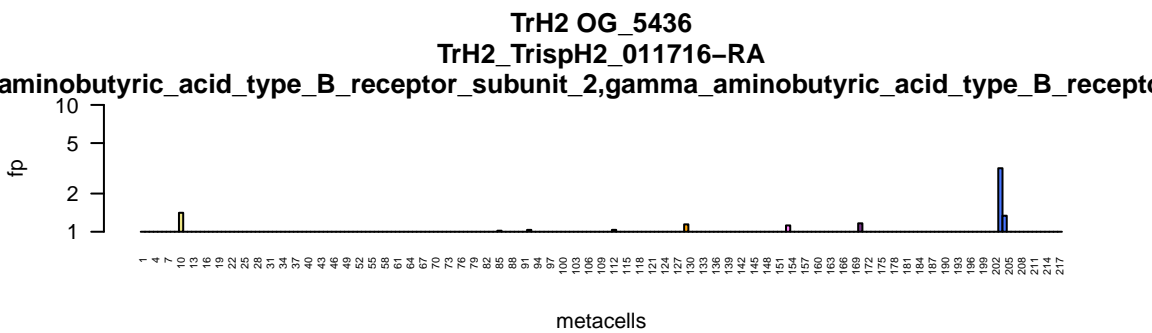
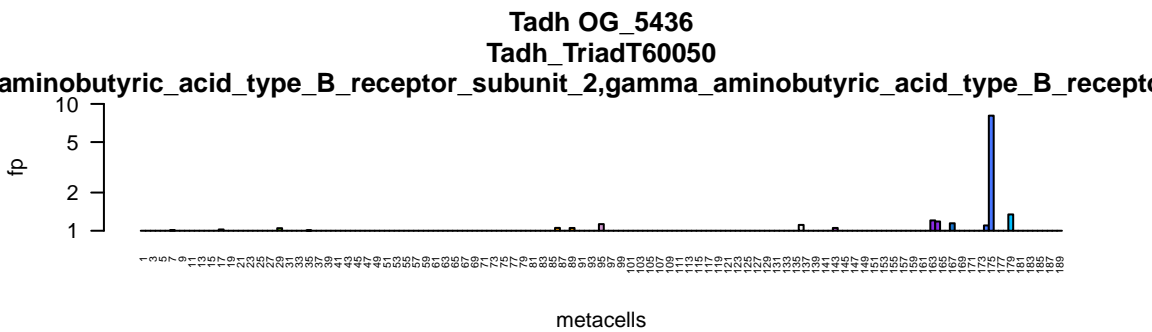


metacells

HoiH23 OG_5432
HoiH23_PIH23_010495-RA



metacells



Bar chart showing the frequency of metacells (x-axis) versus the frequency of pairs (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 frequencies, with most metacells having a frequency of 1, and a few having higher frequencies up to 6.

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, 177, 178, and 179 show higher counts, with 178 having the highest count of 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	2
176	2
177	2
178	4
179	2
180	1
181	1
182	1
183	1
184	1
185	1
186	1
187	1
188	1
189	1

Bar chart showing the frequency of metacells (x-axis) versus frequency (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 frequencies across the metacells, with a peak around metacell 75.

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 161 through 189 show a higher frequency of false positives, with some reaching 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
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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

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 a distribution of metacell frequencies, with most metacells having a frequency of 1, and a few having higher frequencies up to 5.

A bar chart showing the frequency of metacells (x-axis) versus the frequency of pairs (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows a distribution of metacells with varying frequencies, with a peak around metacell 130.

Bar chart showing the frequency (fp) of metacells. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists indices from 1 to 217. Most metacells have a frequency of 1. Notable outliers include metacell 202 with a frequency of 4, metacell 205 with a frequency of 3, and metacell 214 with a frequency of 3.

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	1
199	1
202	4
205	3
208	1
211	1
214	3
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	1.5
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.

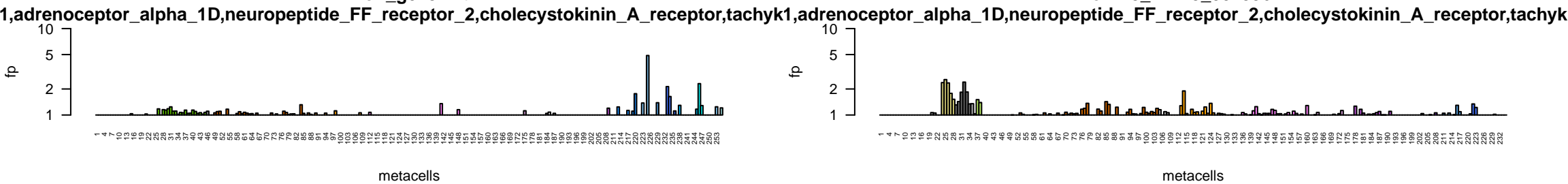
metacells	fp
1	1
4	1
10	1
13	1
16	1
22	1
23	1
25	1
28	1
31	1
32	1
37	1
40	1
43	2
44	1
48	1
49	1
52	1
55	1
59	1
61	1
64	1
67	1
73	1
76	1
79	1
82	1
83	1
88	1
91	1
94	1
97	1
100	1
103	1
108	1
110	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	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 6.

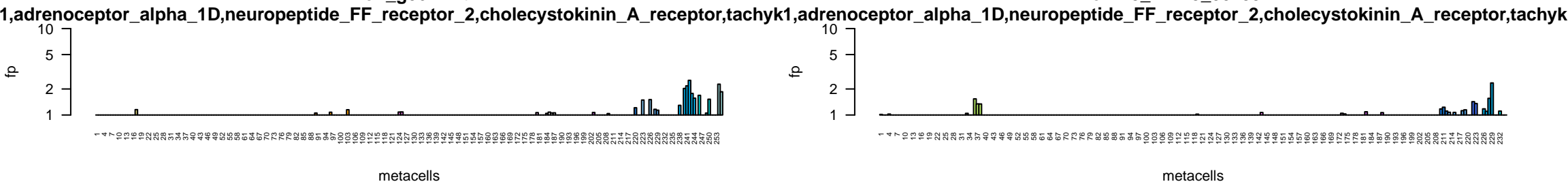
metacells	fp
1	1
4	1
10	1
13	1
16	1
19	1
22	1
25	1
28	1
31	1
37	1
40	2
41	3
42	4
43	3
44	2
49	1
52	1
55	1
58	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	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

metacells	fp
1	1
7	1
10	1
11	1
18	1
19	1
25	1
28	1
34	1
35	1
43	1
46	1
52	1
55	1
61	1
64	1
70	1
77	1
85	1
88	1
94	1
97	1
103	1
108	1
112	1
115	1
121	1
127	1
130	1
139	1
145	1
148	1
154	1
157	1
166	1
172	1
175	1
181	1
187	1
190	1
193	1
199	1
205	1
208	1
214	1
217	1
223	1
226	1
232	1
235	1
241	2
242	3
243	4
244	5
247	1
250	1
253	2

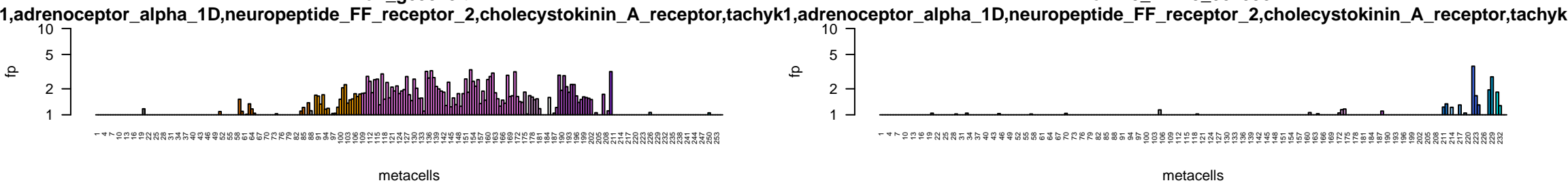
Hhon OG_5725
Hhon_g04574.t1



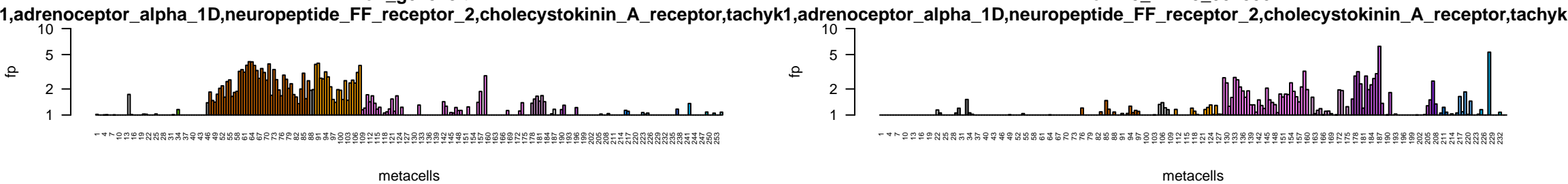
Hhon OG_5725
Hhon_g06114.t1



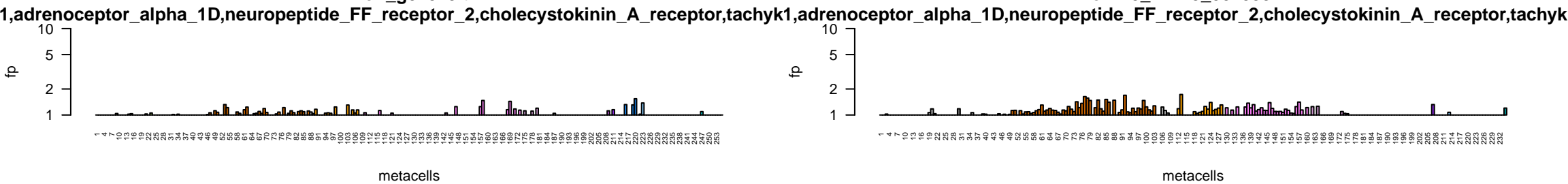
Hhon OG_5725
Hhon_g09573.t1



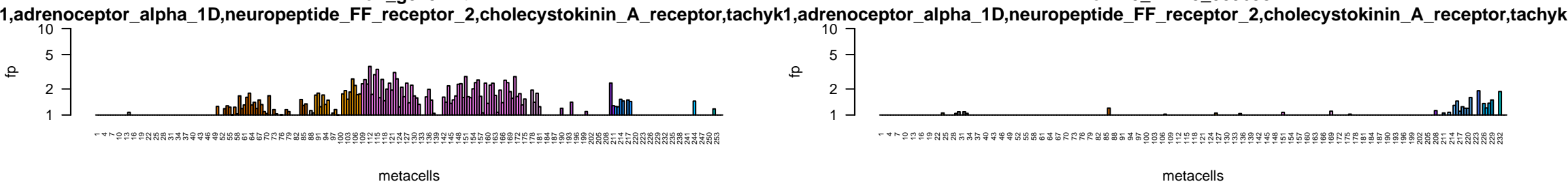
Hhon OG_5725
Hhon_g07528.t1



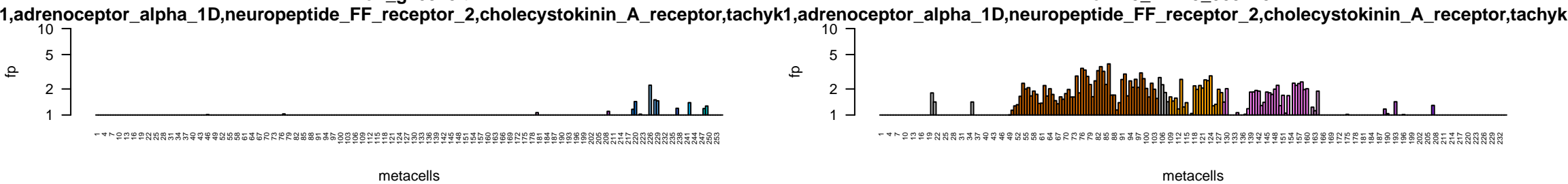
Hhon OG_5725
Hhon_g07529.t1



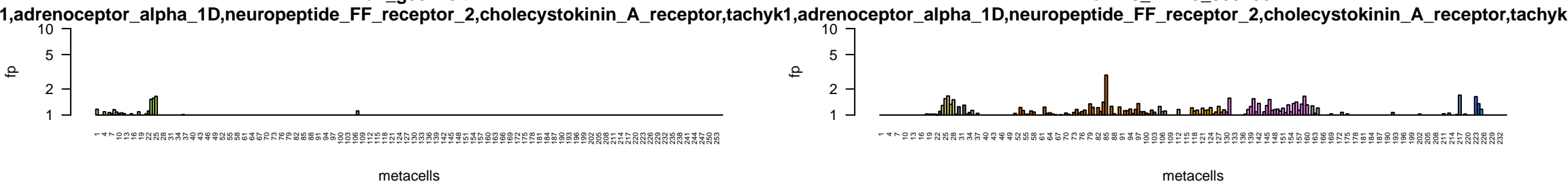
Hhon OG_5725
Hhon_g07527.t1

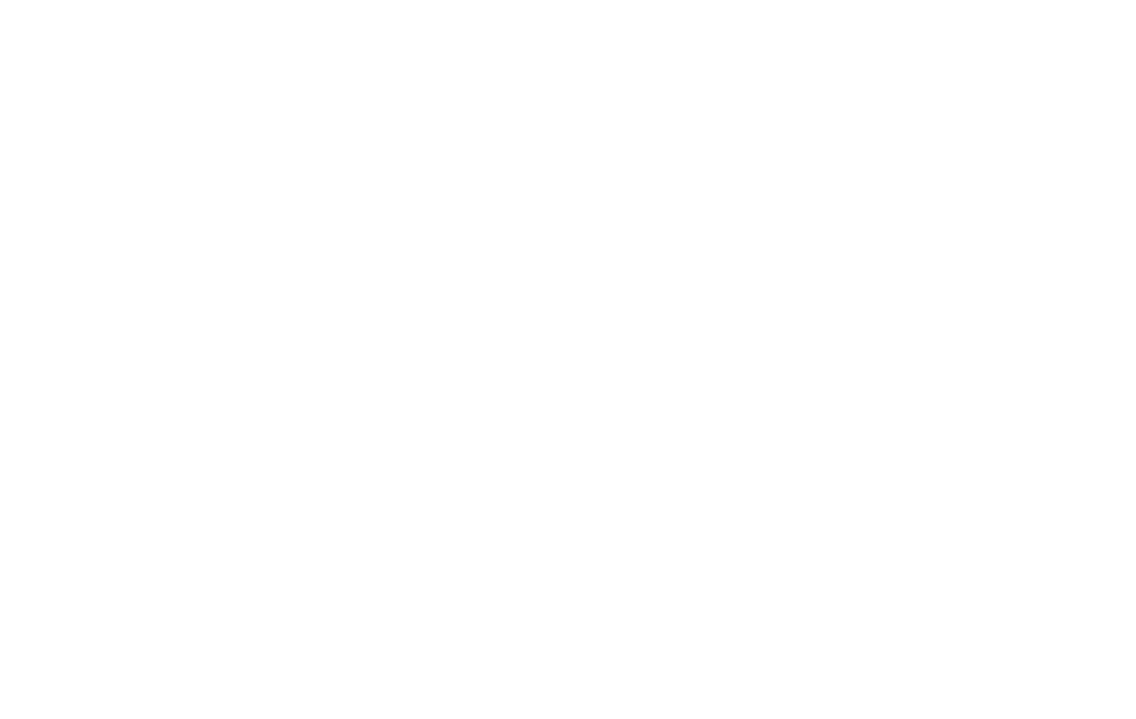
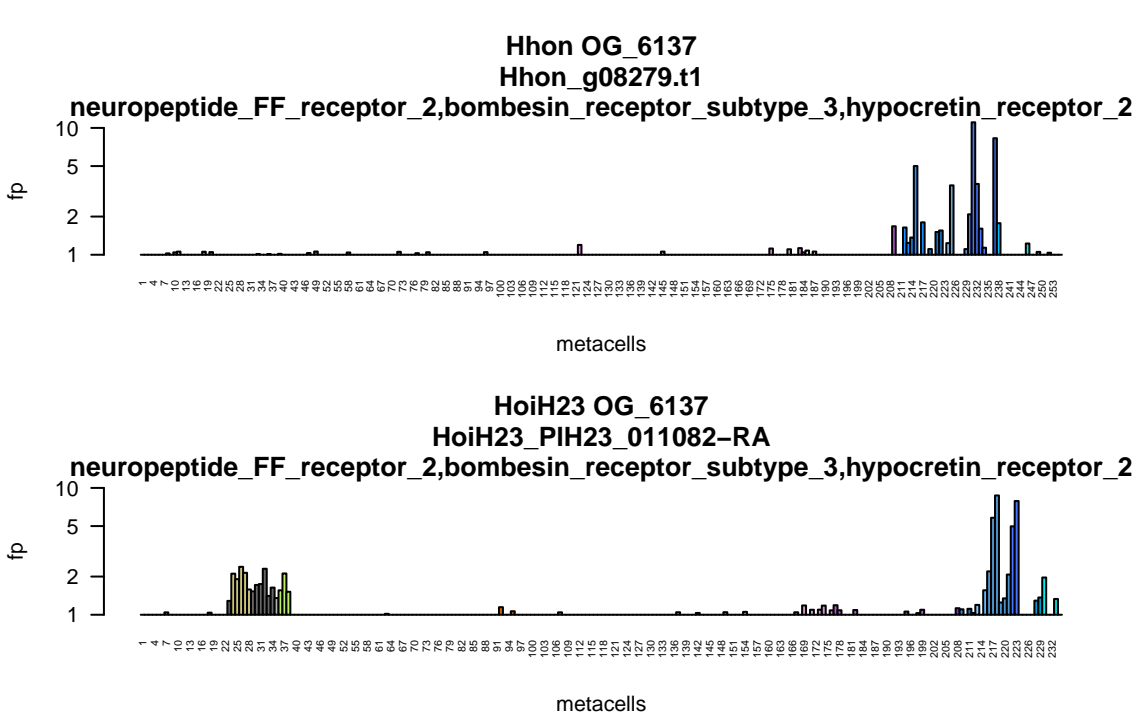
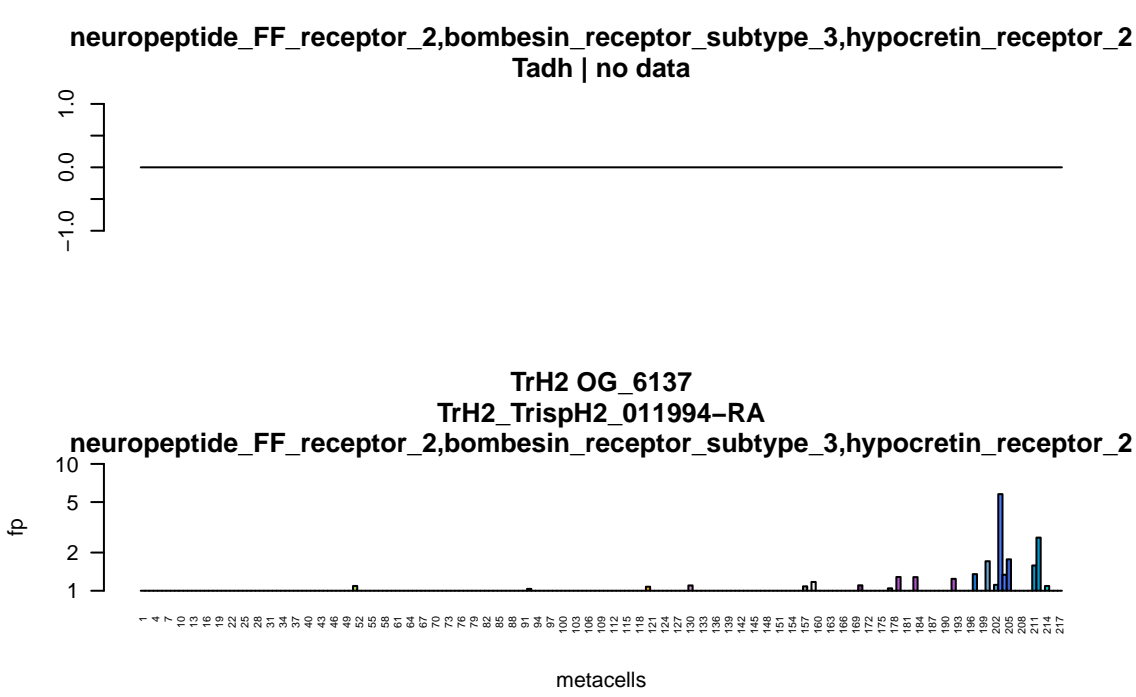


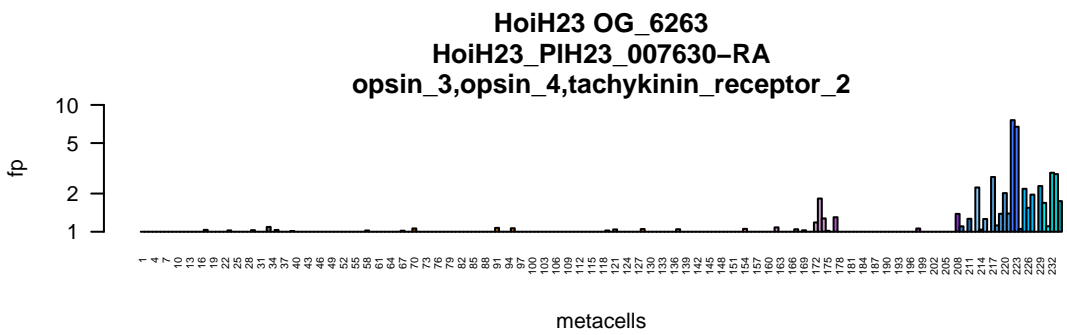
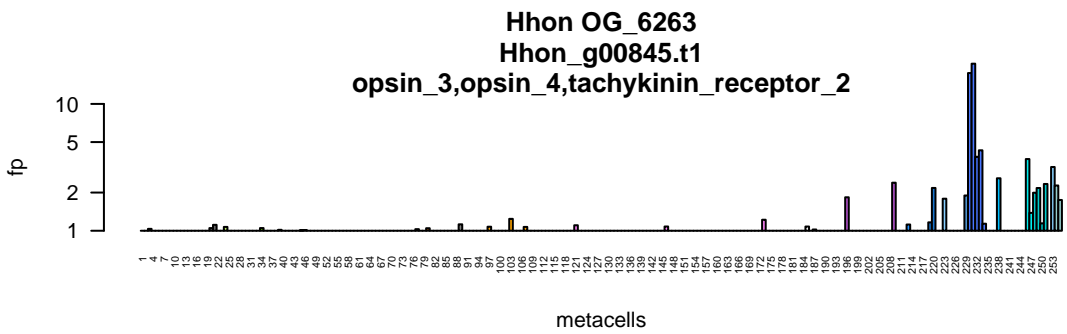
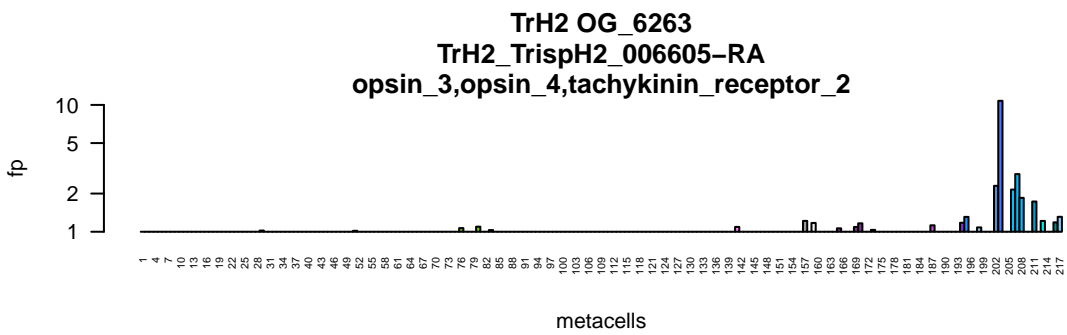
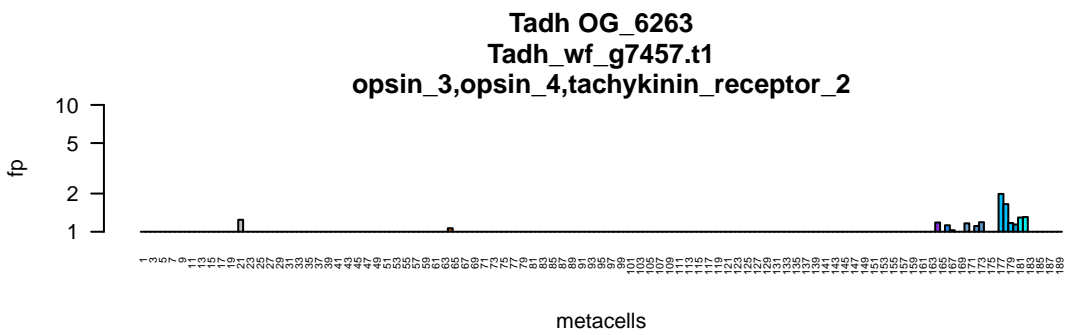
Hhon OG_5725
Hhon_g10923.t1

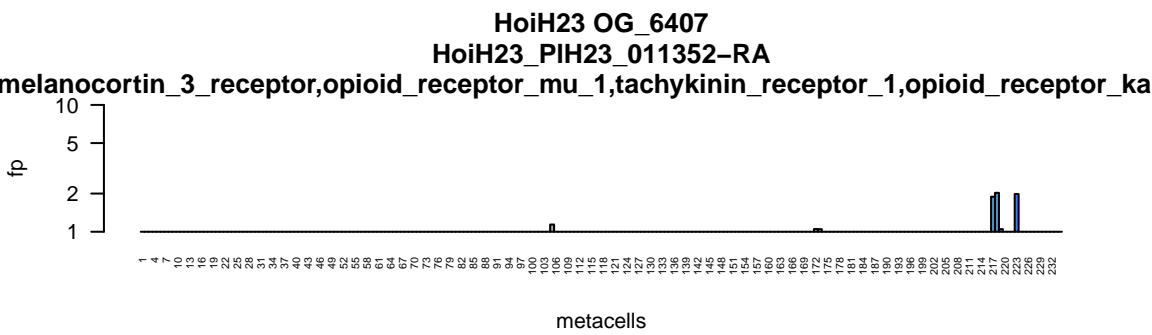
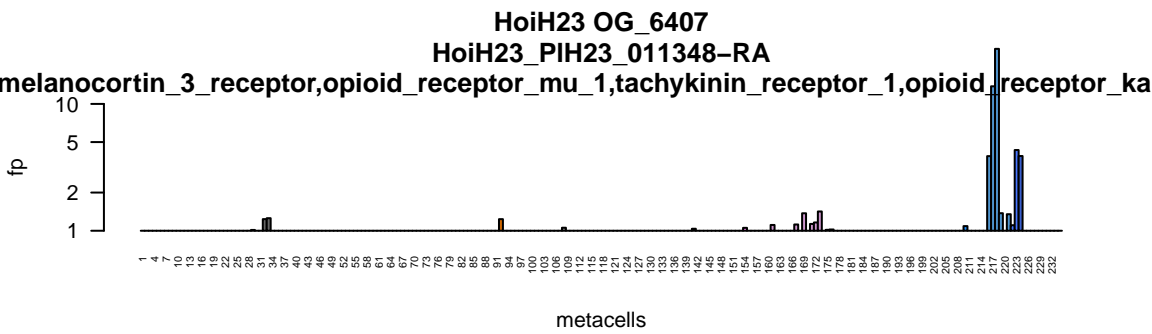
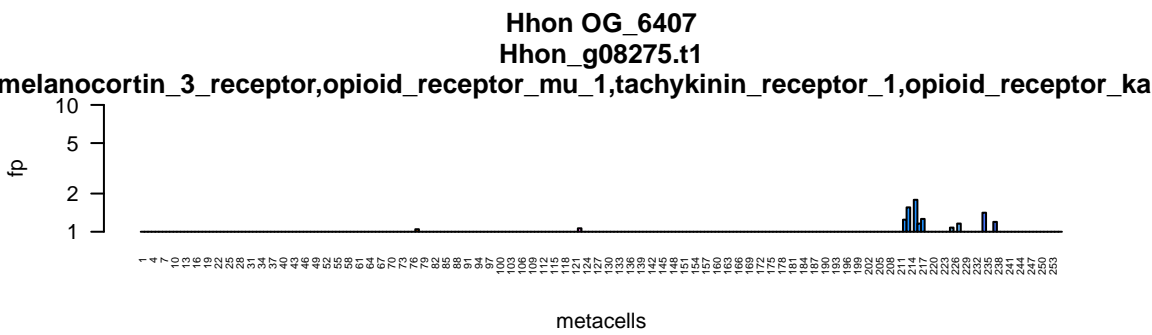
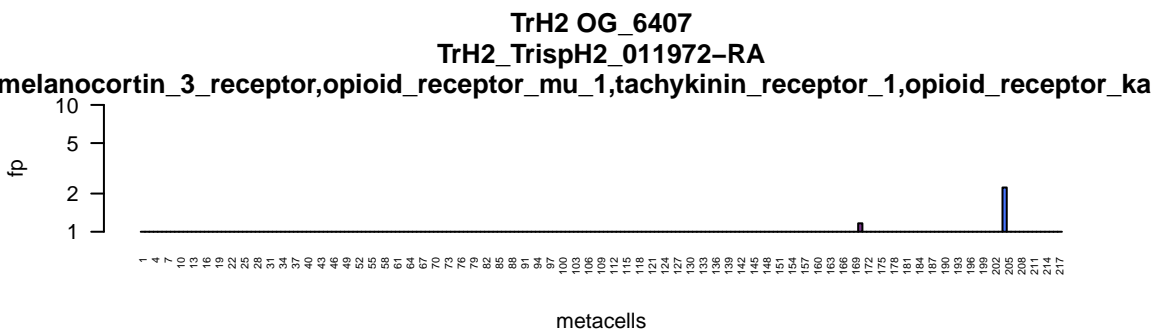
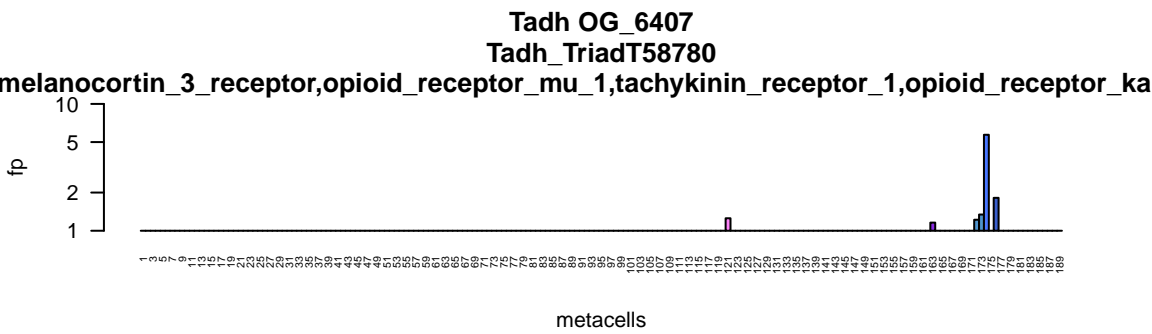


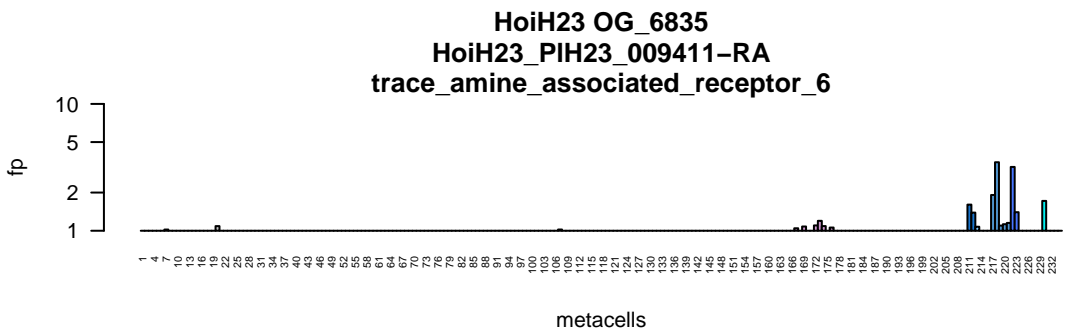
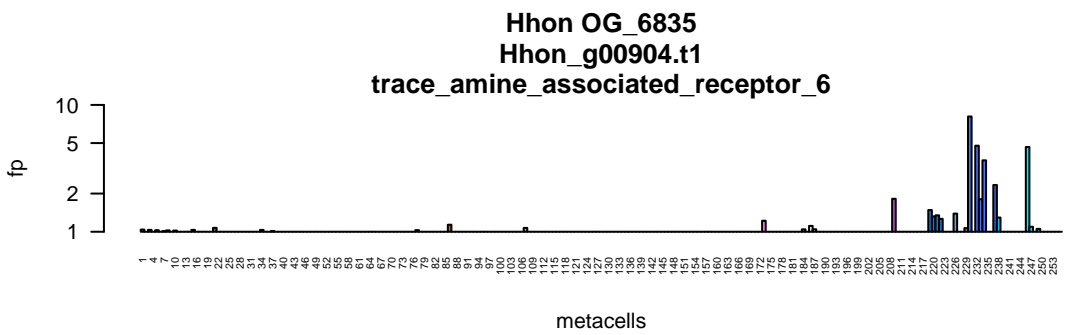
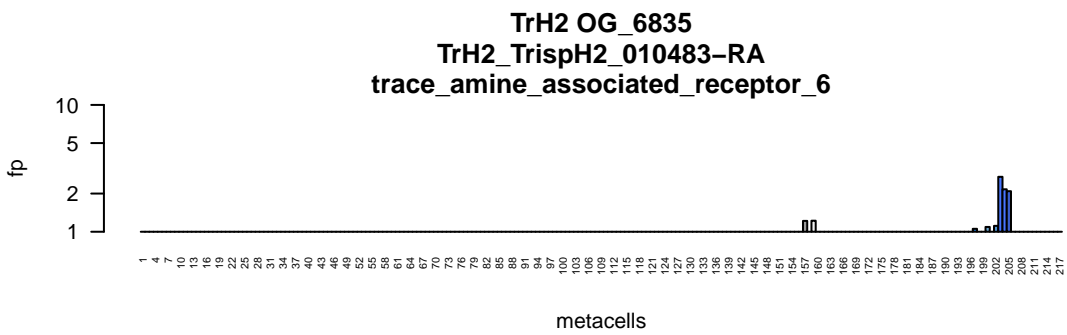
Hhon OG_5725
Hhon_g08418.t1

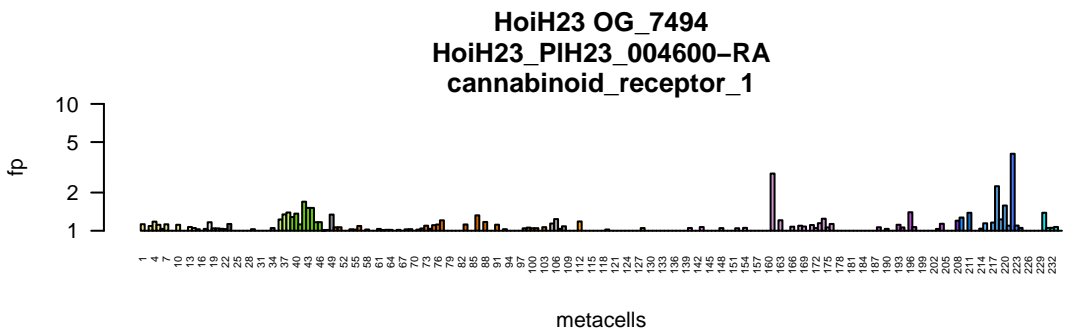
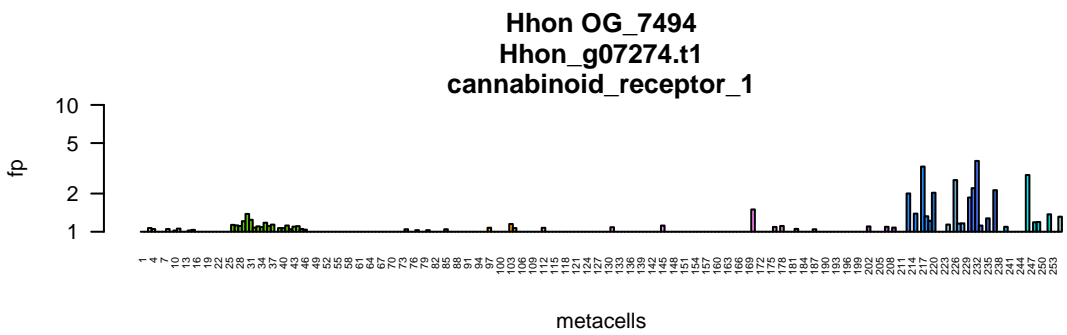
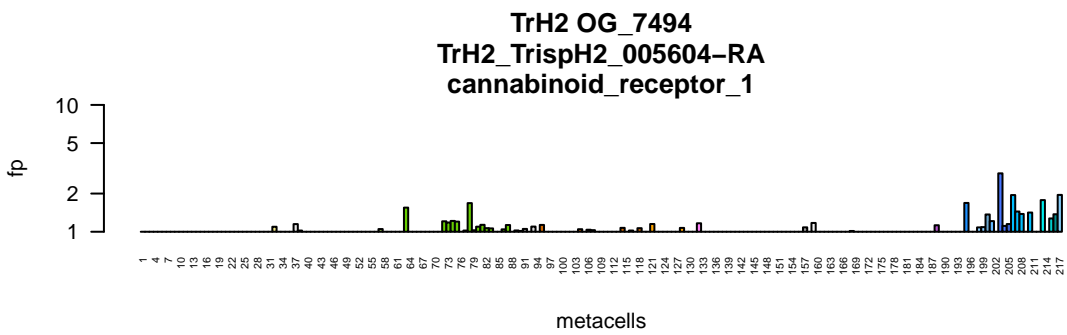
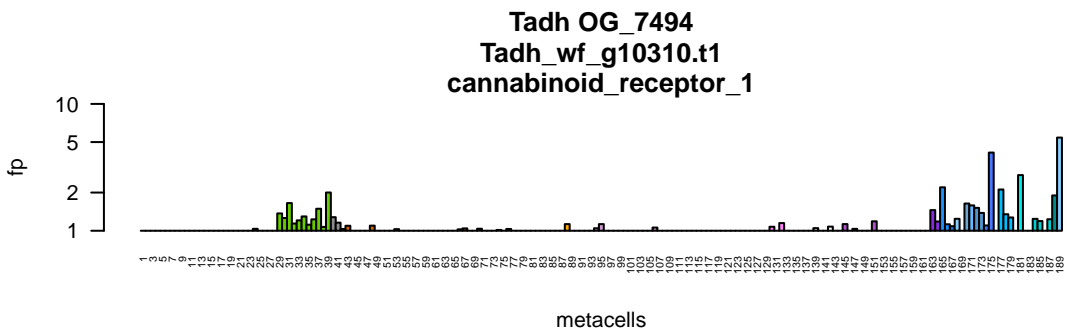


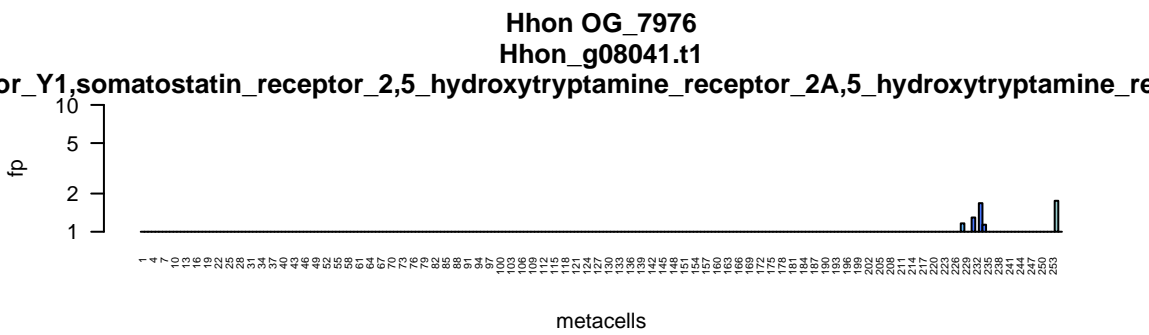
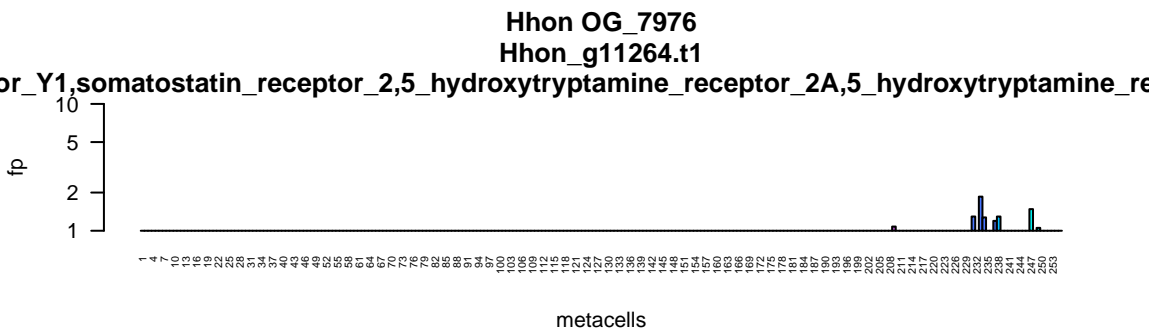
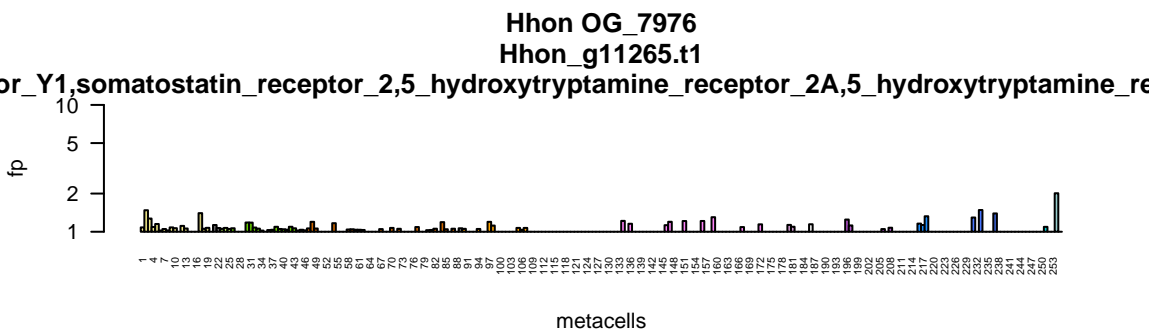
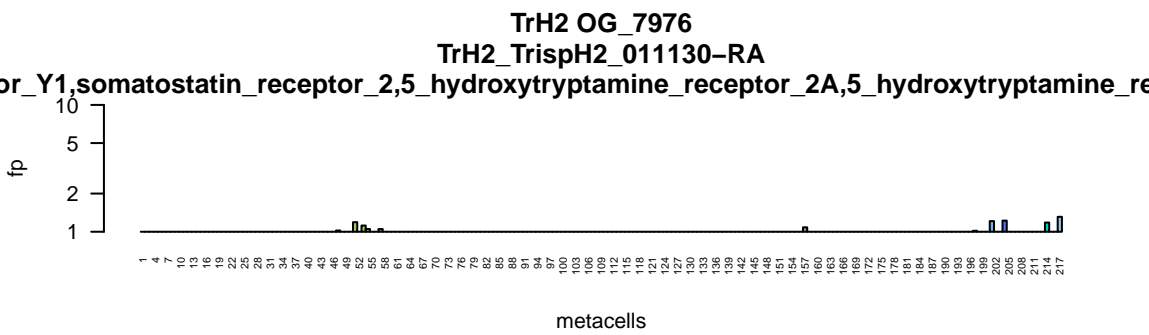
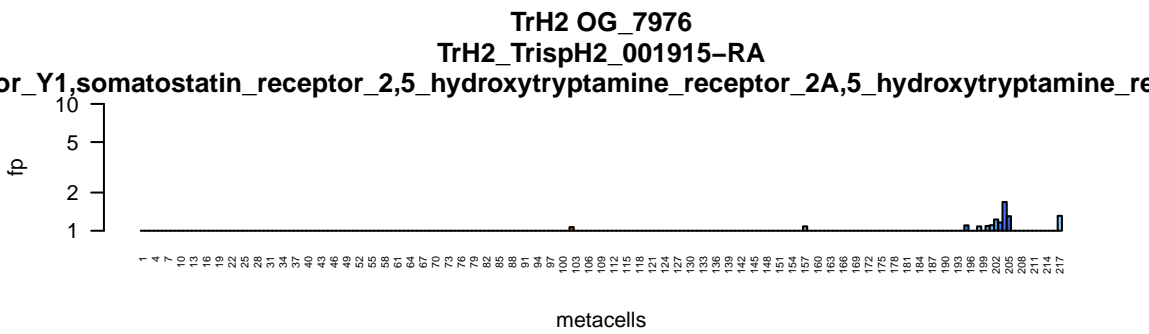
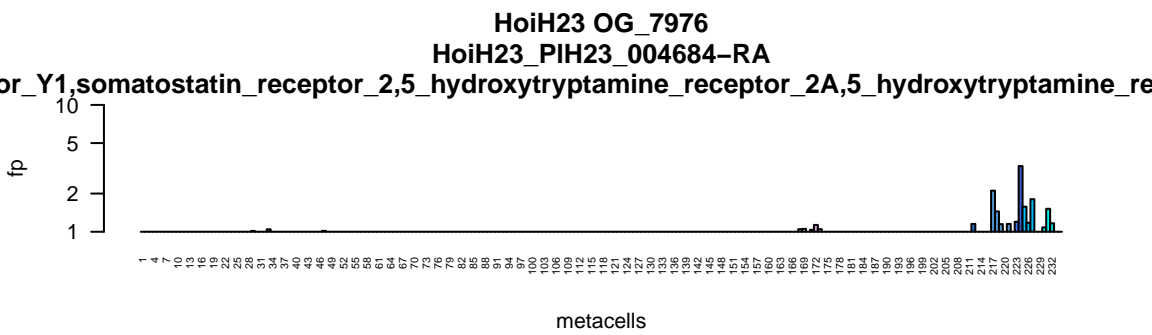
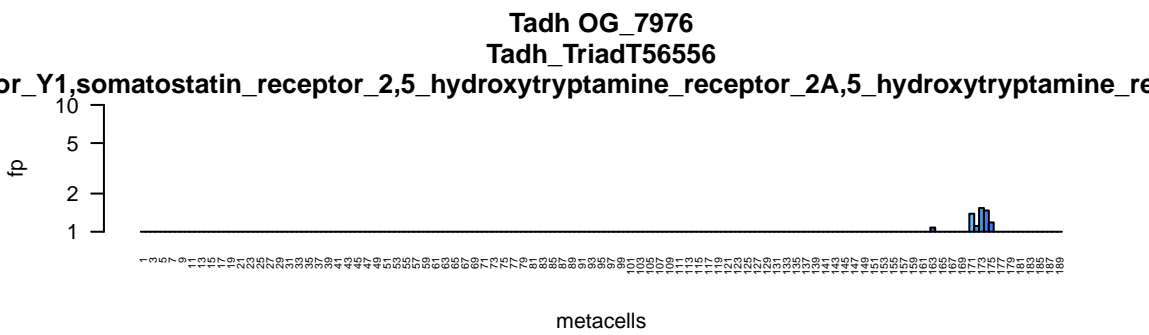
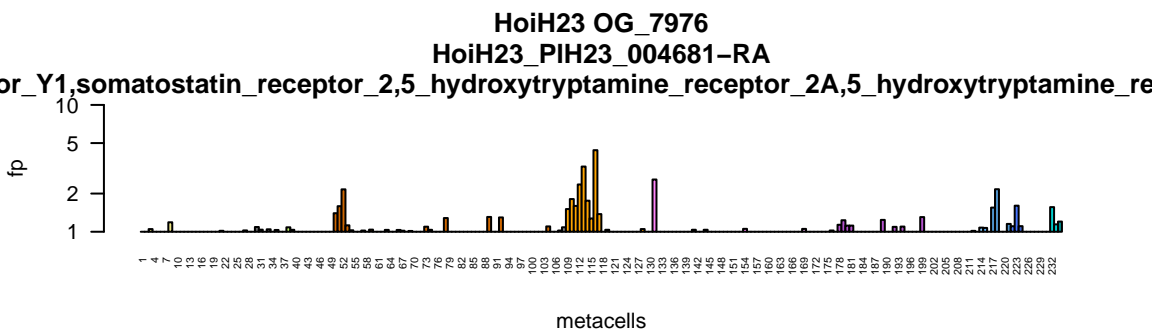
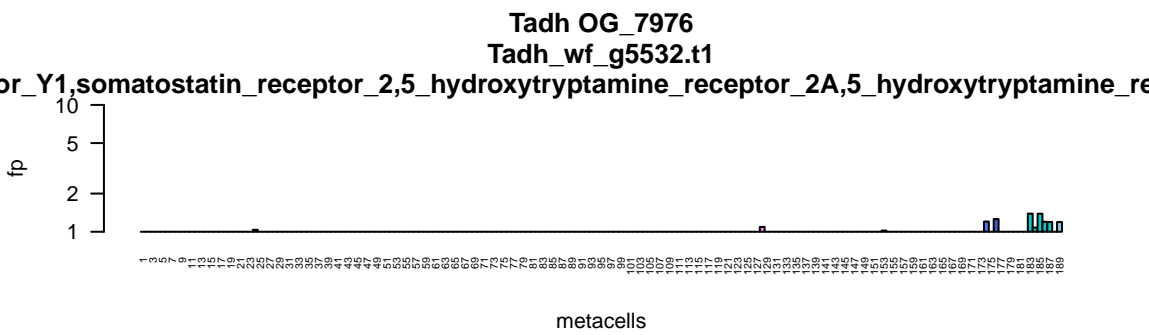
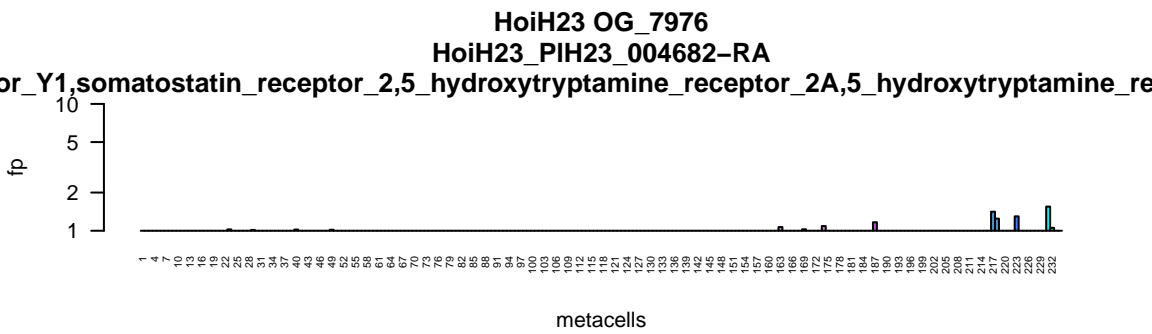
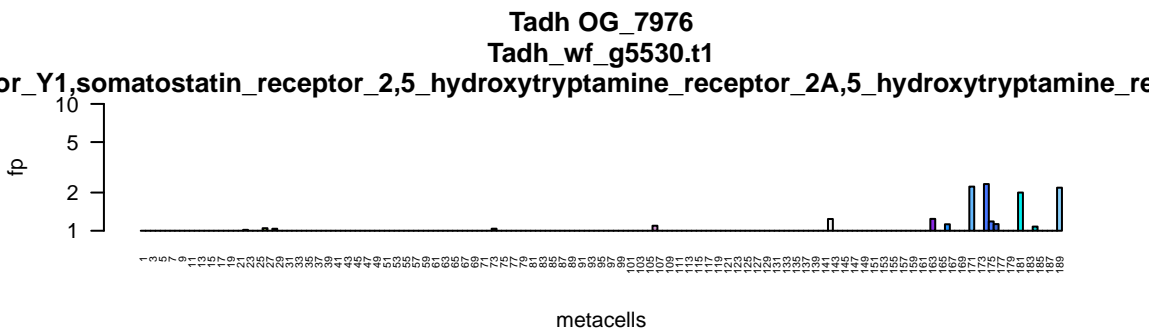


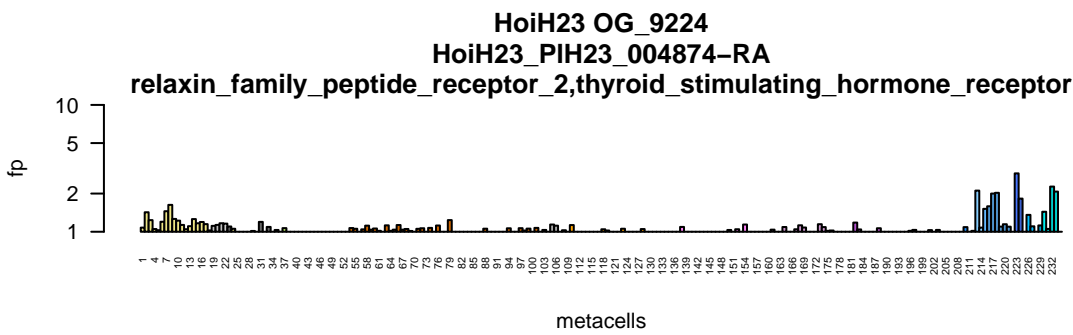
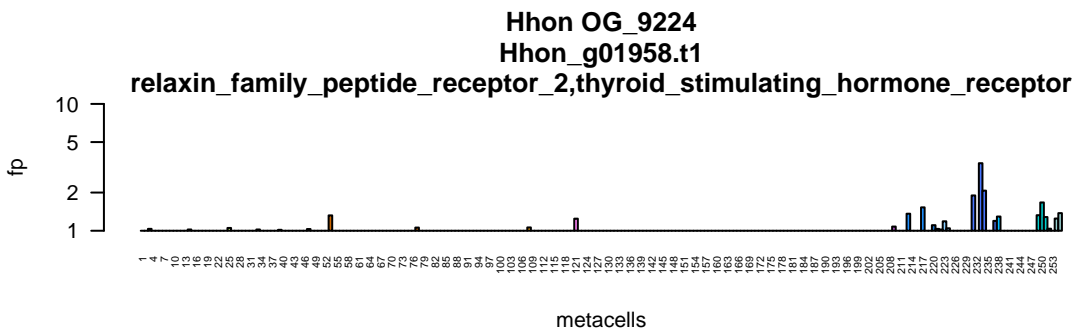
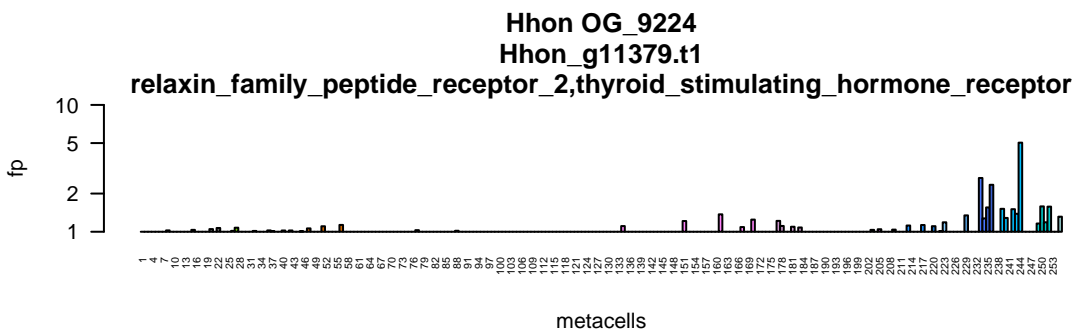
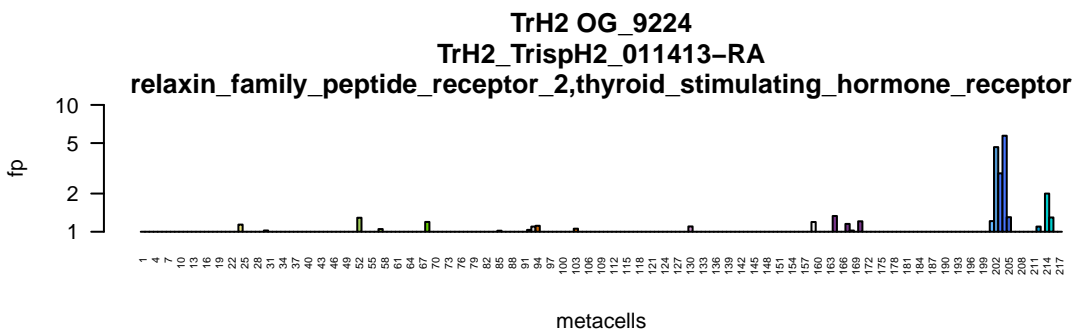
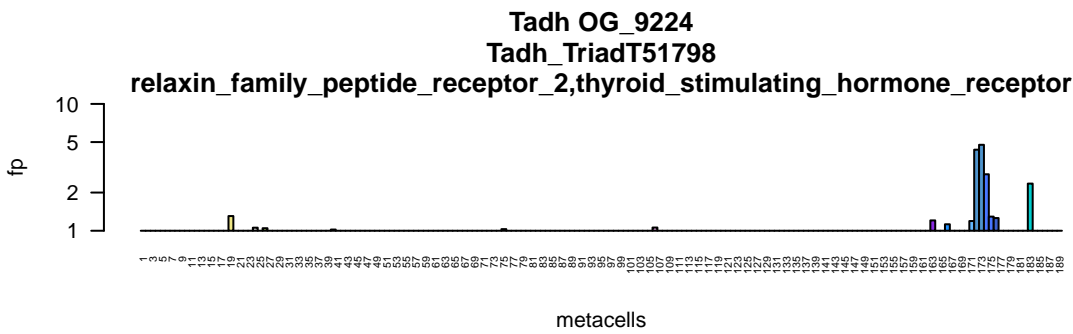


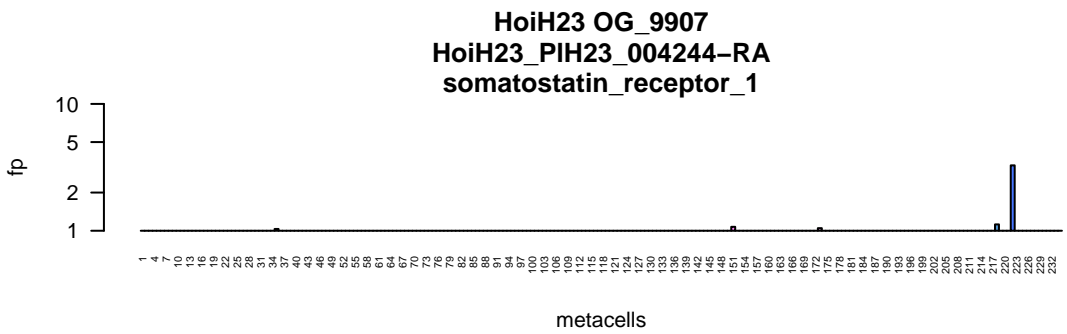
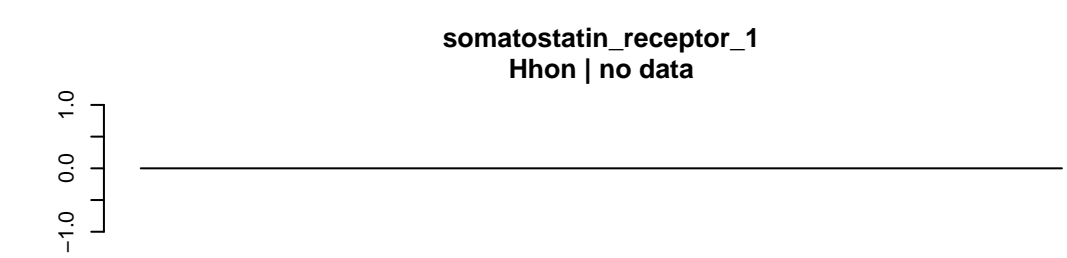
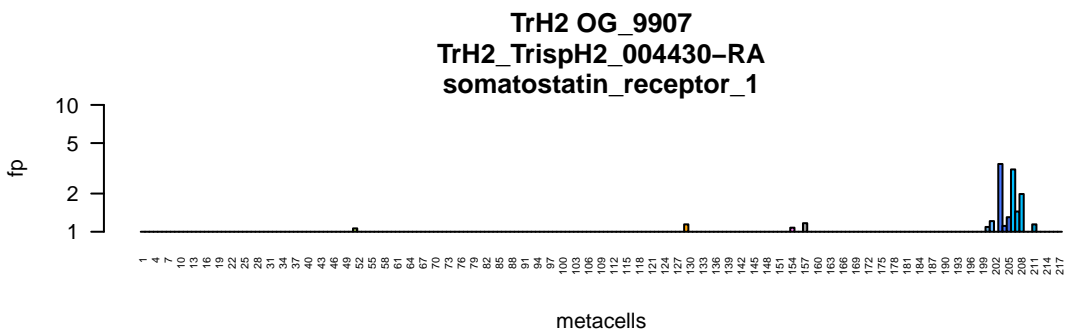
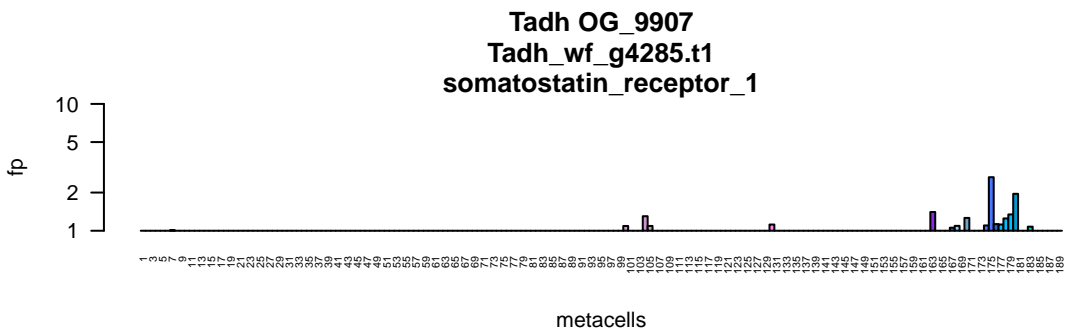


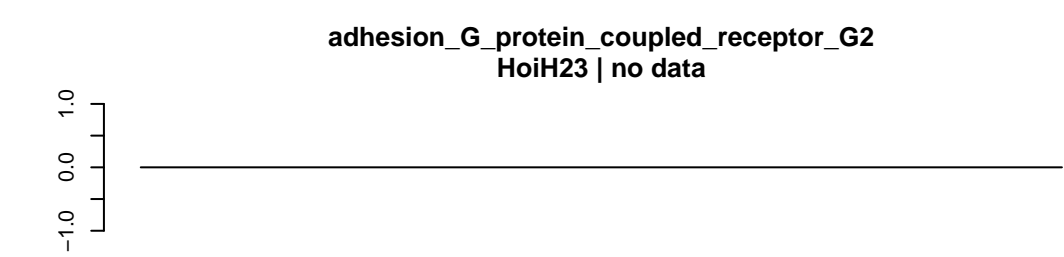
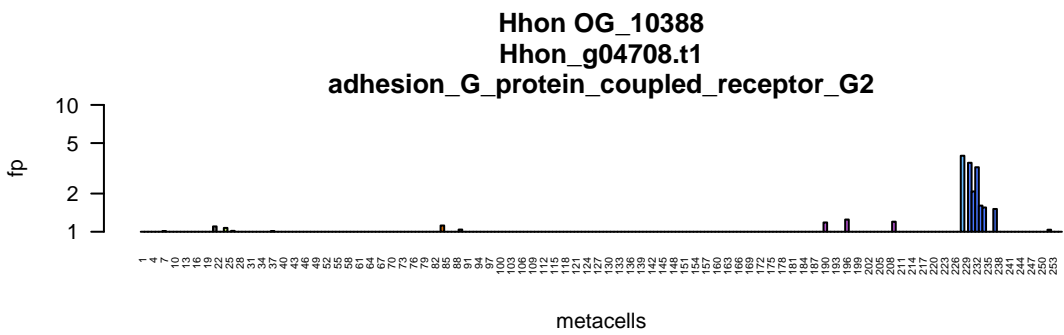
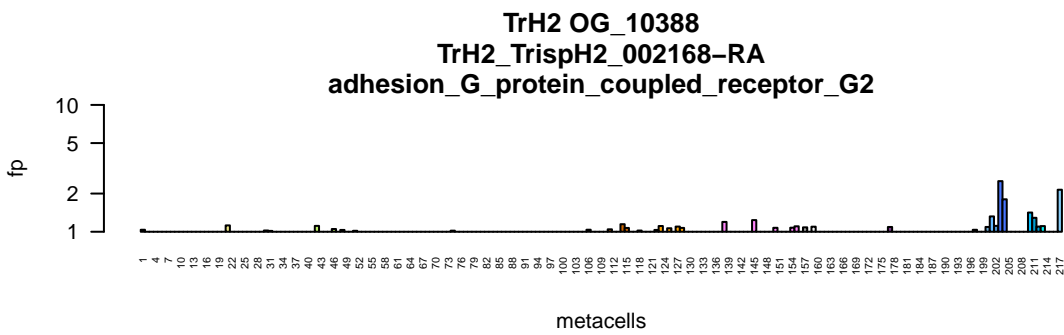
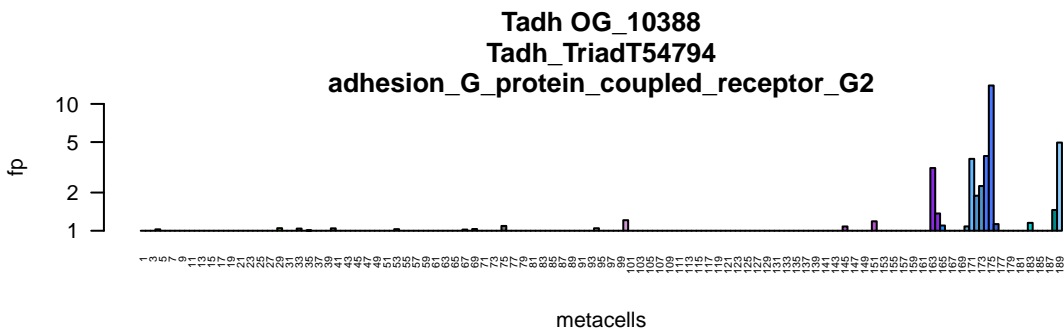




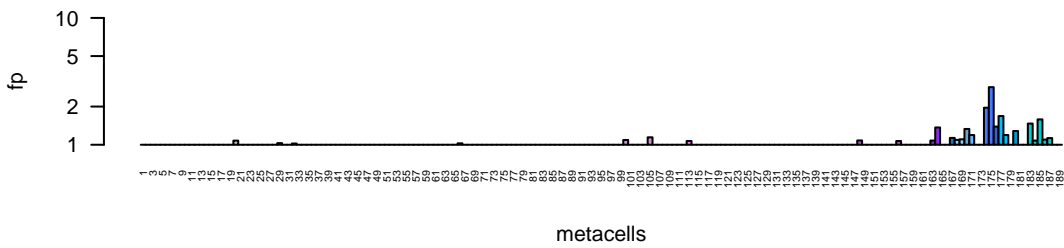




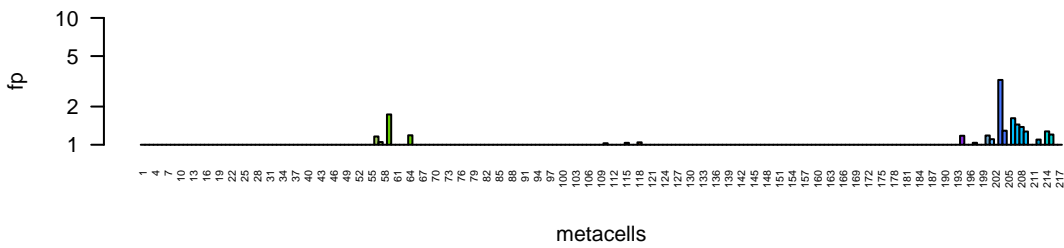




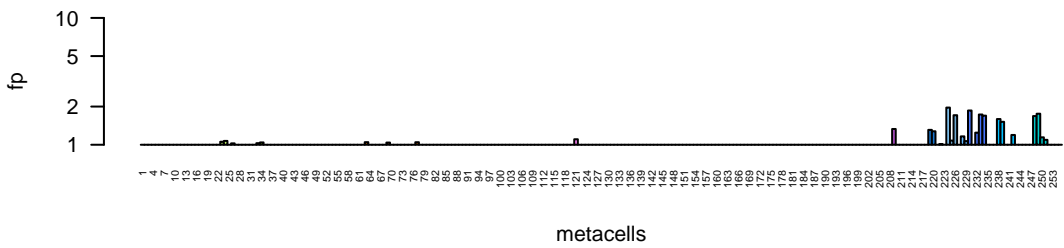
Tadh OG_515
Tadh_TriadT57467



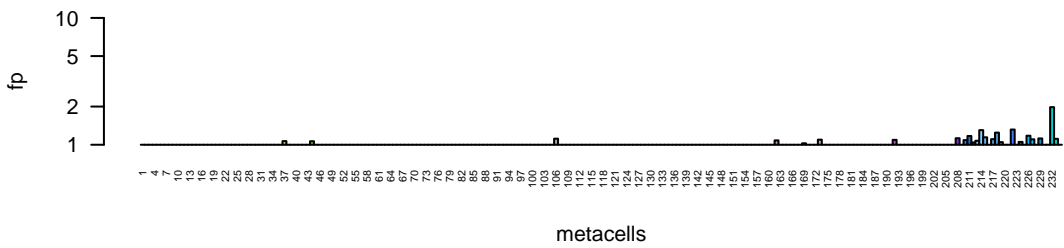
TrH2 OG_515
TrH2_TrispH2_007315-RA

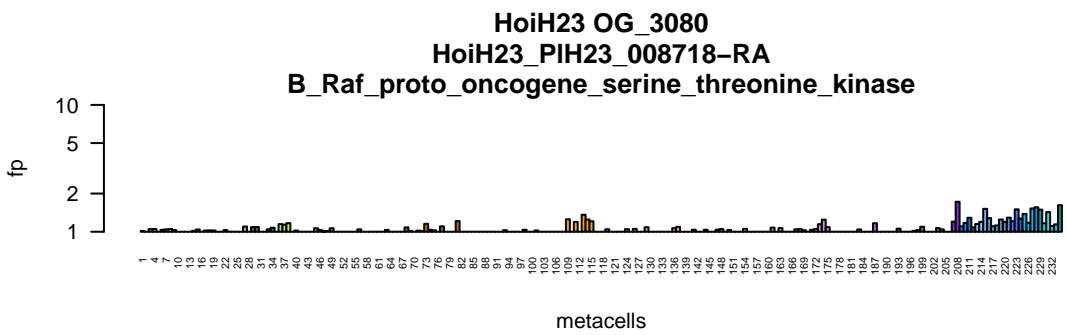
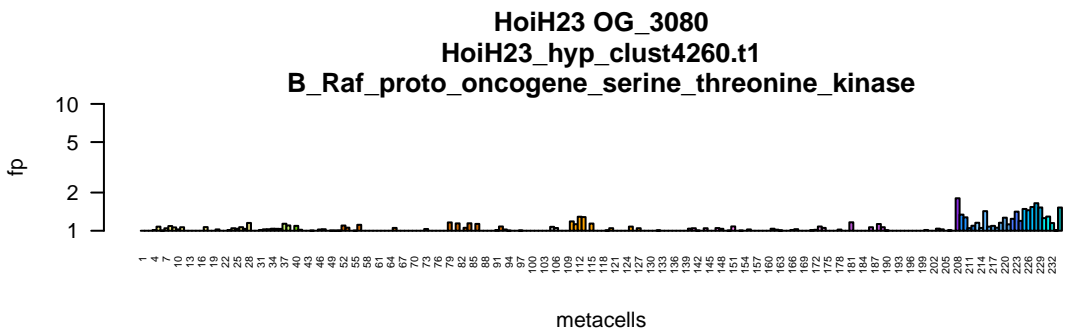
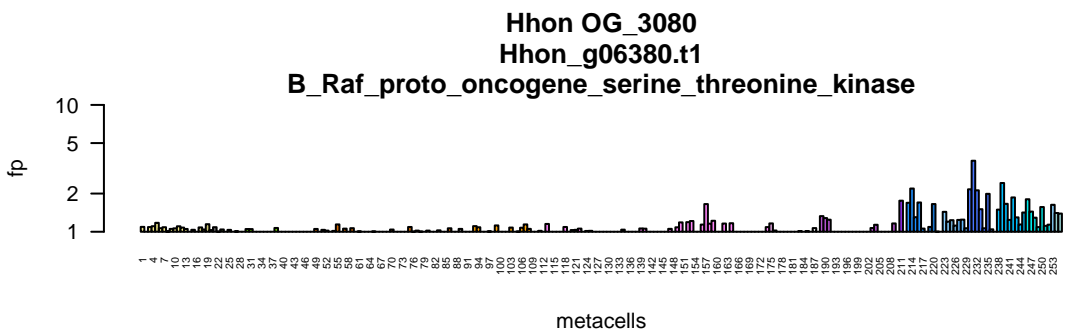
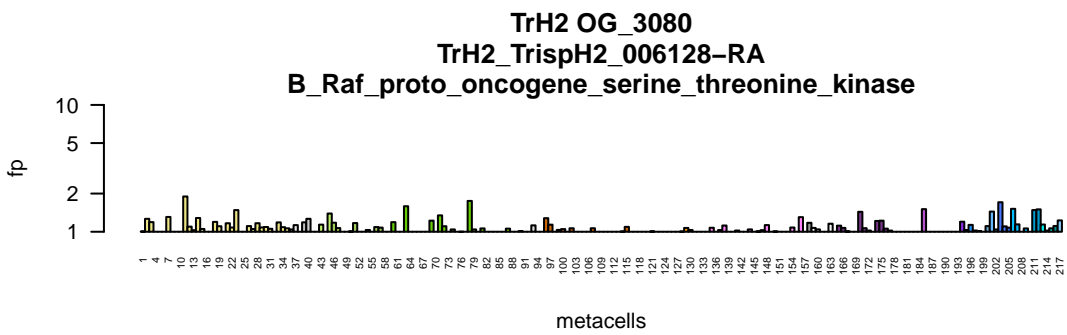
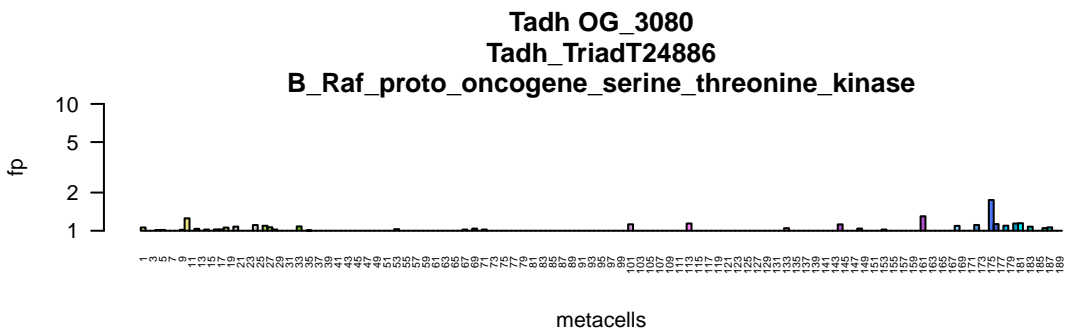


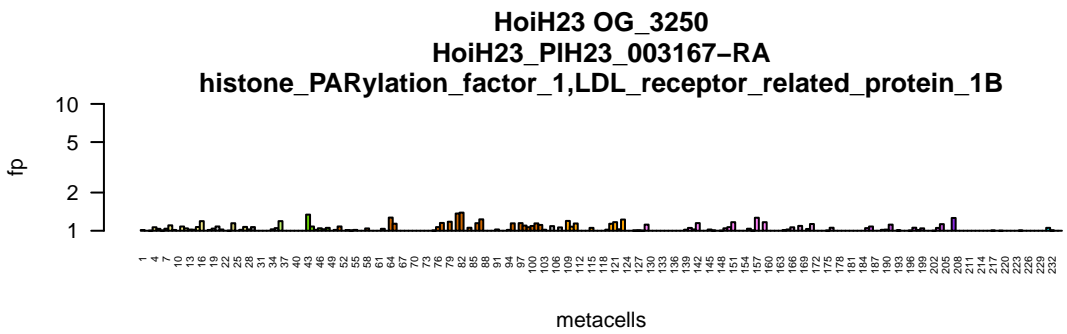
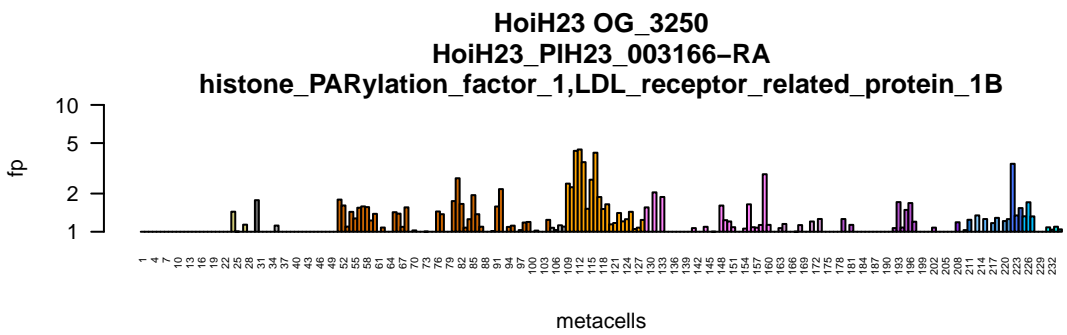
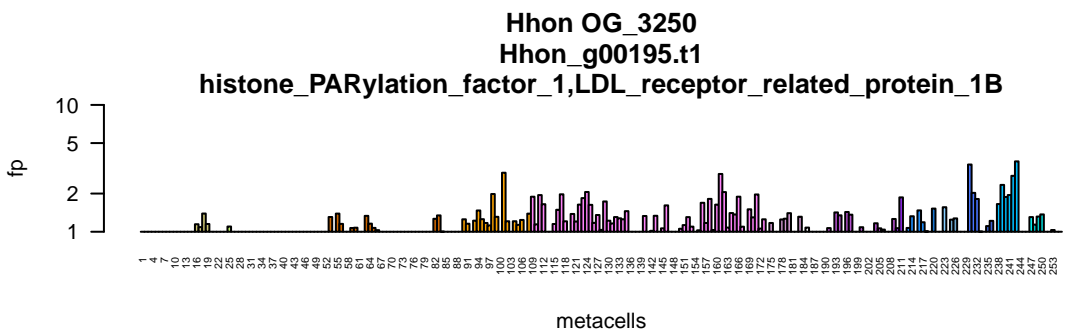
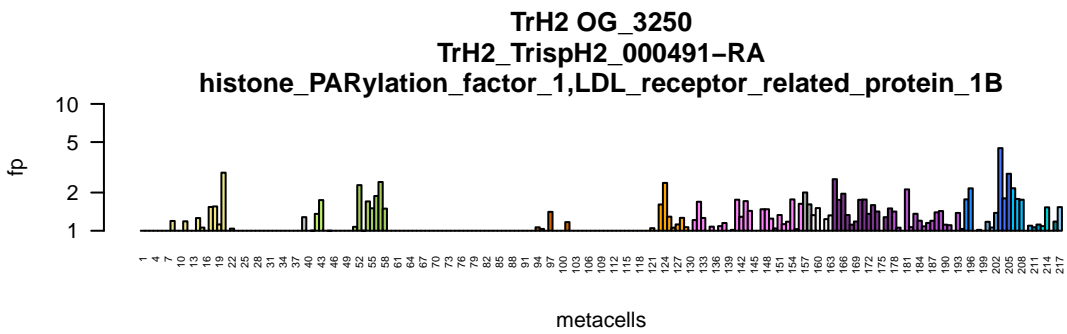
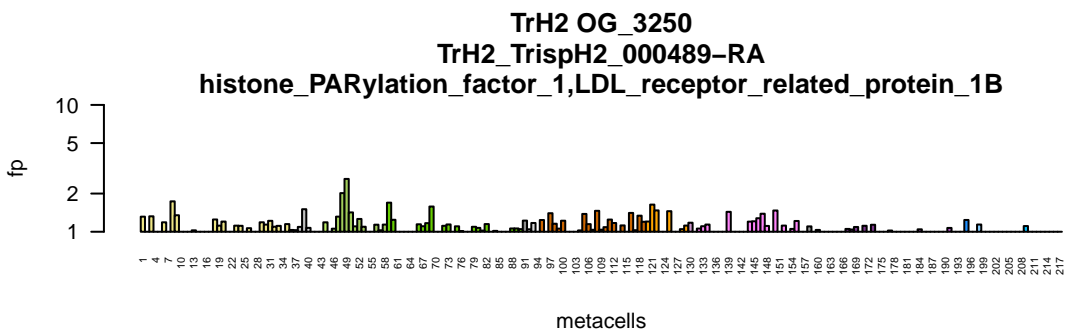
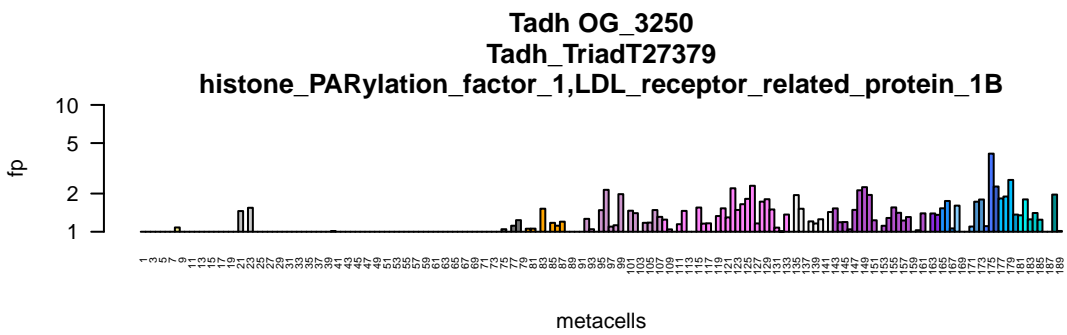
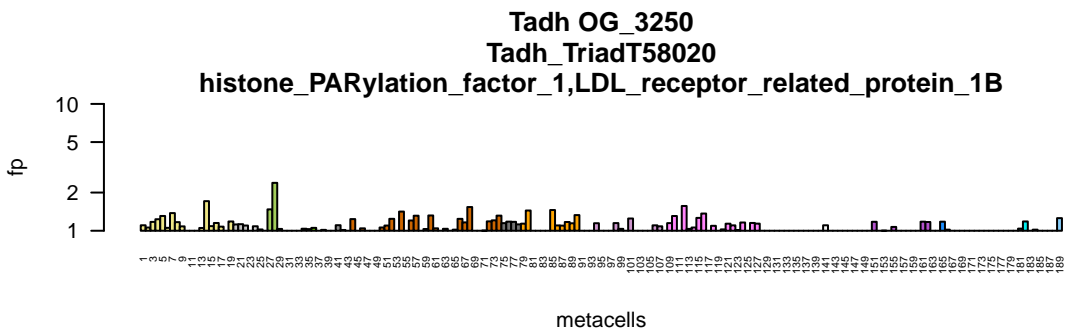
Hhon OG_515
Hhon_g01896.t1



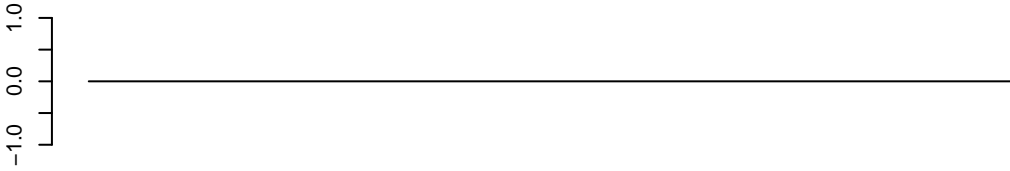
HoiH23 OG_515
HoiH23_PIH23_001462-RA



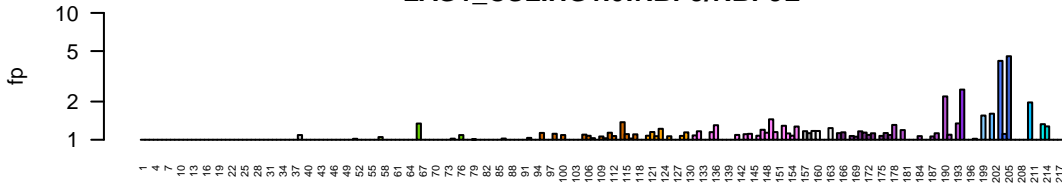




LAG1_CSL.HG1.0:RBPJ/RBPJL
Tadh | no data

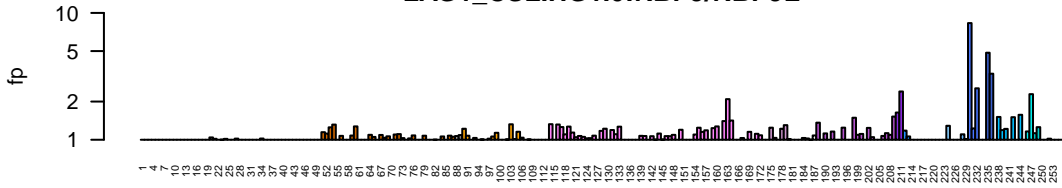


TrH2 OG_3606
TrH2_TrispH2_009810-RA
LAG1_CSL.HG1.0:RBPJ/RBPJL



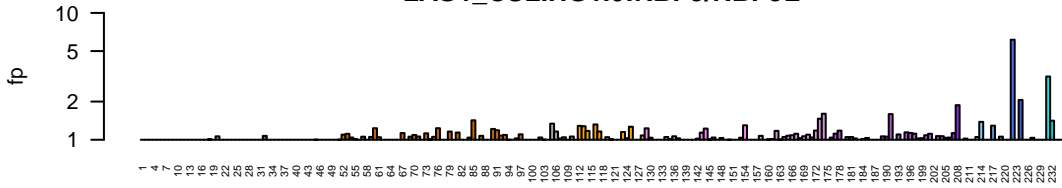
metacells

Hhon OG_3606
Hhon_g05183.t1
LAG1_CSL.HG1.0:RBPJ/RBPJL

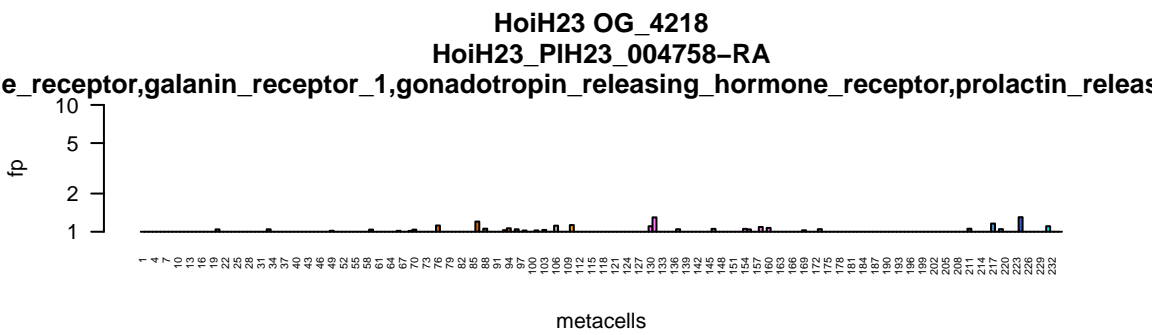
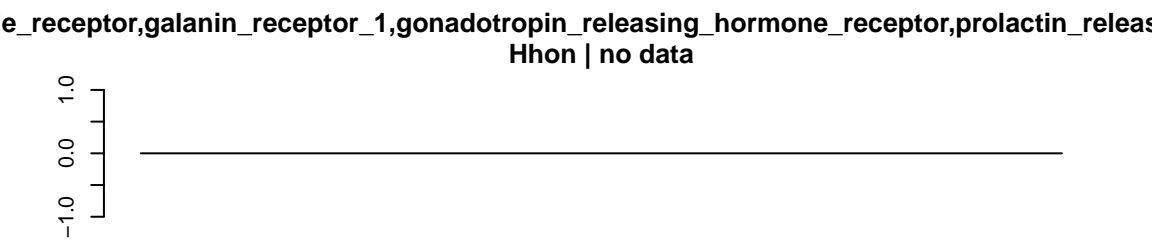
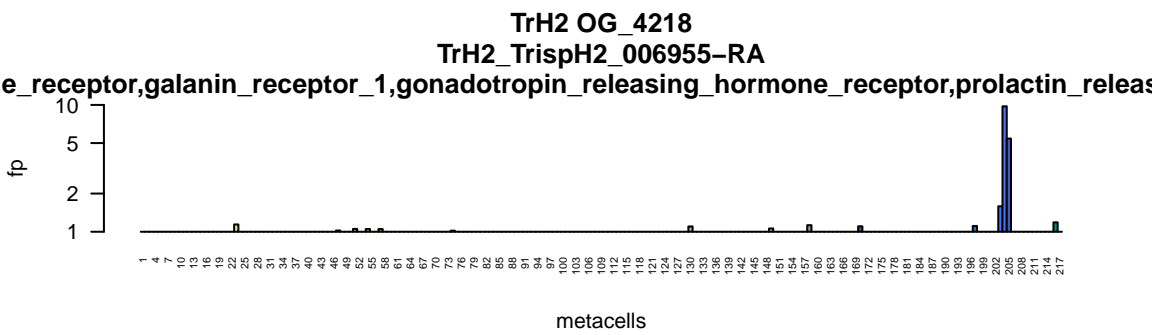
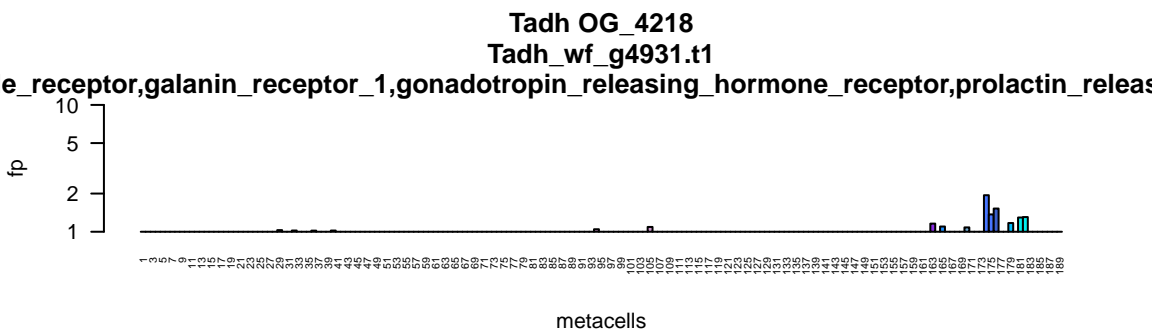
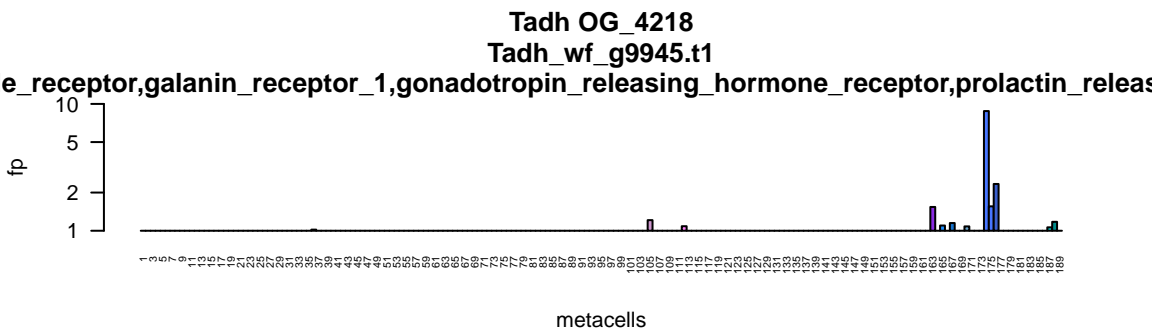


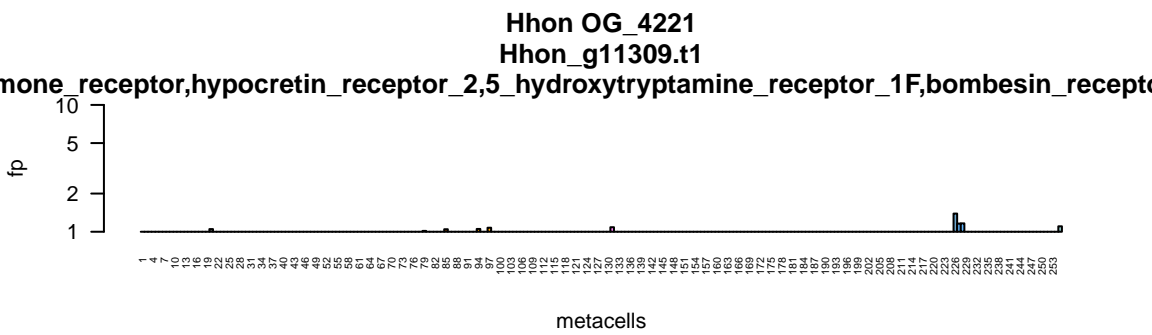
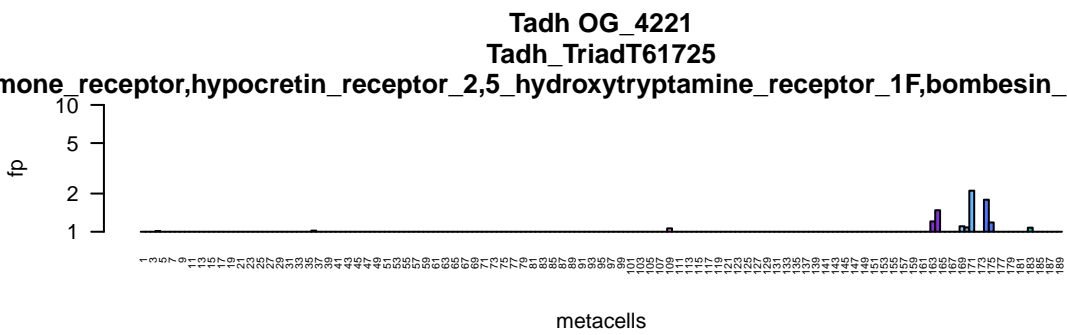
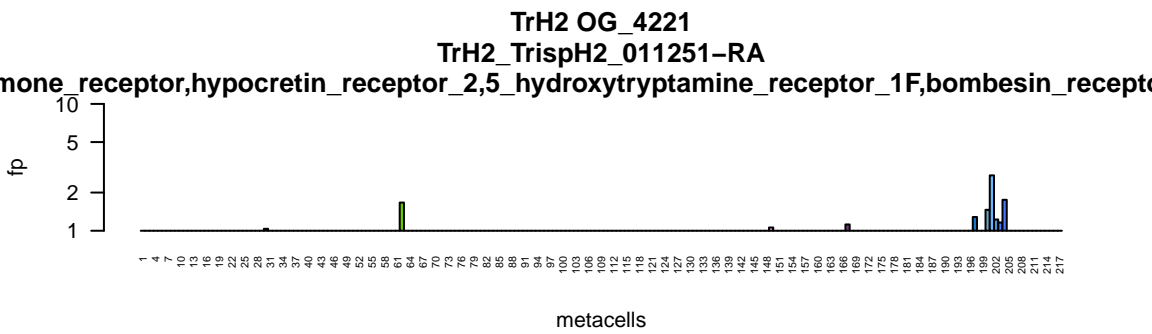
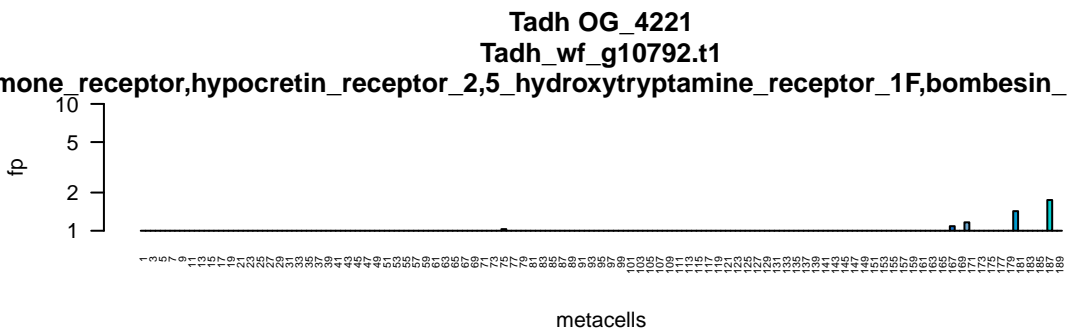
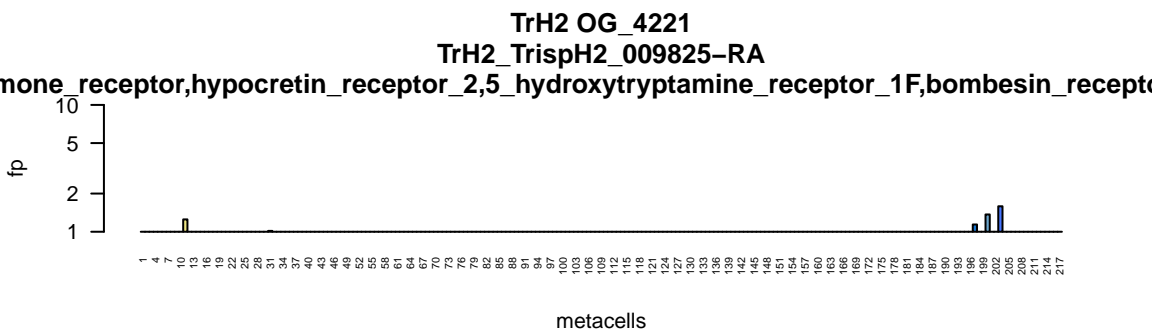
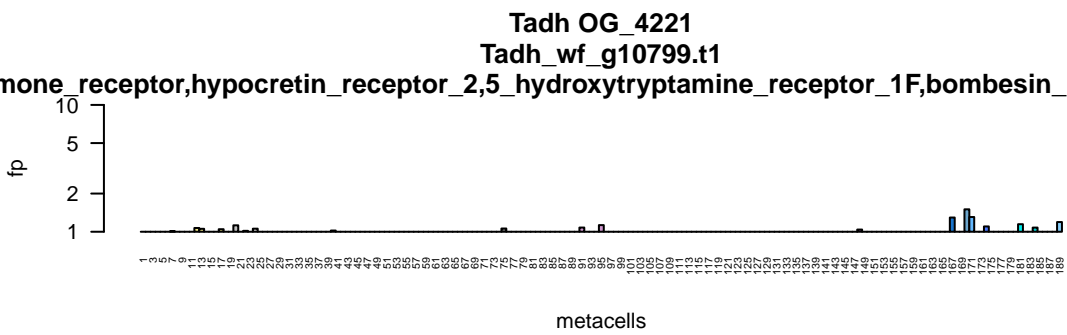
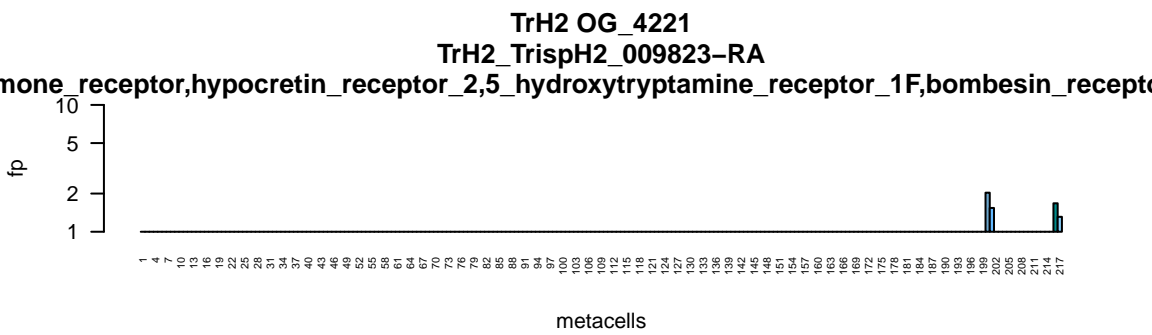
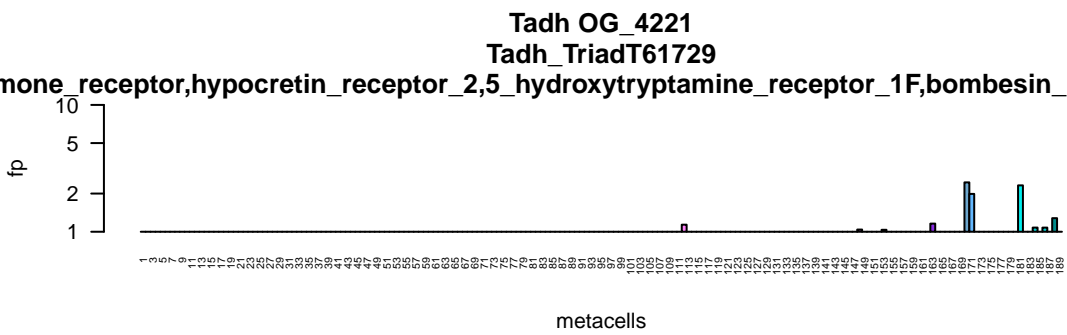
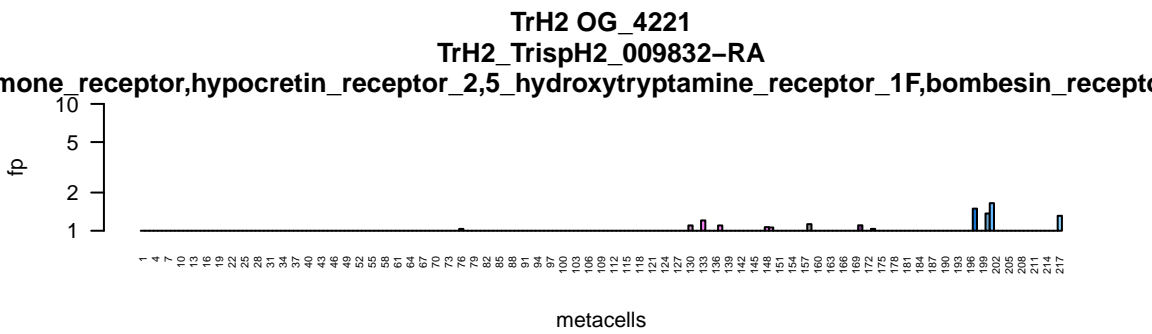
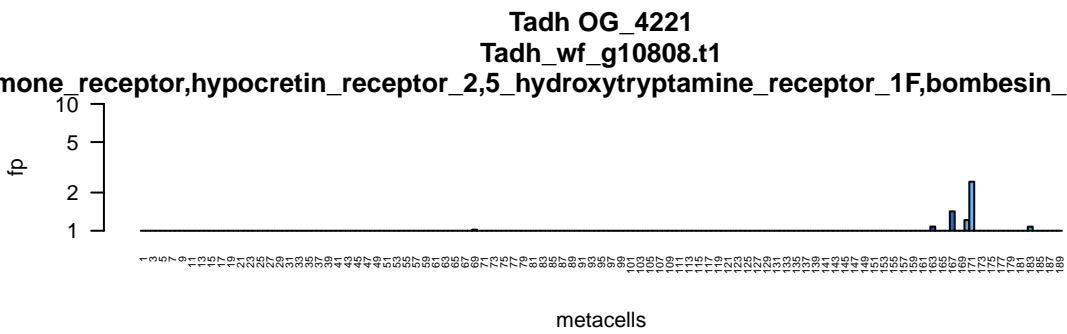
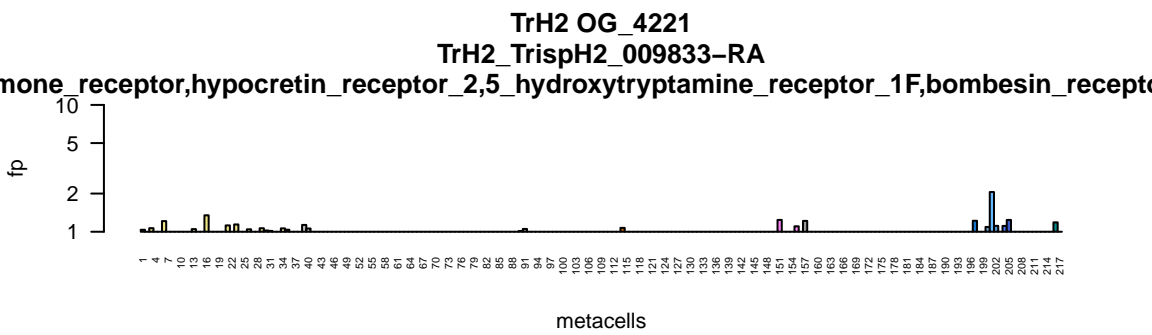
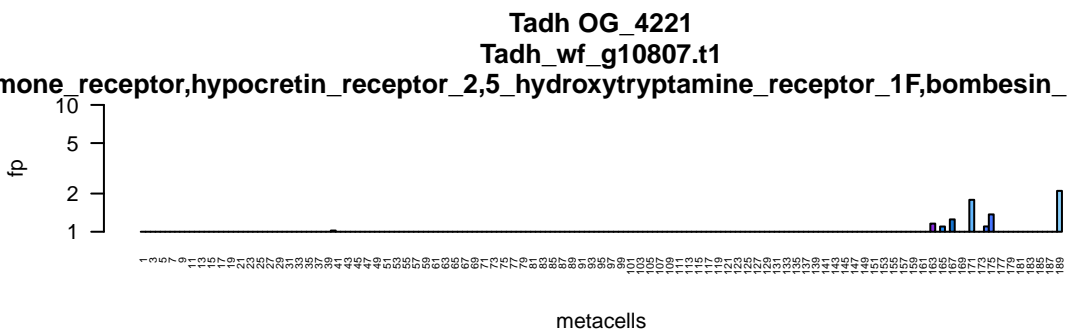
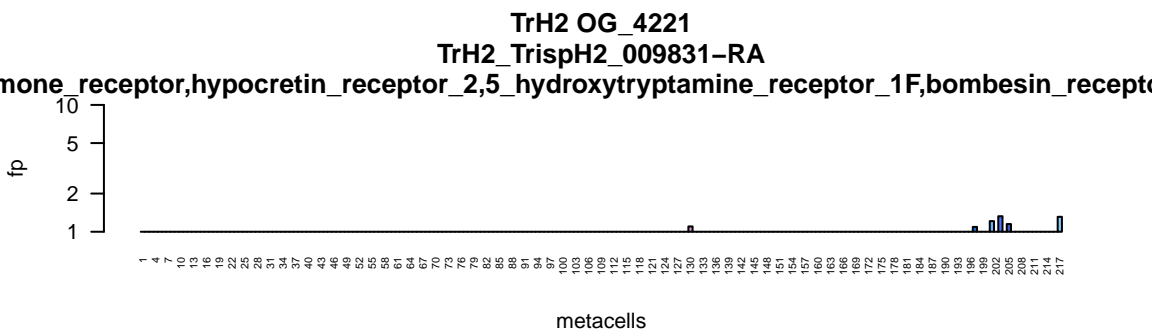
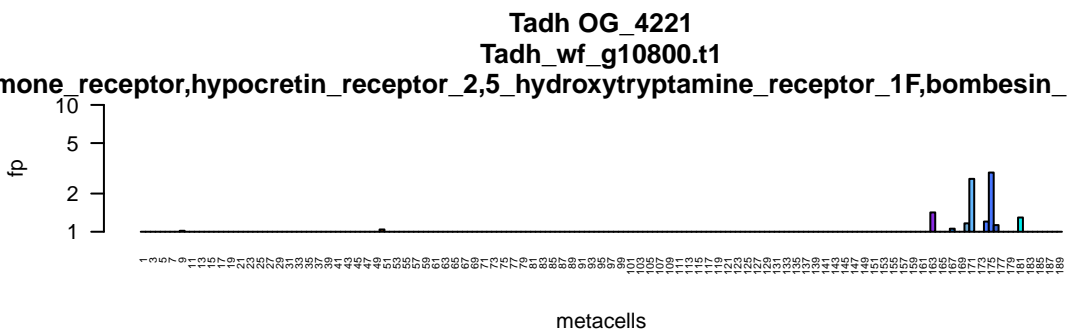
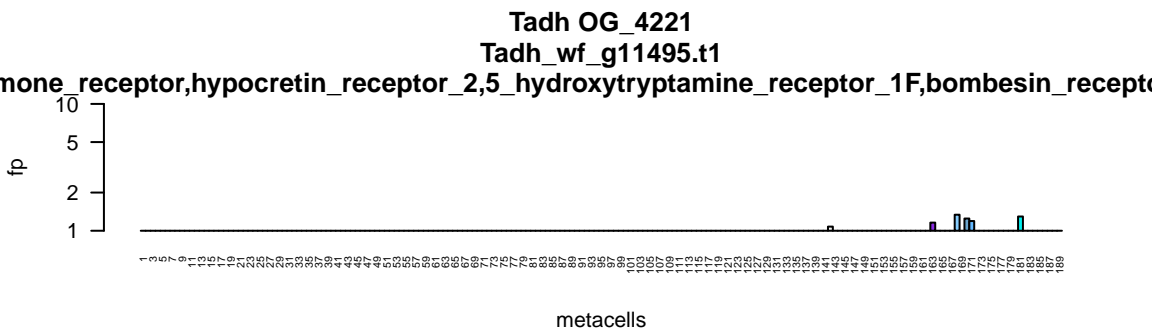
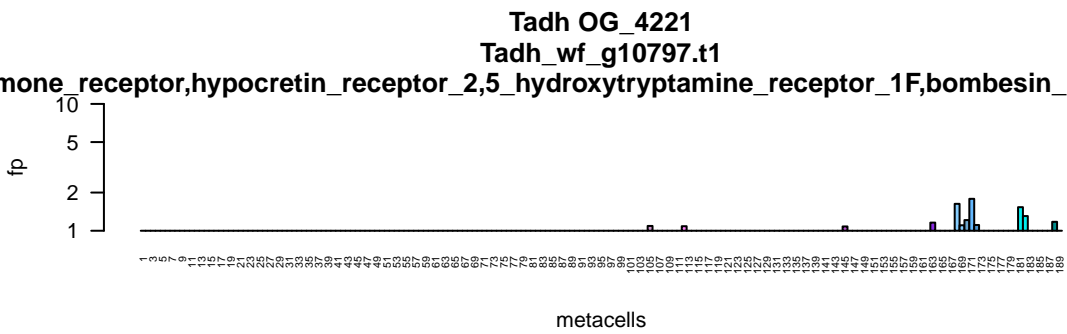
metacells

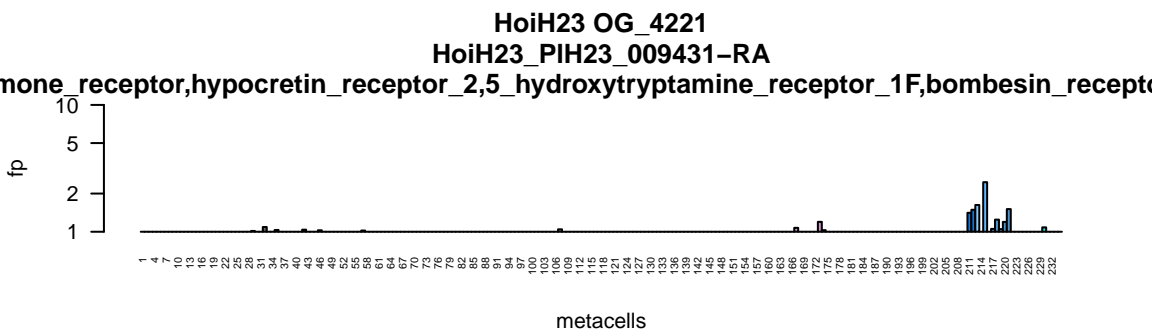
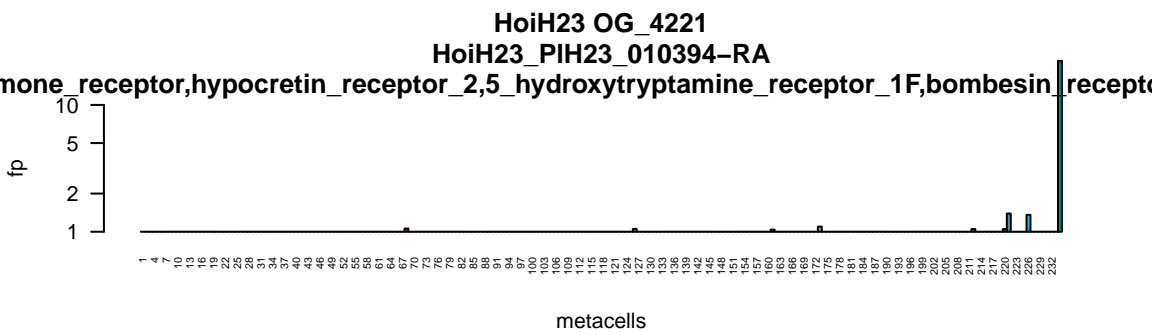
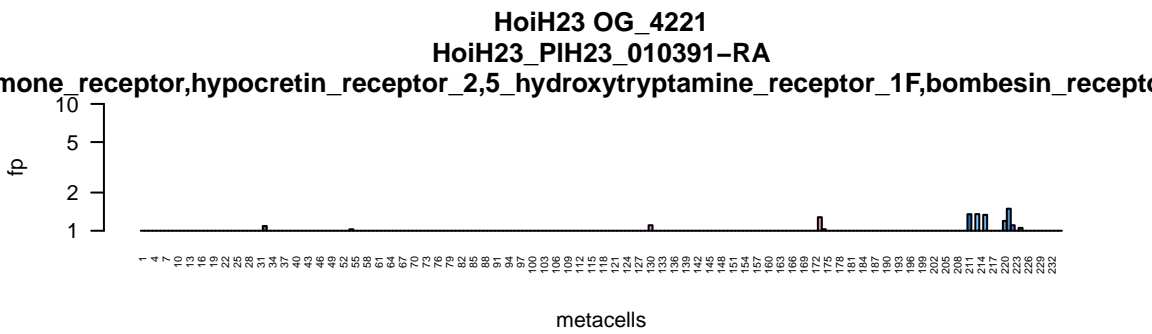
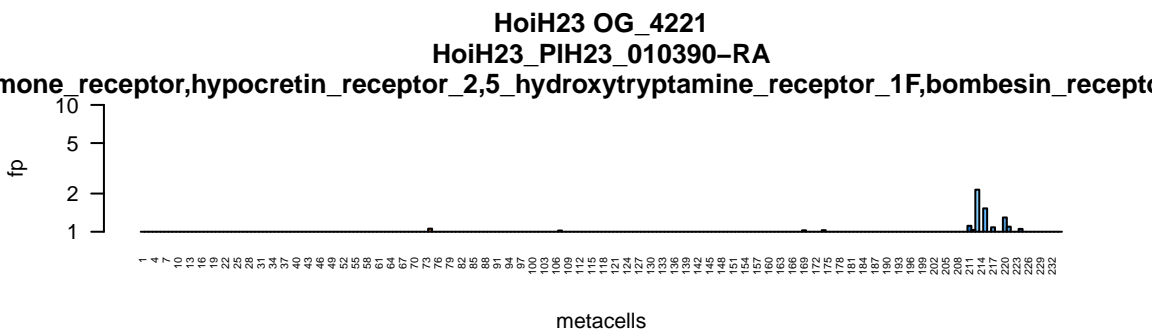
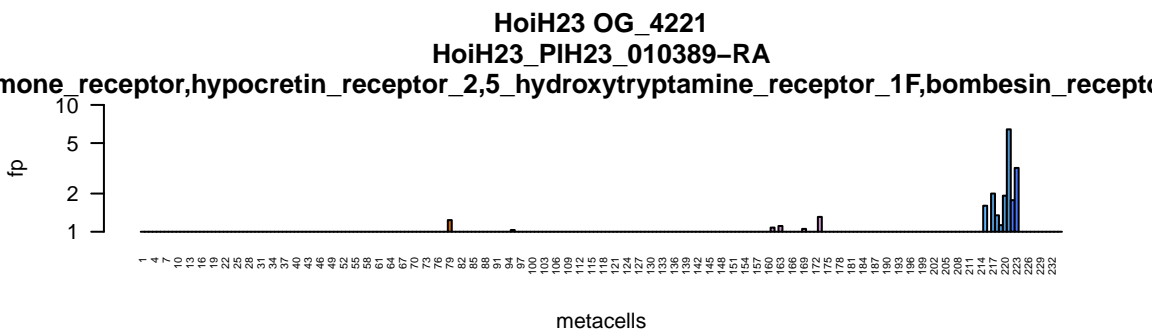
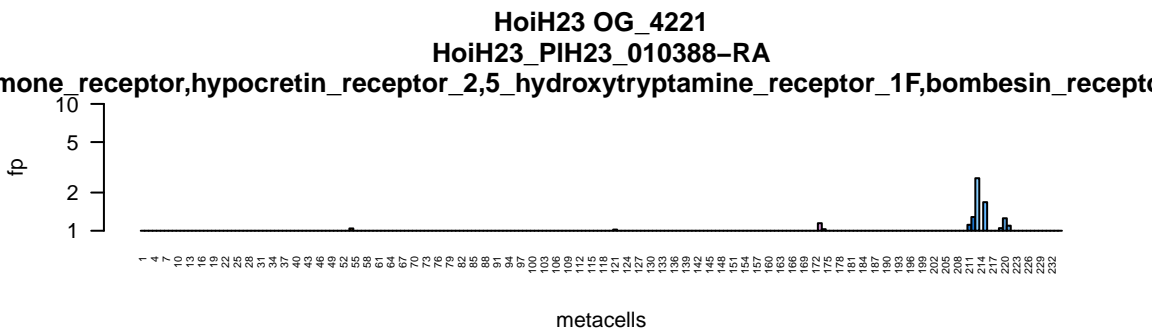
HoiH23 OG_3606
HoiH23_PIH23_001259-RA
LAG1_CSL.HG1.0:RBPJ/RBPJL

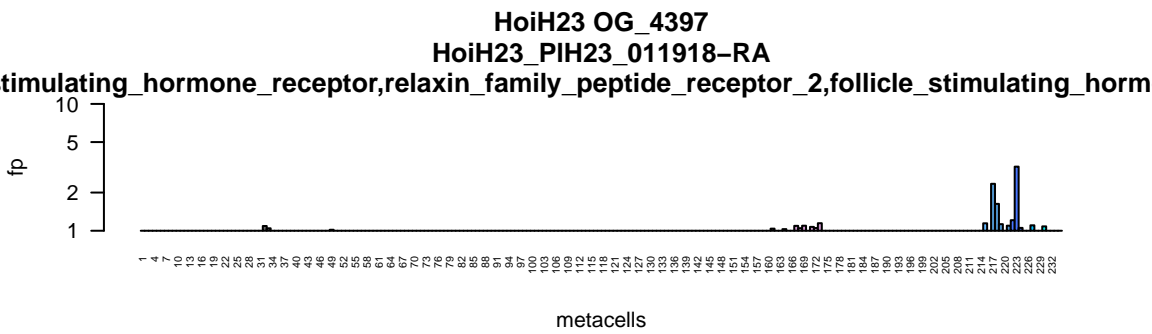
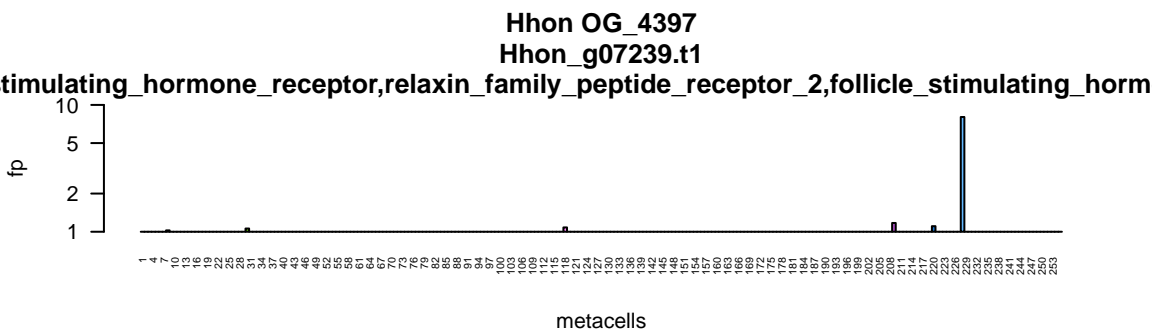
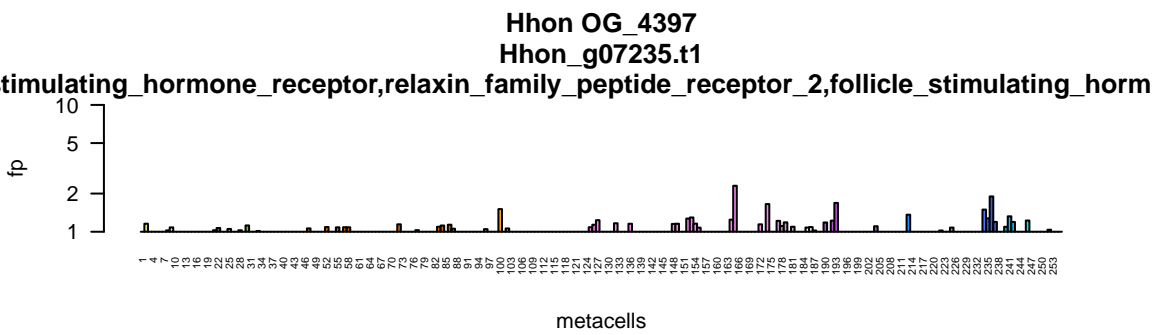
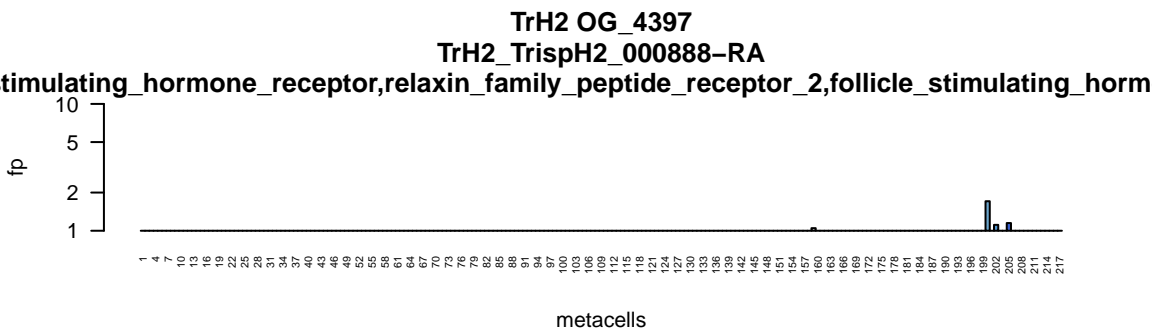
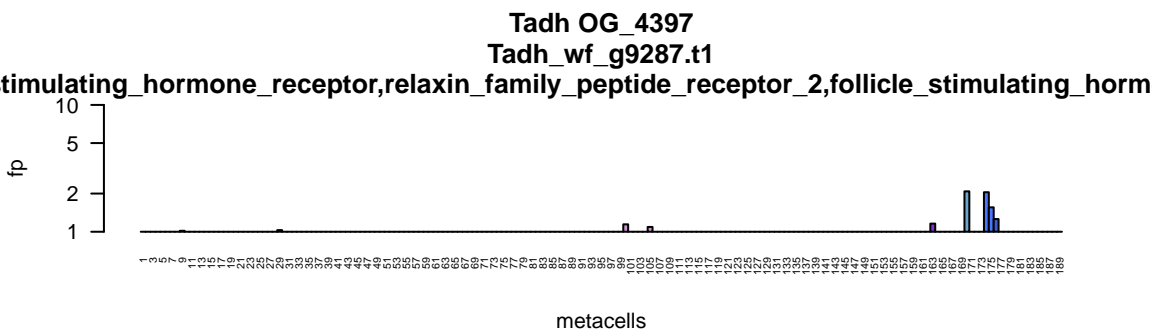
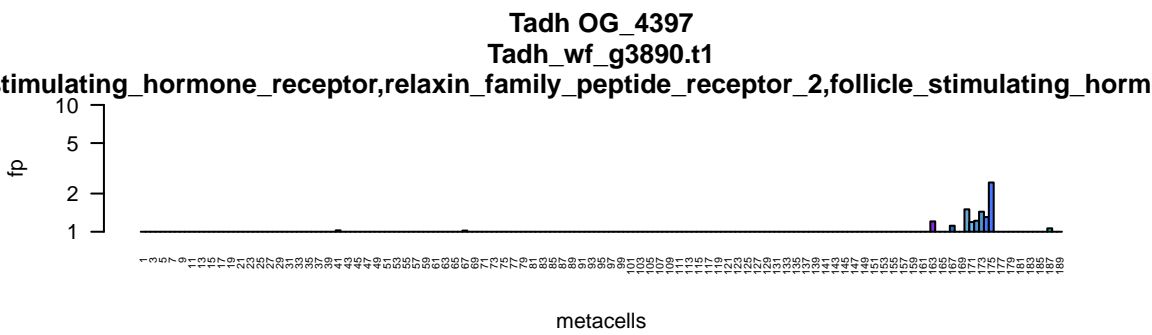
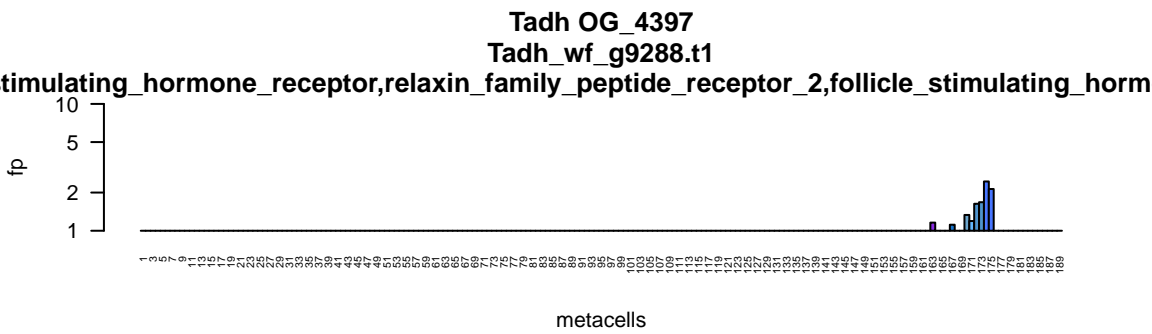


metacells

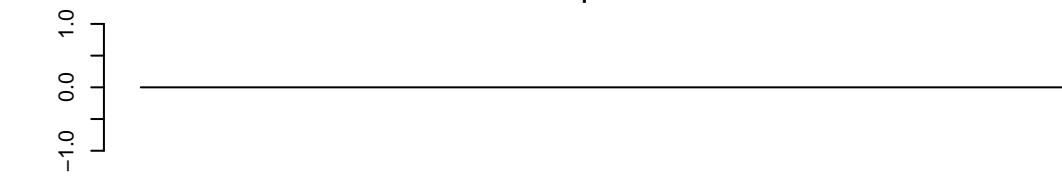




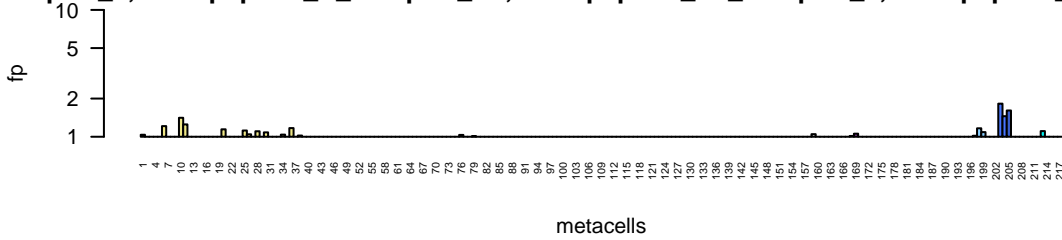




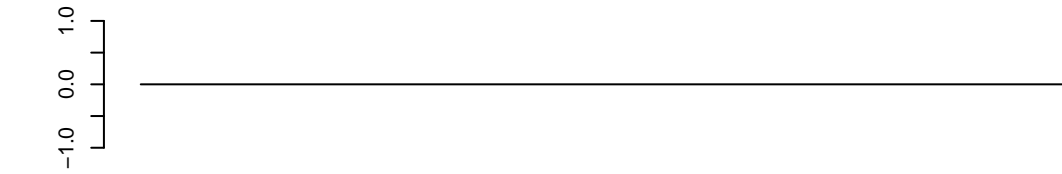
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec
Tadh | no data



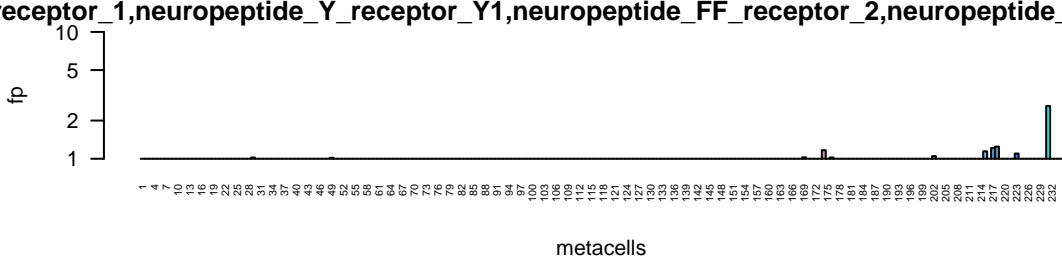
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec
TrH2 OG_4548
TrH2_TrispH2_007473-RA



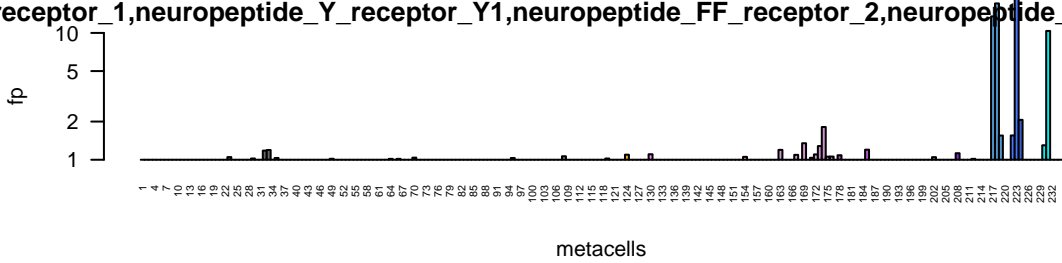
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec
Hhon | no data



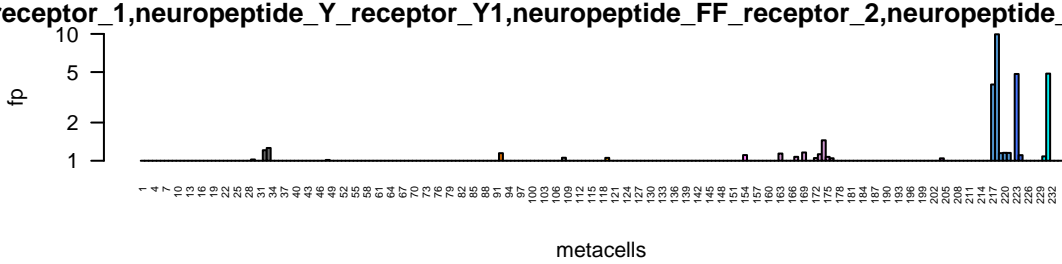
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec
HoiH23 OG_4548
HoiH23_PIH23_006484-RA



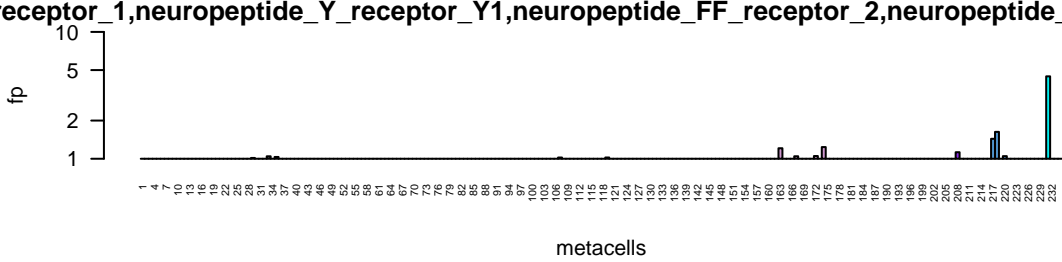
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec
HoiH23 OG_4548
HoiH23_PIH23_011478-RA

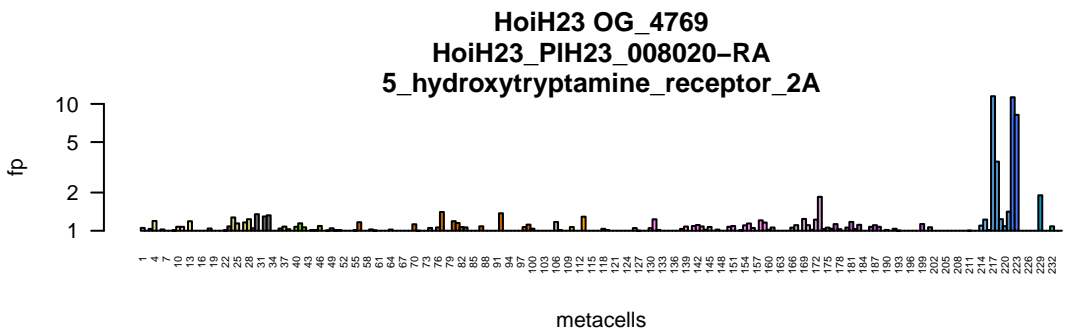
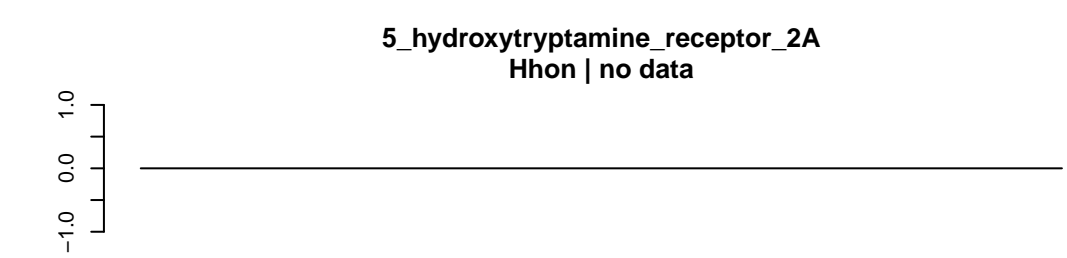
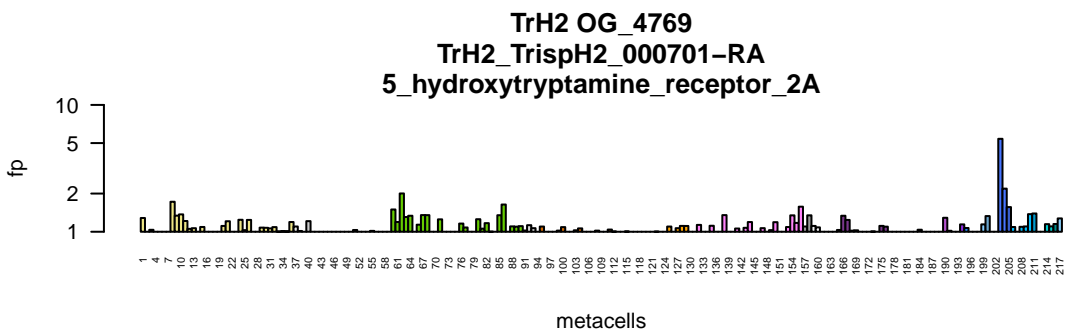
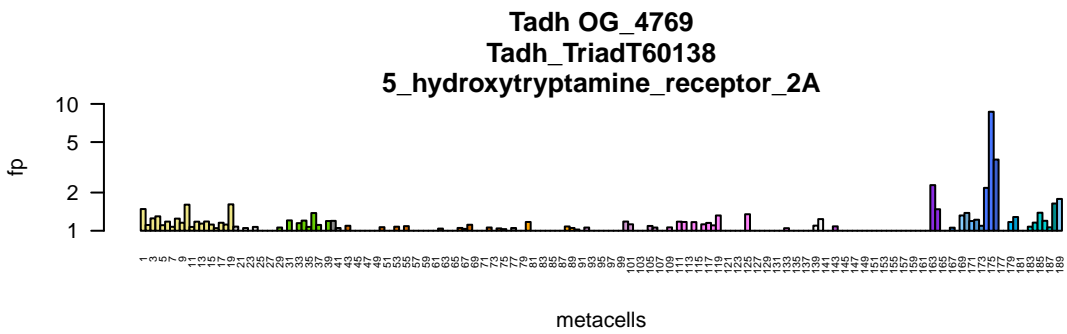


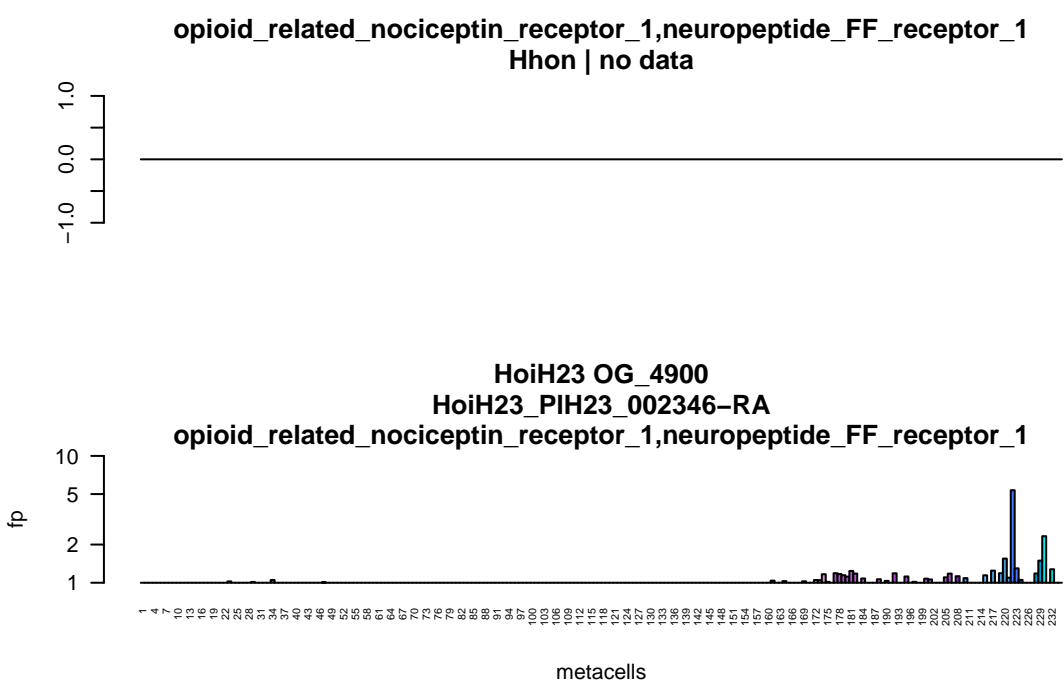
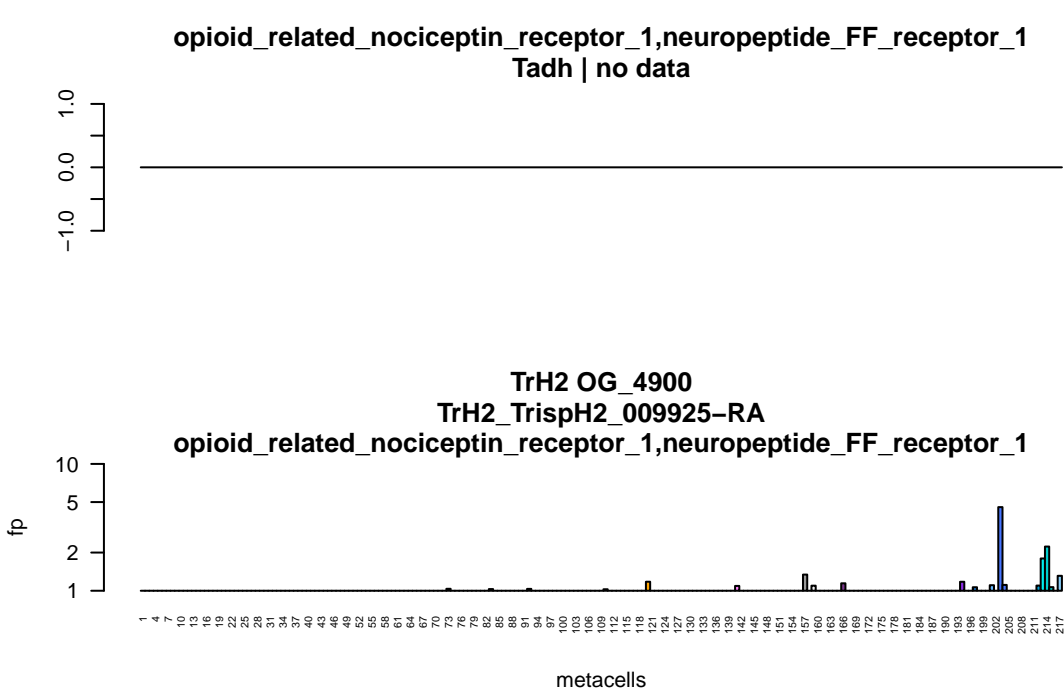
receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec
HoiH23 OG_4548
HoiH23_PIH23_011479-RA

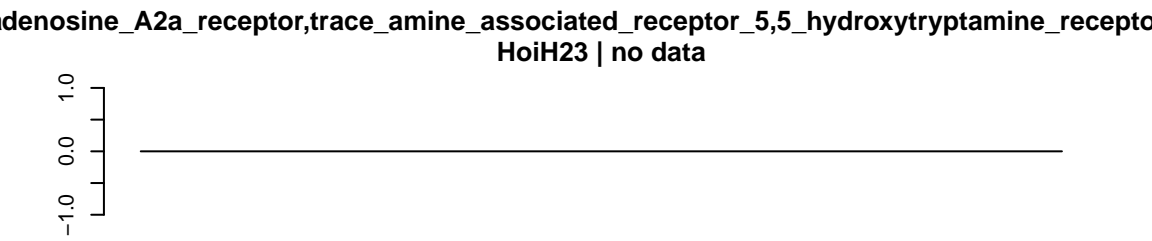
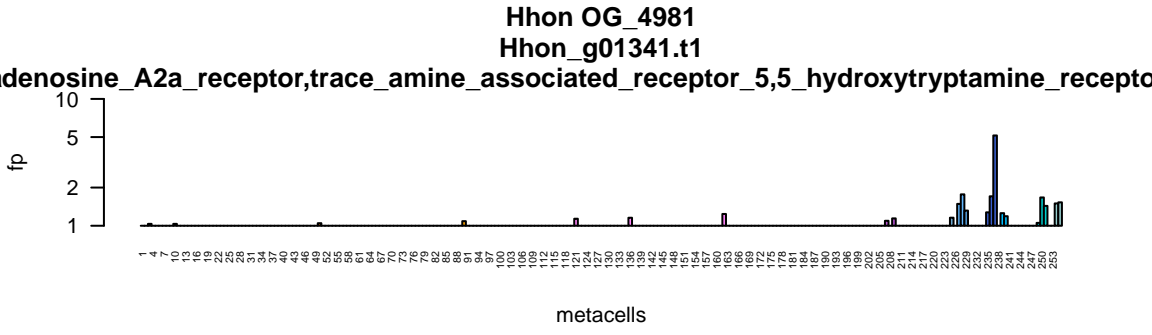
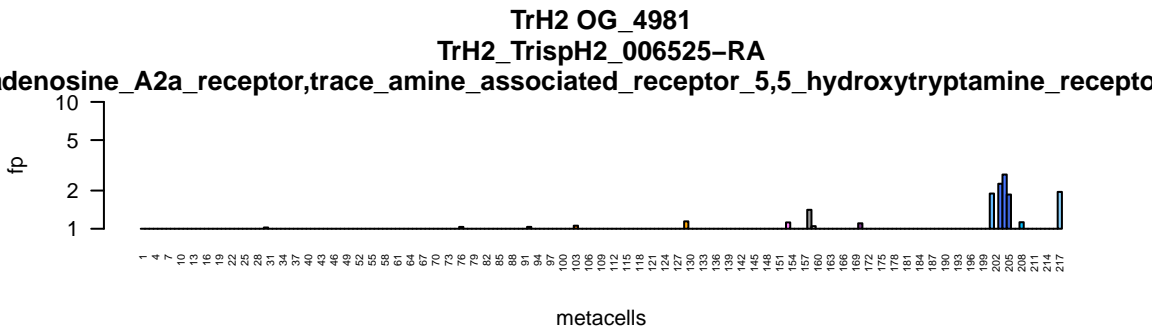
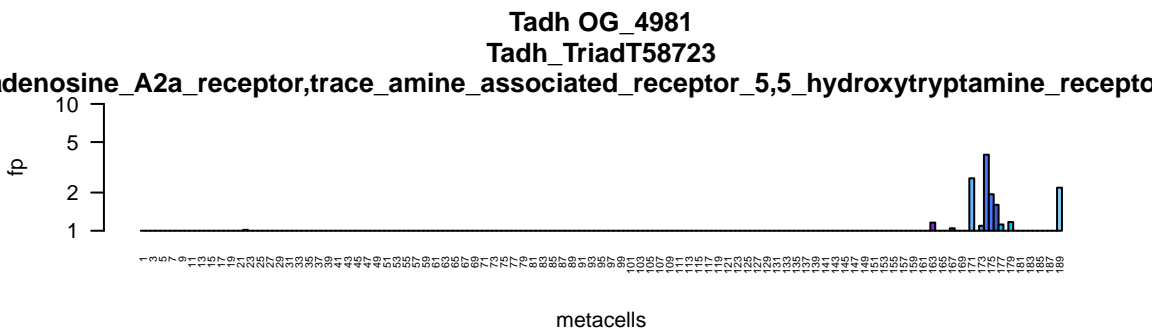


receptor_1,neuropeptide_Y_receptor_Y1,neuropeptide_FF_receptor_2,neuropeptide_FF_rec
HoiH23 OG_4548
HoiH23_PIH23_011481-RA

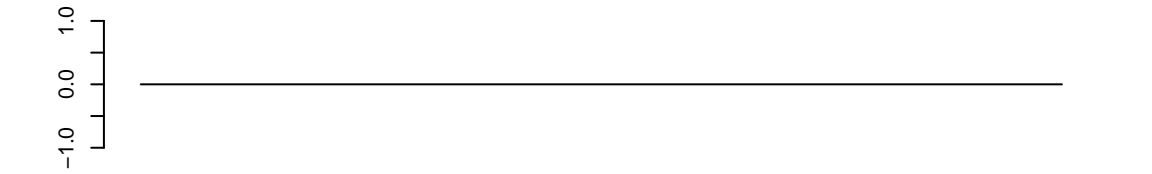




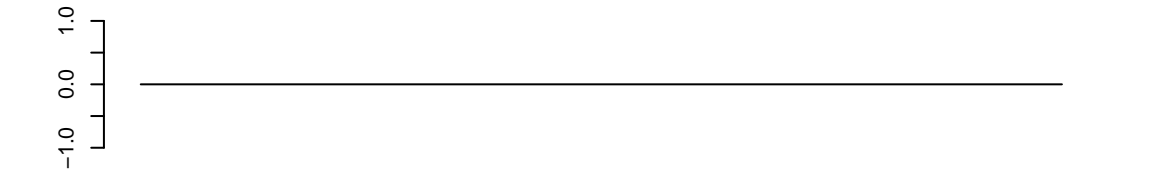




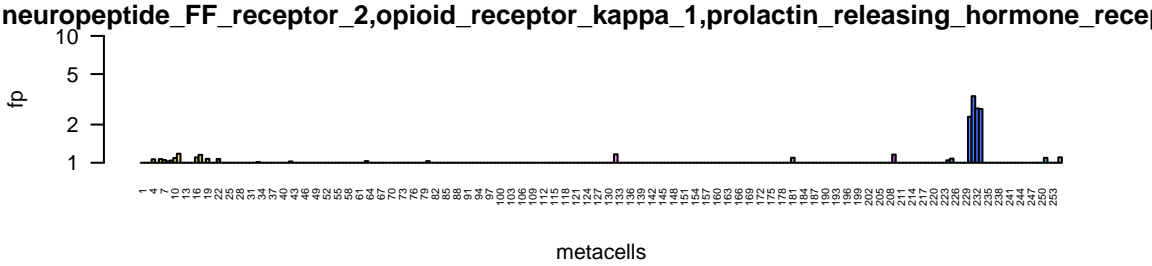
neuropeptide_FF_receptor_2,opioid_receptor_kappa_1,prolactin_releasing_hormone_rece
Tadh | no data



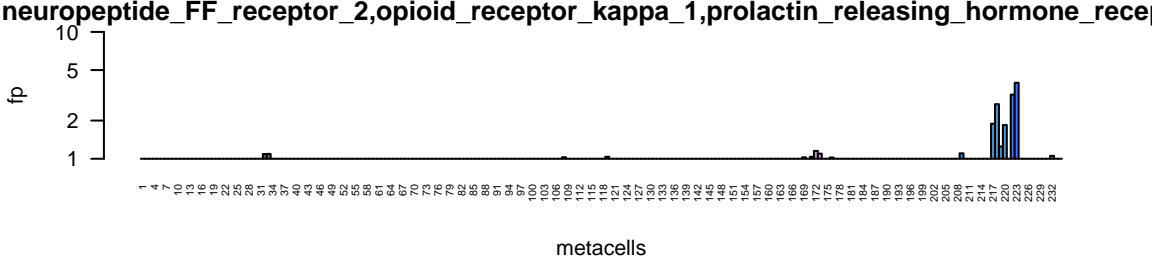
neuropeptide_FF_receptor_2,opioid_receptor_kappa_1,prolactin_releasing_hormone_rece
TrH2 | no data

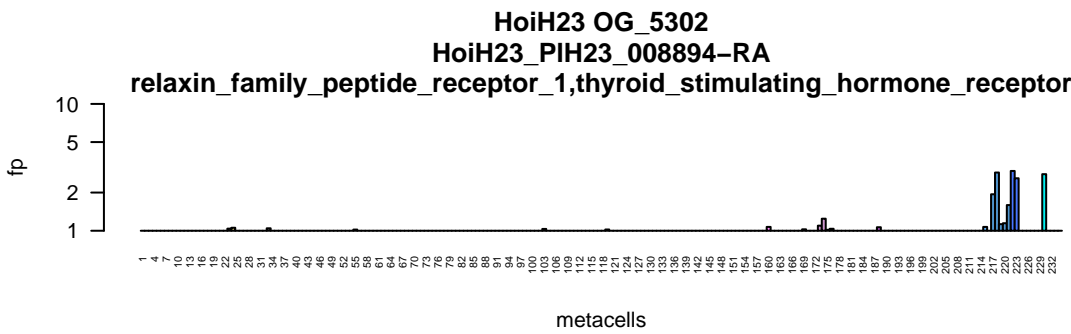
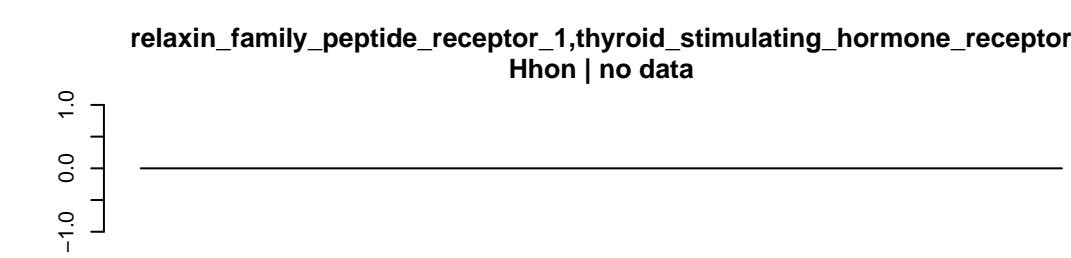
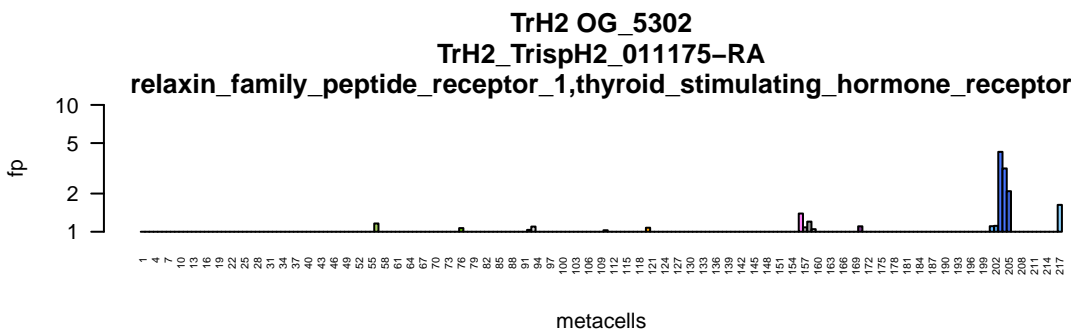
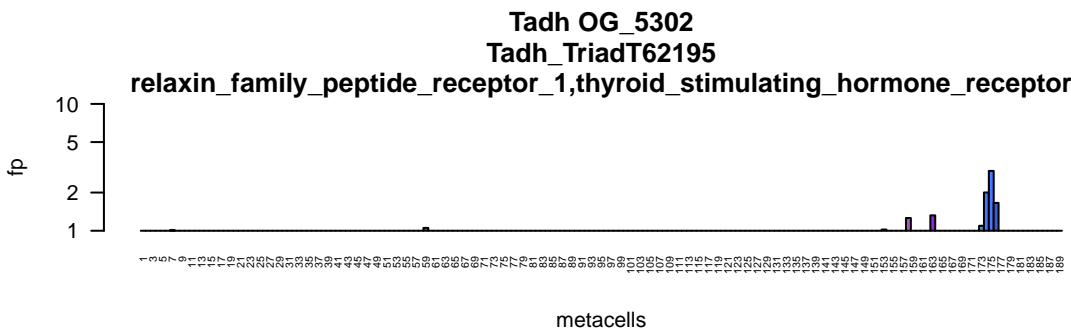


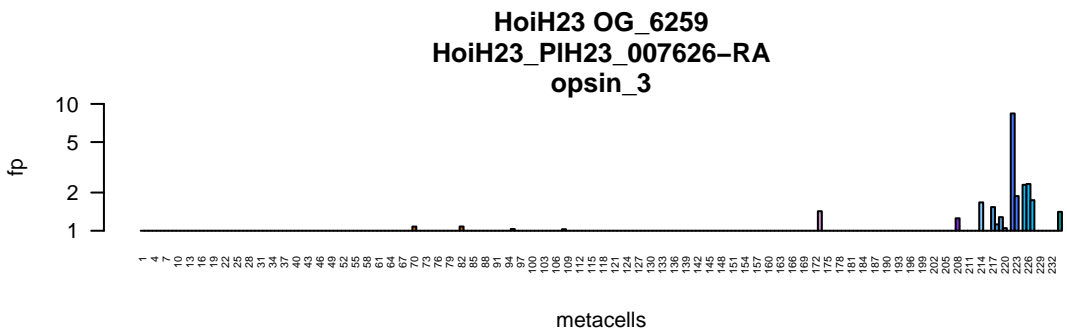
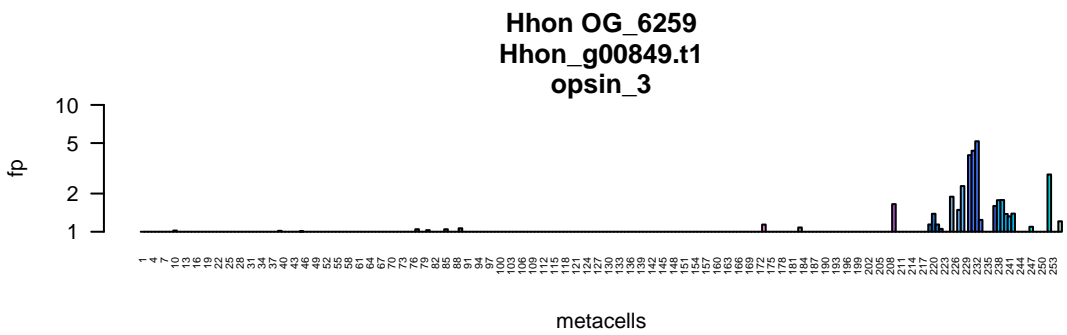
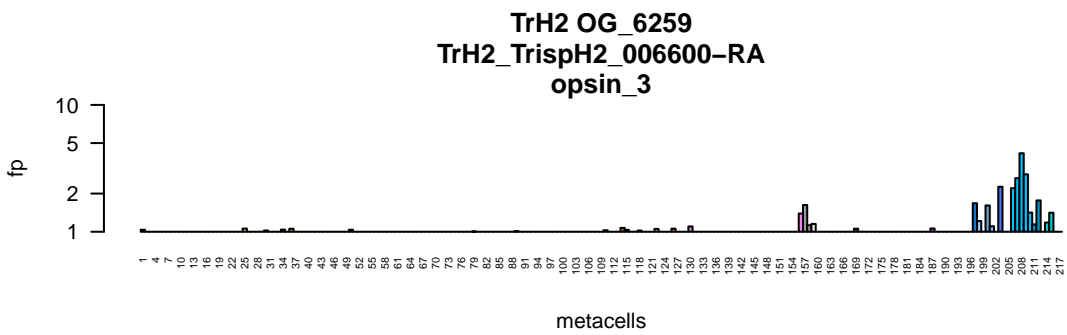
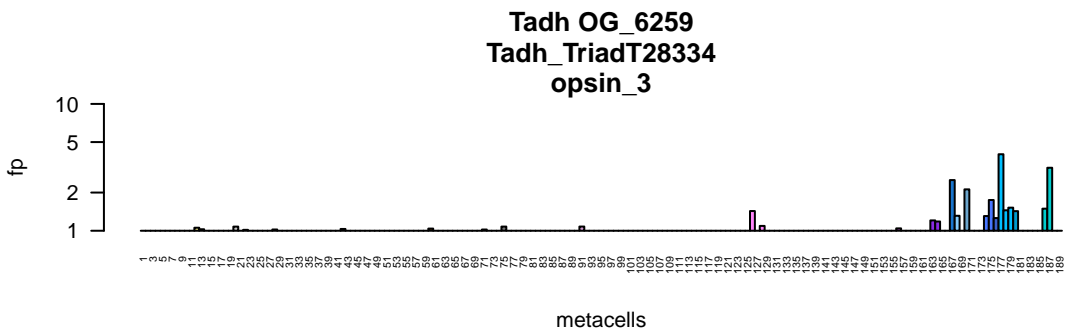
Hhon OG_5254
Hhon_g01826.t1
neuropeptide_FF_receptor_2,opioid_receptor_kappa_1,prolactin_releasing_hormone_rece



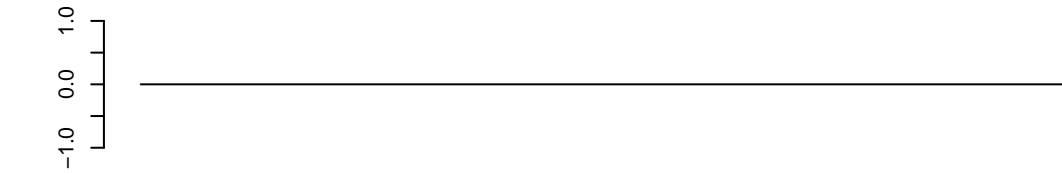
HoiH23 OG_5254
HoiH23_PIH23_011836-RA
neuropeptide_FF_receptor_2,opioid_receptor_kappa_1,prolactin_releasing_hormone_rece



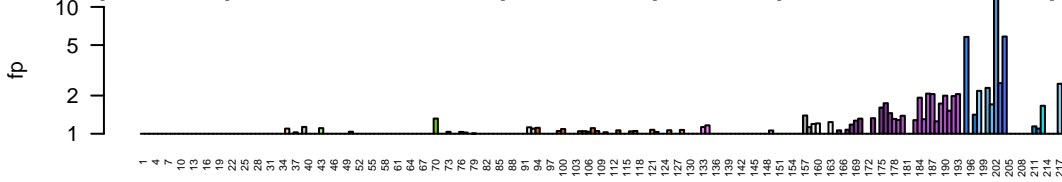




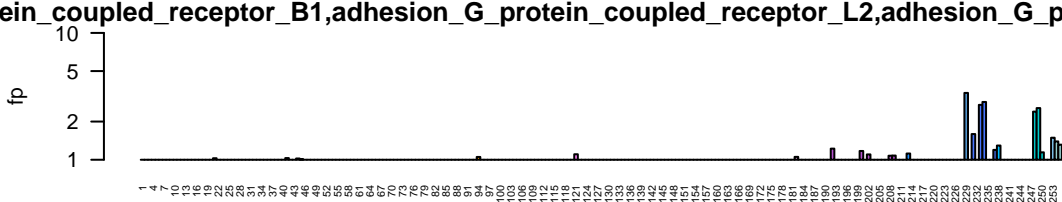
ein_coupled_receptor_B1,adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_
Tadh | no data



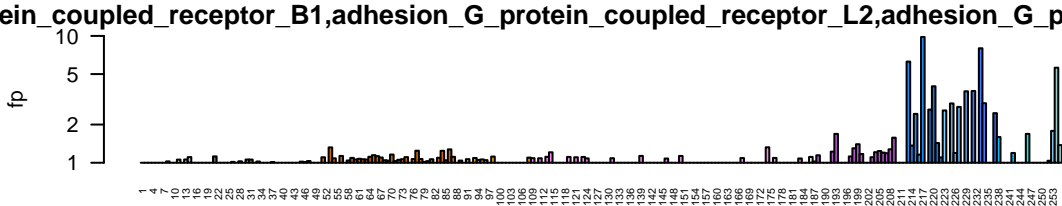
TrH2 OG_6302
TrH2_TrispH2_005655-RA
ein_coupled_receptor_B1,adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_



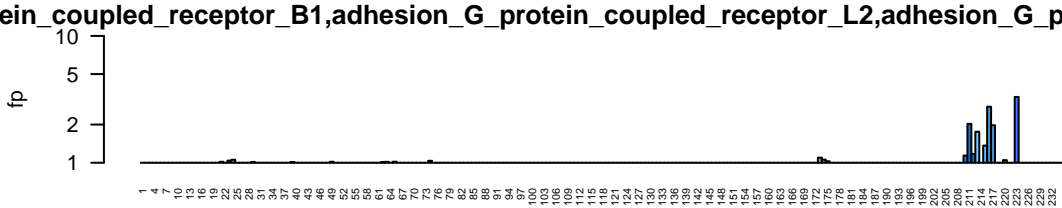
metacells
Hhon OG_6302
Hhon_g06052.t1
ein_coupled_receptor_B1,adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_



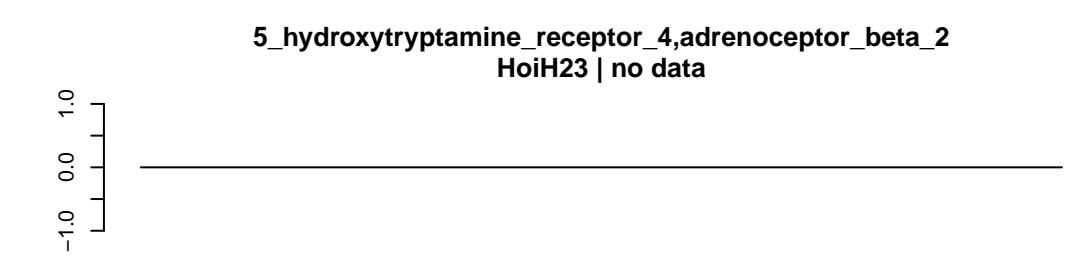
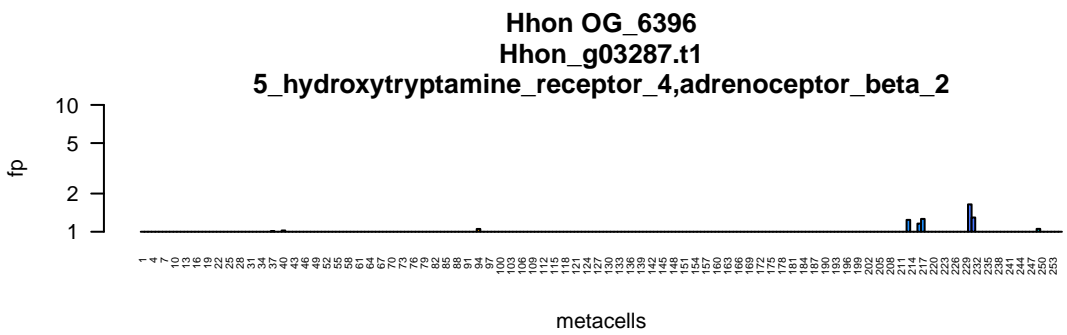
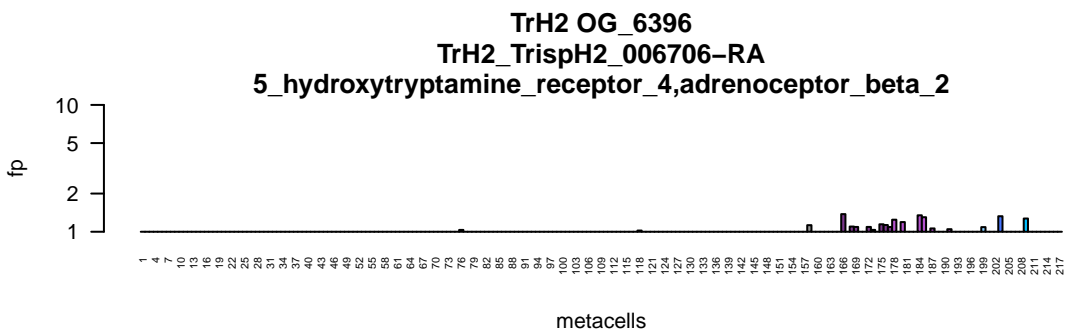
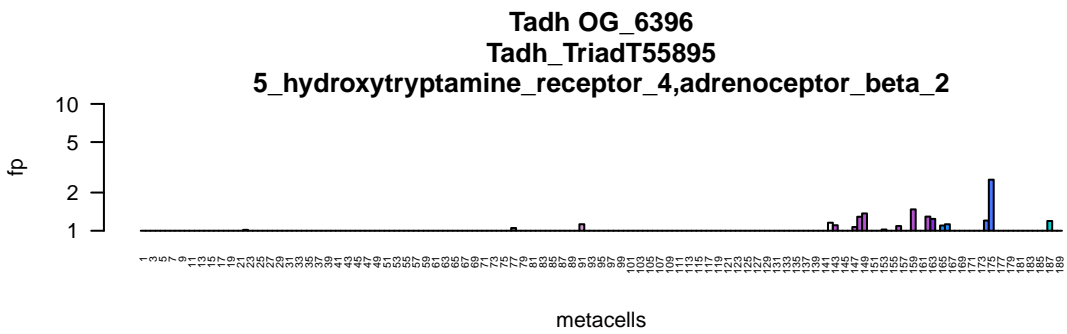
metacells
Hhon OG_6302
Hhon_g06053.t1
ein_coupled_receptor_B1,adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_



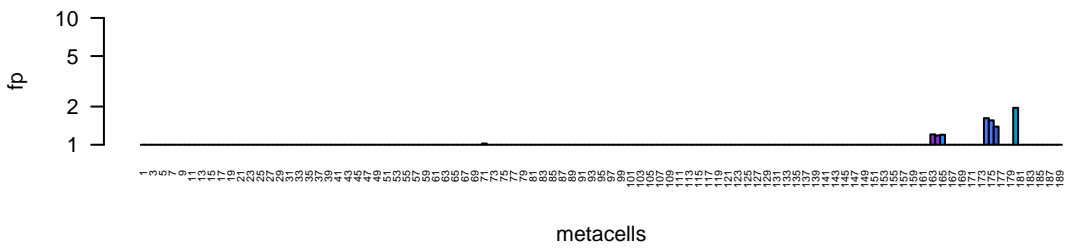
metacells
HoiH23 OG_6302
HoiH23_PIH23_004487-RA
ein_coupled_receptor_B1,adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_



metacells



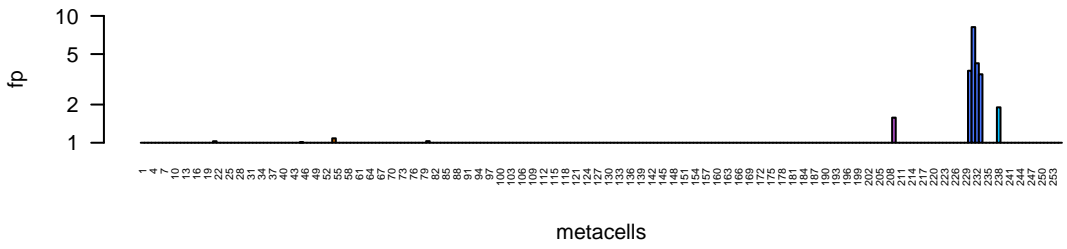
Tadh OG_6447
Tadh_TriadT60874



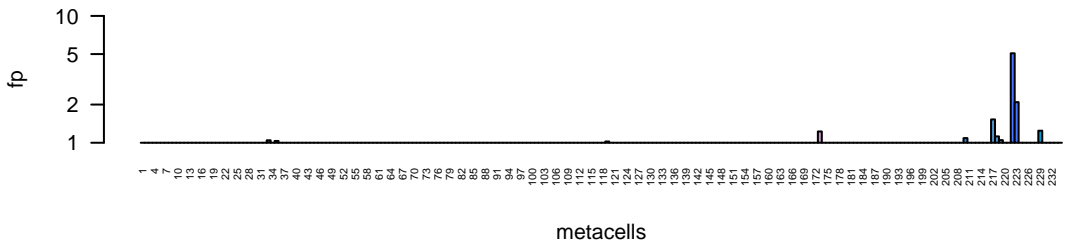
TrH2 | no data

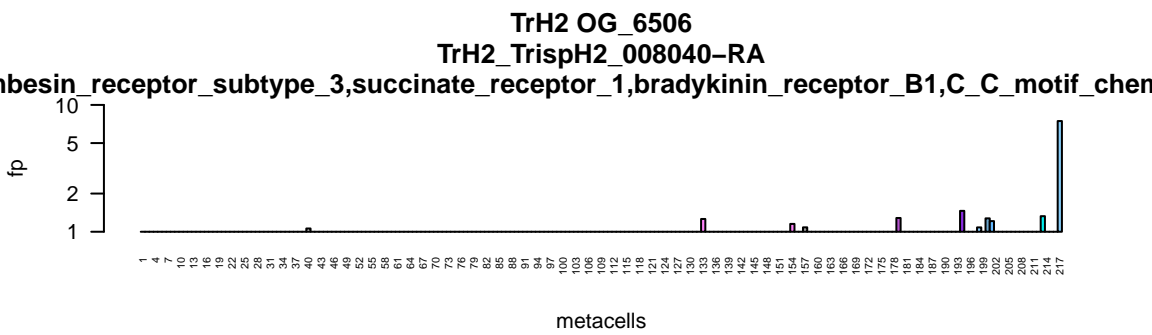
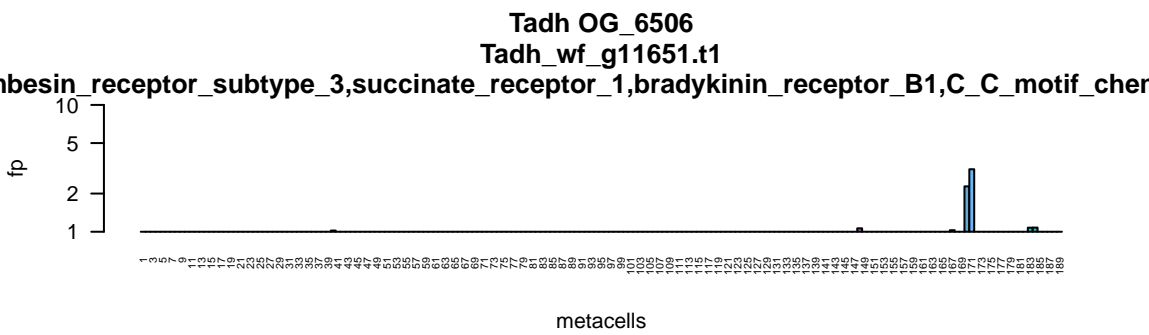
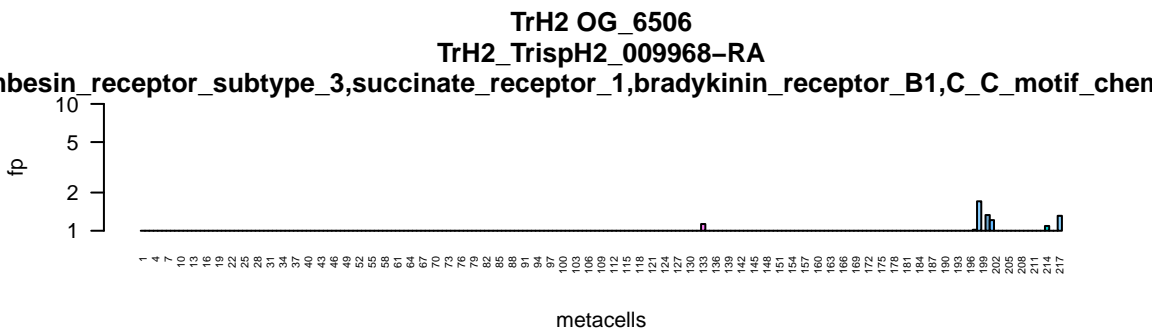
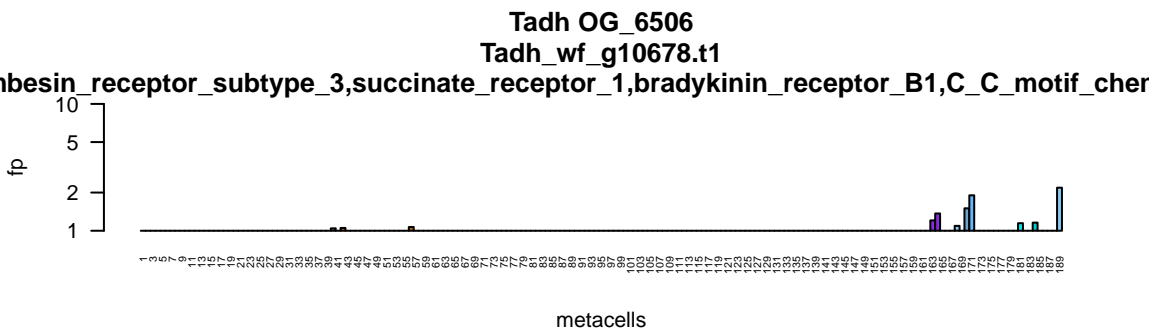
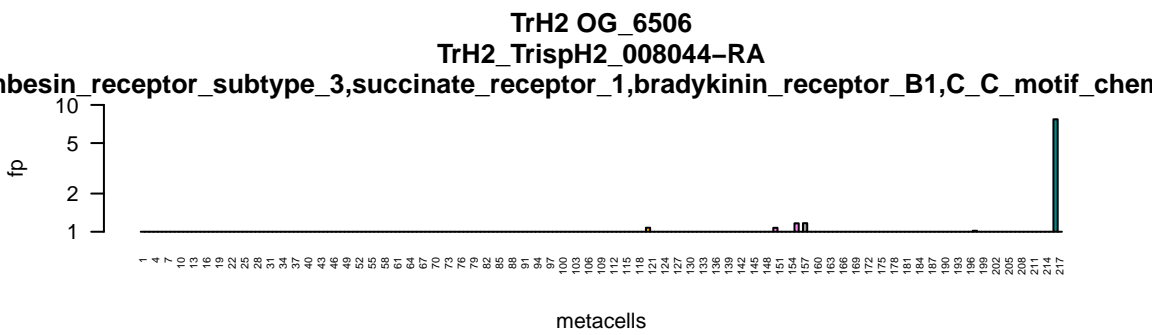
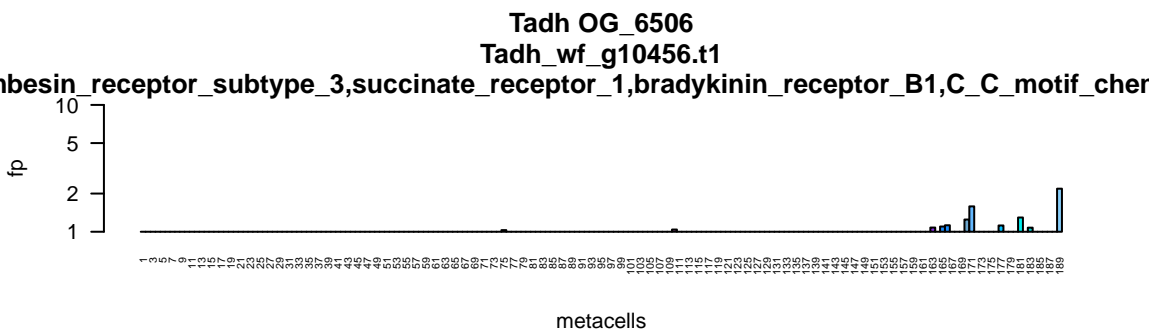
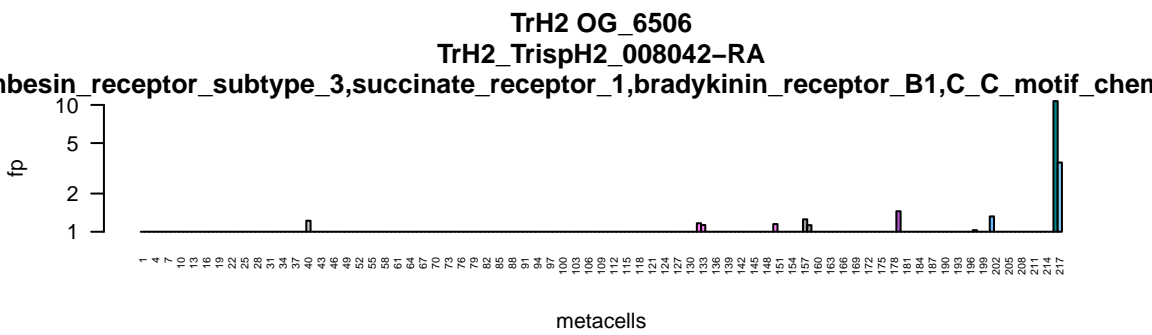
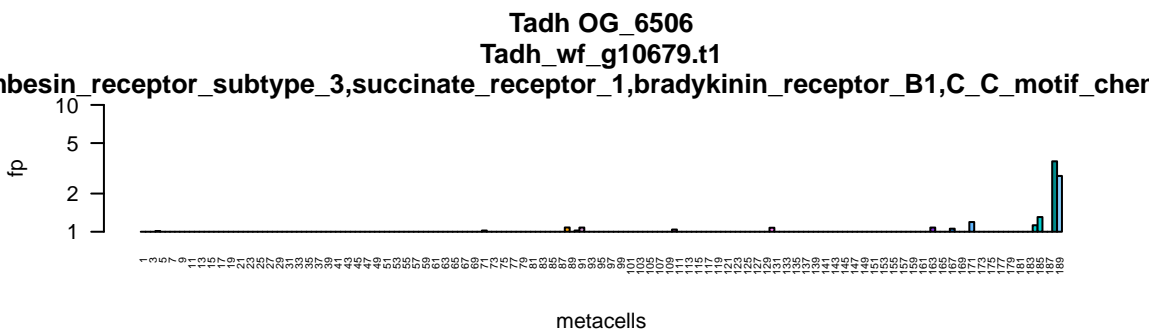
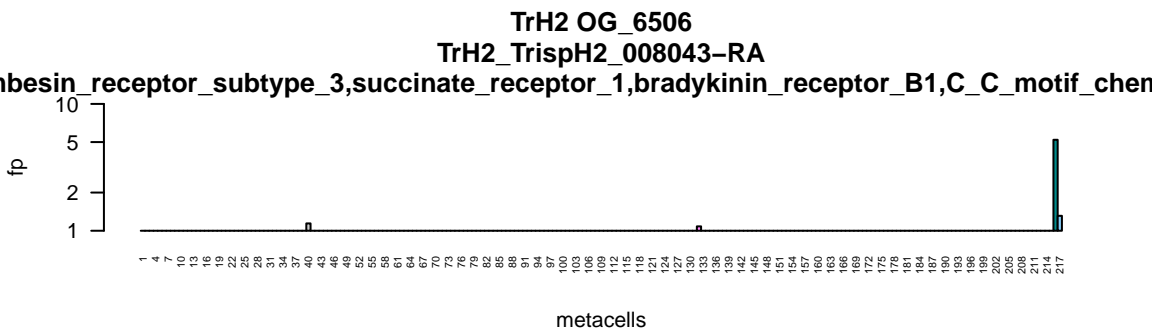
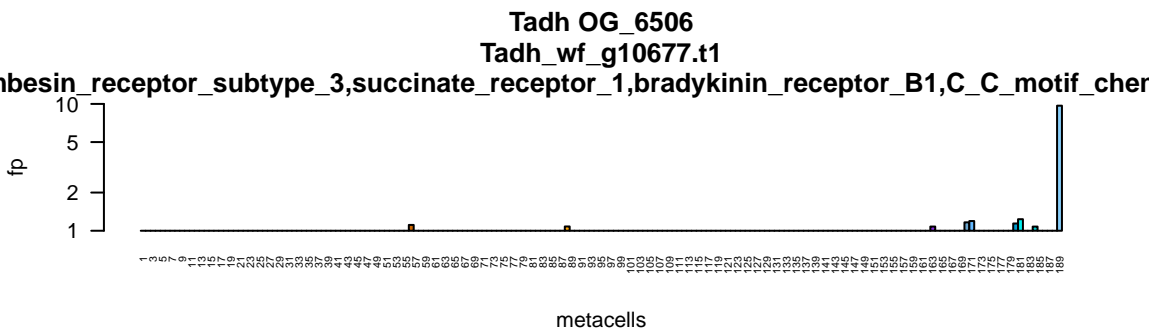
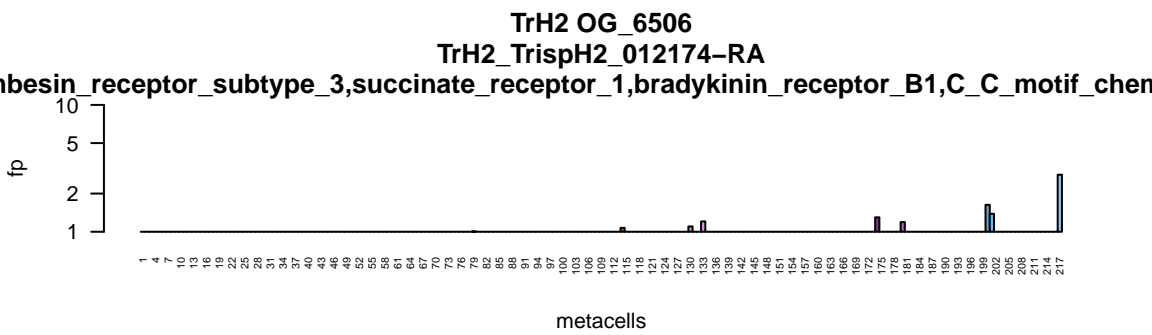
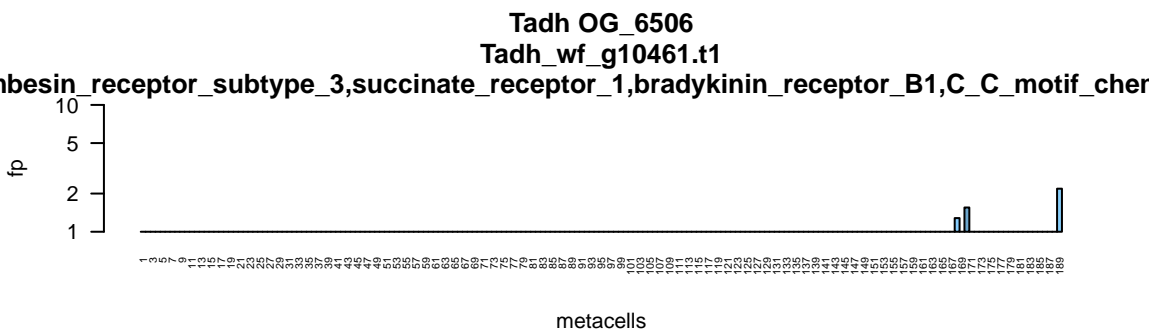
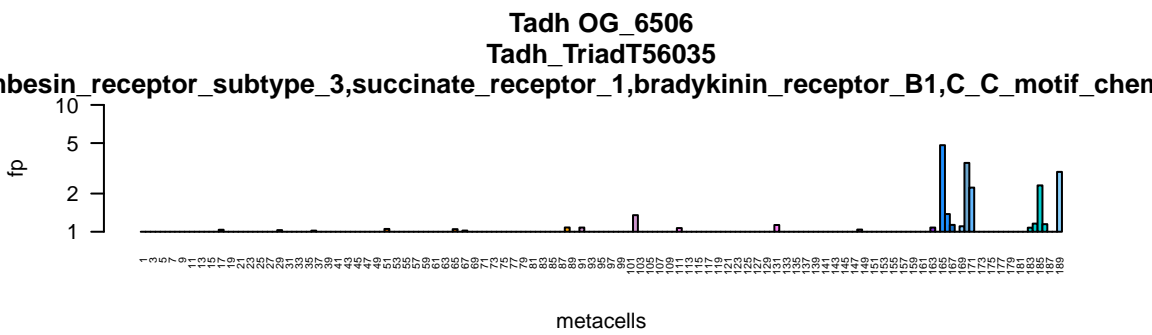
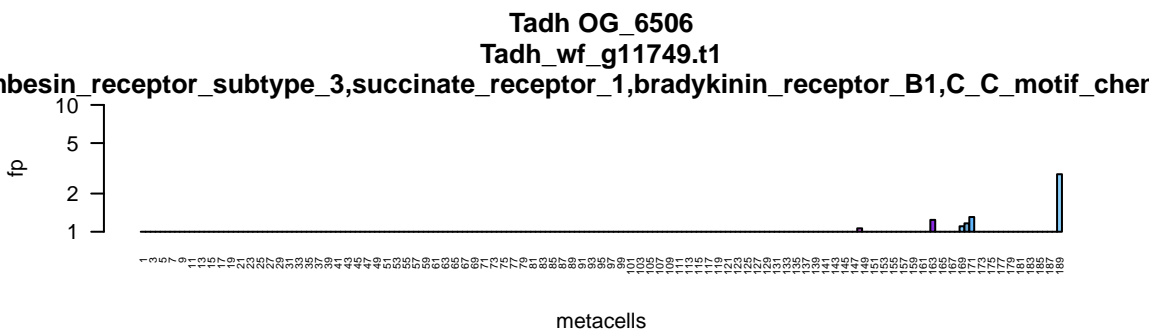
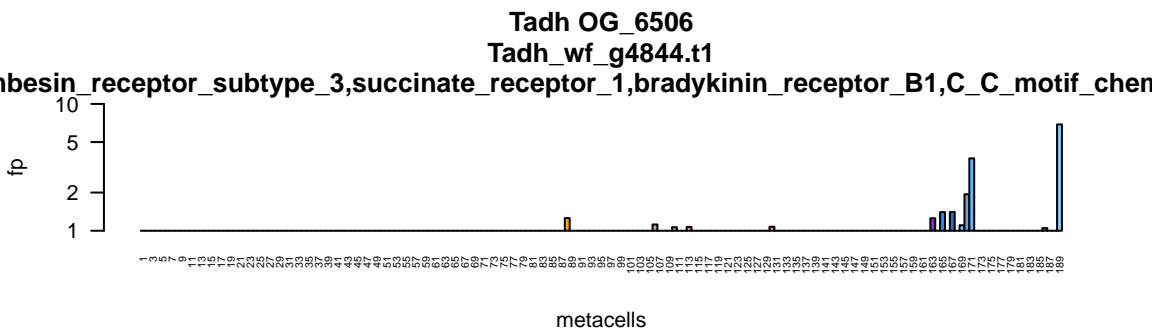
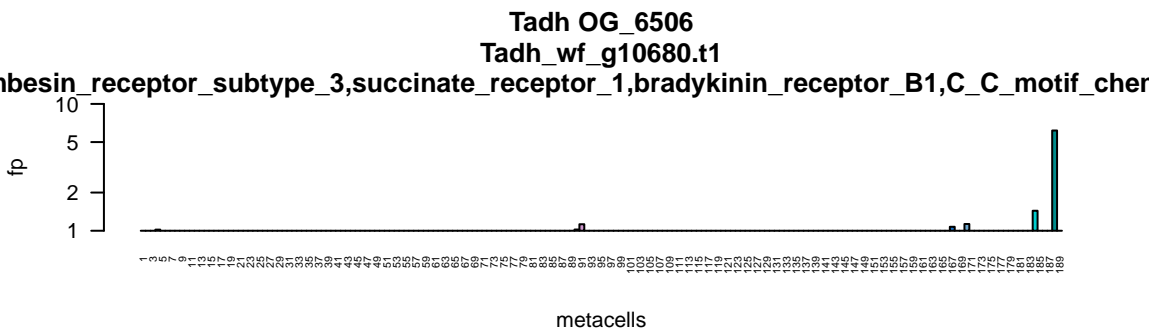


Hhon OG_6447
Hhon_g10773.t1

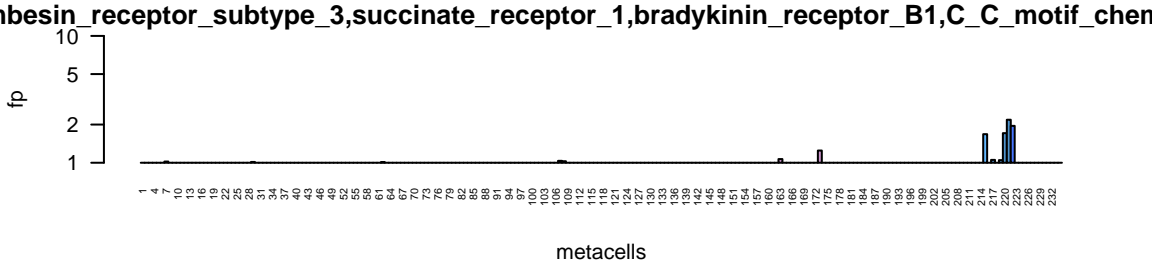


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HoiH23_PIH23_005834-RA

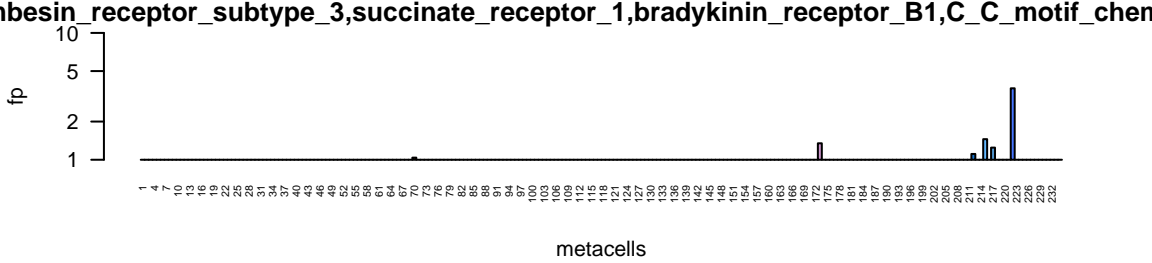


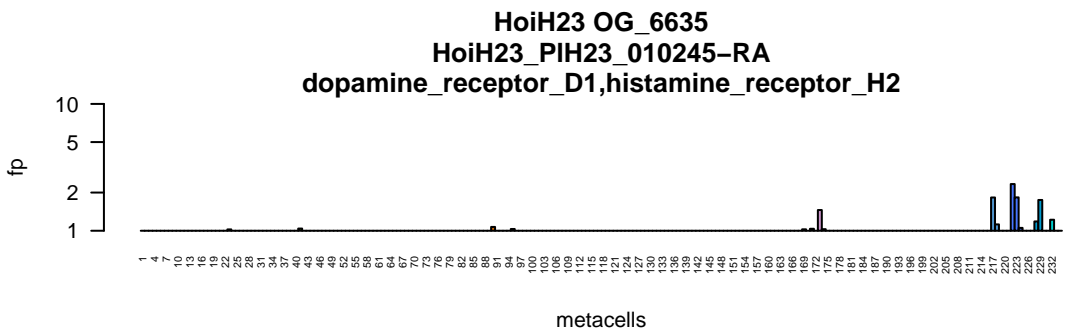
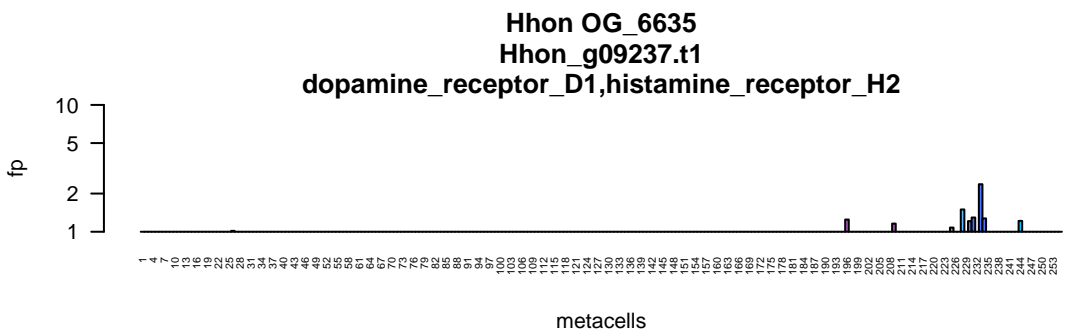
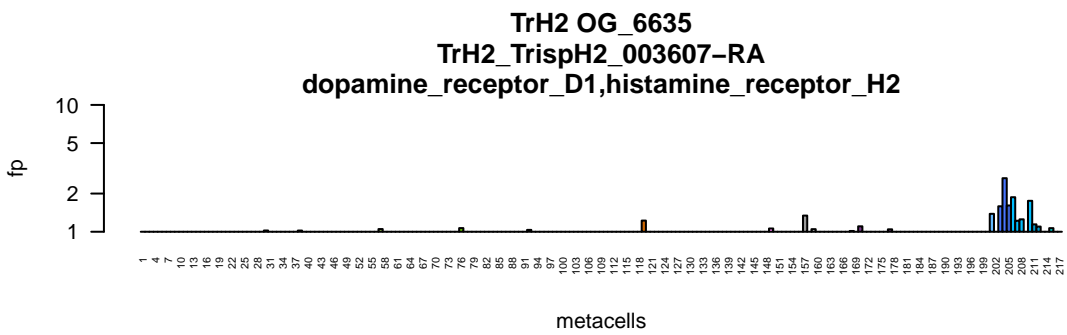
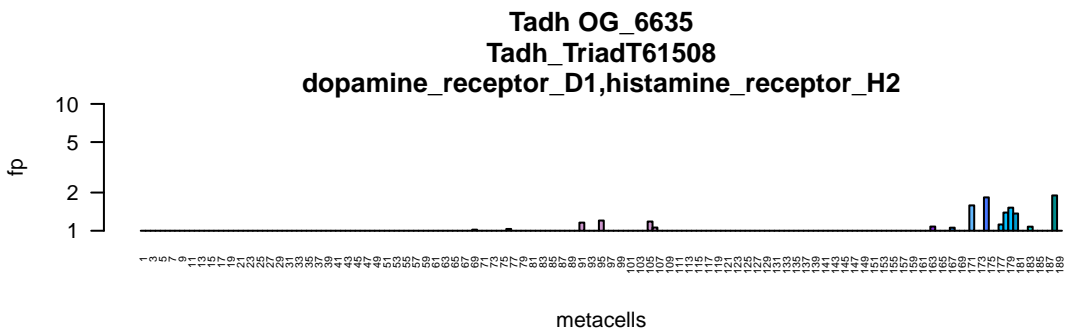


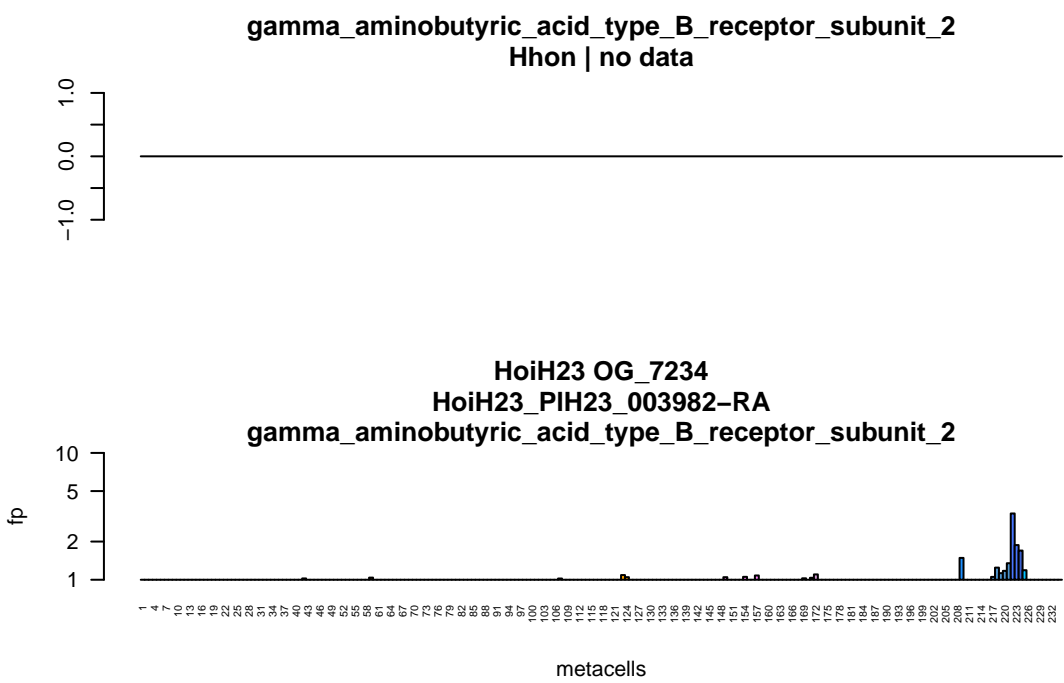
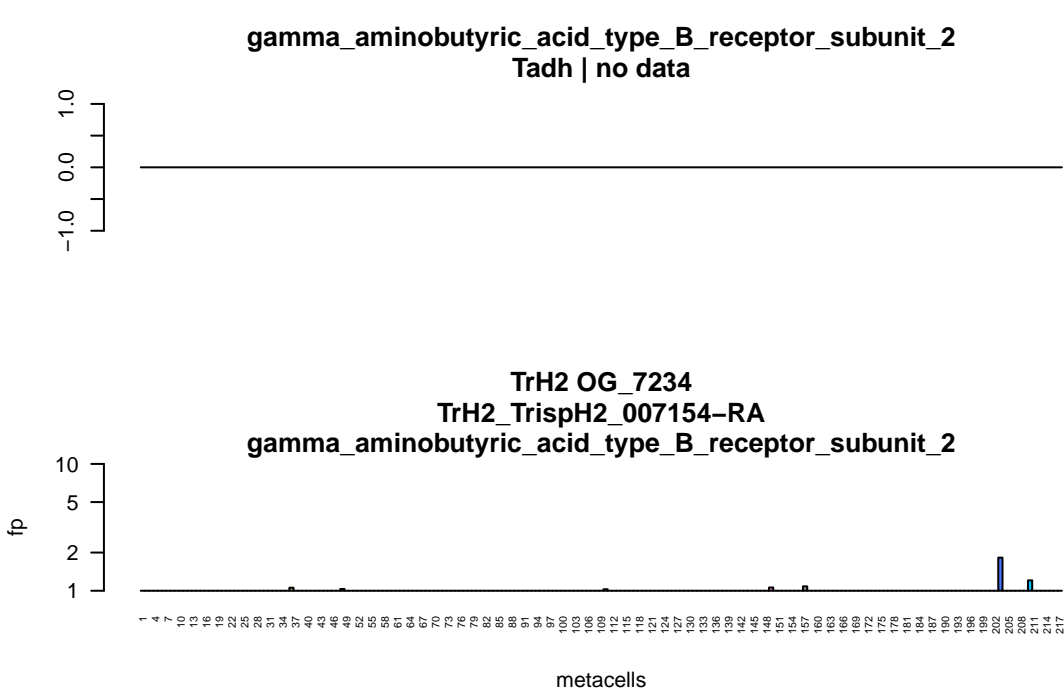
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HoiH23_PIH23_008211-RA

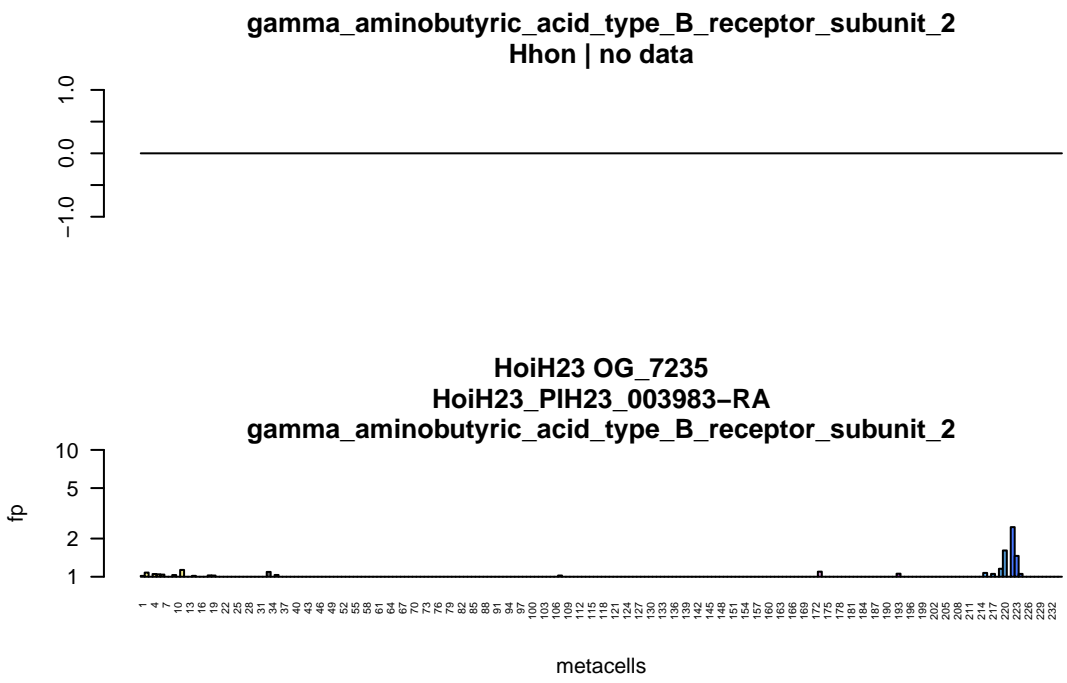
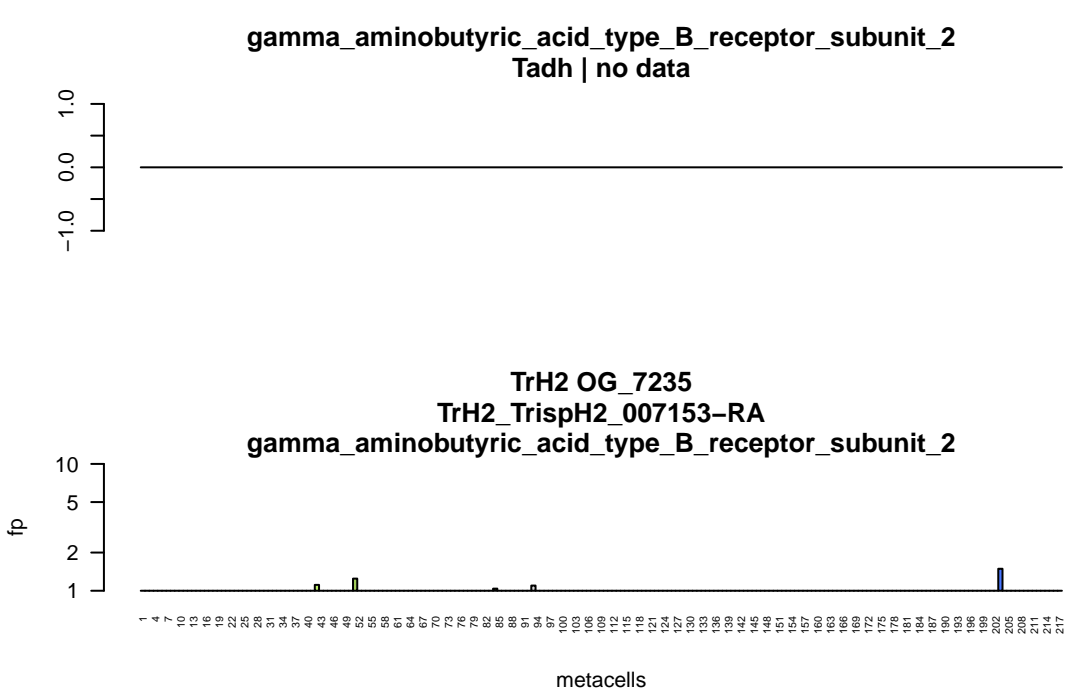


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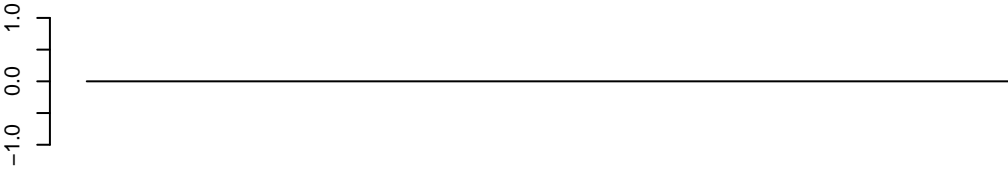








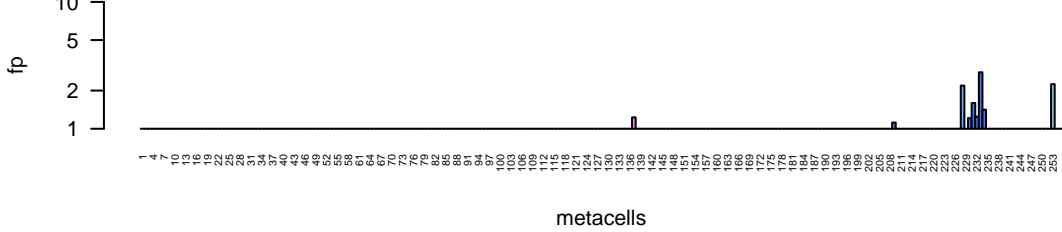
gamma_aminobutyric_acid_type_B_receptor_subunit_2
Tadh | no data



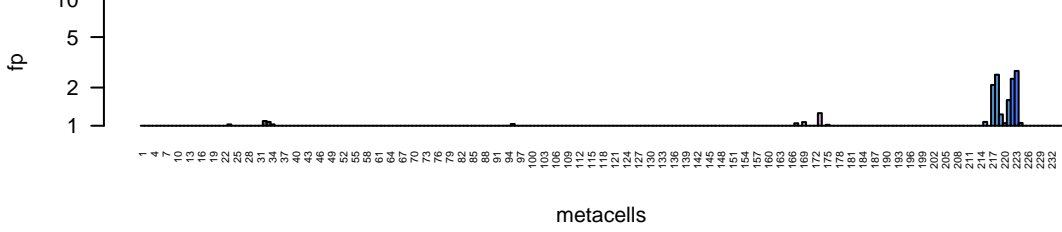
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TrH2 | no data

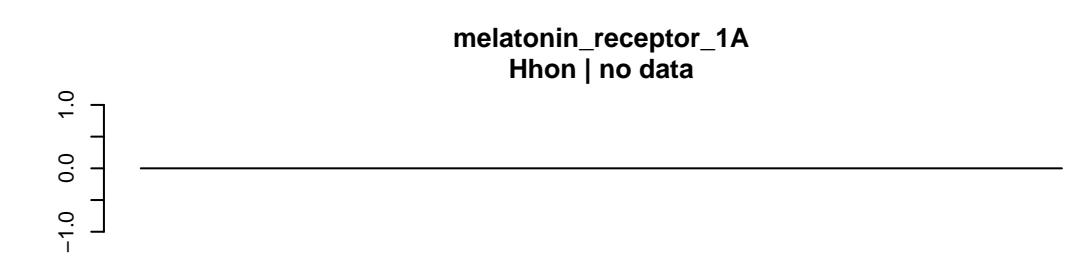
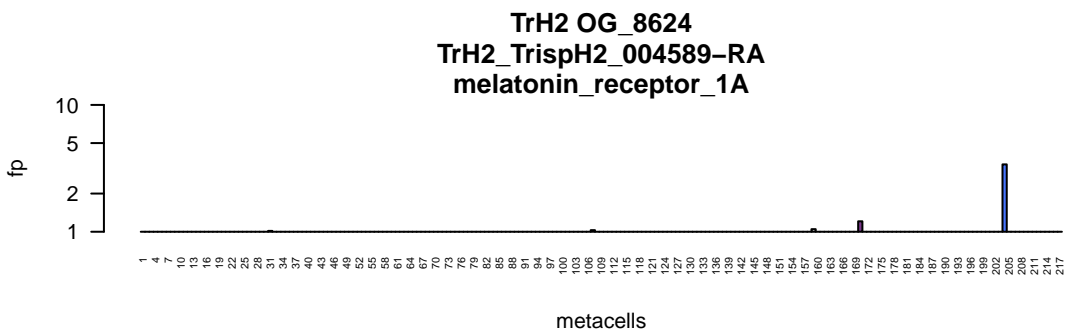
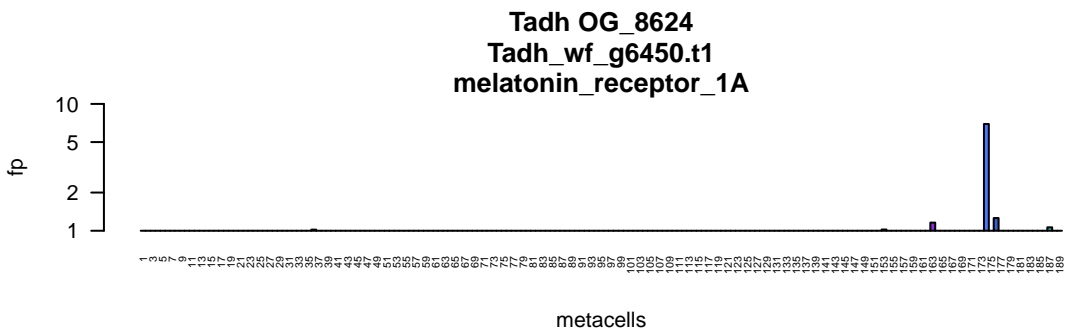


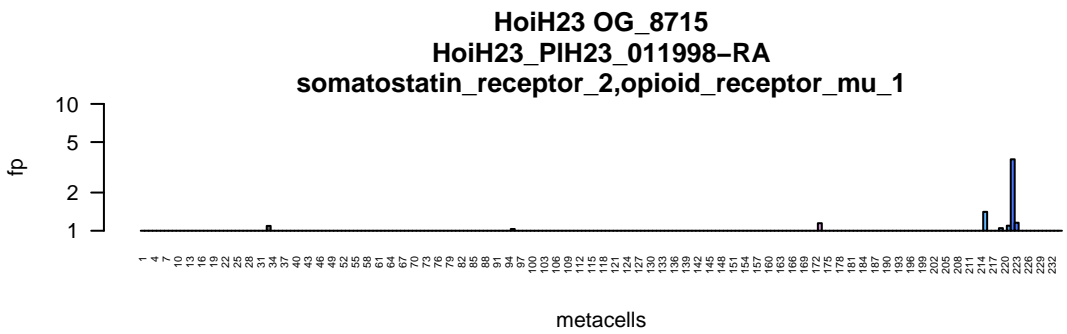
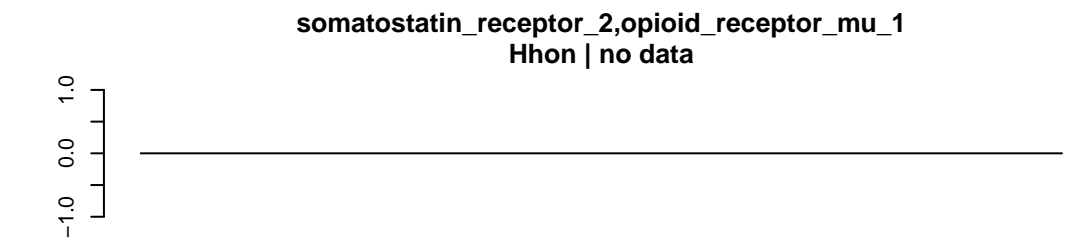
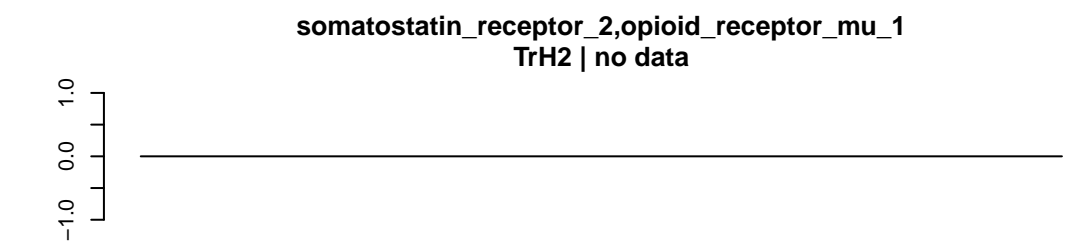
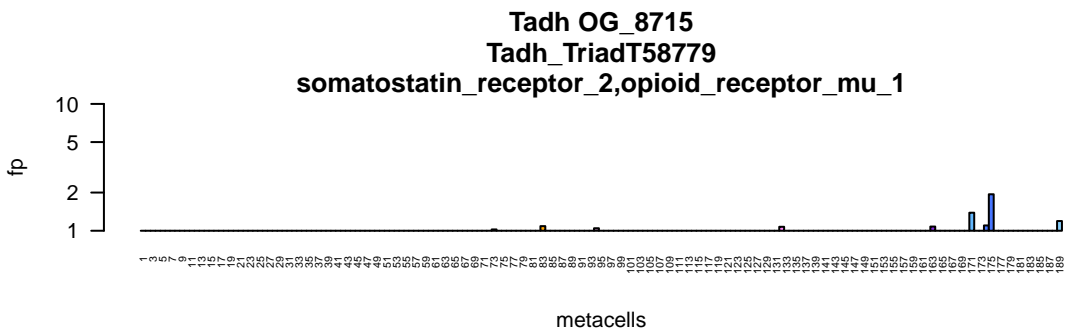
Hhon OG_7236
Hhon_g07333.t1
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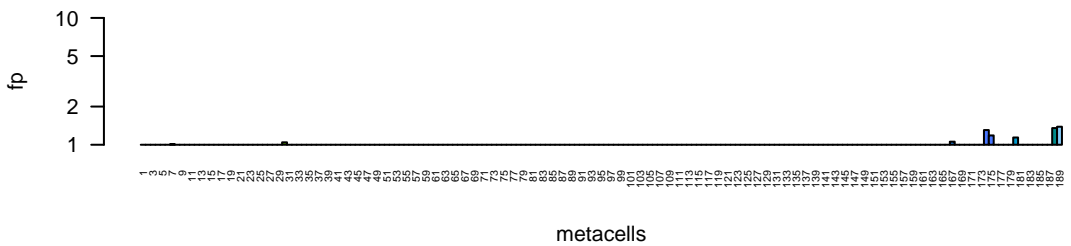
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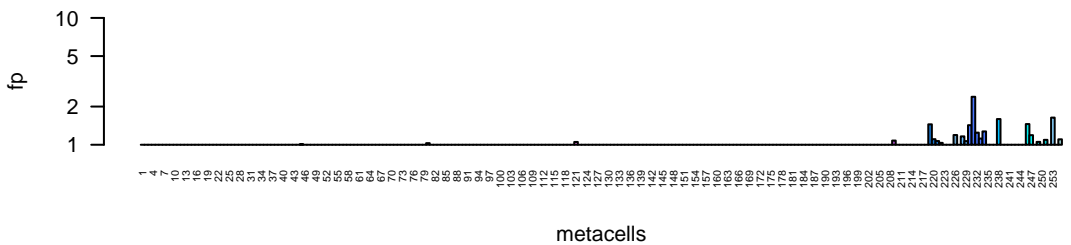
Tadh OG_9610
Tadh_TriadT60281



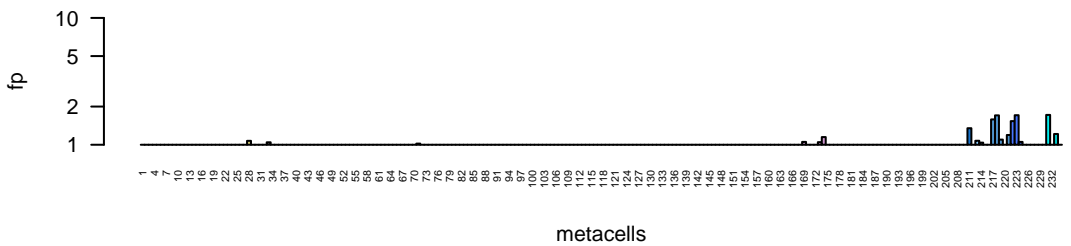
TrH2 | no data

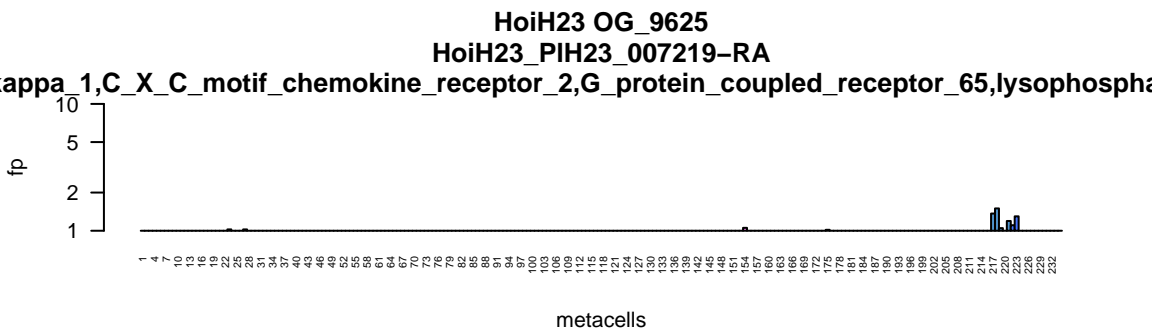
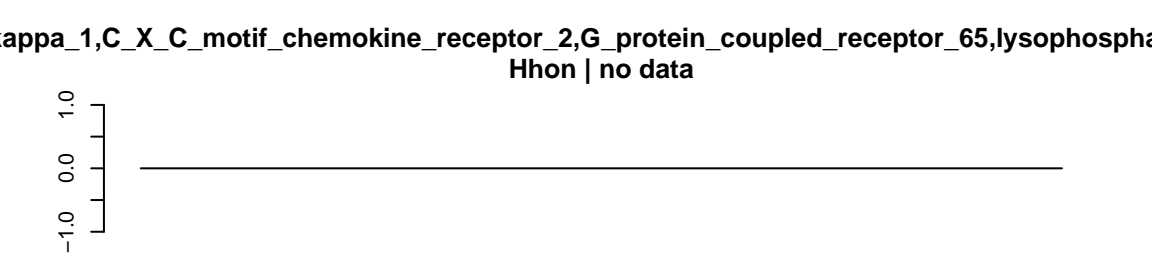
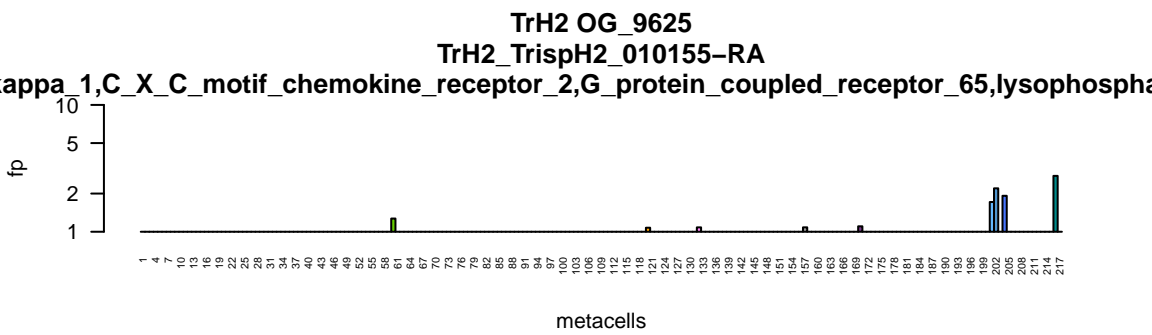
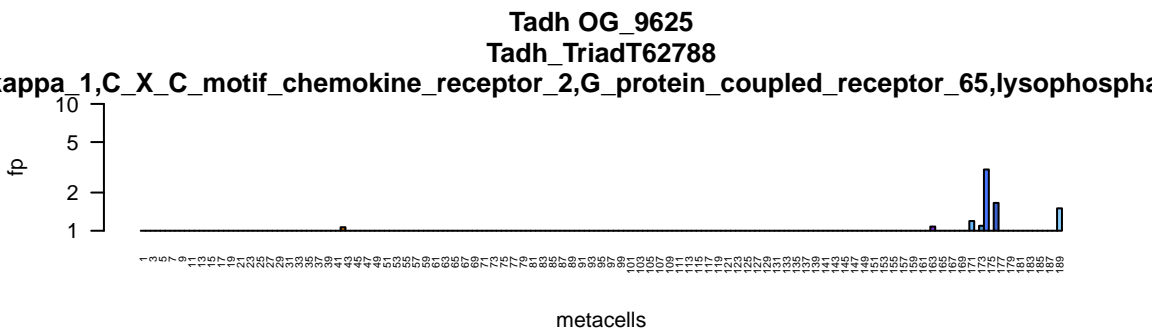


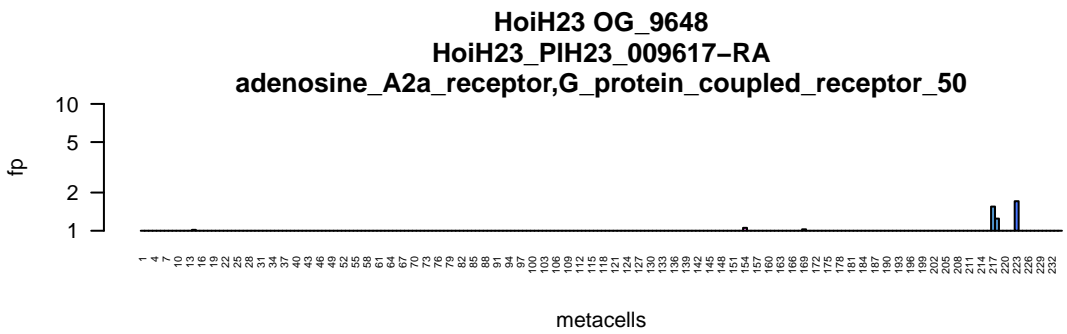
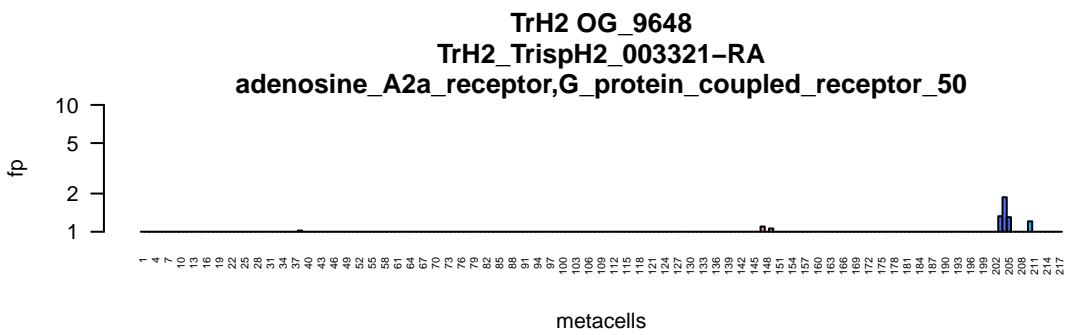
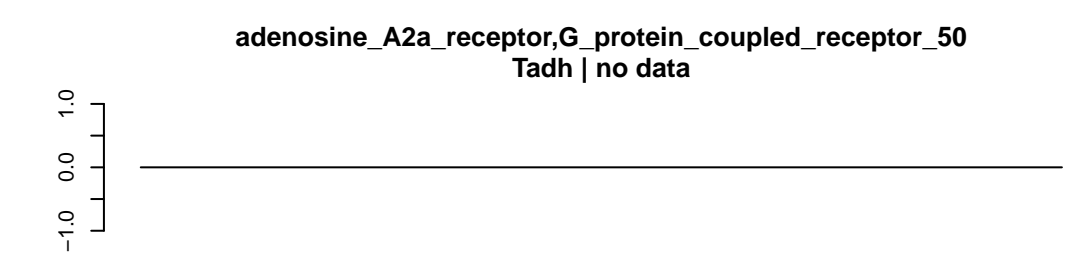
Hhon OG_9610
Hhon_g08104.t1

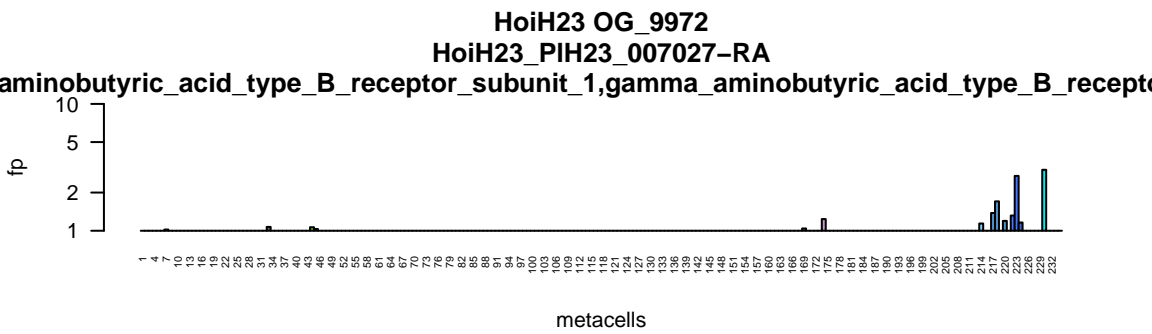
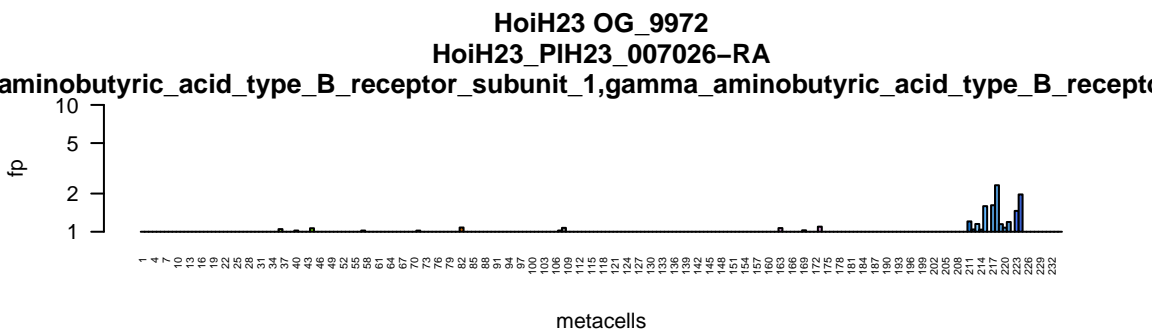
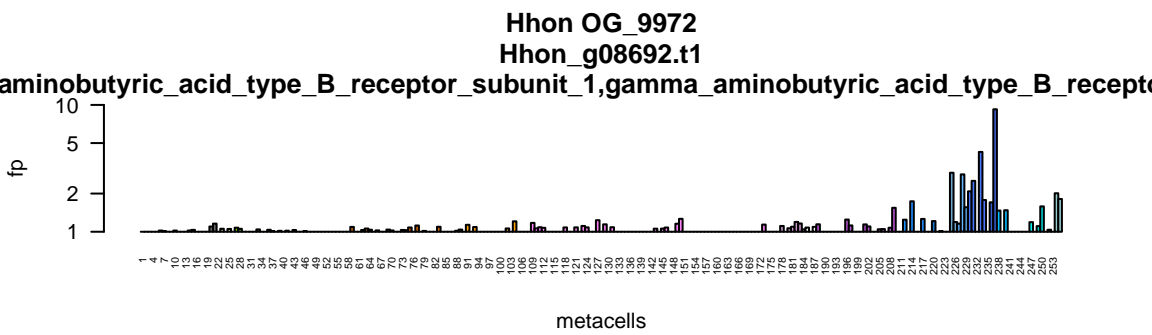
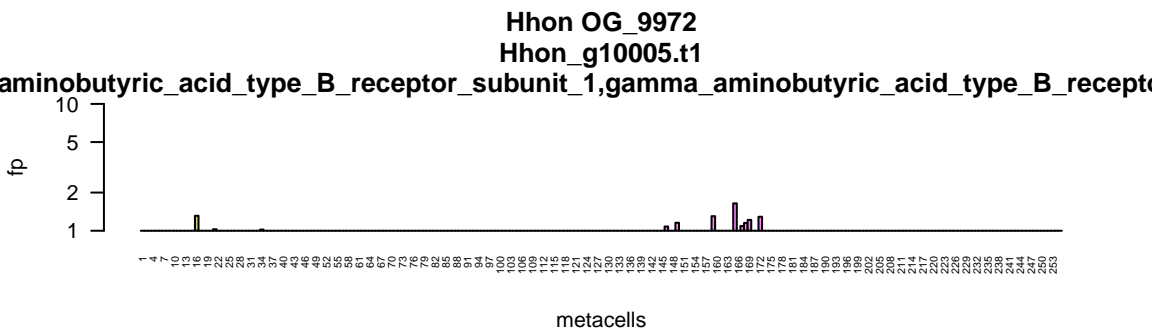
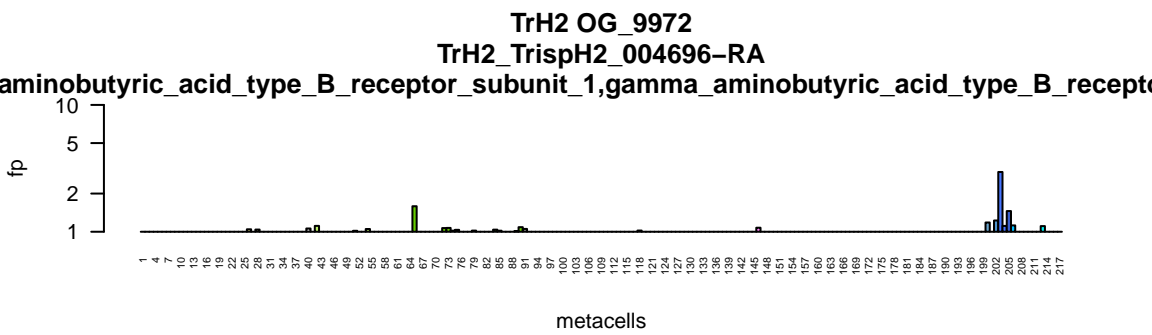
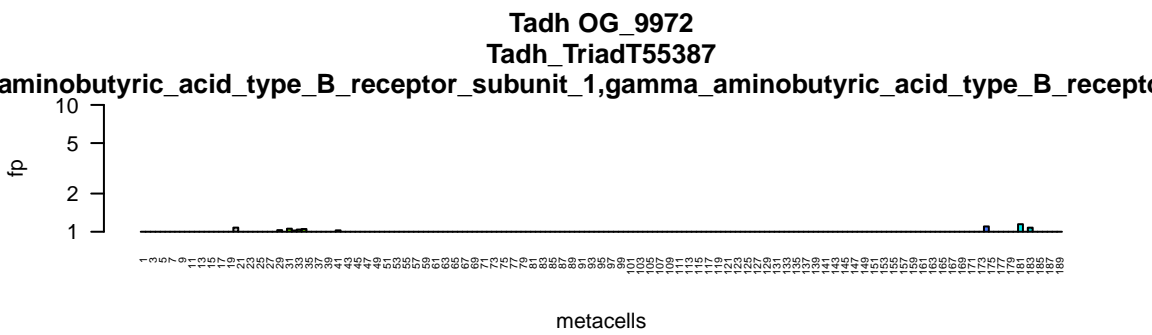
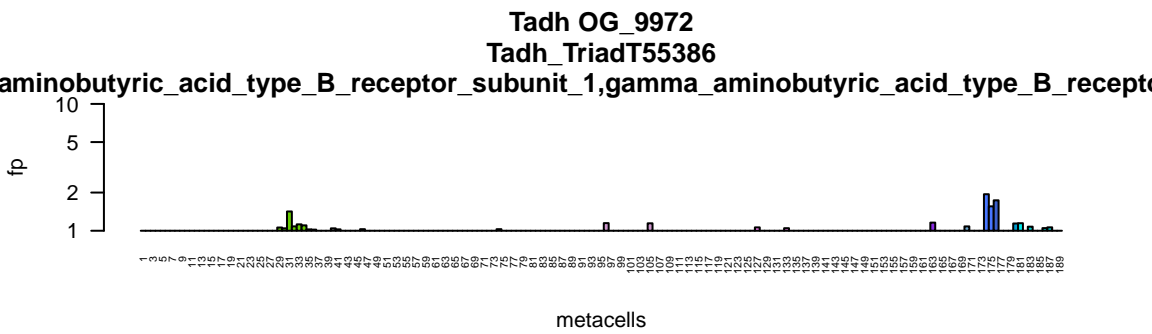


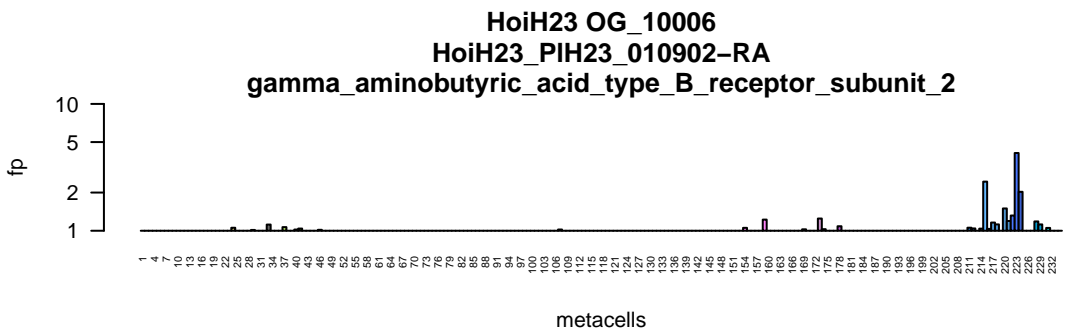
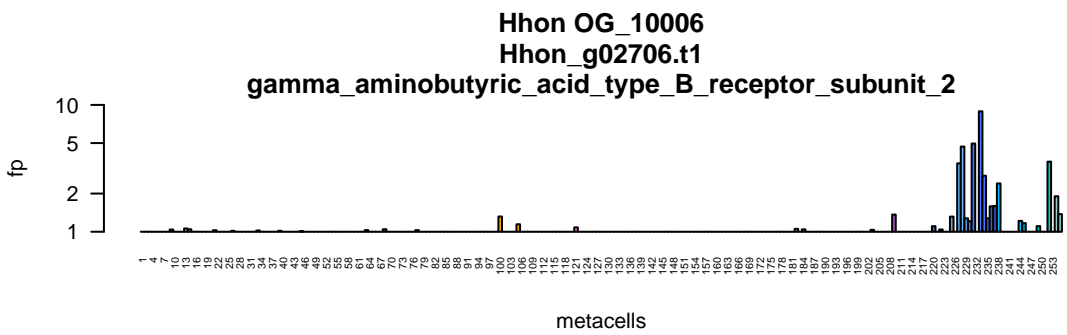
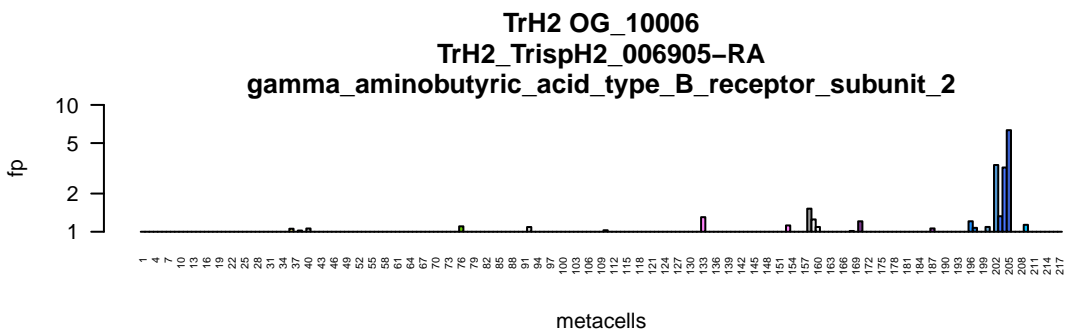
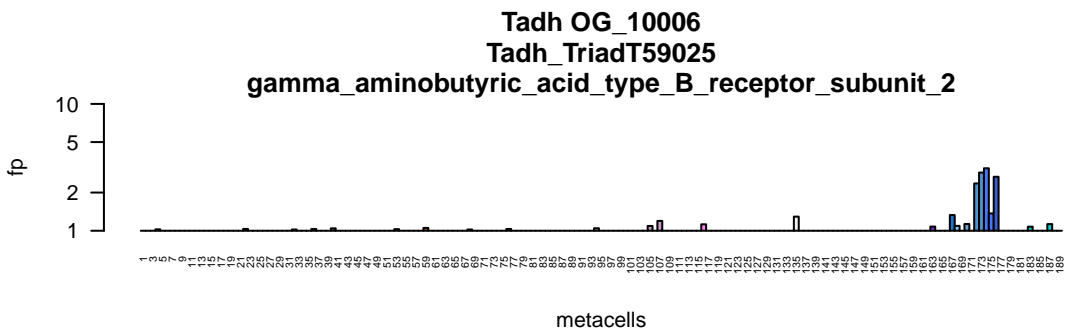
HoiH23 OG_9610
HoiH23_PIH23_005684-RA

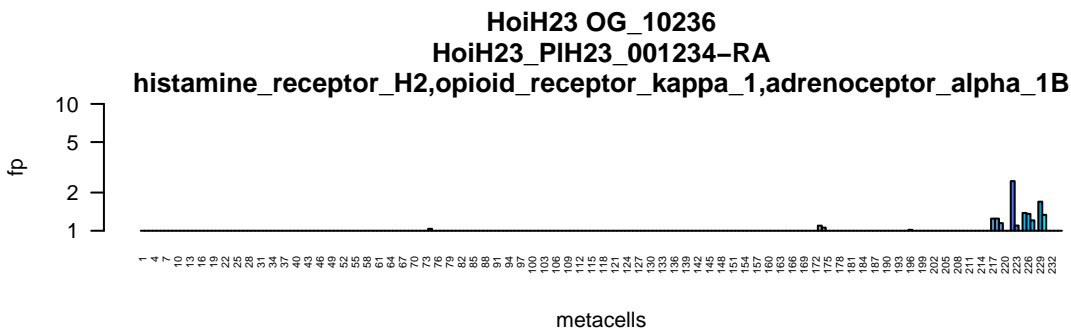
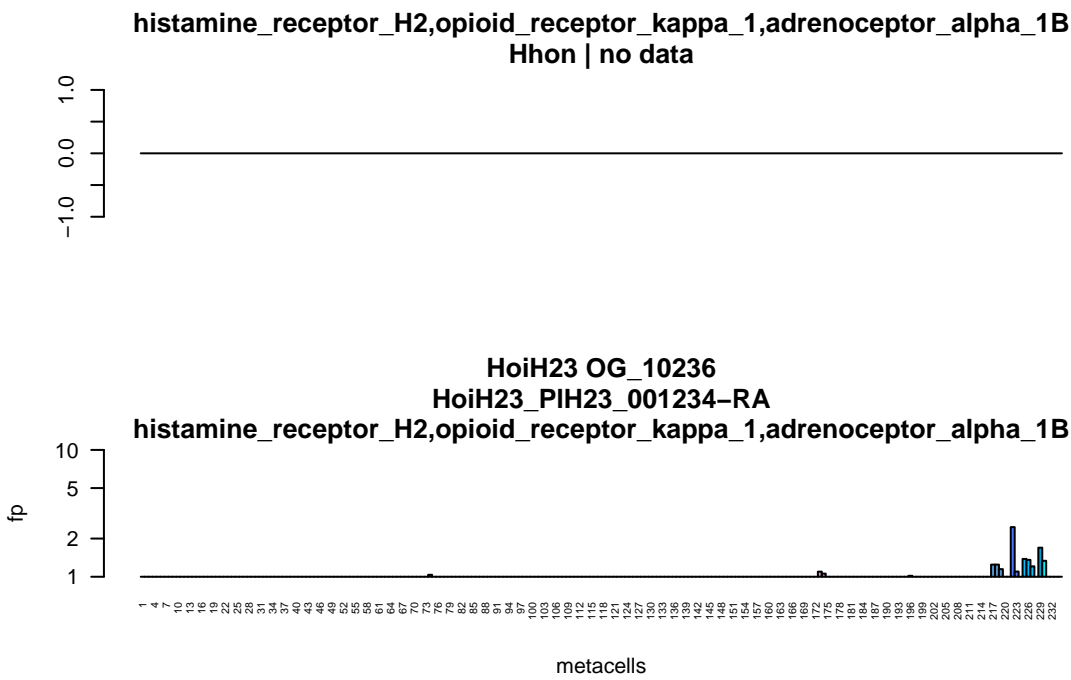
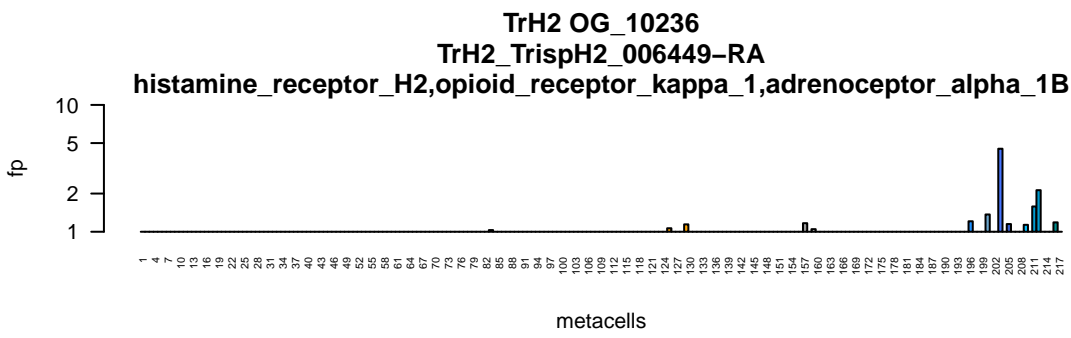
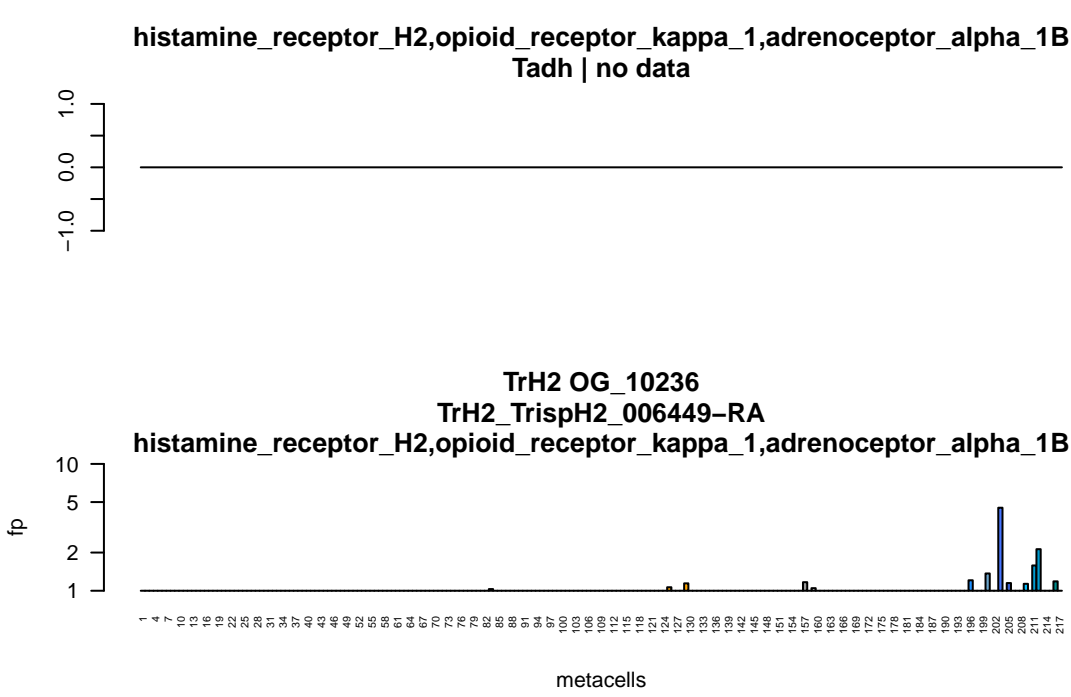




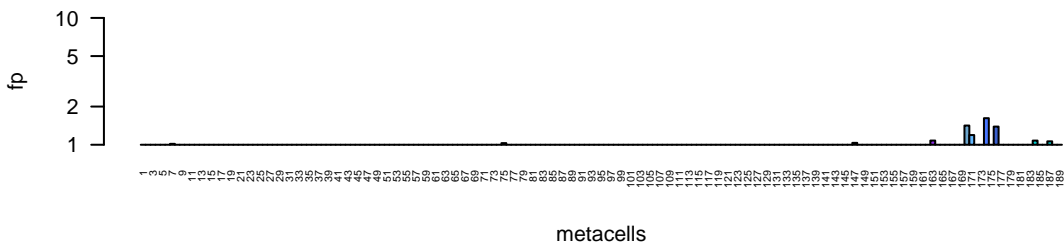




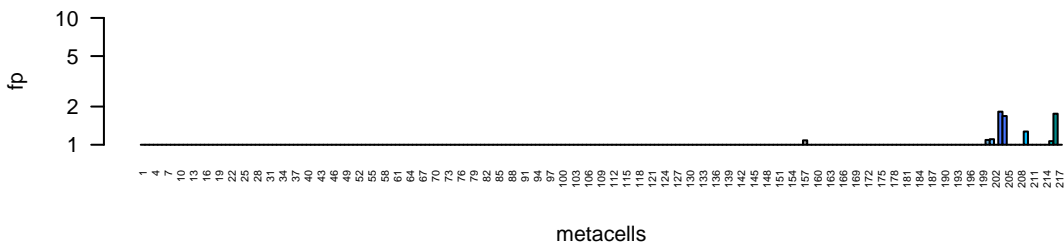




Tadh OG_10359
Tadh_wf_g9895.t1



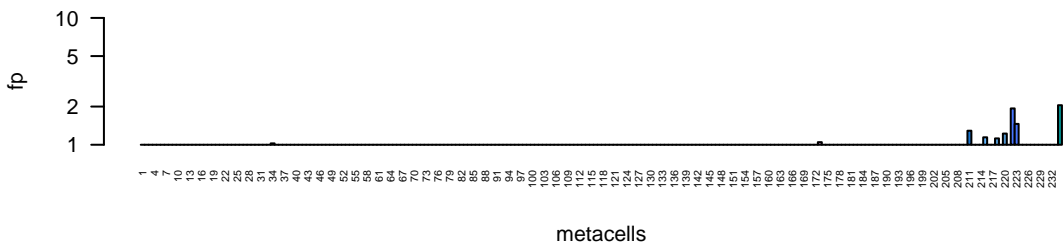
TrH2 OG_10359
TrH2_TrispH2_008852-RA

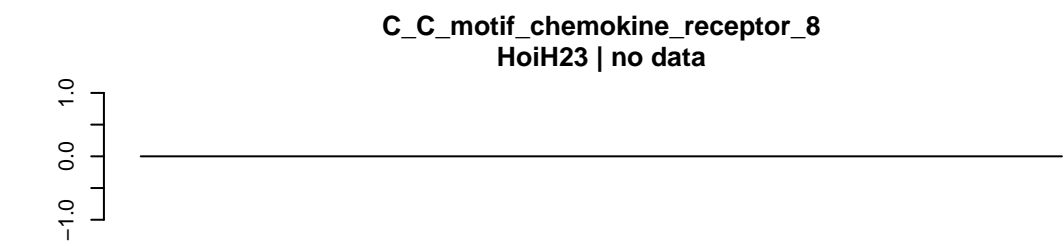
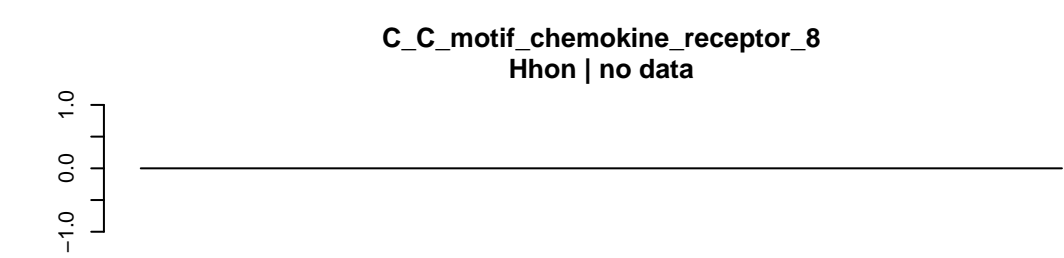
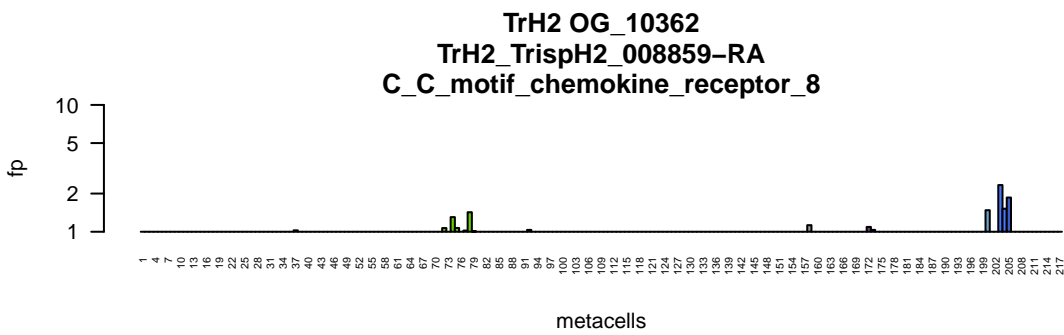
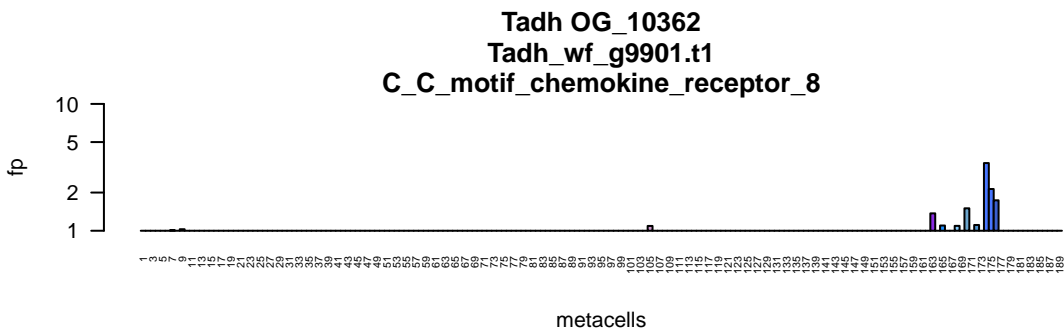


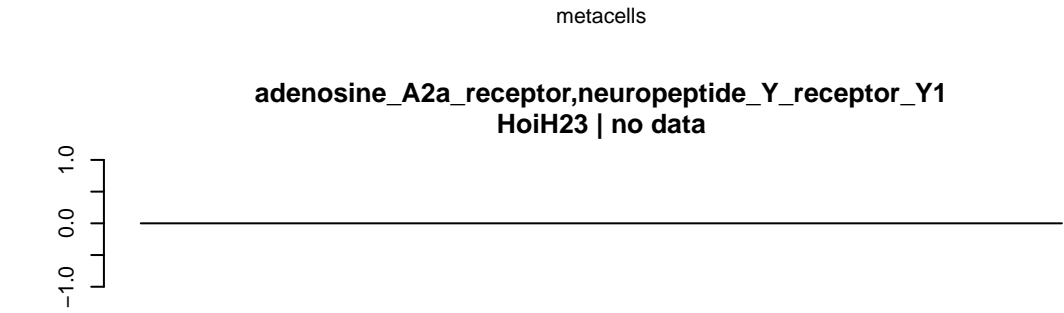
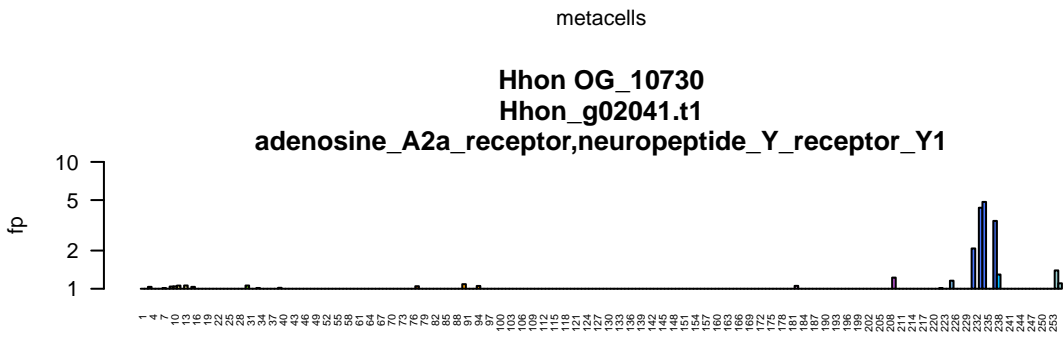
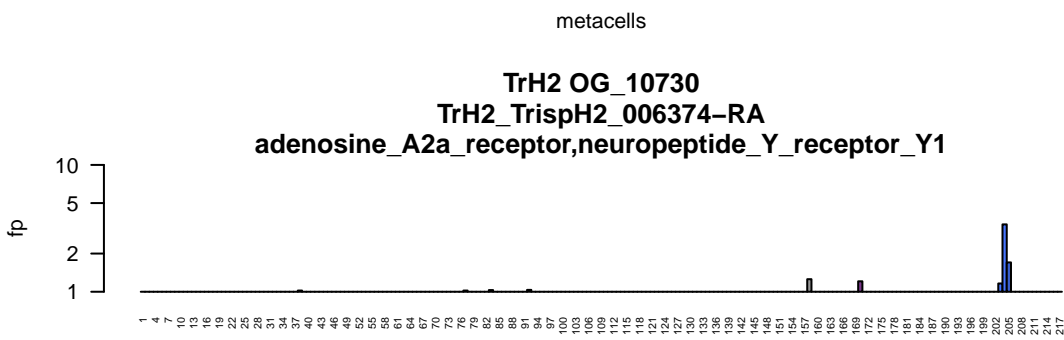
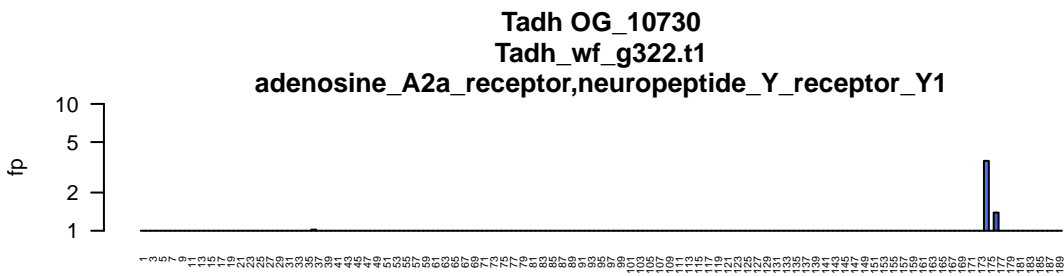
Hhon | no data

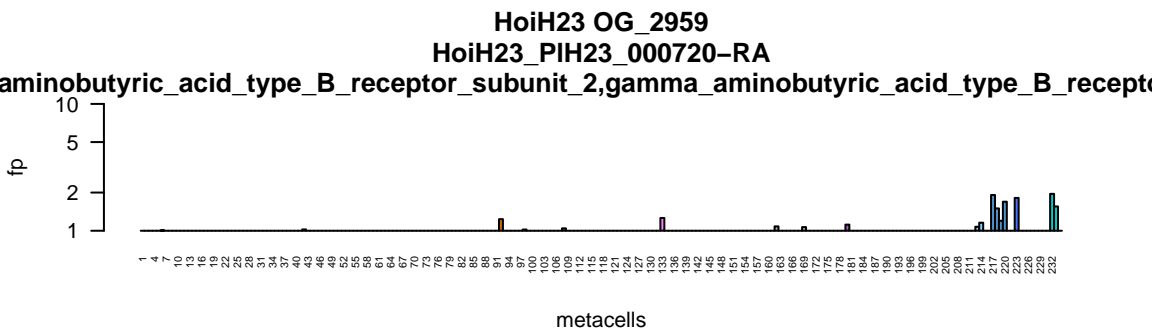
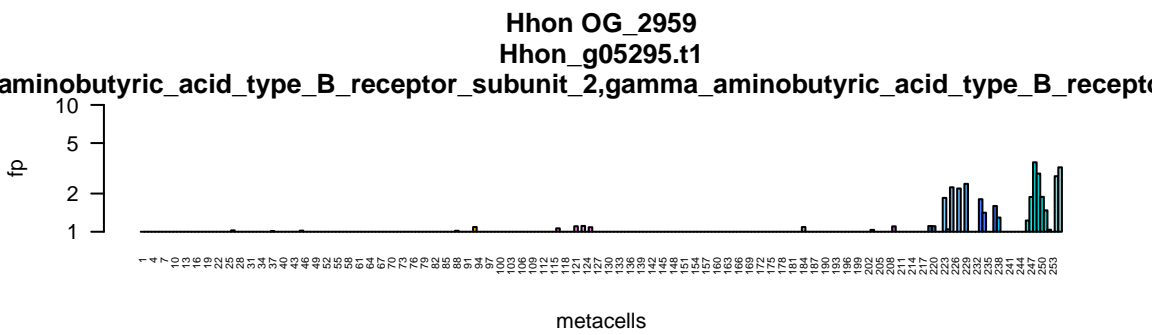
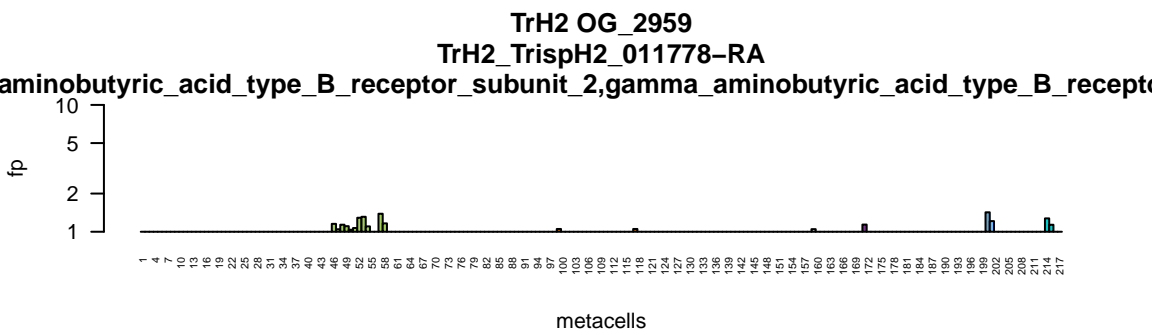
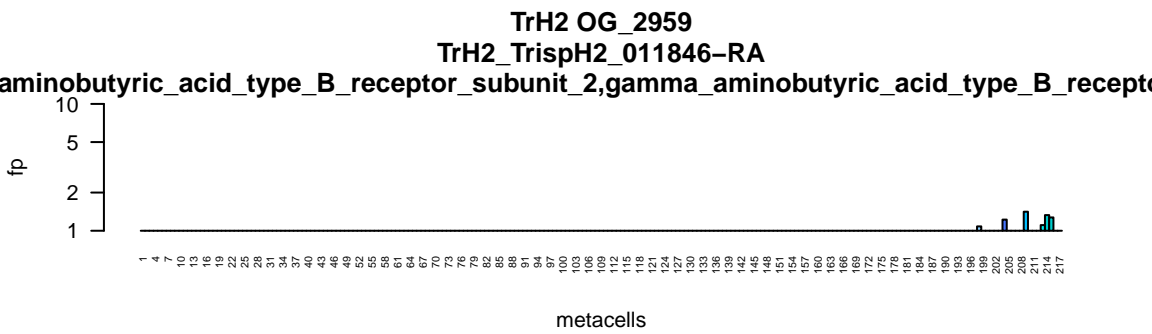
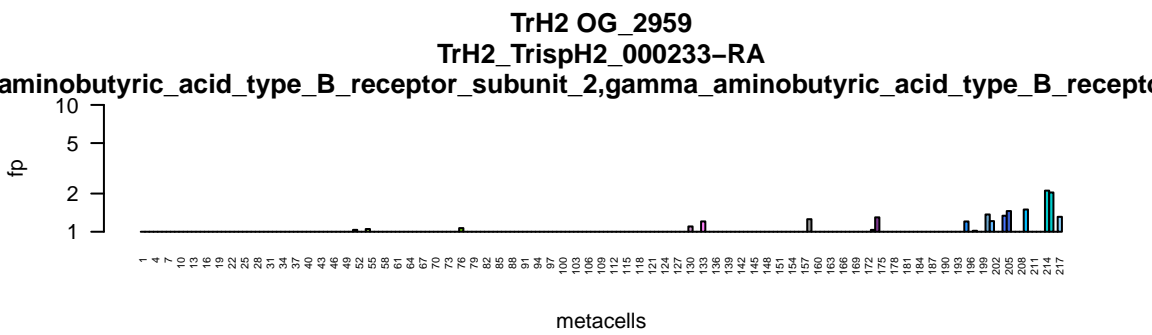
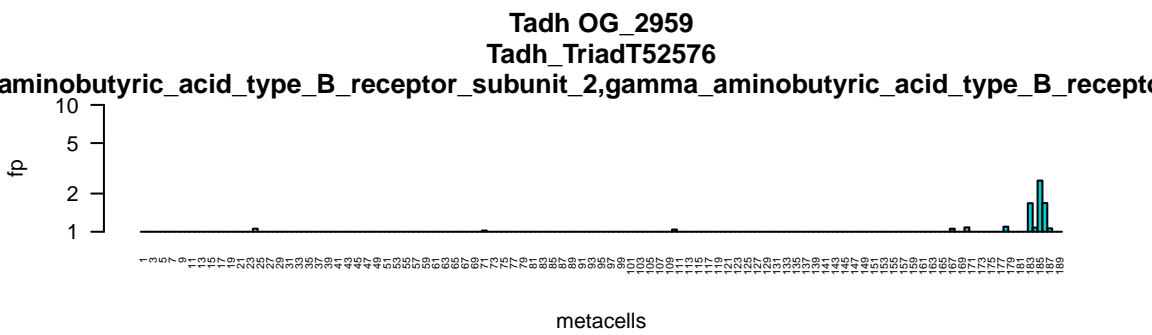
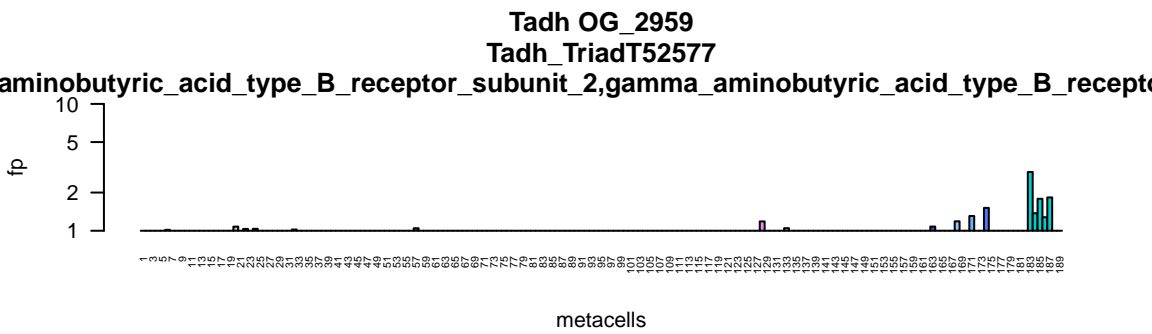


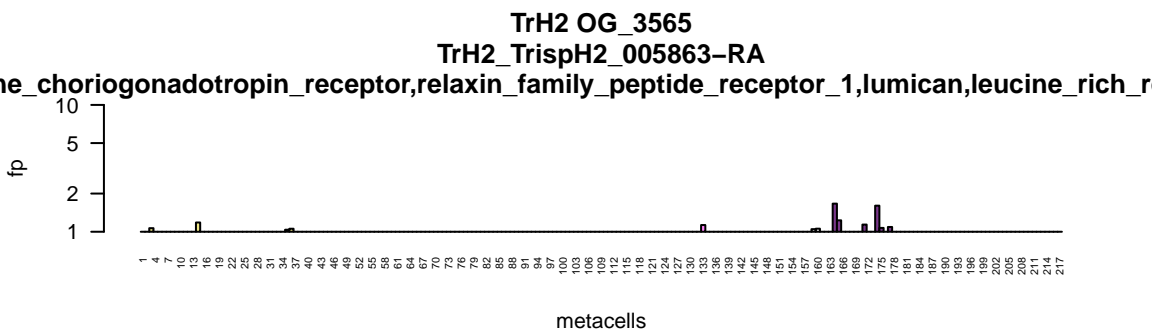
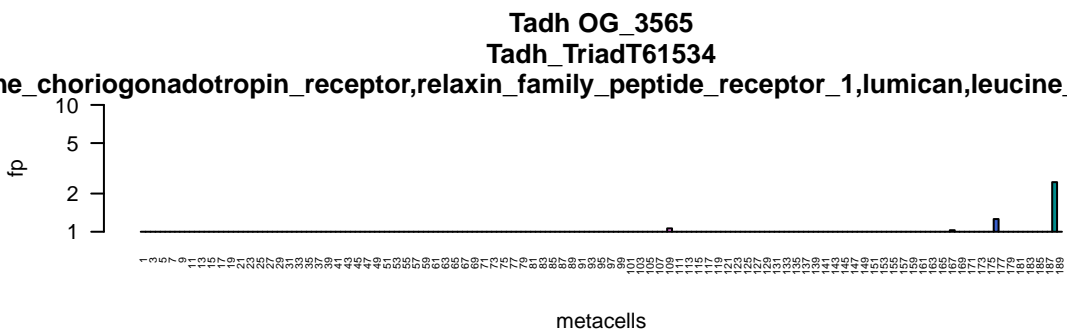
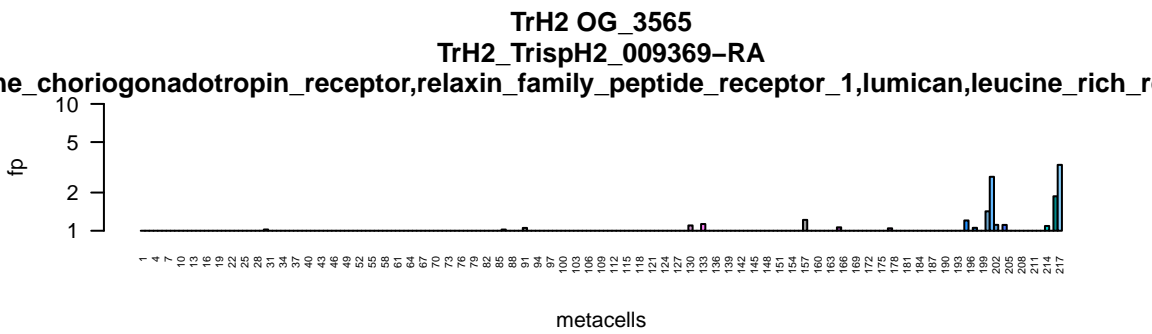
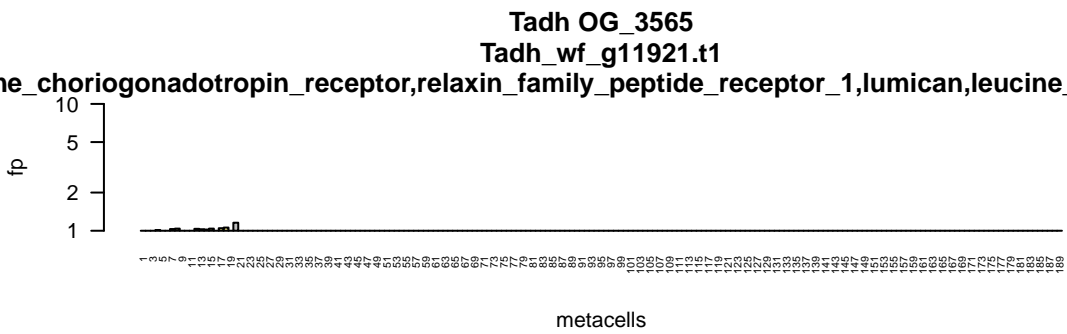
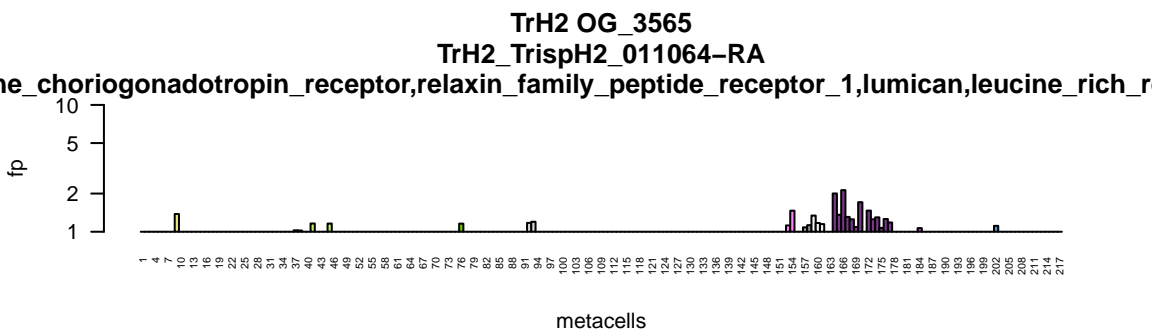
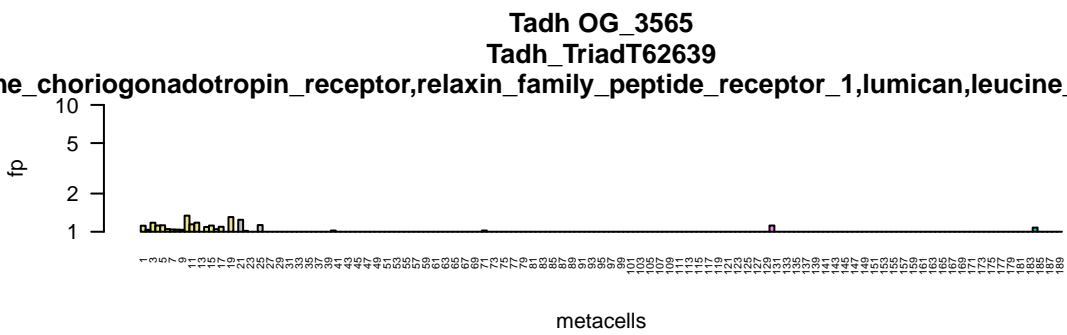
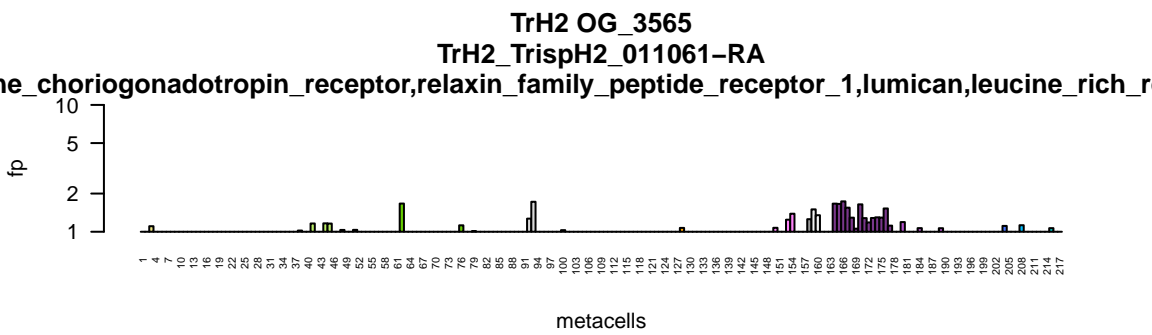
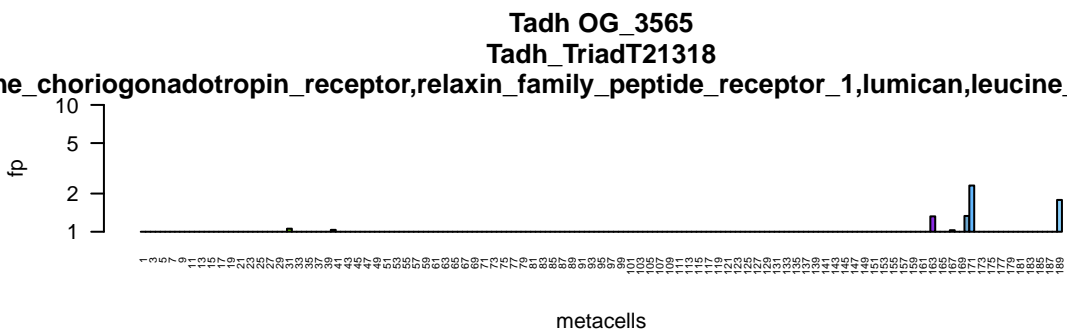
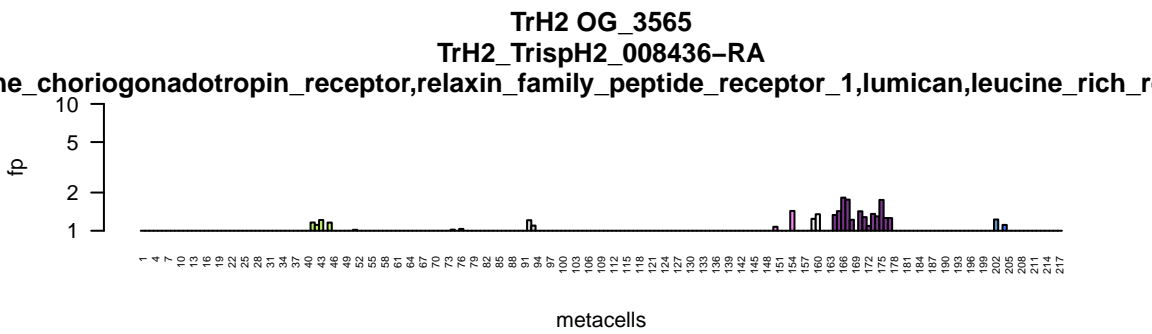
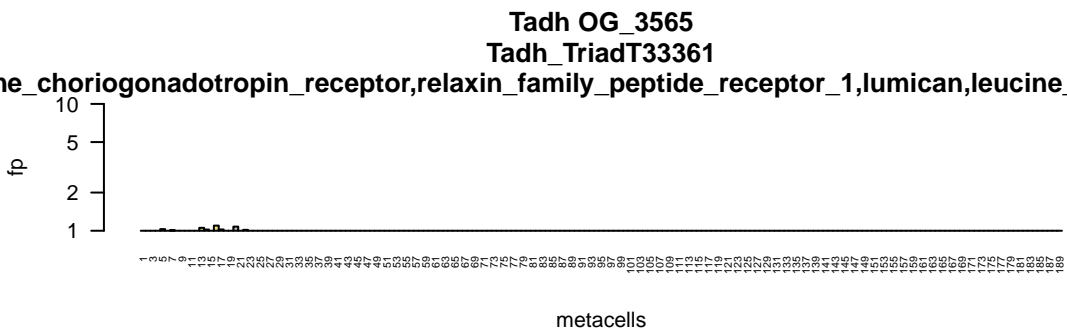
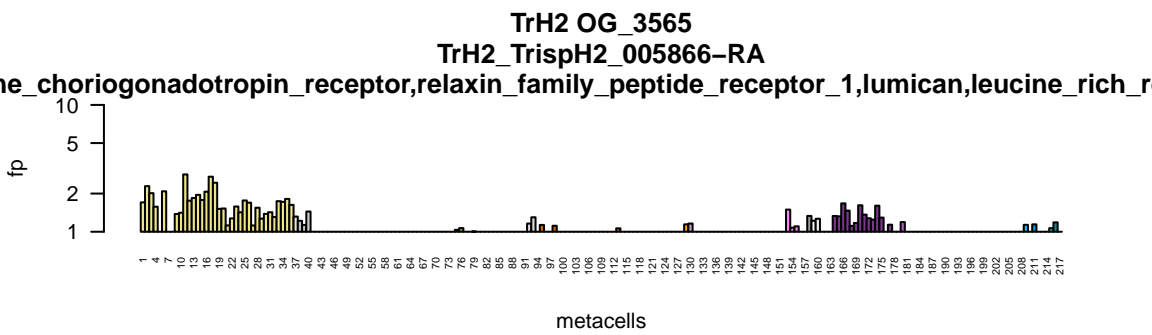
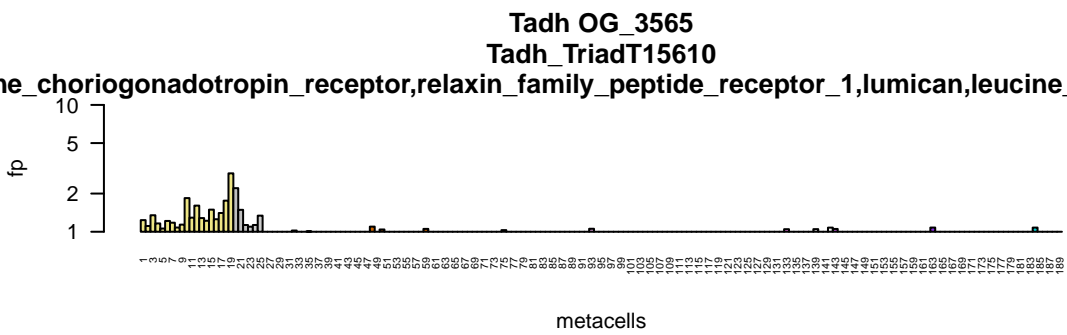
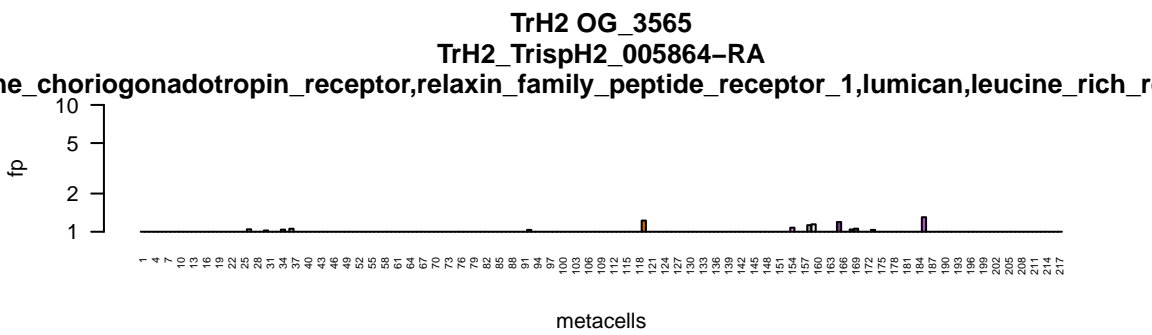
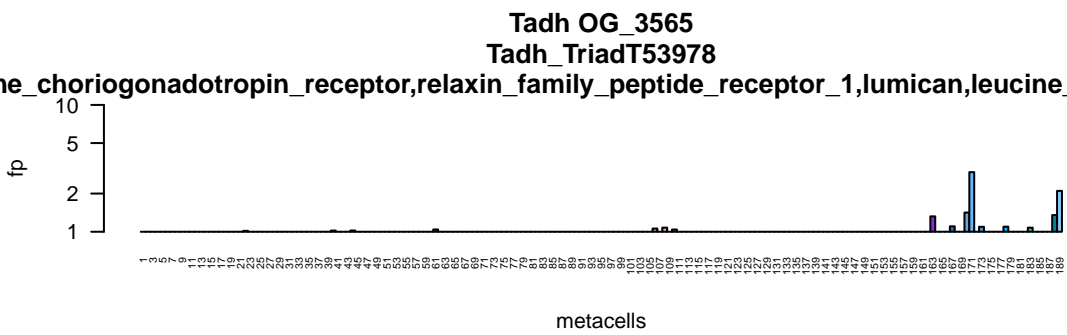
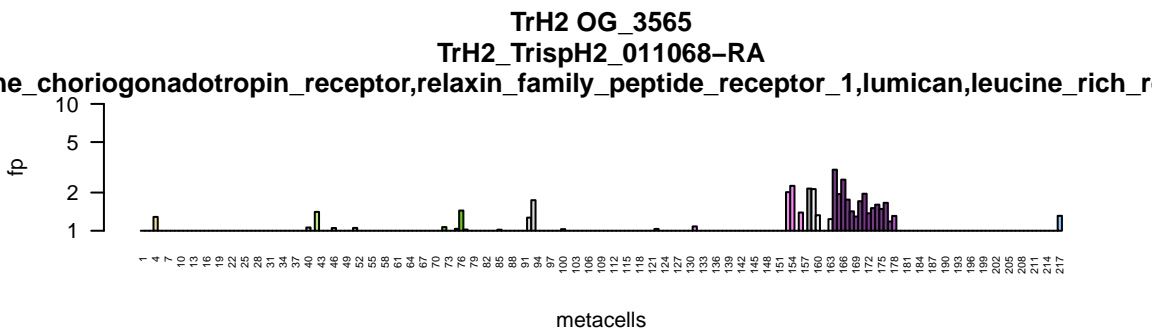
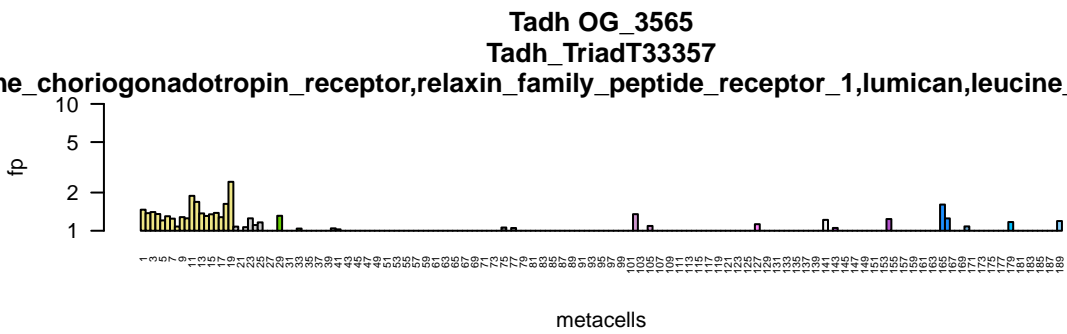
HoiH23 OG_10359
HoiH23_PIH23_008471-RA

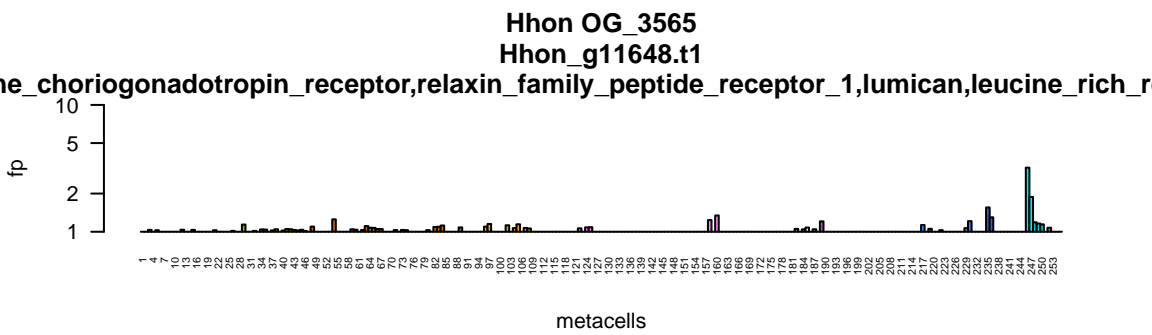
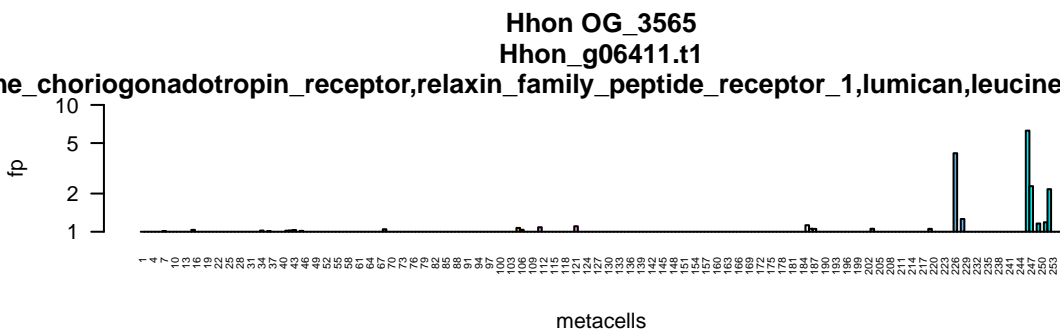
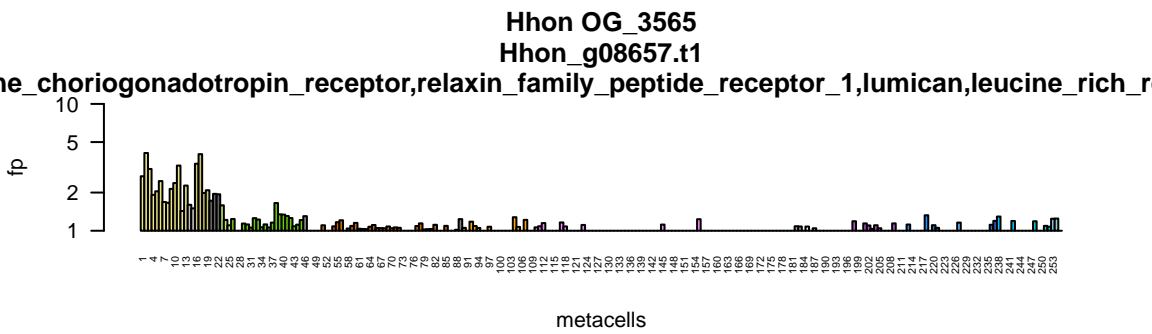
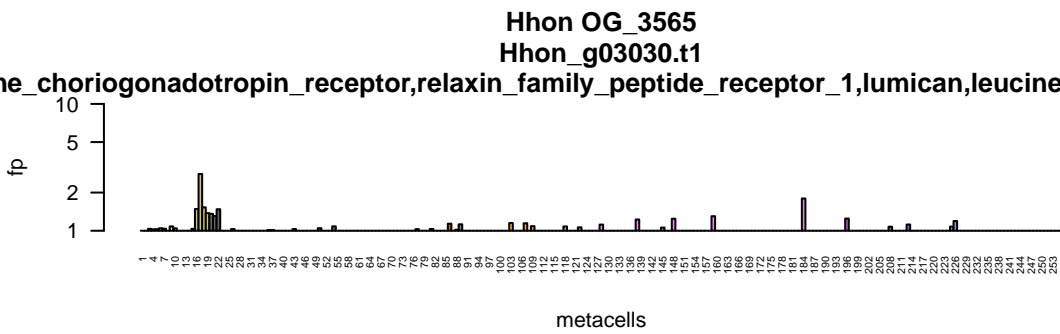
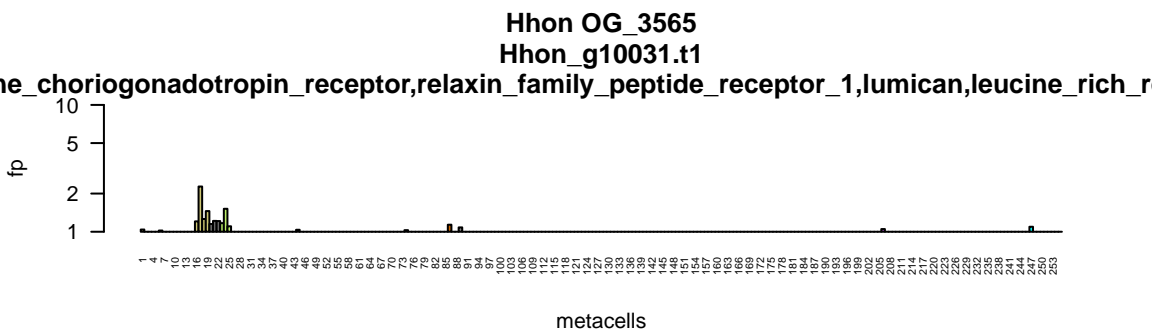
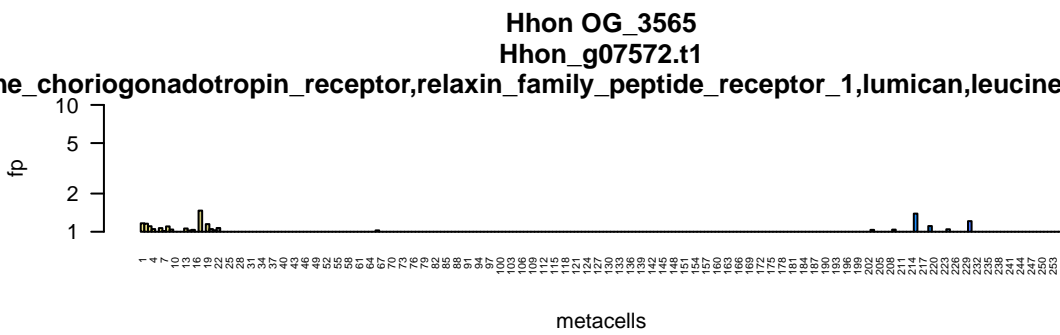
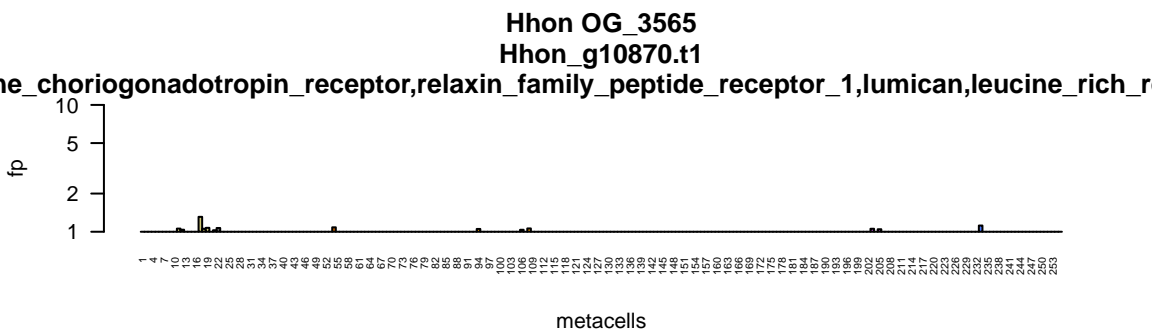
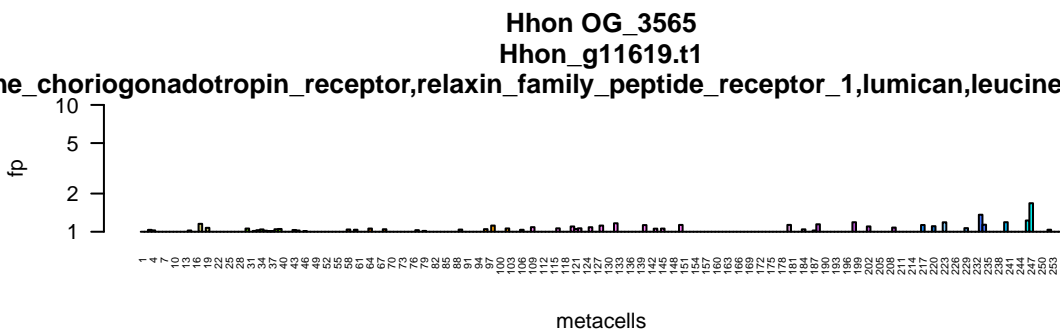
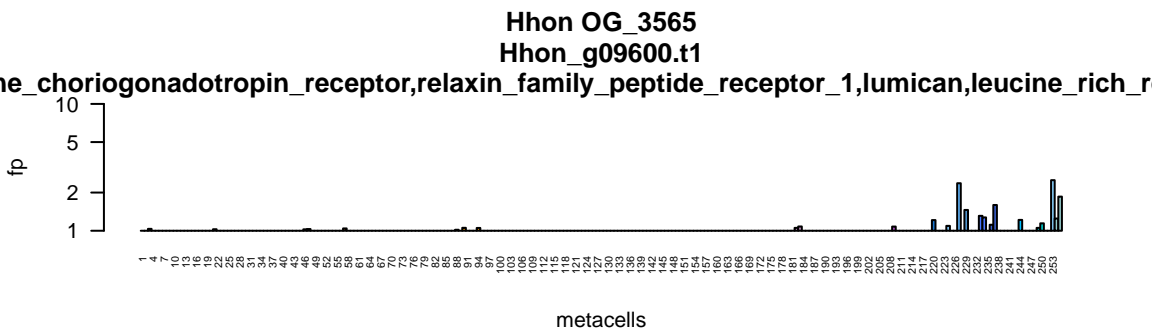
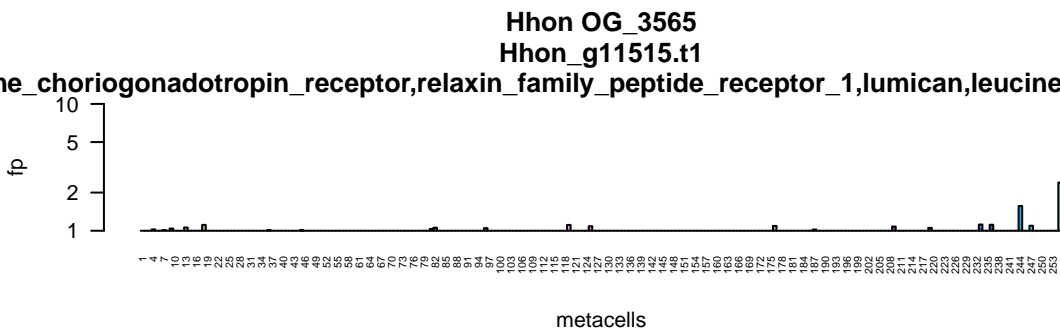
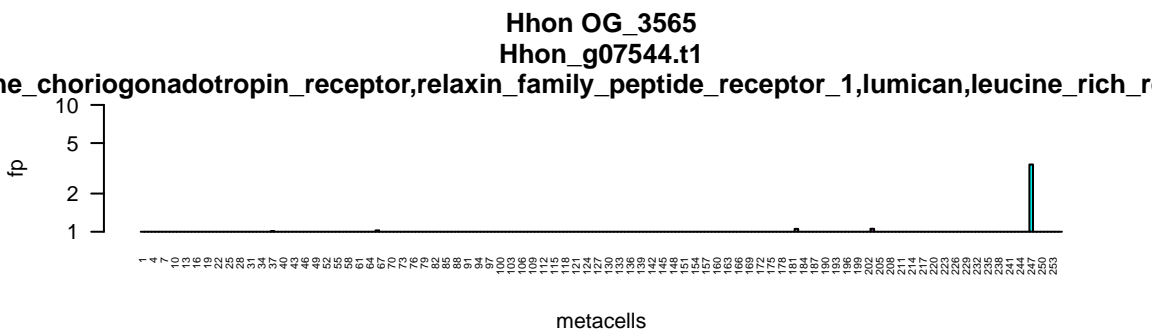
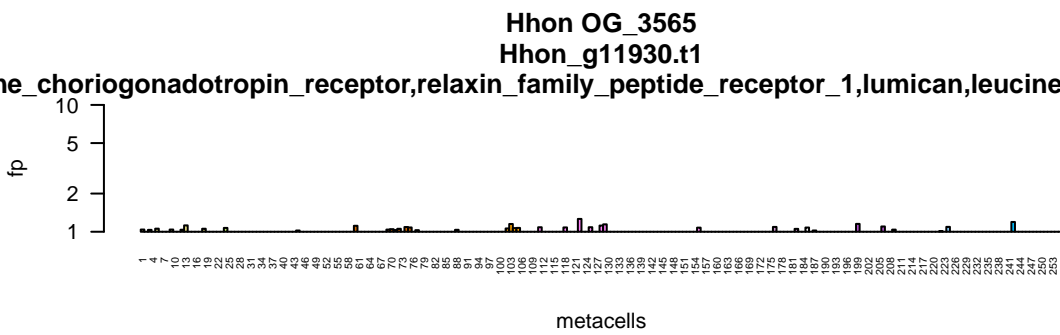
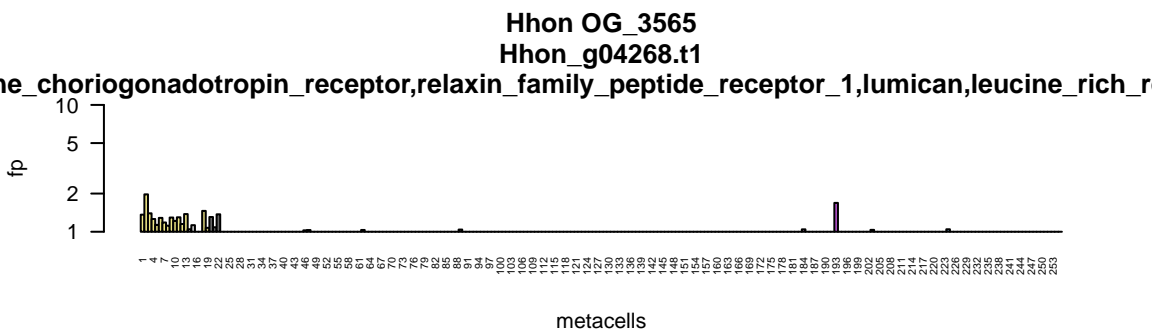
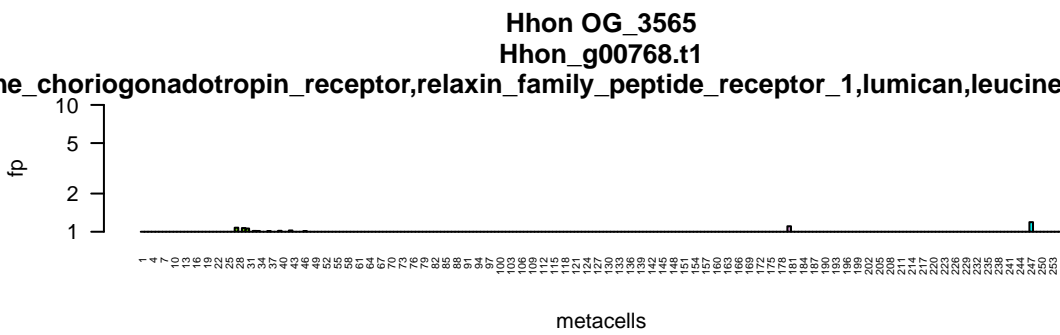
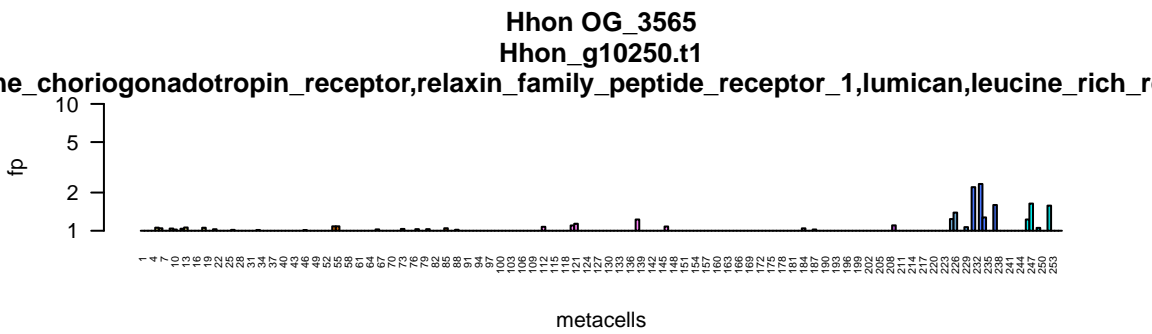
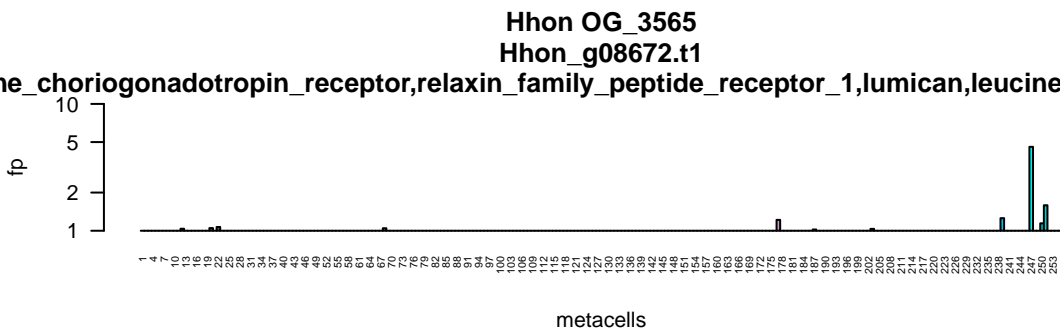


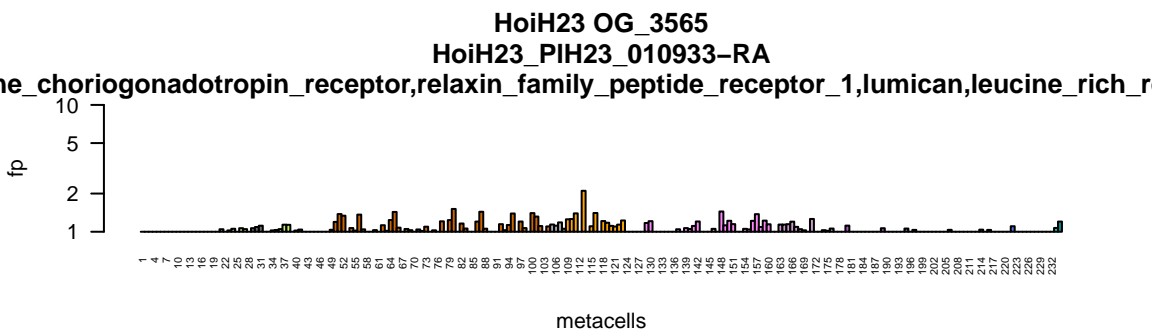
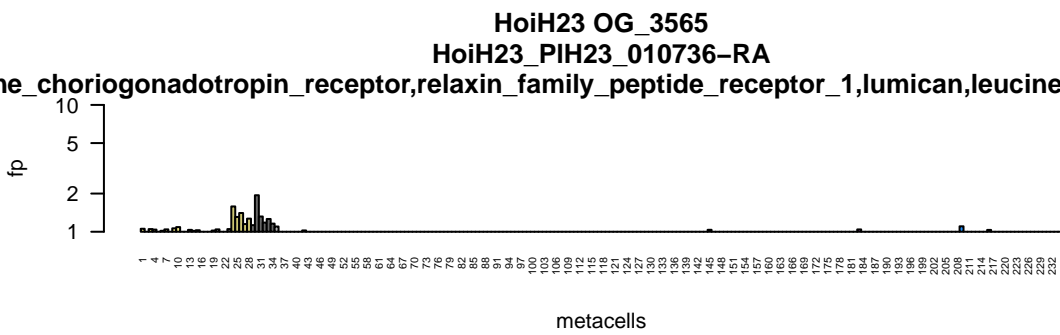
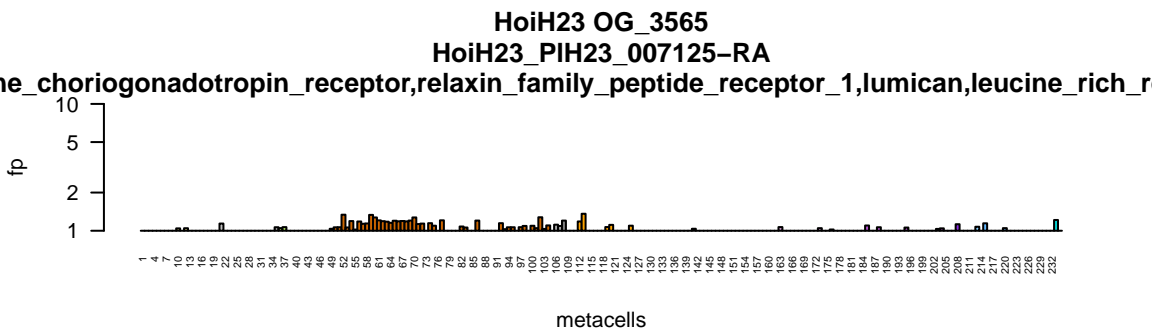
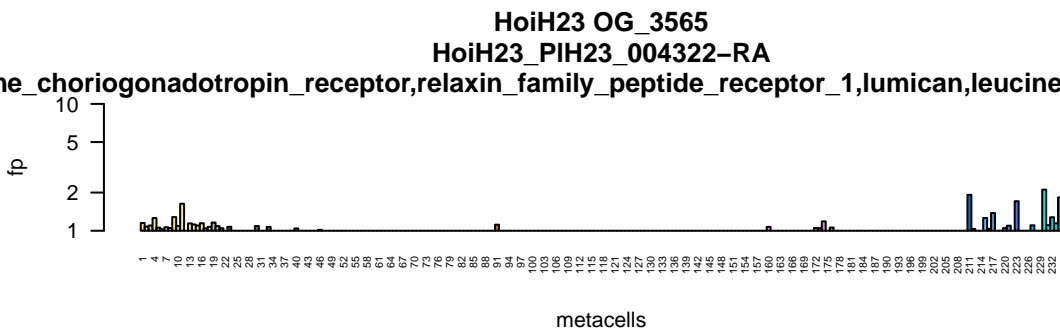
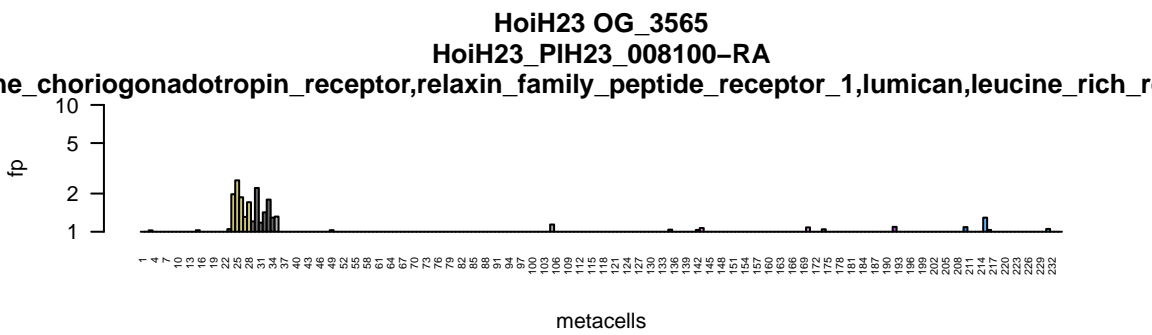
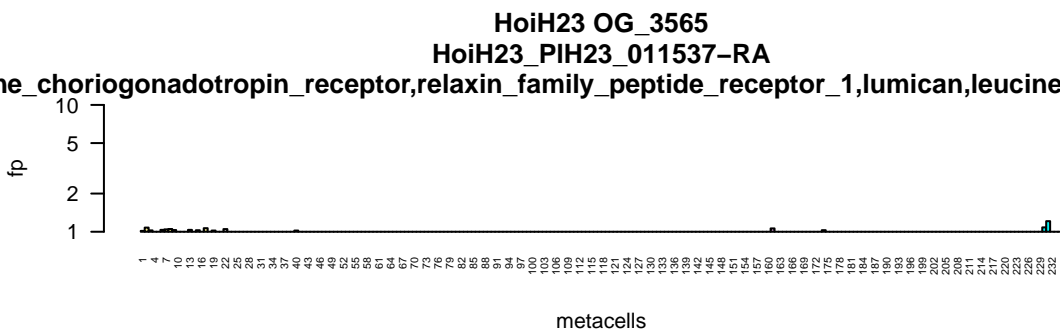
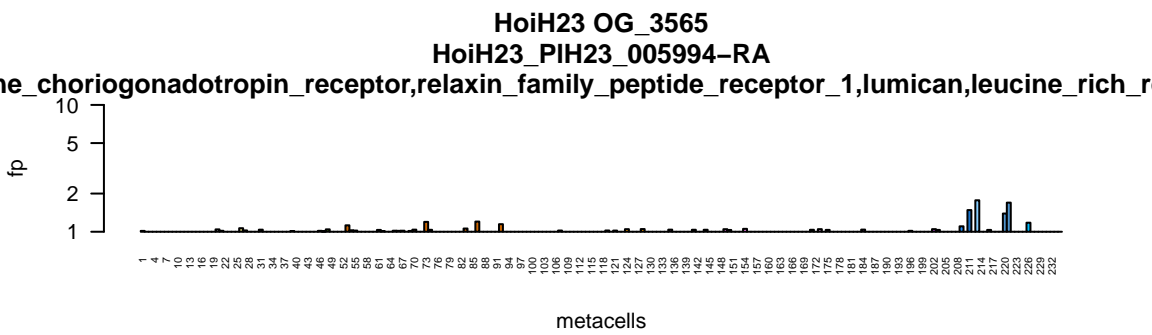
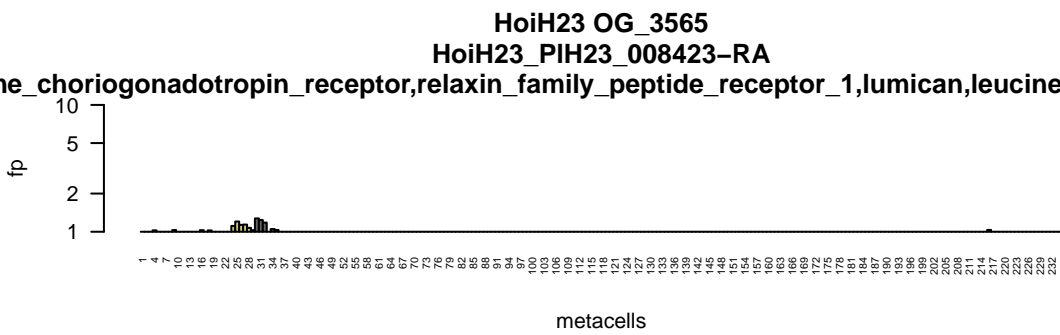
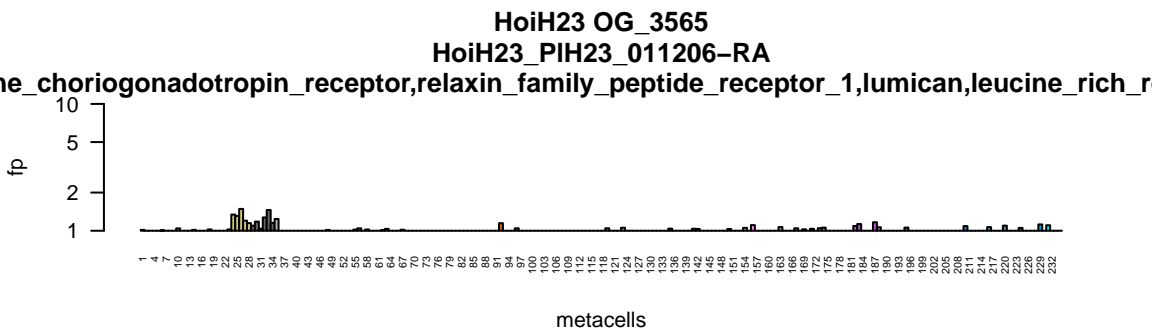
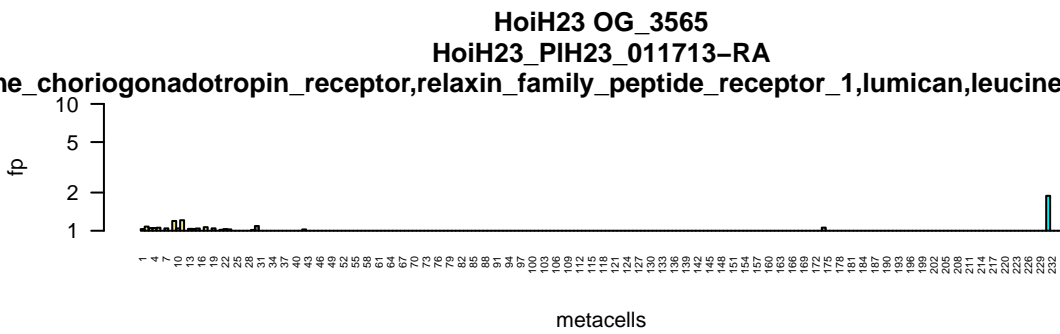
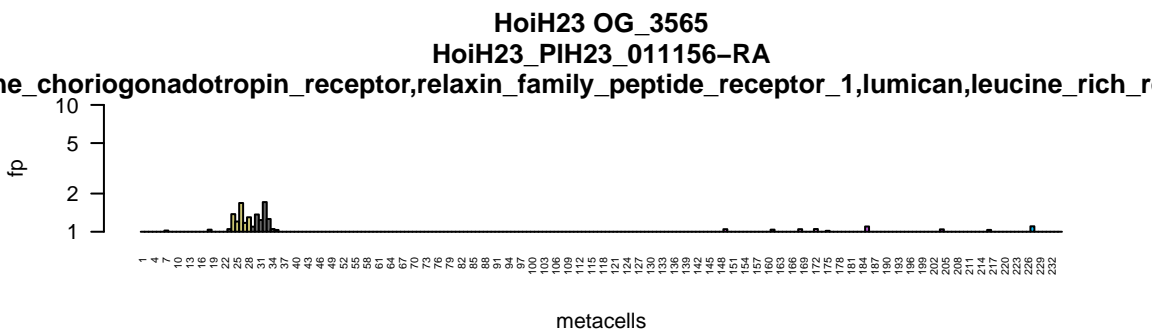
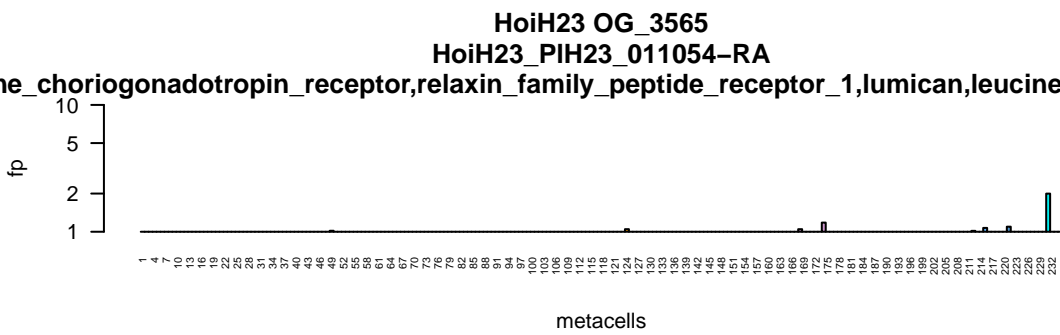
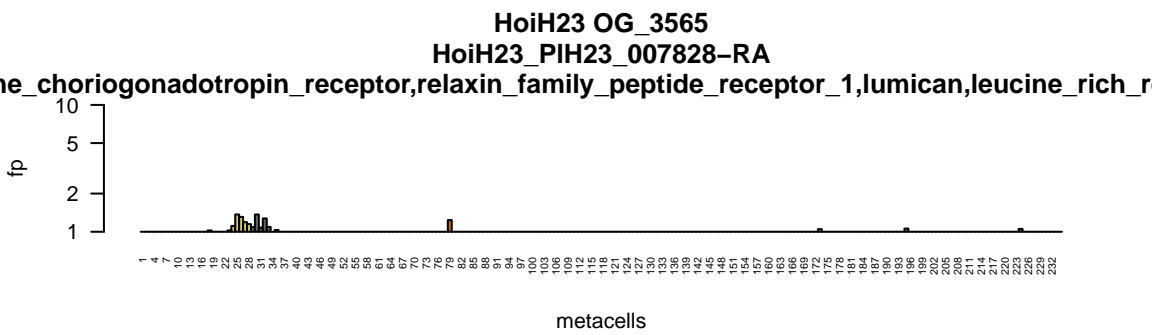
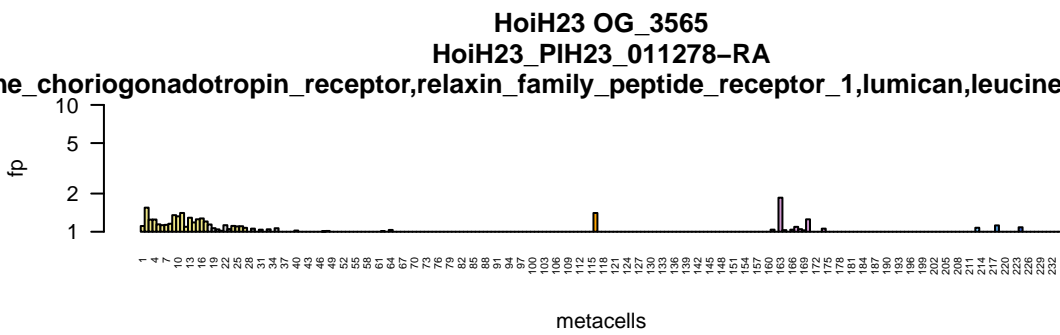
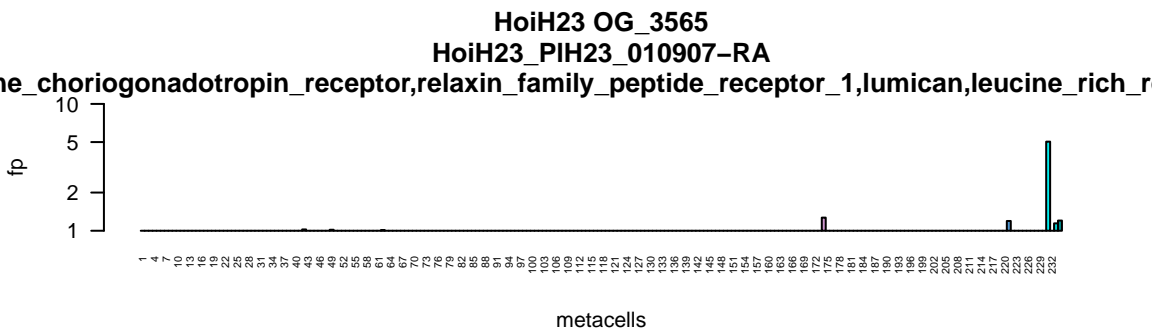
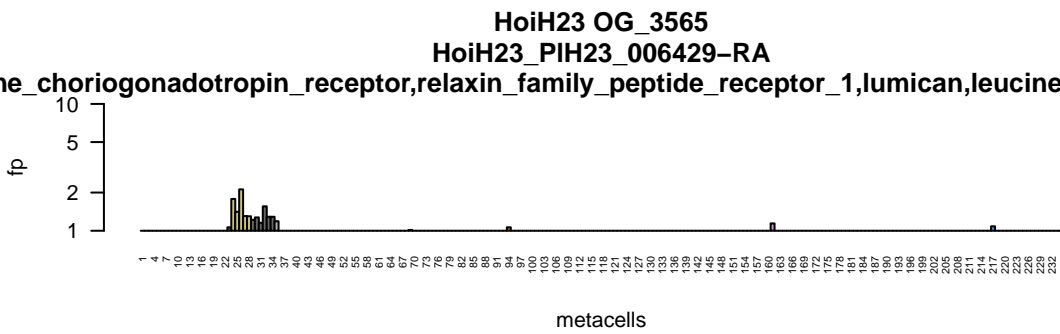


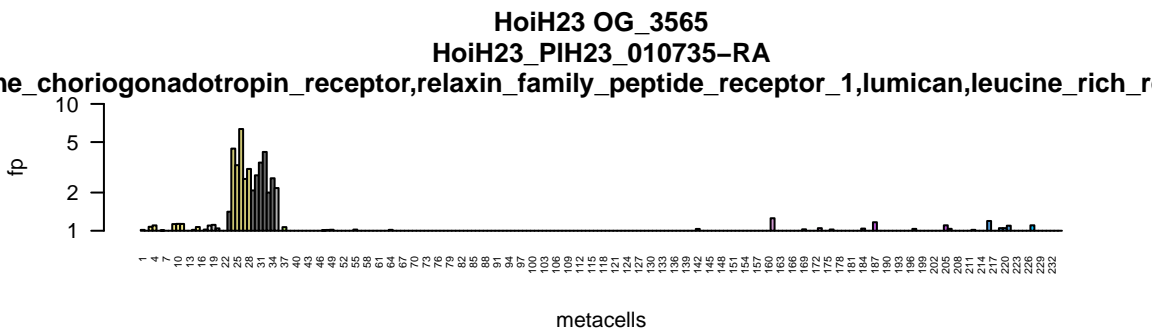
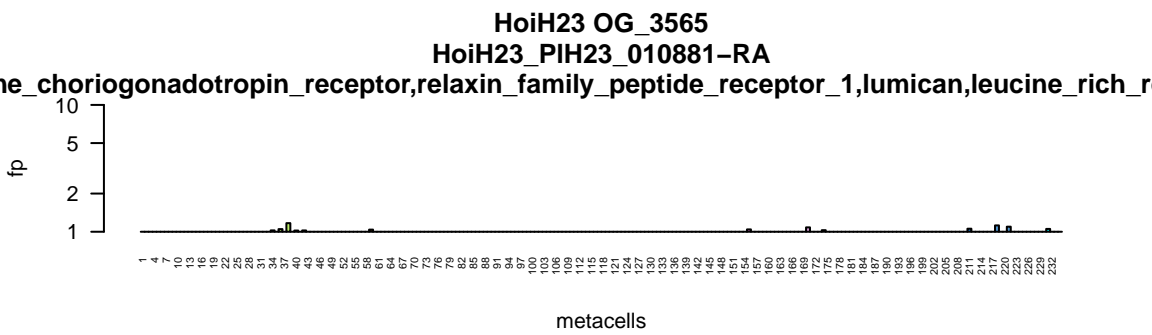
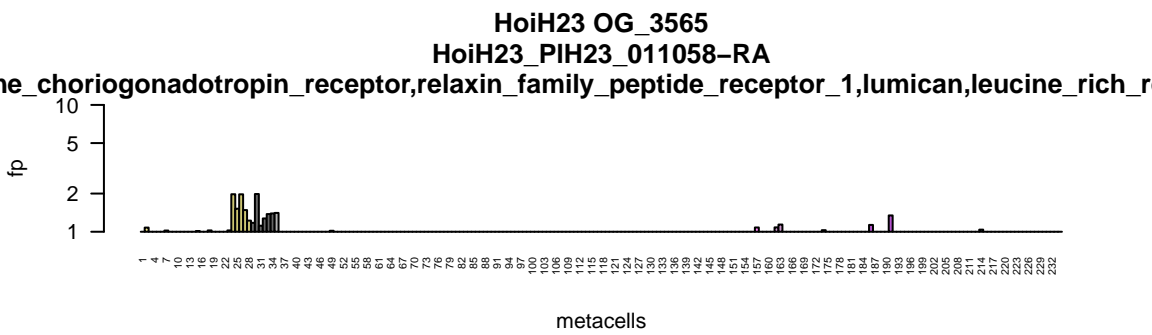
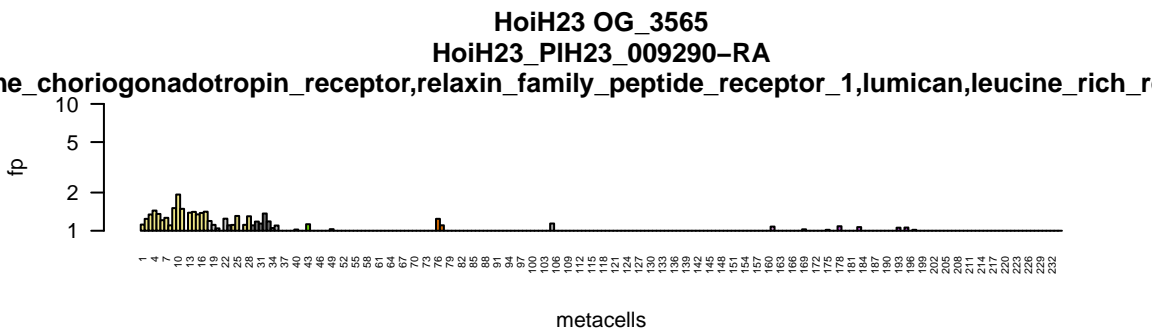


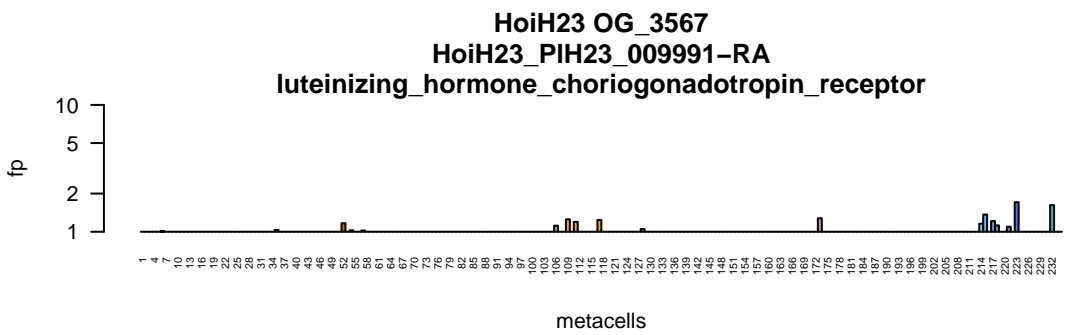
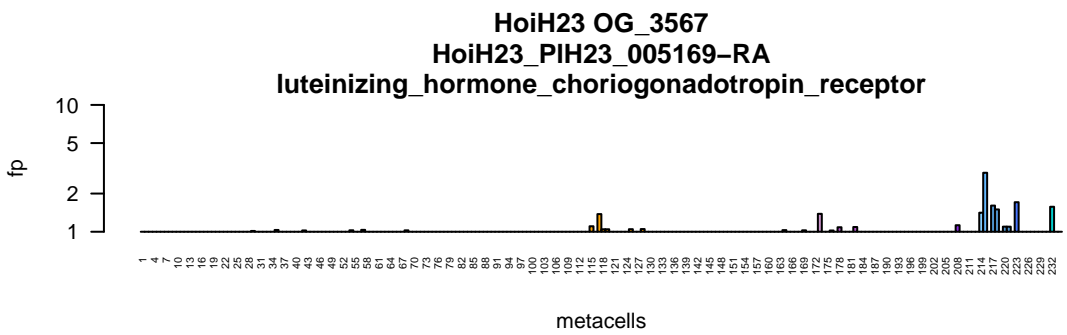
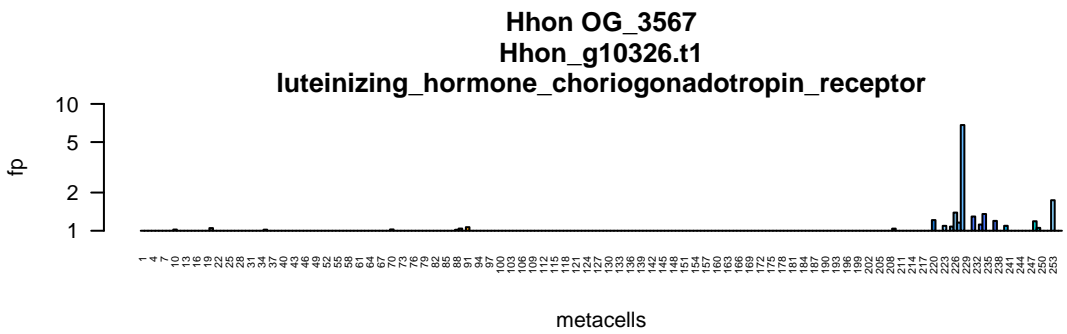
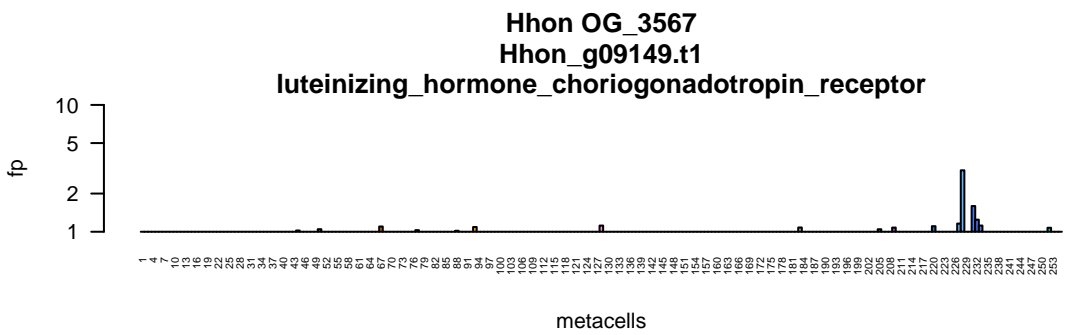
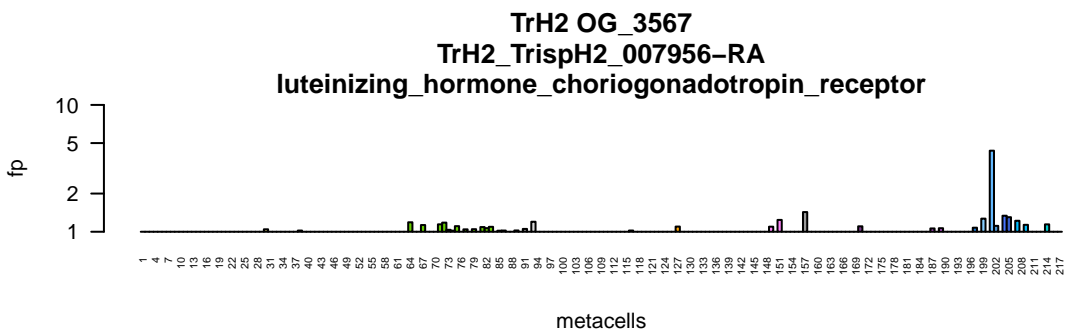
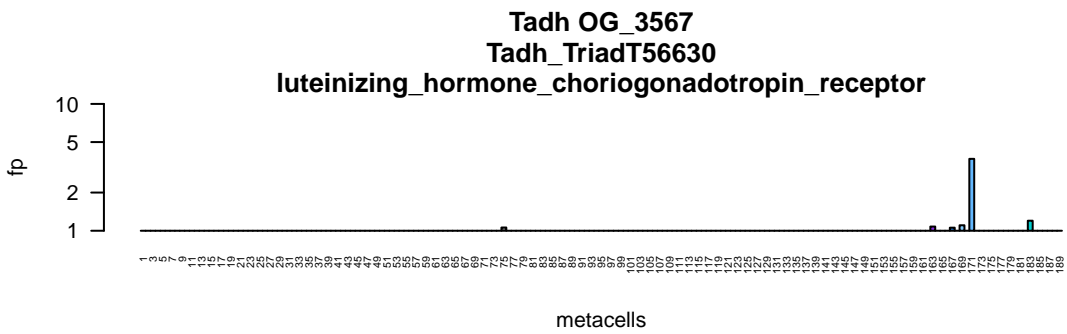








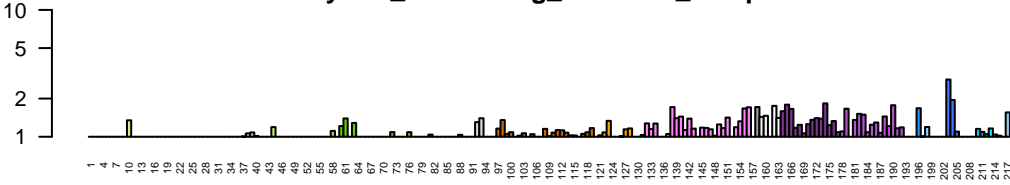




thyroid_stimulating_hormone_receptor
Tadh | no data

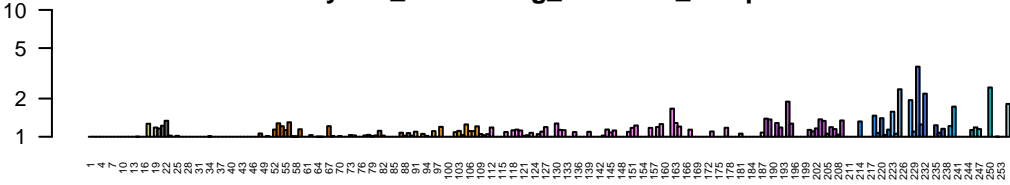


TrH2 OG_3593
TrH2_TrispH2_009078-RA
thyroid_stimulating_hormone_receptor



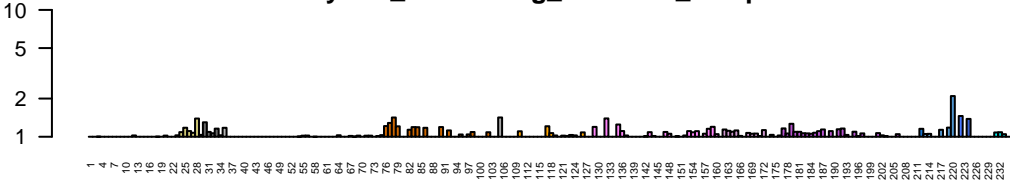
metacells

Hhon OG_3593
Hhon_g02067.t1
thyroid_stimulating_hormone_receptor

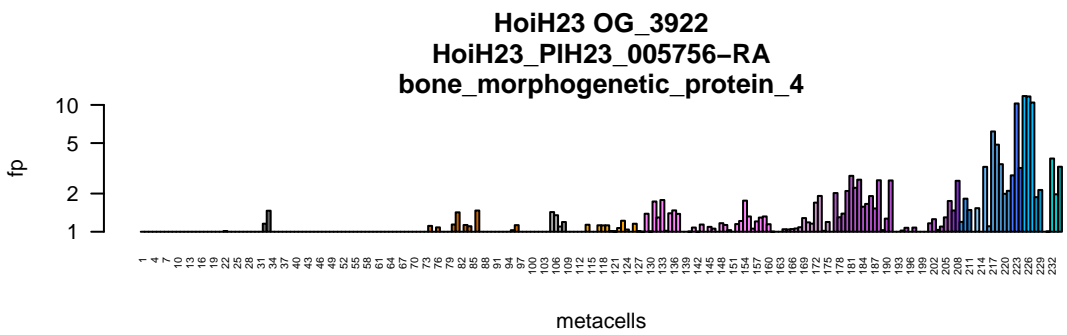
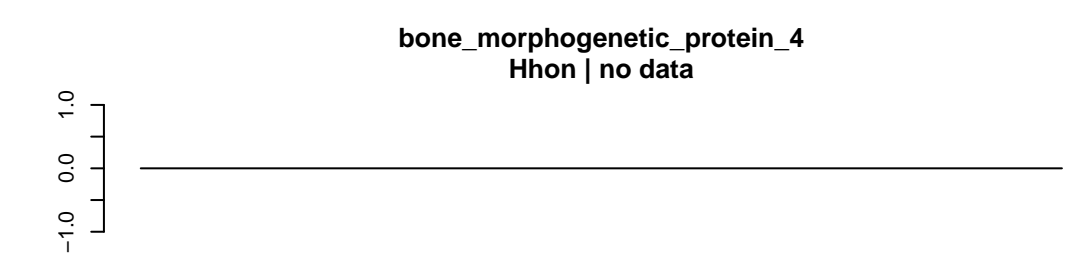
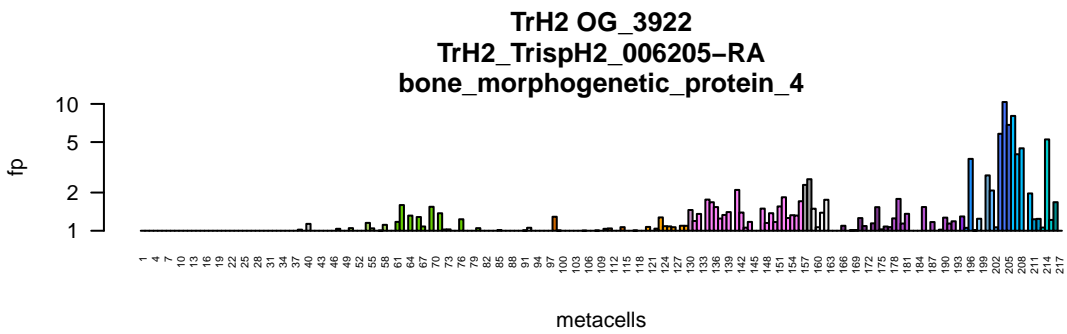
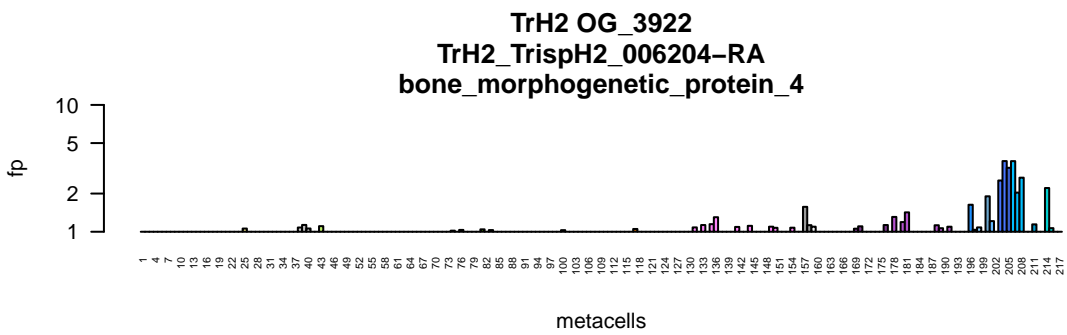
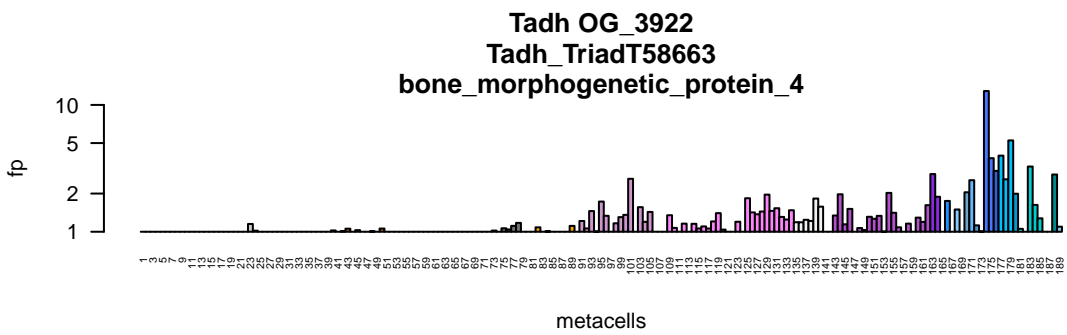
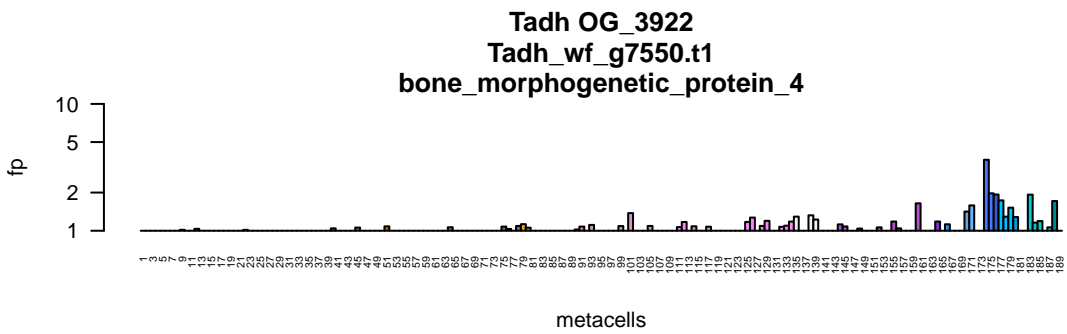


metacells

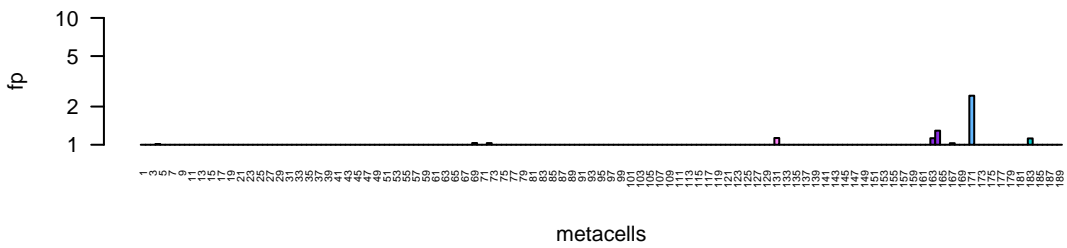
HoiH23 OG_3593
HoiH23_PIH23_010007-RA
thyroid_stimulating_hormone_receptor



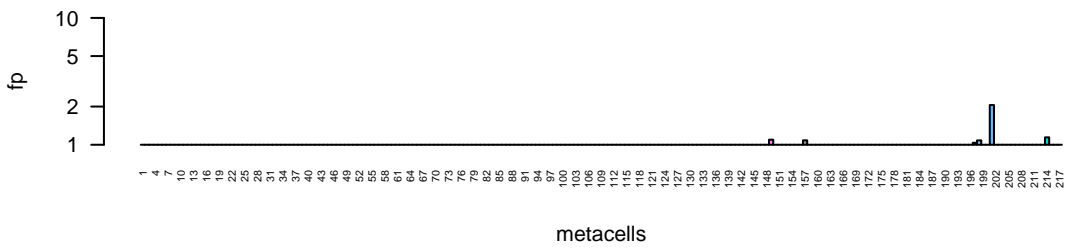
metacells



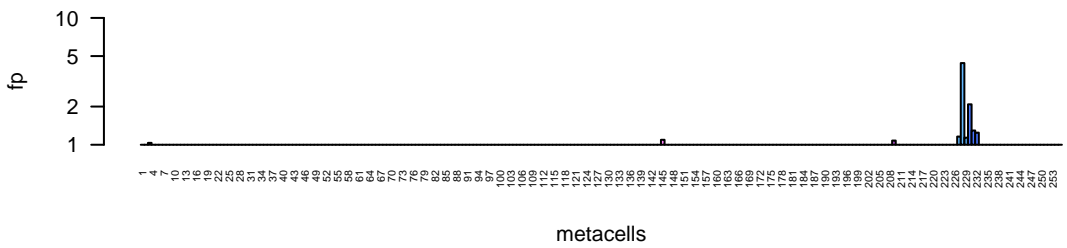
Tadh OG_4120
Tadh_TriadT57620



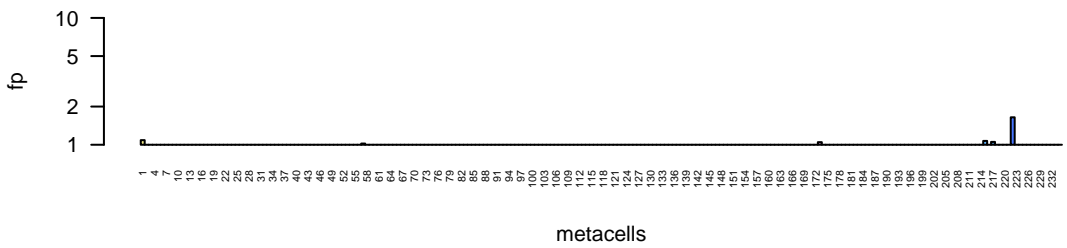
TrH2 OG_4120
TrH2_TrispH2_007435-RA



Hhon OG_4120
Hhon_g03439.t1



HoiH23 OG_4120
HoiH23_PIH23_007154-RA



Tadh_TriadT62237



Tadh_TriadT62236



Tadh_wf_g11342.t1



Tadh_TriadT62238



Tadh_wf_g11350.t1



Tadh_TriadT3775



Tadh_wf_g11493.t1



2_TrispH2_008282-RA



TrH2_TrispH2_010704-RA



TrH2_TrispH2_010702-RA



TrH2_TrispH2_010703-RA



TrH2_TrispH2_010706-RA



TrH2_TrispH2_011713-RA



TrH2_TrispH2_007495-RA

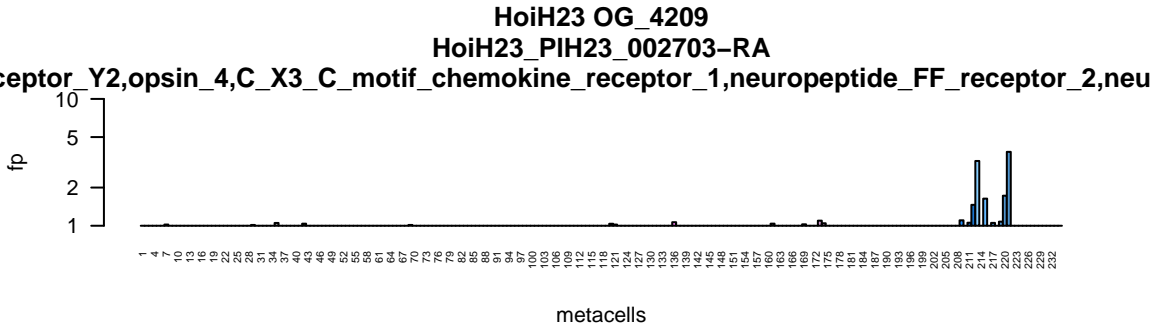
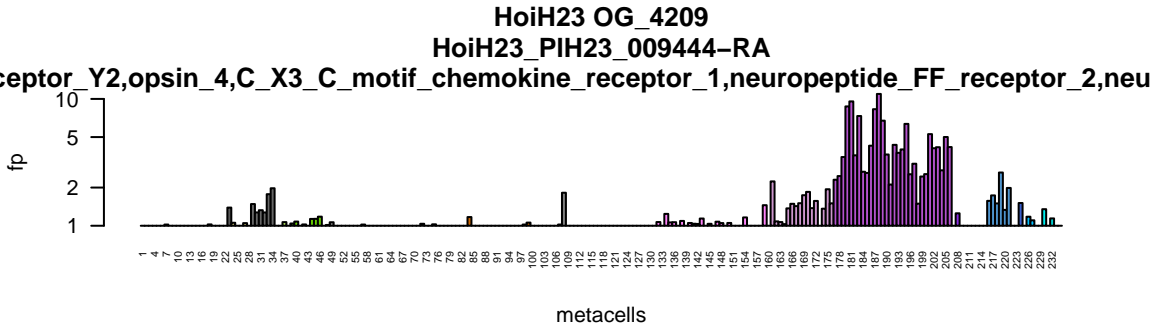
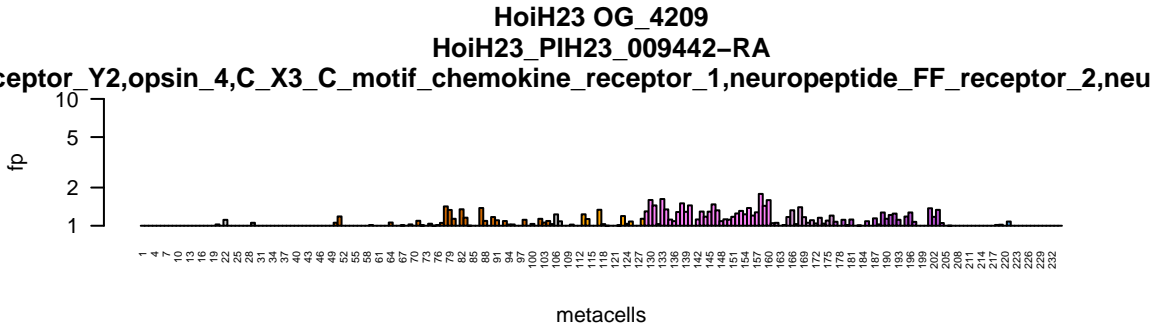
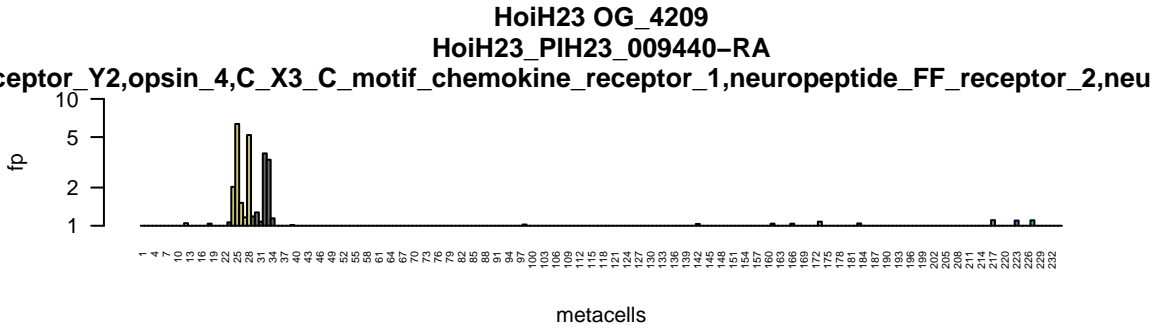
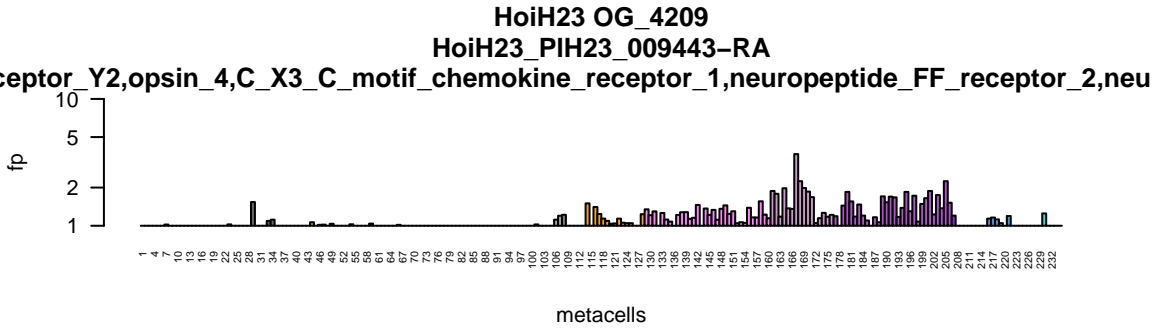
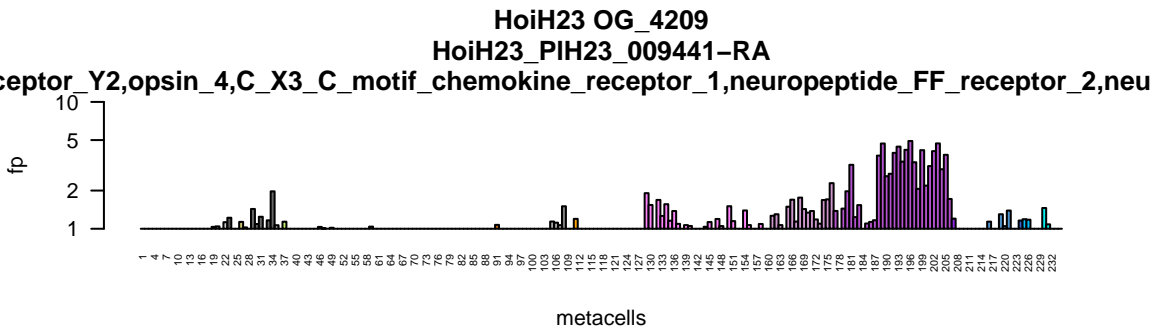
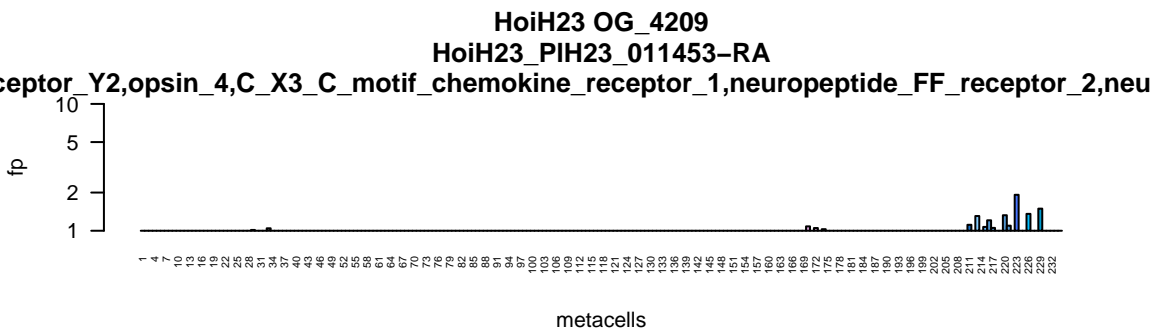


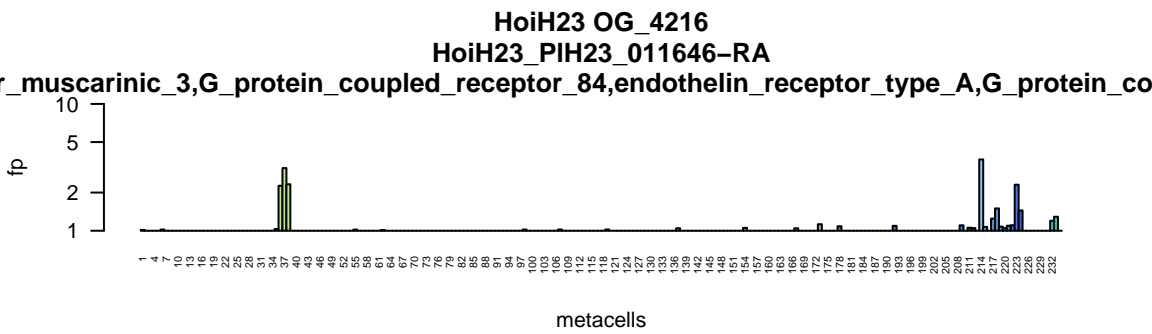
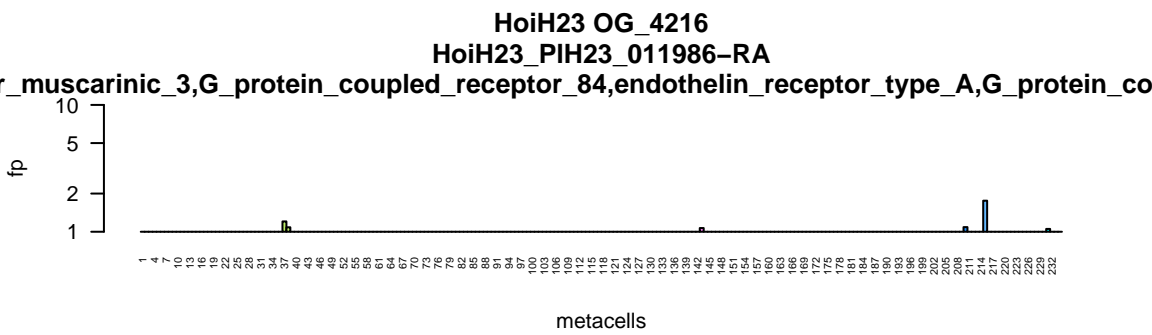
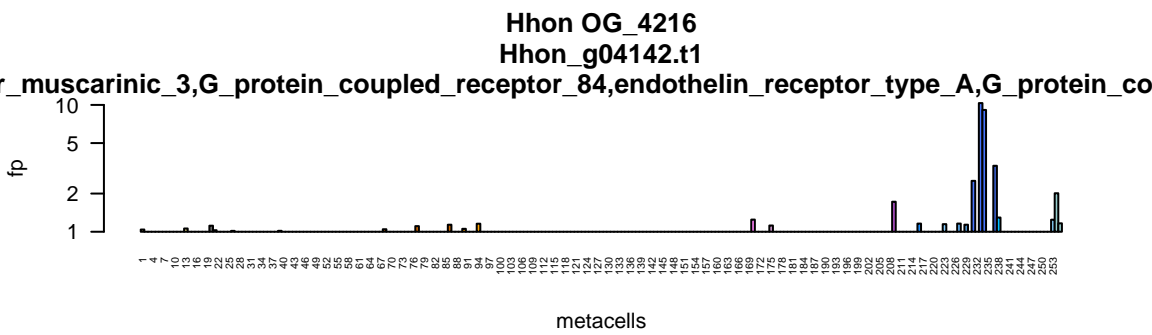
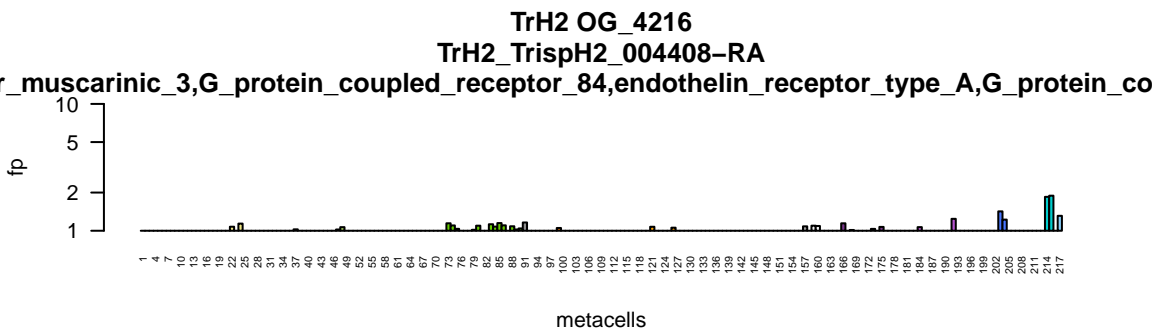
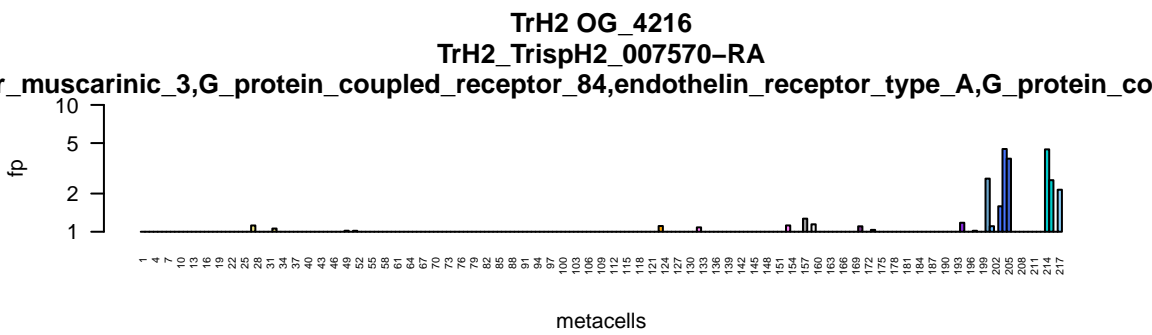
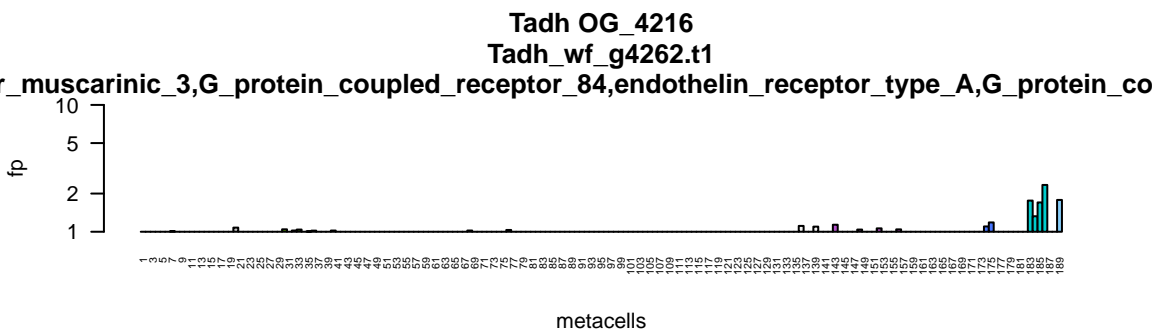
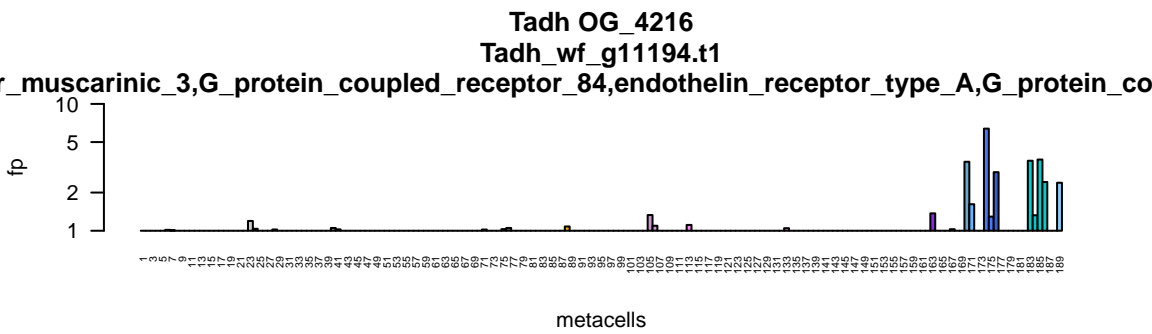
Hhon_g11321.t1

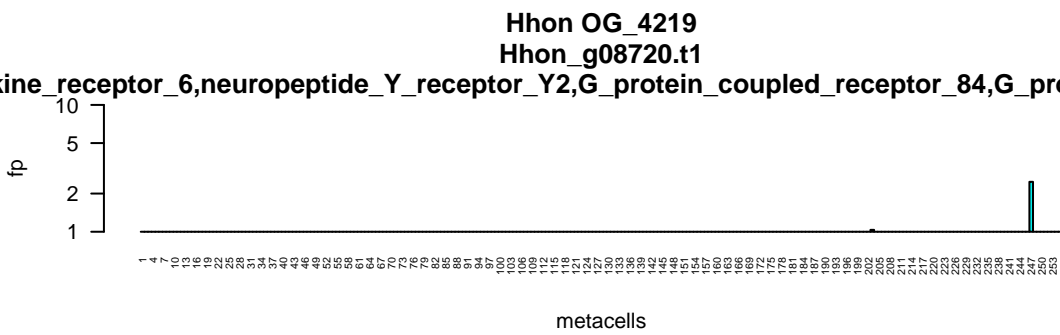
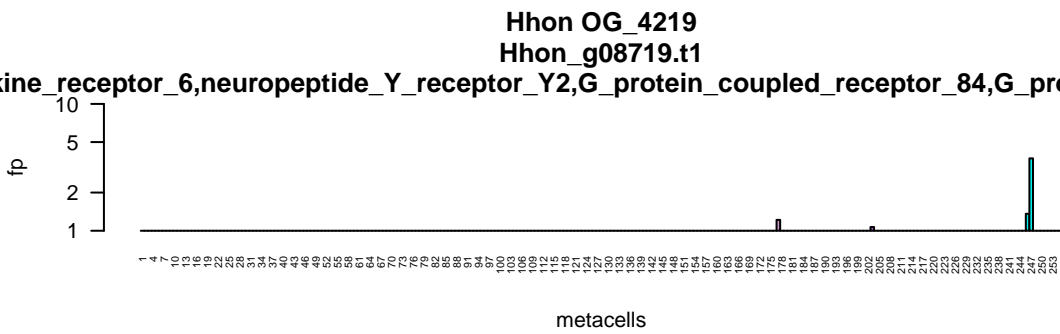
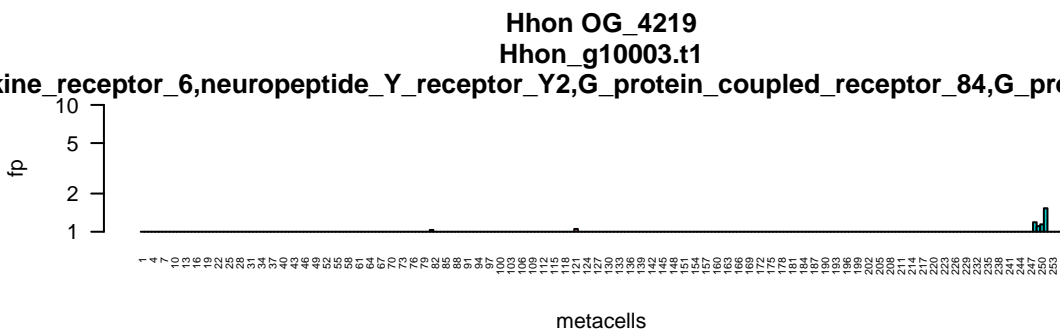
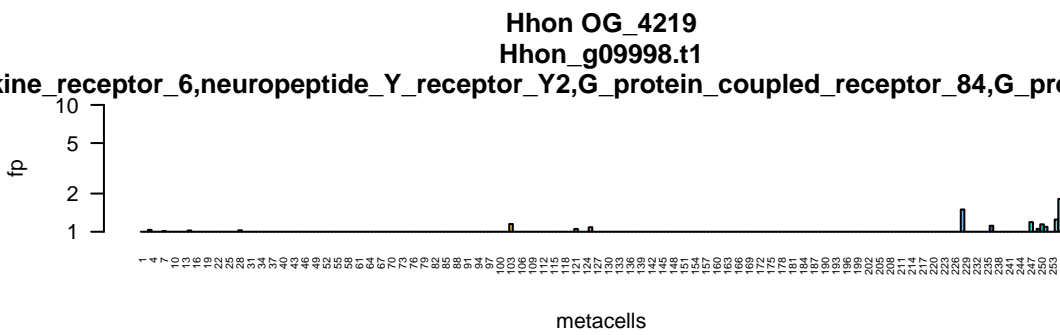
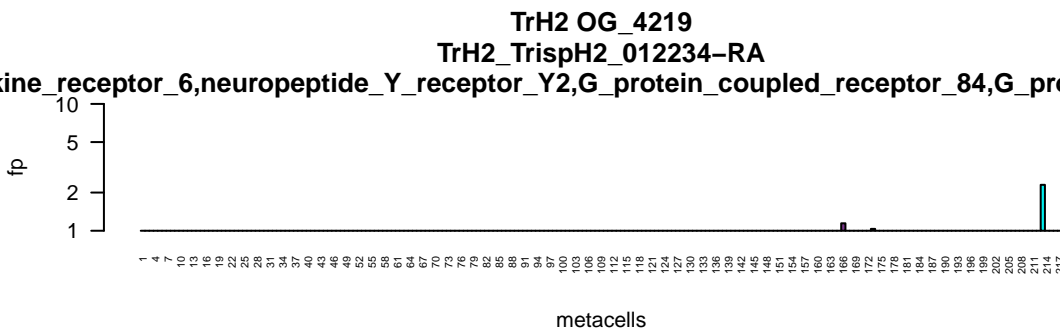
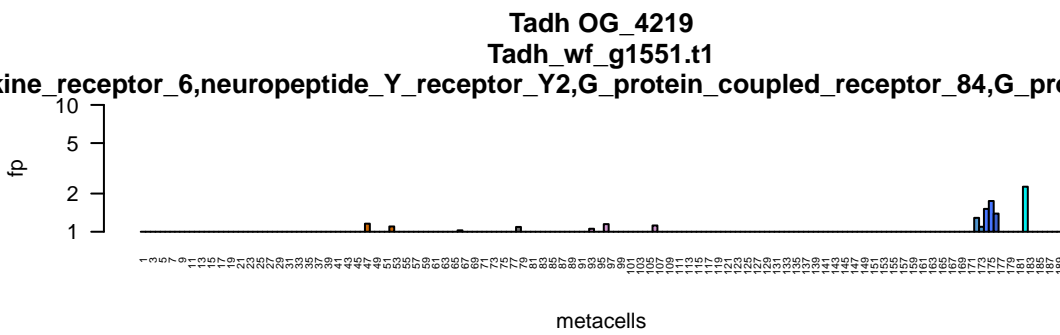
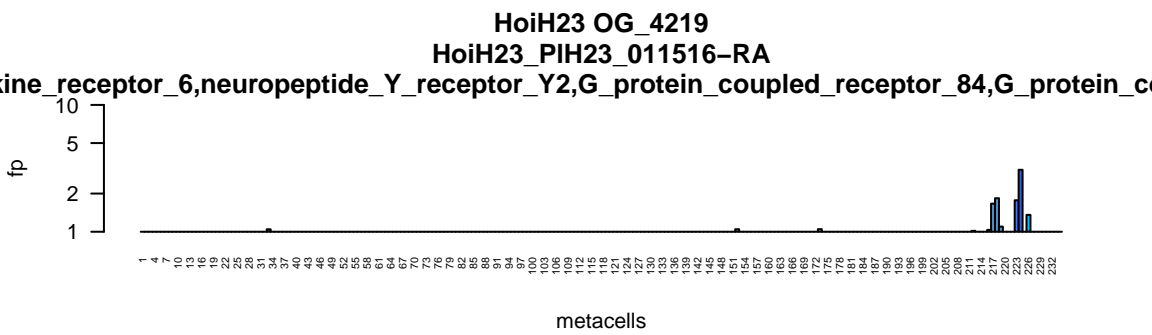
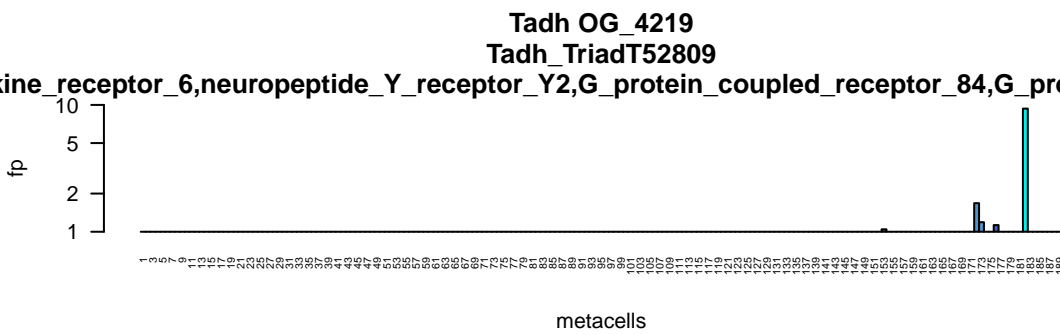
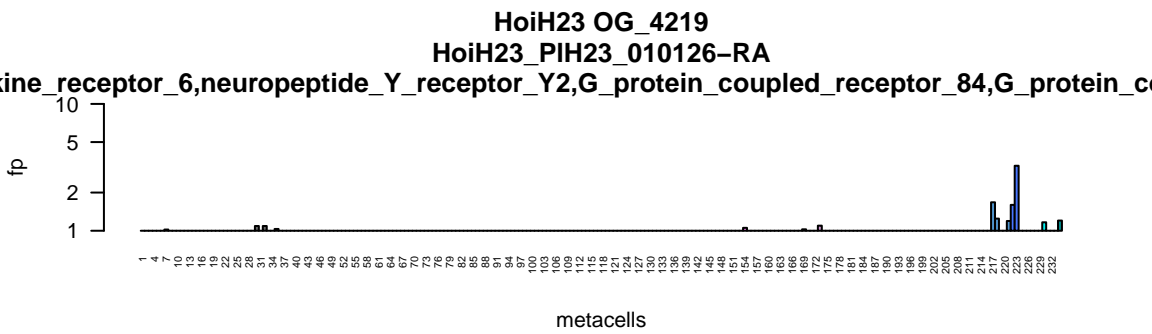
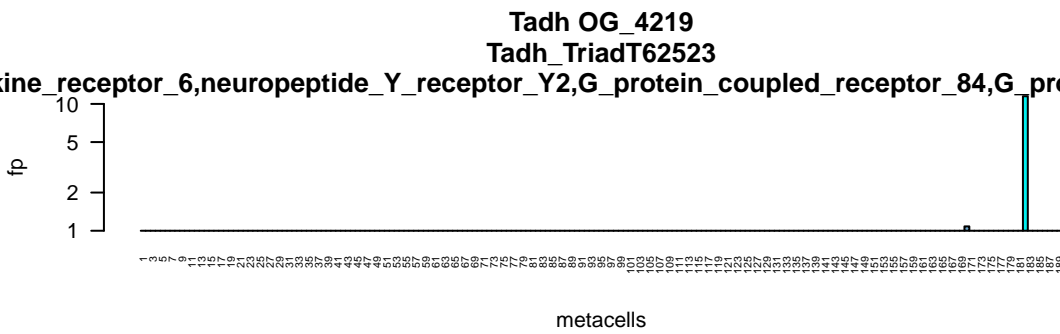


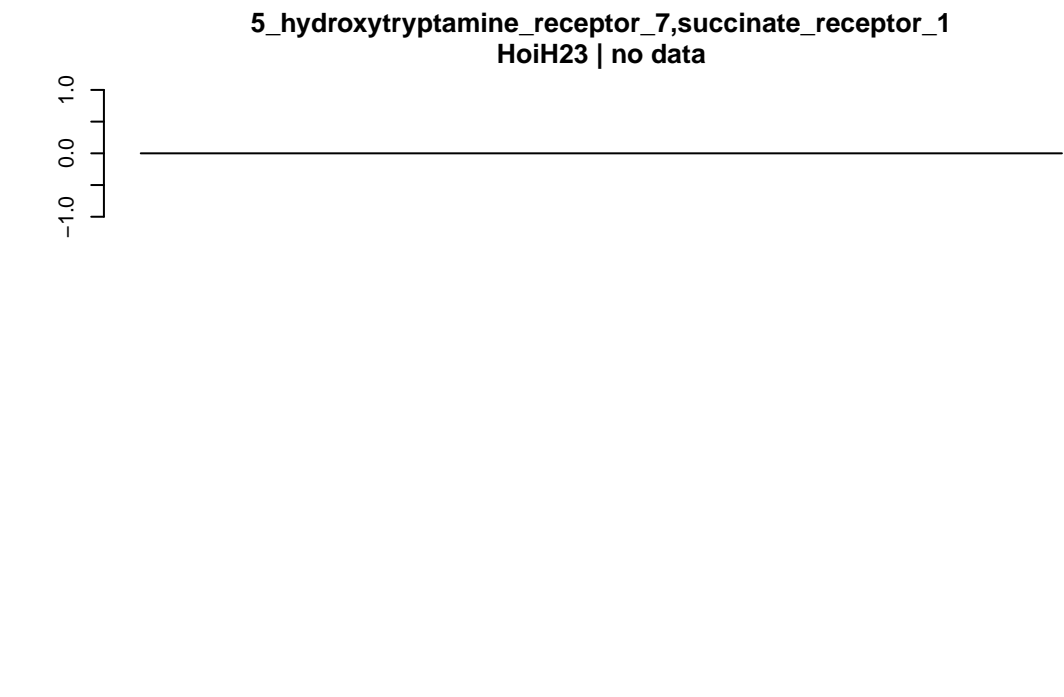
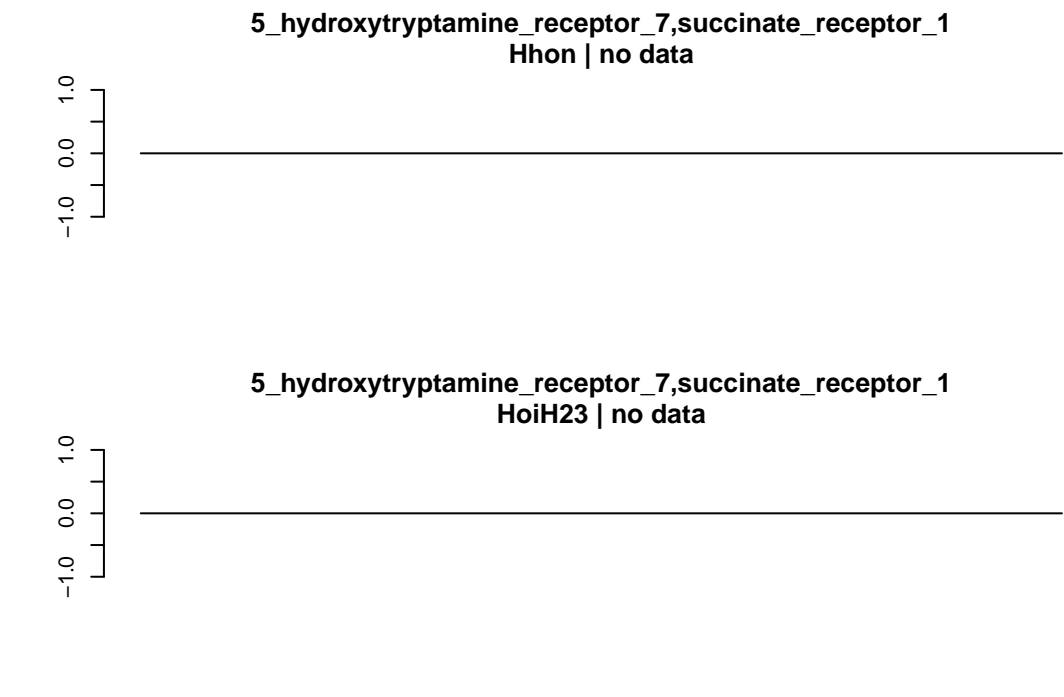
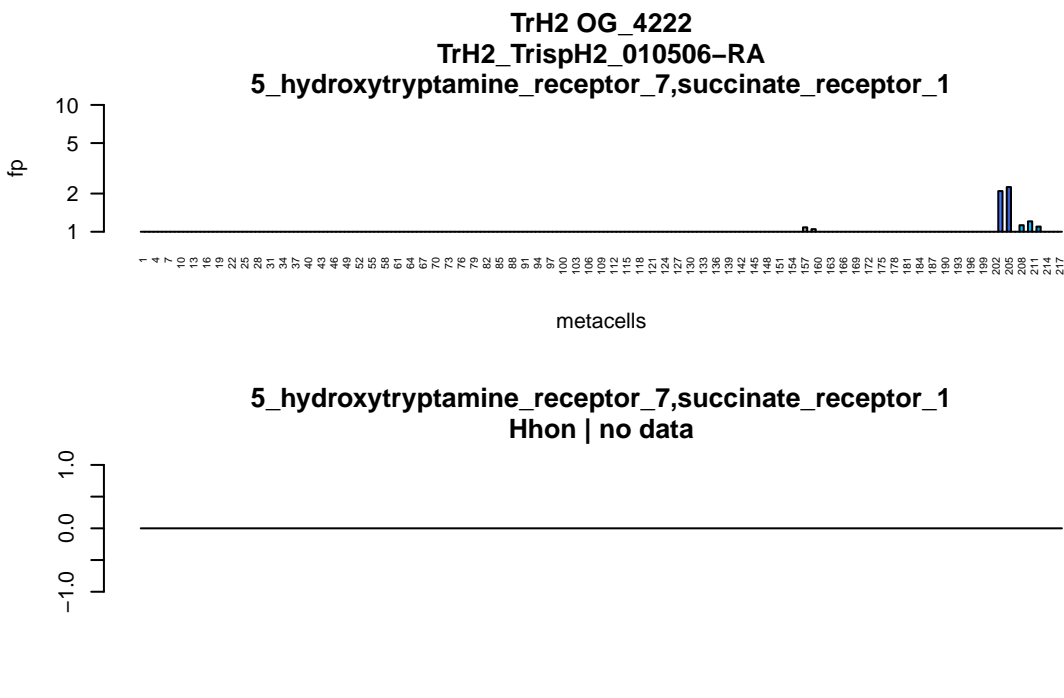
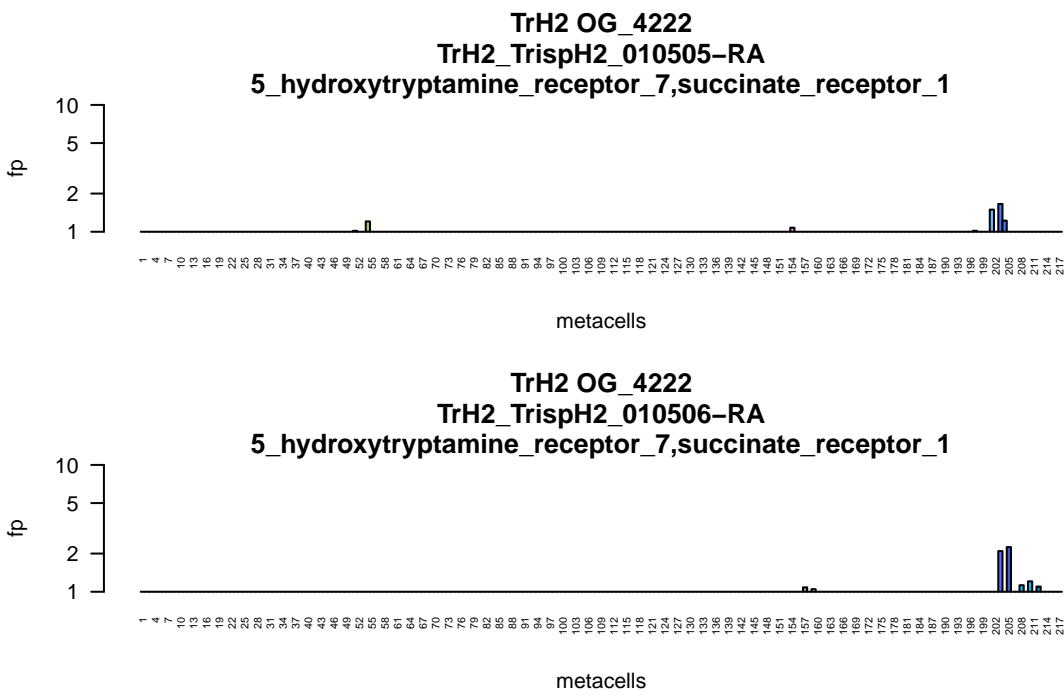
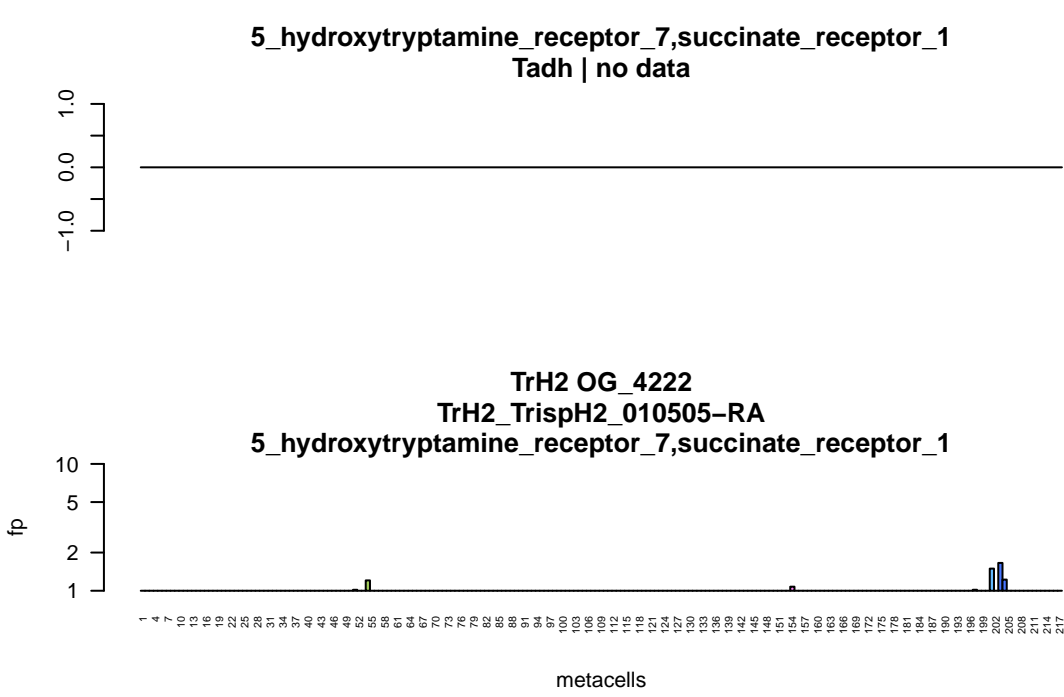
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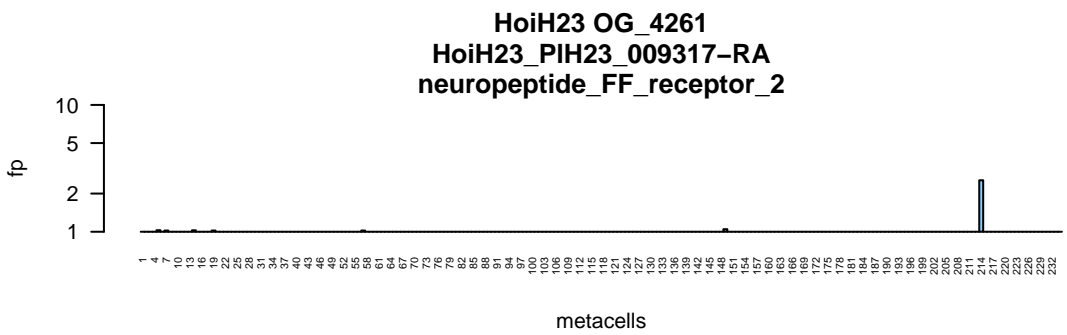
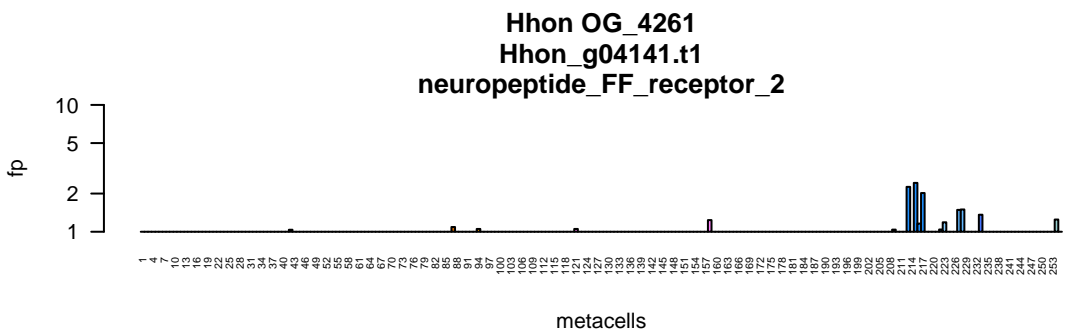
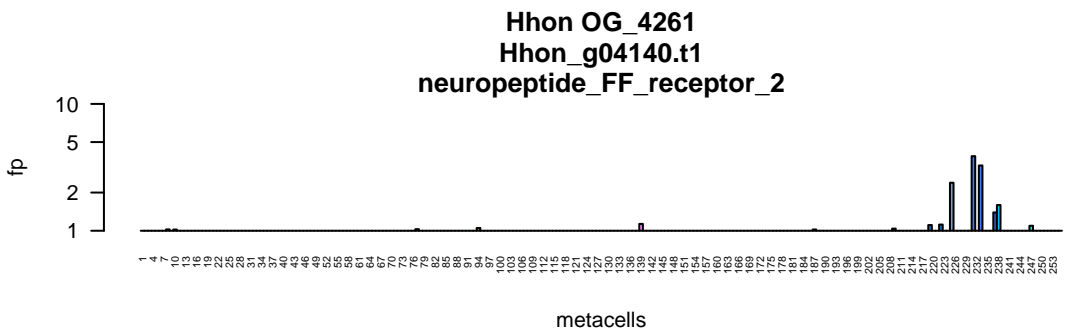
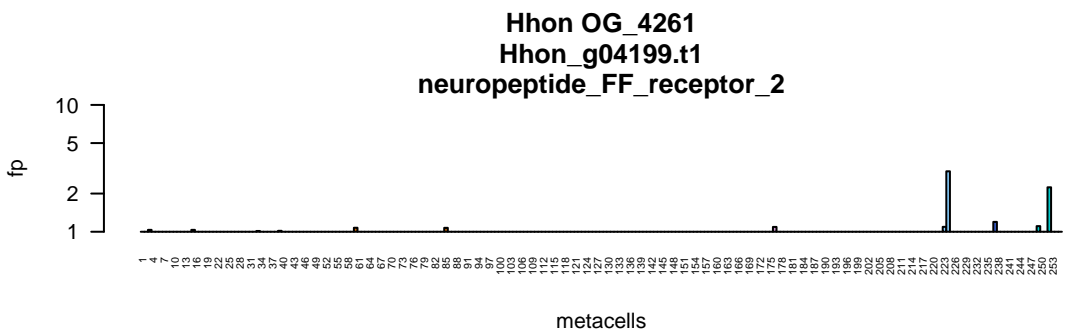
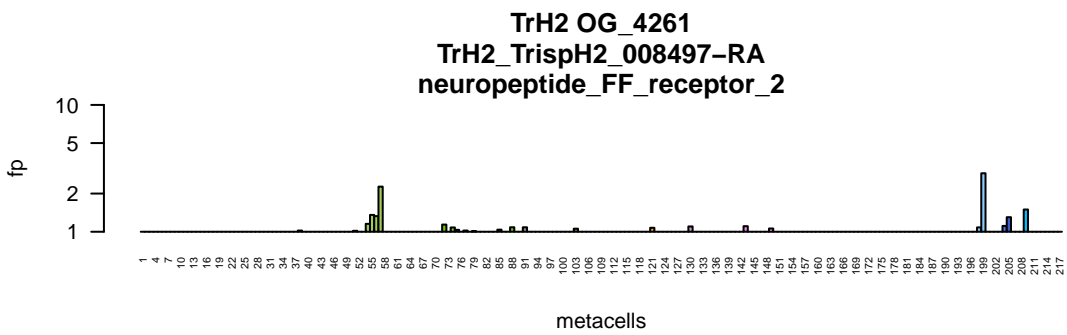
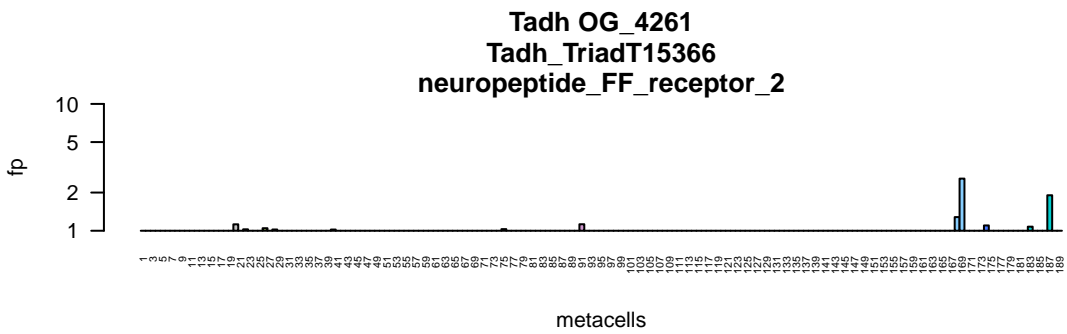


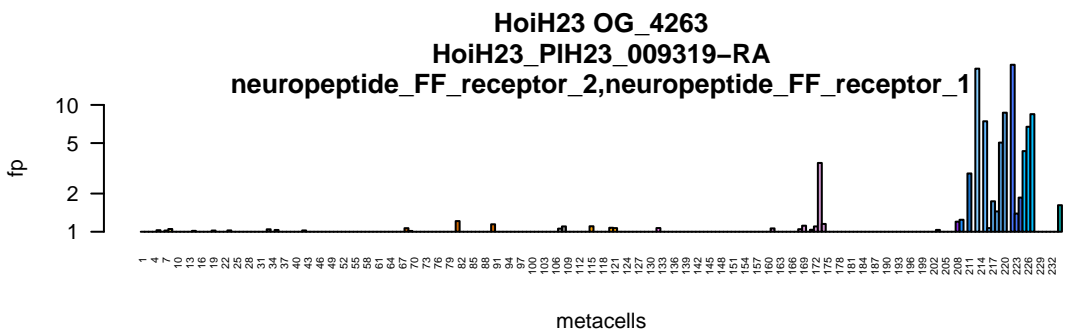
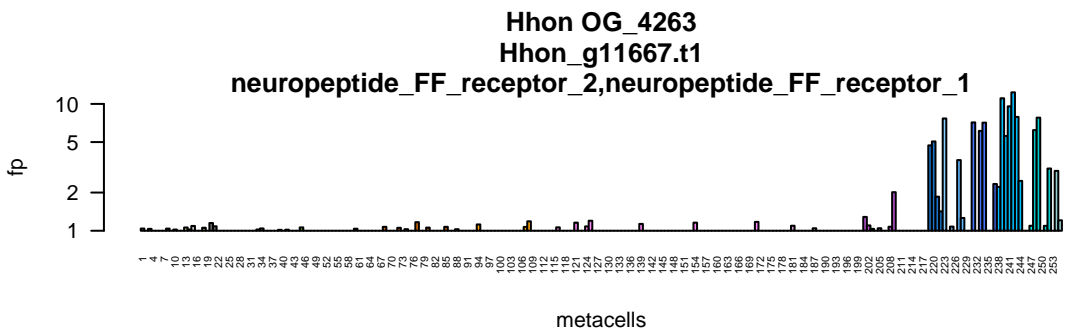
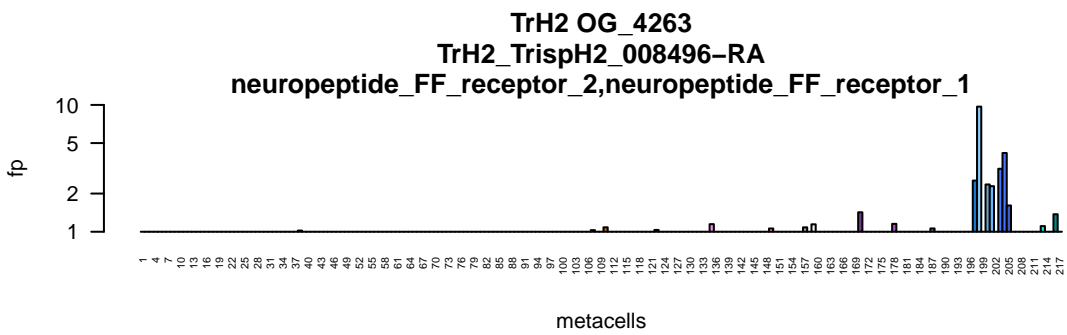
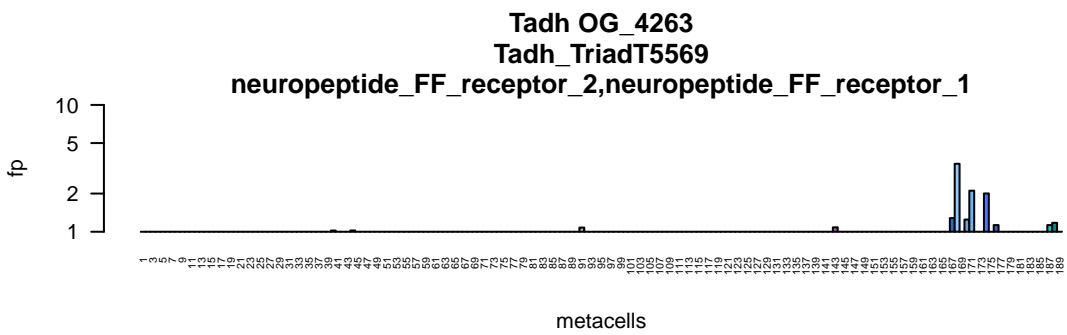
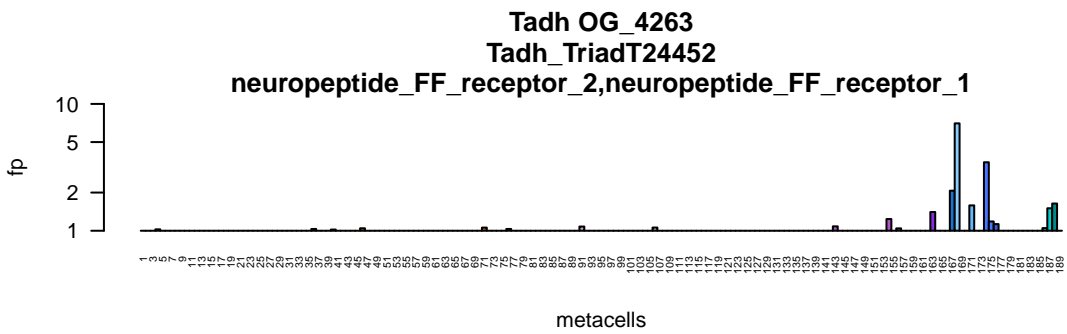












A bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a few outliers reaching up to 4.

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

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacells	fp
1	1
4	1
17	1
13	1
16	1
19	1
25	1
28	1
31	1
34	1
40	1
43	1
46	1
52	1.5
55	1.2
58	1
61	1
67	1
70	1
73	1
77	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1.2
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1.2
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1.2
163	1
166	1
169	1
172	1
175	1.5
179	1
181	1
184	1
187	1
190	1
193	1
199	1.8
202	1.8
205	1.5
208	1
211	1
214	1.2
217	1.8

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

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacell	fp
1	1
4	1
13	1
16	2
17	1
19	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
52	1
55	1
58	1
61	1
67	1
70	1
73	1
77	1
79	1
82	1
85	1
88	1
94	1
97	1
100	2
106	1
109	1
112	1
118	1
121	1
124	1
127	1
133	2
136	1
139	1
142	1
145	1
148	1
151	1
154	2
157	1
160	2
163	1
166	1
168	1
172	1
175	1
179	1
181	1
184	1
187	1
190	1
193	1
198	1.5
199	3
202	3
205	3
206	1.5
207	1.5
208	1.5
209	1.5
210	1.5
211	1.5
212	1.5
213	1.5
214	1.5
215	1.5
216	1.5
217	1.5

metacells

A bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists indices from 1 to 217. Most metacells have 1 false positive, but metacells 202, 205, and 214 have 2 false positives.

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

cd_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

fp

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 253. The y-axis shows fp values from 1 to 10. Most metacells have a frequency of 1, with a few having higher frequencies (e.g., metacell 220 has a frequency of 2).

Hhon OG_4305
Hhon_g09074.t1

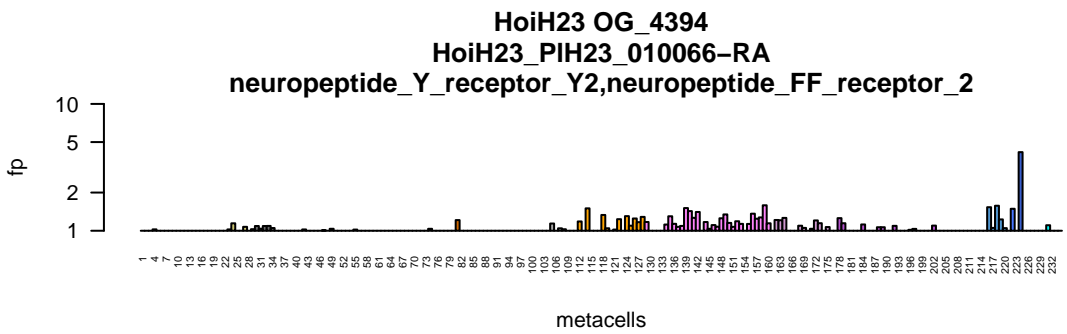
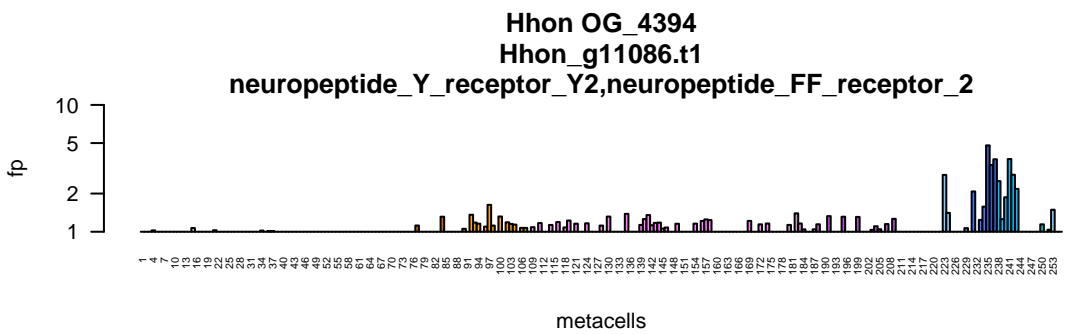
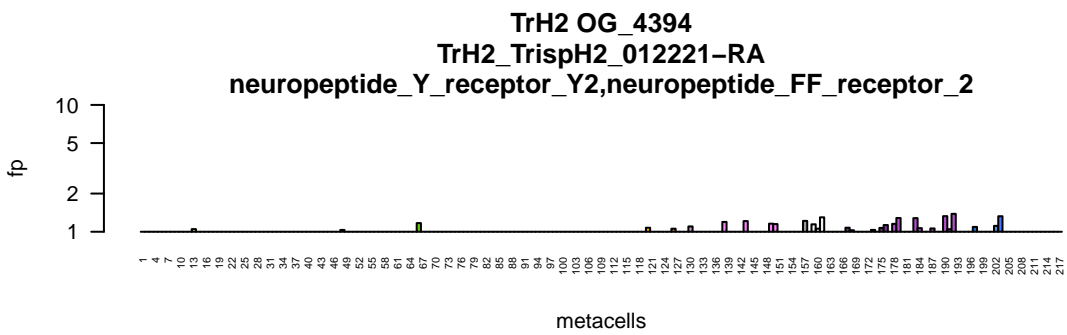
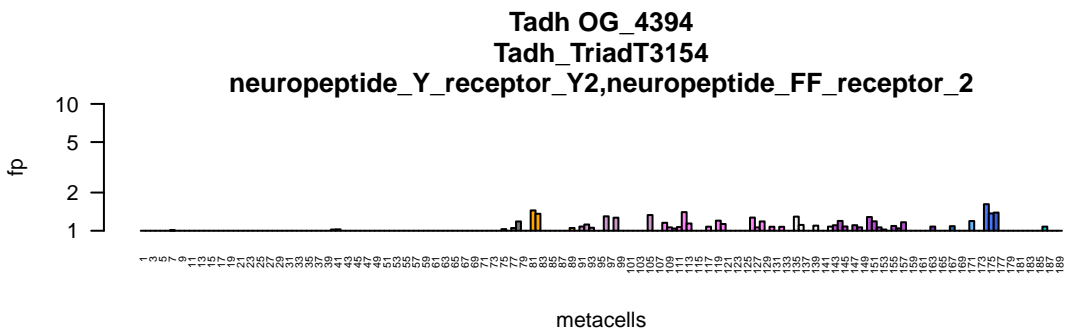
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

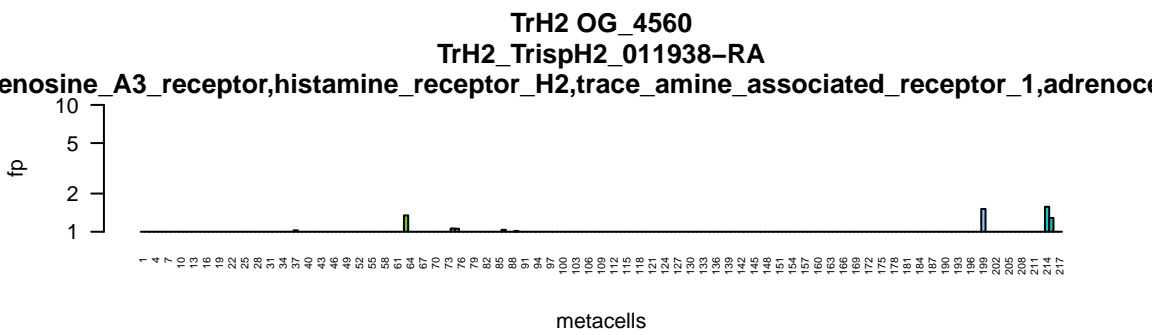
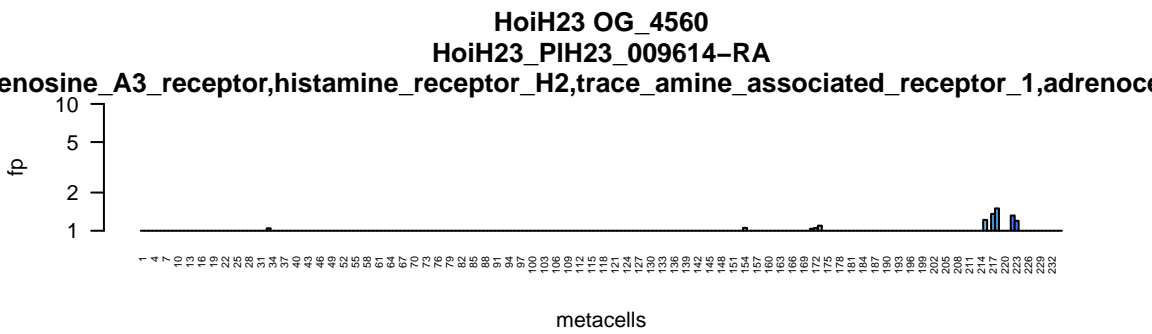
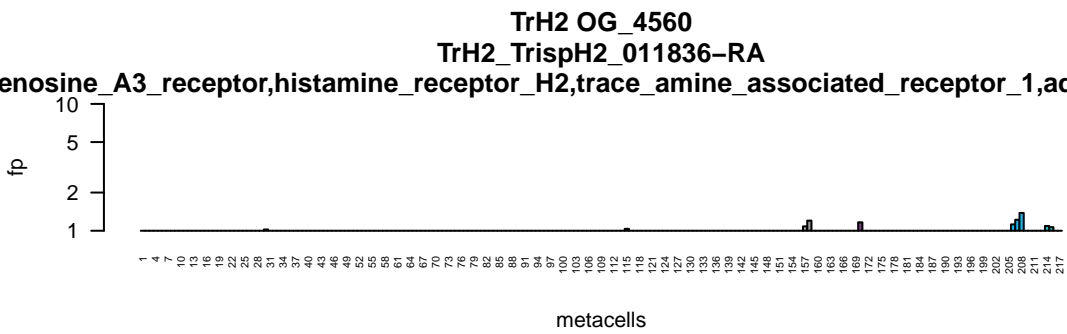
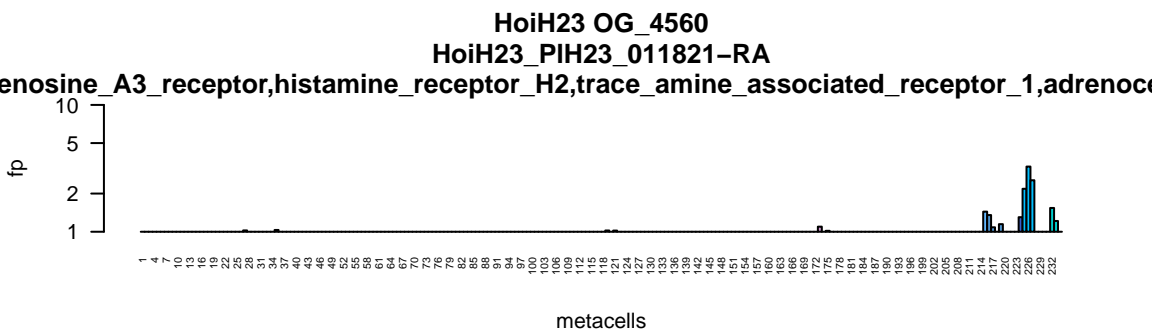
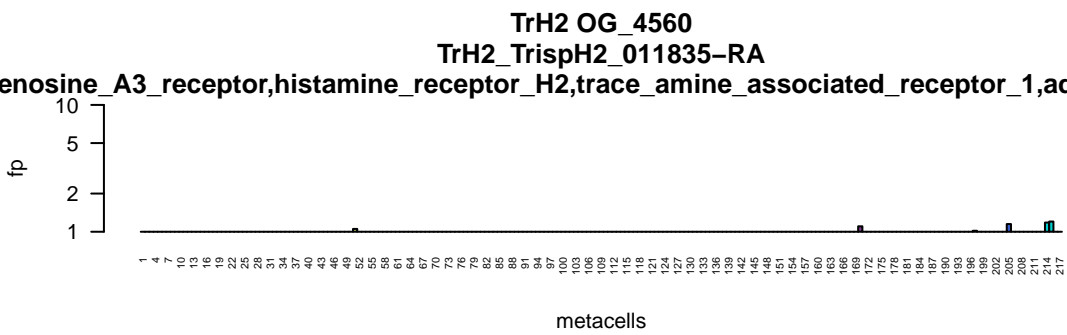
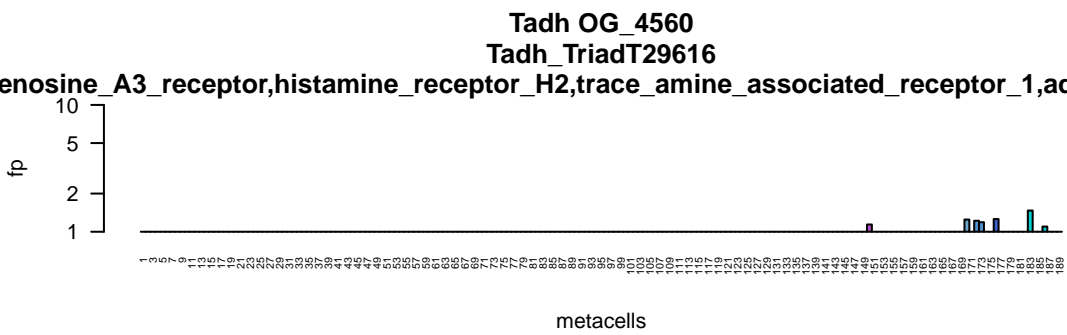
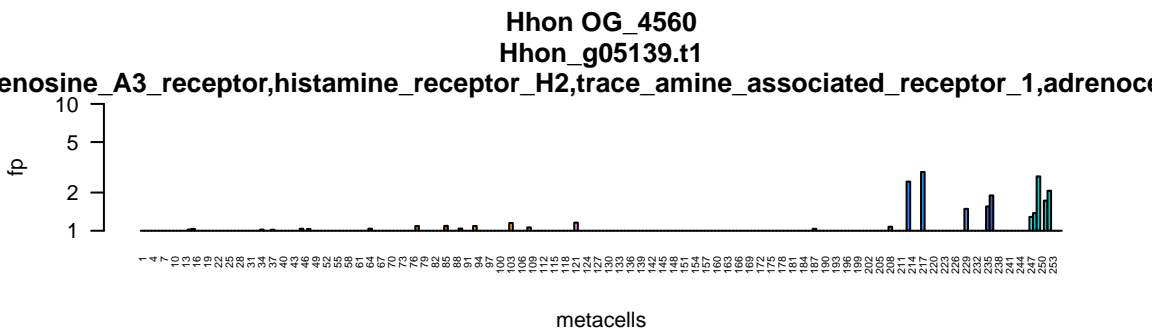
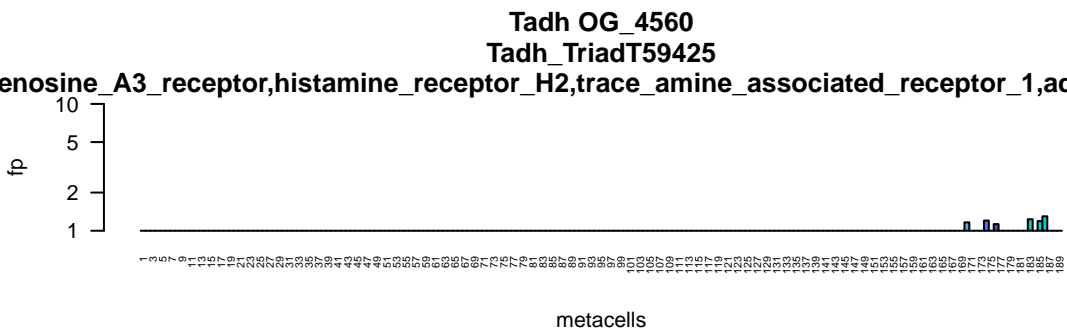
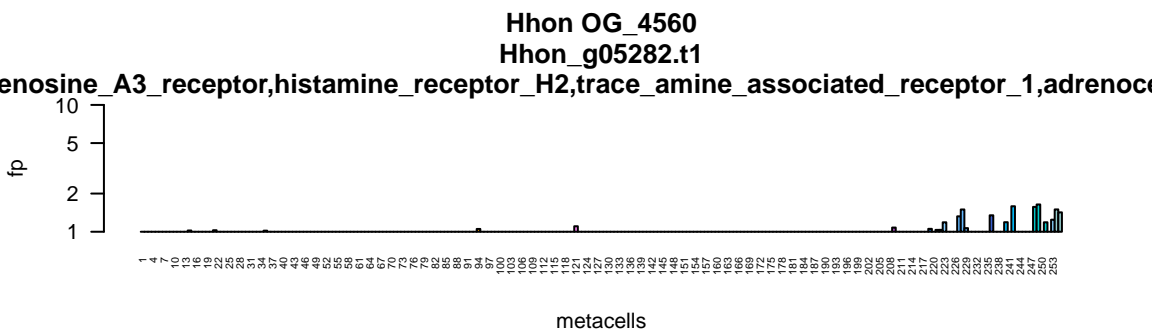
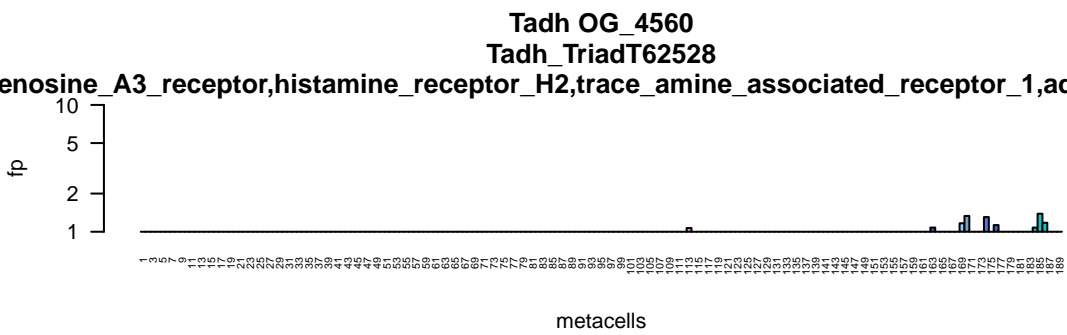
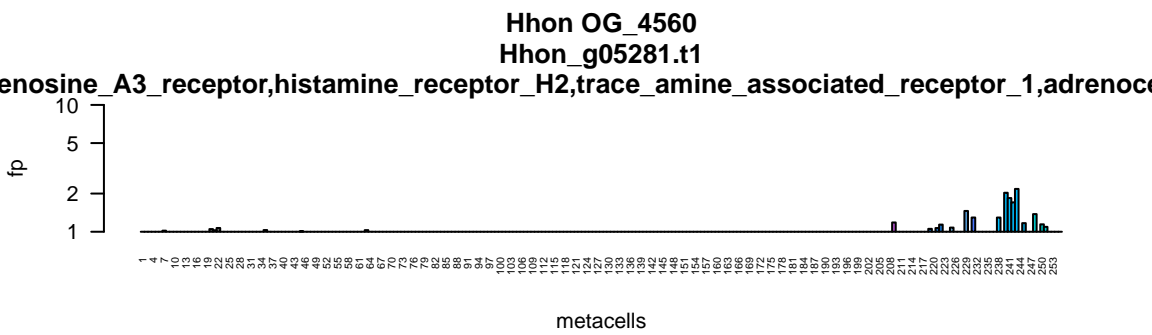
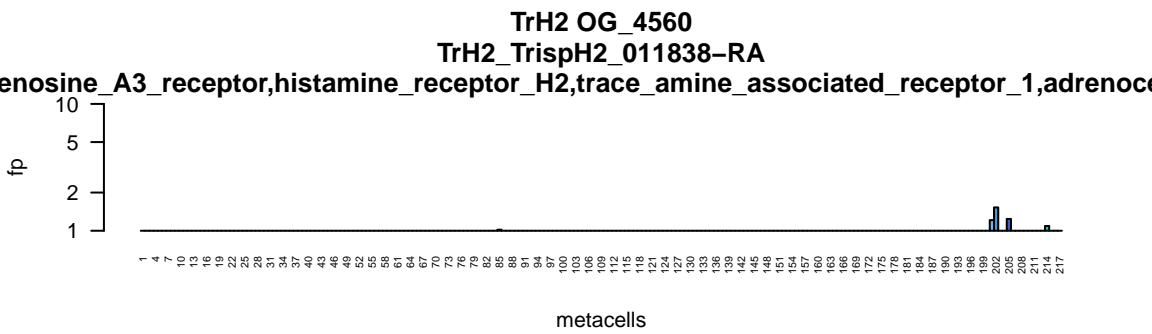
fp

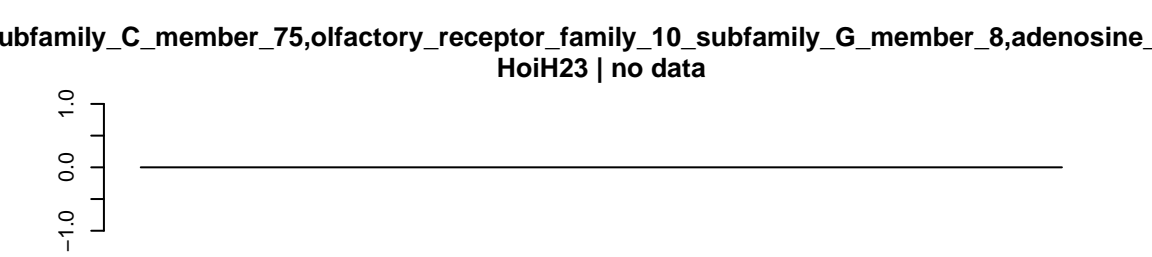
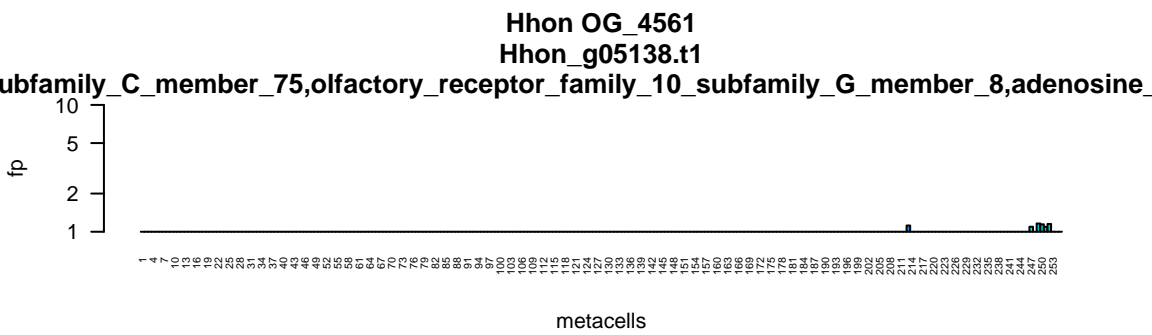
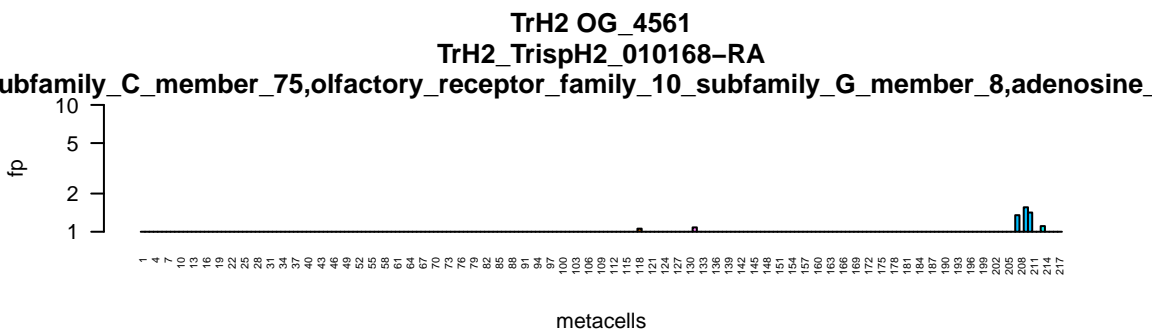
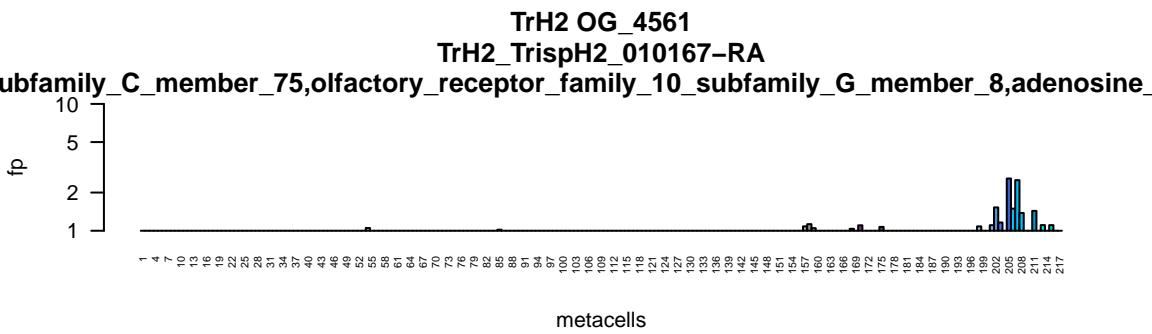
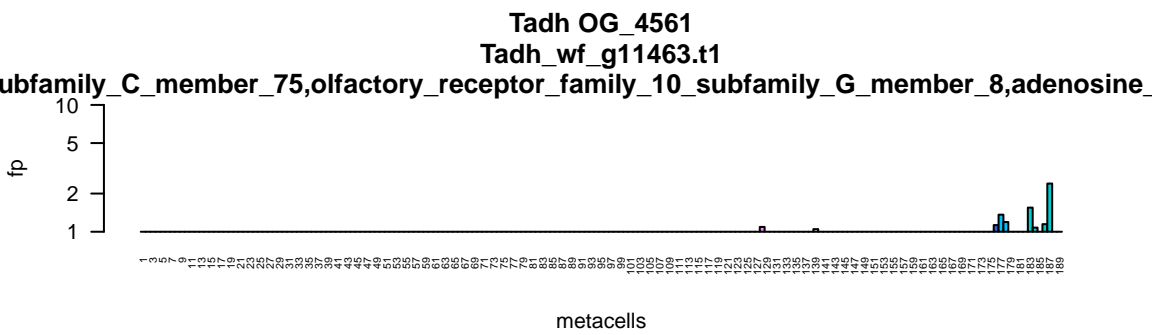
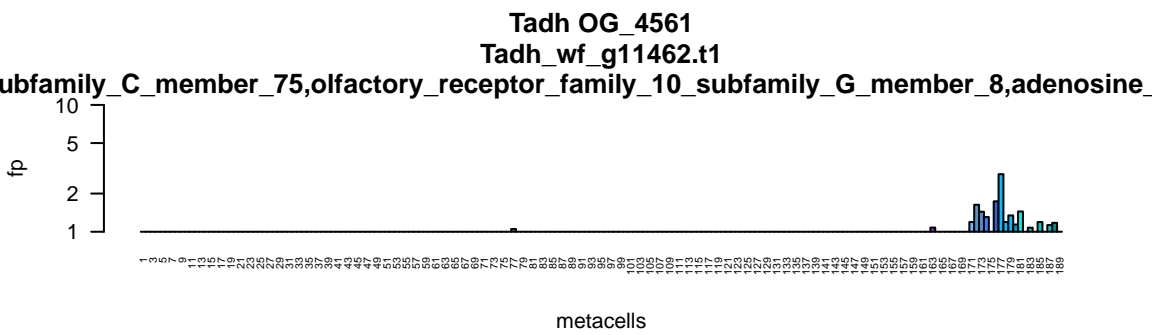
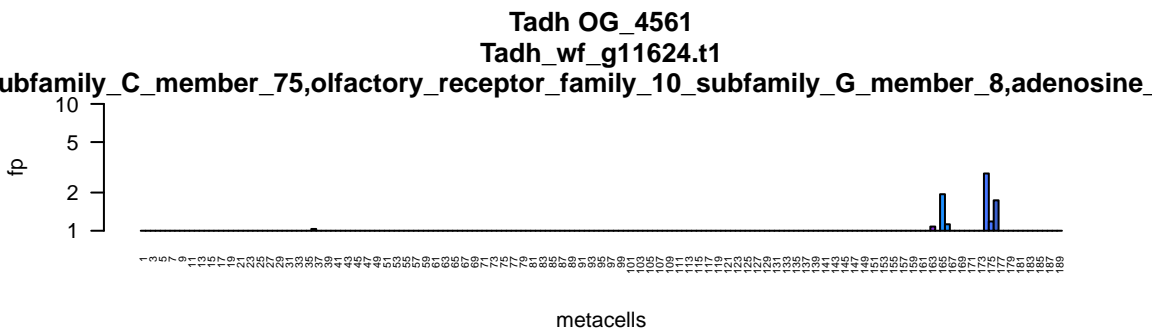
metacells

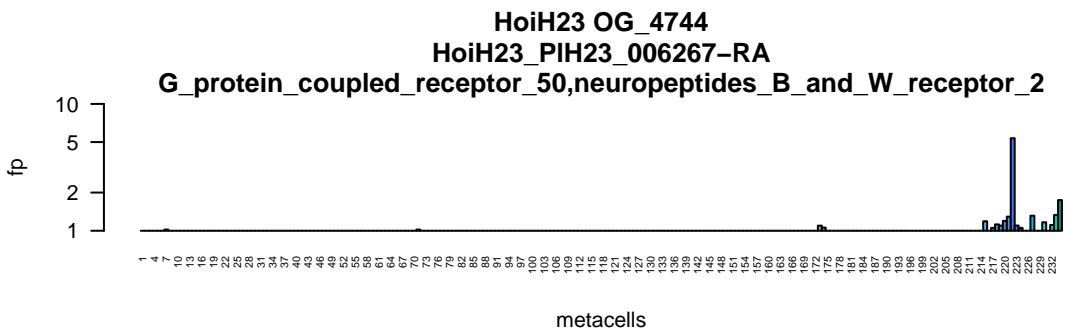
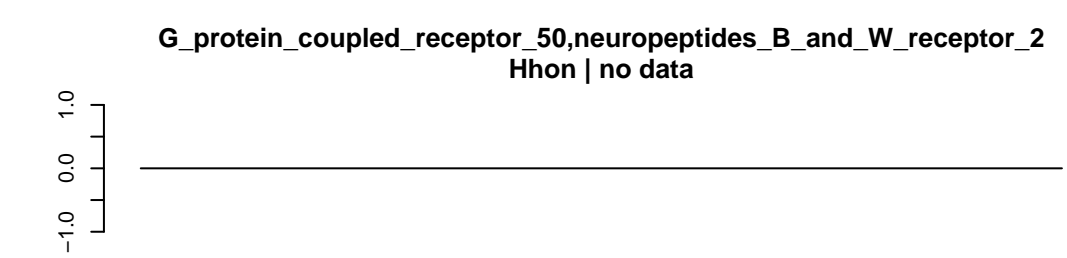
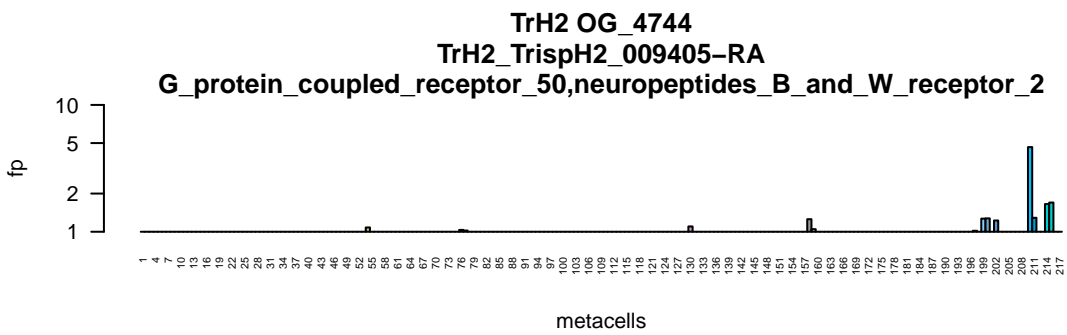
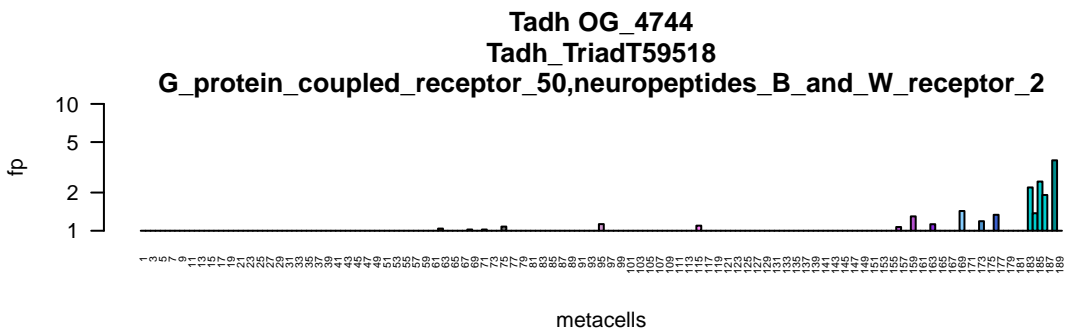
Bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and lists indices from 1 to 232. The y-axis is labeled 'fp' and has a logarithmic scale with values 1, 2, 5, and 10. Most metacells have a frequency of 1. Metacells 214, 223, and 232 show higher frequencies, with 214 reaching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

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









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
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	2
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
214	2
215	1
216	1
217	1

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

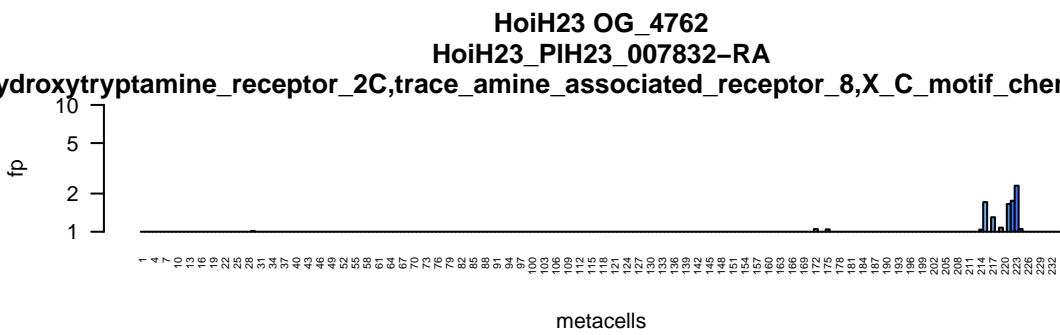
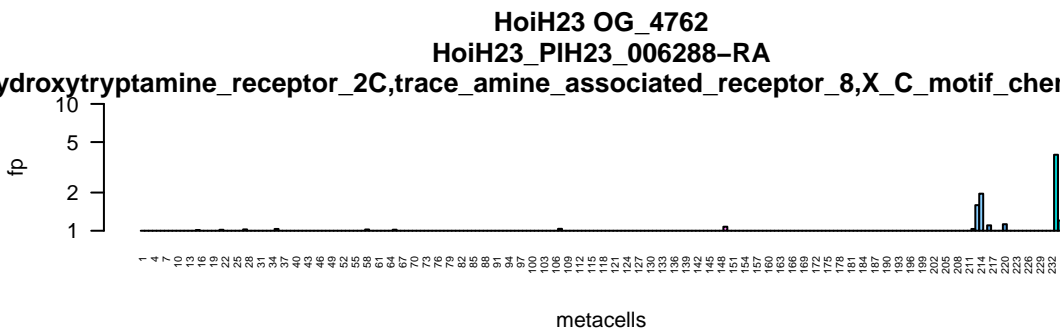
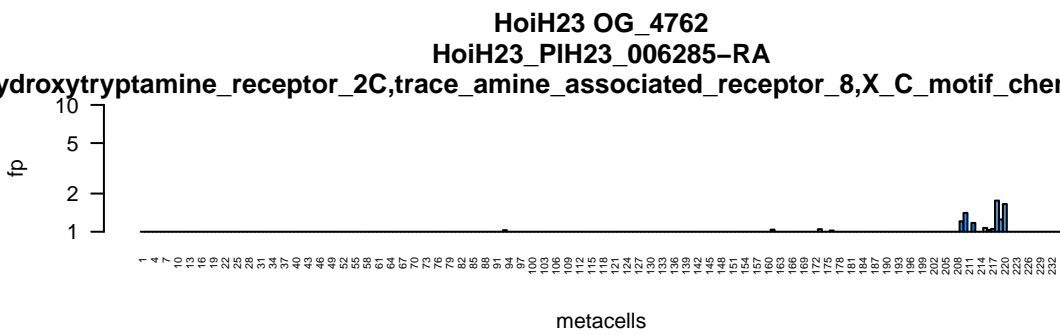
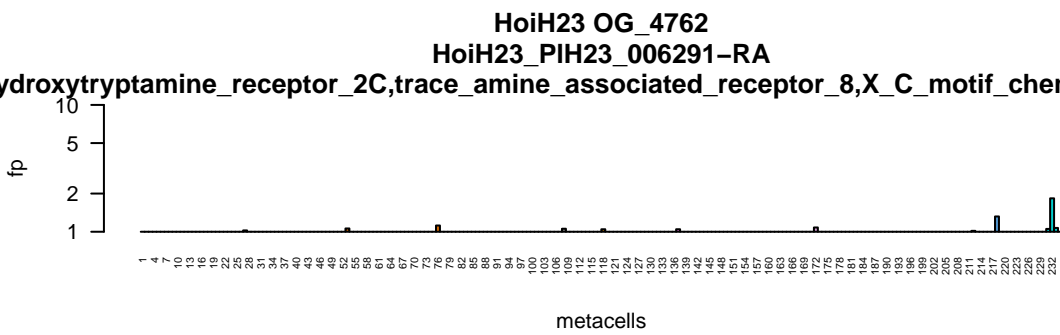
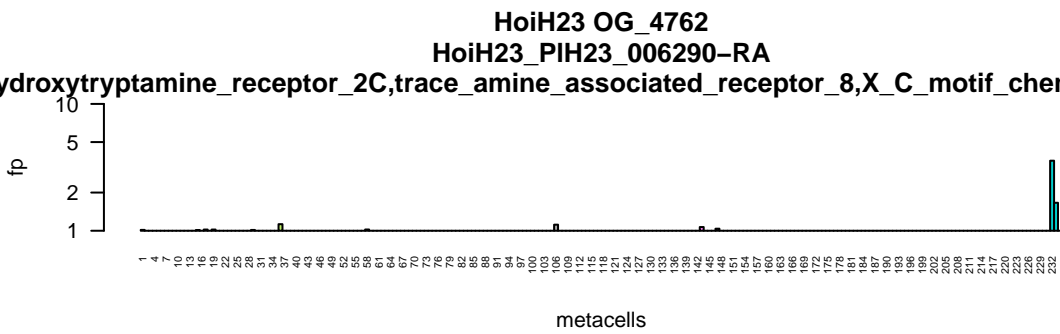
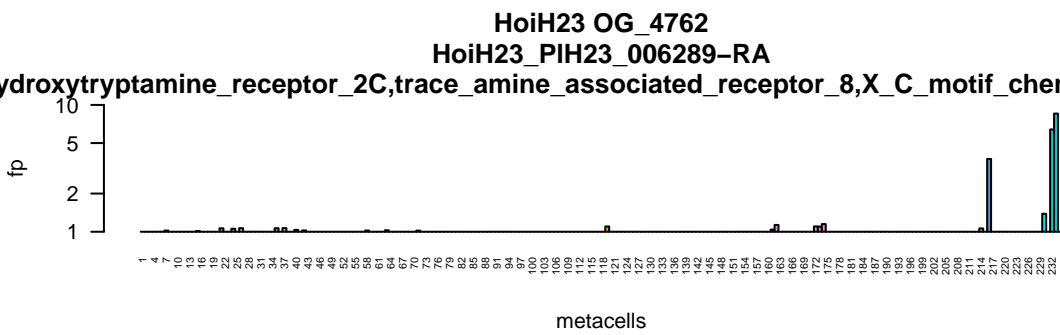
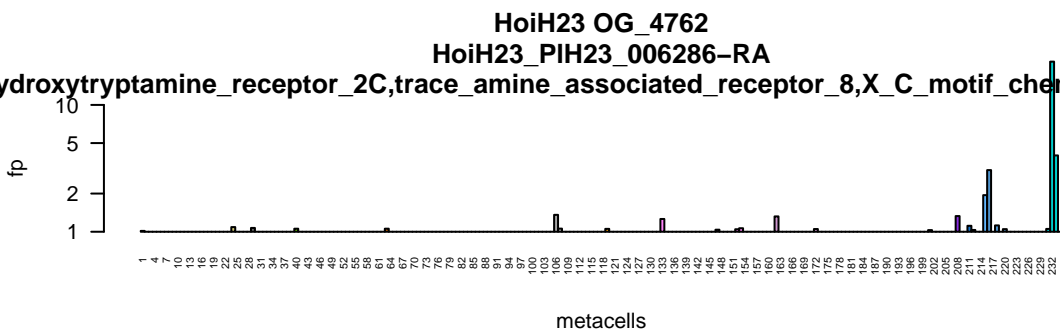
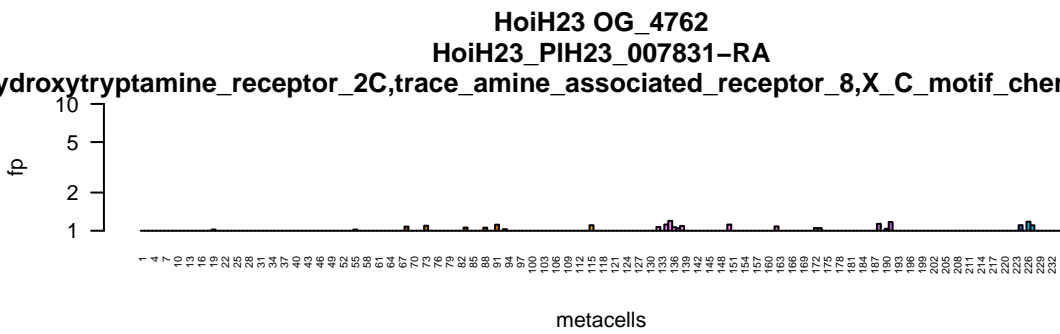
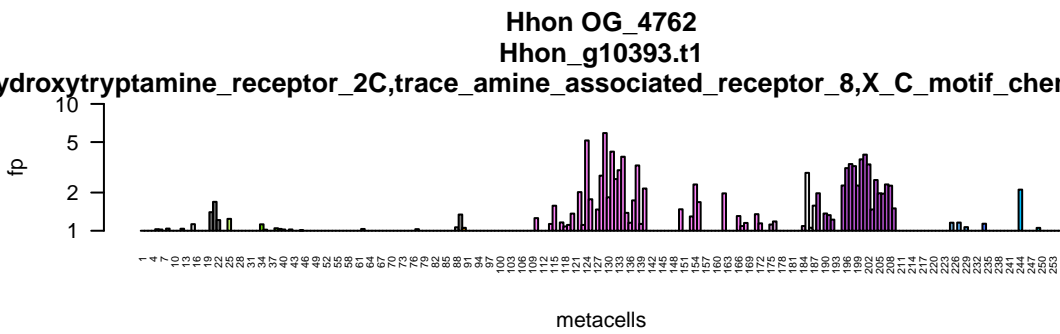
Hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

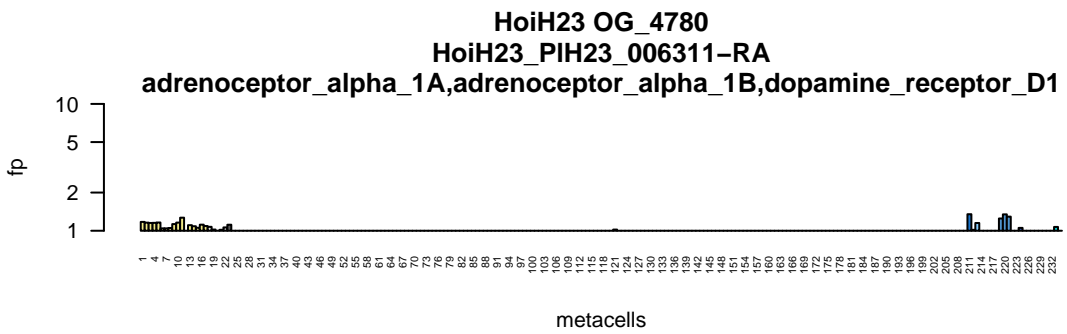
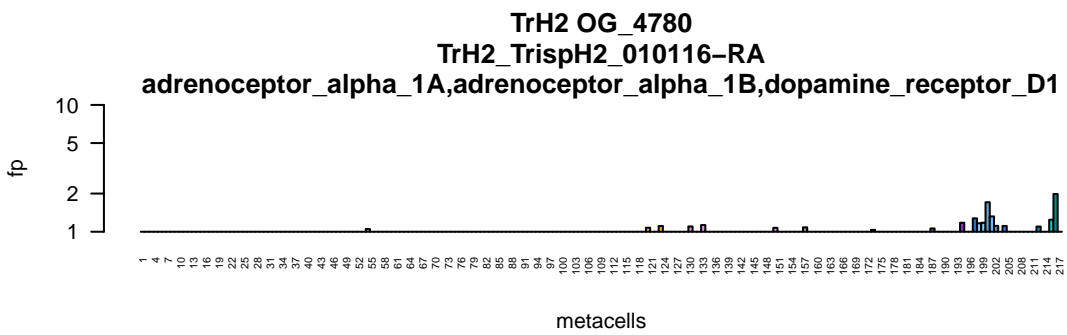
Hhon OG_4762

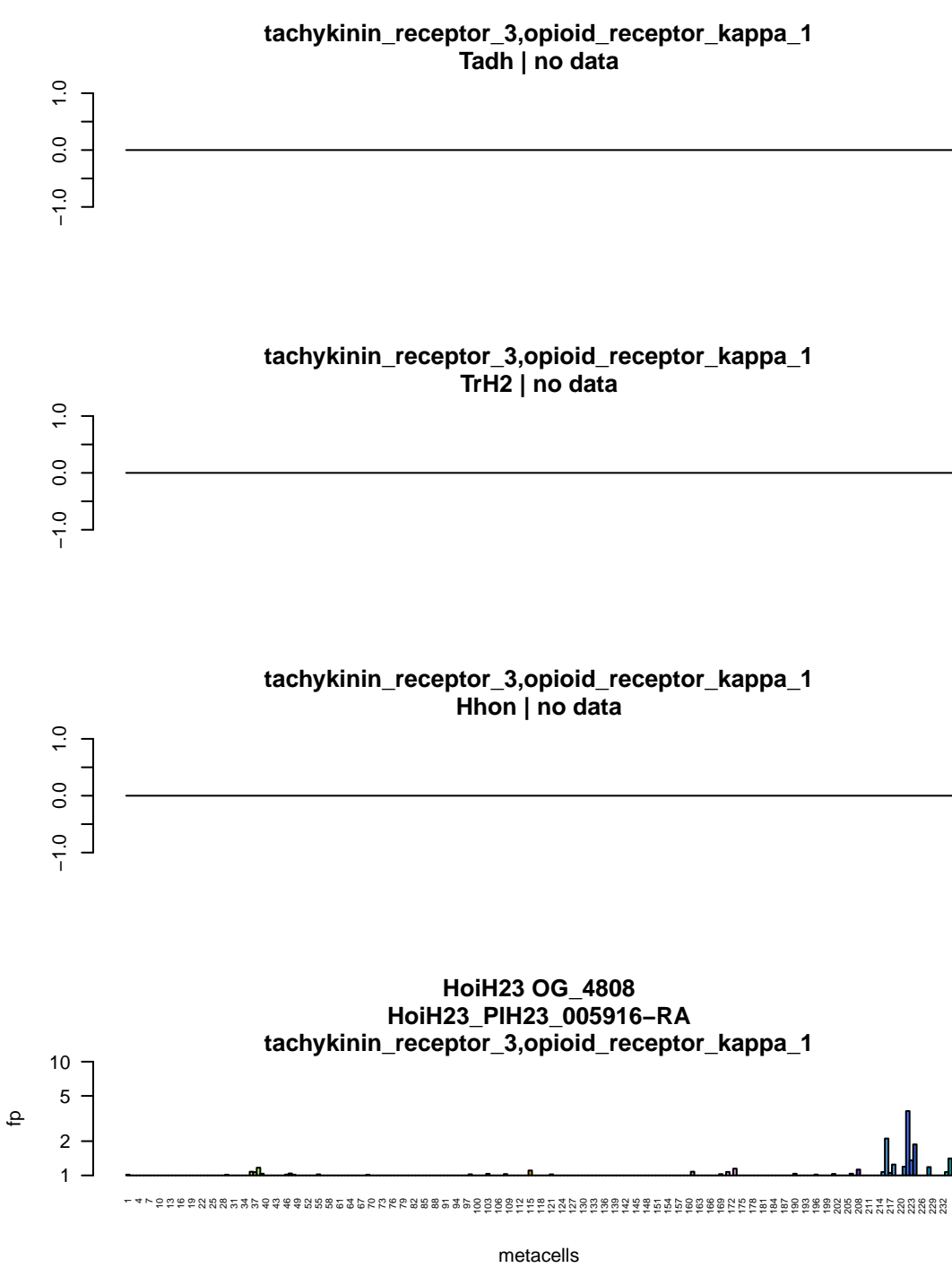
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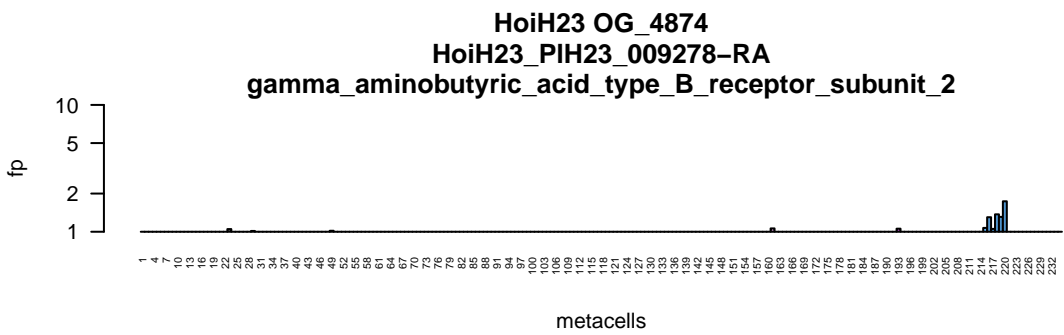
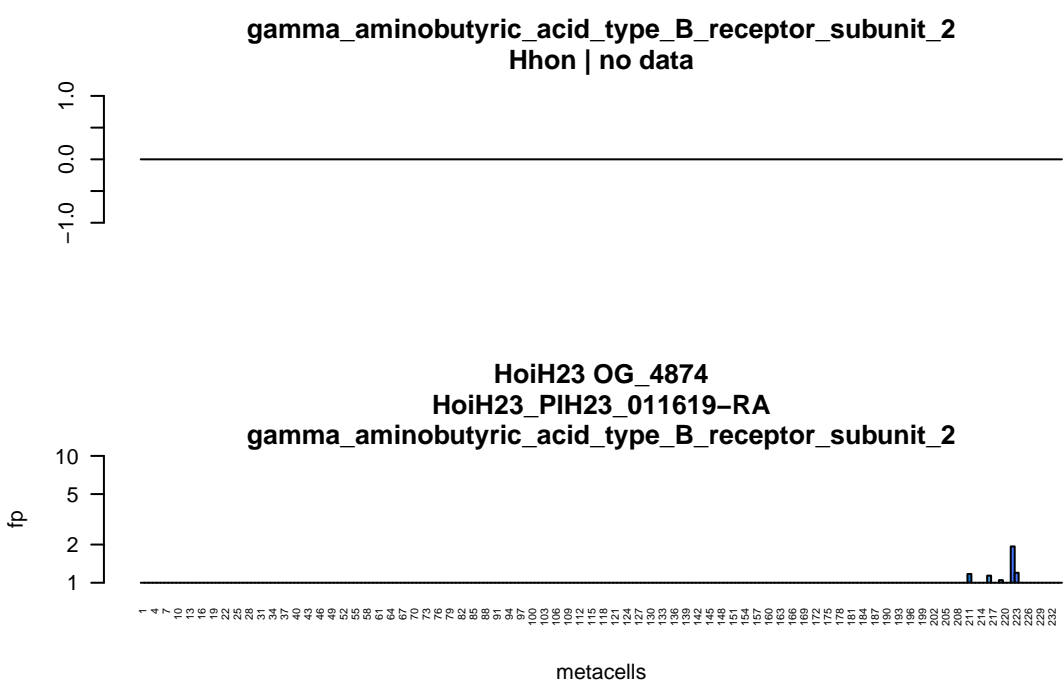
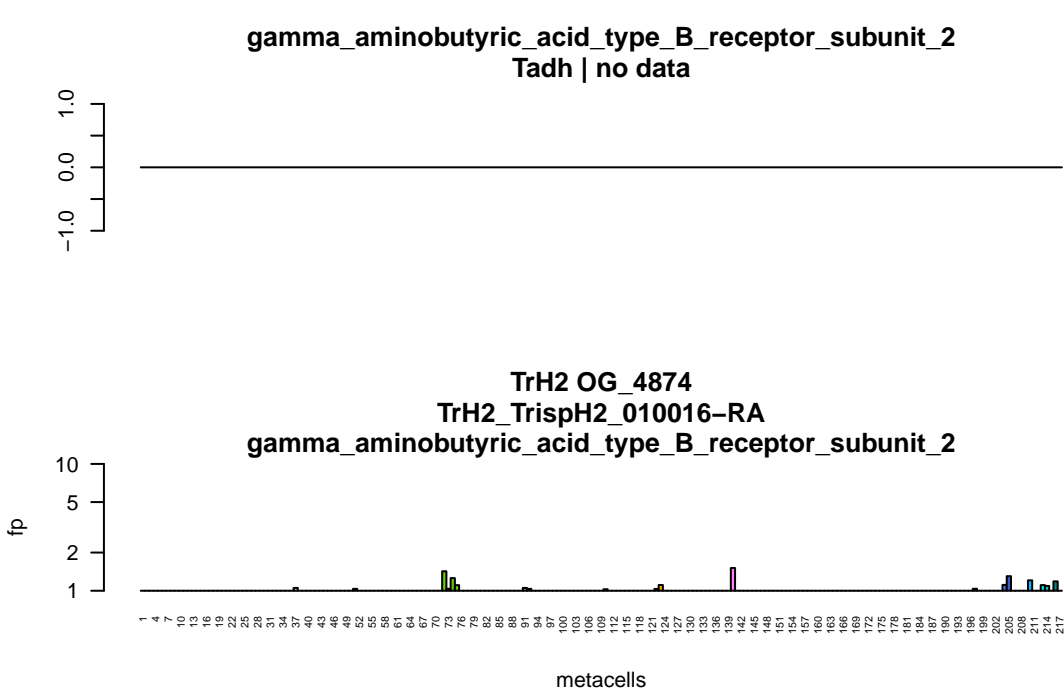
fp

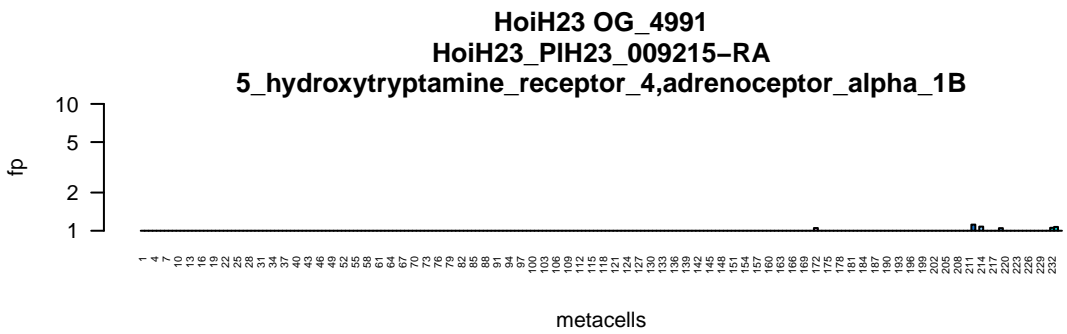
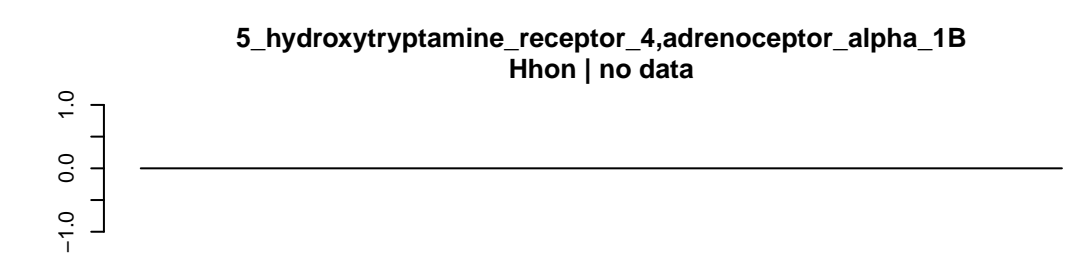
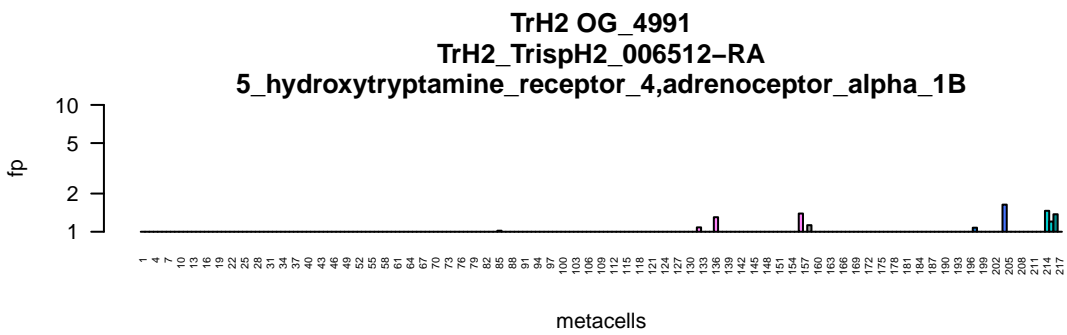
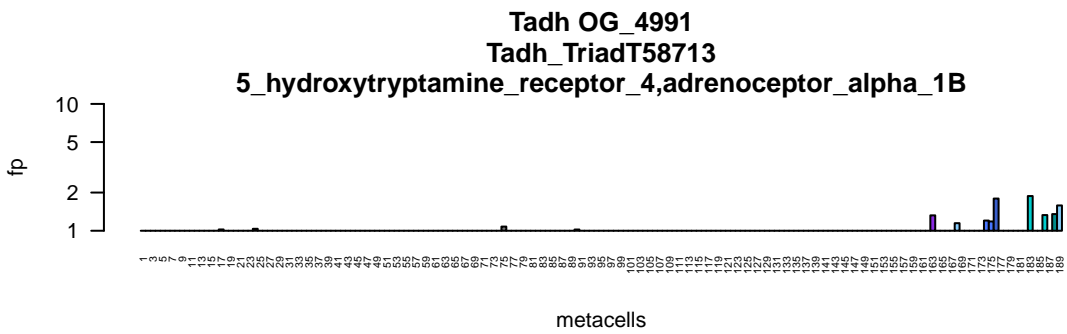
metacells



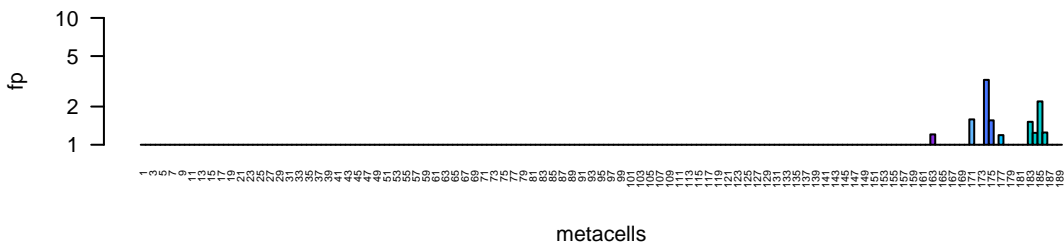




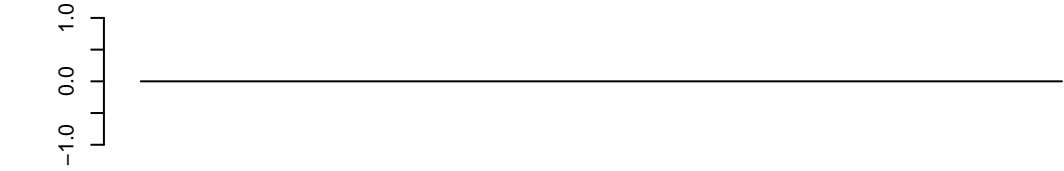




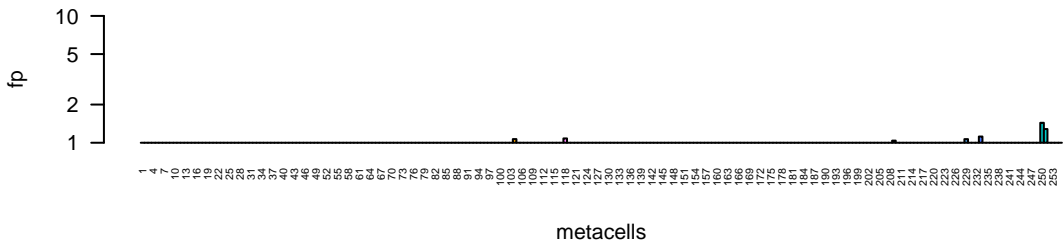
Tadh OG_5118
Tadh_wf_g10490.t1



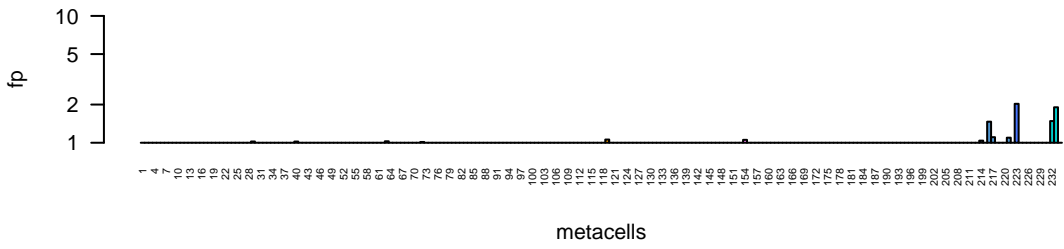
TrH2 | no data

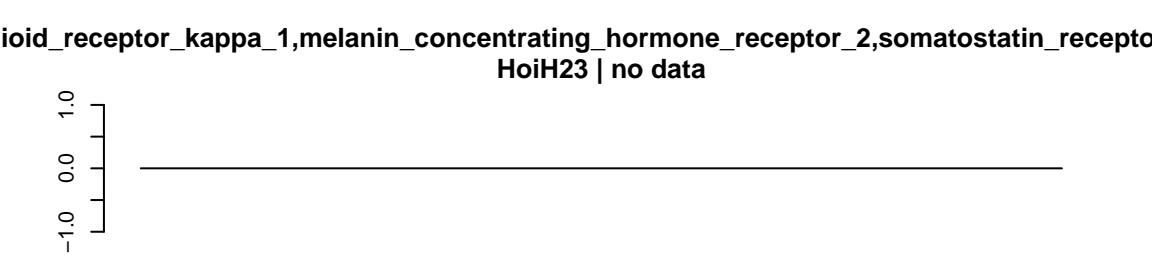
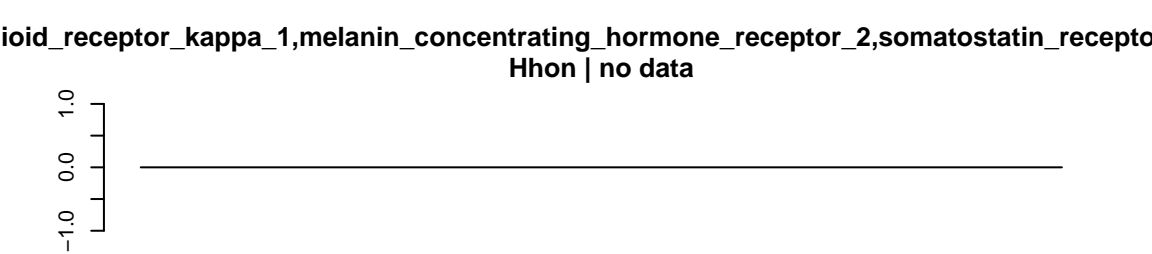
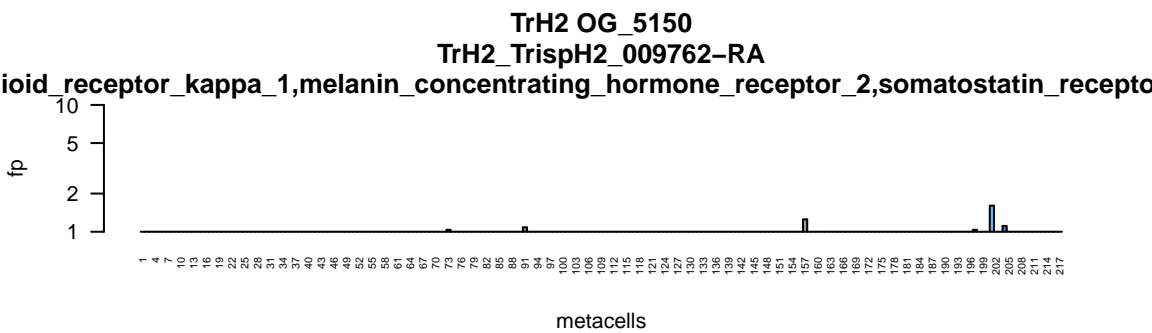
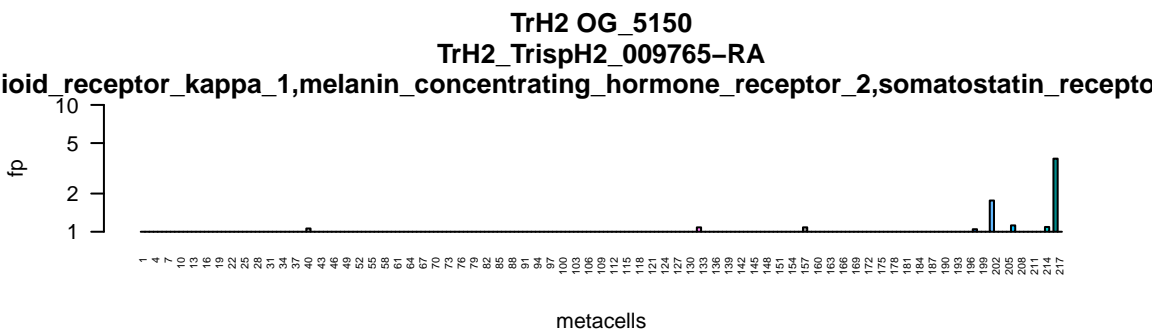
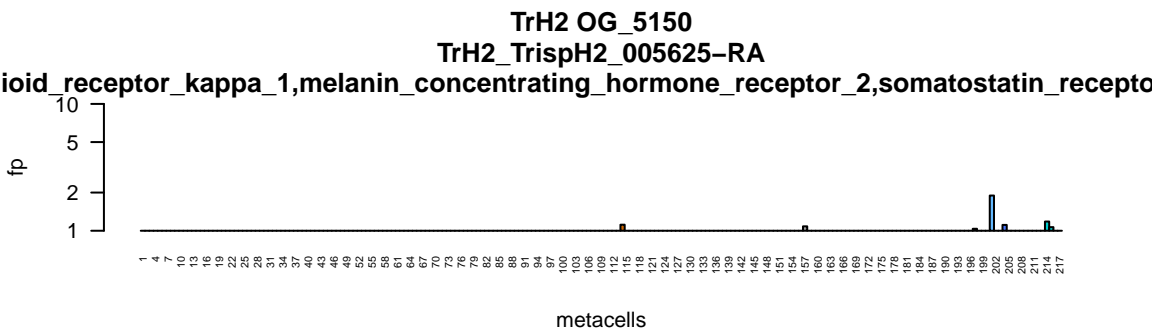
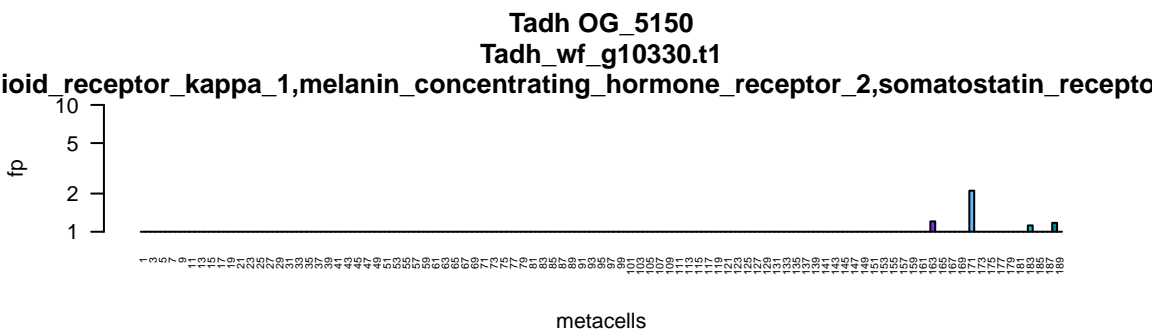
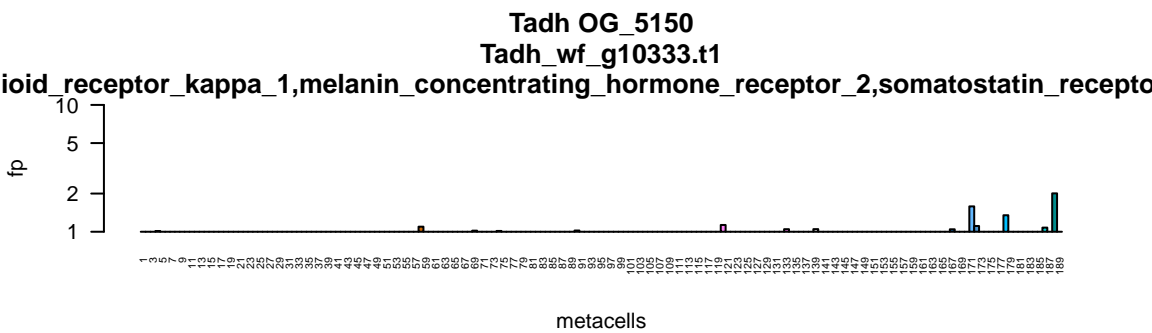
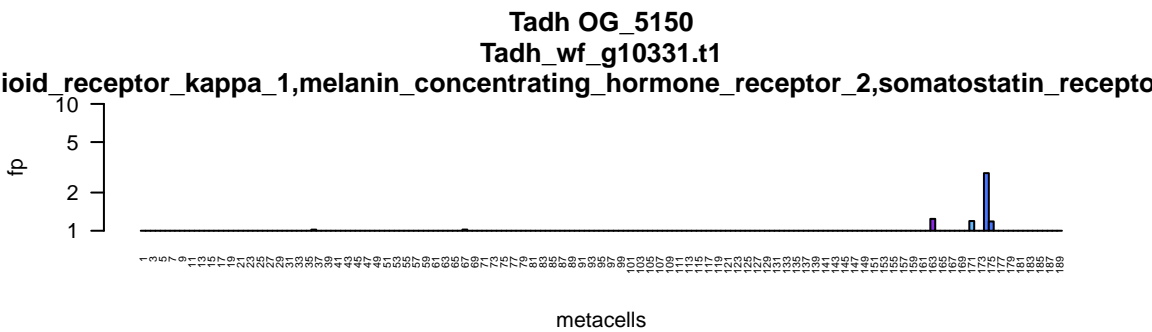


Hhon OG_5118
Hhon_g07855.t1



HoiH23 OG_5118
HoiH23_PIH23_007704-RA





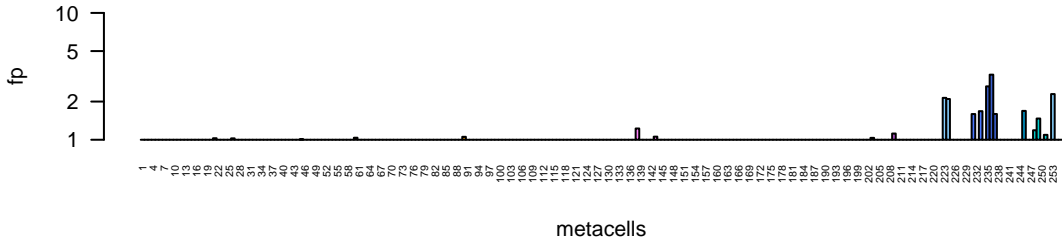
Tadh | no data



TrH2 | no data

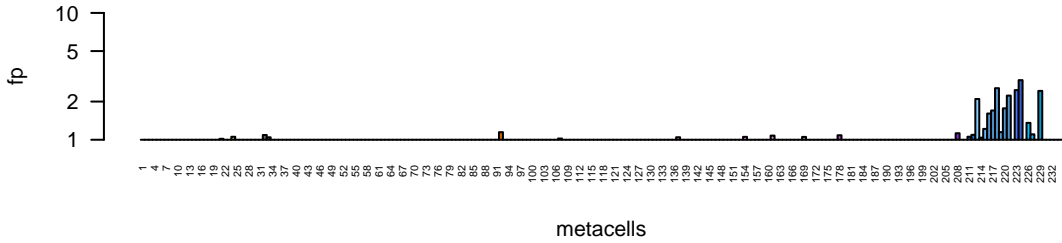


Hhon OG_5311
Hhon_g07916.t1



metacells

HoiH23 OG_5311
HoiH23_PIH23_006433-RA

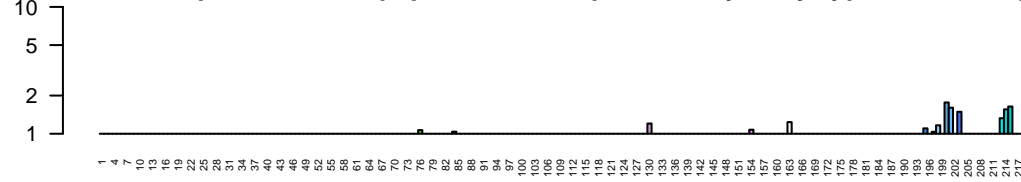


metacells

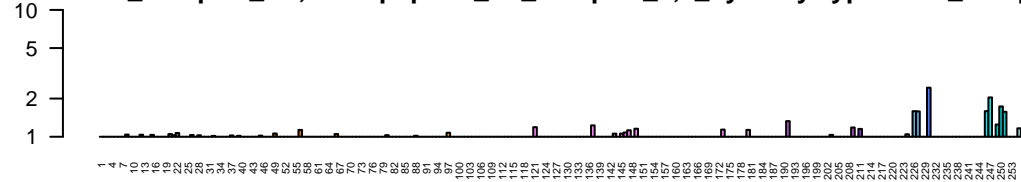
melatonin_receptor_1A,neuropeptide_FF_receptor_1,5_hydroxytryptamine_receptor_1A
Tadh | no data



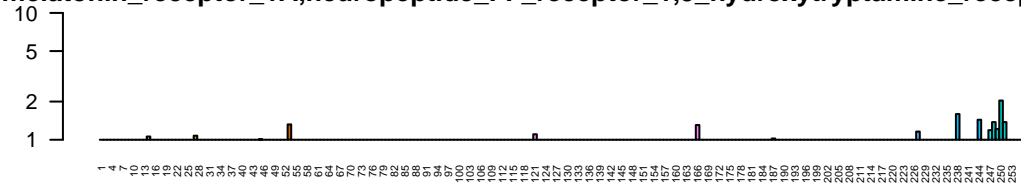
TrH2 OG_5421
TrH2_TrispH2_008291-RA
melatonin_receptor_1A,neuropeptide_FF_receptor_1,5_hydroxytryptamine_receptor_1A



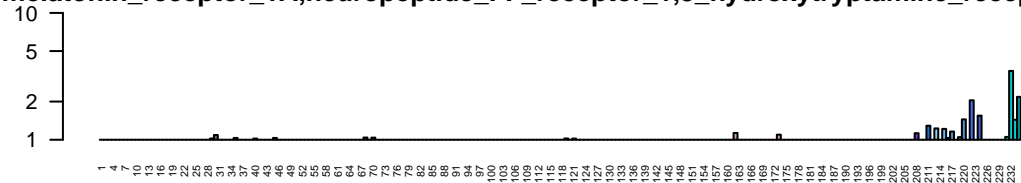
Hhon OG_5421
Hhon_g11348.t1
melatonin_receptor_1A,neuropeptide_FF_receptor_1,5_hydroxytryptamine_receptor_1A

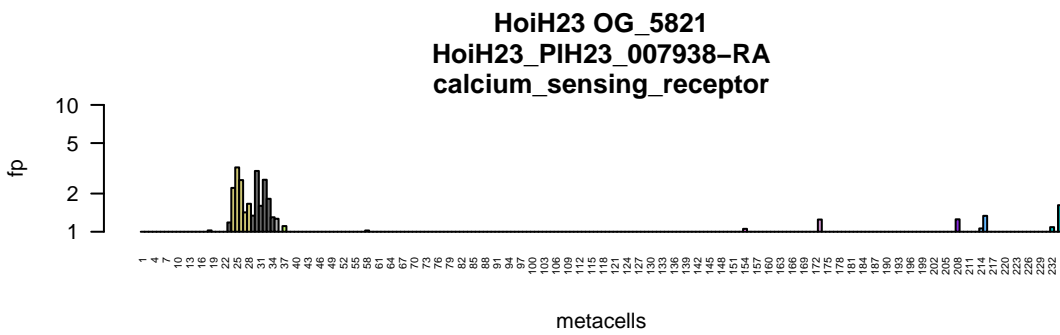
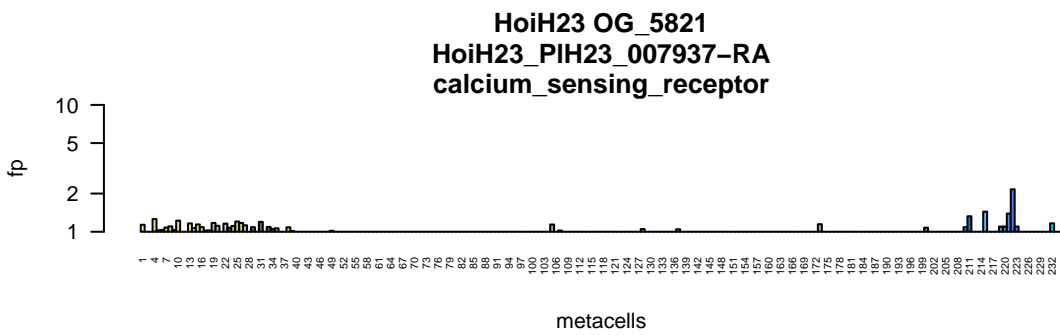
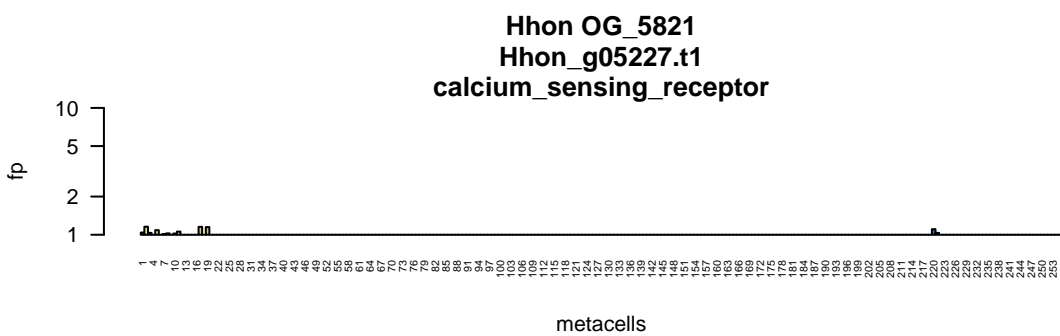
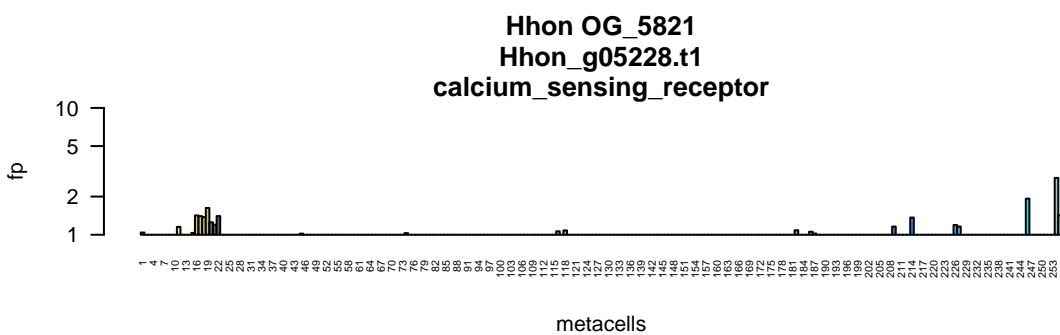
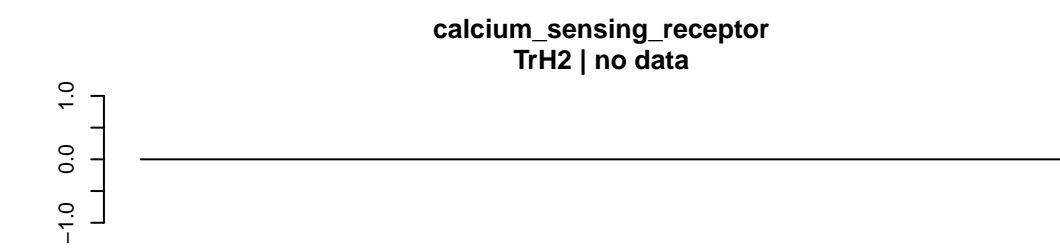
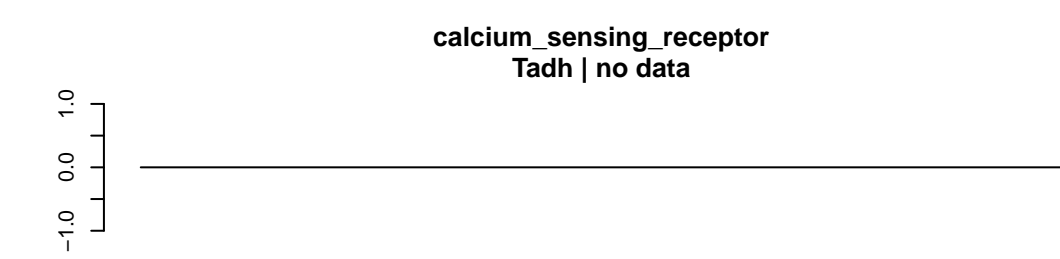


Hhon OG_5421
Hhon_g05484.t1
melatonin_receptor_1A,neuropeptide_FF_receptor_1,5_hydroxytryptamine_receptor_1A



HoiH23 OG_5421
HoiH23_PIH23_006174-RA
melatonin_receptor_1A,neuropeptide_FF_receptor_1,5_hydroxytryptamine_receptor_1A

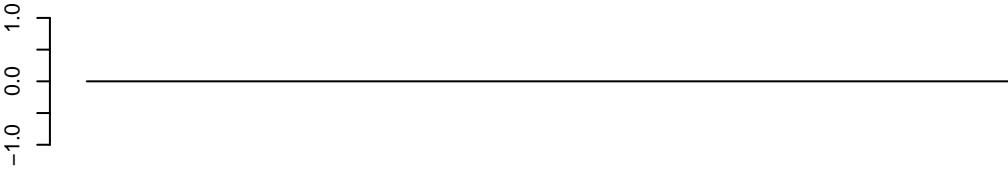




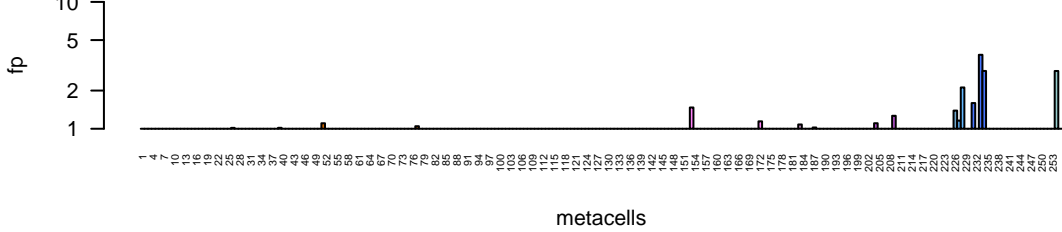
gamma_aminobutyric_acid_type_B_receptor_subunit_2
Tadh | no data



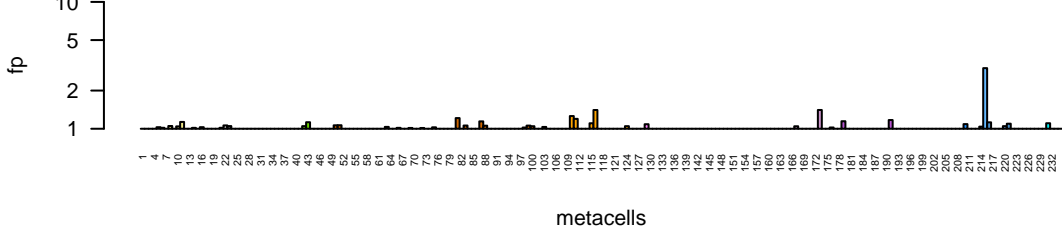
gamma_aminobutyric_acid_type_B_receptor_subunit_2
TrH2 | no data

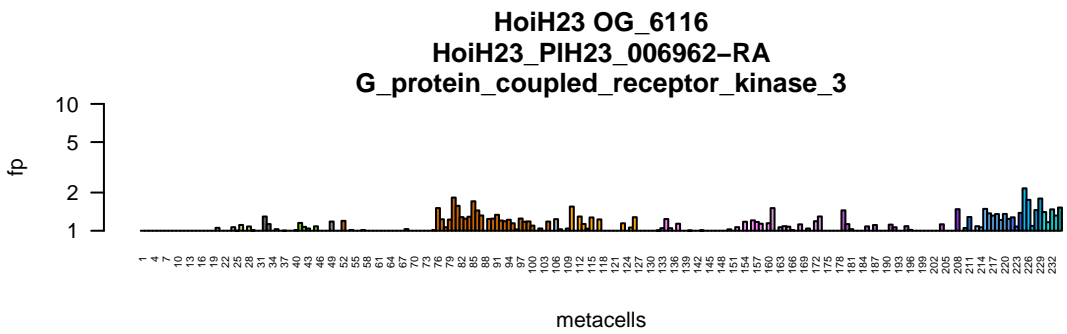
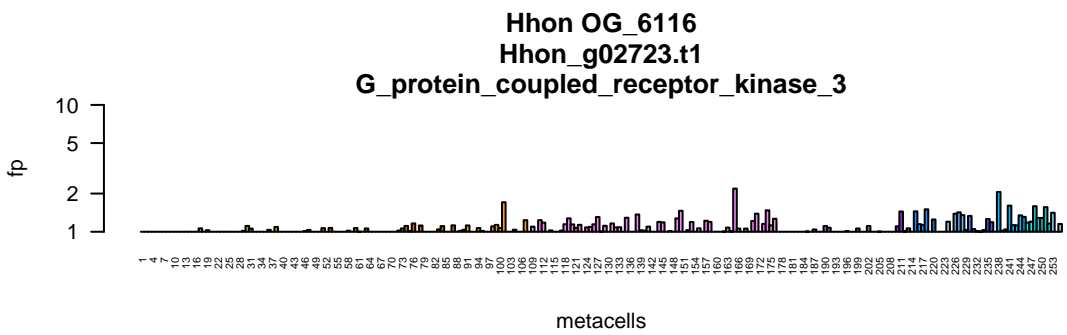
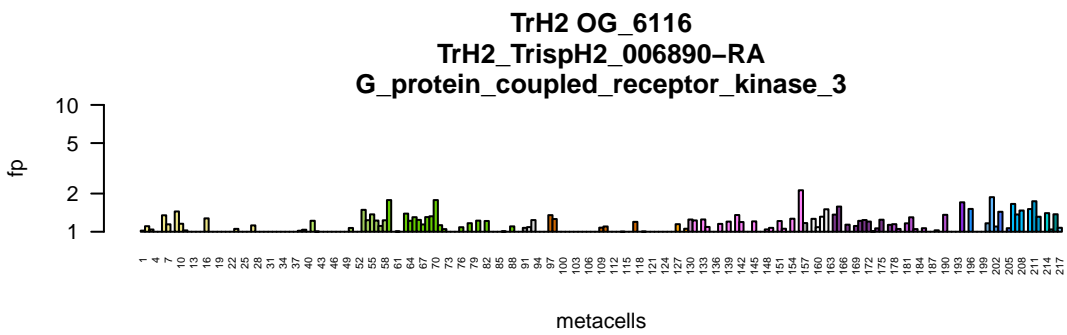
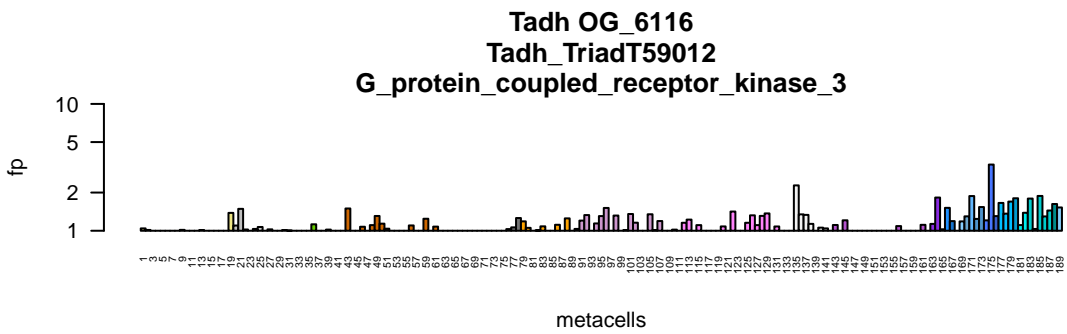


Hhon OG_6010
Hhon_g01250.t1
gamma_aminobutyric_acid_type_B_receptor_subunit_2

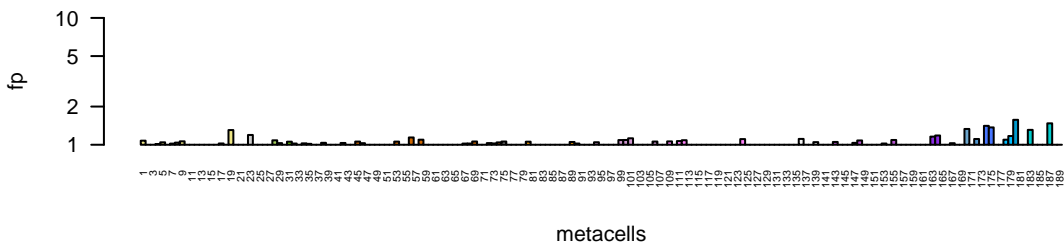


HoiH23 OG_6010
HoiH23_PIH23_000005-RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2

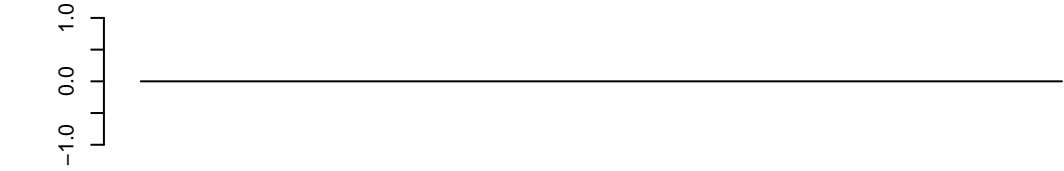




Tadh OG_6121
Tadh_wf_g9892.t1



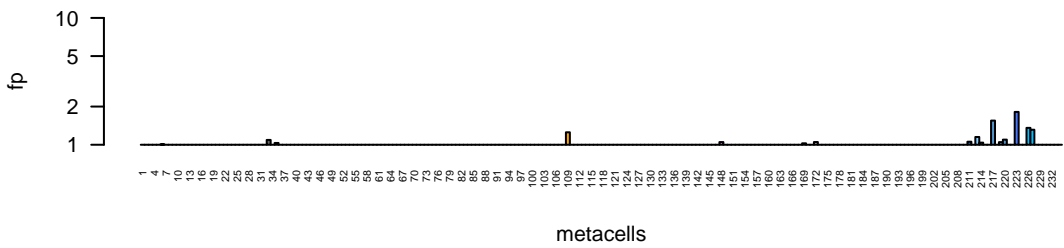
TrH2 | no data

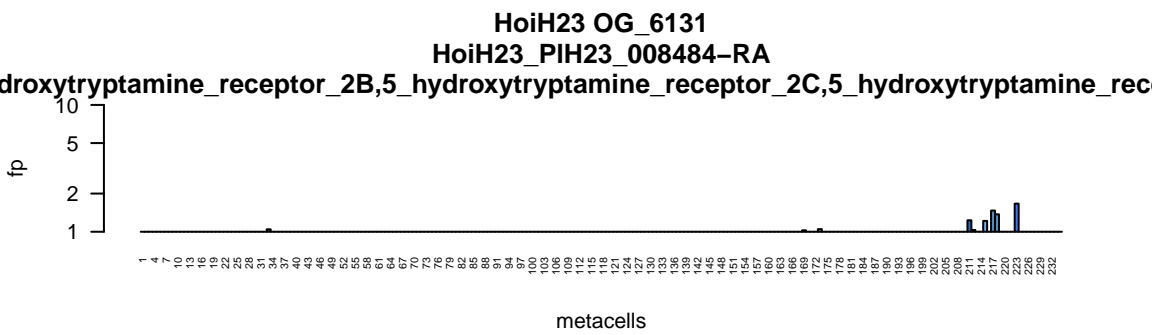
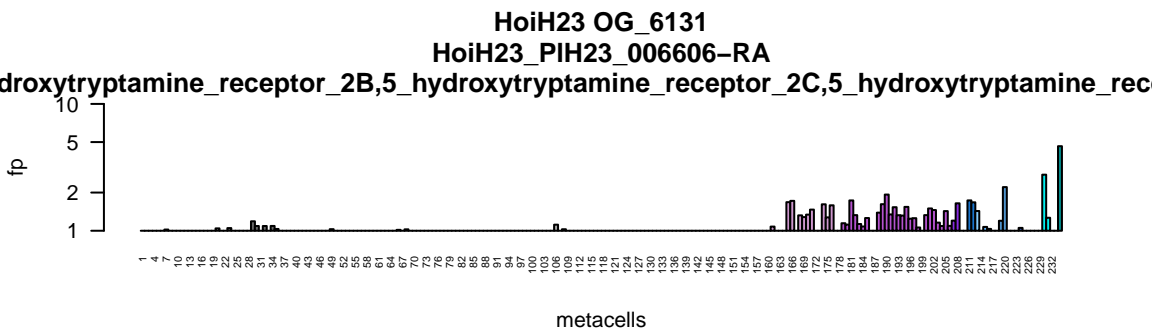
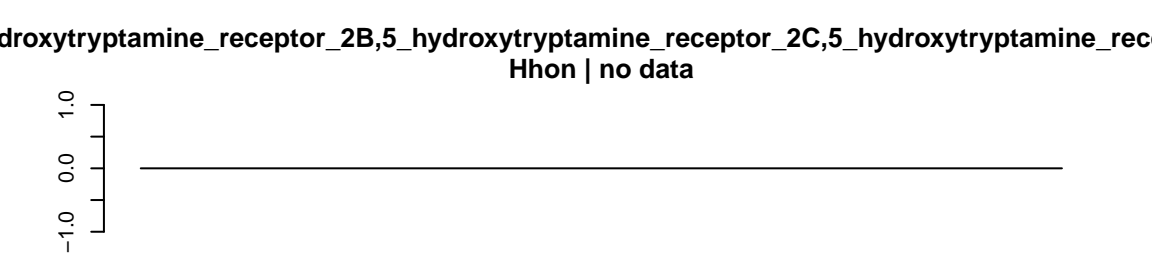
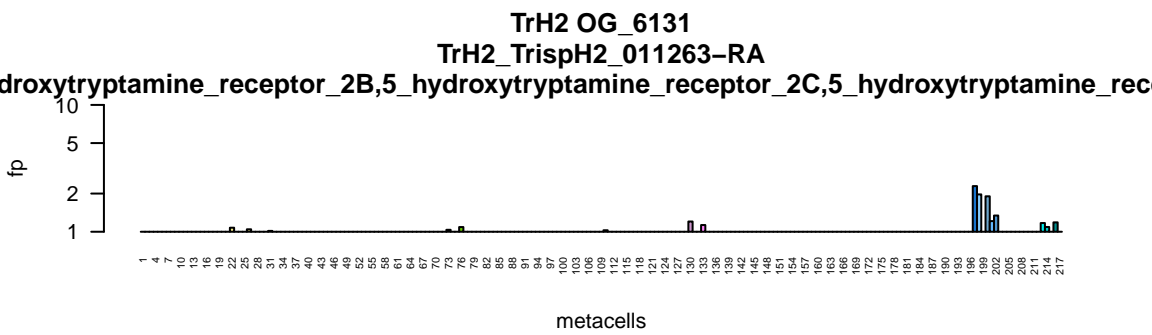
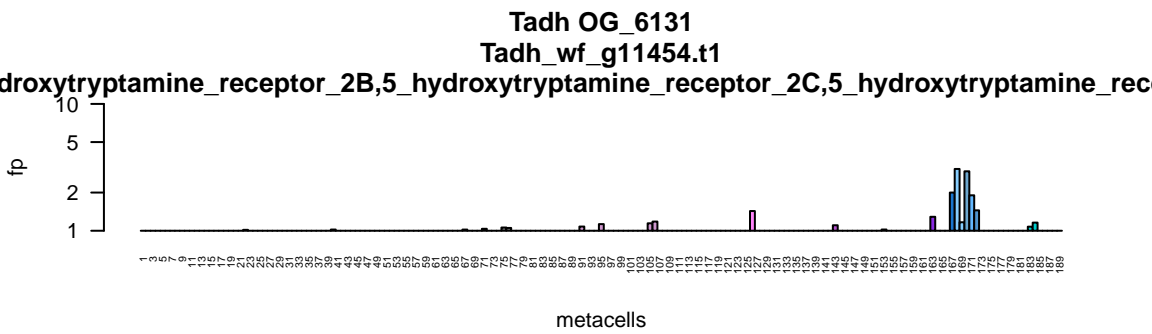


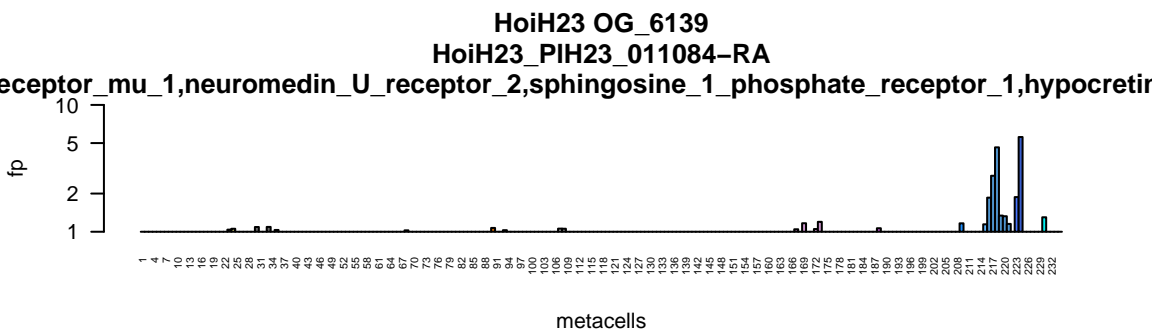
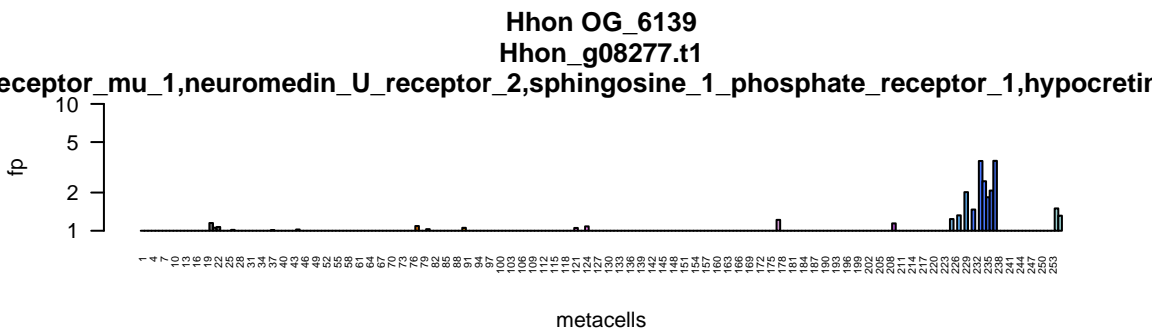
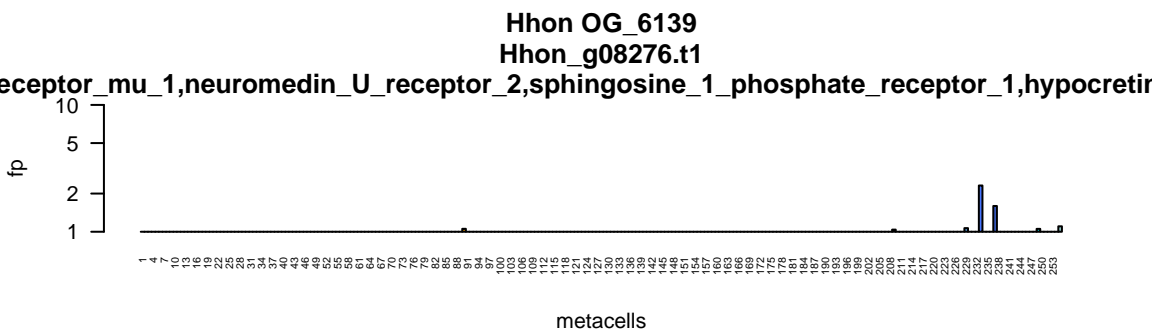
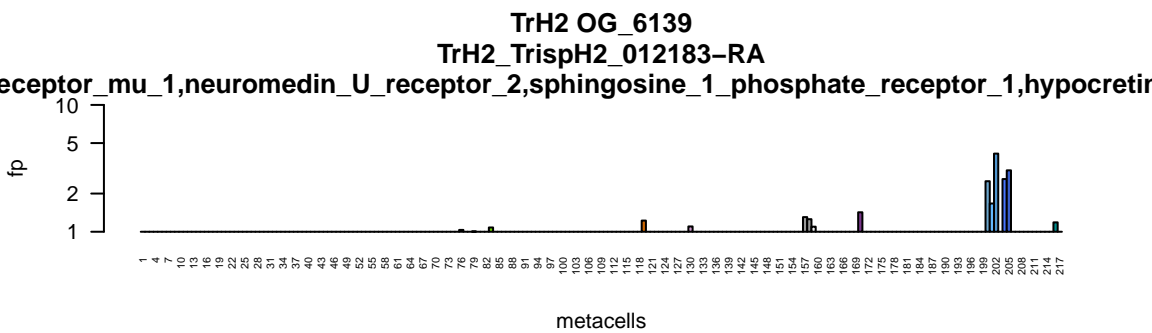
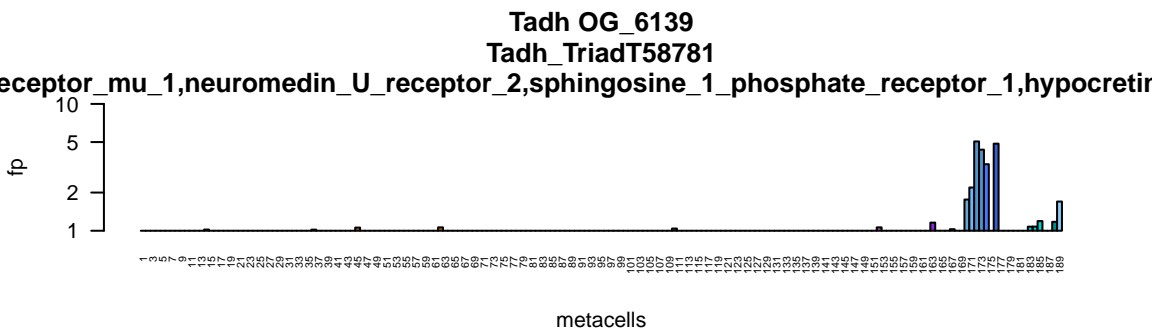
Hhon | no data

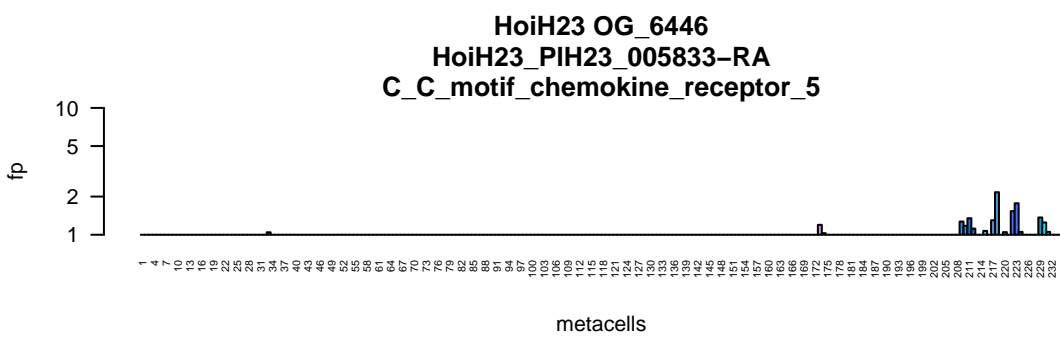
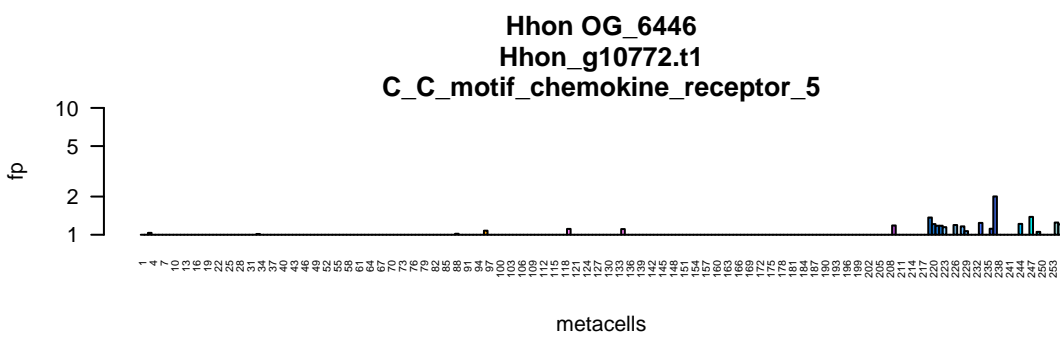
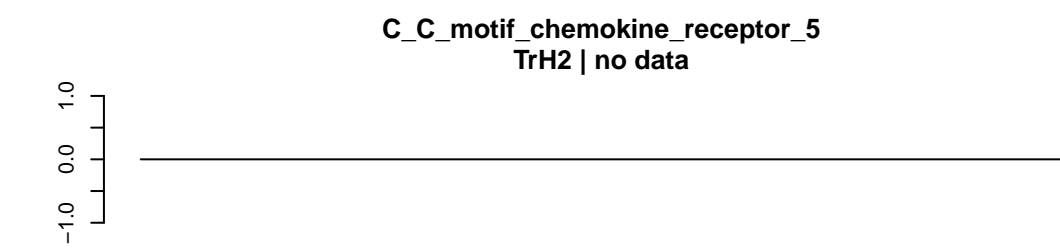
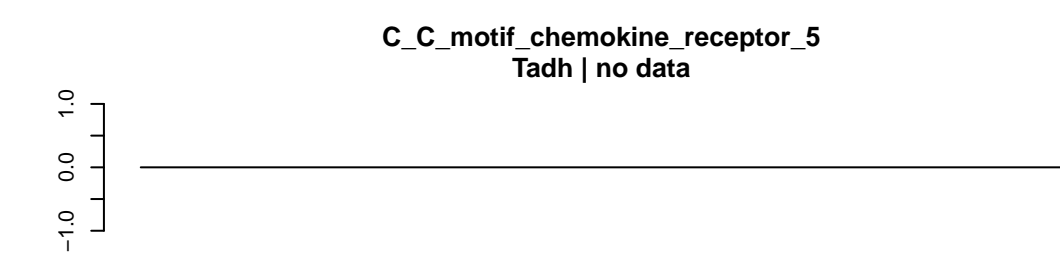


HoiH23 OG_6121
HoiH23_PIH23_008468-RA

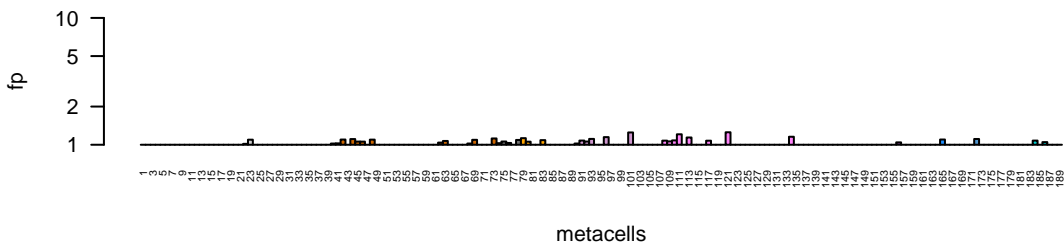








Tadh OG_6636
Tadh_TriadT61507



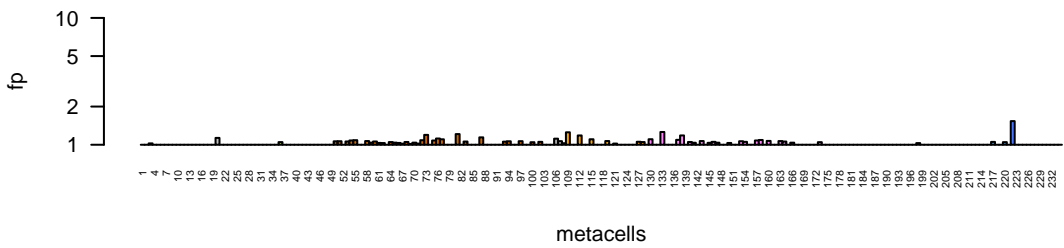
TrH2 | no data

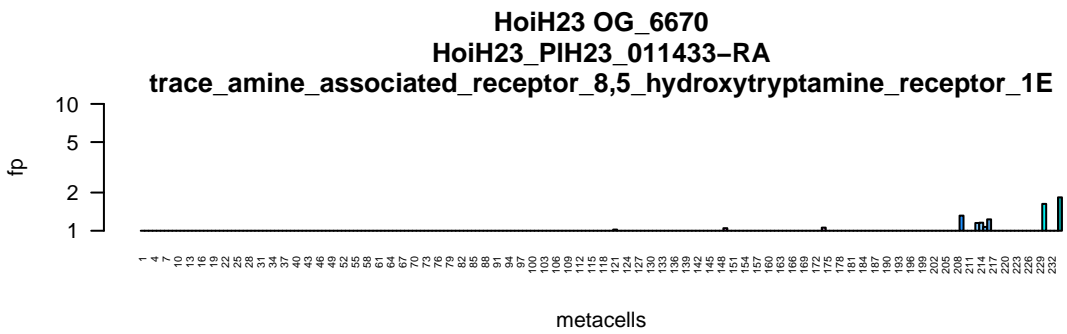
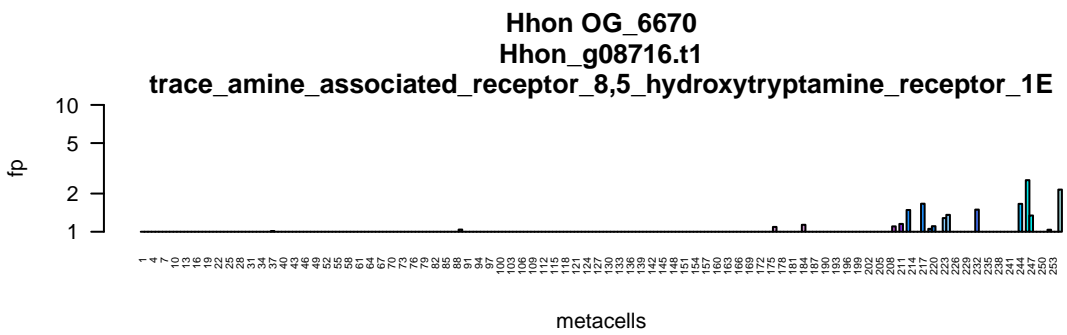
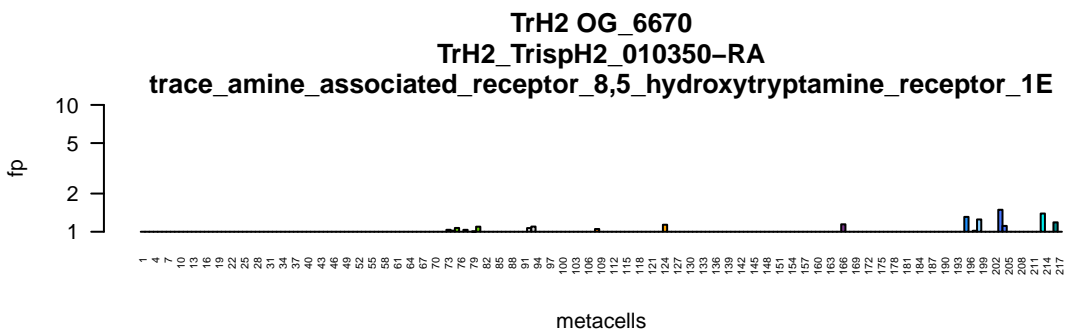
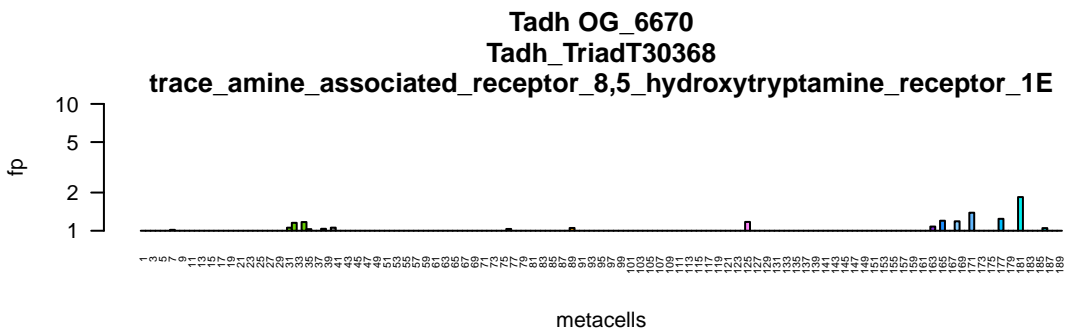


Hhon | no data

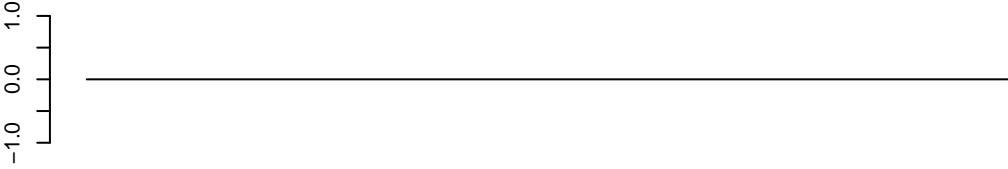


HoiH23 OG_6636
HoiH23_PIH23_010246-RA

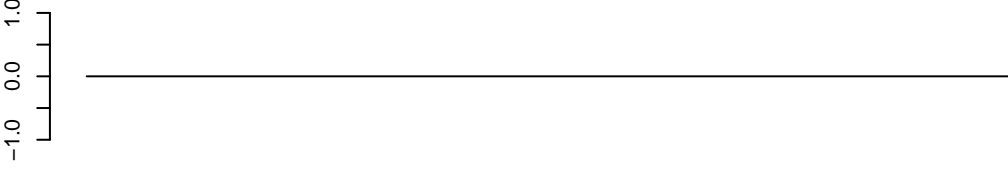




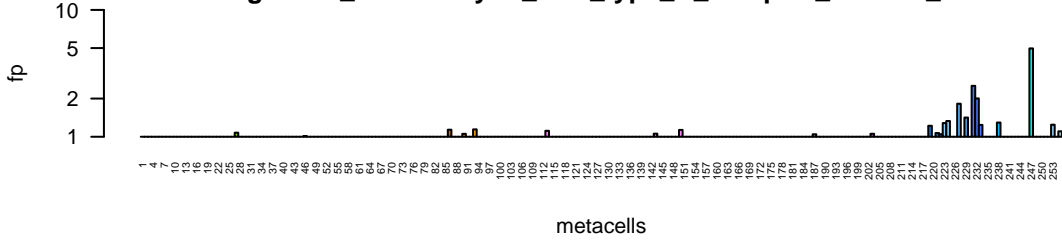
gamma_aminobutyric_acid_type_B_receptor_subunit_1
Tadh | no data



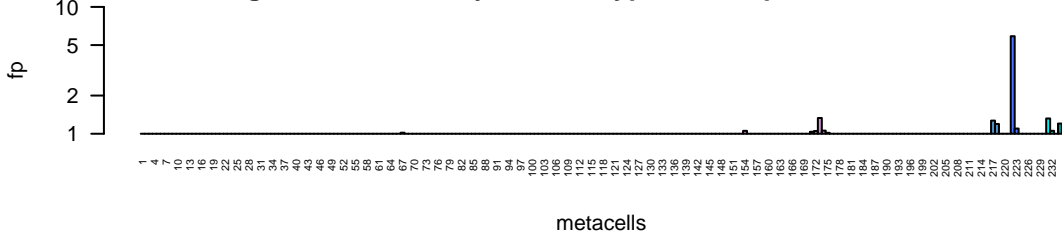
gamma_aminobutyric_acid_type_B_receptor_subunit_1
TrH2 | no data



Hhon OG_6851
Hhon_g09350.t1
gamma_aminobutyric_acid_type_B_receptor_subunit_1



HoiH23 OG_6851
HoiH23_PIH23_008905-RA
gamma_aminobutyric_acid_type_B_receptor_subunit_1



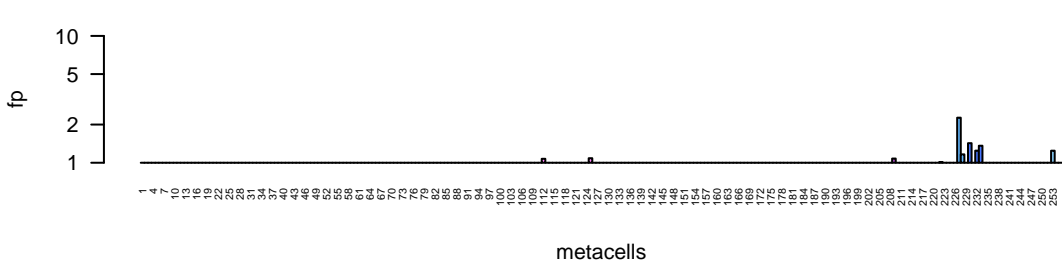
Tadh | no data



TrH2 | no data

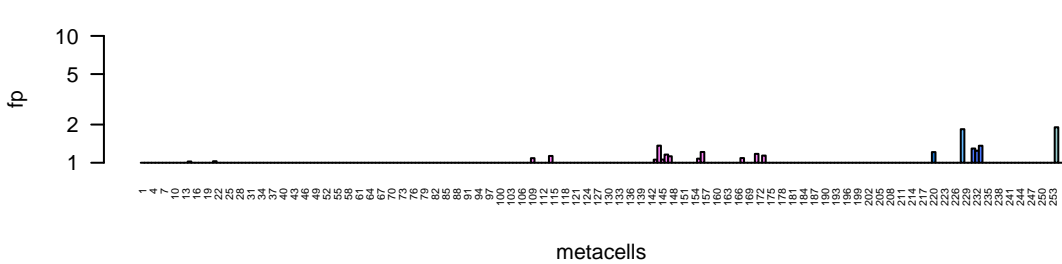


Hhon OG_6868
Hhon_g08412.t1



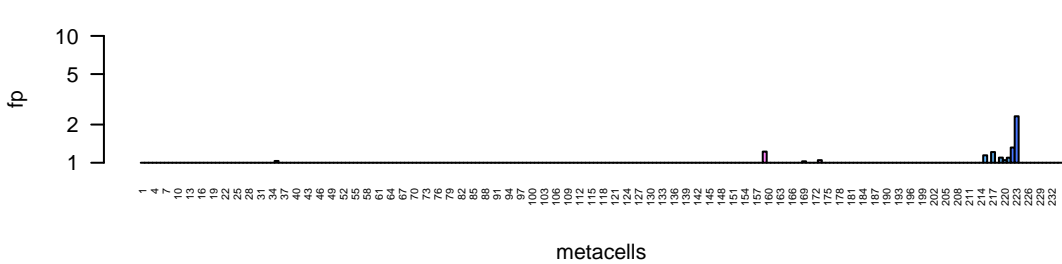
metacells

Hhon OG_6868
Hhon_g07653.t1



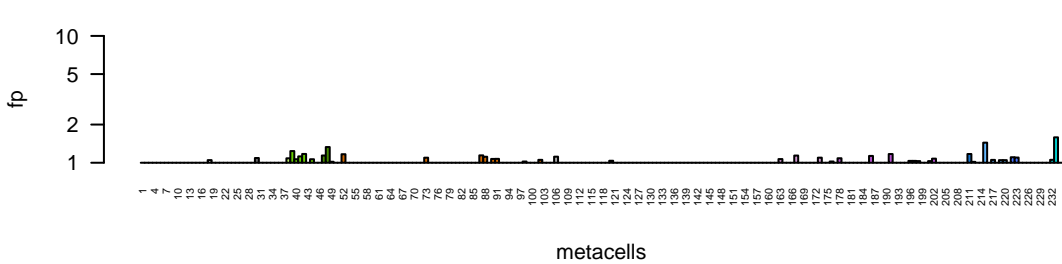
metacells

HoiH23 OG_6868
HoiH23_PIH23_011708-RA

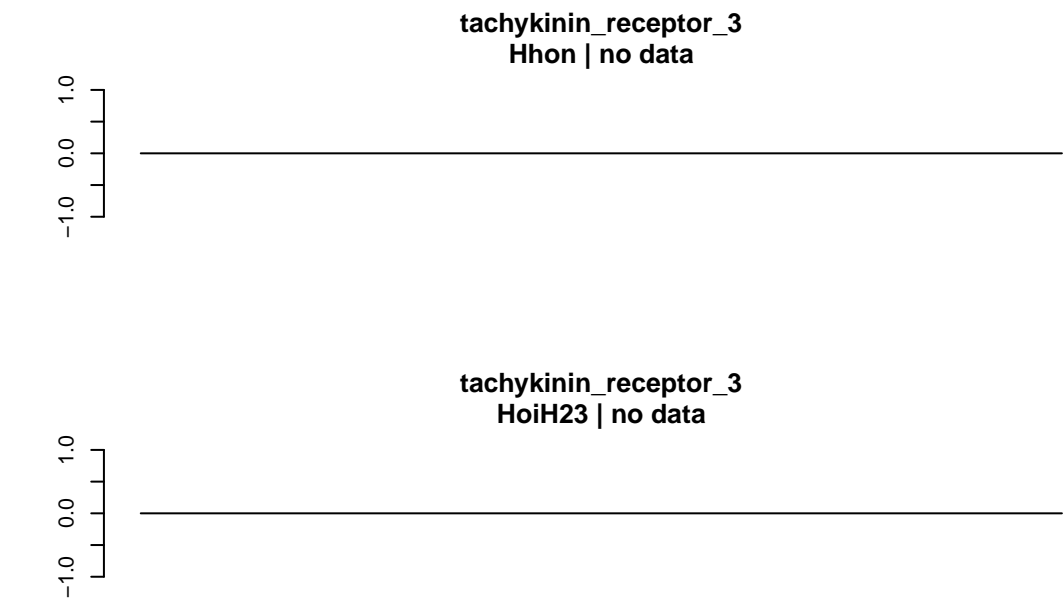
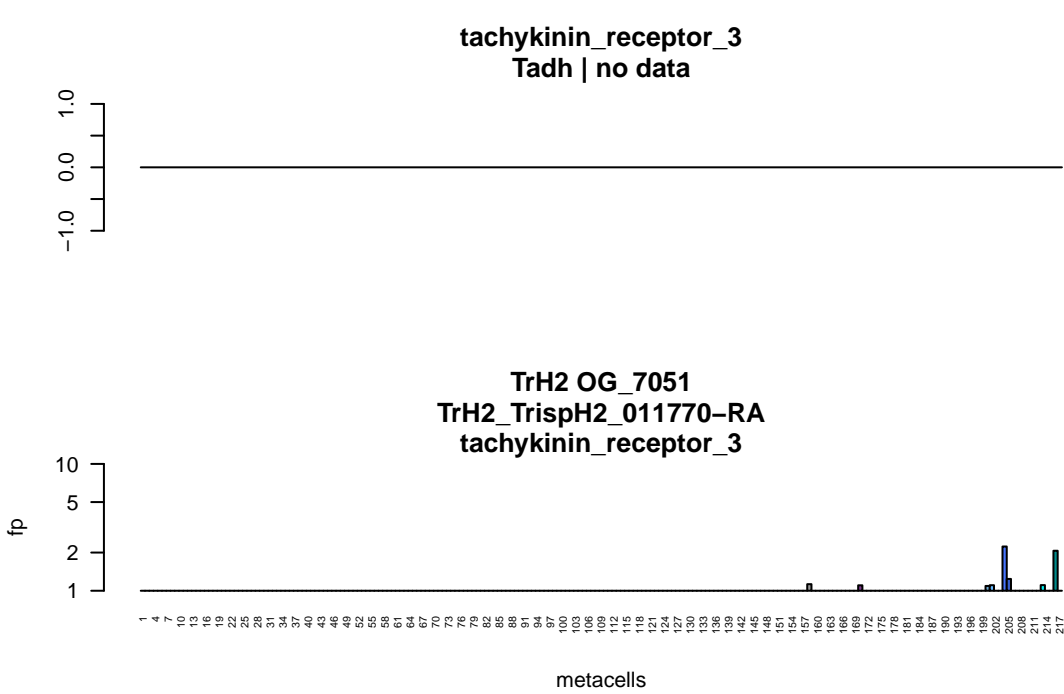


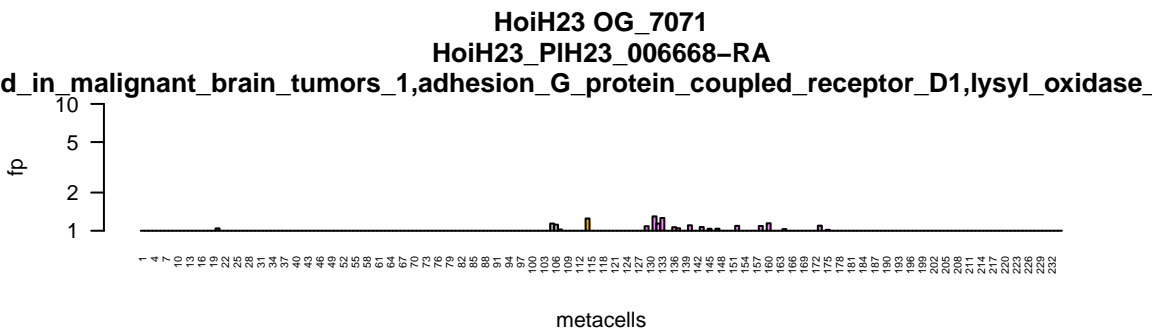
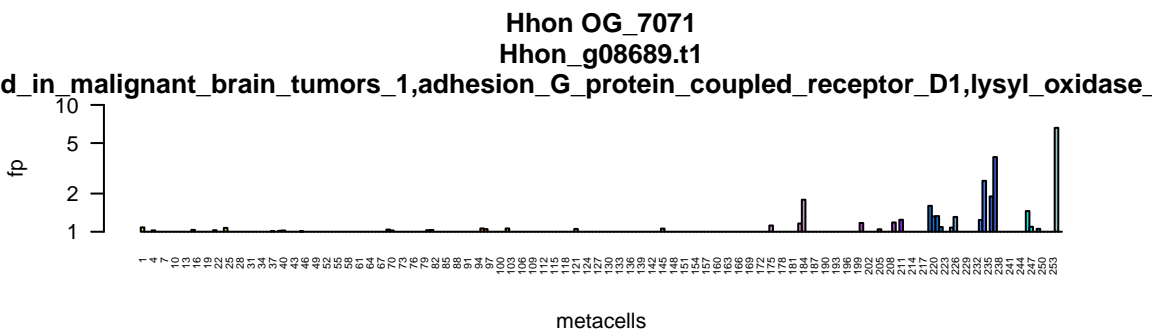
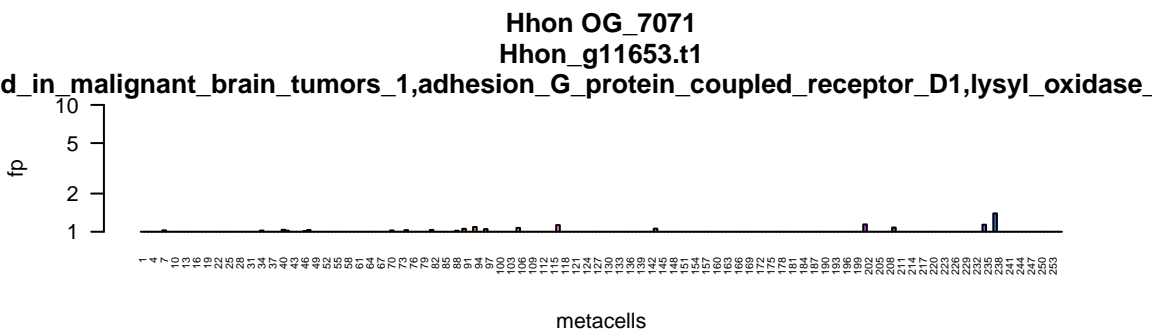
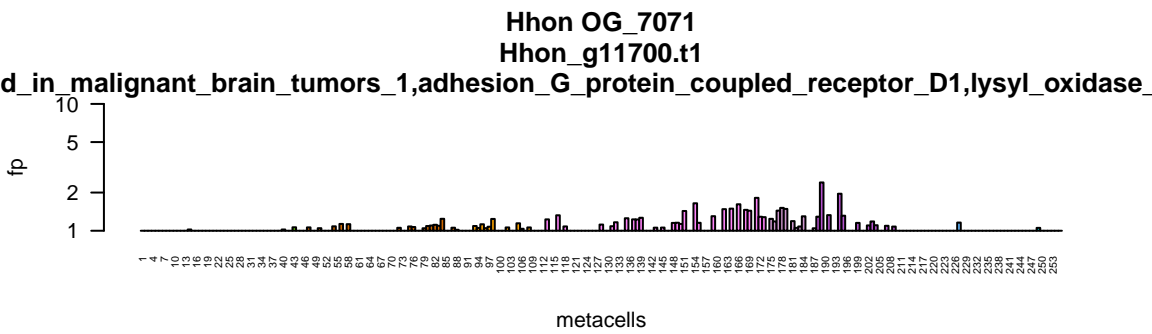
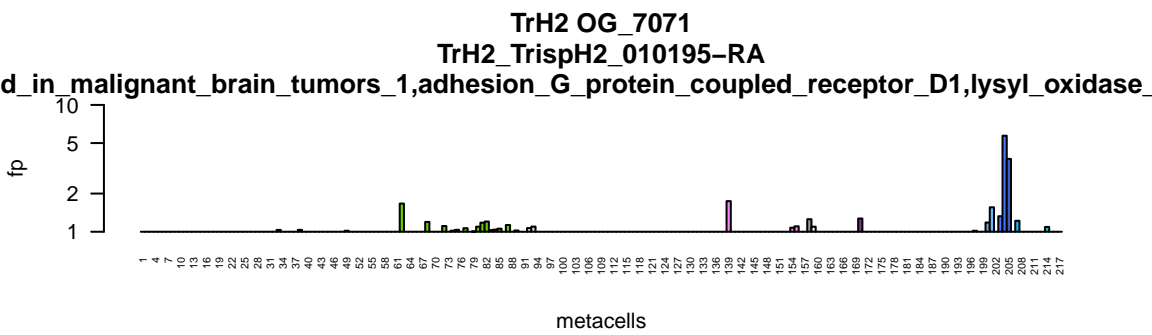
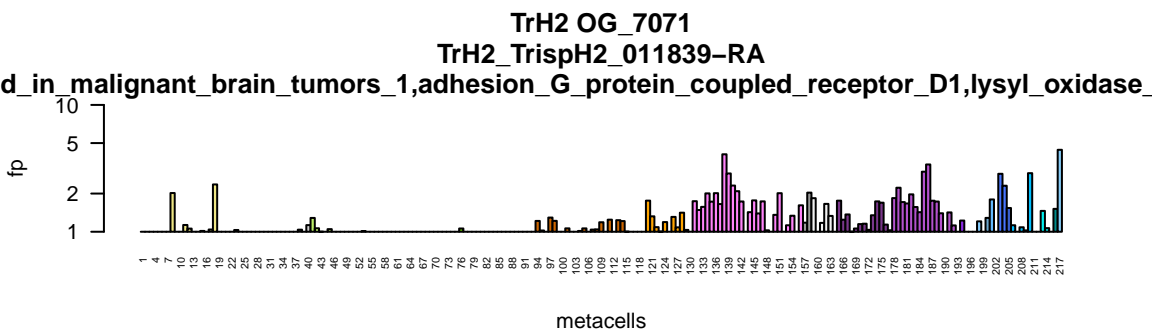
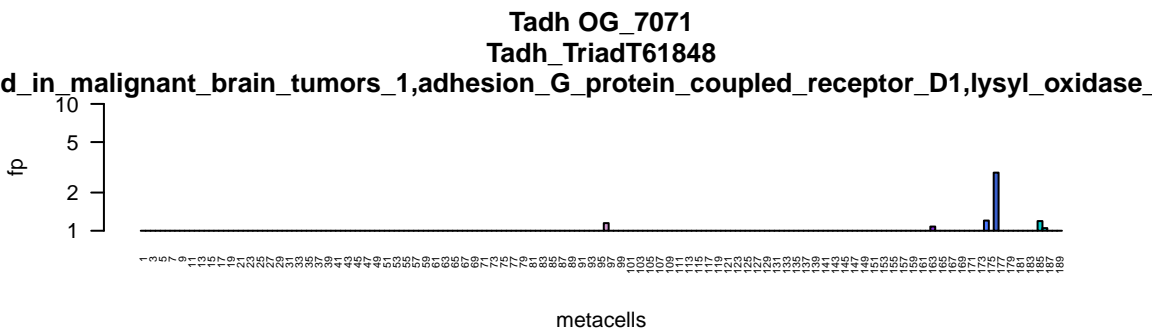
metacells

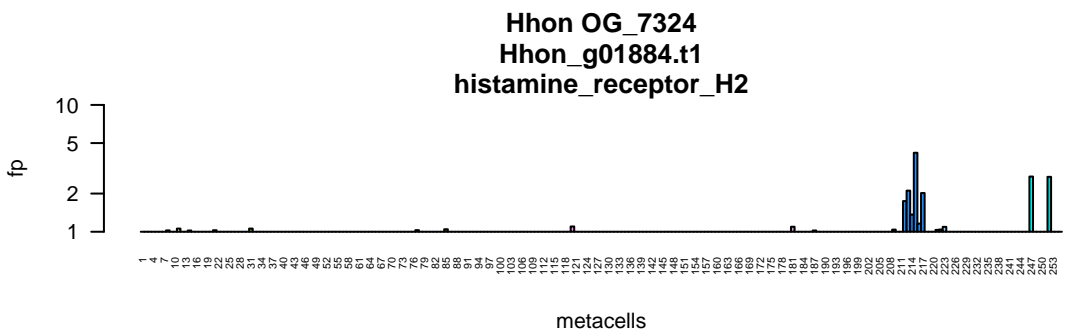
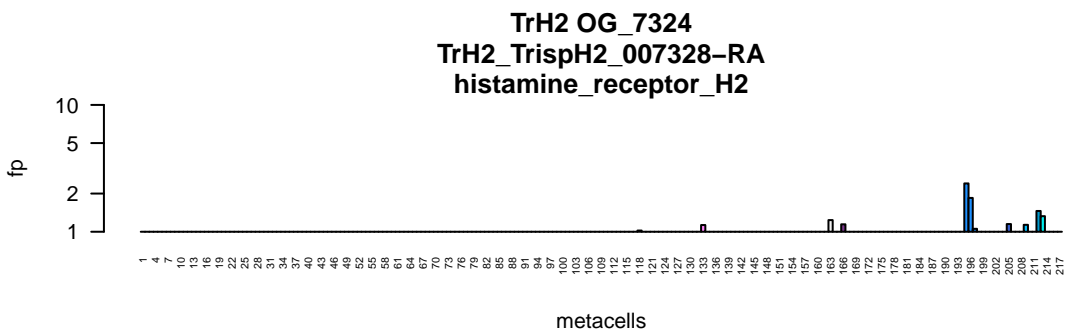
HoiH23 OG_6868
HoiH23_PIH23_011709-RA

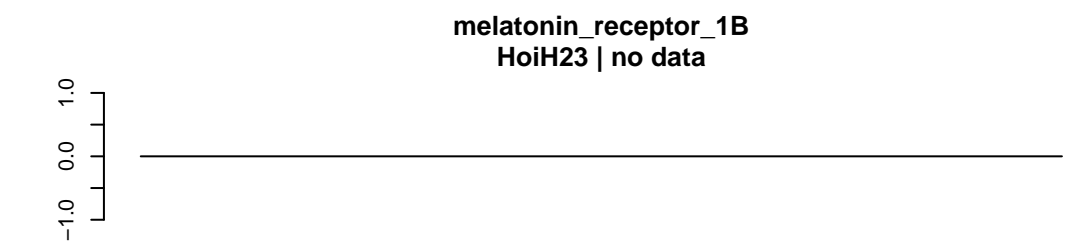
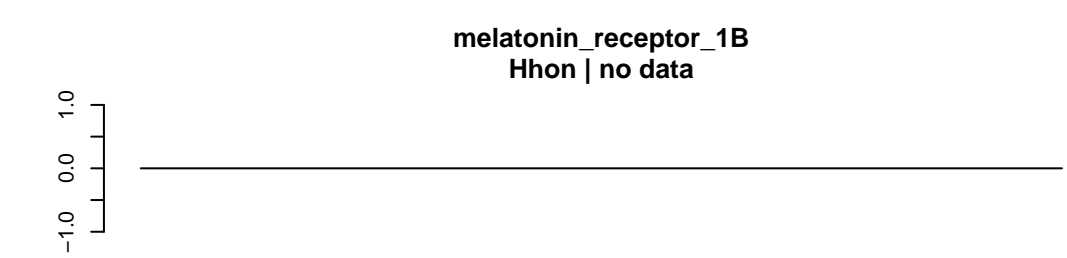
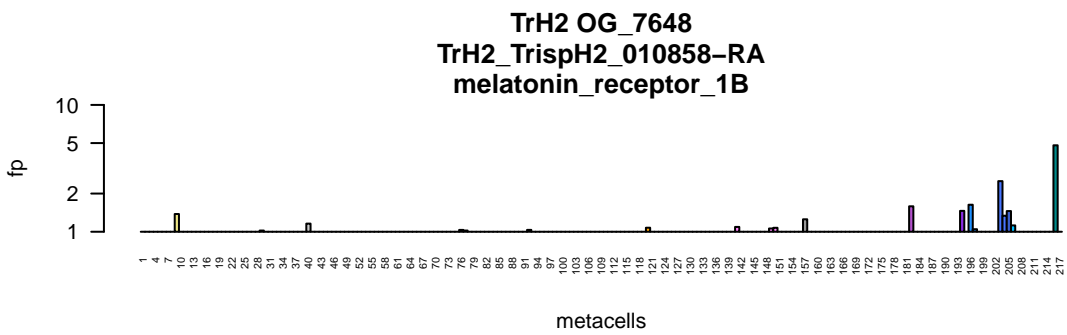
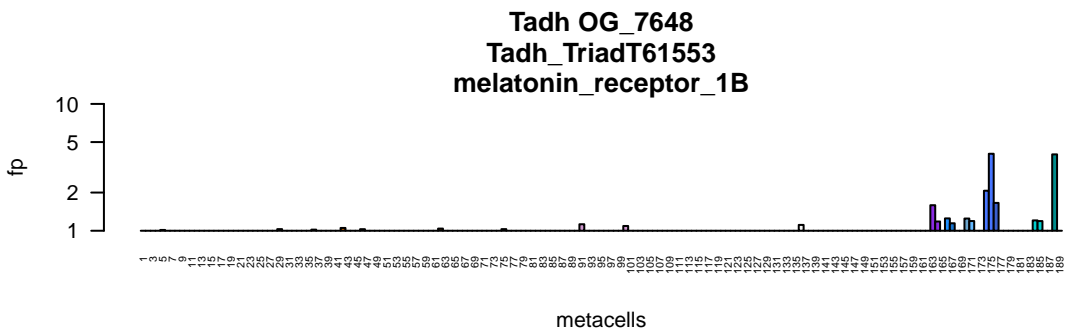


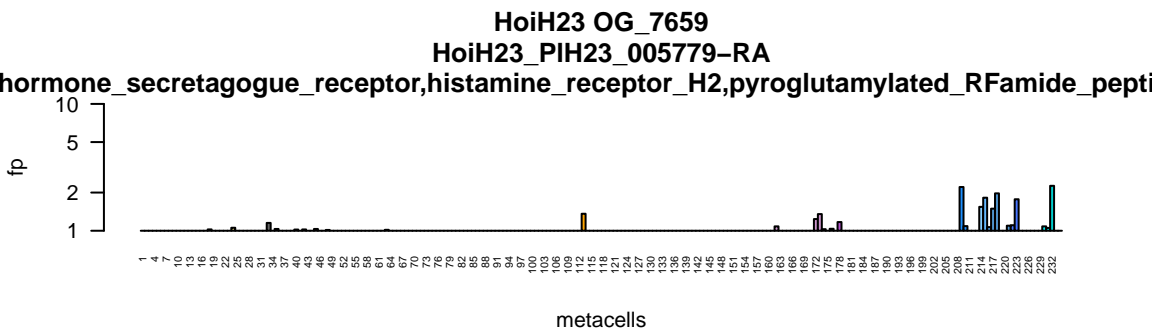
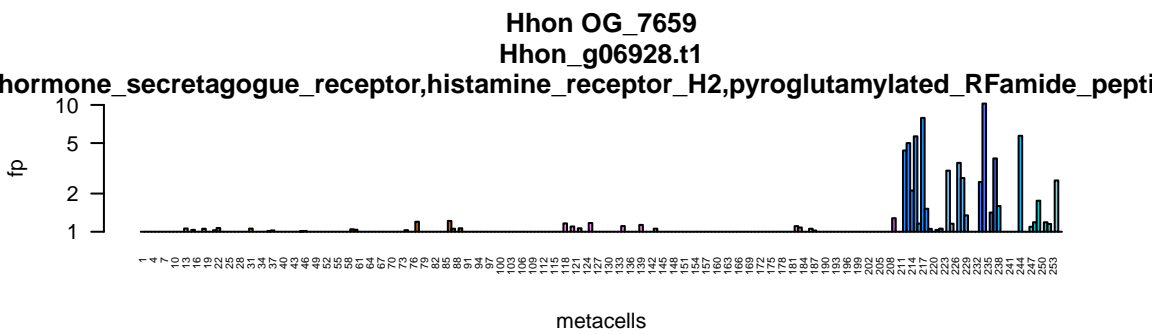
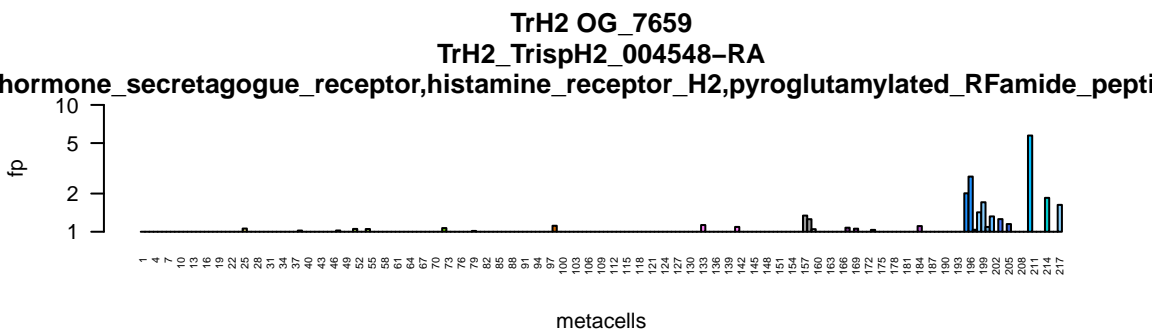
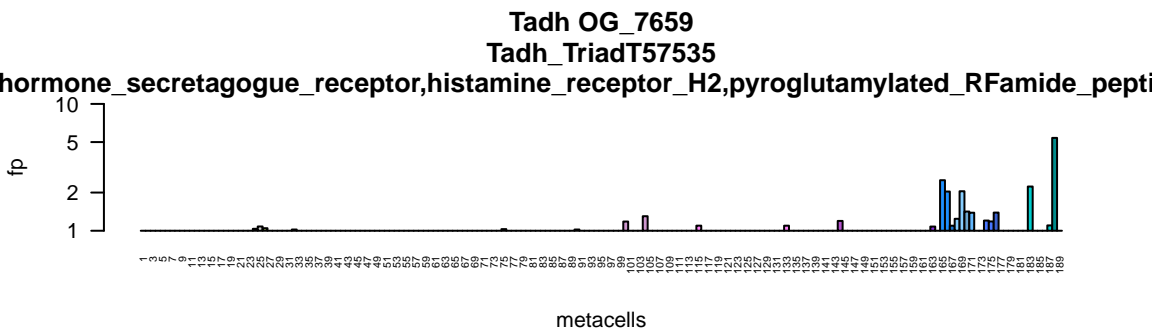
metacells



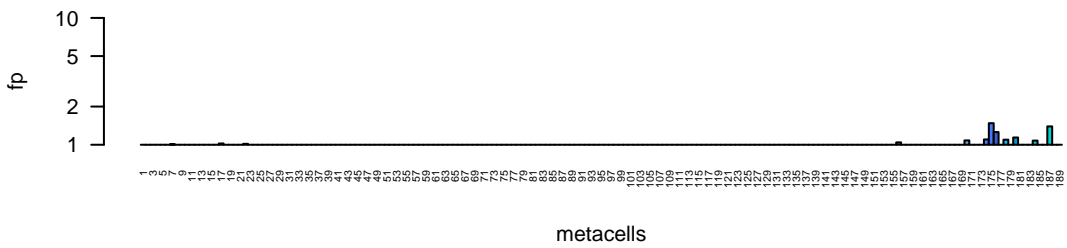




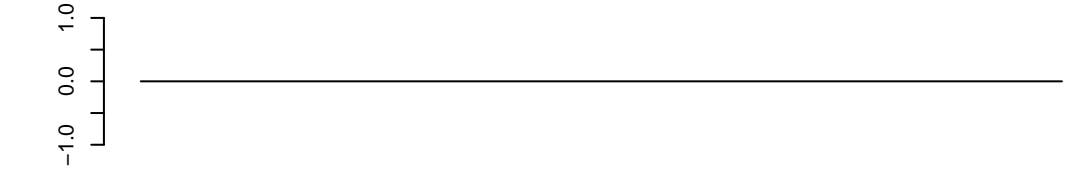




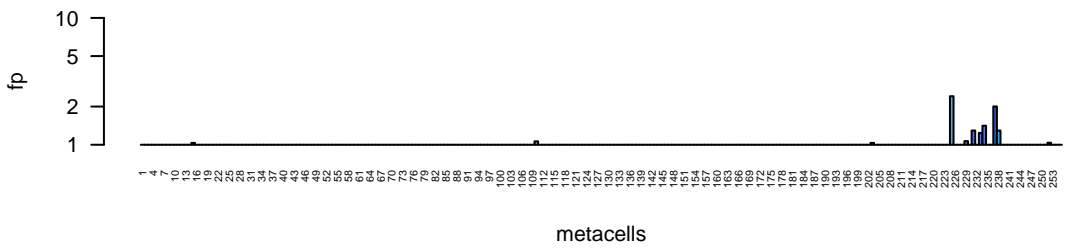
Tadh OG_7675
Tadh_wf_g9283.t1



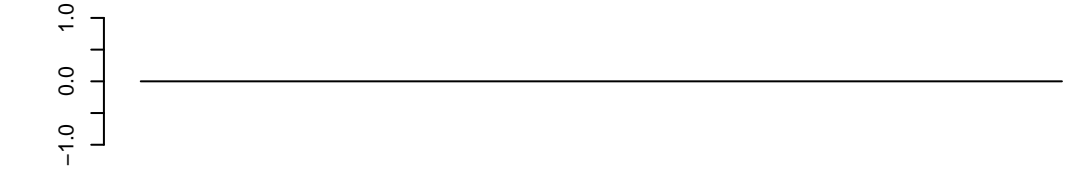
TrH2 | no data



Hhon OG_7675
Hhon_g09919.t1

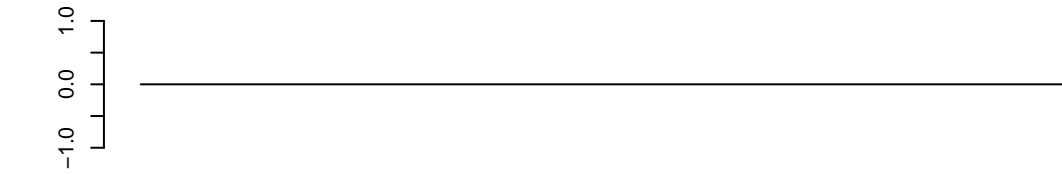


HoiH23 | no data



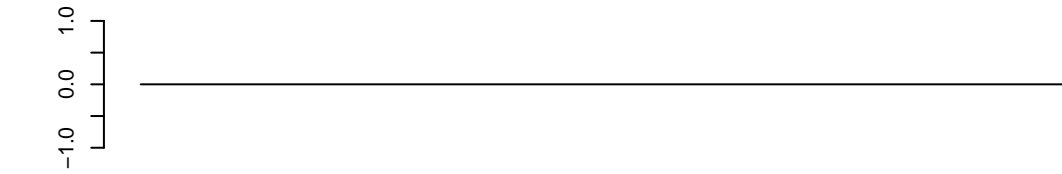
C_X_C_motif_chemokine_receptor_2,opsin_1_short_wave_sensitive,opioid_receptor_kappa

Tadh | no data



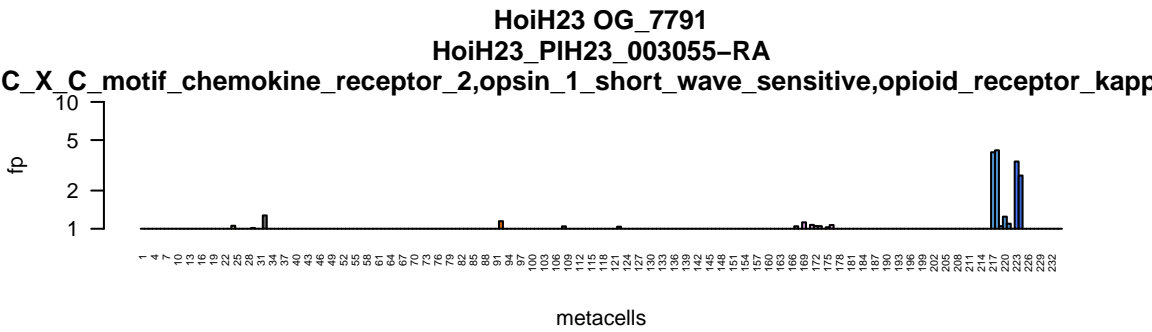
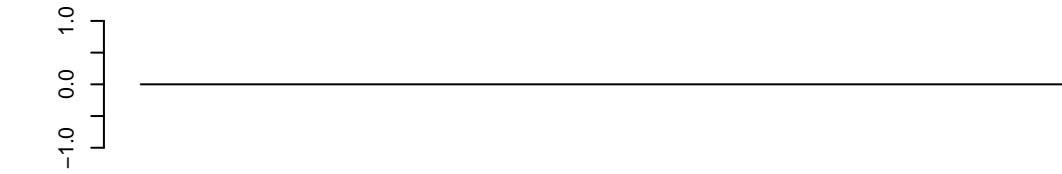
C_X_C_motif_chemokine_receptor_2,opsin_1_short_wave_sensitive,opioid_receptor_kappa

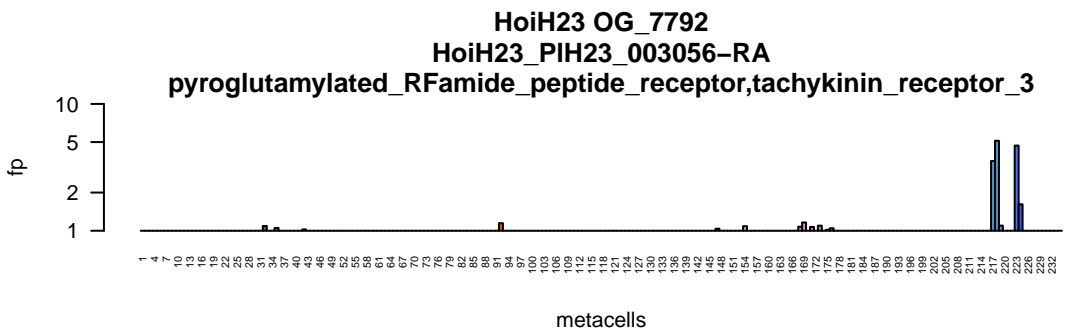
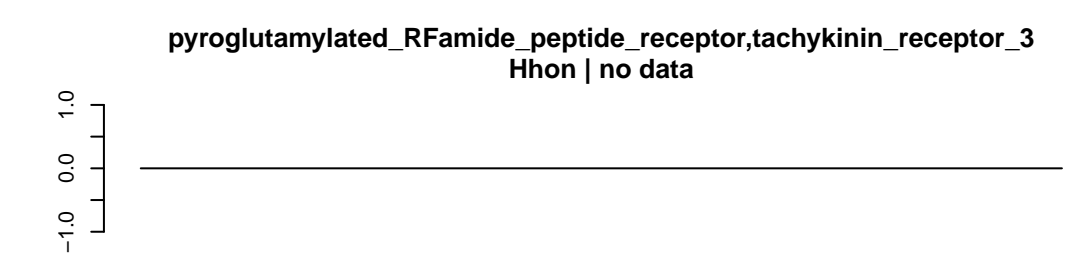
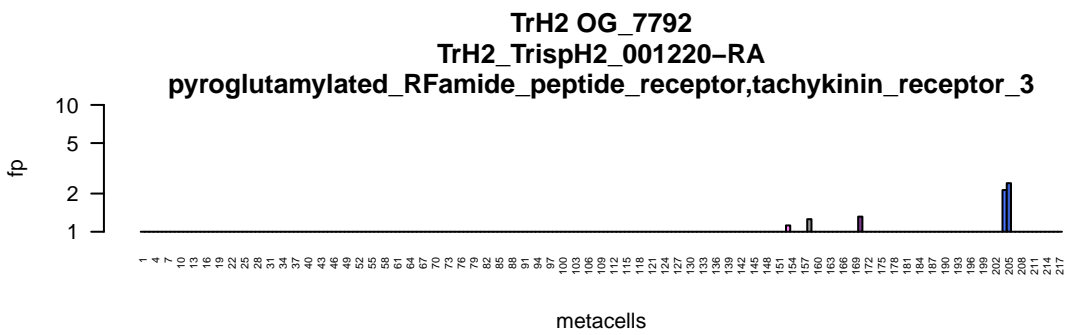
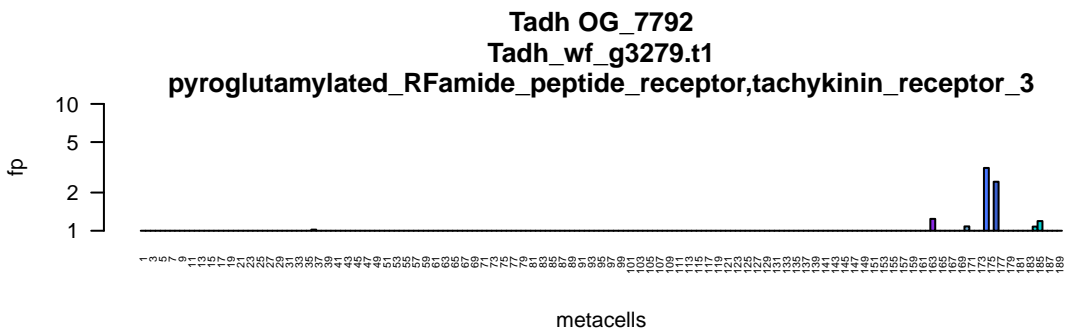
TrH2 | no data

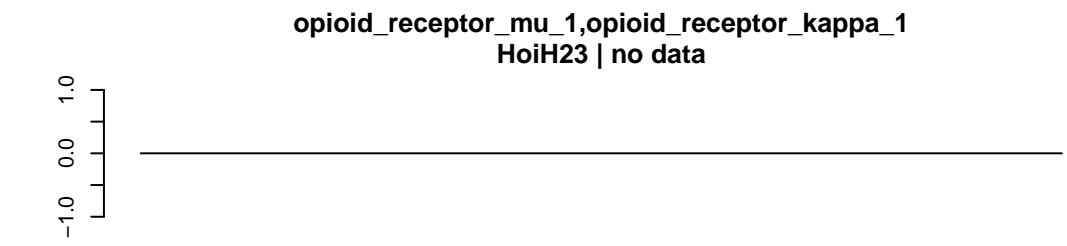
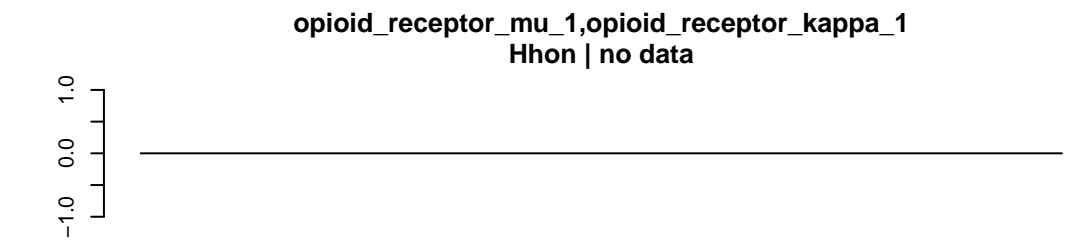
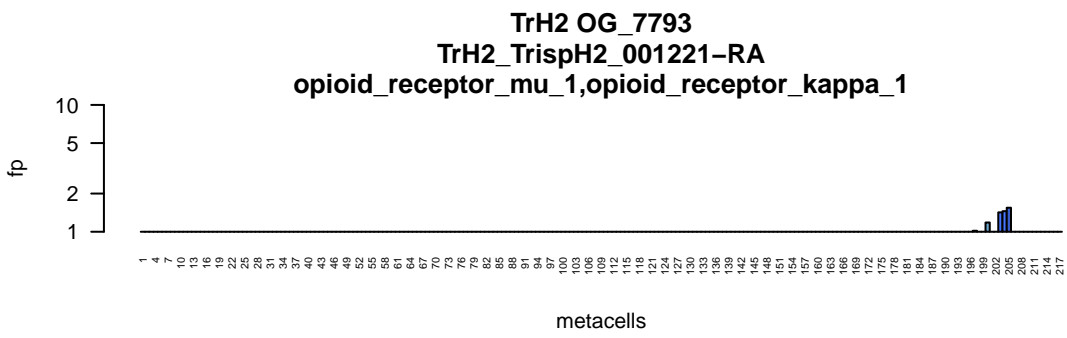
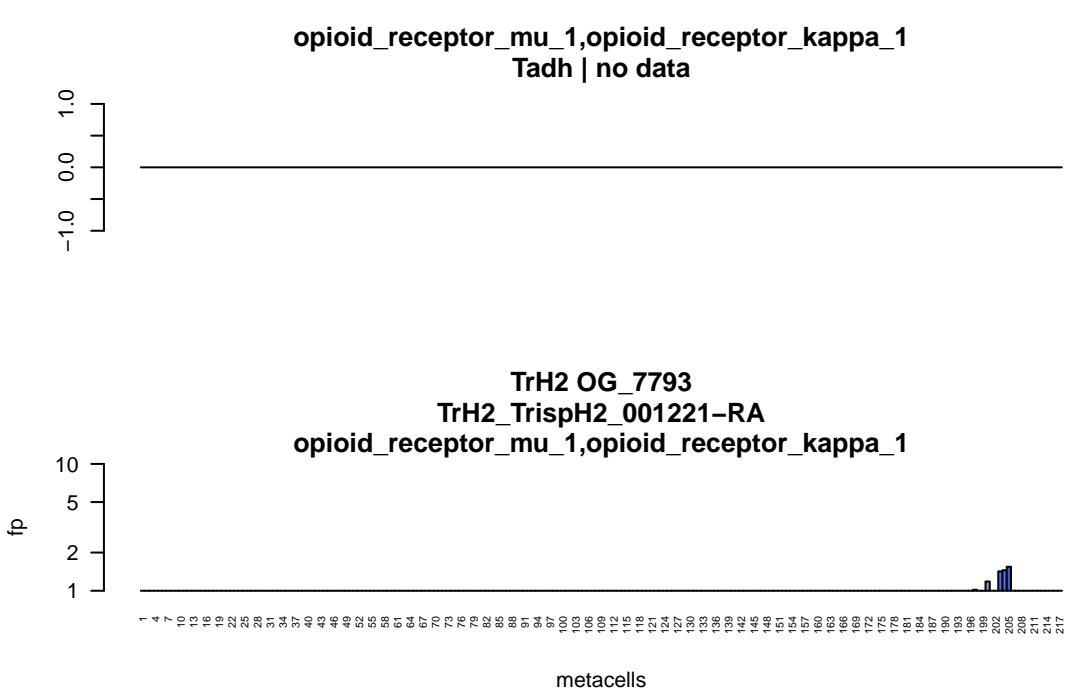


C_X_C_motif_chemokine_receptor_2,opsin_1_short_wave_sensitive,opioid_receptor_kappa

Hhon | no data



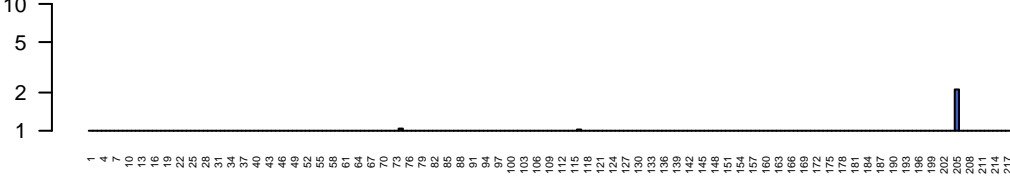




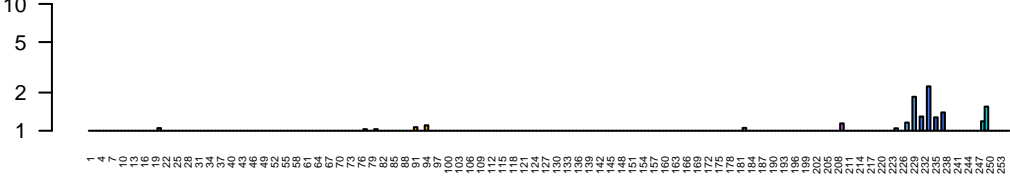
adhesion_G_protein_coupled_receptor_G4,adhesion_G_protein_coupled_receptor_G2
Tadh | no data



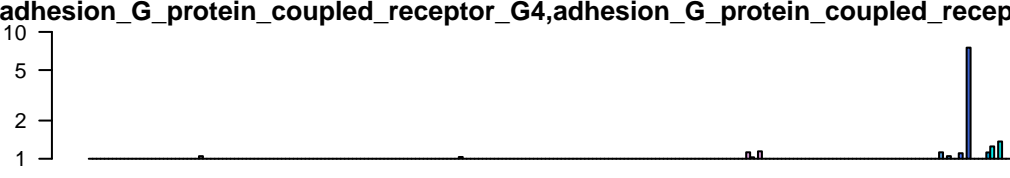
TrH2 OG_8020
TrH2_TrispH2_004376-RA
adhesion_G_protein_coupled_receptor_G4,adhesion_G_protein_coupled_receptor_G2



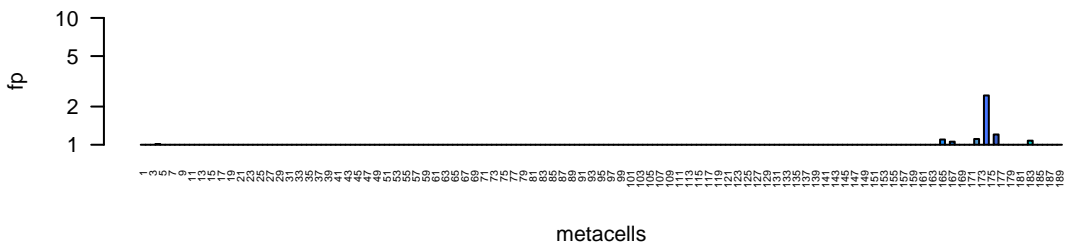
Hhon OG_8020
Hhon_g07984.t1
adhesion_G_protein_coupled_receptor_G4,adhesion_G_protein_coupled_receptor_G2



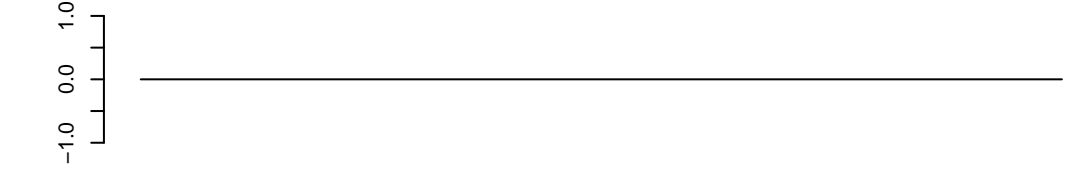
HoiH23 OG_8020
HoiH23_PIH23_006151-RA
adhesion_G_protein_coupled_receptor_G4,adhesion_G_protein_coupled_receptor_G2



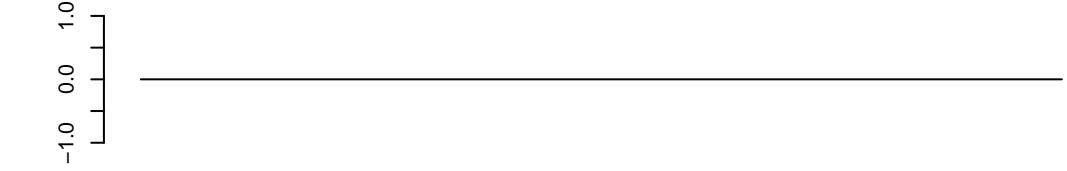
Tadh OG_8095
Tadh_TriadT57627



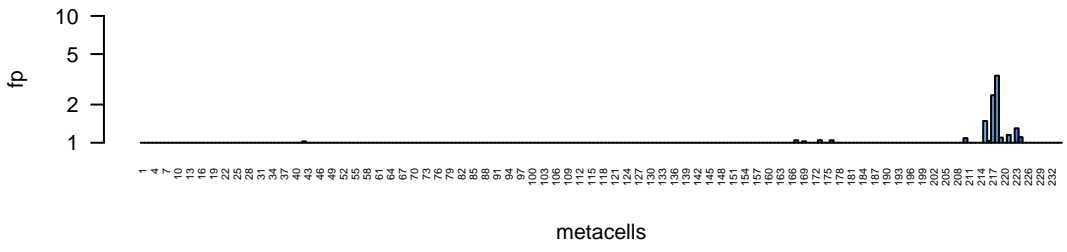
TrH2 | no data



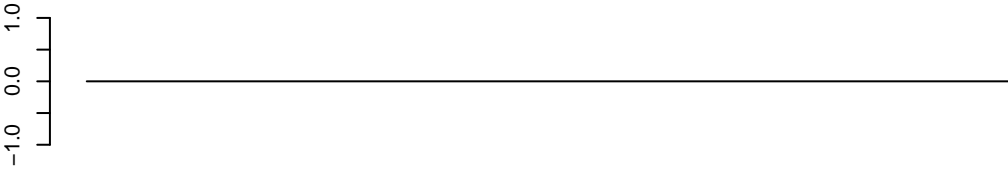
Hhon | no data



HoiH23 OG_8095
HoiH23_PIH23_007148-RA



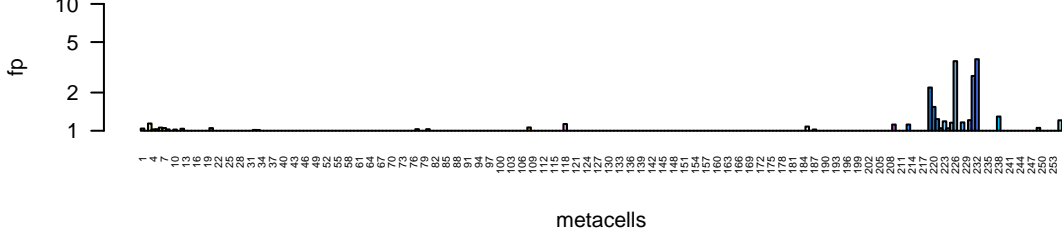
adrenoceptor_beta_2,histamine_receptor_H2
Tadh | no data



adrenoceptor_beta_2,histamine_receptor_H2
TrH2 | no data

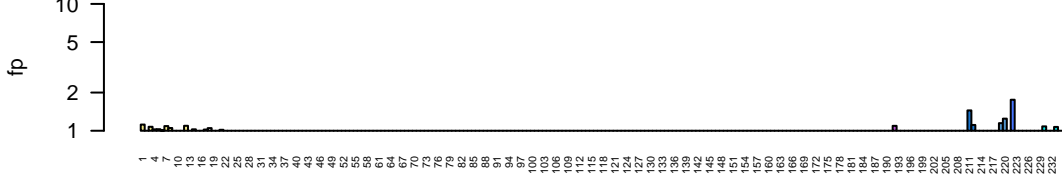


Hhon OG_8098
Hhon_g03438.t1
adrenoceptor_beta_2,histamine_receptor_H2



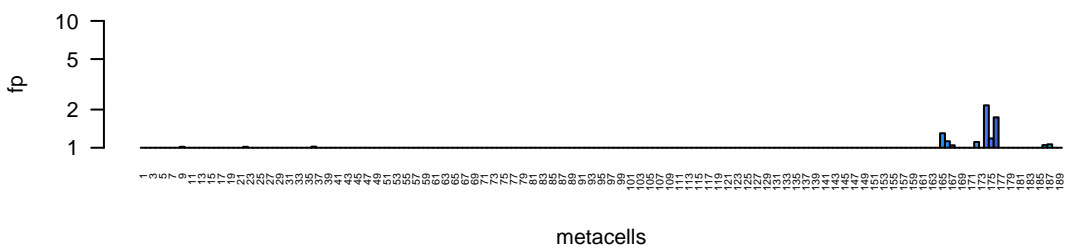
metacells

HoiH23 OG_8098
HoiH23_PIH23_007155-RA
adrenoceptor_beta_2,histamine_receptor_H2

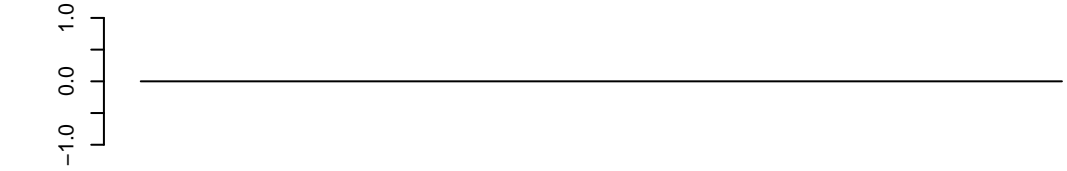


metacells

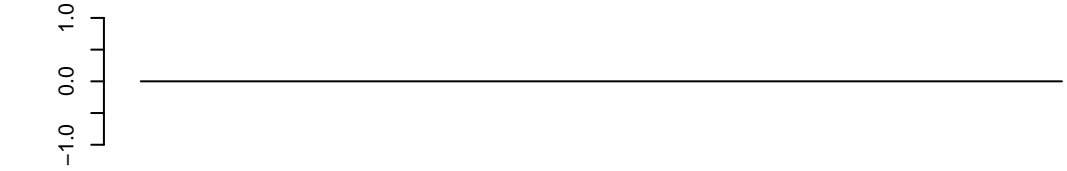
Tadh OG_8191
Tadh_TriadT60875



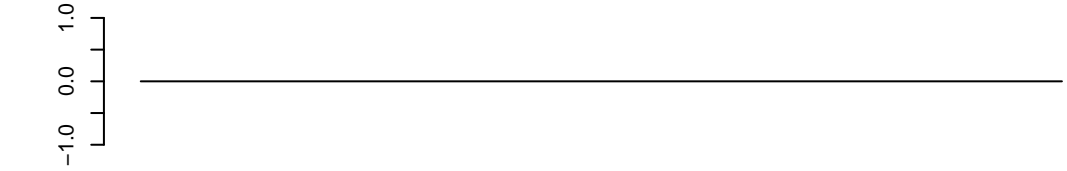
TrH2 | no data

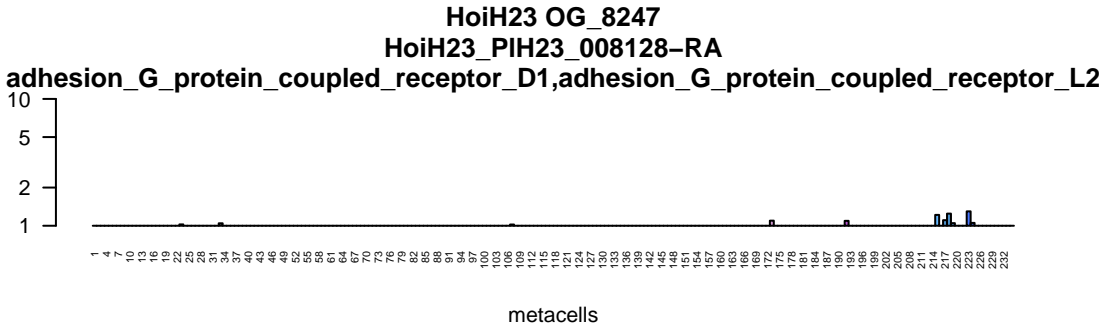
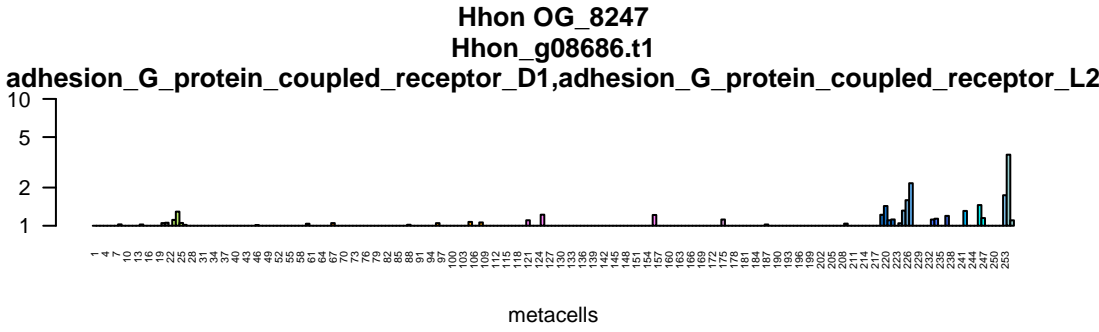
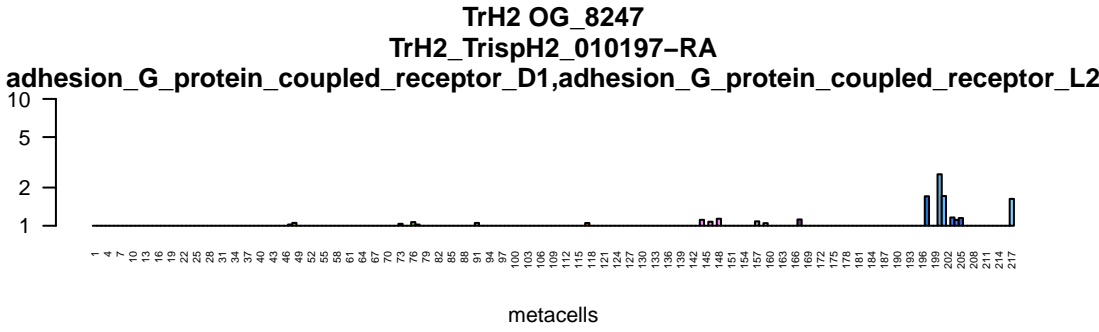
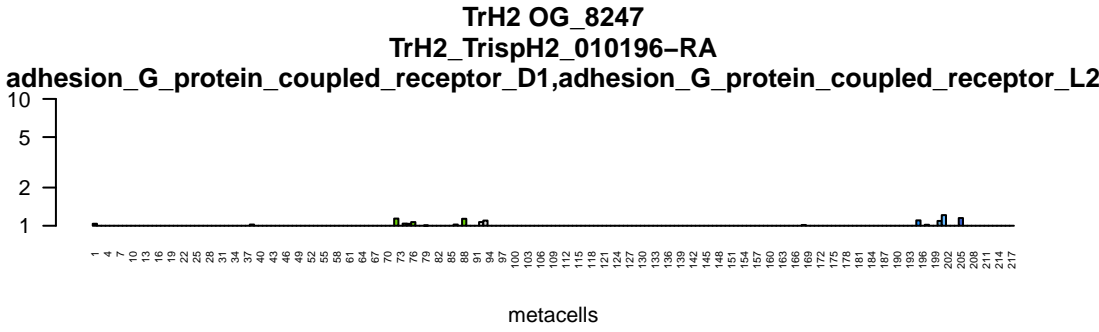
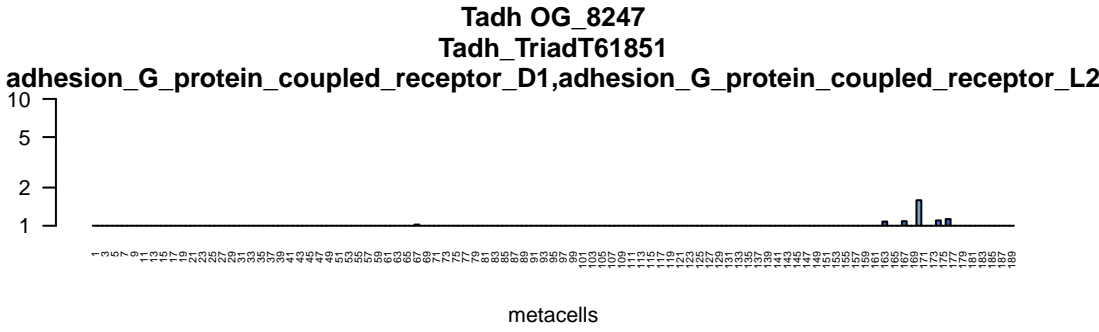
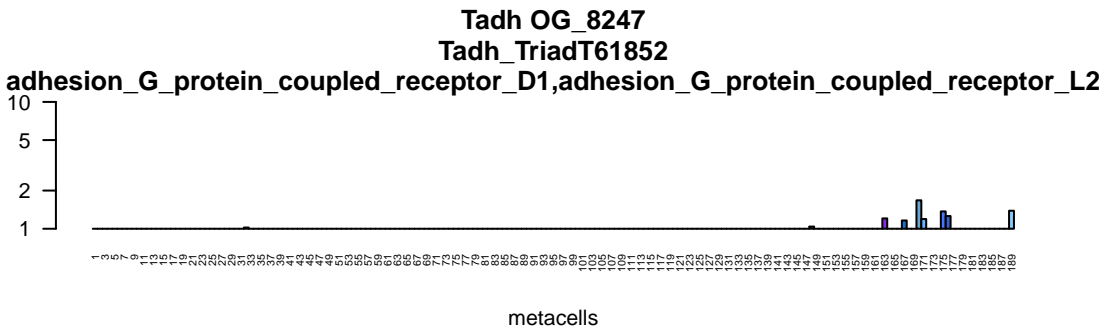
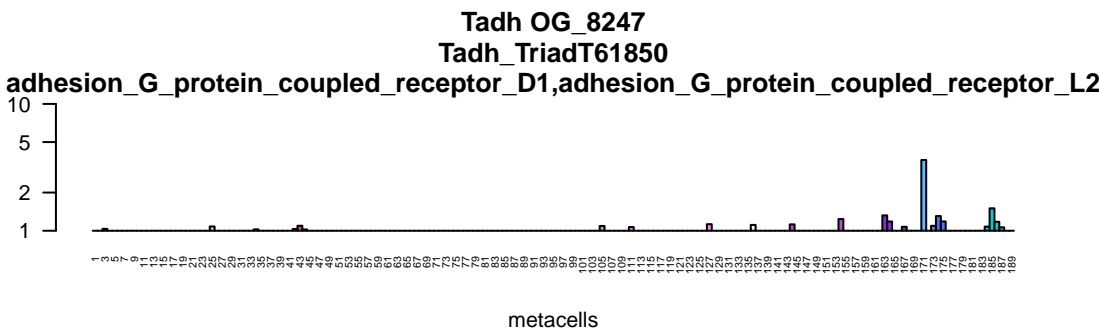


Hhon | no data



HoiH23 | no data

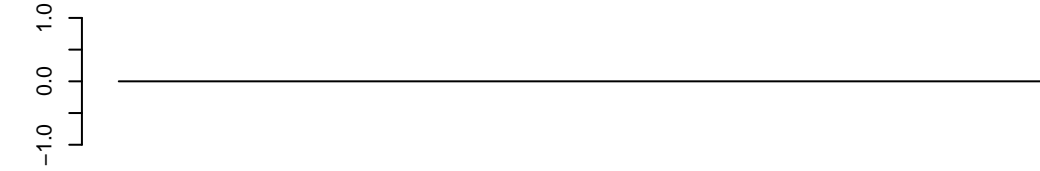




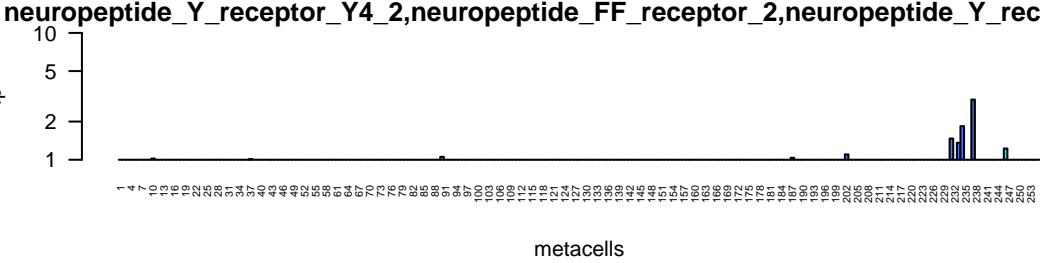
neuropeptide_Y_receptor_Y4_2,neuropeptide_FF_receptor_2,neuropeptide_Y_receptor_\
Tadh | no data



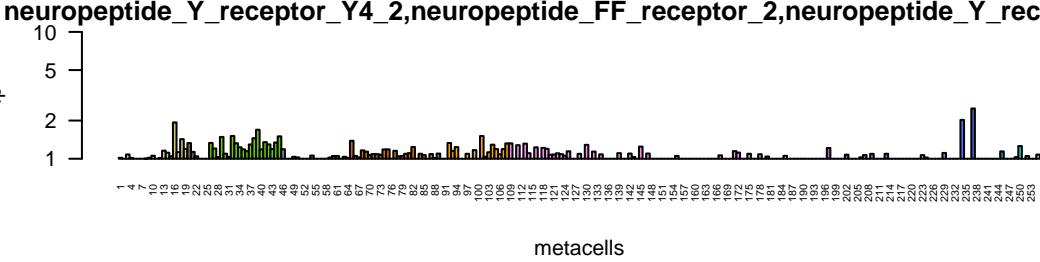
neuropeptide_Y_receptor_Y4_2,neuropeptide_FF_receptor_2,neuropeptide_Y_receptor_\
TrH2 | no data



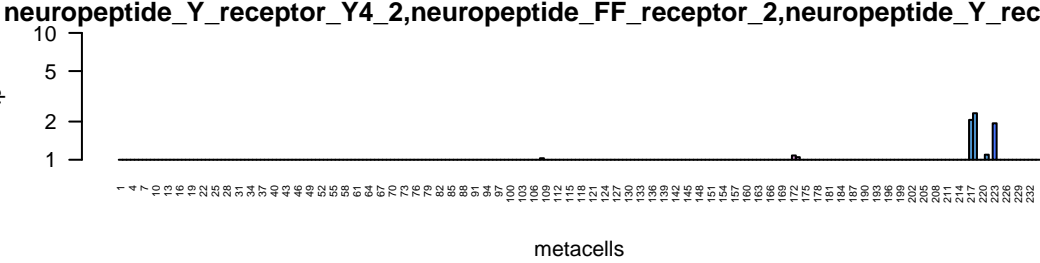
Hhon OG_8289
Hhon_g07093.t1
neuropeptide_Y_receptor_Y4_2,neuropeptide_FF_receptor_2,neuropeptide_Y_receptor_\
metacells

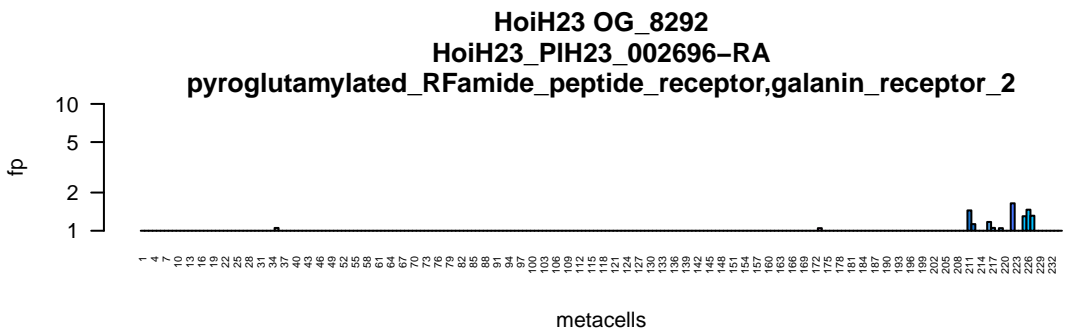
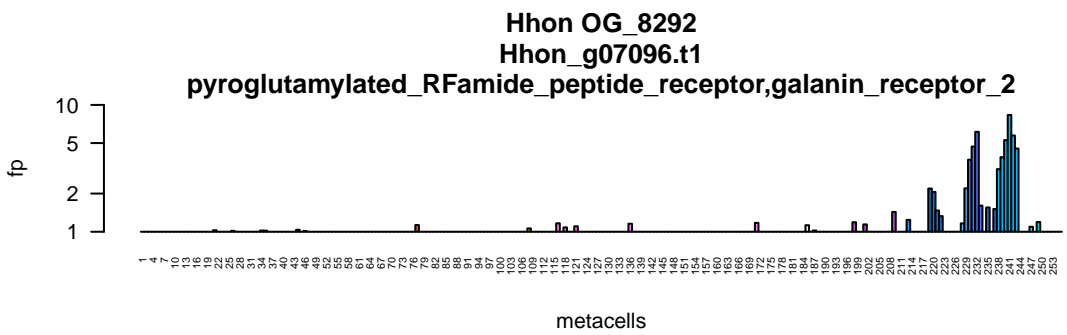
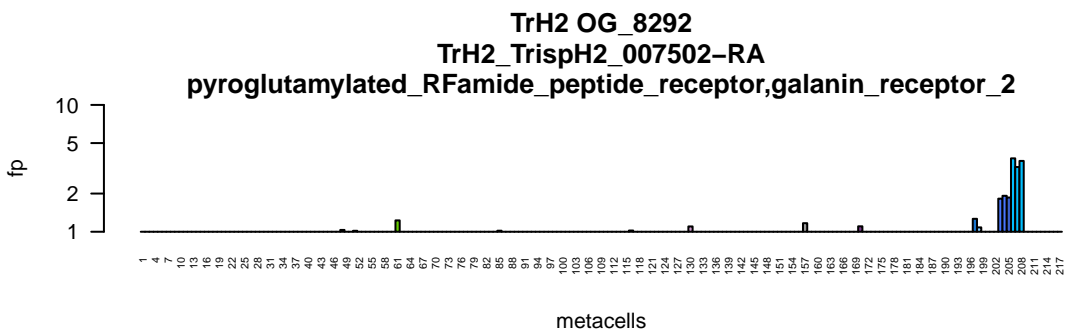
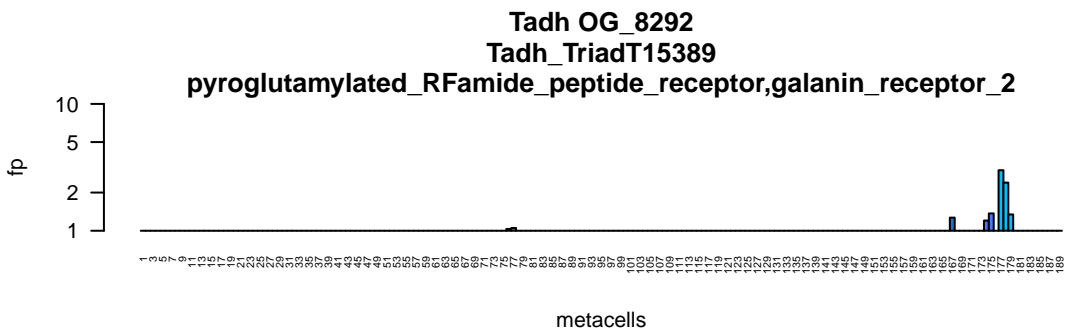


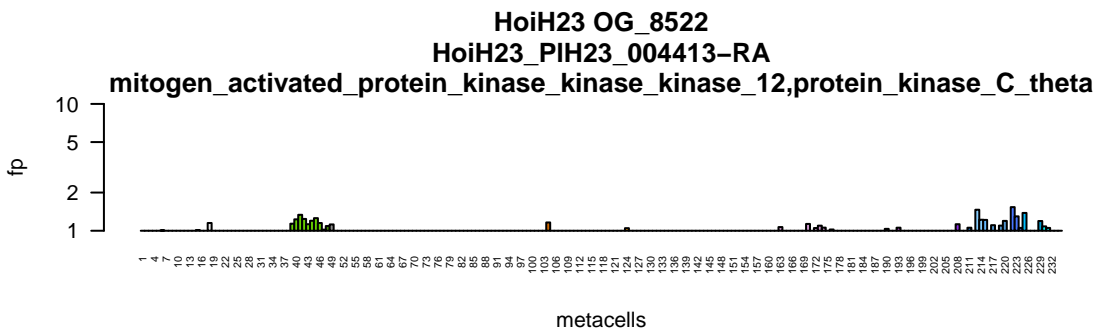
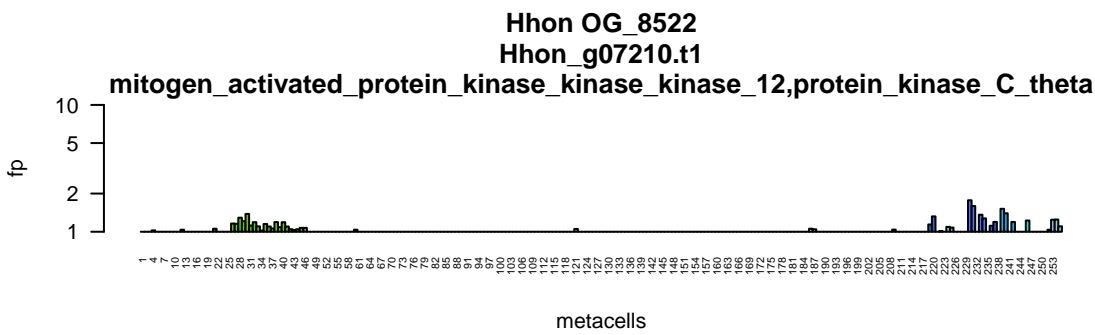
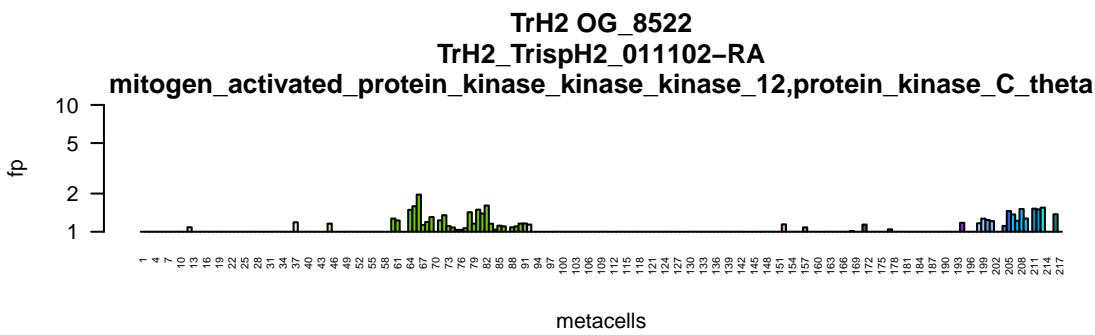
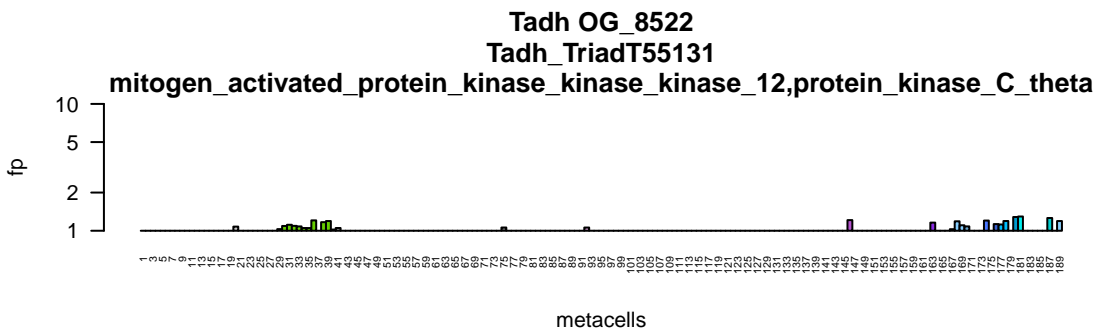
Hhon OG_8289
Hhon_g11435.t1
neuropeptide_Y_receptor_Y4_2,neuropeptide_FF_receptor_2,neuropeptide_Y_receptor_\
metacells



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



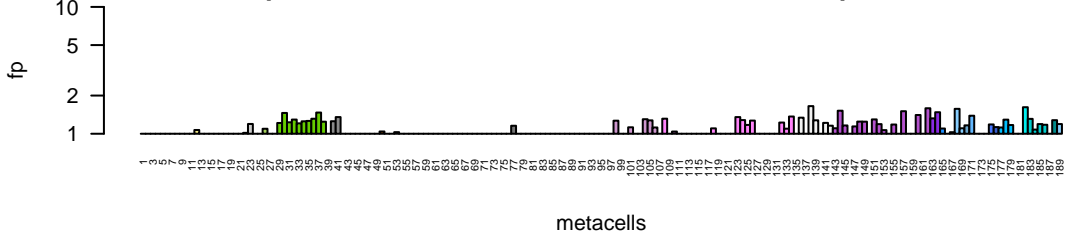




Tadh OG_8598

Tadh_TriadT51564

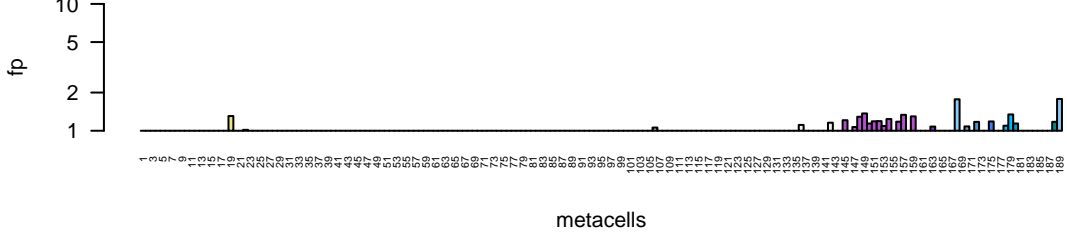
interleukin_1_receptor_associated_kinase_1,interleukin_1_receptor_associated_kinase_1



Tadh OG_8598

Tadh_TriadT51565

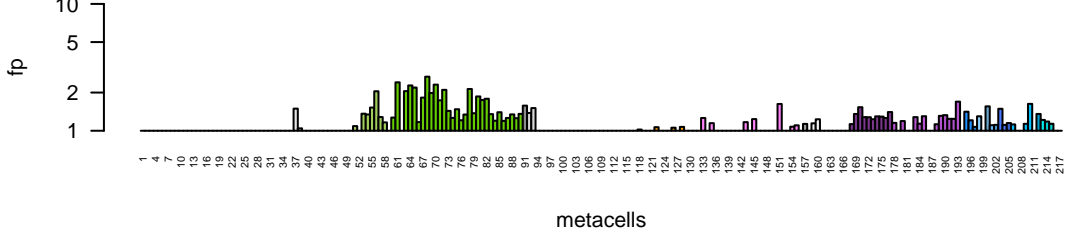
interleukin_1_receptor_associated_kinase_1,interleukin_1_receptor_associated_kinase_1



TrH2 OG_8598

TrH2_TrispH2_009241-RA

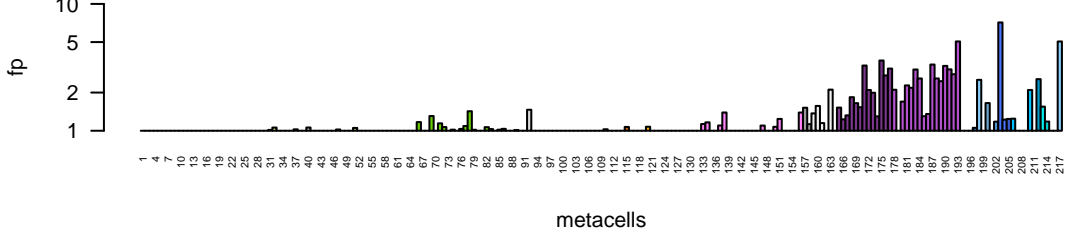
interleukin_1_receptor_associated_kinase_1,interleukin_1_receptor_associated_kinase_1



TrH2 OG_8598

TrH2_TrispH2_009242-RA

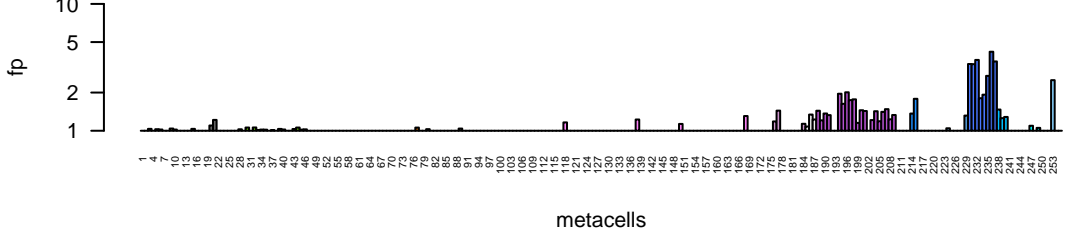
interleukin_1_receptor_associated_kinase_1,interleukin_1_receptor_associated_kinase_1



Hhon OG_8598

Hhon_g02060.t1

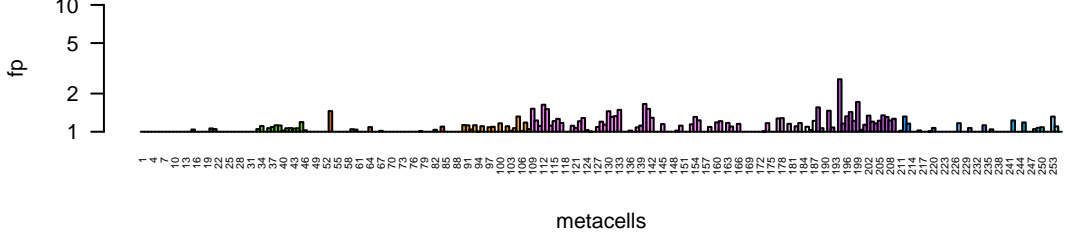
interleukin_1_receptor_associated_kinase_1,interleukin_1_receptor_associated_kinase_1



Hhon OG_8598

Hhon_g02061.t1

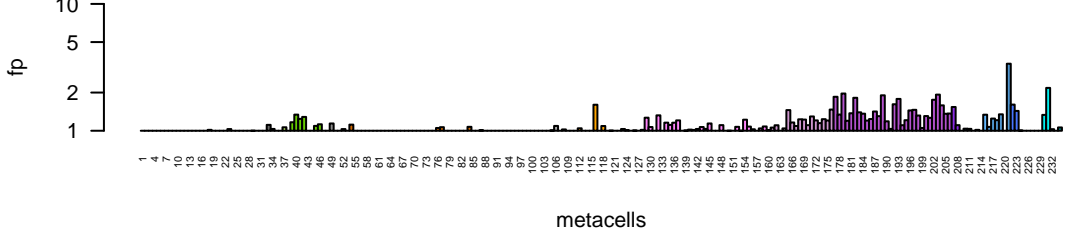
interleukin_1_receptor_associated_kinase_1,interleukin_1_receptor_associated_kinase_1

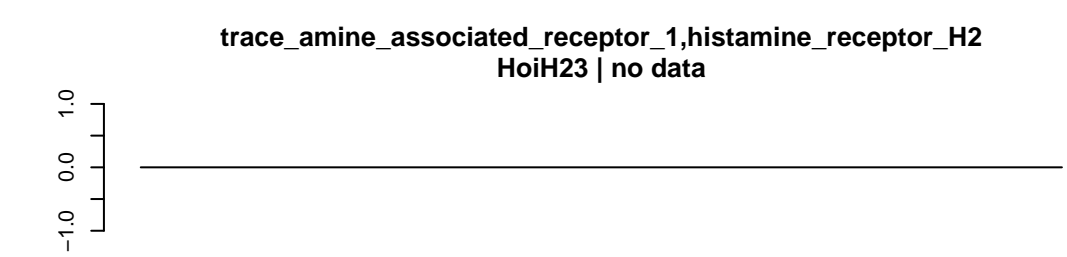
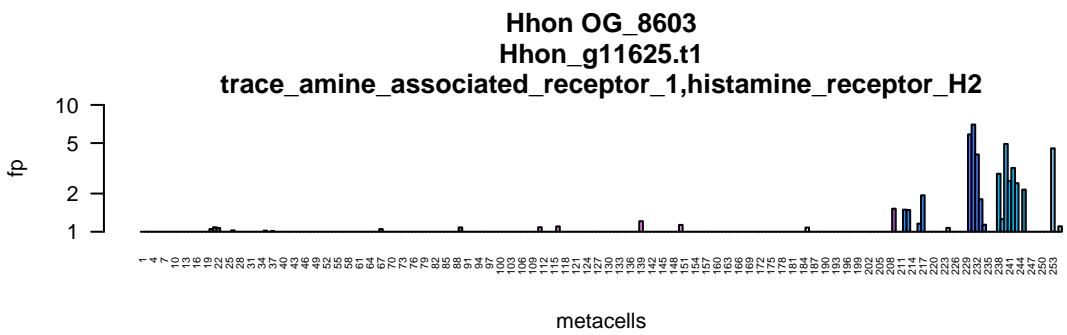
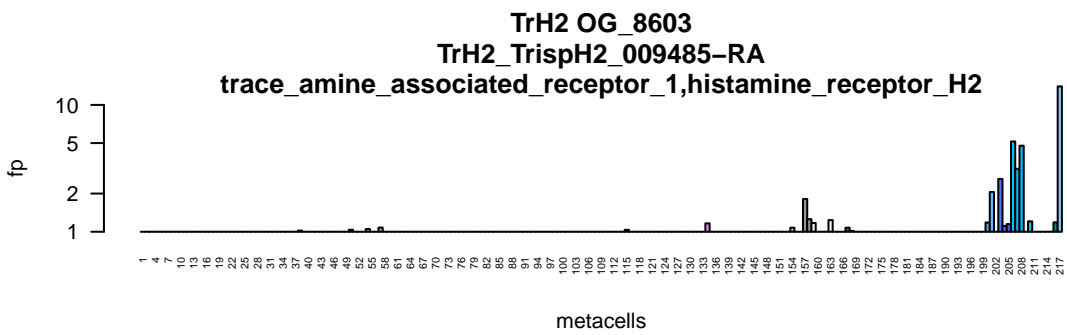
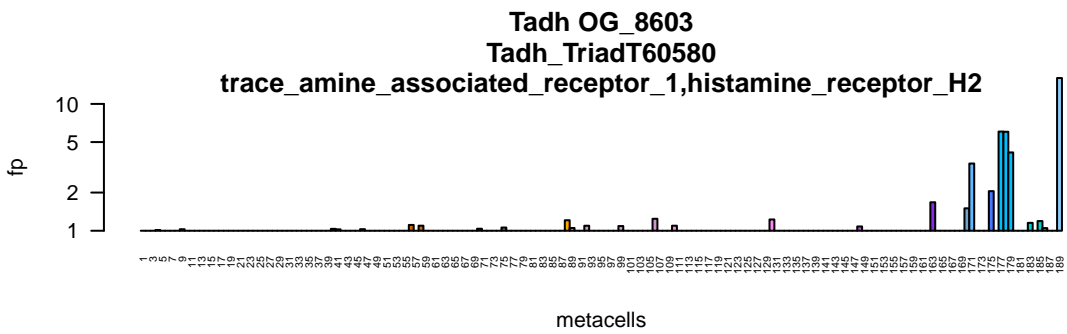


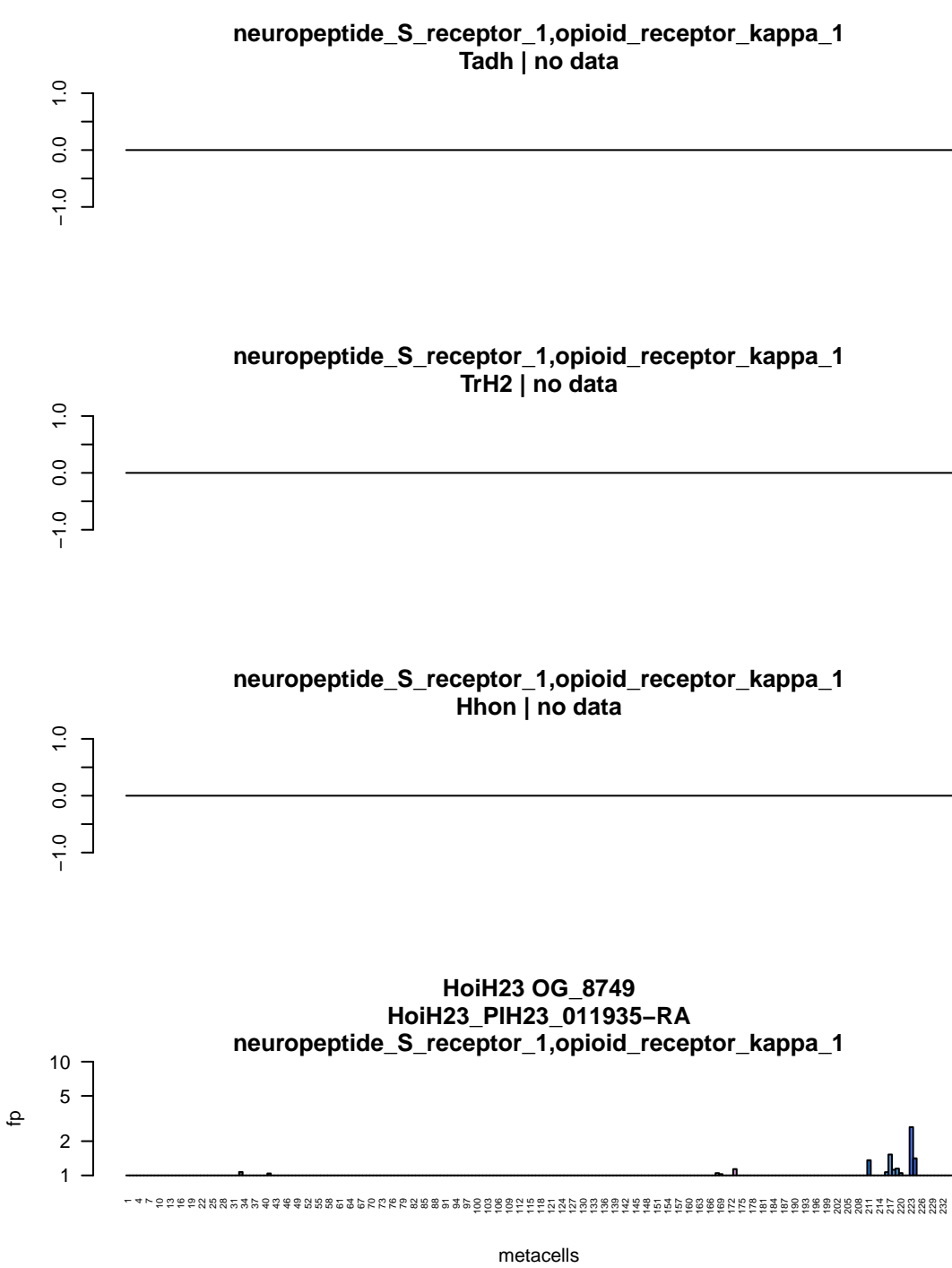
HoiH23 OG_8598

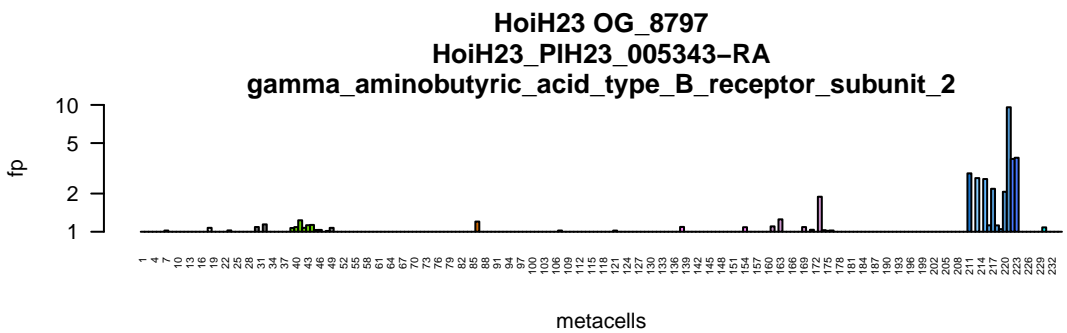
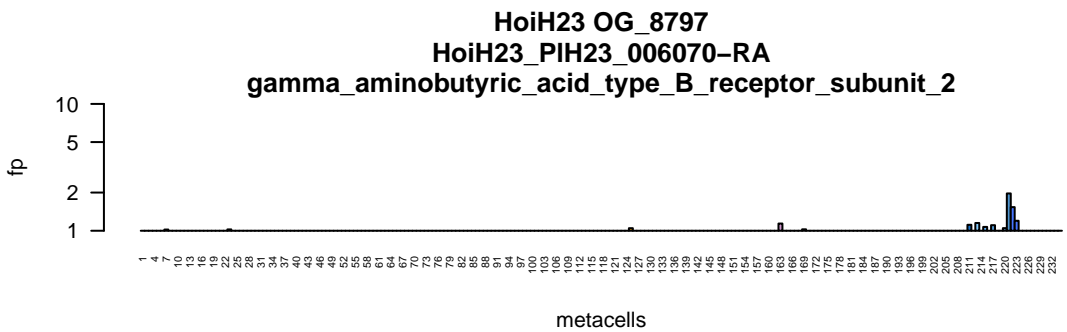
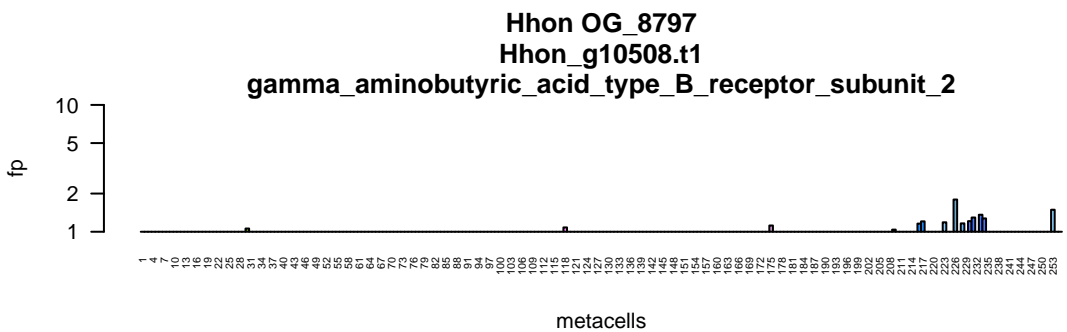
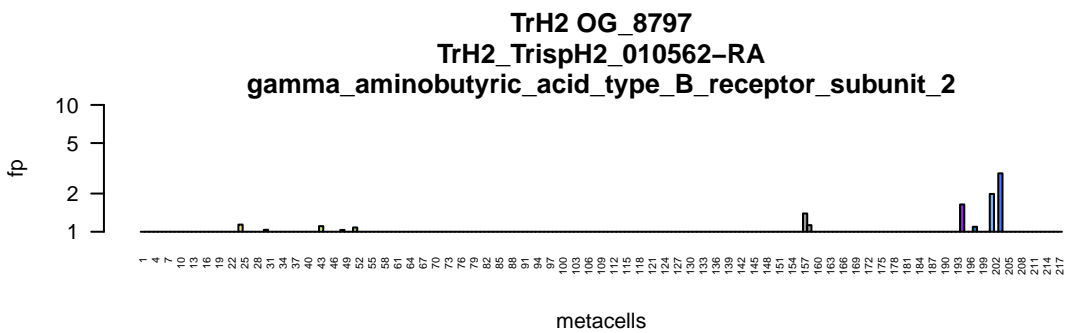
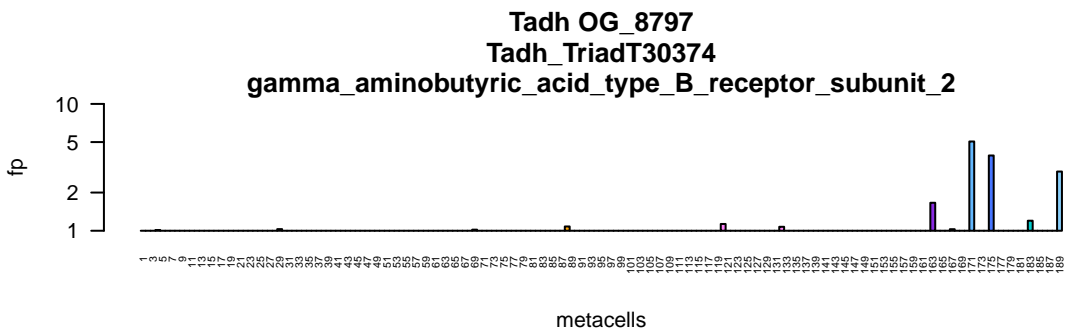
HoiH23_PIH23_010002-RA

interleukin_1_receptor_associated_kinase_1,interleukin_1_receptor_associated_kinase_1



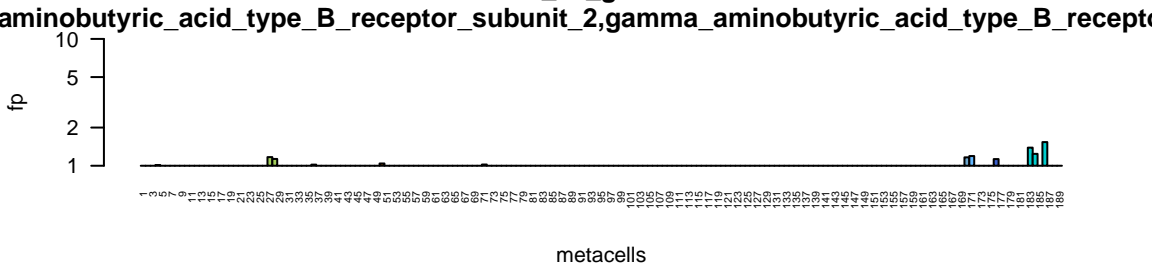






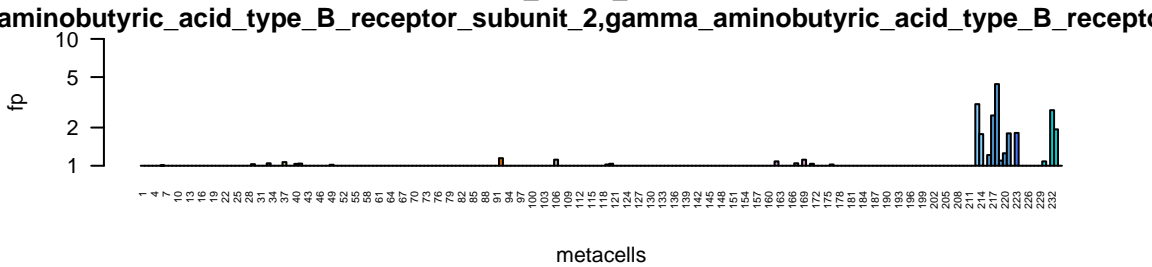
Tadh OG 8960

Tadh_wf_g11896.t1



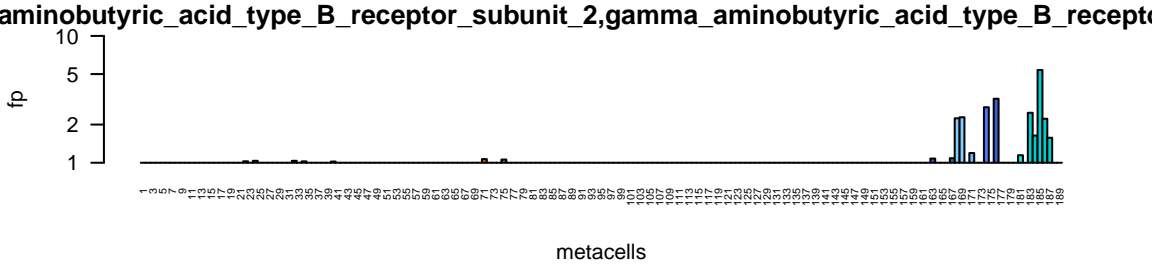
HoiH23 OG 8960

HoiH23_PIH23_000719-RA



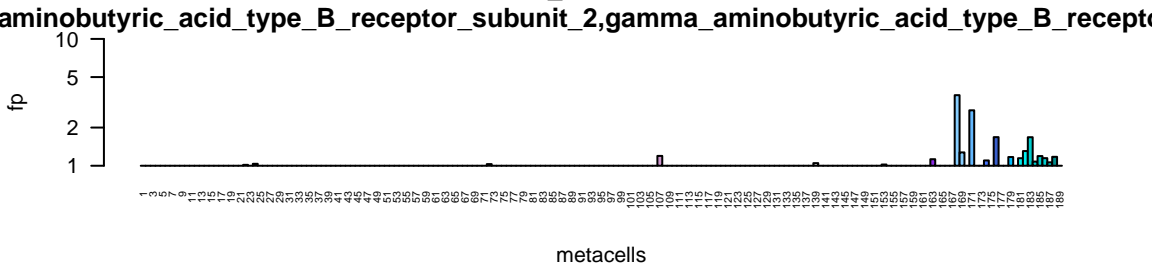
Tadh OG_8960

Tadh_TriadT52578



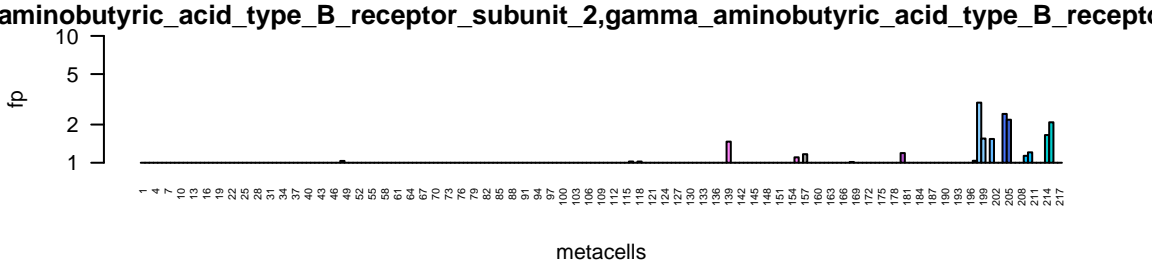
Tadh OG_8960

Tadh_TriadT52579



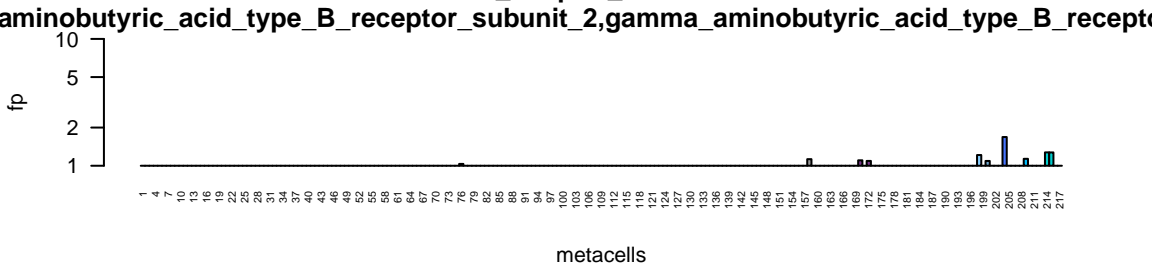
TrH2 OG_8960

TrH2_TrispH2_012100-RA



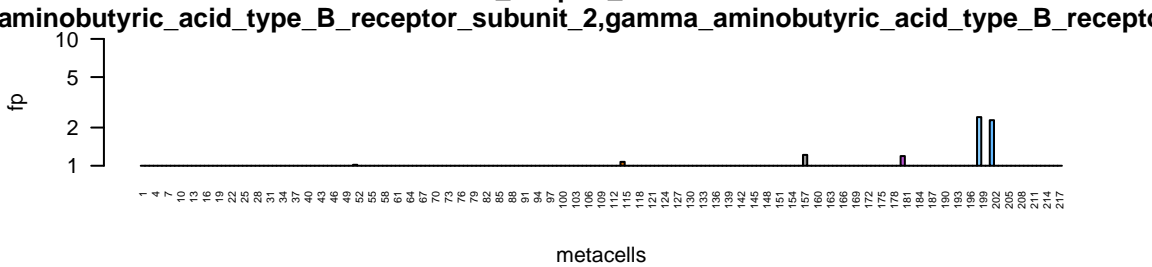
TrH2 OG_8960

TrH2_TrispH2_000234-RA



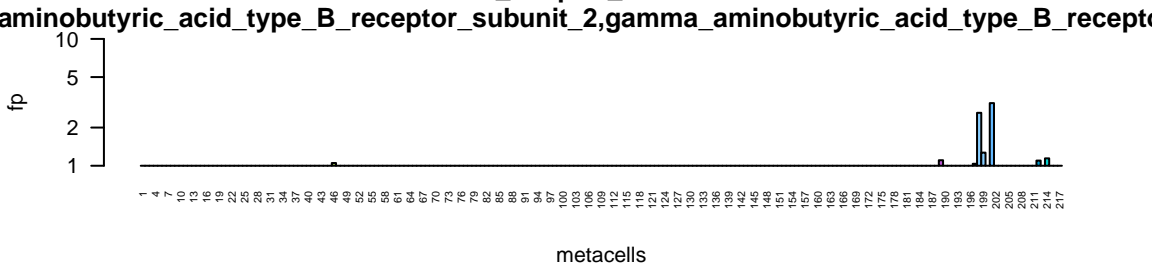
TrH2 OG_8960

TrH2_TrispH2_011964-RA



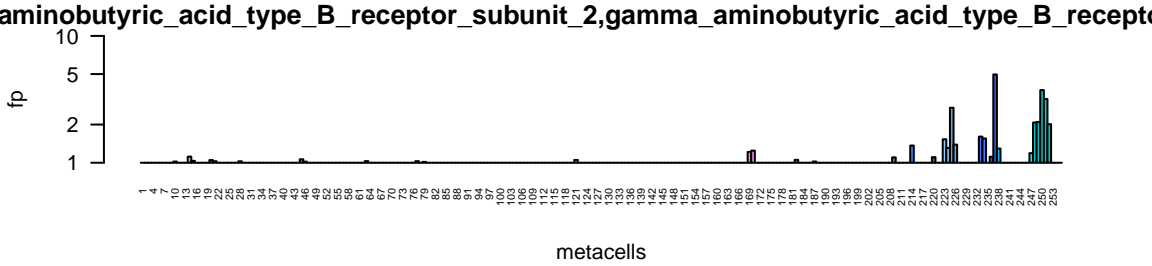
TrH2 OG_8960

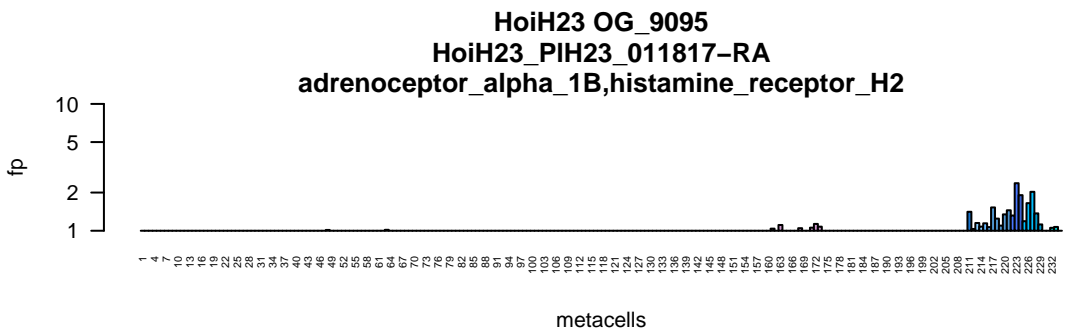
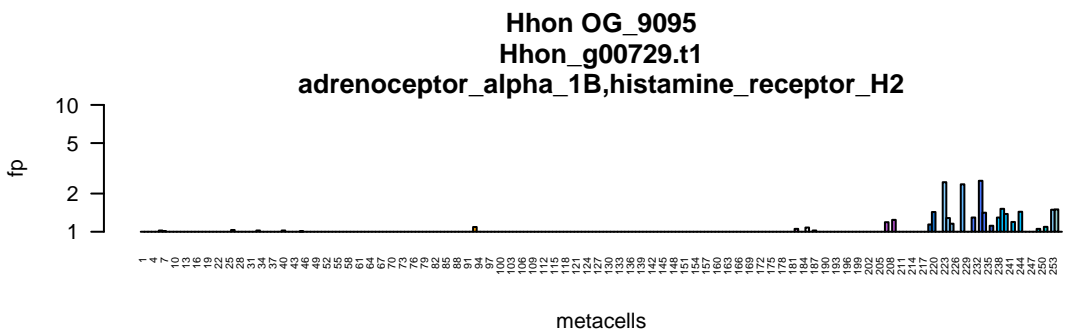
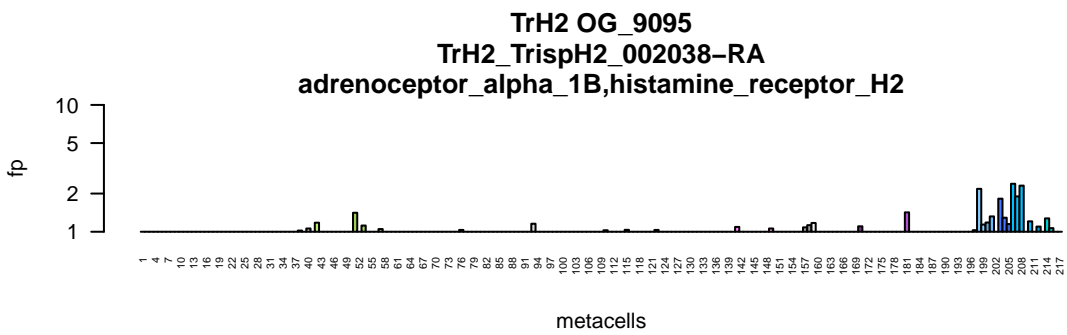
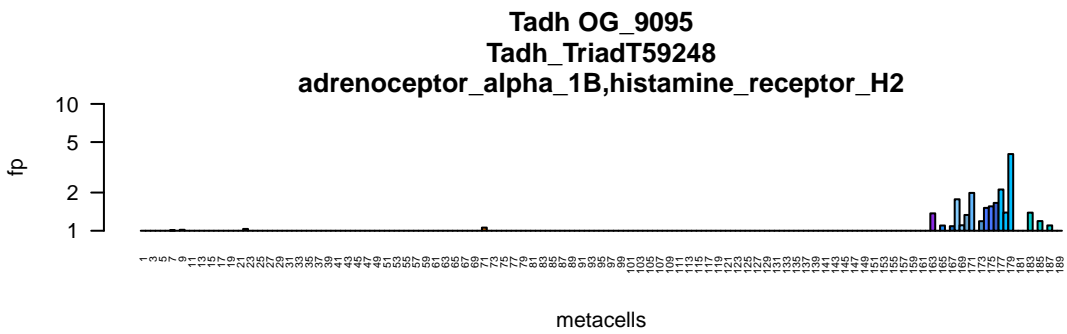
TrH2_TrispH2_011847-RA

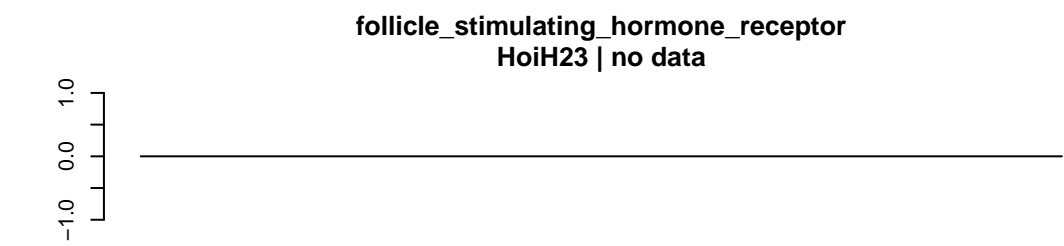
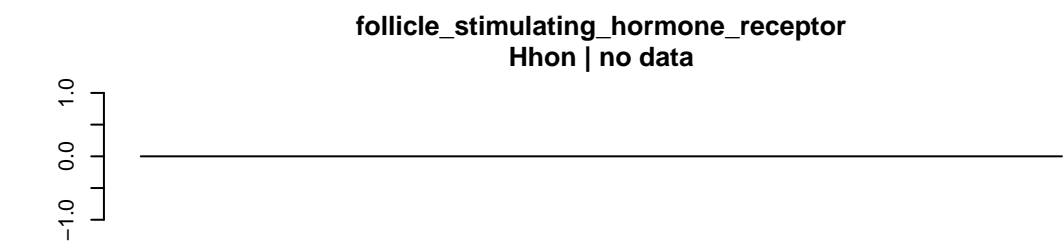
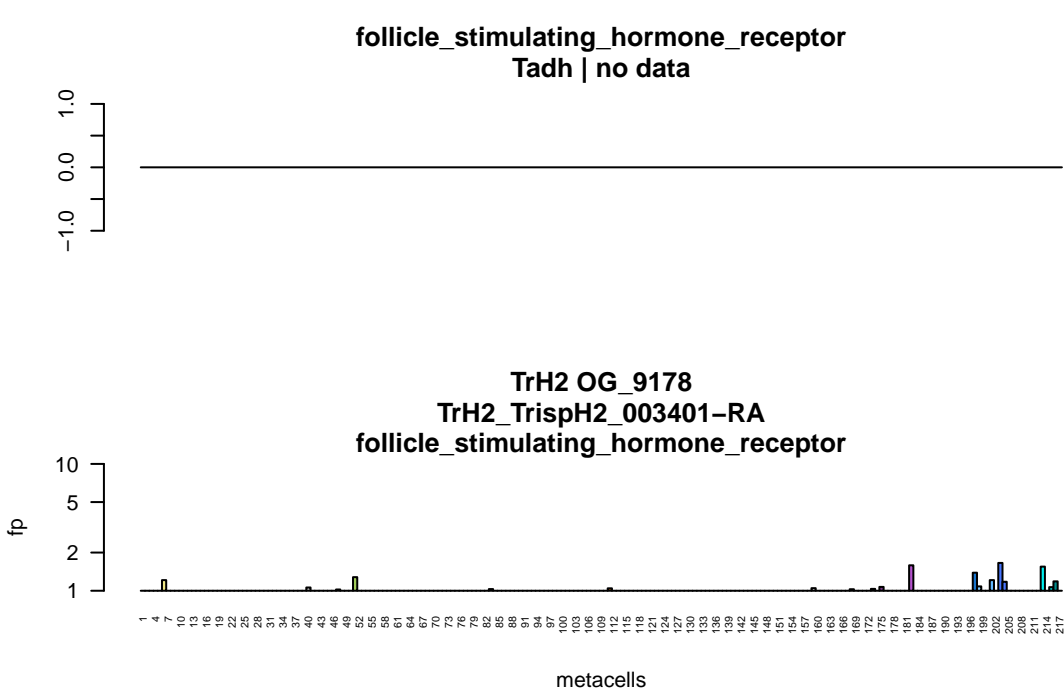


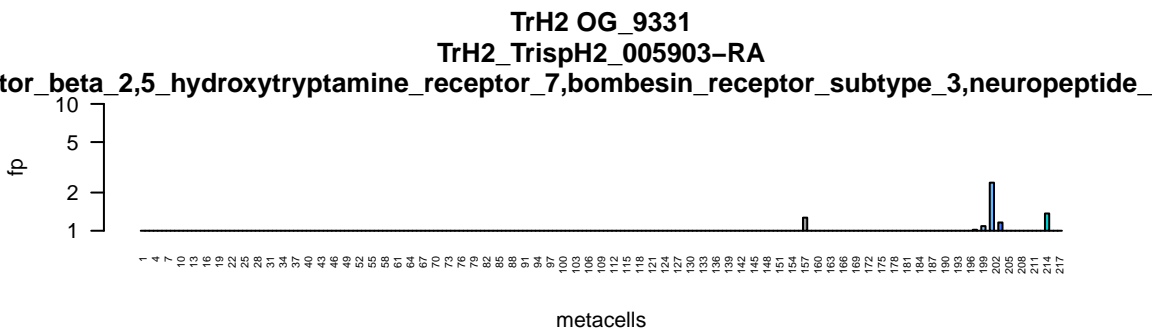
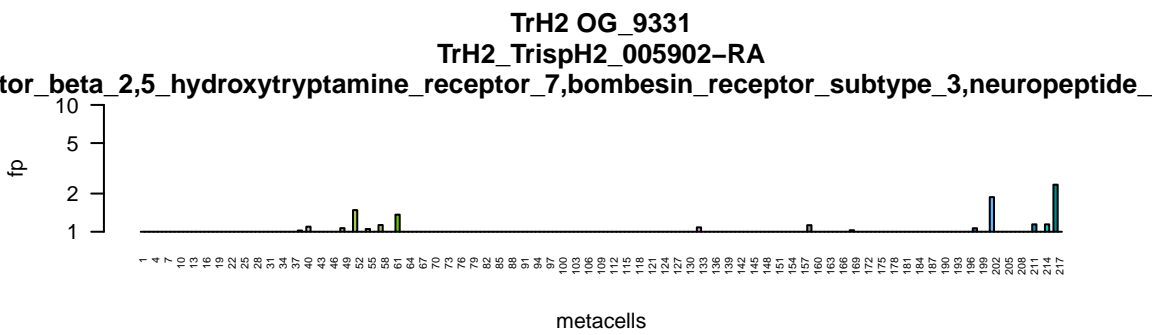
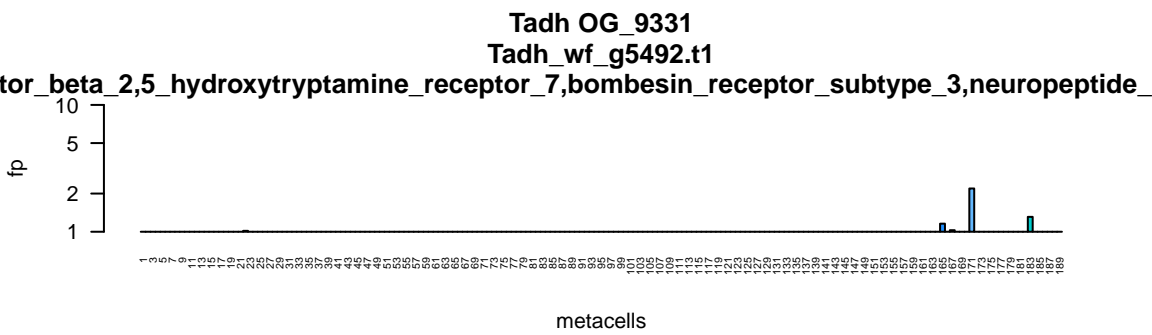
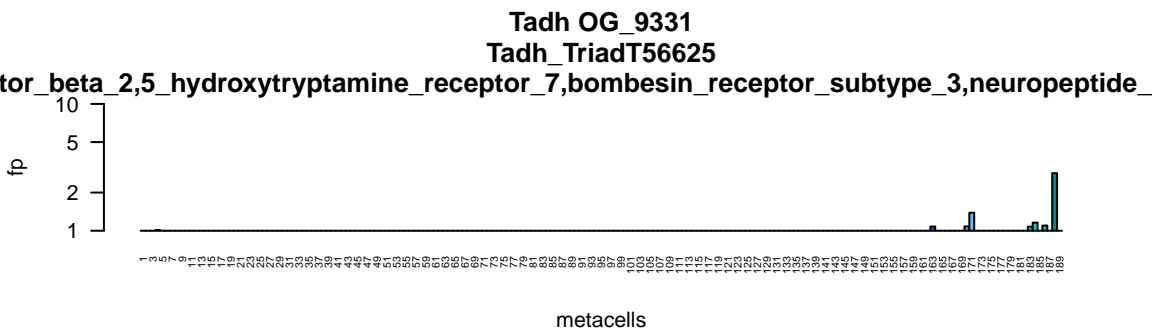
Hhon OG_8960

Hhon_g05296.t1





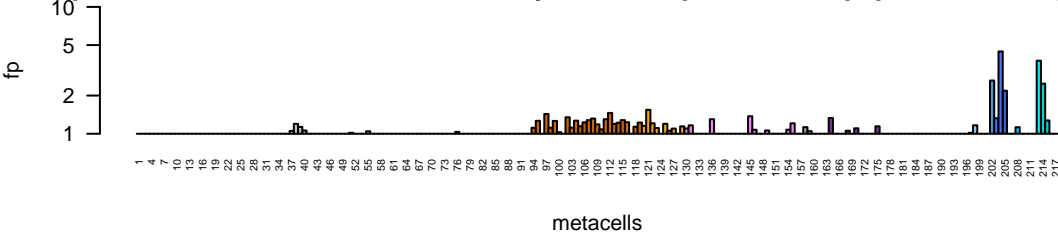




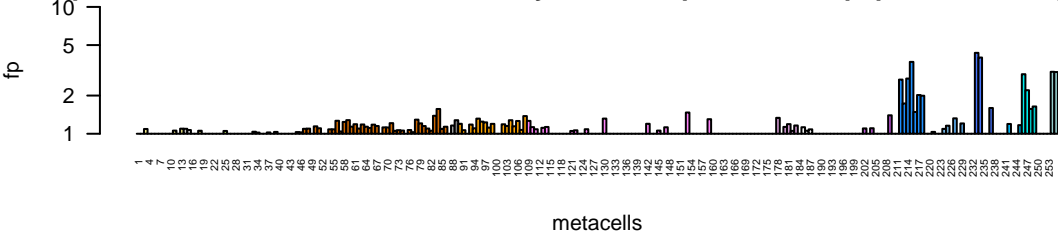
aprataxin_and_PNKP_like_factor,tachykinin_receptor_2,neuropeptide_FF_receptor_1
Tadh | no data



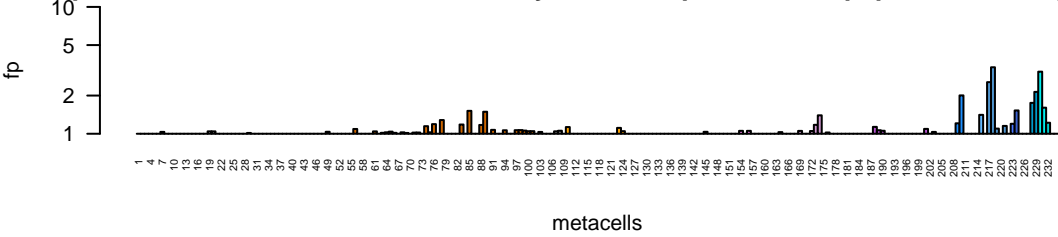
TrH2 OG_9368
TrH2_TrispH2_010619-RA
aprataxin_and_PNKP_like_factor,tachykinin_receptor_2,neuropeptide_FF_receptor_1

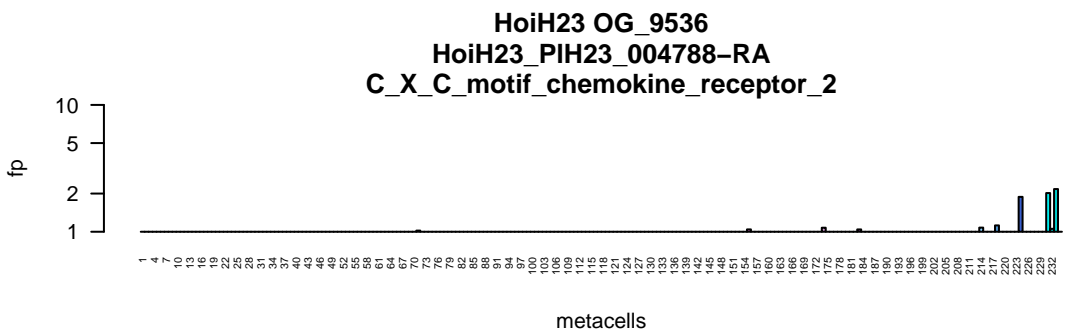
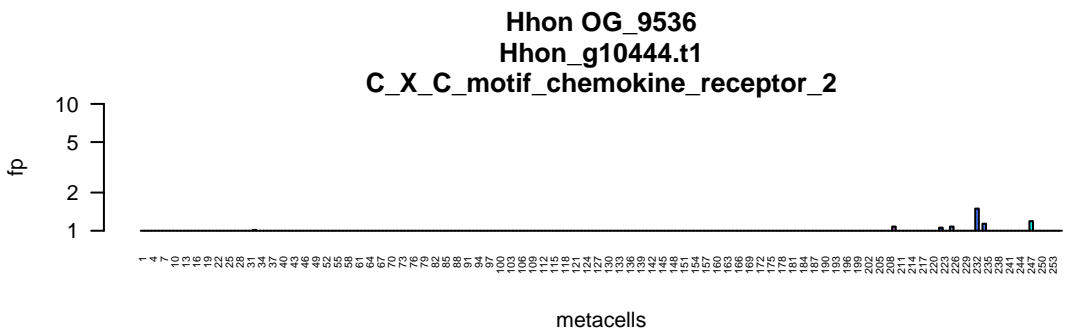
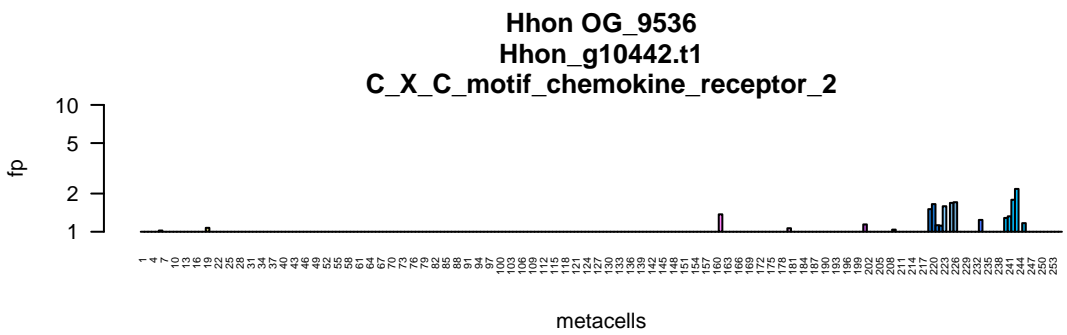
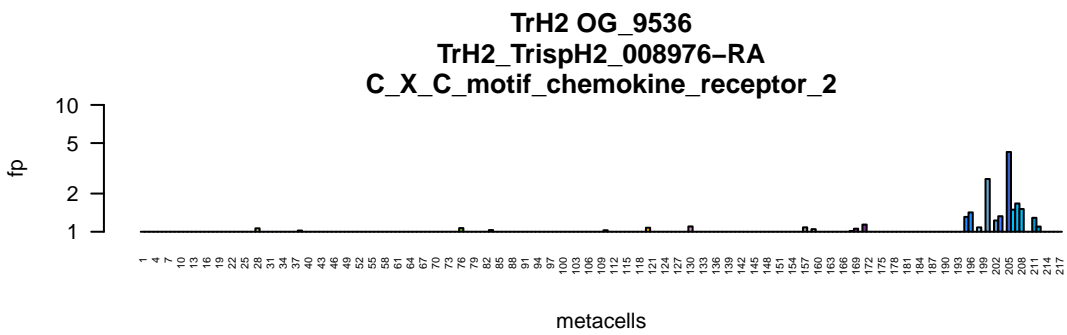
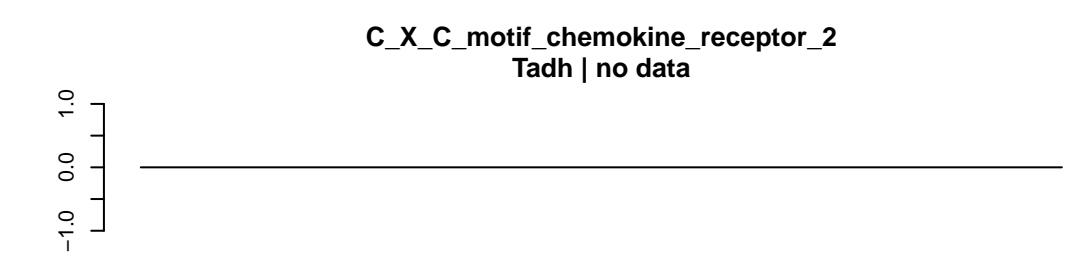


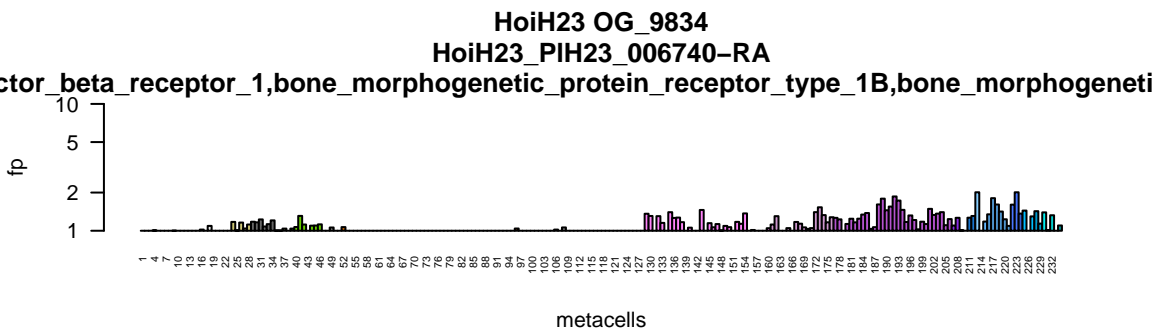
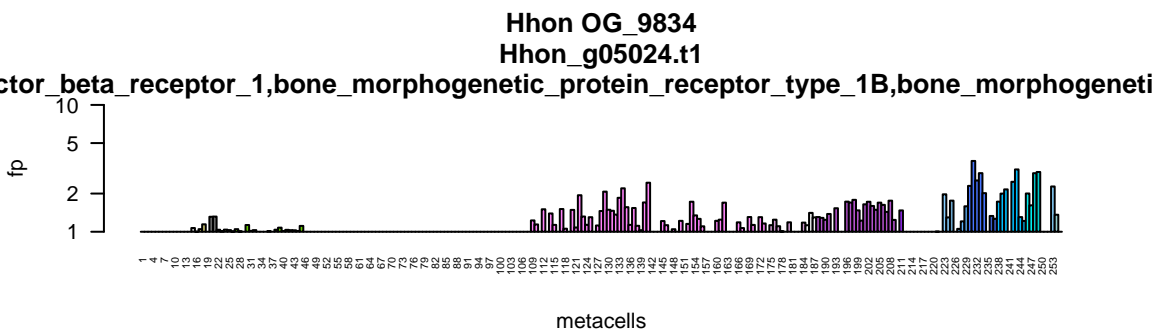
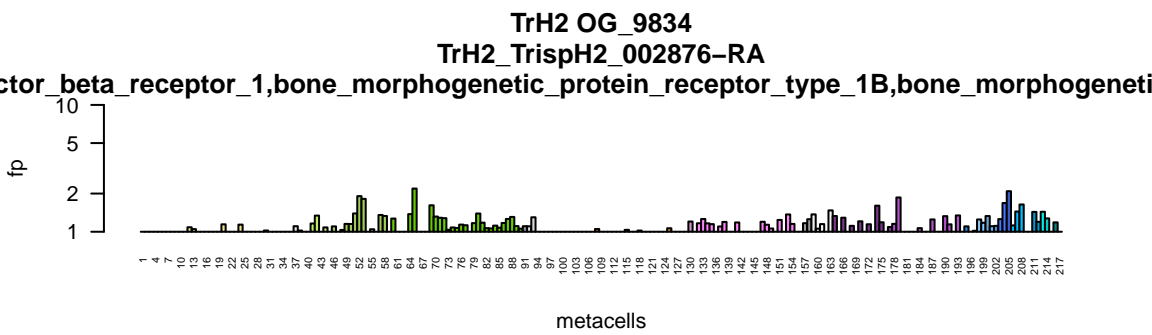
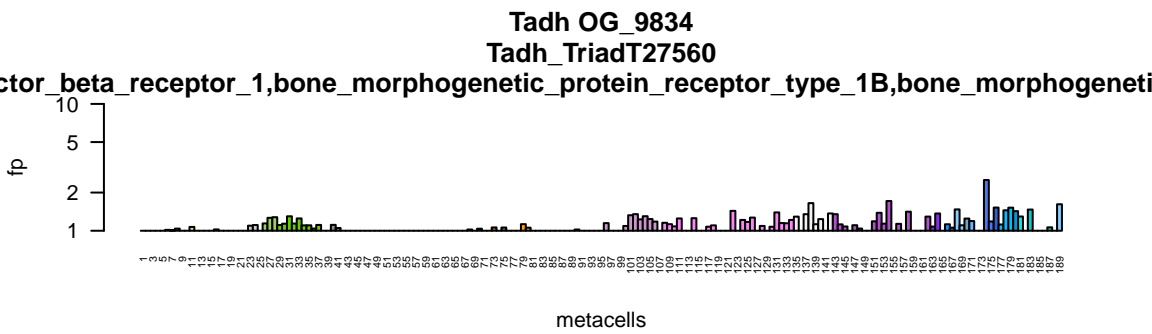
Hhon OG_9368
Hhon_g04575.t1
aprataxin_and_PNKP_like_factor,tachykinin_receptor_2,neuropeptide_FF_receptor_1



HoiH23 OG_9368
HoiH23_PIH23_003409-RA
aprataxin_and_PNKP_like_factor,tachykinin_receptor_2,neuropeptide_FF_receptor_1







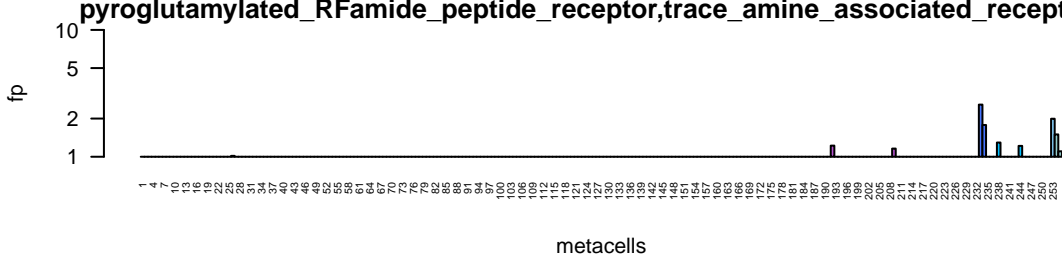
pyroglutamylated_RFamide_peptide_receptor,trace_amine_associated_receptor_2
Tadh | no data



pyroglutamylated_RFamide_peptide_receptor,trace_amine_associated_receptor_2
TrH2 | no data

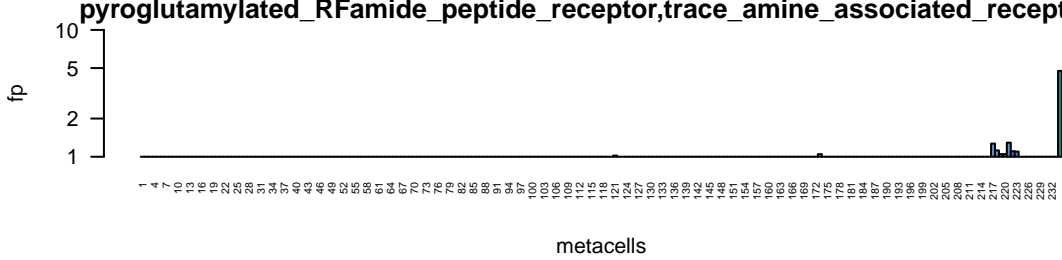


Hhon OG_10042
Hhon_g08855.t1
pyroglutamylated_RFamide_peptide_receptor,trace_amine_associated_receptor_2



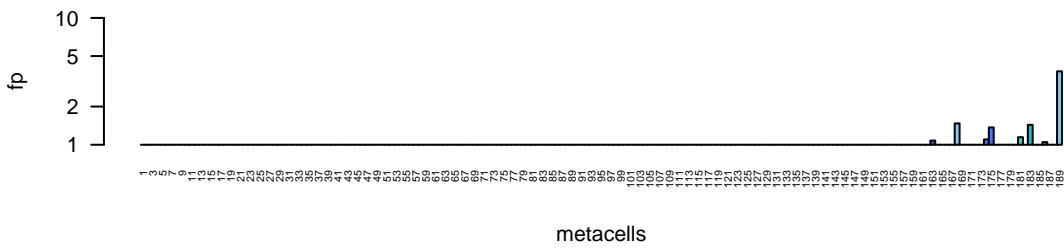
metacells

HoiH23 OG_10042
HoiH23_PIH23_007371-RA
pyroglutamylated_RFamide_peptide_receptor,trace_amine_associated_receptor_2

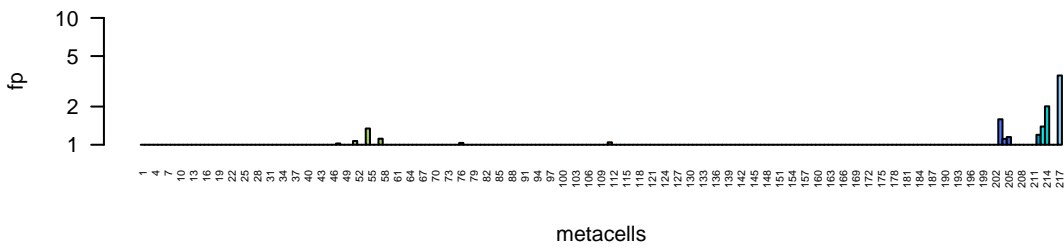


metacells

Tadh OG_10104
Tadh_TriadT60048



TrH2 OG_10104
TrH2_TrispH2_011300-RA



Hhon | no data



HoiH23 OG_10104
HoiH23_PIH23_006092-RA

