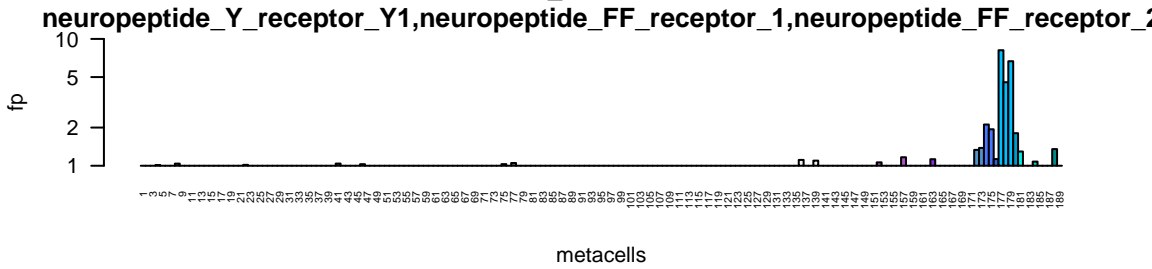


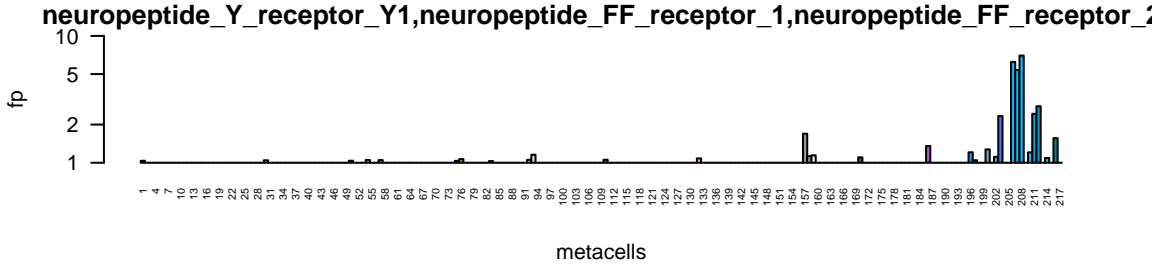
Tadh OG\_6978

Tadh\_TriadT32926



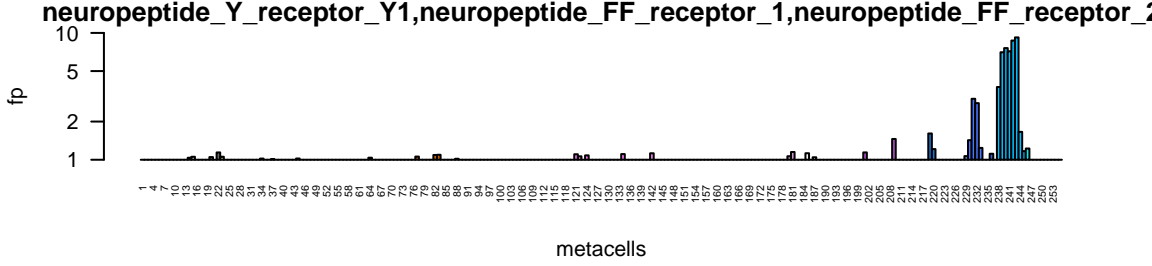
TrH2 OG\_6978

TrH2\_TrispH2\_009841-RA



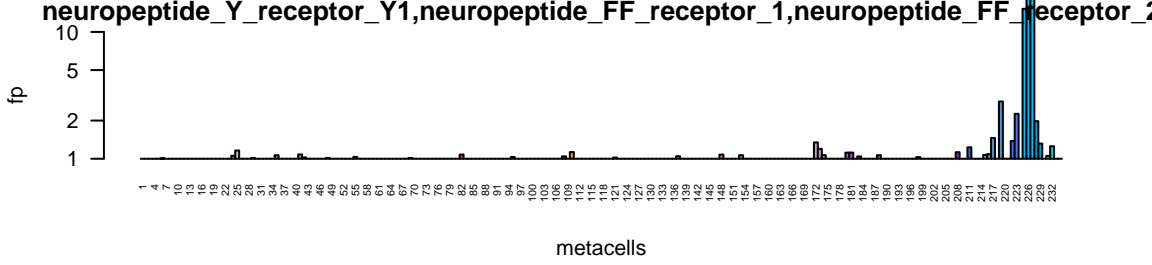
Hhon OG\_6978

Hhon\_g04425.t1

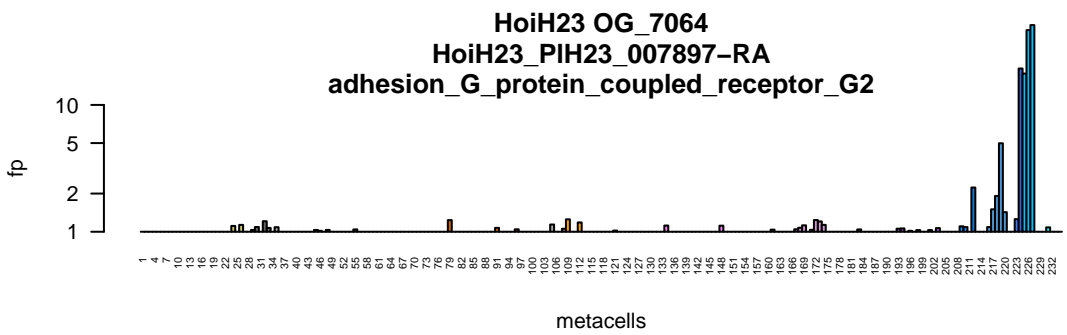
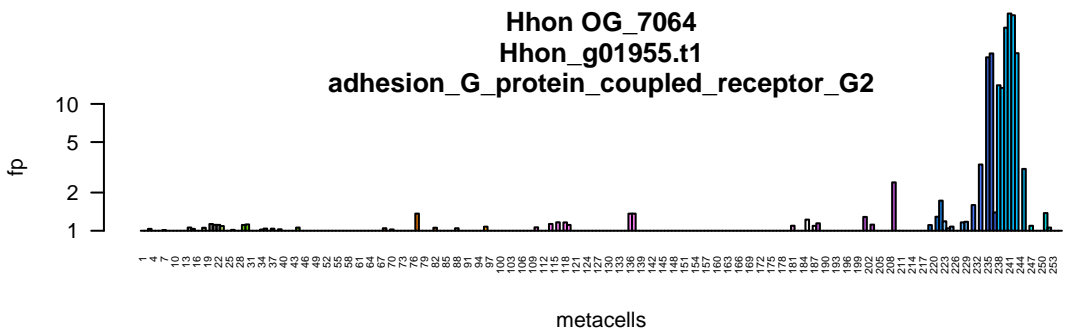
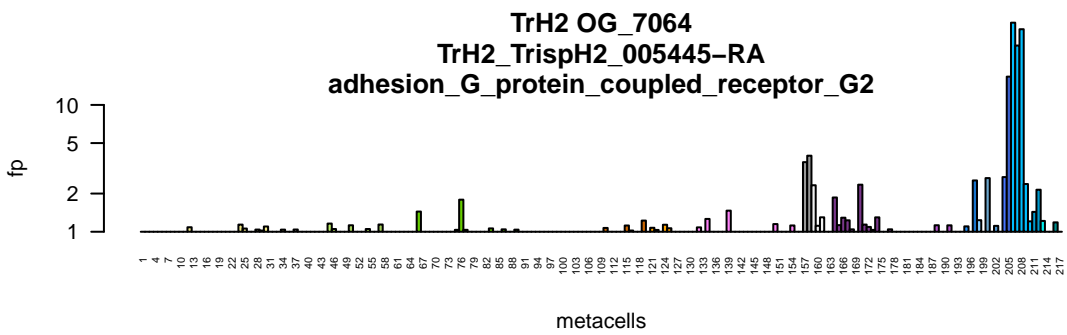
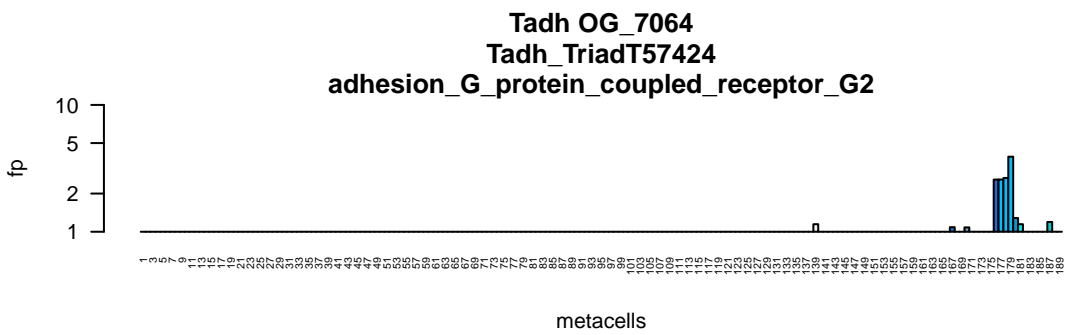
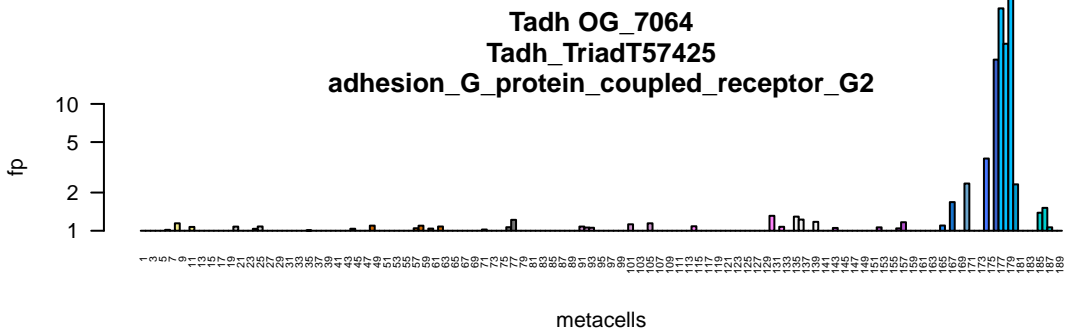


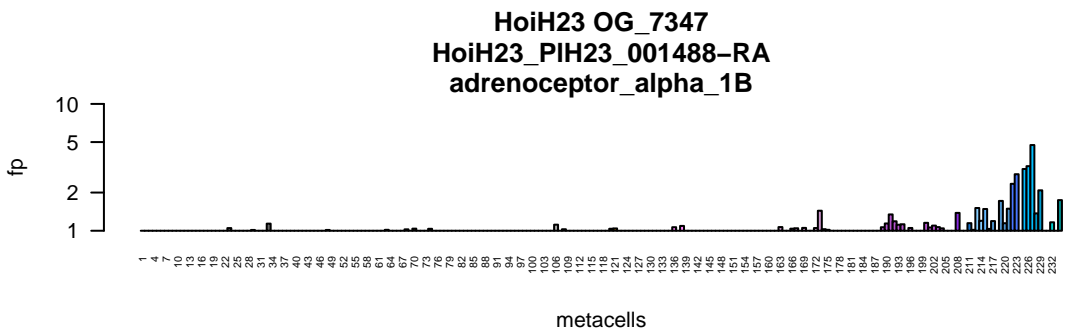
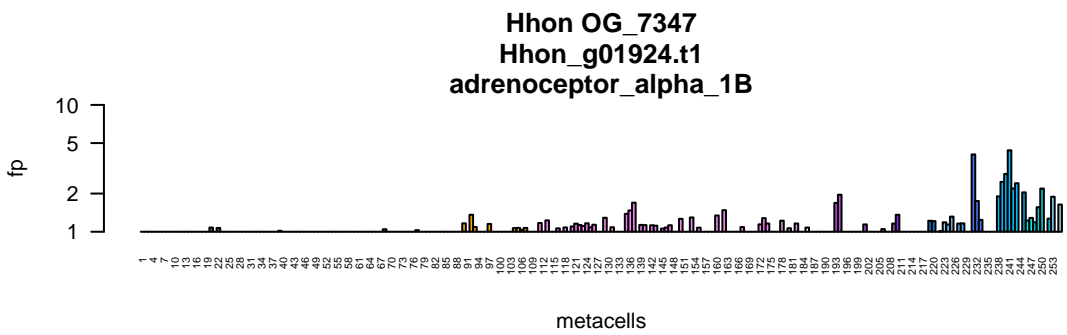
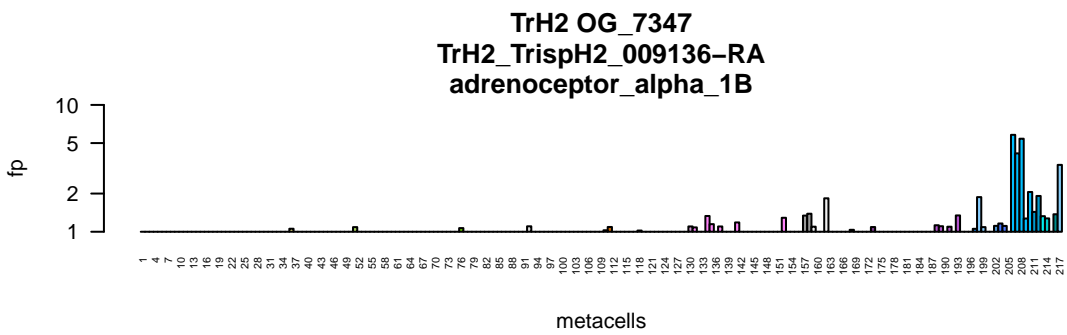
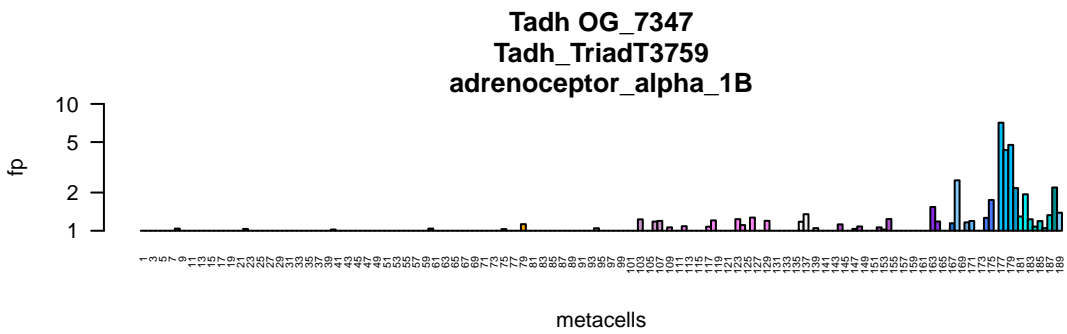
HoiH23 OG\_6978

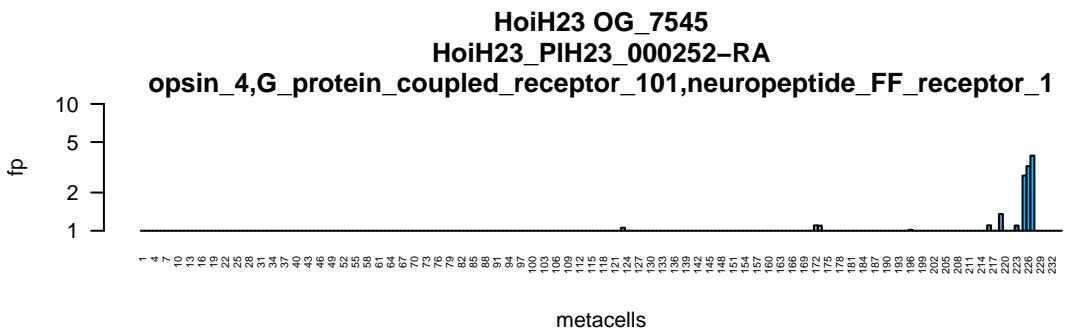
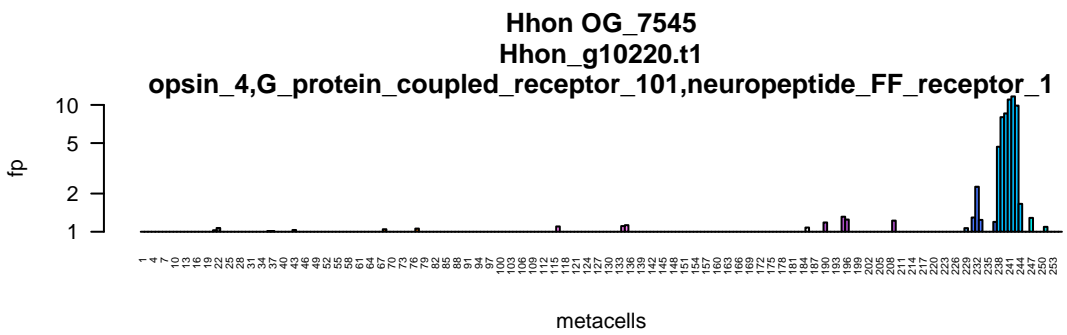
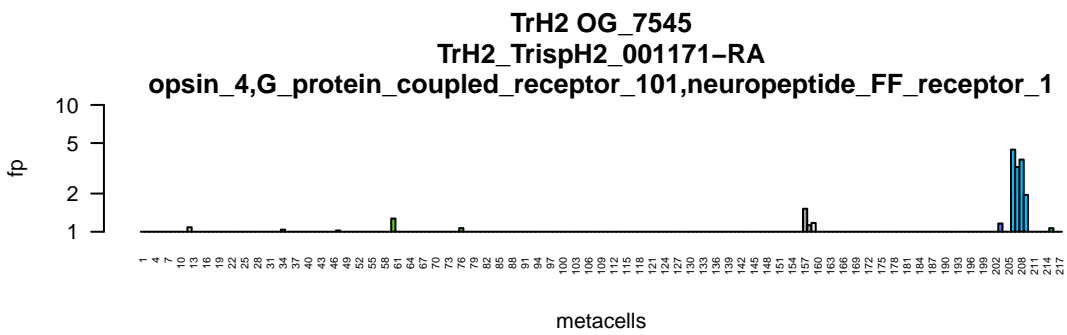
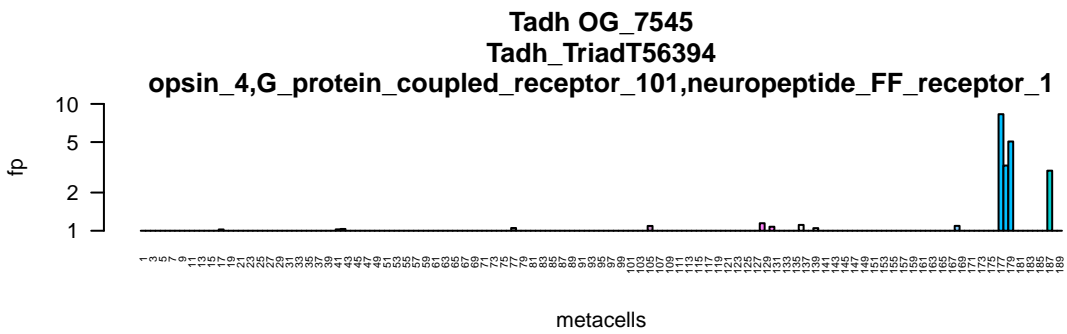
HoiH23\_PIH23\_011166-RA







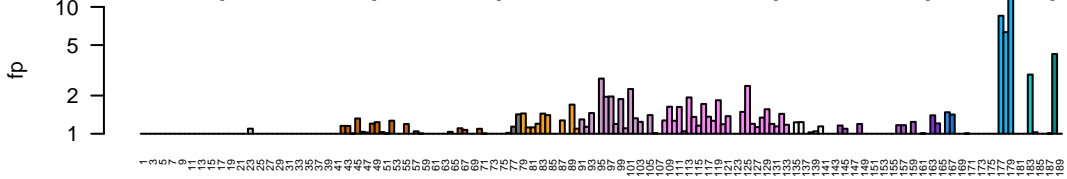




Tadh OG\_7669

Tadh\_TriadT57516

adhesion\_G\_protein\_coupled\_receptor\_G7,adhesion\_G\_protein\_coupled\_receptor\_G2

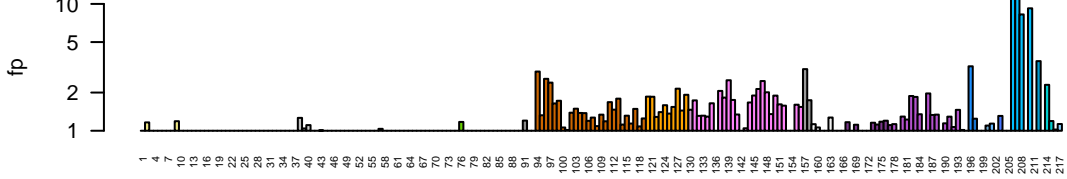


metacells

TrH2 OG\_7669

TrH2\_TrispH2\_004529-RA

adhesion\_G\_protein\_coupled\_receptor\_G7,adhesion\_G\_protein\_coupled\_receptor\_G2

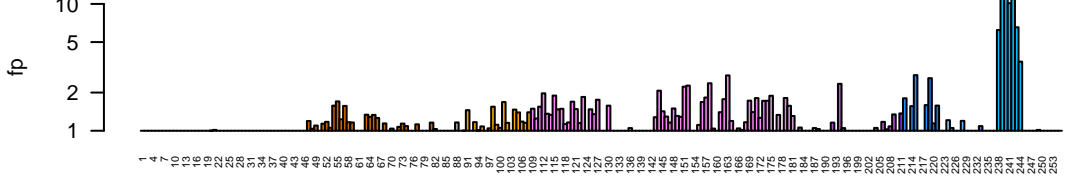


metacells

Hhon OG\_7669

Hhon\_g09503.t1

adhesion\_G\_protein\_coupled\_receptor\_G7,adhesion\_G\_protein\_coupled\_receptor\_G2

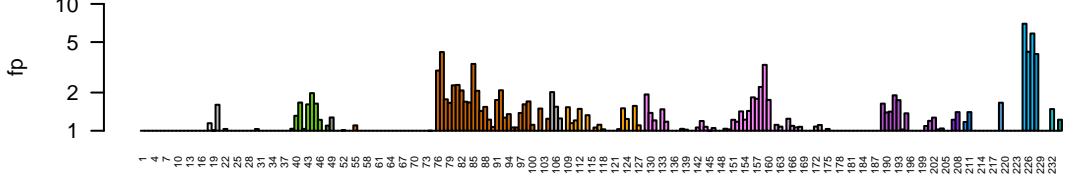


metacells

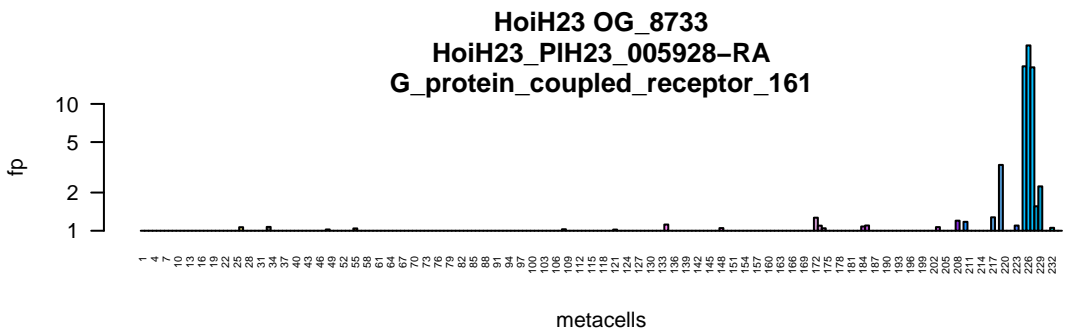
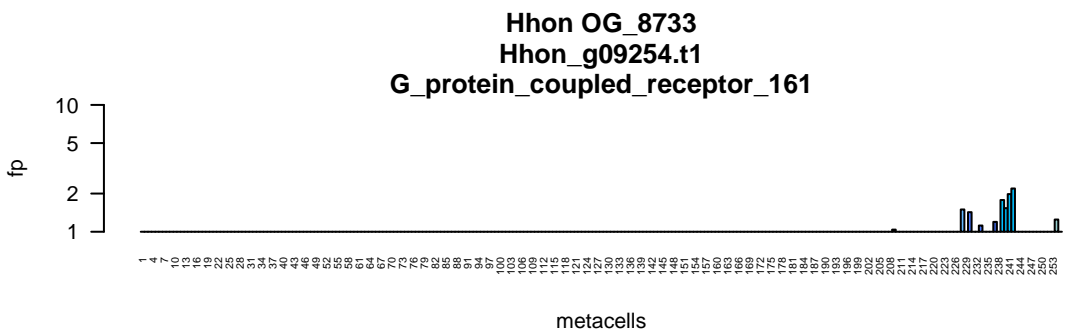
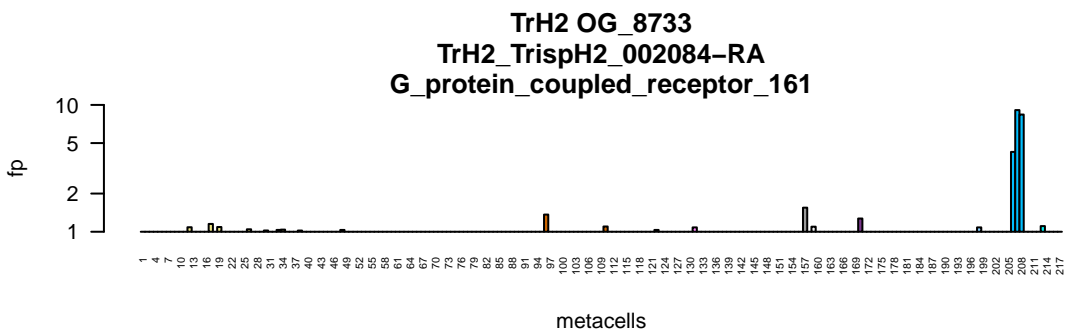
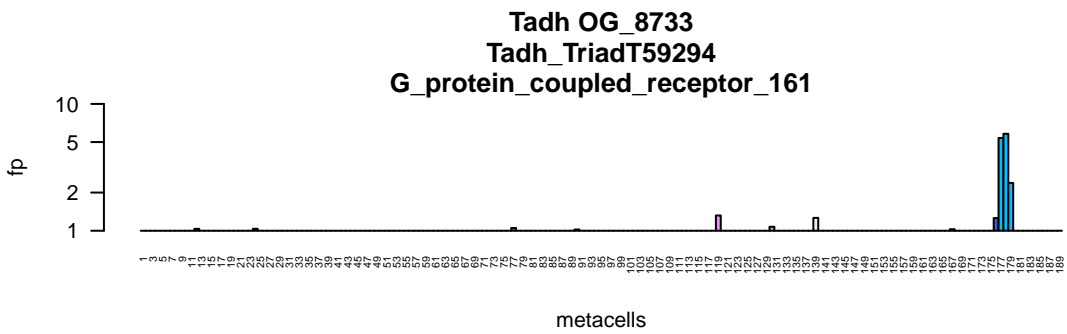
HoiH23 OG\_7669

HoiH23\_PIH23\_005798-RA

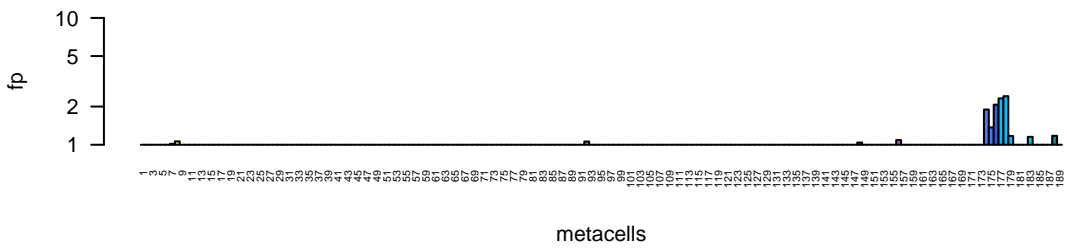
adhesion\_G\_protein\_coupled\_receptor\_G7,adhesion\_G\_protein\_coupled\_receptor\_G2



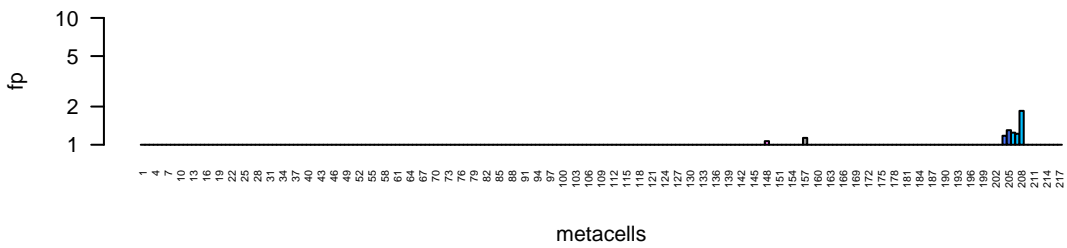
metacells



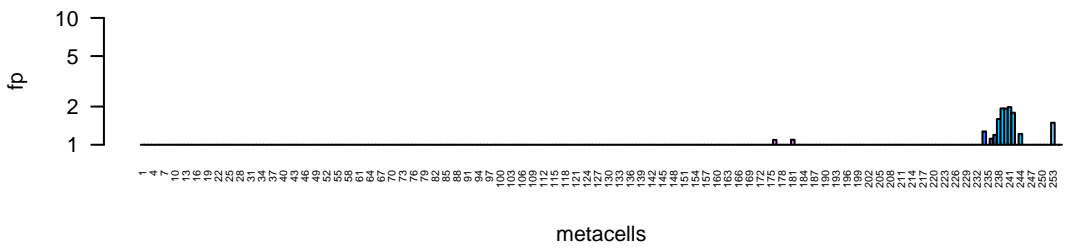
Tadh OG\_8863  
Tadh\_TriadT56181



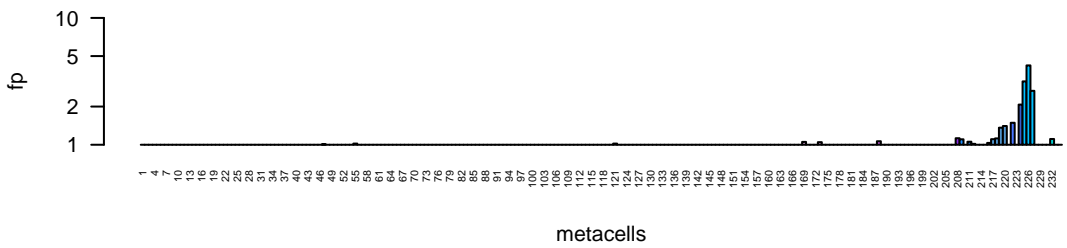
TrH2 OG\_8863  
TrH2\_TrispH2\_005495-RA

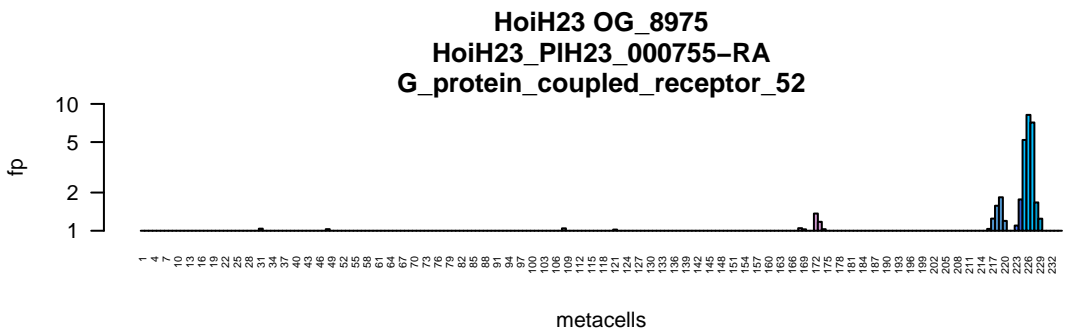
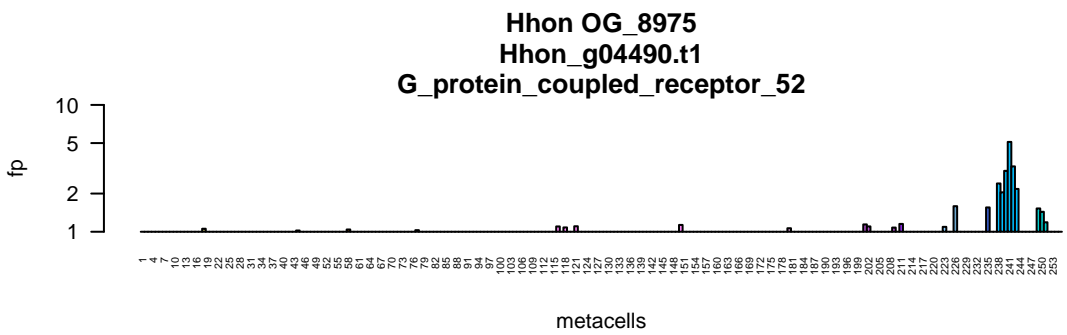
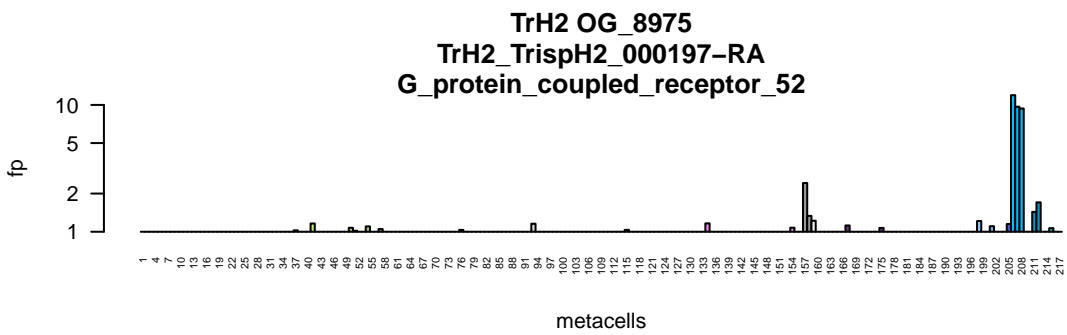
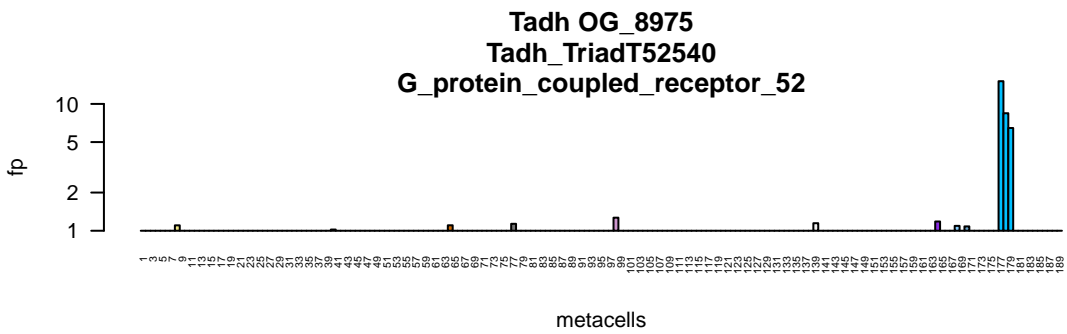


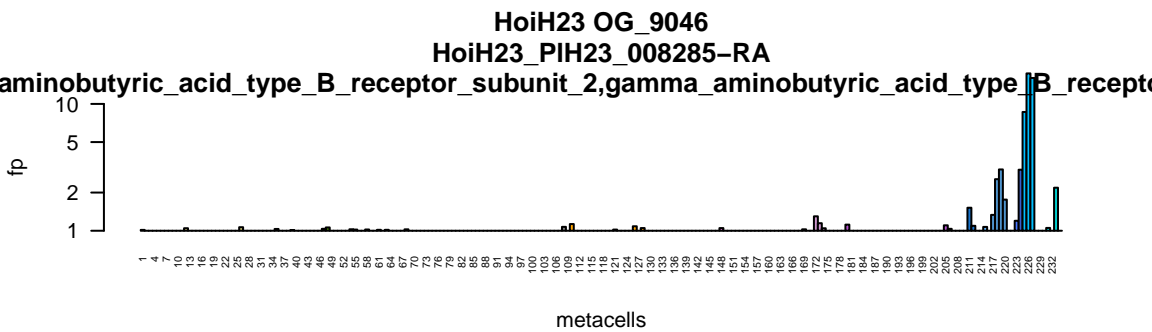
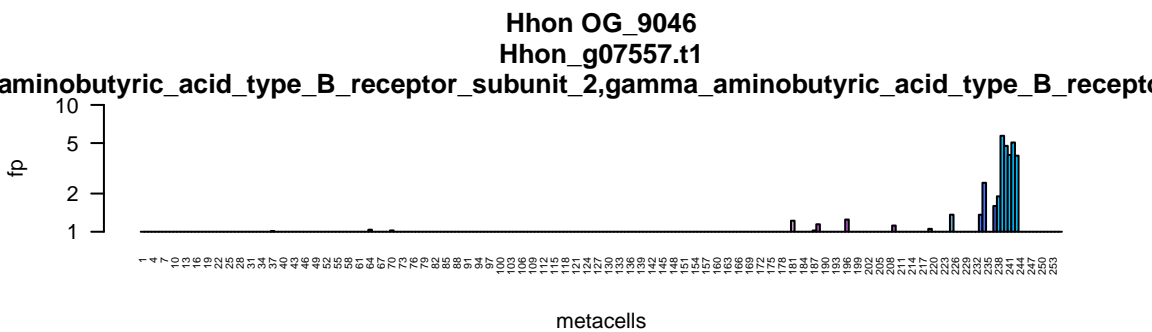
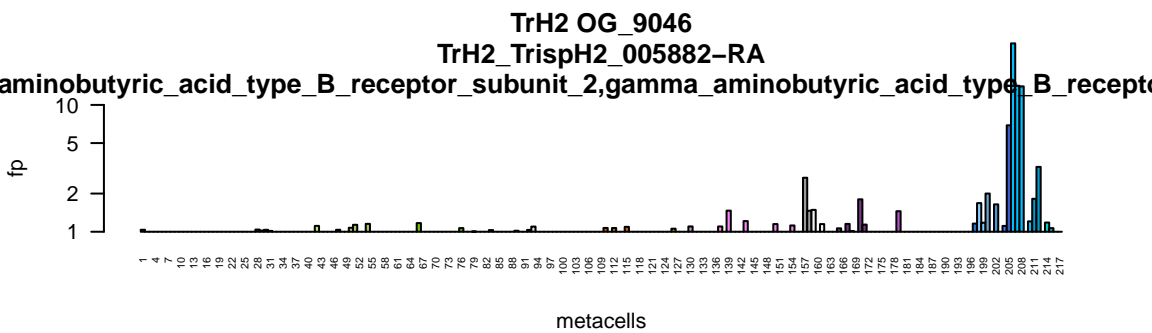
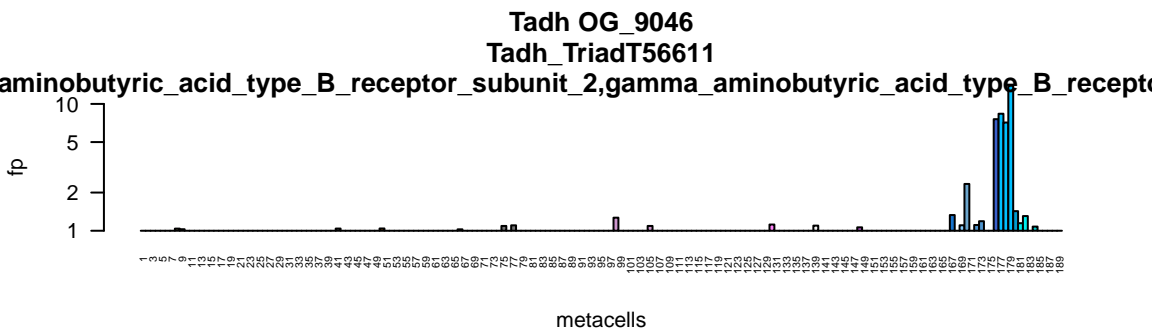
Hhon OG\_8863  
Hhon\_g10926.t1



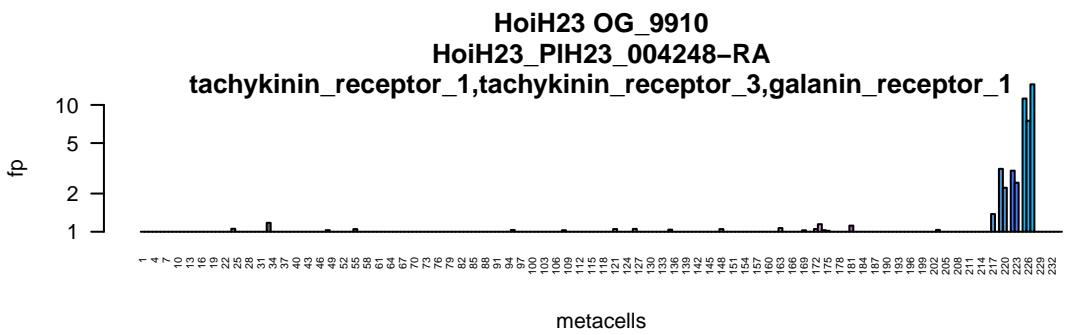
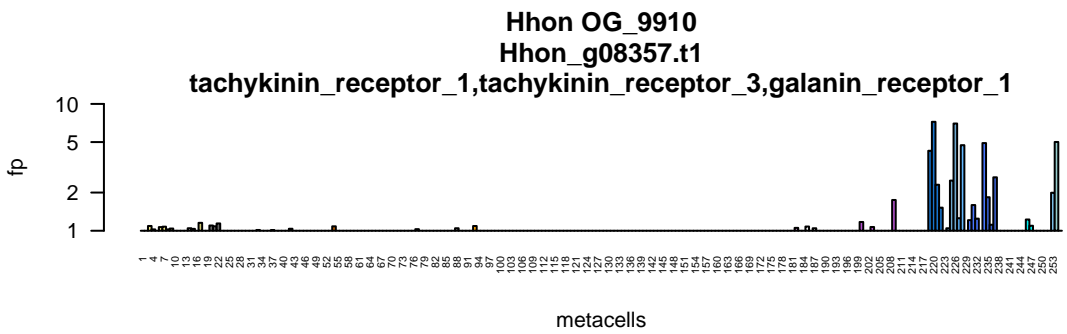
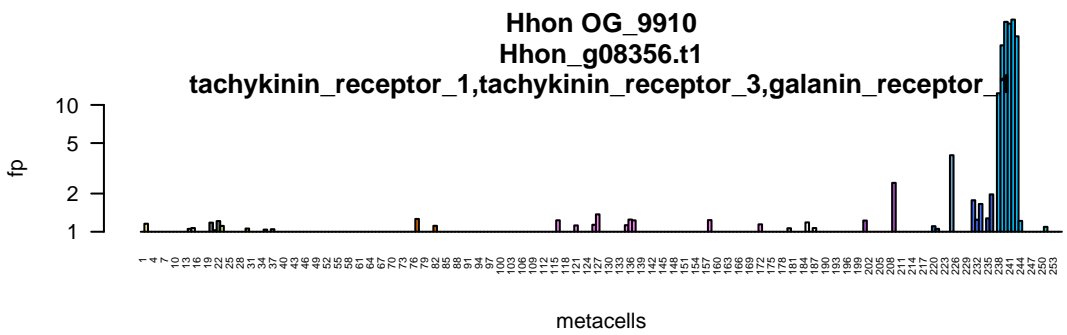
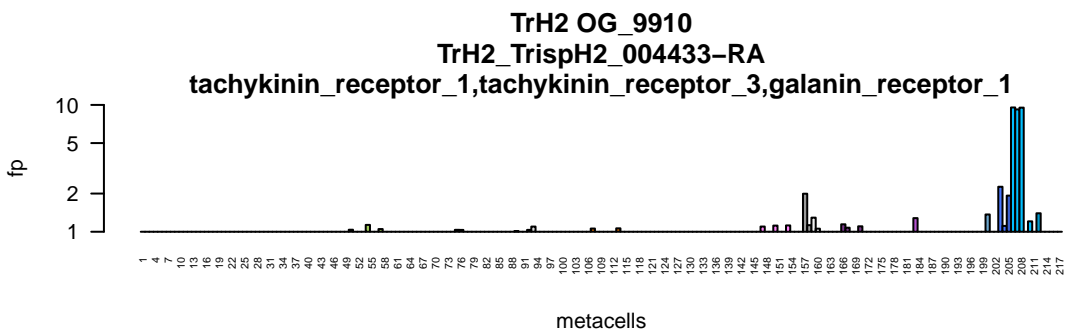
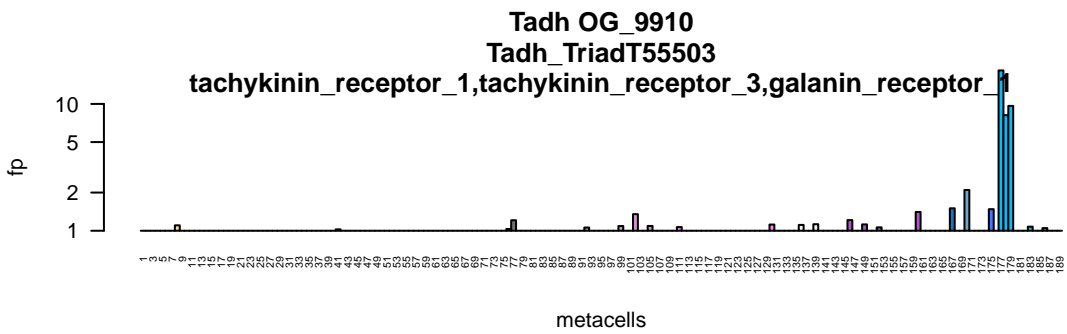
HoiH23 OG\_8863  
HoiH23\_PIH23\_009695-RA





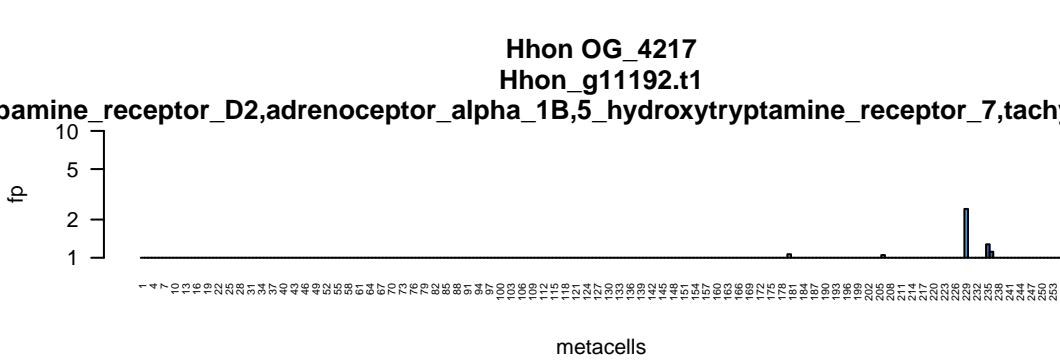
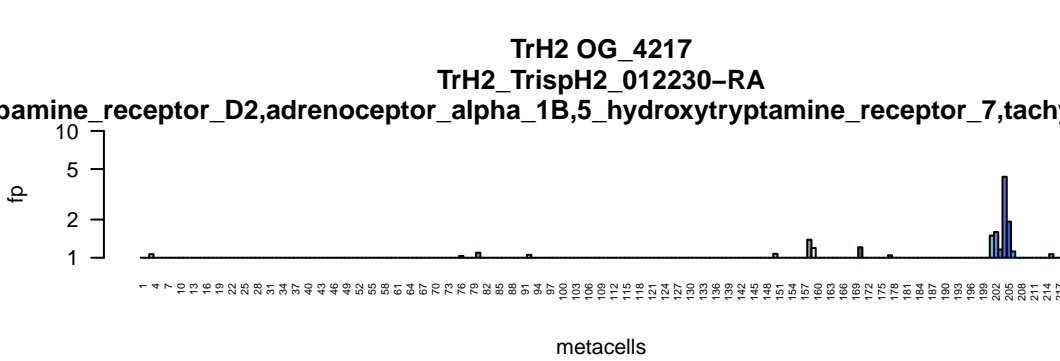
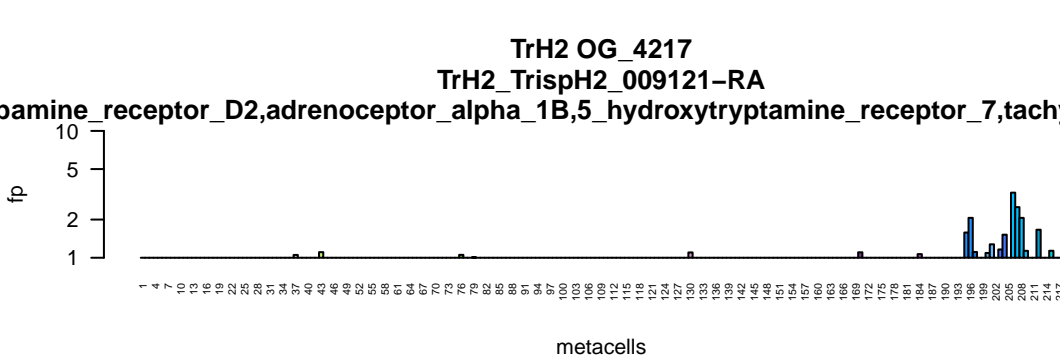
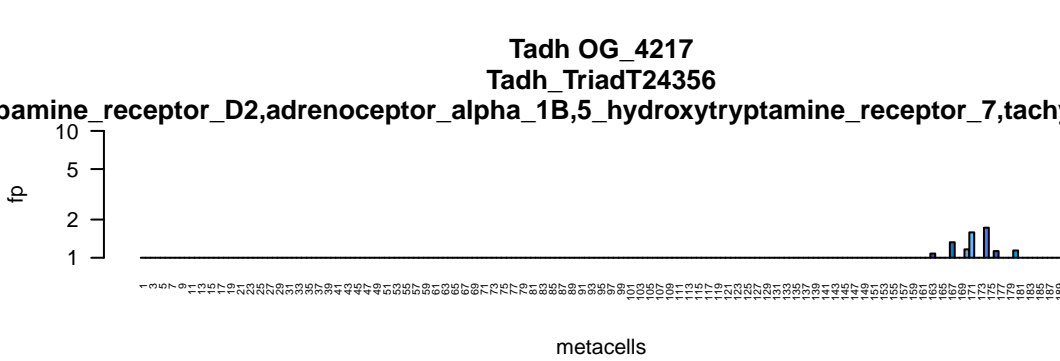




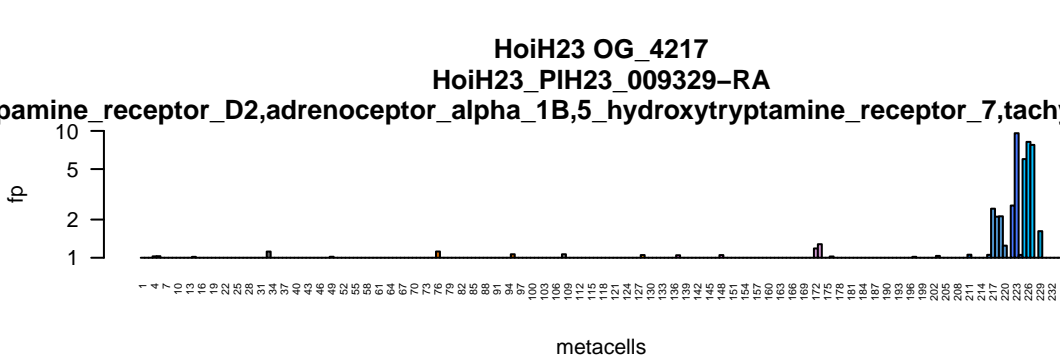


Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists 100 metacells. Most metacells have 0 false positives, but a few have 1, and a small cluster of metacells (173-187) has 2, 3, or 4 false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

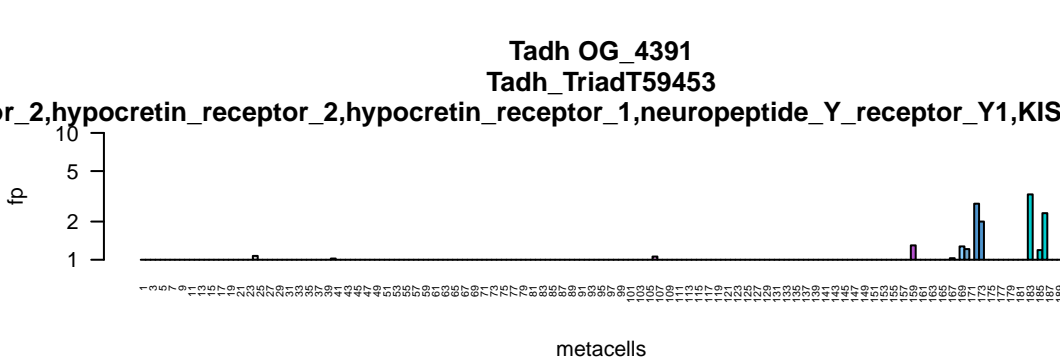
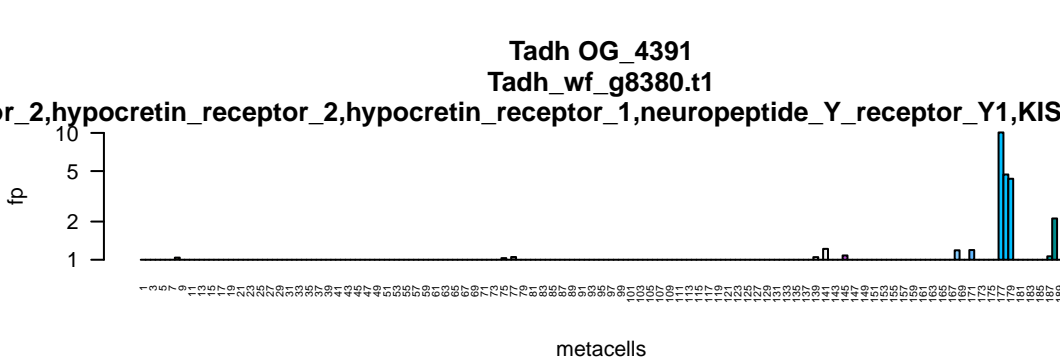


Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists indices from 1 to 232. Most metacells have a false positive count of 1. Metacells 169, 170, 171, 172, 173, 174, 175, 176, 177, 178, 179,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



or\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KIS

metacells	fp
11	1
13	1
17	1
21	1
22	1
23	1
27	1
31	1
33	1
35	1
39	1
41	1
45	1
49	1
51	1
53	1
55	1
59	1
61	1
63	1
65	1
69	1
73	1
75	1
79	1
83	1
85	1
89	1
93	1
95	1
99	1
103	1
107	1
109	1
113	1
117	1
119	1
123	1
127	1
129	1
133	1
137	1
139	1
143	1
147	1
149	1
153	1
157	1
161	1
163	1
167	1
171	1
173	1
177	2
178	2
179	1
183	1
187	1

[illegible]

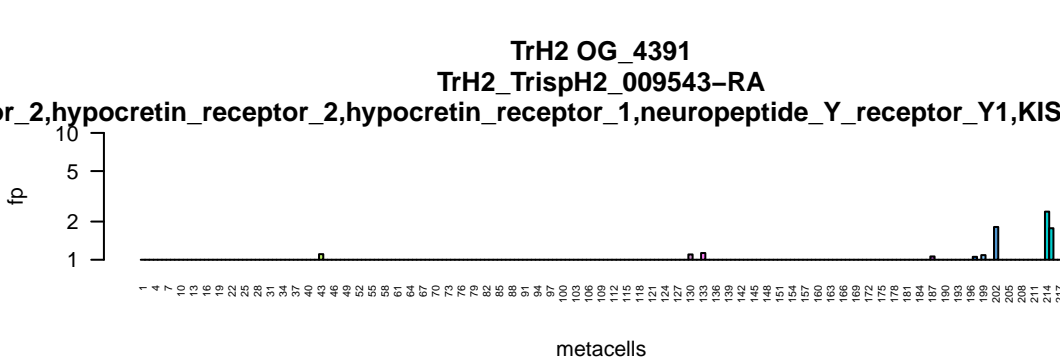
r\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KIS

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

or\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KIS

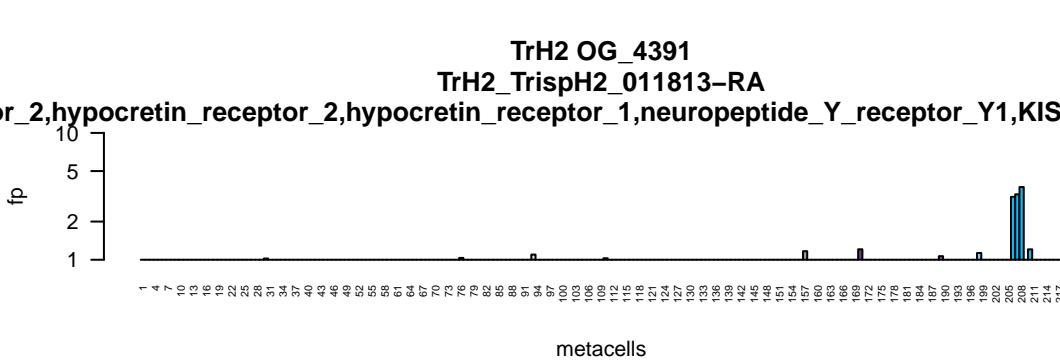
fp

metacells



or\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KIS

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	1
205	1
208	3
209	4
210	3
211	2
214	1



r\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KIS

metacell	fp
1	1
4	1
10	1
13	1
16	1
19	1
22	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	1
169	1.5
172	1.5
175	1
178	1
181	1.5
184	1
187	1
190	1
193	1
196	1
199	1
202	1
205	1
208	3
209	3
210	3
211	2
212	2

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is logarithmic, ranging from 1 to 10. The x-axis lists metacells from 1 to 232. Most metacells have a false positive count of 1, with a small cluster of higher counts (2, 3, 4, 5) for metacells 228 through 232.

Bar chart showing the number of reads (fp) for each metacell. The y-axis is logarithmic, ranging from 1 to 10. The x-axis lists metacells from 1 to 217. Most metacells have 1 read, with a few having 2 reads. Metacells 205 and 214 show significantly higher read counts, around 4 and 2 respectively.

metacell	fp
1	1
4	1
10	1
13	1
16	1
22	1
23	1
25	1
28	1
31	1
32	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
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	4
208	4
211	2
214	2
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 on a logarithmic scale. The x-axis is labeled 'metacells' and lists 50 metacells by index. Most metacells have a false positive count of 1, with a few having 2 or 5.

metacell index	fp
1	1
4	1
10	1
13	1
19	1
22	1
25	1
31	1
34	1
40	1
43	1
46	1
52	1
55	1
58	1
61	1
64	1
70	1
73	1
76	1
82	1
85	1
88	1
91	1
94	1
97	1
103	1
106	1
112	1
115	1
118	1
121	1
124	1
127	1
133	1
136	1
142	1
145	1
148	1
154	1
157	1
163	1
166	1
169	1
175	1
181	1
184	1
187	1
190	1
196	1
199	1
205	1
208	1
211	1
214	1
217	1
220	5
223	2
229	2

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is logarithmic, ranging from 1 to 10. The x-axis lists metacells from 1 to 253. Most metacells have a false positive count of 1, with a few outliers reaching up to 4.

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is logarithmic, ranging from 1 to 10. The x-axis lists metacells from 1 to 232. Most metacells have a false positive count of 1, with a small cluster of higher counts (2, 3, 4, 5) for metacells 228 through 232.

A bar chart showing the number of false positives (fp) for each metacell. The y-axis is logarithmic, ranging from 1 to 10. The x-axis lists metacells from 1 to 232. Most metacells have 1 false positive, with a small cluster of higher values around metacell 228.

metacell	fp
1	1
4	1
10	1
13	1
19	1
22	1
25	1
28	1
31	1
34	1
40	1
43	1
46	1
52	1
55	1
58	1
61	1
64	1
70	1
73	1
76	1
82	1
85	1
88	1
91	1
94	1
97	1
102	1
103	1
106	1
108	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
142	1
145	1
148	1
154	1
157	1
163	1
166	1
169	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
205	1
208	1
211	1
214	1
217	1
220	1
223	1
226	1
228	3
229	2
232	1

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is logarithmic, ranging from 1 to 10. The x-axis lists metacells from 1 to 232. Most metacells have 1 false positive, with a few having 2 or 3. Metacells 228 and 229 have significantly higher counts, around 8 and 6 respectively.

metacell	fp
1	1
4	1
10	1
13	1
19	1
22	1
25	1
28	1
31	1
34	1
40	1
43	1
46	1
55	1
58	1
61	1
64	1
73	1
76	1
82	2
85	2
88	1
91	1
94	1
97	1
102	1
103	1
106	1
108	1
112	1
115	1
118	1
119	1
124	1
127	1
133	1
136	1
142	1
145	1
148	1
154	1
157	1
163	1
166	1
169	1
175	1
178	1
181	1
184	1
187	1
190	1
196	1
199	1
205	1
208	1
211	1
214	1
217	2
220	3
223	3
228	8
229	6

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is logarithmic, ranging from 1 to 10. The x-axis lists metacells from 1 to 232. Most metacells have 1 false positive, with a few having 2 or 3. Metacells 228 and 229 have 10 false positives.

metacell	fp
1	1
4	1
10	1
13	1
19	1
22	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
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
220	1
223	1
226	1
228	10
229	10

A bar chart showing the number of false positives (fp) for each metacell. The y-axis is logarithmic, ranging from 1 to 10. The x-axis lists metacells from 1 to 232. Most metacells have 1 false positive, with a small cluster of higher values around metacell 228.

metacell	fp
1	1
4	1
10	1
13	1
19	1
22	1
25	1
28	1
31	1
34	1
40	1
43	1
46	1
55	1
61	1
64	1
70	1
73	1
76	1
82	1
85	1
88	1
91	1
94	1
97	1
102	1
103	1
106	1
112	1
115	1
119	1
121	1
124	1
127	1
133	1
136	1
142	1
145	1
148	1
154	1
157	1
163	1
166	1
169	1
175	1
184	1
187	1
190	1
196	1
199	1
205	1
208	1
211	1
214	1
217	1
220	1
223	1
228	2
229	1
232	1

TrH2 OG\_4560  
TrH2\_TrispH2\_011838-RA  
adenosine\_A3\_receptor,histamine\_receptor\_H2,trace\_amine\_associated\_receptor\_1,adrenoceptor

metacells	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	1.5
202	1.8
205	1.2
208	1
211	1
214	1.1
217	1

**Hhon OG\_4560**  
**Hhon\_g05281.t1**  
**adenosine\_A3\_receptor,histamine\_receptor\_H2,trace\_amine\_associated\_receptor\_1,adrenoceptor**

fp

metacells

Hhon OG\_4560  
Hhon\_g05282.t1  
adenosine\_A3\_receptor,histamine\_receptor\_H2,trace\_amine\_associated\_receptor\_1,adrenocortical\_11b\_hydroxylase

fp

metacells

Hhox OG\_4560  
Hhox\_g05139.t1

adenosine\_A3\_receptor,histamine\_receptor\_H2,trace\_amine\_associated\_receptor\_1,adrenoceptor\_1

fp

metacells

metacell	Hhox OG_4560 (fp)	Hhox_g05139.t1 (fp)
1	1	1
10	1	1
13	1	1
19	1	1
22	1	1
28	1	1
31	1	1
37	1	1
43	1	1
49	1	1
52	1	1
55	1	1
59	1	1
64	1	1
70	1	1
73	1	1
79	1	1
82	1	1
88	1	1
91	1	1
97	1	1
100	1	1
106	1	1
109	1	1
115	1	1
118	1	1
124	1	1
127	1	1
136	1	1
142	1	1
146	1	1
151	1	1
154	1	1
160	1	1
163	1	1
169	1	1
175	1	1
178	1	1
184	1	1
187	1	1
193	1	1
196	1	1
202	1	1
205	1	1
211	1	1
214	3	3
215	3	3
220	3	3
223	1	1
229	1	1
232	1	1
238	1	1
241	1	1
247	1	1
250	1	1
253	1	1

**HoiH23 OG\_4560**  
**HoiH23\_PIH23\_011580-RA**  
**adenosine\_A3\_receptor,histamine\_receptor\_H2,trace\_amine\_associated\_receptor\_1,adrenocortical\_11beta\_hydroxylase**

metacells	fp
1	1
4	1
10	1
13	1
16	1
22	1
25	1
31	1
34	1
37	1
43	1
46	1
52	1
55	1
58	1
64	1
67	1
73	1
76	1
79	1
85	1
88	1
91	1
97	1
100	1
103	1
106	1
109	1
112	1
118	1
121	1
127	1
130	1
133	1
139	1
142	1
148	1
151	1
154	1
160	1
163	1
169	1
172	1
175	1
181	1
184	1
187	1
190	1
193	1
196	1
202	1
205	1
209	1
214	1
217	1
223	1
226	1
229	1
232	2

HoiH23 OG\_4560  
 HoiH23\_PIH23\_011821-RA  
 enosine\_A3\_receptor,histamine\_receptor\_H2,trace\_amine\_associated\_receptor\_1,adreno

fp

metacells

fp

metacells

metacell	fp
1	1
4	1
10	1
13	1
16	1
22	1
25	1
34	1
37	1
43	1
46	1
52	1
55	1
58	1
64	1
67	1
73	1
76	1
79	1
85	1
88	1
91	1
97	1
100	1
103	1
106	1
109	1
112	1
118	1
121	1
127	1
130	1
133	1
139	1
142	1
148	1
151	1
154	1
160	1
163	1
169	1
172	1
175	1
181	1
184	1
187	1
190	1
193	1
196	1
202	1
205	1
209	1
214	1
217	1
223	1
226	1
229	1
236	1

Bar chart showing the number of reads (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 217. Most metacells have 1 read, with a few having 2 reads (metacells 64, 76, 196, 214, 217).

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 shows frequency from 1 to 10. The chart displays a distribution of frequencies across the metacells, 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.

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

A bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and the y-axis is labeled 'fp'. The y-axis has a non-linear scale with ticks at 1, 2, 5, and 10. The x-axis lists 189 metacells, with some labels truncated. The bars represent the frequency of each metacell, with most having a frequency of 1, and a few having higher frequencies up to 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
188	1
189	1

fp

metacells

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists 189 metacells. Most metacells have 1 false positive, with a few having 2 or 4. Metacells 175, 176, 177, 178, and 179 have 4 false positives each.

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.

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 217. The chart shows a distribution of false positives across the metacells, with most having 0 or 1 false positive, and a few having up to 3.

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

metacells	fp
1	1
4	1
10	1
13	1
16	1
22	1
23	1
25	1
28	1
31	1
32	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
202	1
205	1
208	1
211	1
214	1
217	1

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

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

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

metacells	fp
1	1
4	1
10	1
13	1
16	1
22	1
23	1
25	1
28	1
31	1
32	1
37	1
40	1
43	2
44	1
45	1
49	1
52	1
55	1
59	1
61	1
64	1
67	1
73	1
76	1
79	1
82	1
83	1
88	1
91	1
94	1
97	1
100	1
103	1
108	1
110	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	2
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
200	1
202	2
205	1
208	1
211	1
214	1
217	1

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 217. Most metacells have a false positive count of 1, with a few having counts of 2 or 3. Metacells 205 and 206 show the highest counts, around 4.

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 217. Most metacells have 0 false positives, but some have 1 or 2. Metacells 181 and 202 have 2 false positives each.

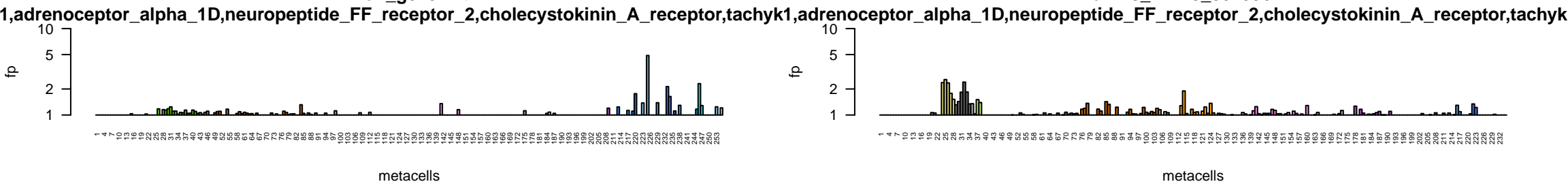
metacell	fp
1	0
4	0
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	0
34	0
37	1
40	1
43	0
46	0
49	0
52	0
55	0
58	0
61	0
64	1
67	0
70	0
73	0
76	0
79	0
82	0
85	0
88	0
91	0
94	1
97	1
100	0
103	0
106	0
109	0
112	1
115	1
118	1
121	1
124	1
127	1
130	0
133	0
136	0
139	0
142	1
145	1
148	0
151	1
154	0
157	0
160	0
163	0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

A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis ranges from 1 to 217, and the y-axis ranges from 1 to 10. The chart shows a distribution of feature counts across metacells, with a peak around metacell 145.

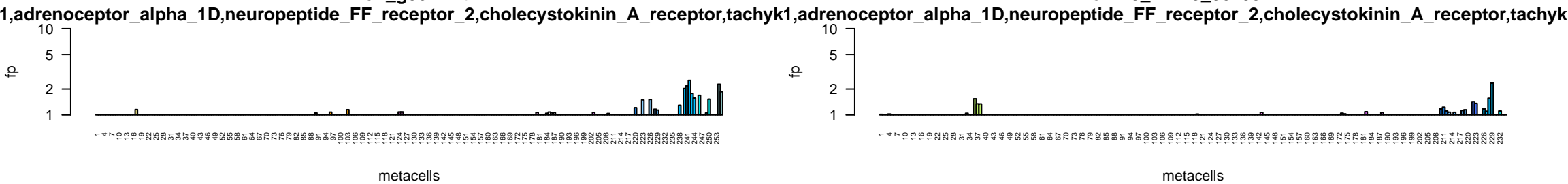
A bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and the y-axis is labeled 'fp'. The x-axis has 253 categories, and the y-axis ranges from 1 to 10. Most categories have a frequency of 1, with a small cluster of higher frequencies around category 241.

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
76	1
79	1
85	1
89	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
236	1
241	2
241	3
241	4
241	5
247	1
250	1
253	1

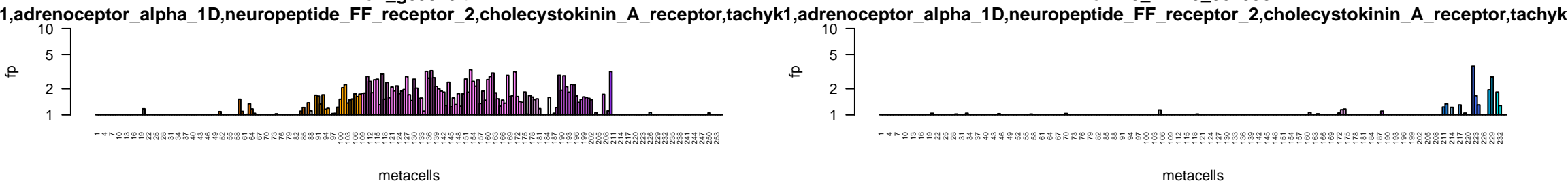
Hhon OG\_5725  
Hhon\_g04574.t1



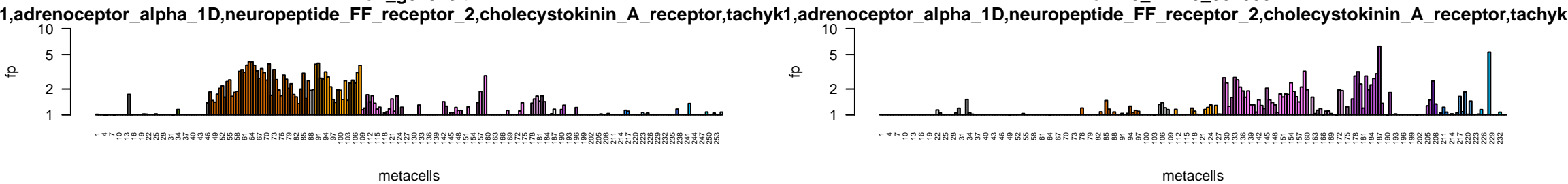
Hhon OG\_5725  
Hhon\_g06114.t1



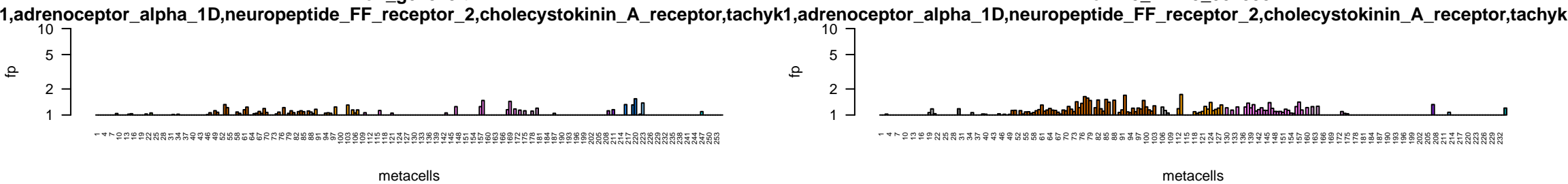
Hhon OG\_5725  
Hhon\_g09573.t1



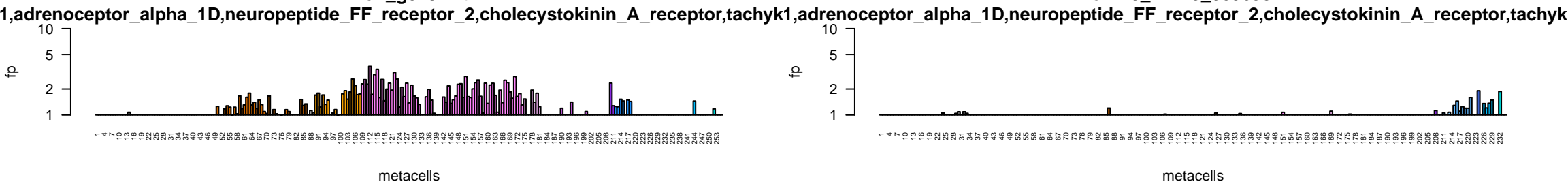
Hhon OG\_5725  
Hhon\_g07528.t1



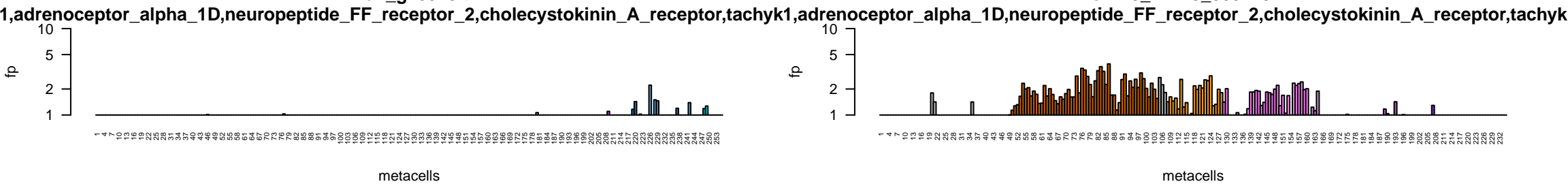
Hhon OG\_5725  
Hhon\_g07529.t1



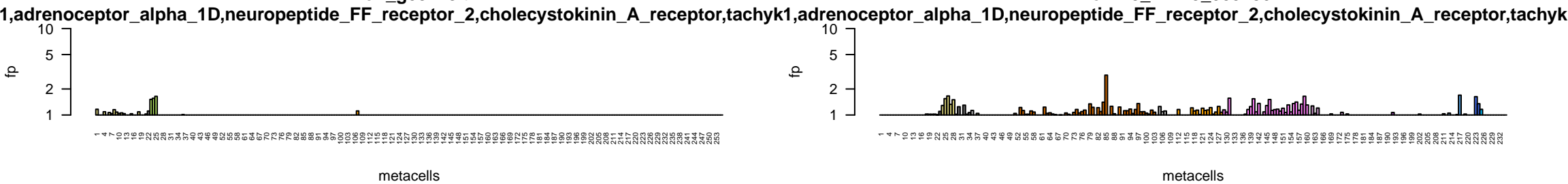
Hhon OG\_5725  
Hhon\_g07527.t1

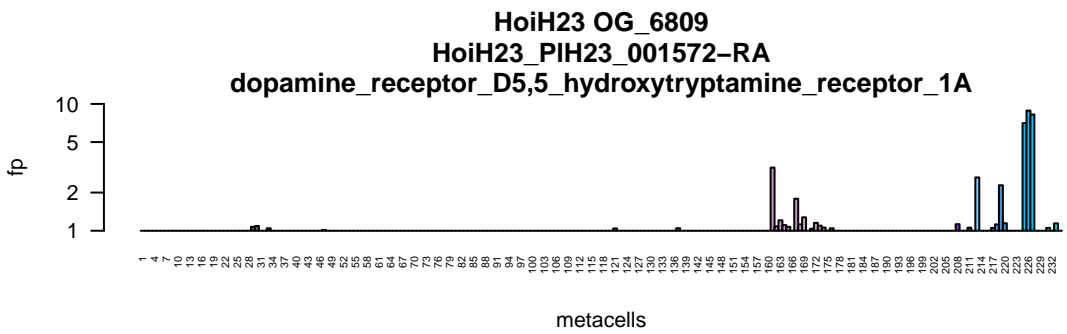
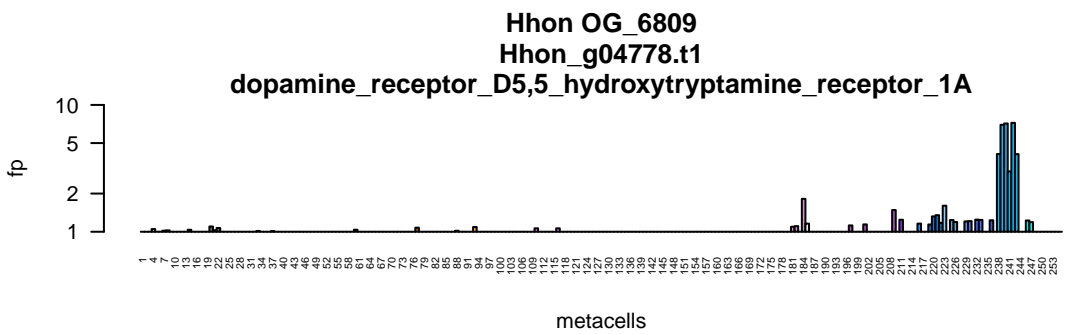
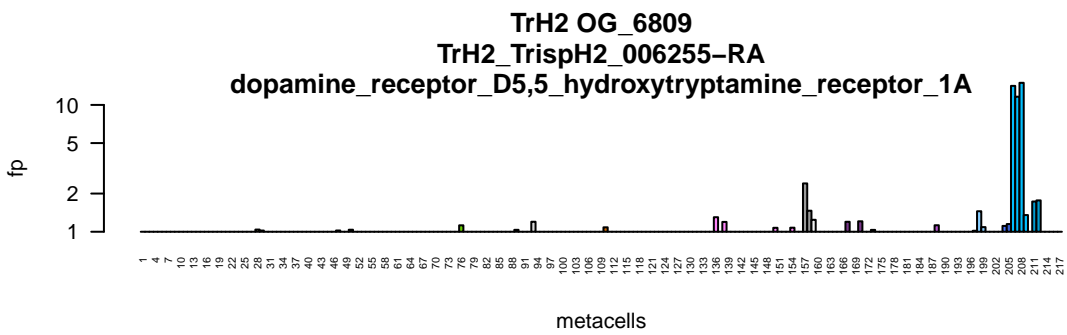
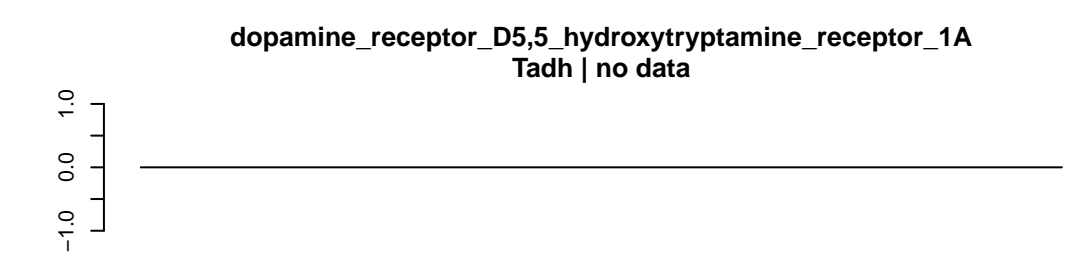


Hhon OG\_5725  
Hhon\_g10923.t1

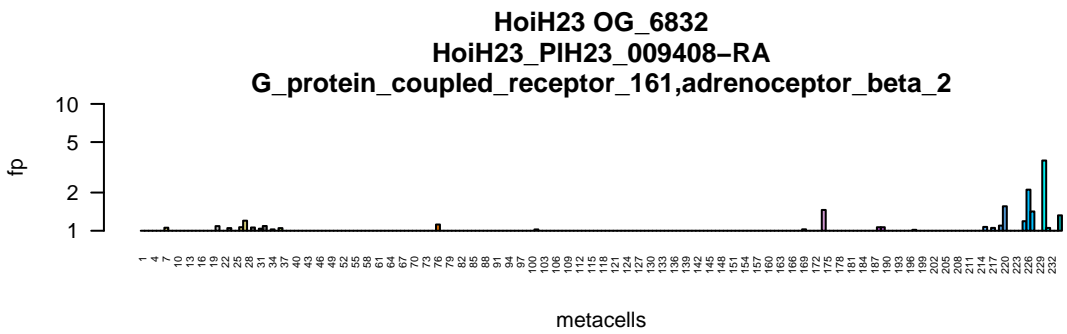
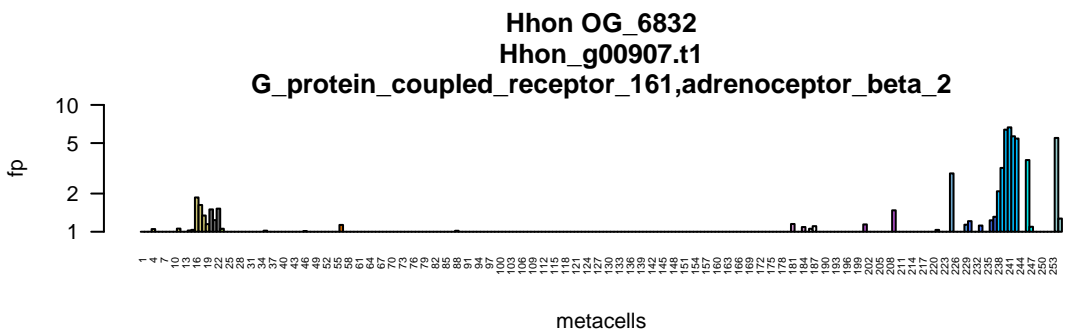
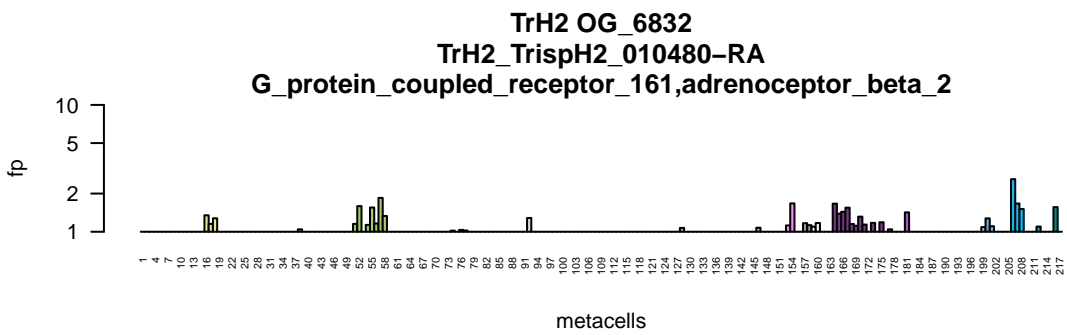
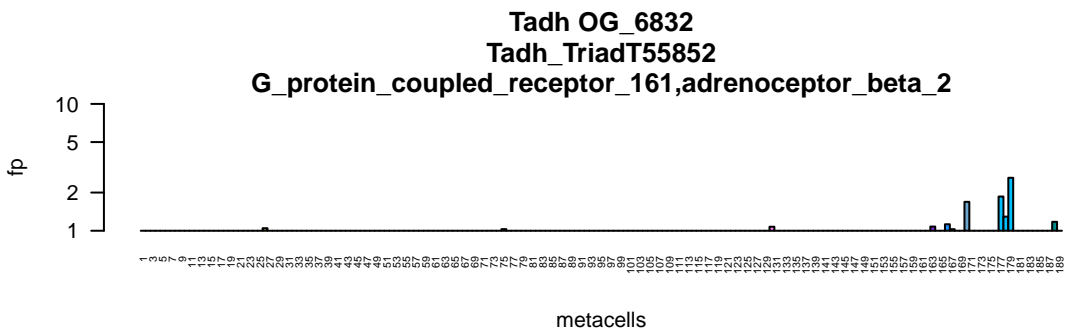


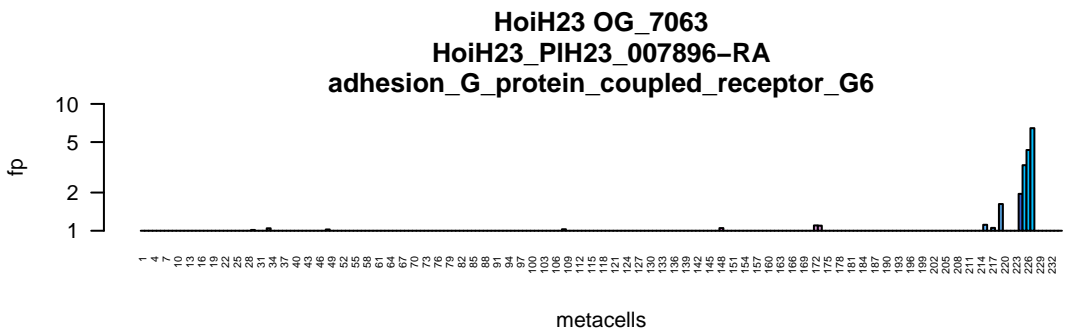
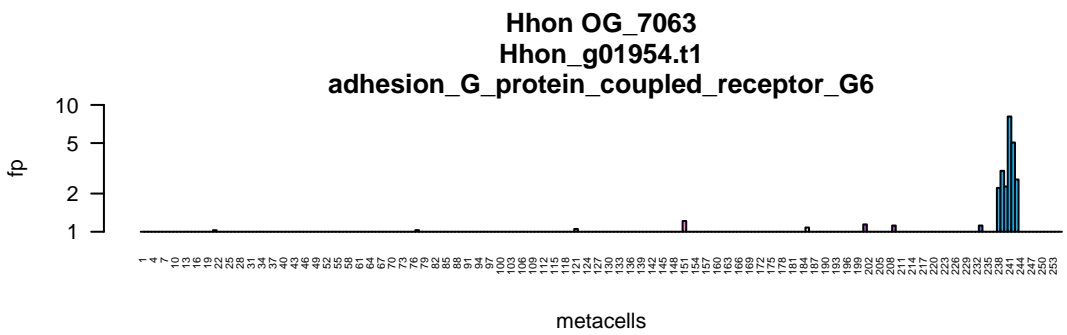
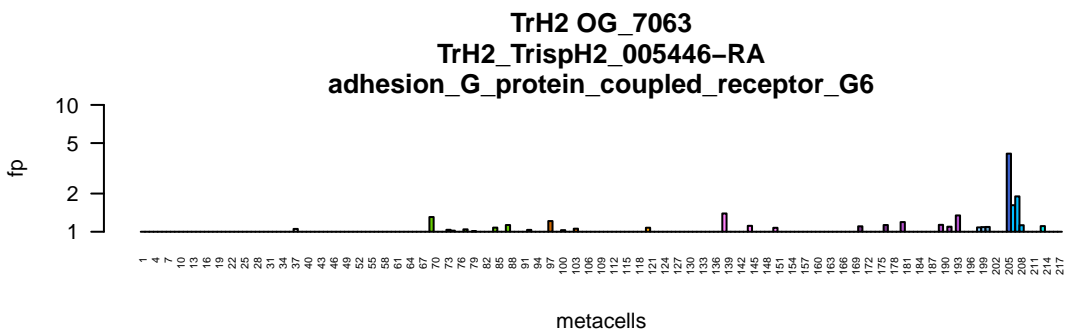
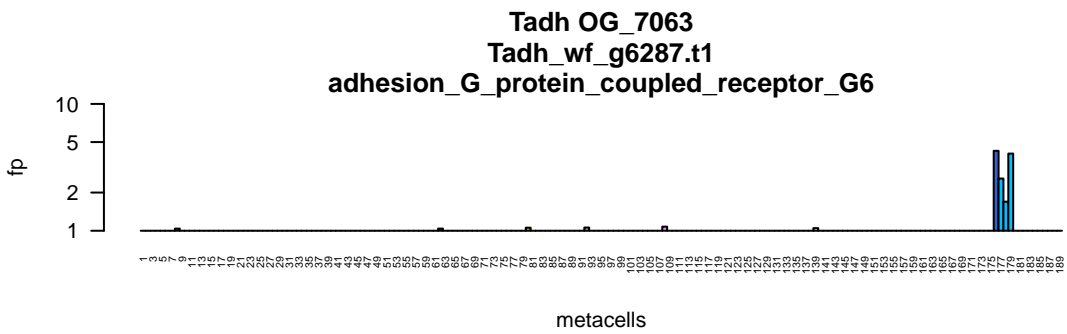
Hhon OG\_5725  
Hhon\_g08418.t1

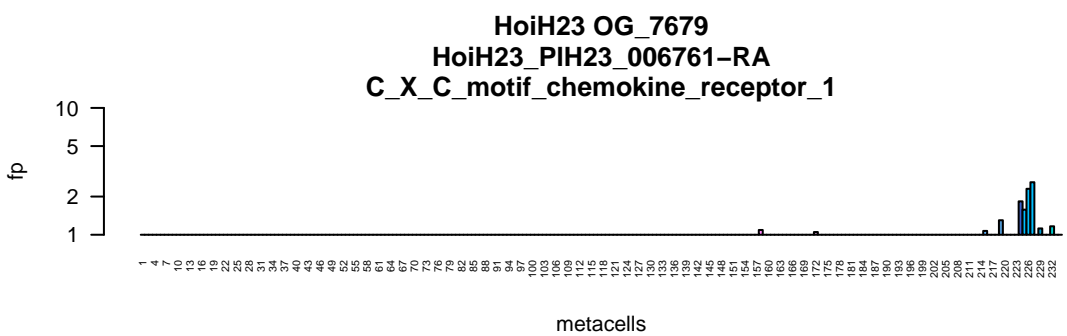
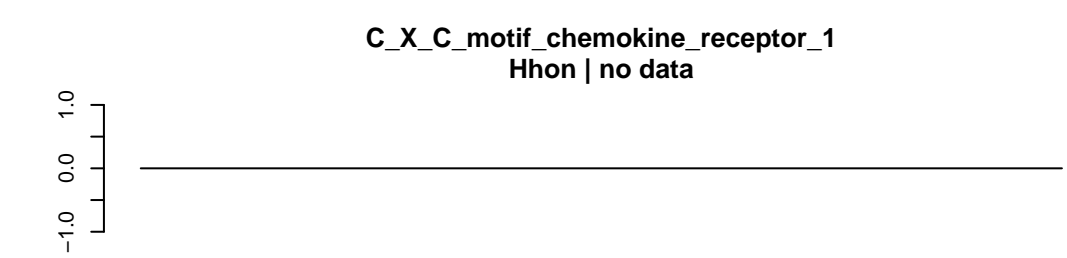
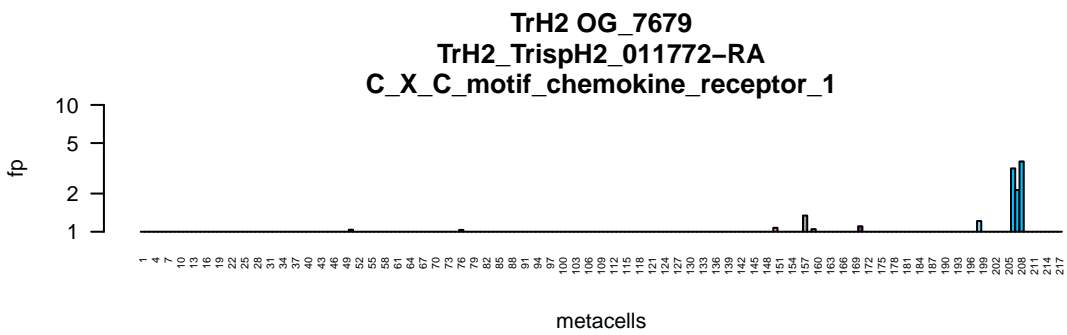
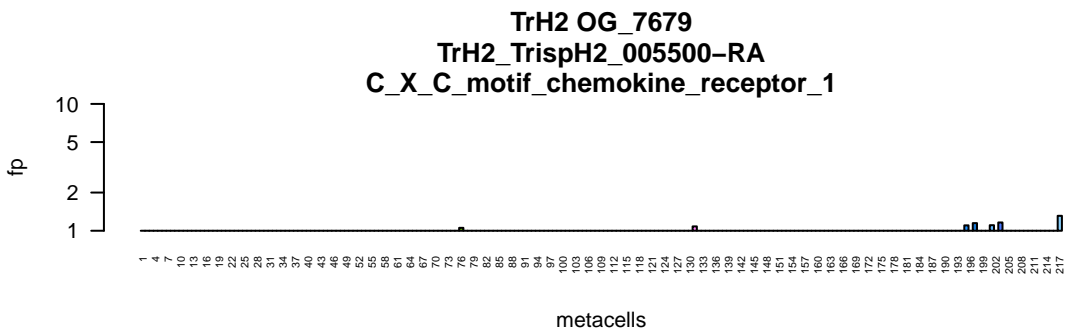
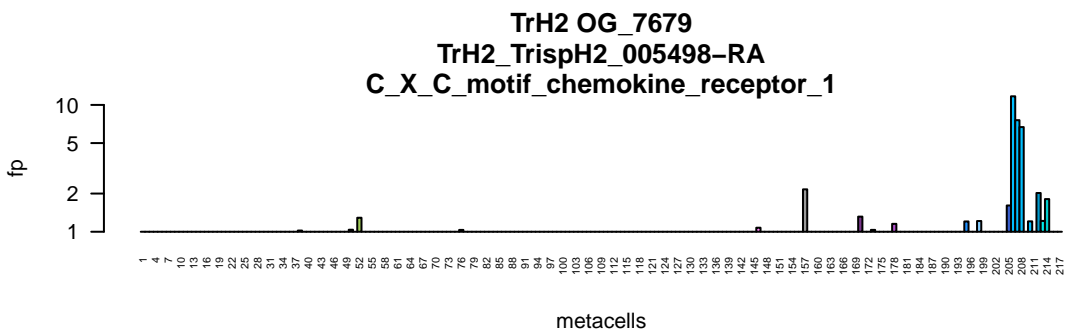
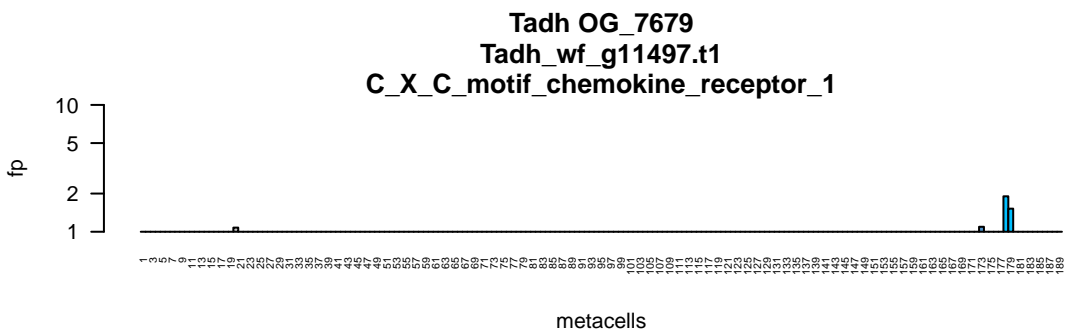
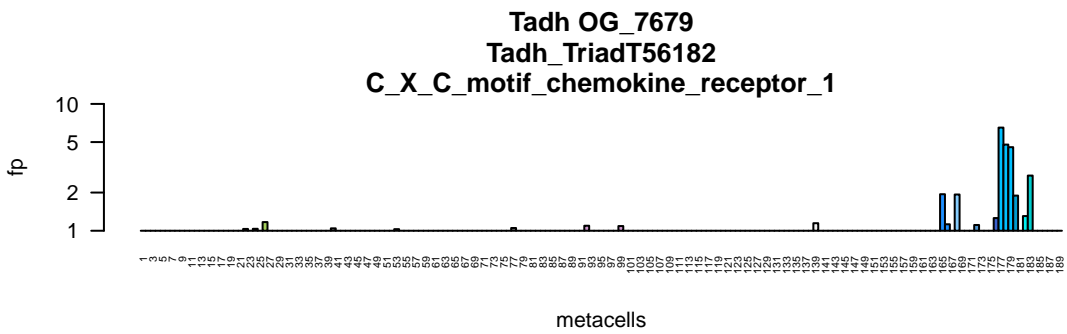


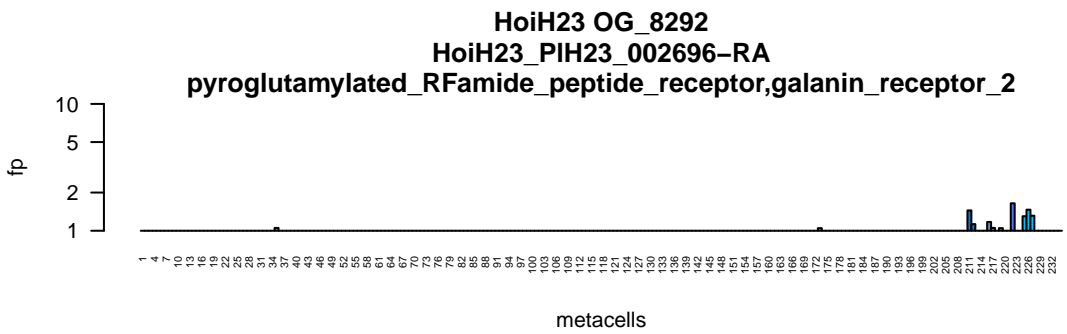
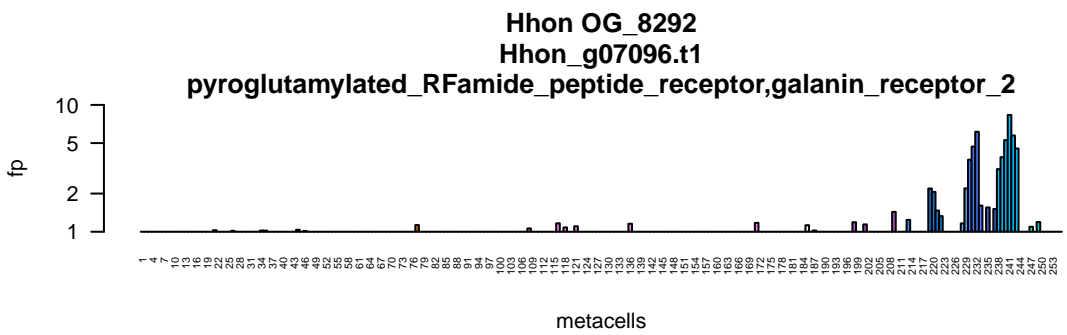
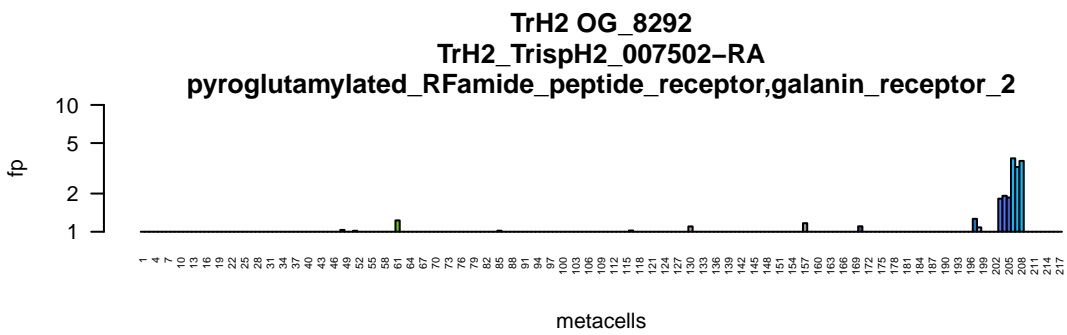
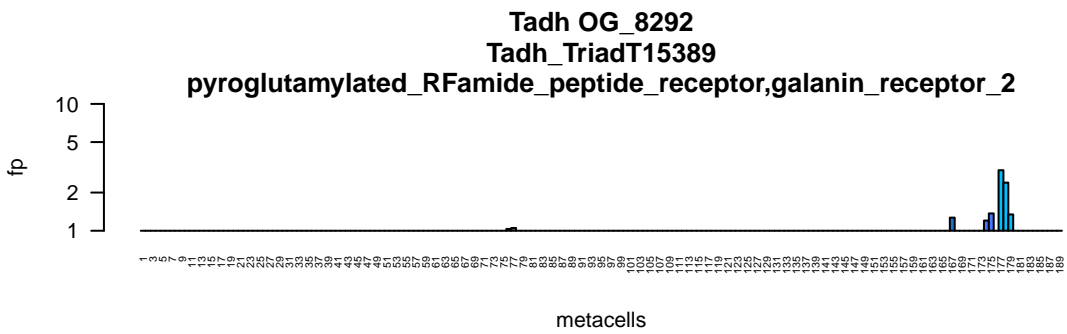


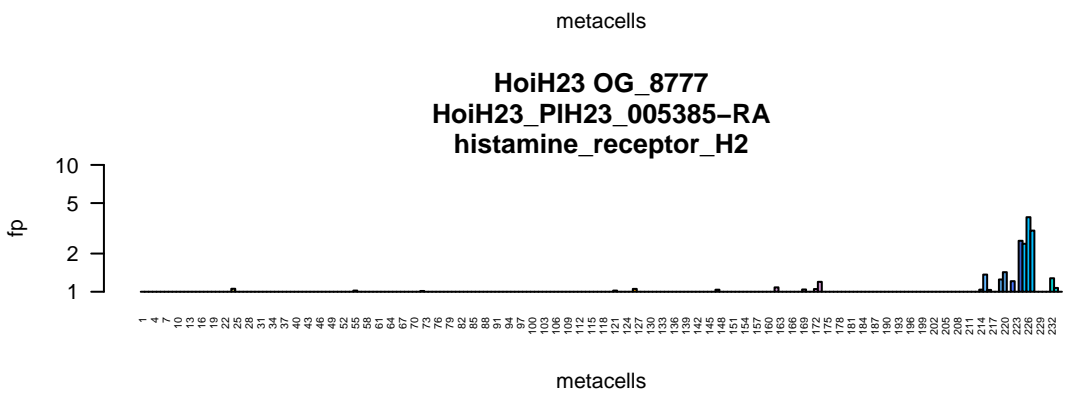
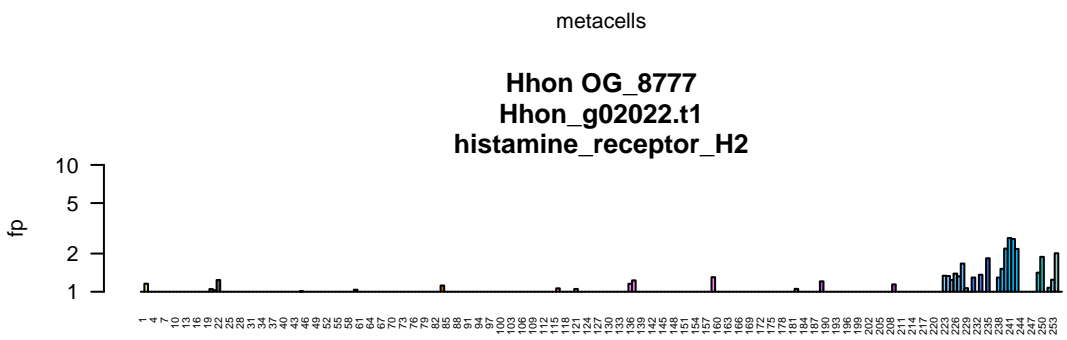
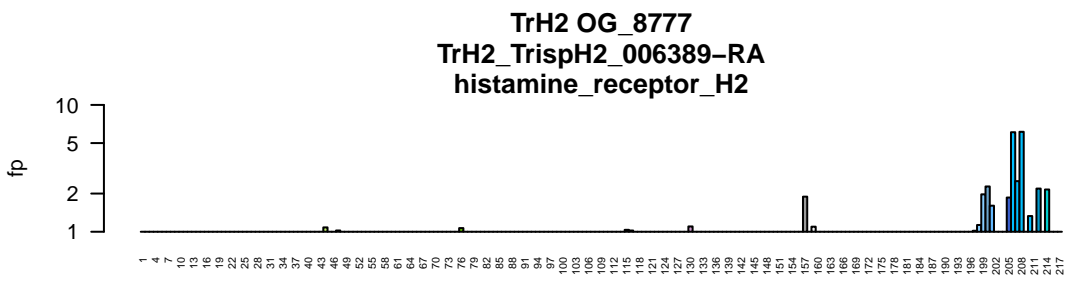
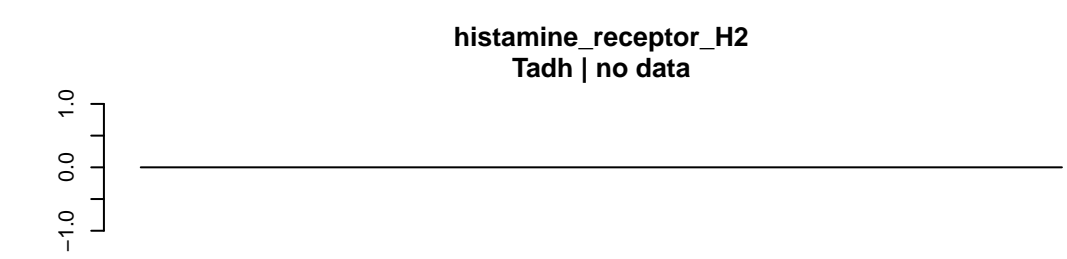


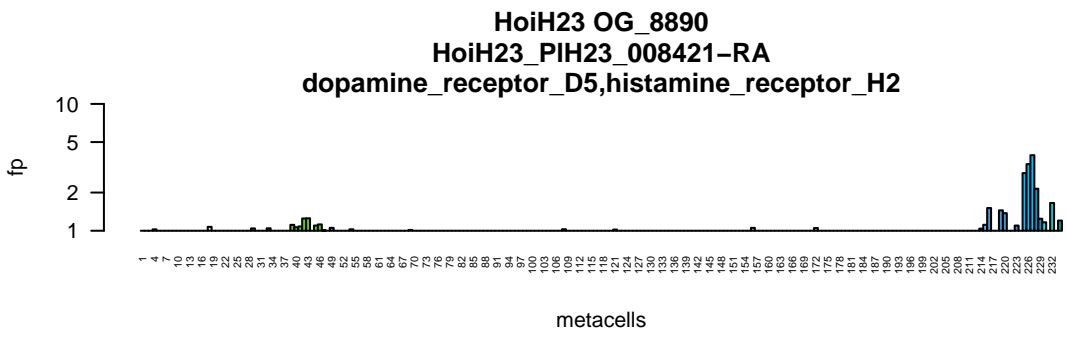
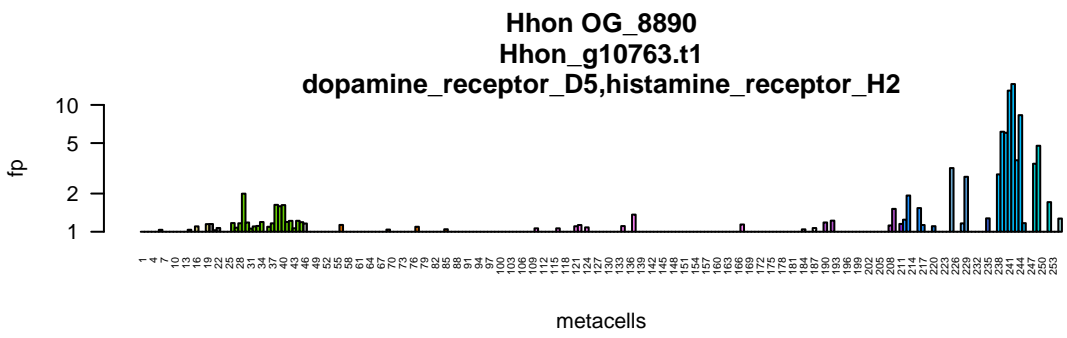
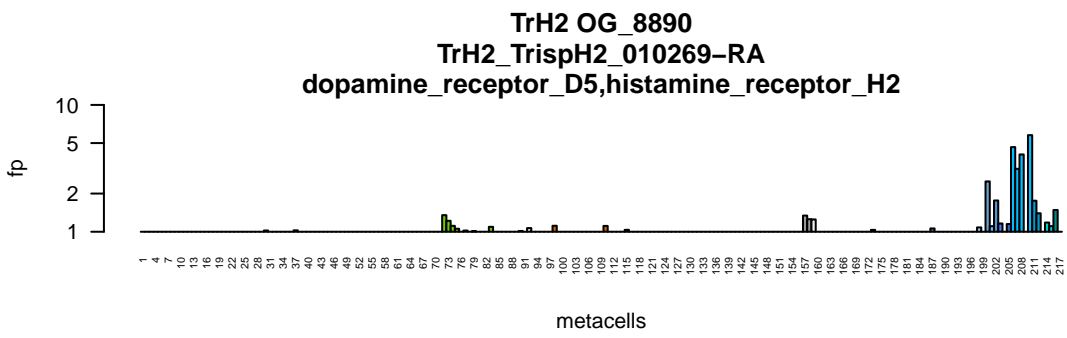
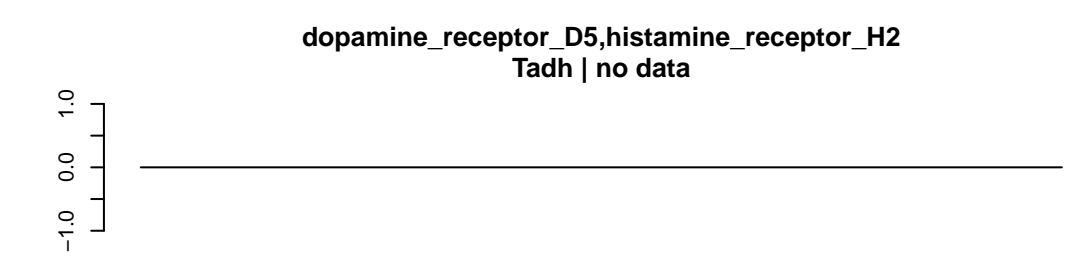


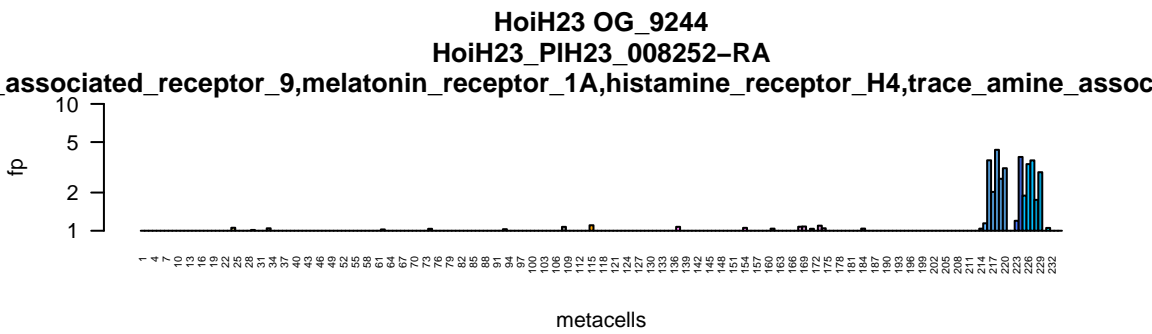
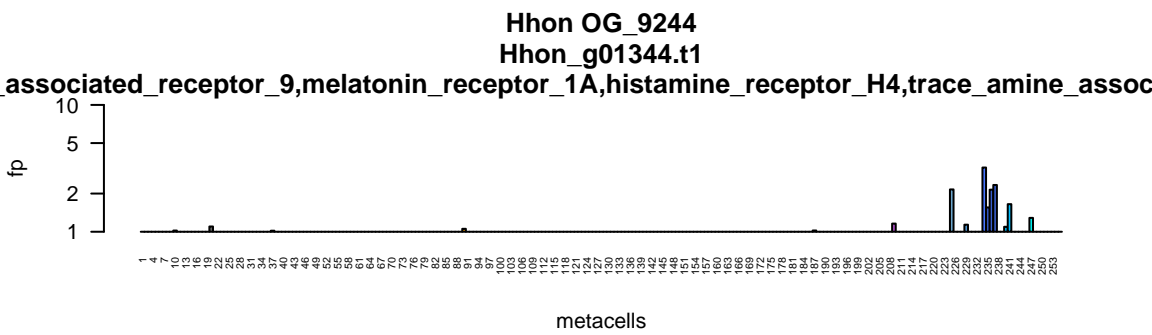
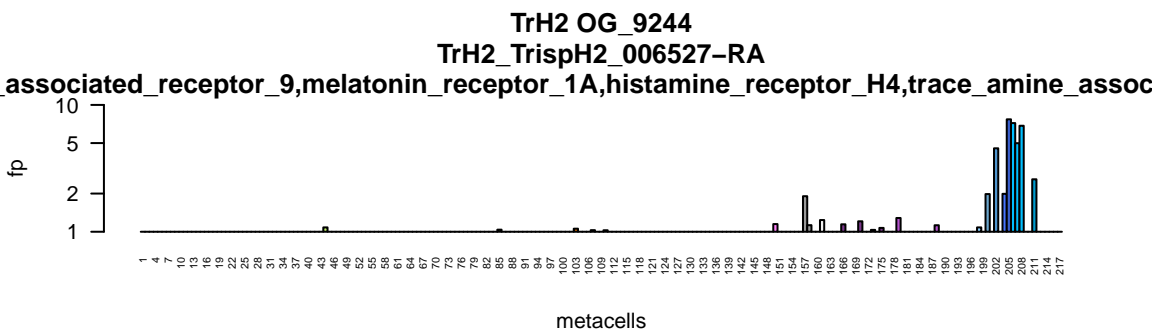
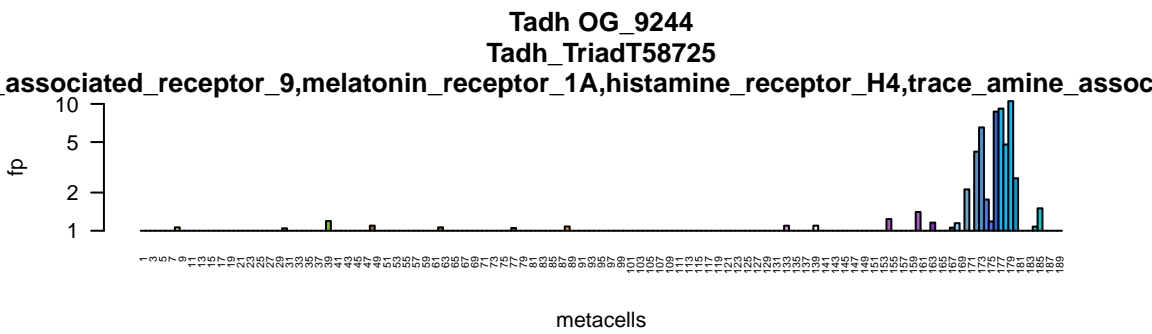


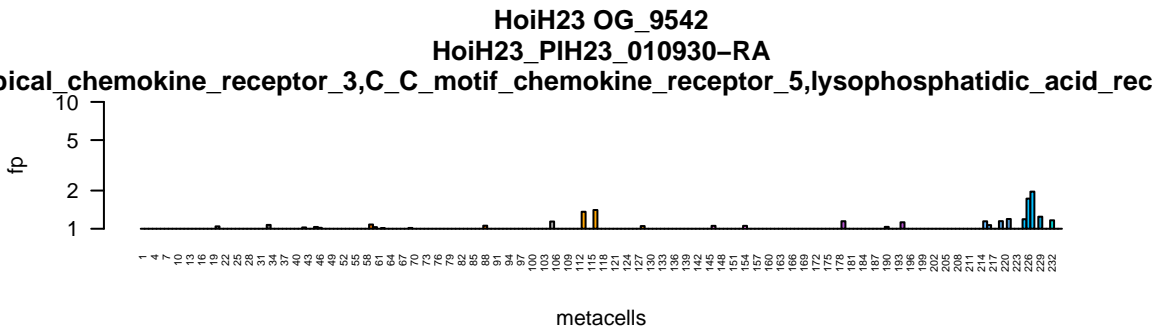
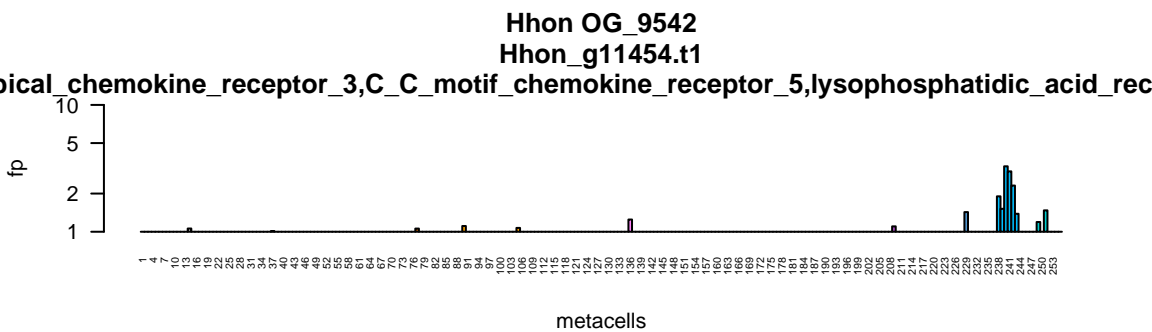
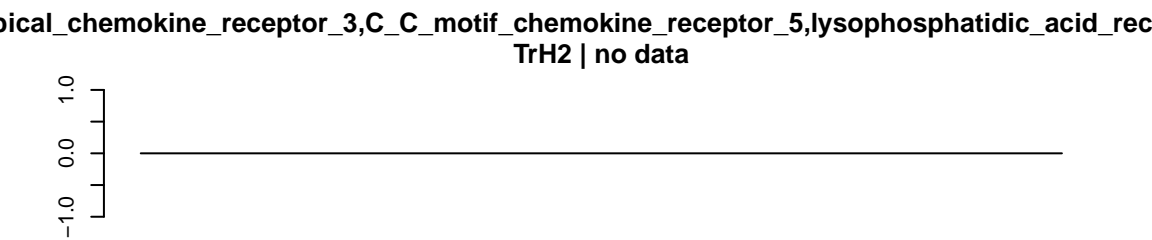
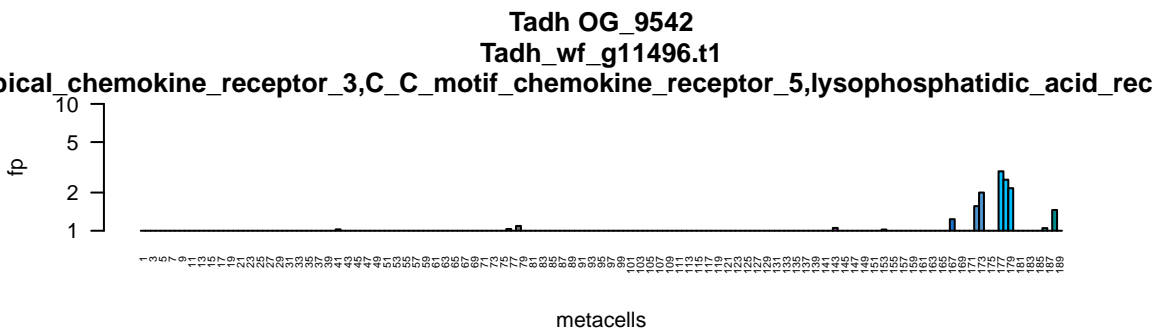




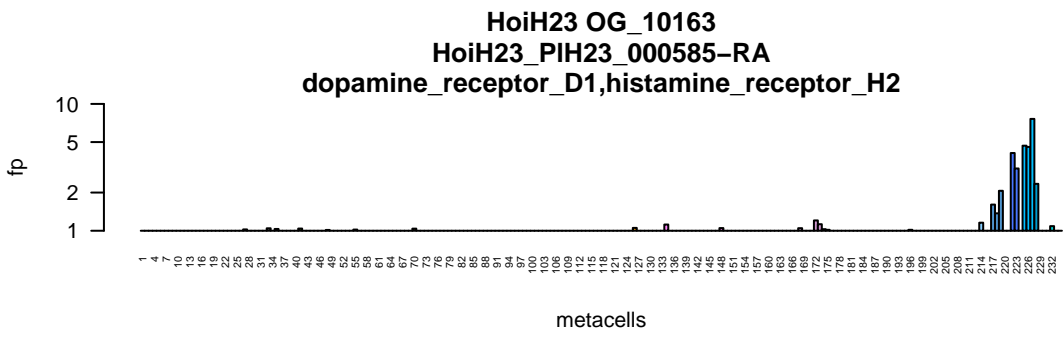
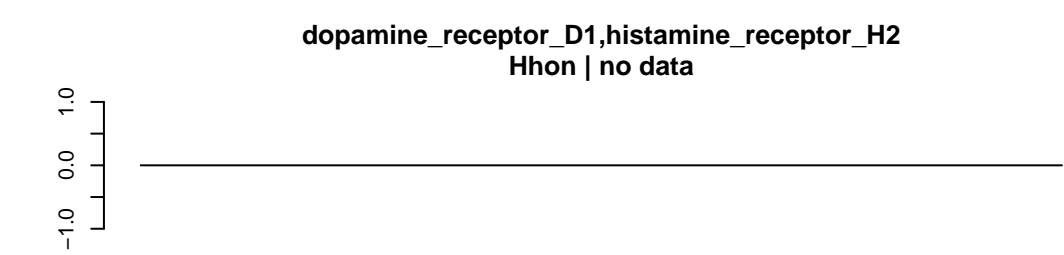
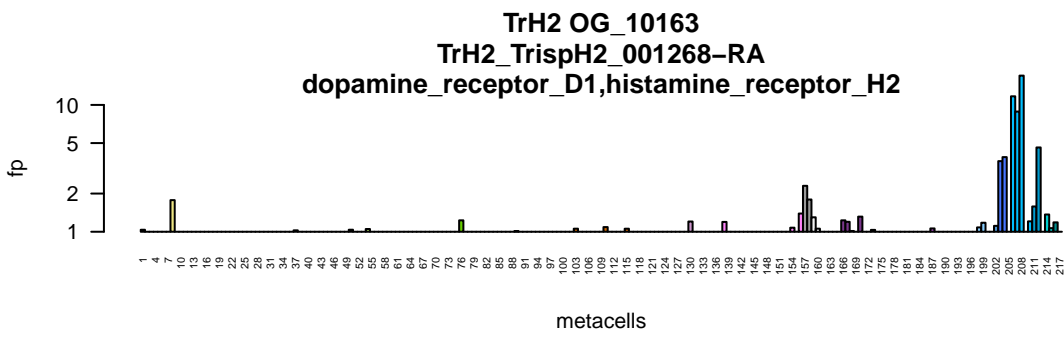
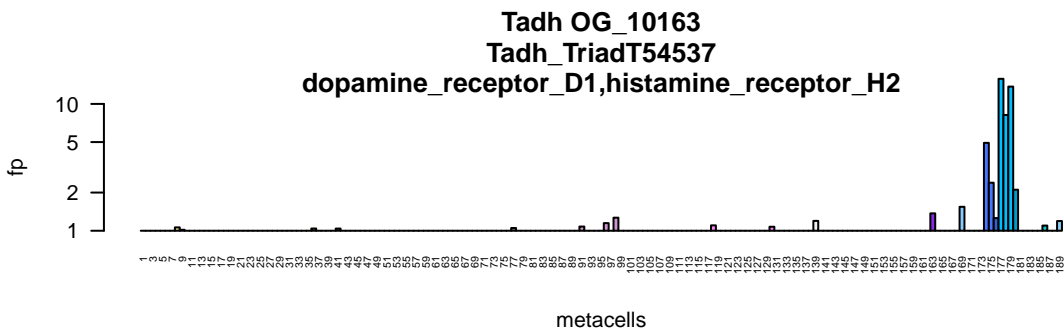




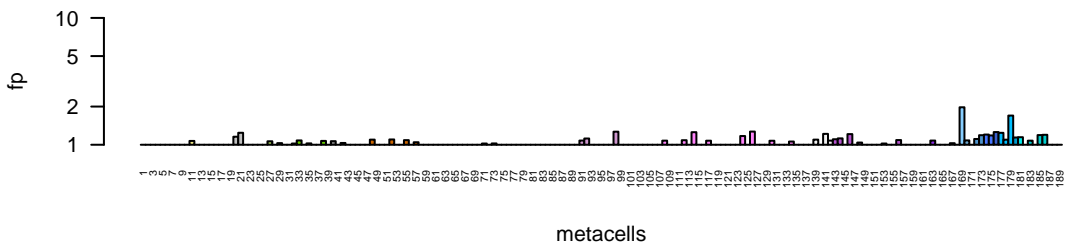






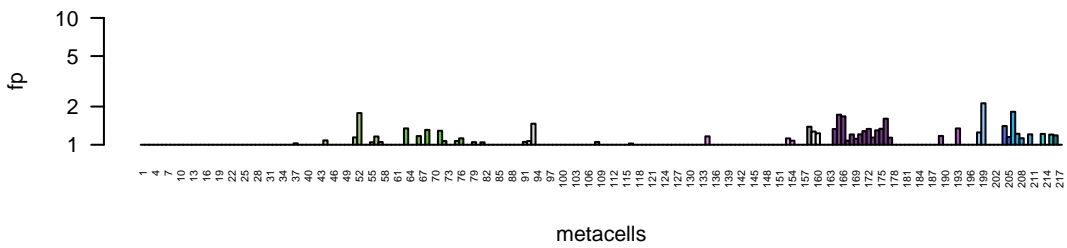


Tadh OG\_1881  
Tadh\_wf\_g2794.t1



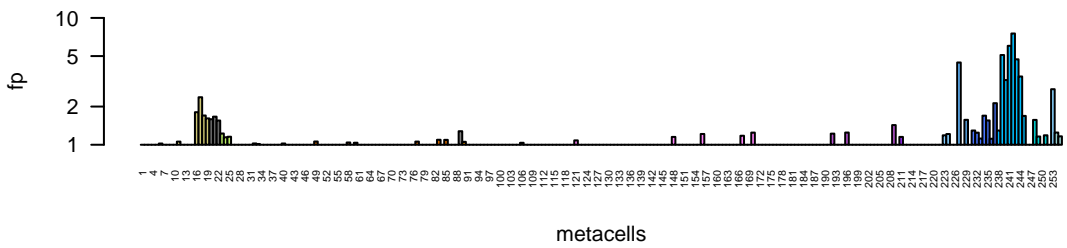
metacells

TrH2 OG\_1881  
TrH2\_TrispH2\_007804-RA



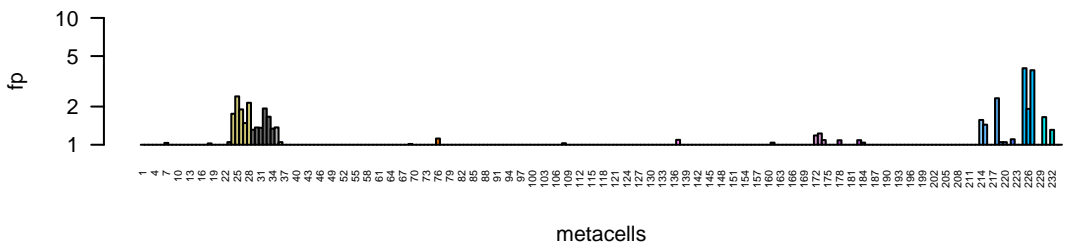
metacells

Hhon OG\_1881  
Hhon\_g06778.t1

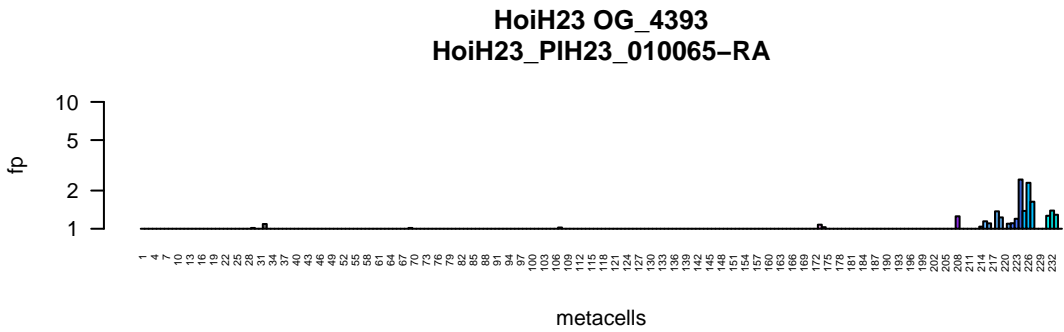
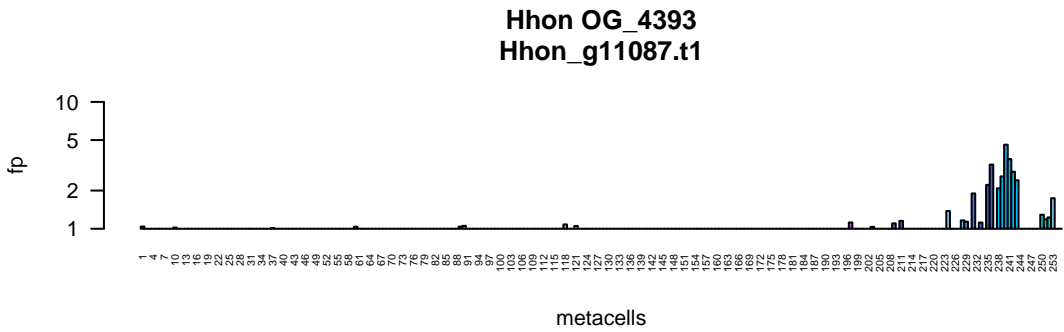
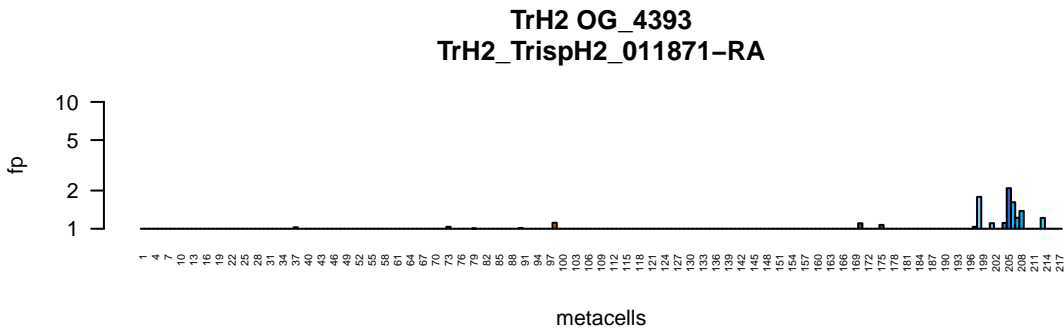
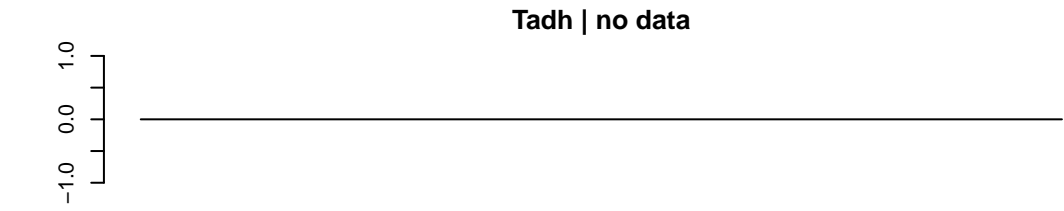


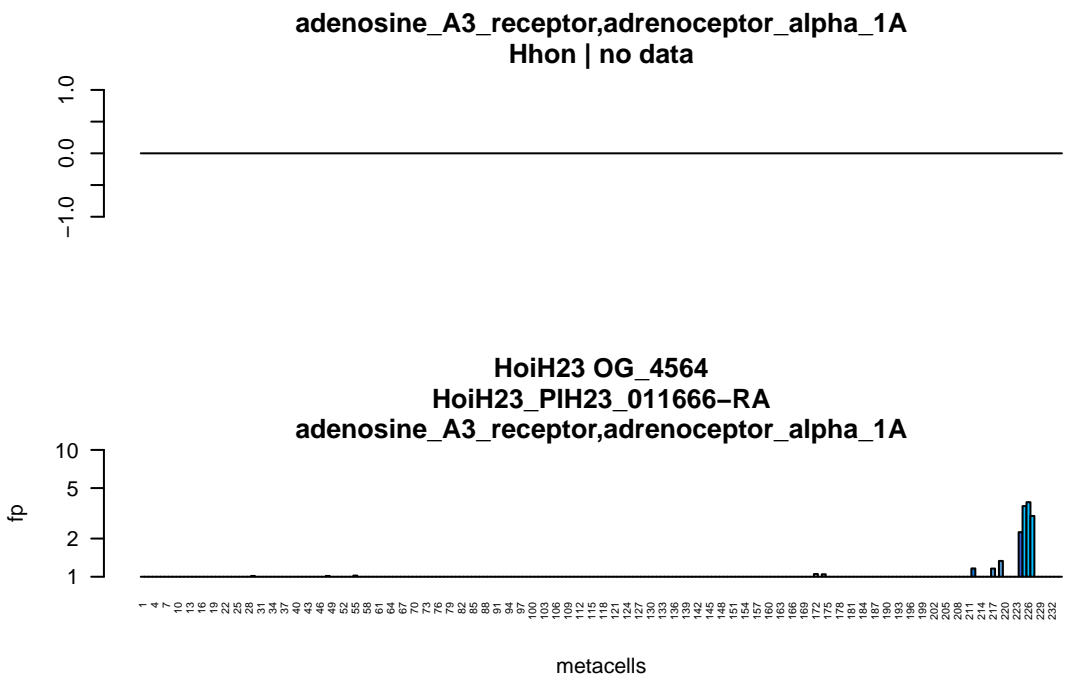
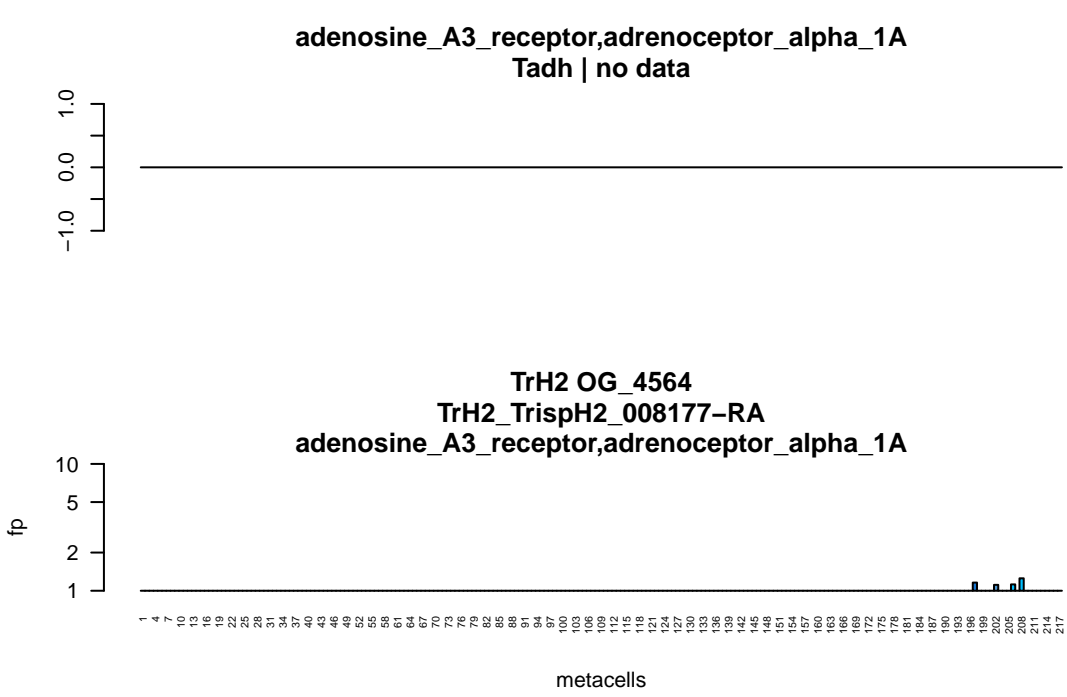
metacells

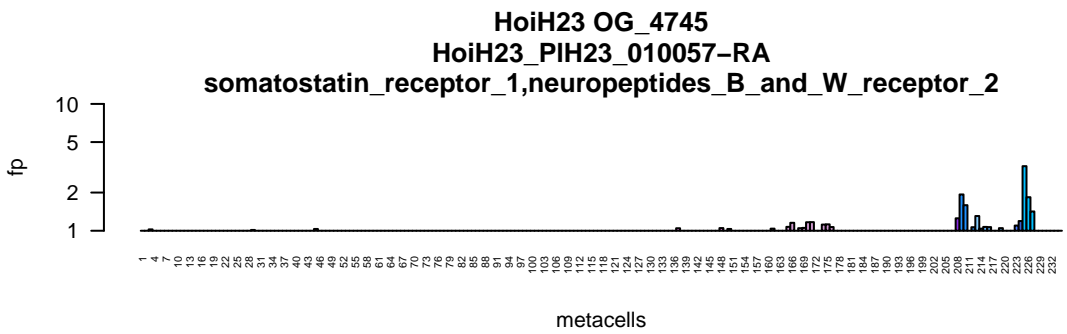
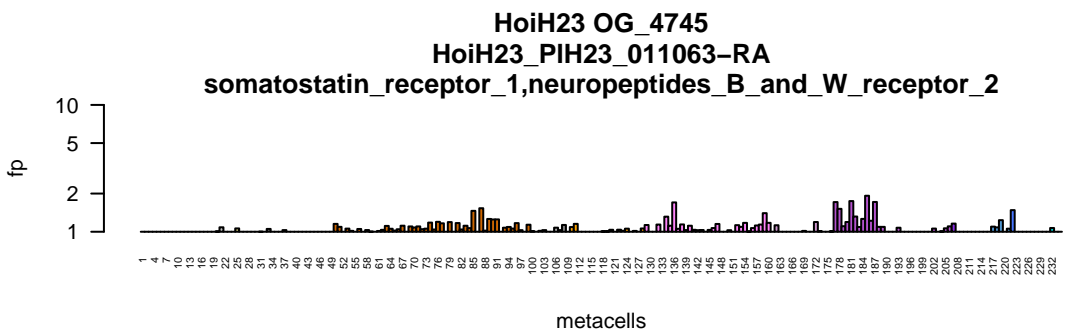
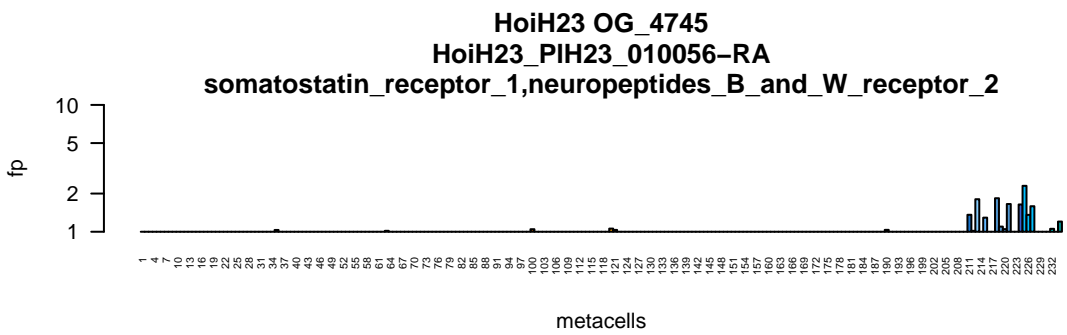
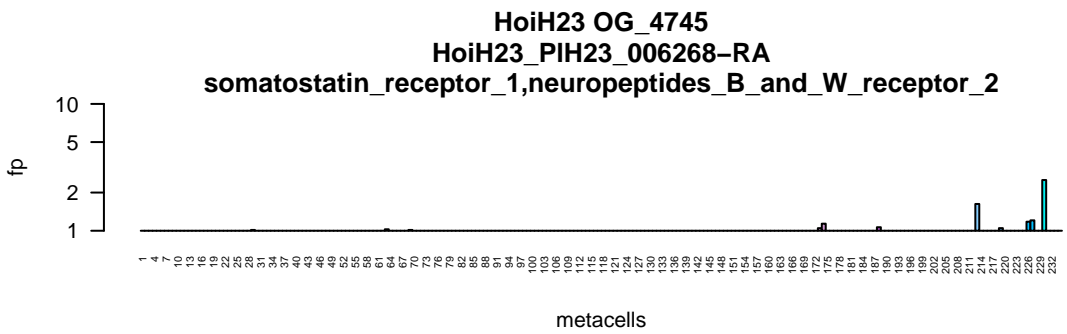
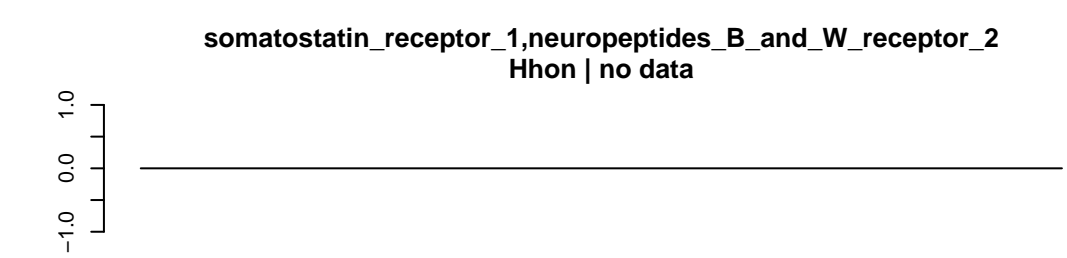
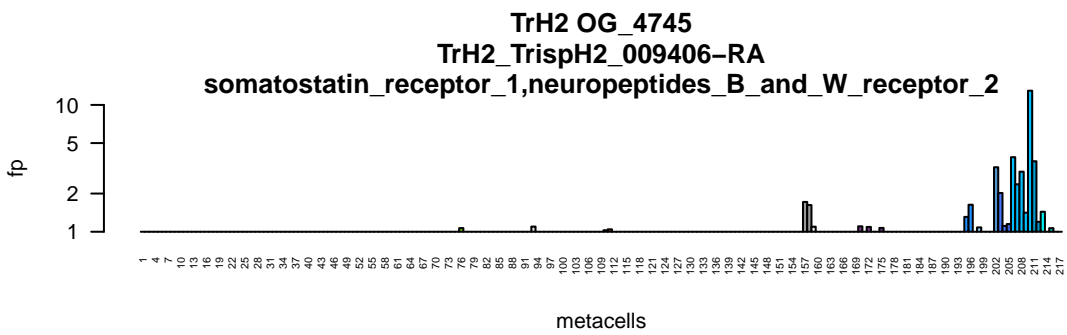
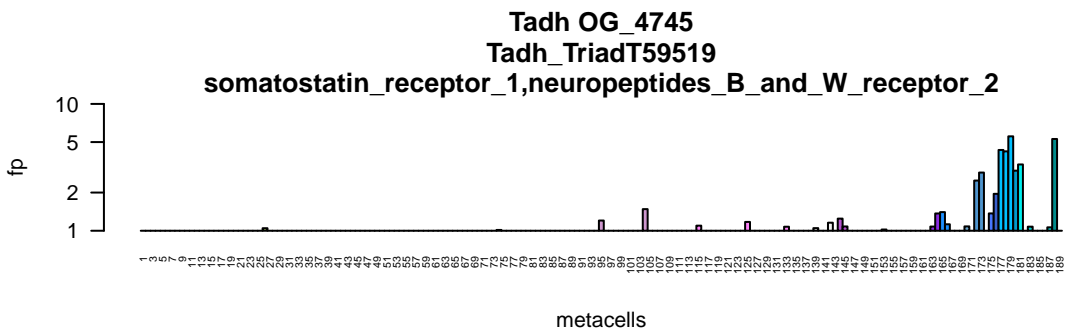
HoiH23 OG\_1881  
HoiH23\_PIH23\_005068-RA

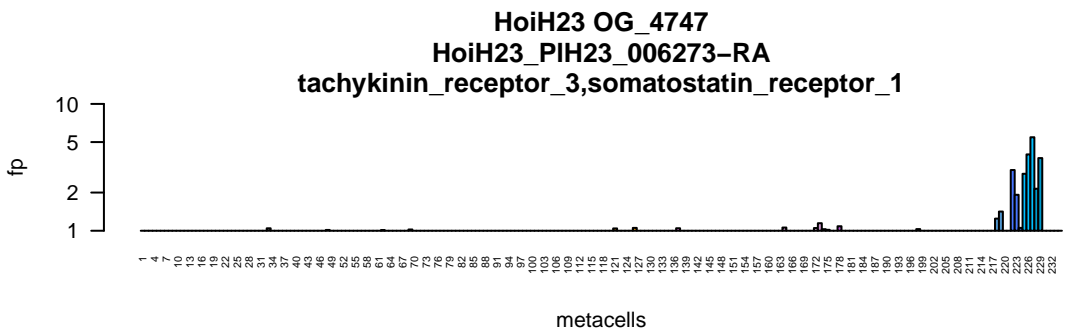
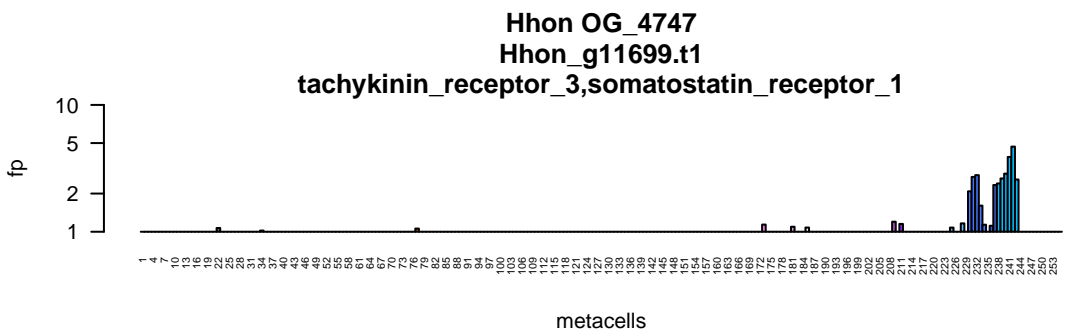
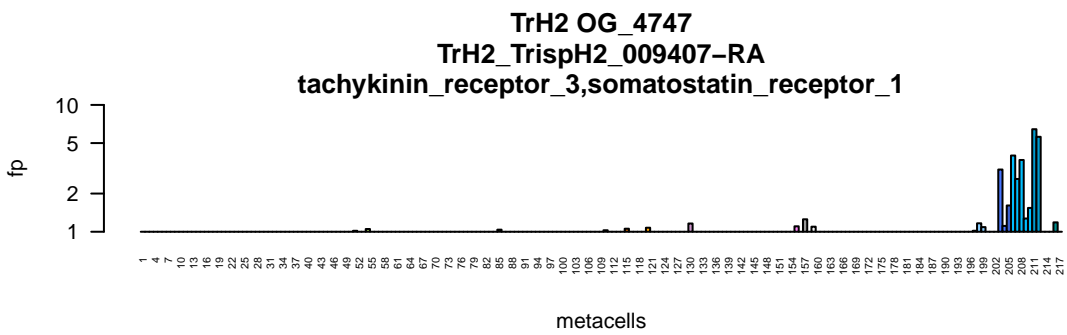
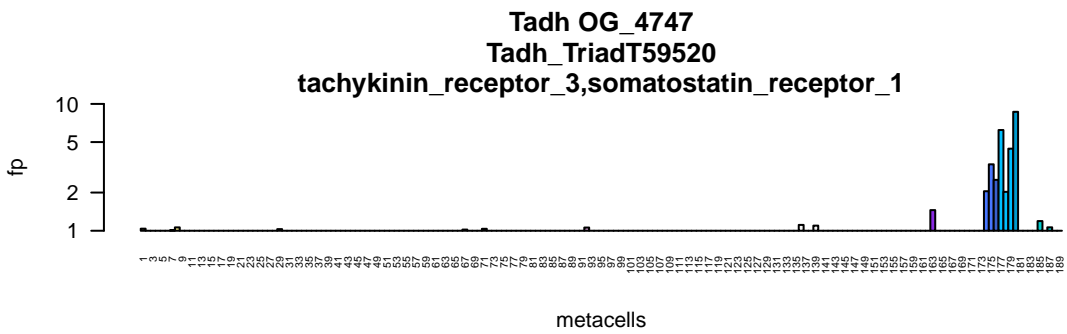


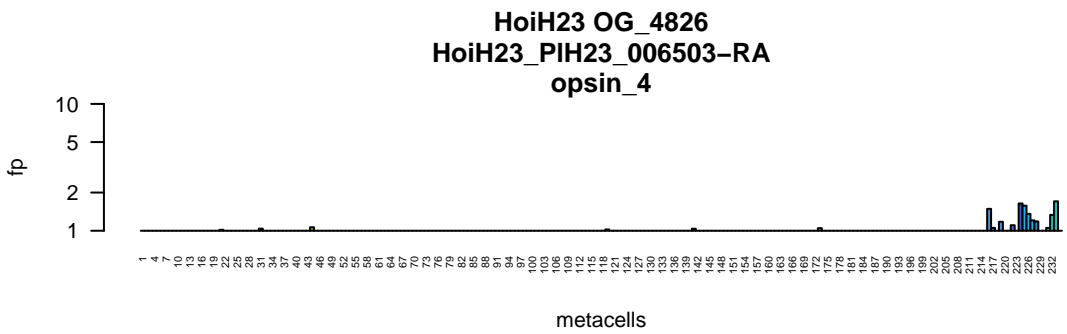
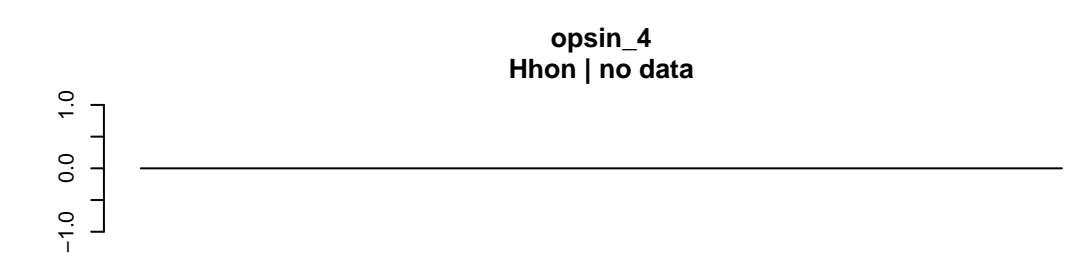
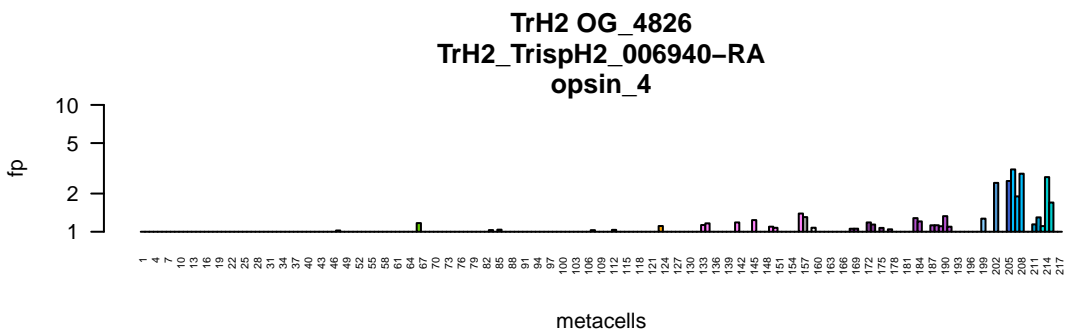
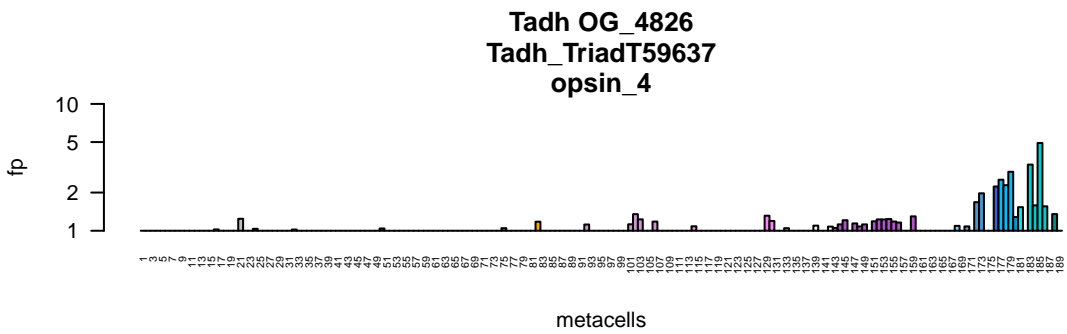
metacells

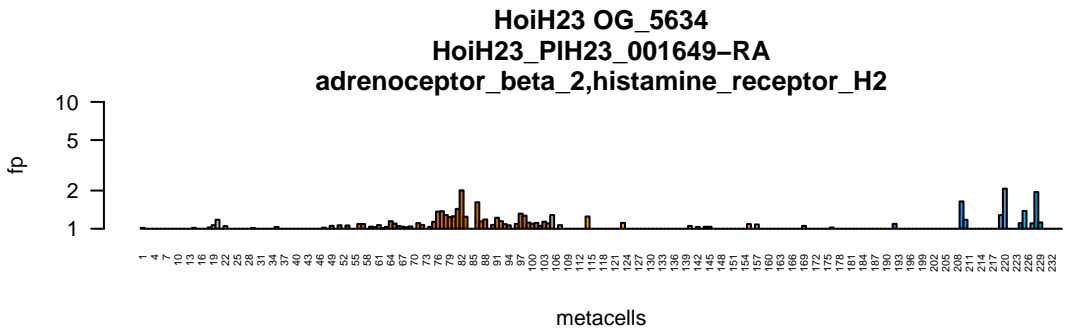
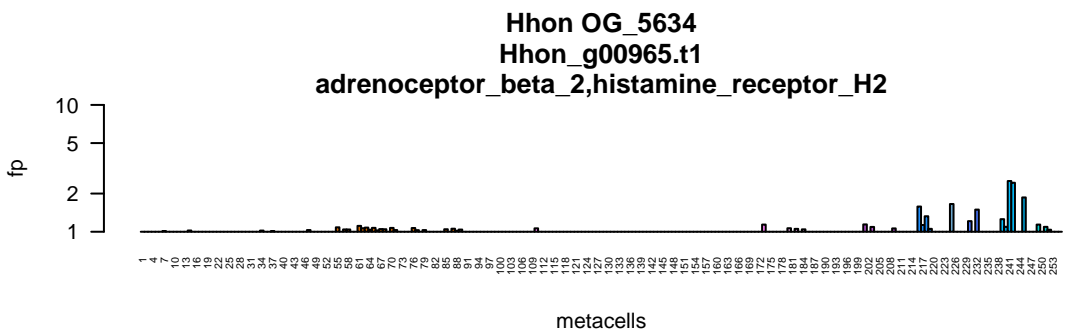
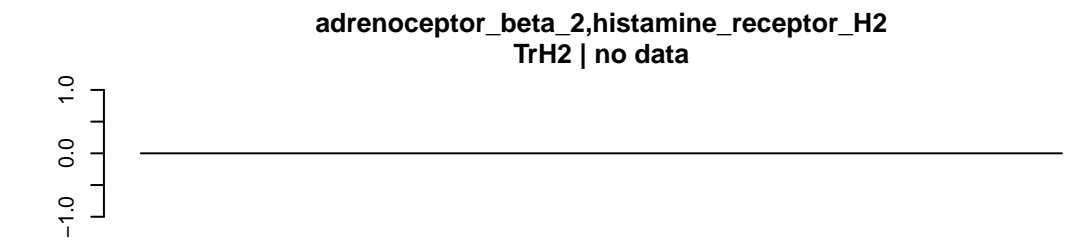
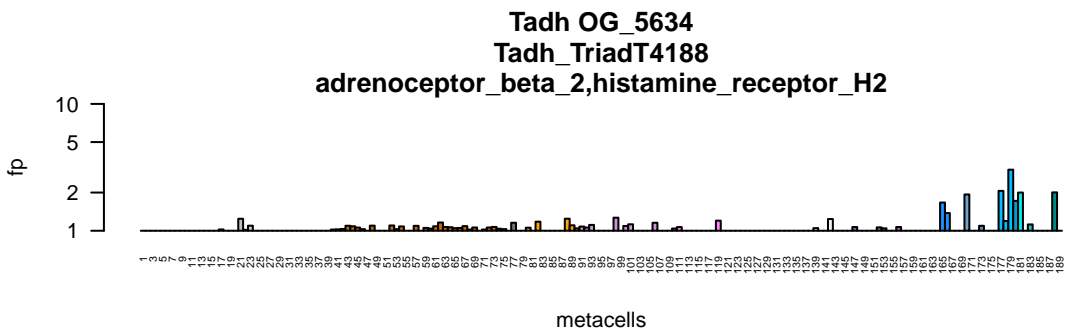








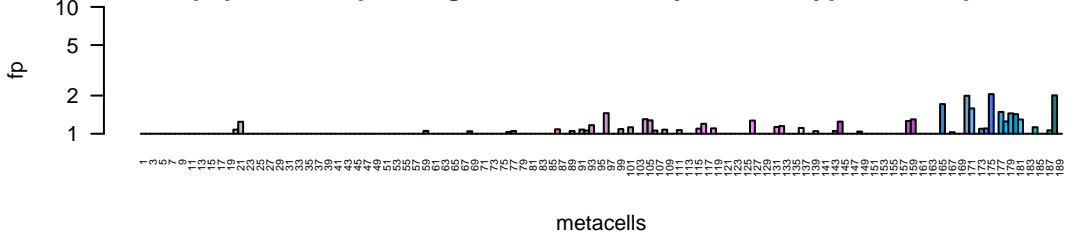






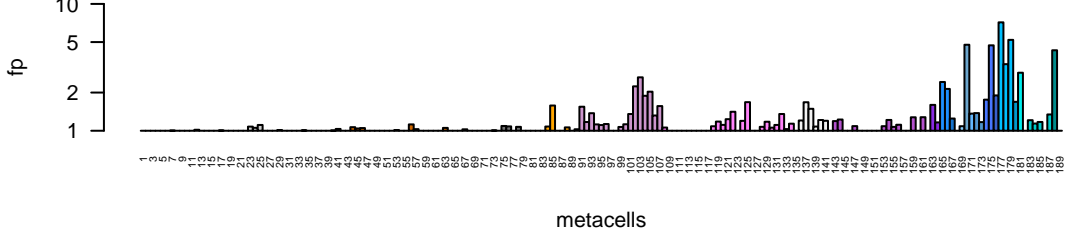
Tadh OG\_5687  
Tadh\_TriadT59259

natriuretic\_peptide\_receptor\_1,gamma\_aminobutyric\_acid\_type\_B\_receptor\_subunit\_1



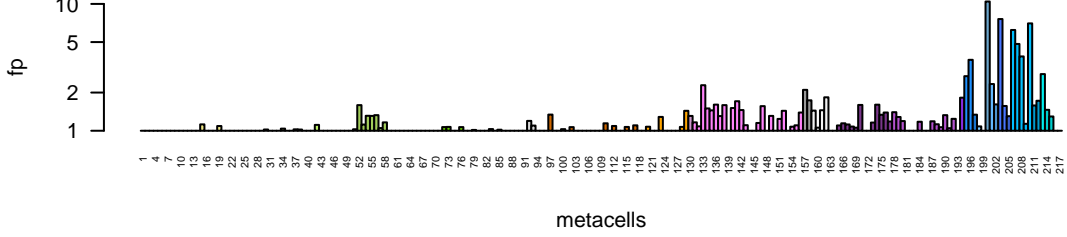
Tadh OG\_5687  
Tadh\_TriadT29353

natriuretic\_peptide\_receptor\_1,gamma\_aminobutyric\_acid\_type\_B\_receptor\_subunit\_1



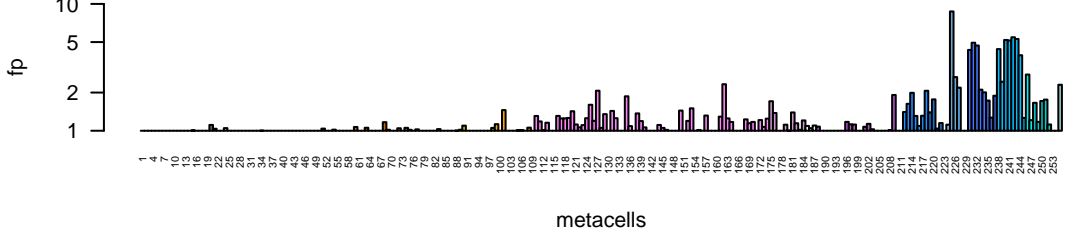
TrH2 OG\_5687  
TrH2\_TrispH2\_002051-RA

natriuretic\_peptide\_receptor\_1,gamma\_aminobutyric\_acid\_type\_B\_receptor\_subunit\_1



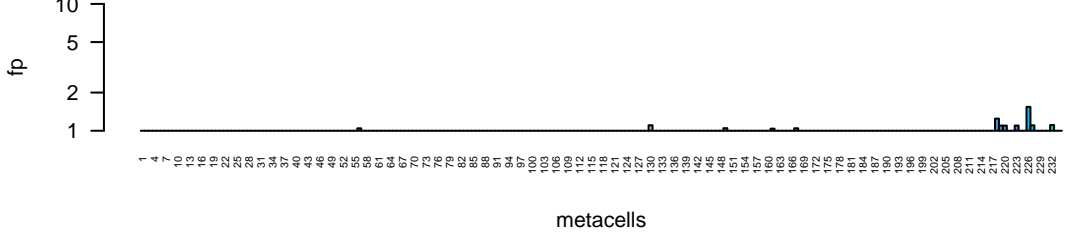
Hhon OG\_5687  
Hhon\_g00715.t1

natriuretic\_peptide\_receptor\_1,gamma\_aminobutyric\_acid\_type\_B\_receptor\_subunit\_1



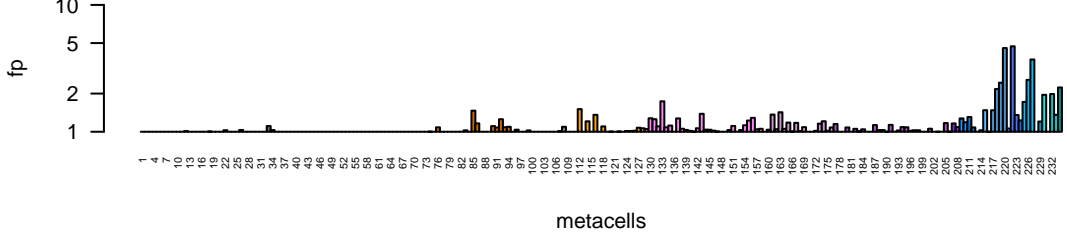
HoiH23 OG\_5687  
HoiH23\_PIH23\_009782-RA

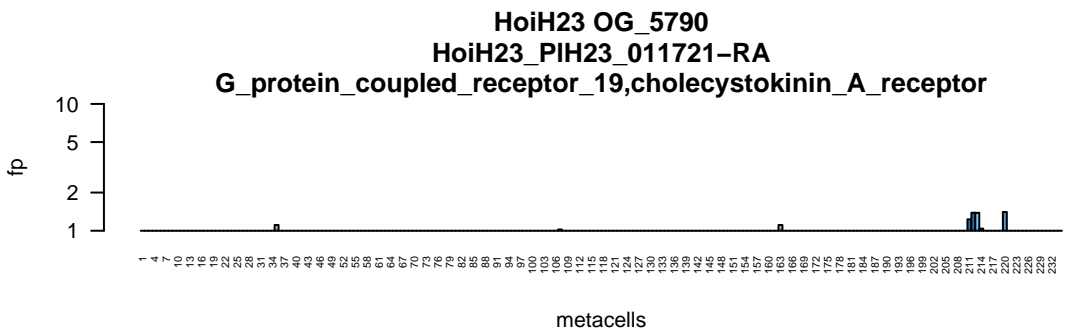
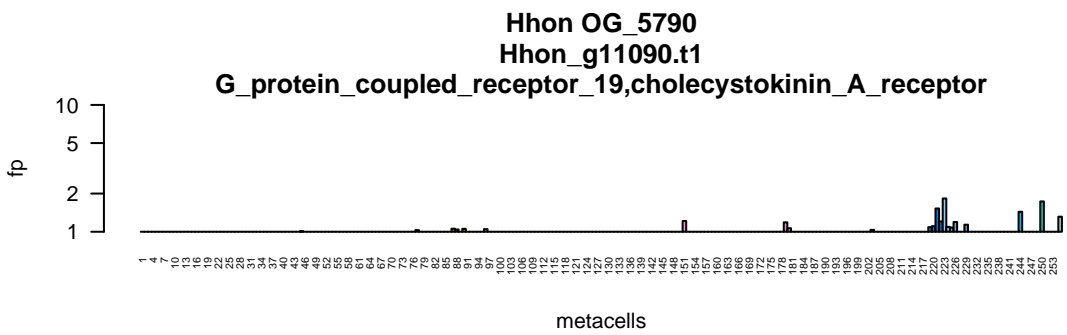
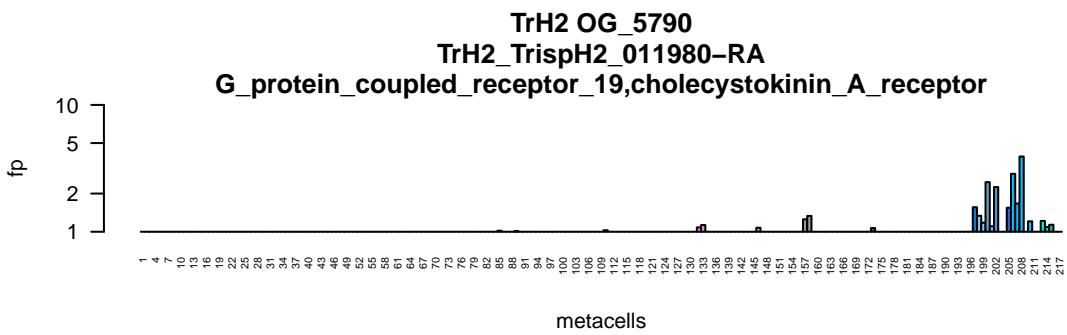
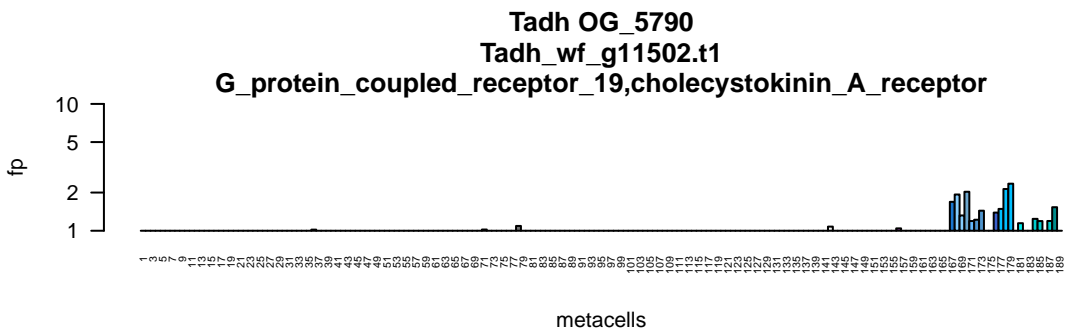
natriuretic\_peptide\_receptor\_1,gamma\_aminobutyric\_acid\_type\_B\_receptor\_subunit\_1



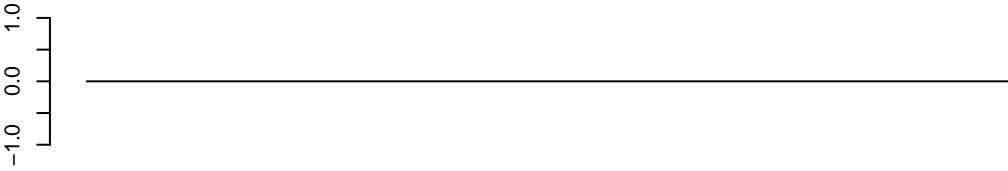
HoiH23 OG\_5687  
HoiH23\_PIH23\_009783-RA

natriuretic\_peptide\_receptor\_1,gamma\_aminobutyric\_acid\_type\_B\_receptor\_subunit\_1

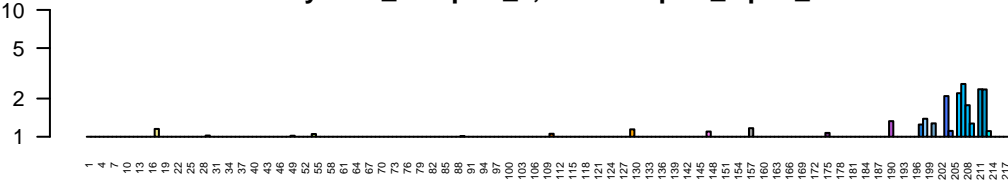




tachykinin\_receptor\_1,adrenoceptor\_alpha\_1B  
Tadh | no data

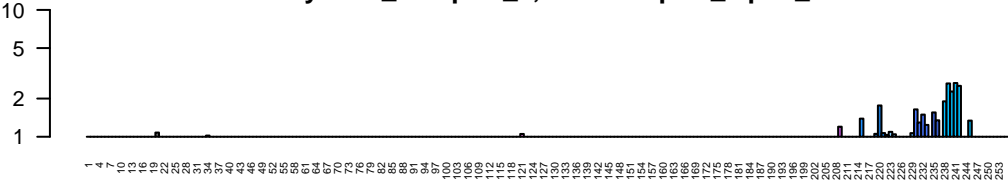


TrH2 OG\_5841  
TrH2\_TrispH2\_002792-RA  
tachykinin\_receptor\_1,adrenoceptor\_alpha\_1B



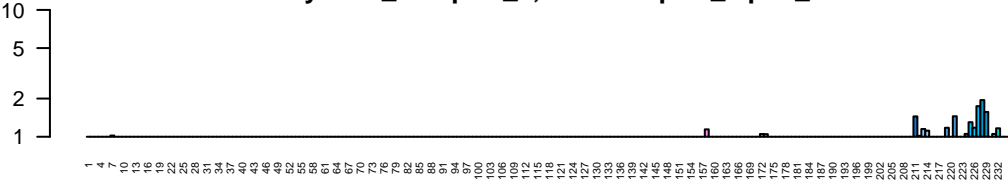
metacells

Hhon OG\_5841  
Hhon\_g00396.t1  
tachykinin\_receptor\_1,adrenoceptor\_alpha\_1B



metacells

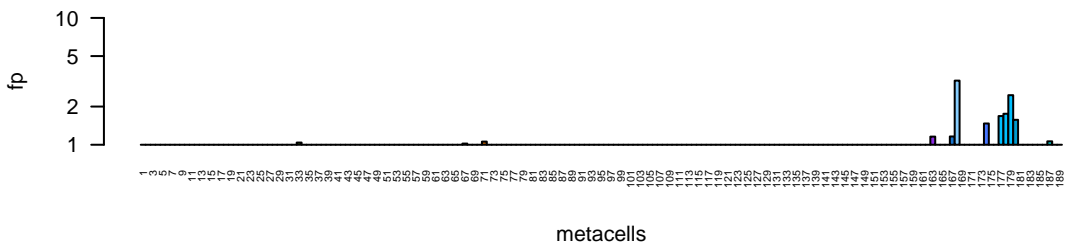
HoiH23 OG\_5841  
HoiH23\_PIH23\_001771-RA  
tachykinin\_receptor\_1,adrenoceptor\_alpha\_1B



metacells

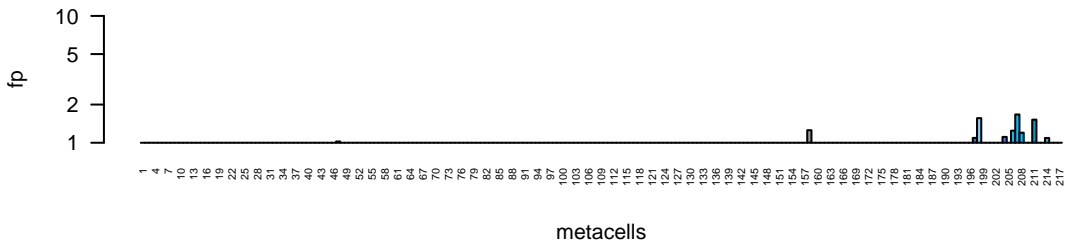
Tadh OG\_6120

Tadh\_TriadT60881



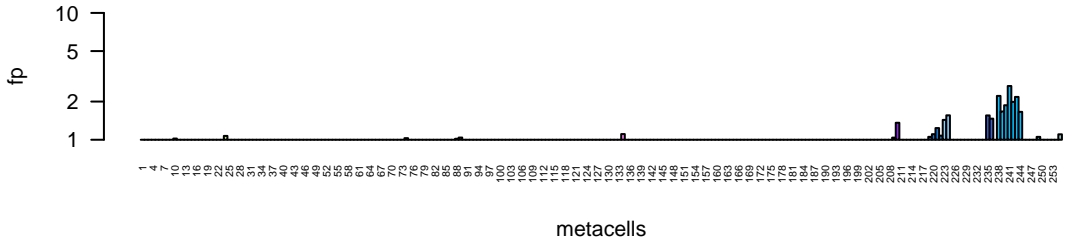
TrH2 OG\_6120

TrH2\_TrispH2\_008845-RA



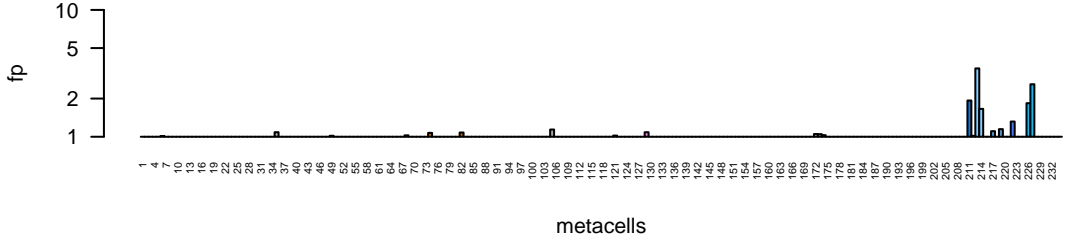
Hhon OG\_6120

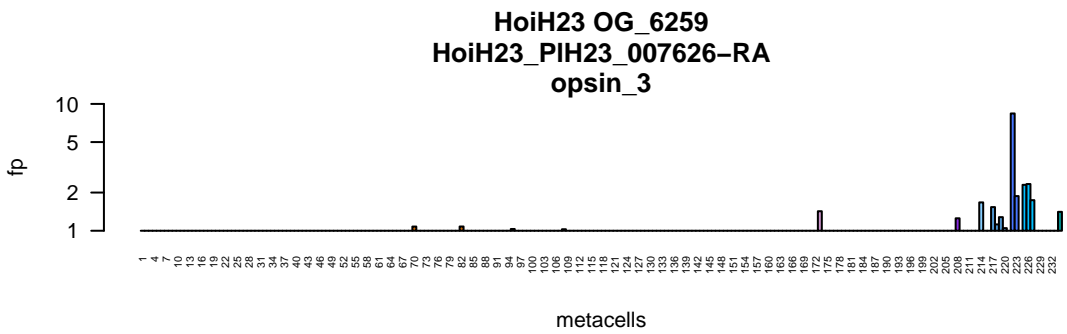
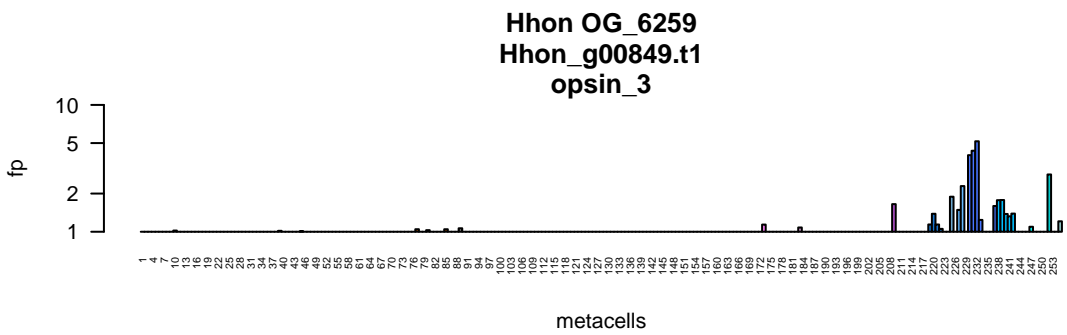
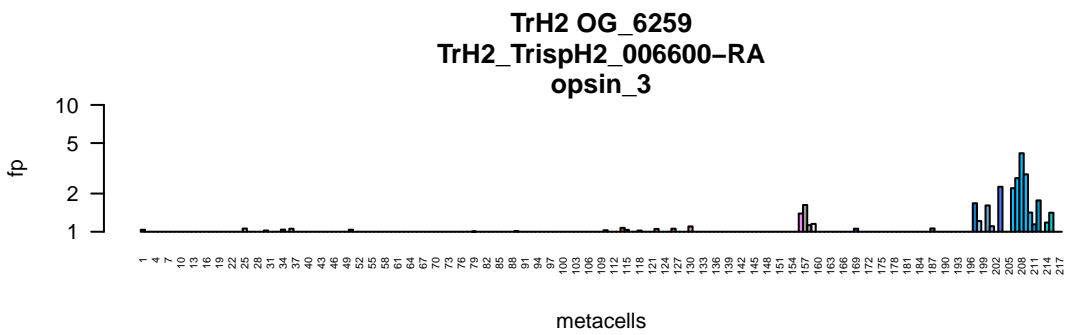
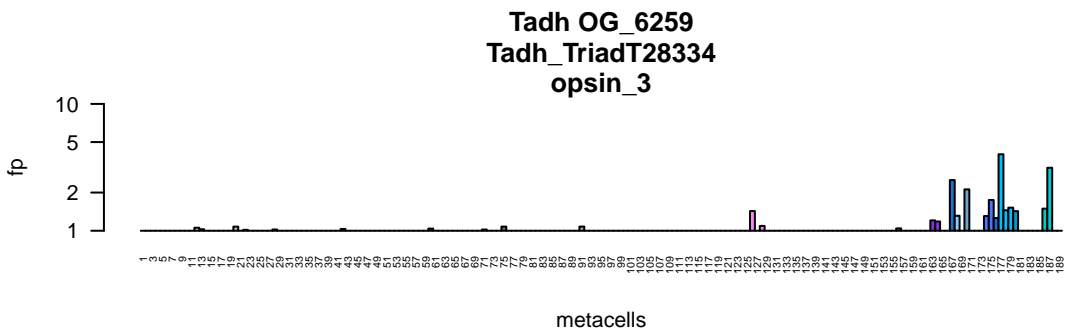
Hhon\_g10766.t1

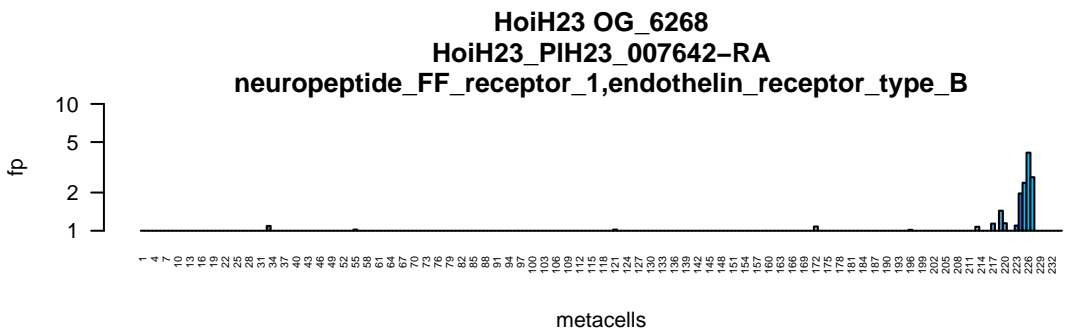
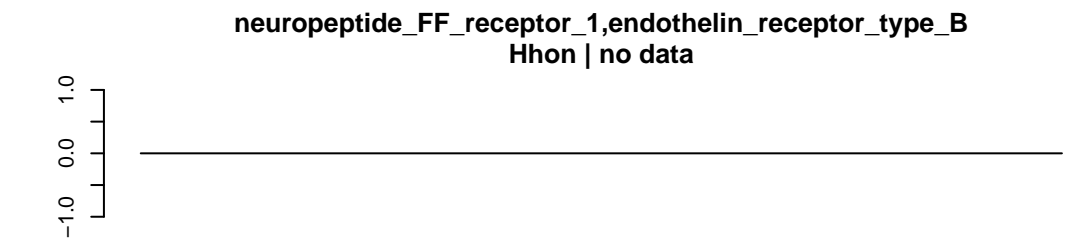
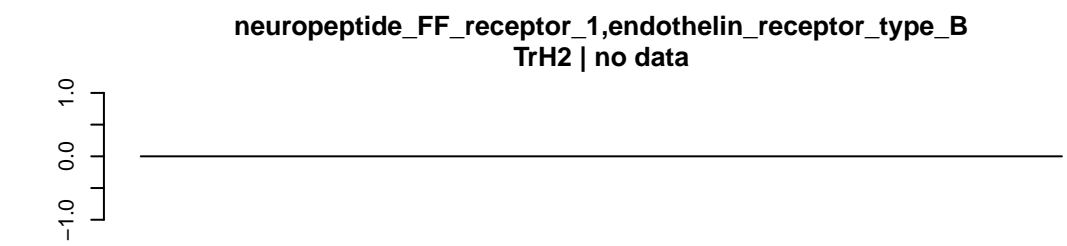
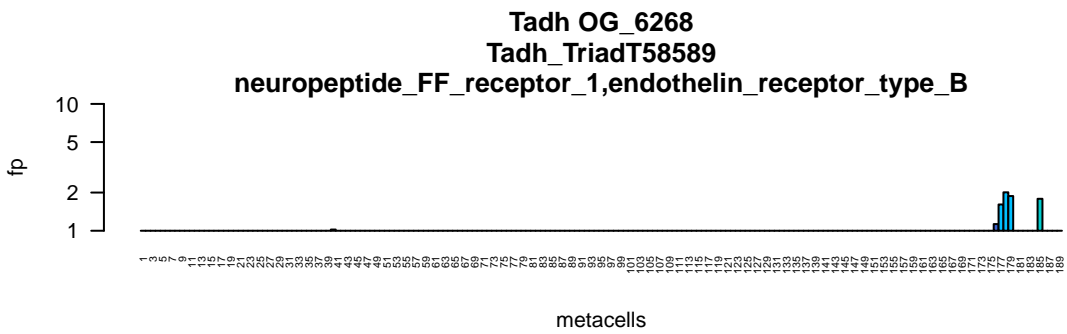


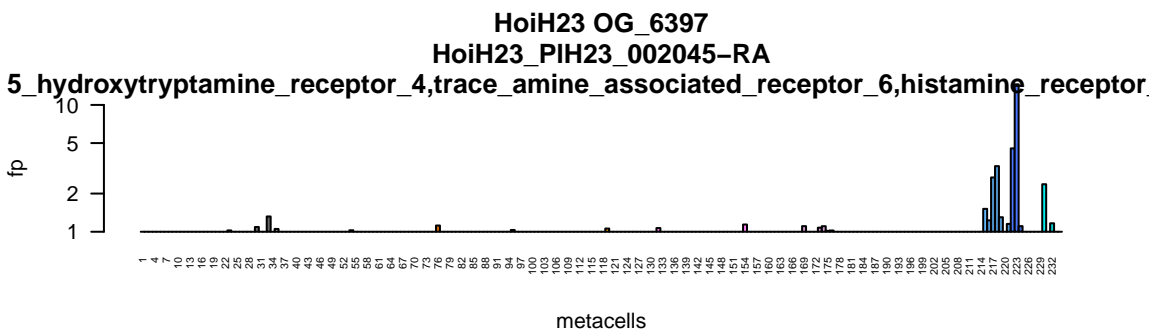
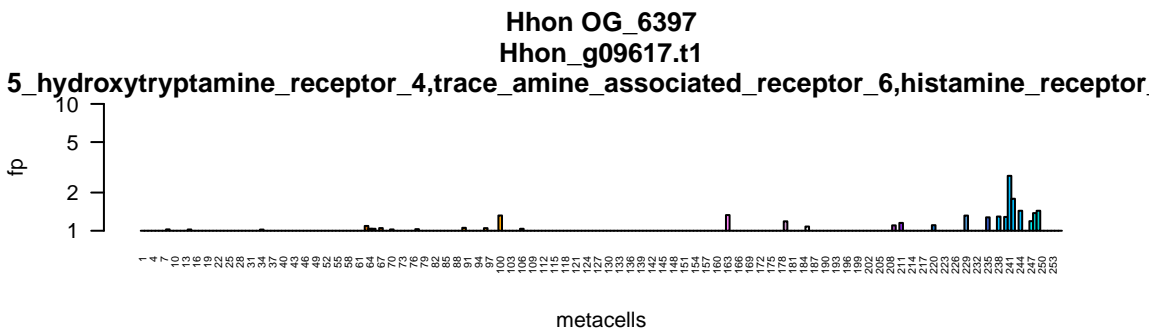
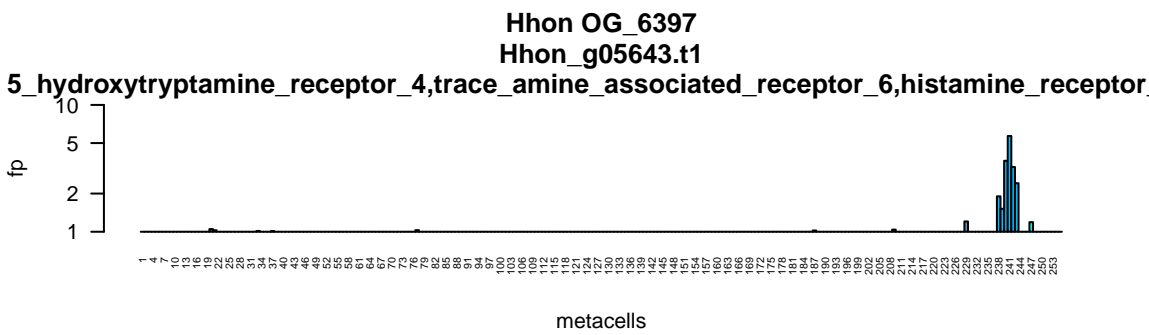
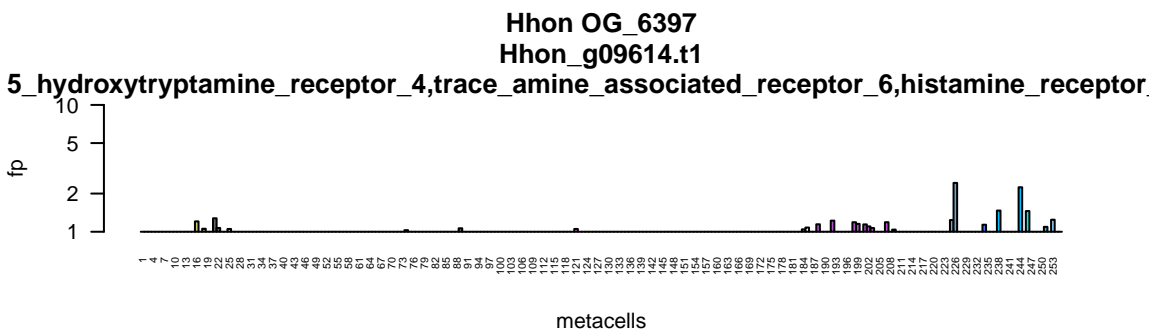
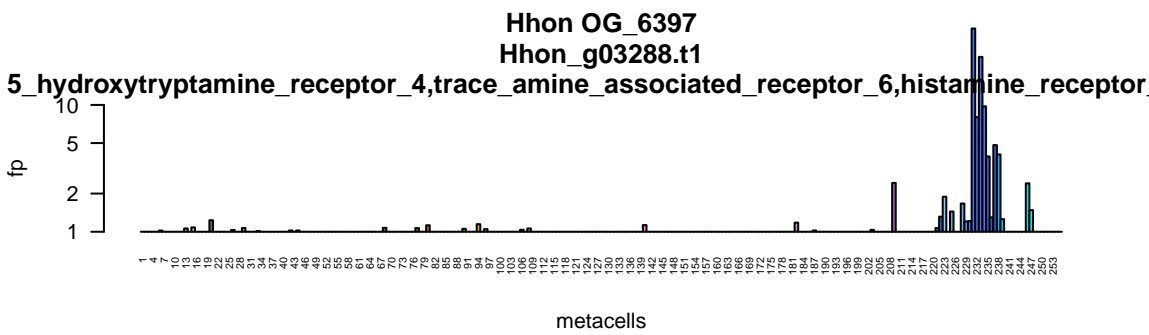
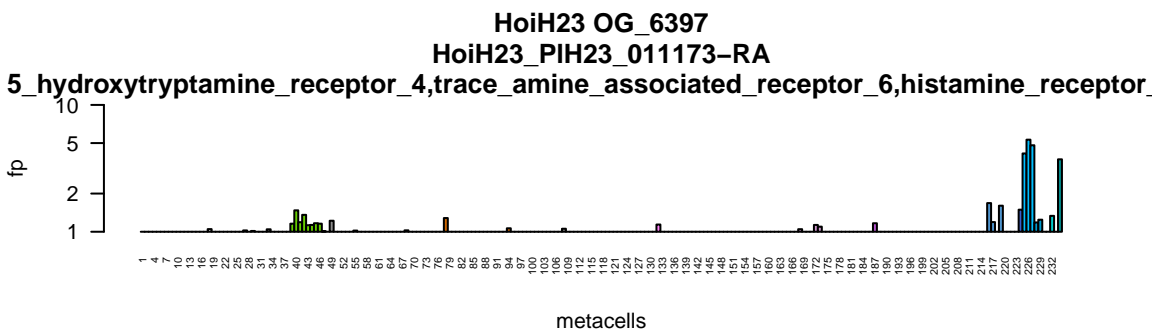
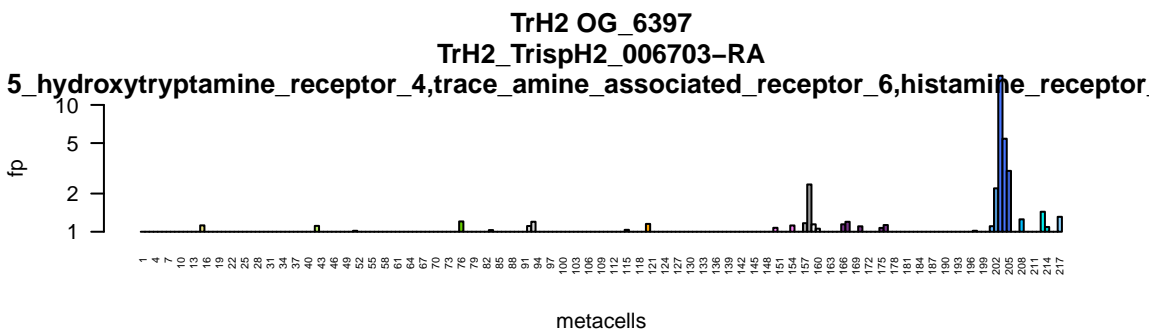
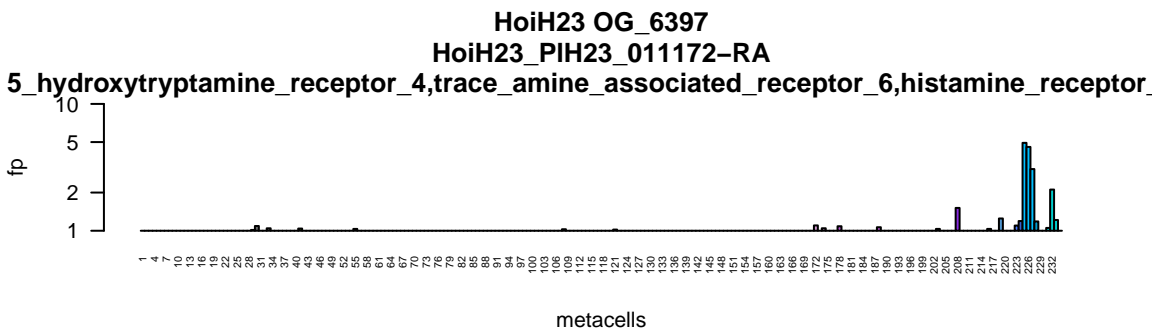
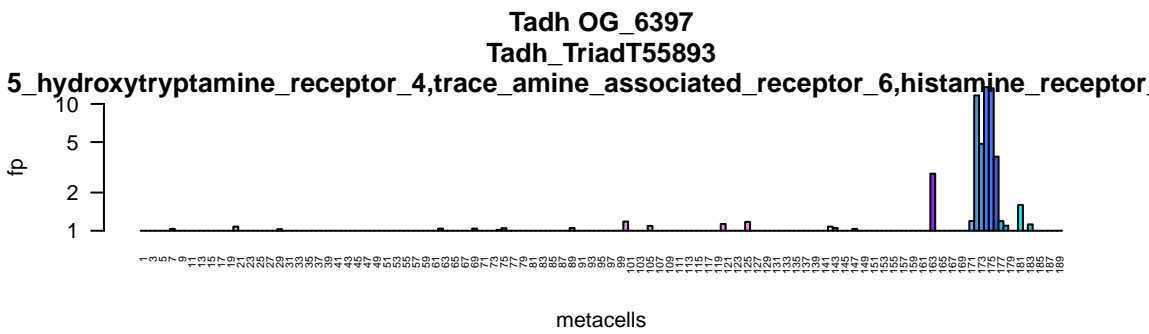
HoiH23 OG\_6120

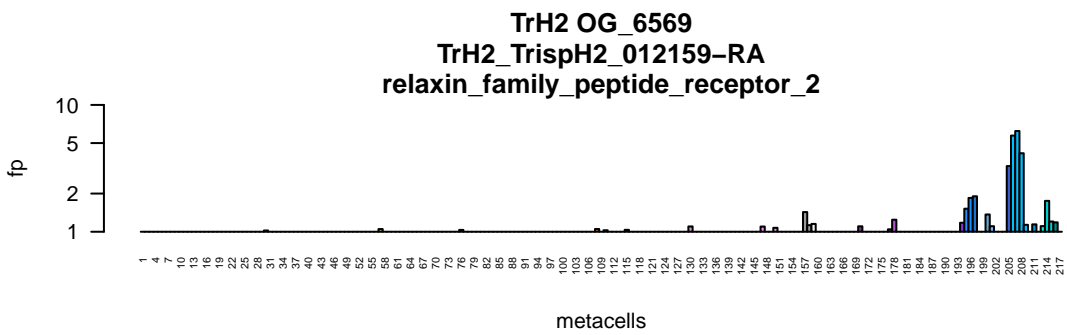
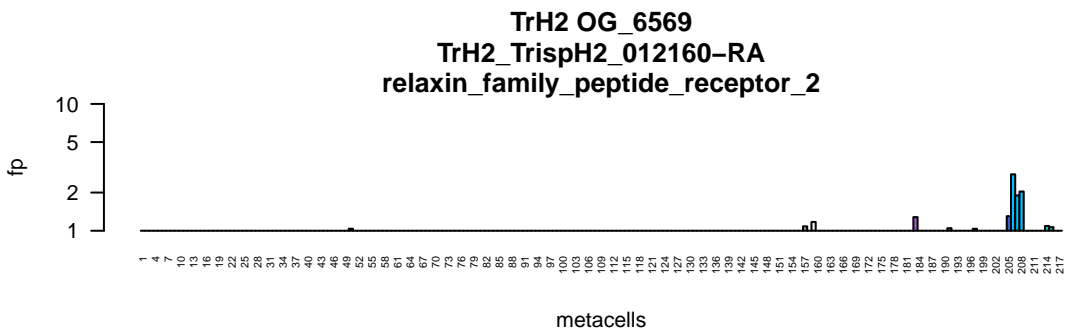
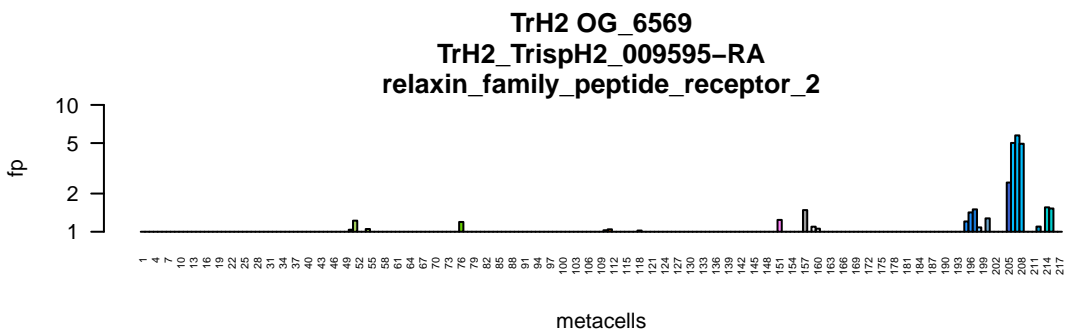
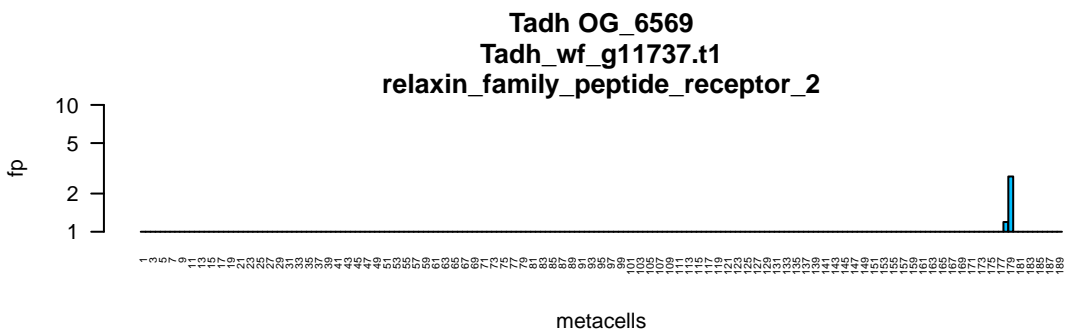
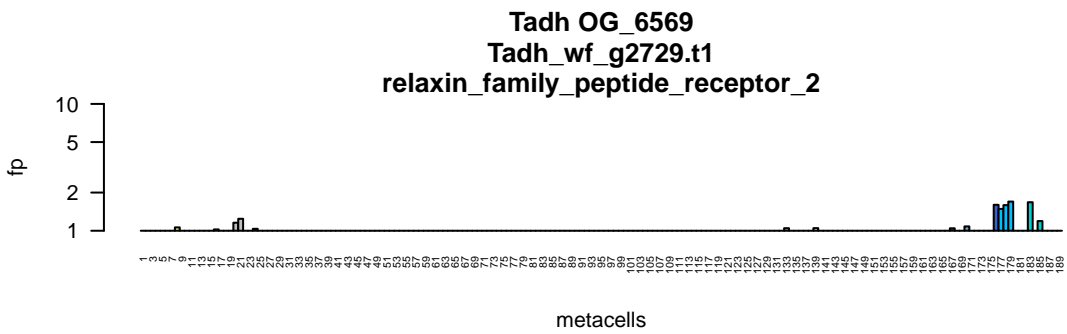
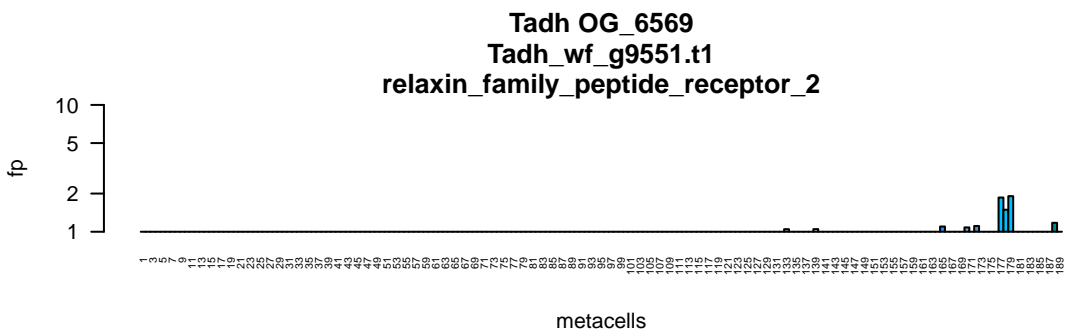
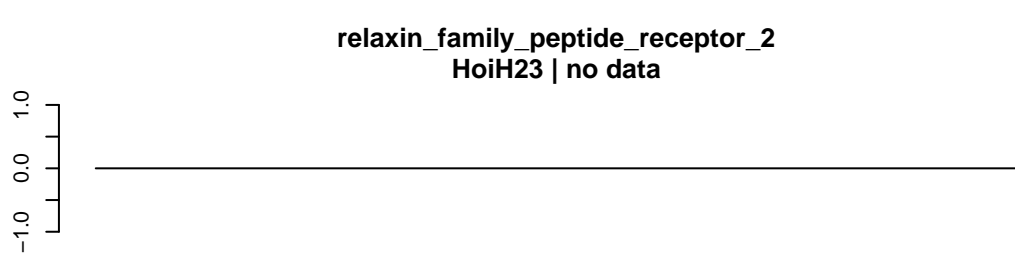
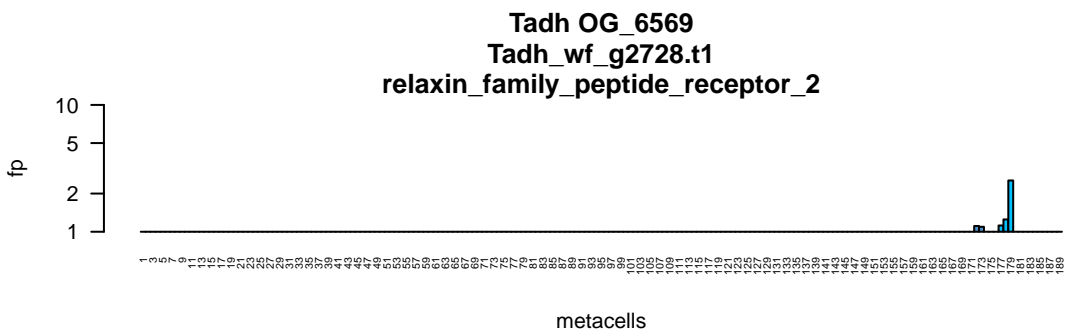
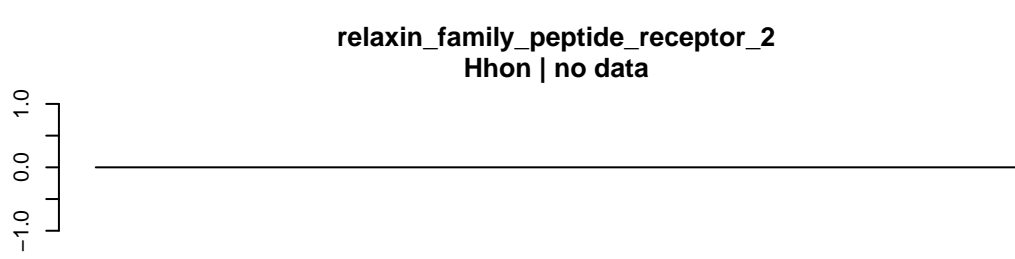
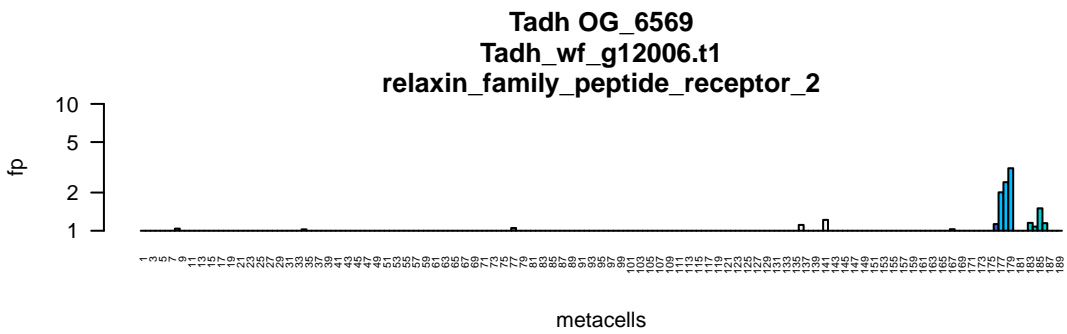
HoiH23\_PIH23\_008464-RA



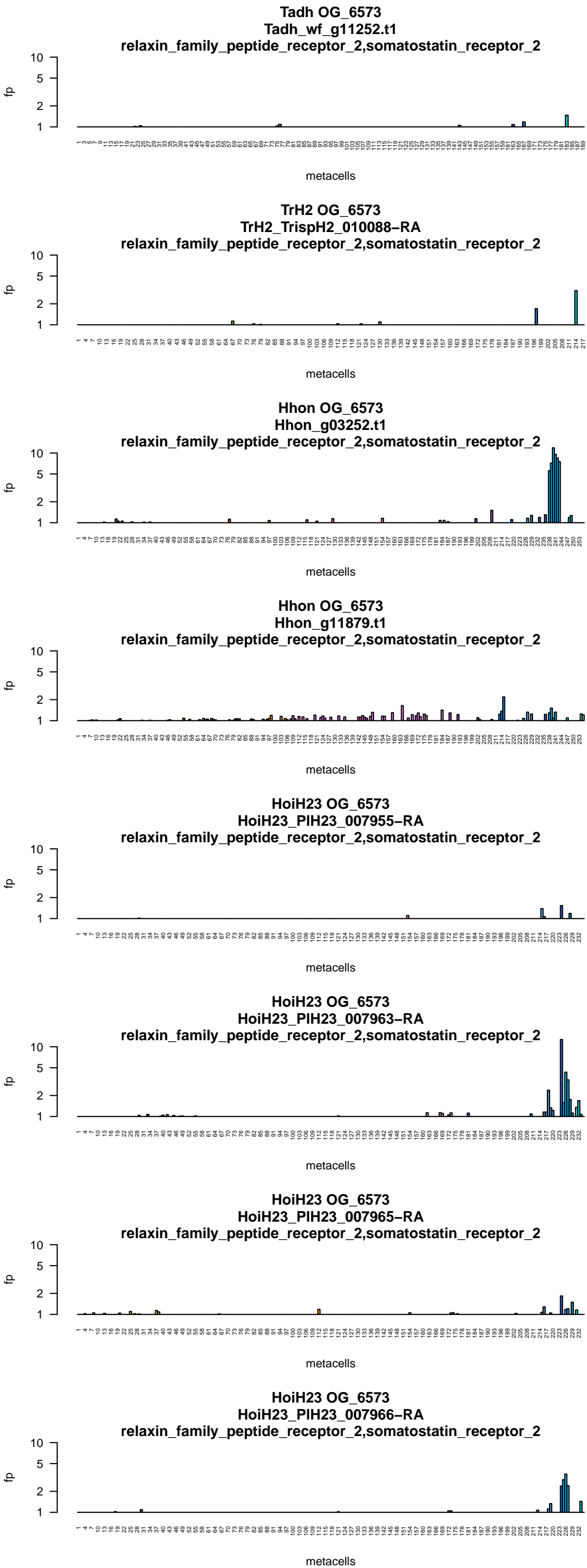


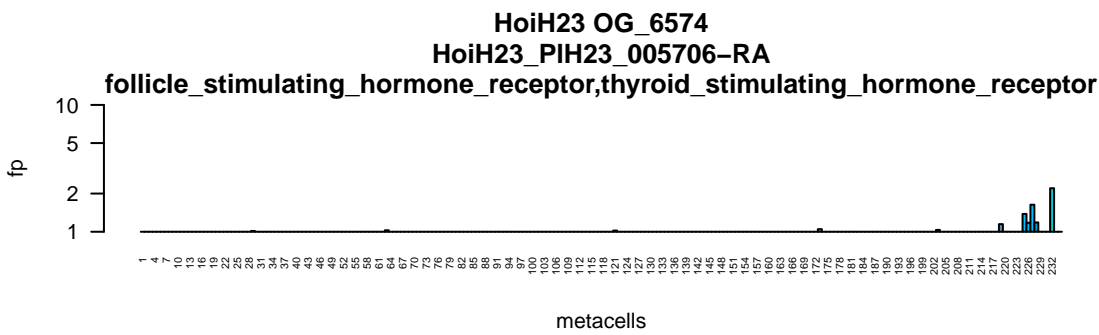
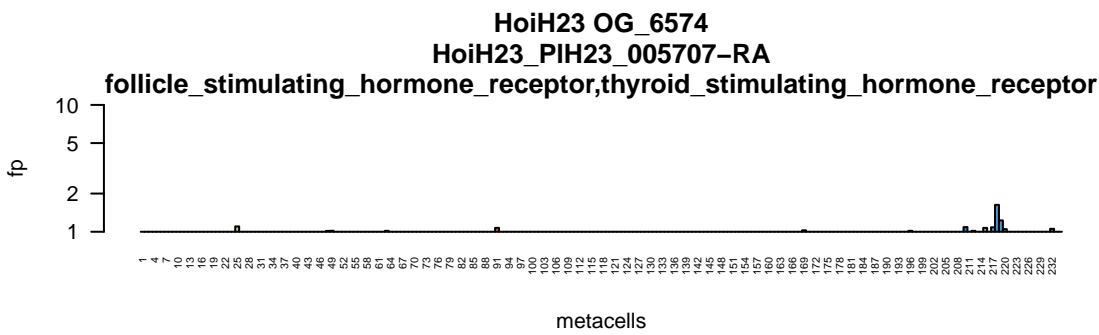
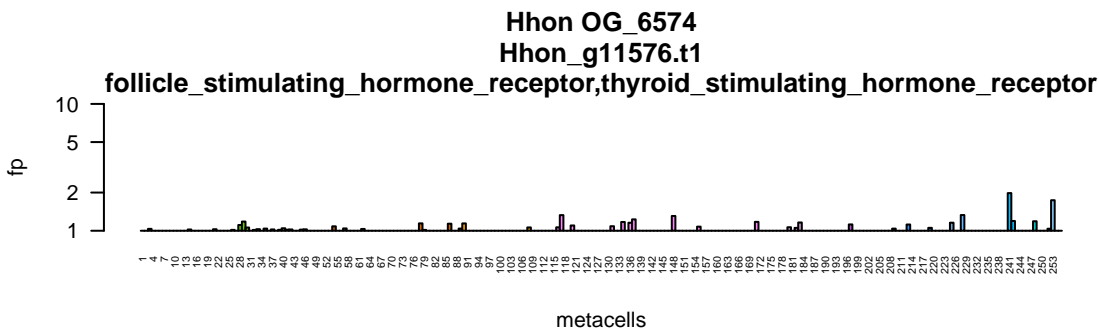
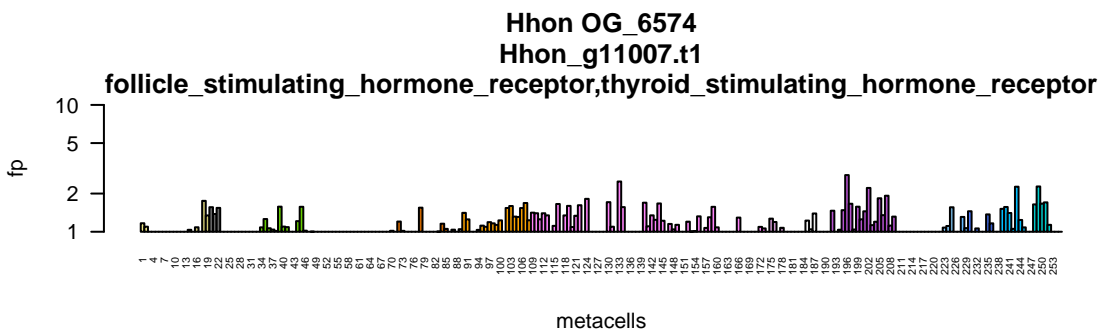
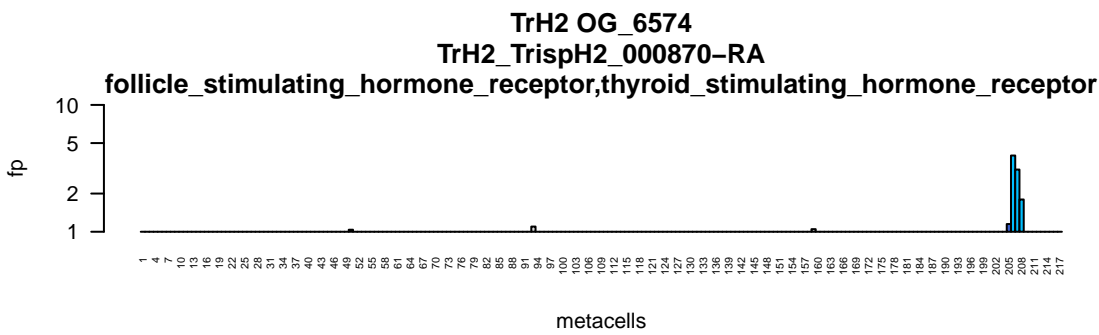
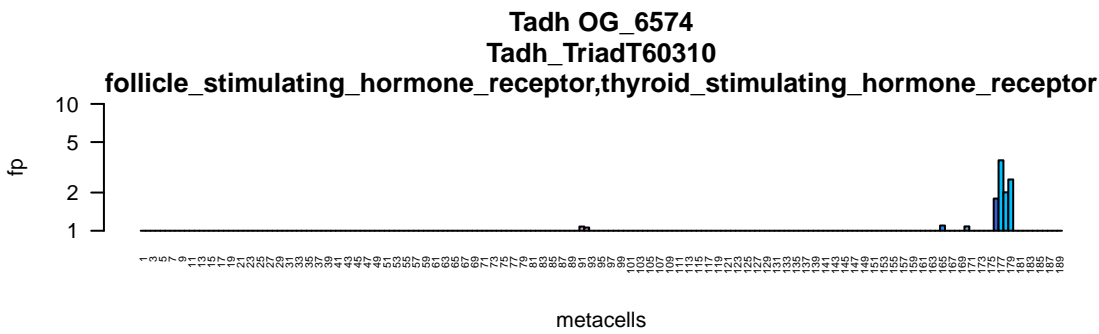


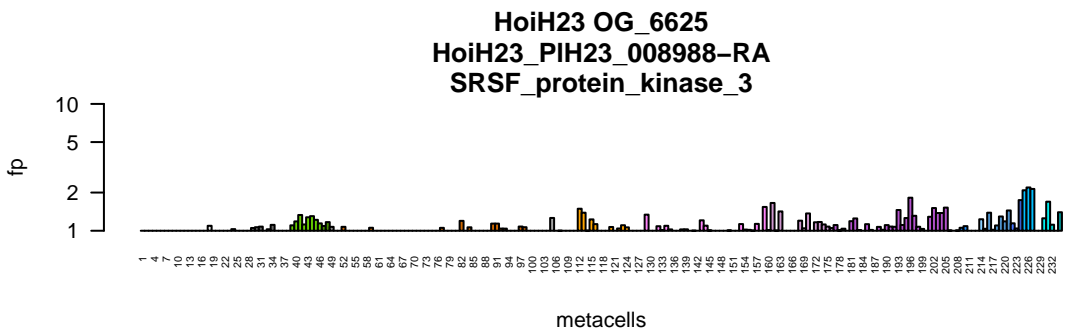
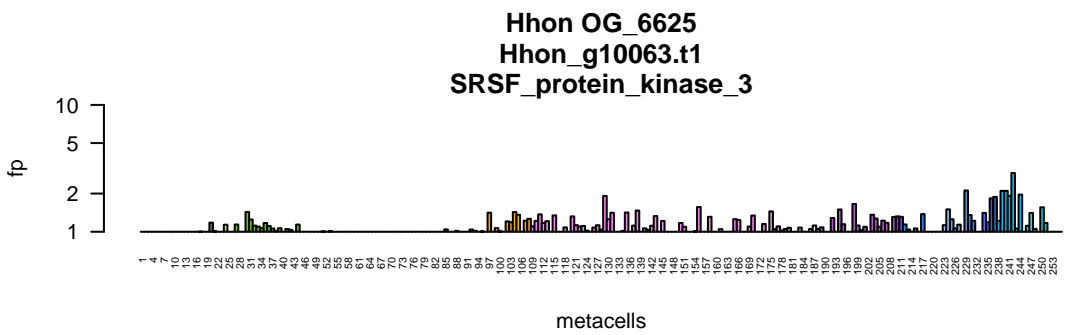
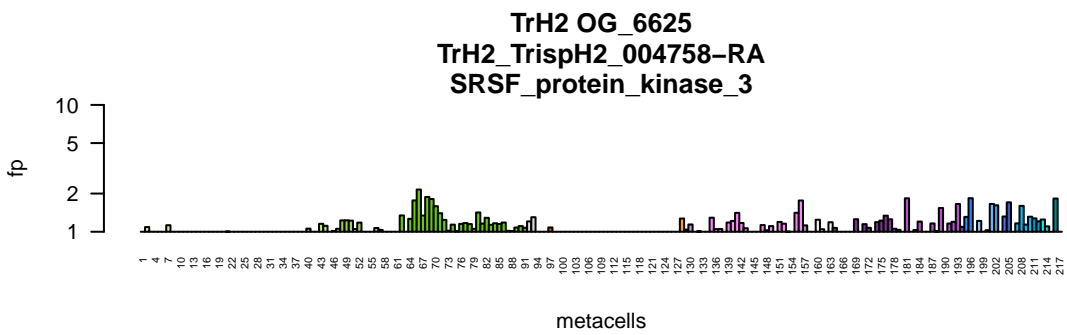
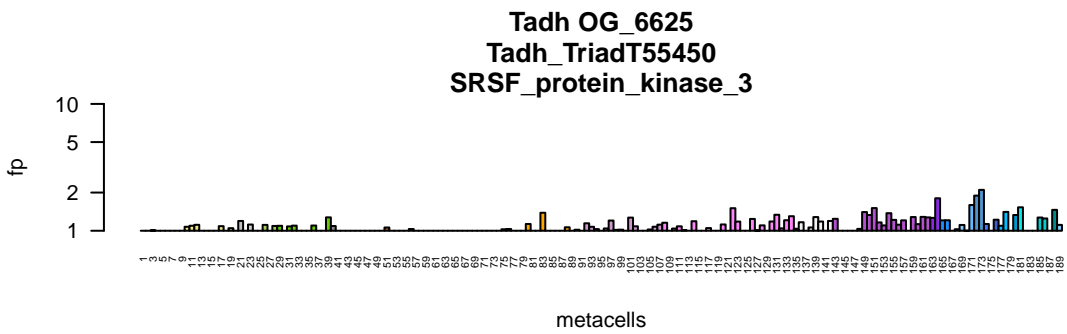


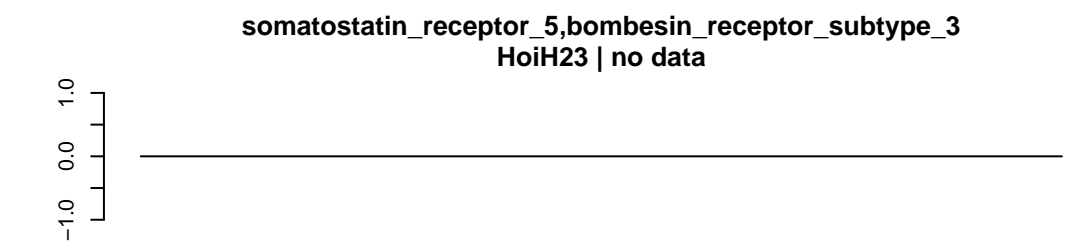
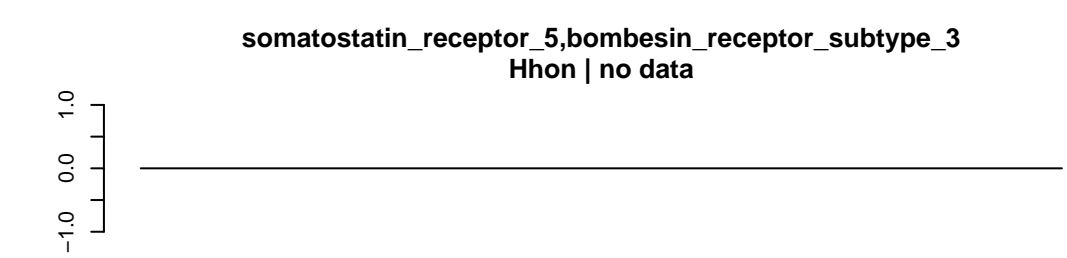
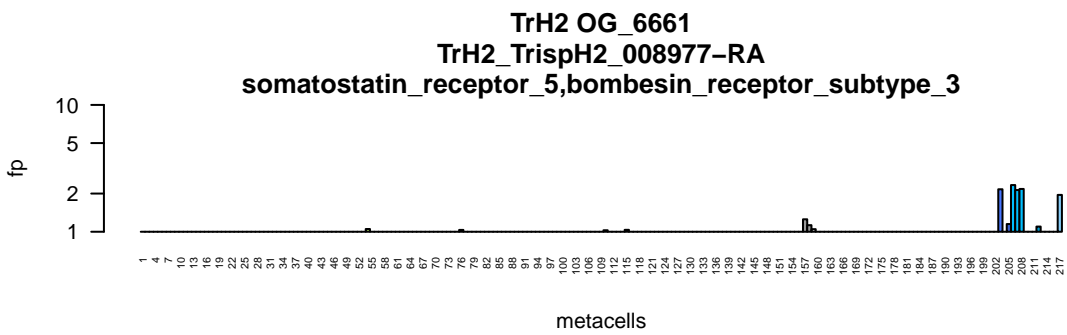
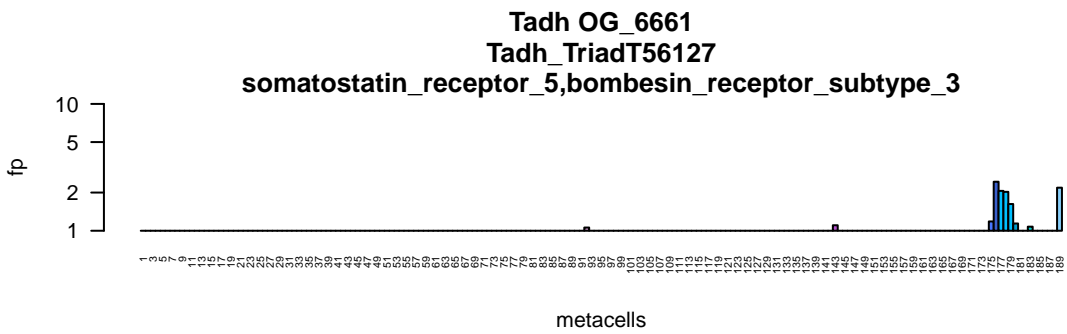


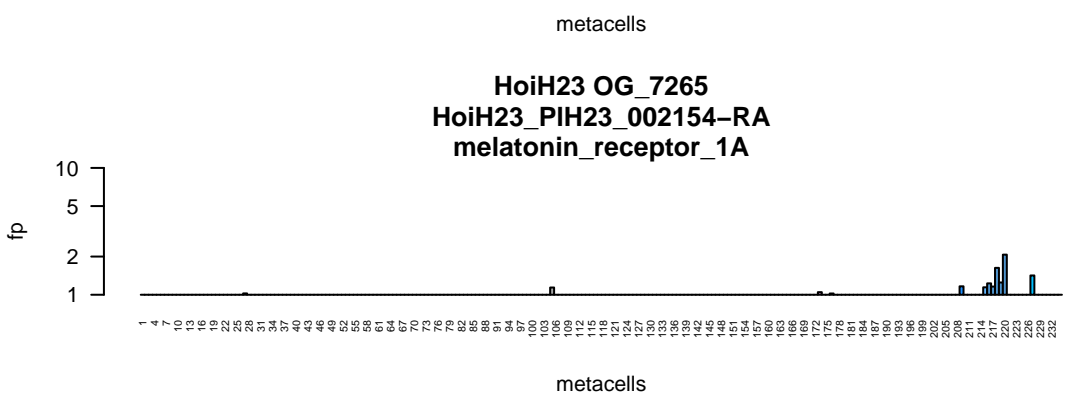
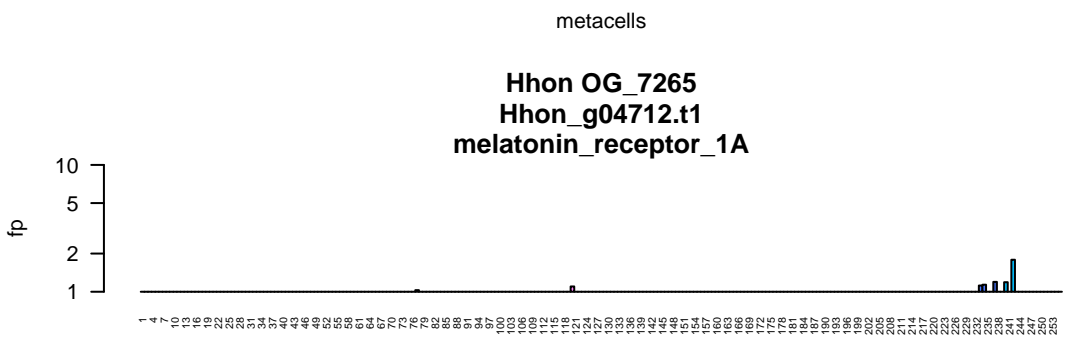
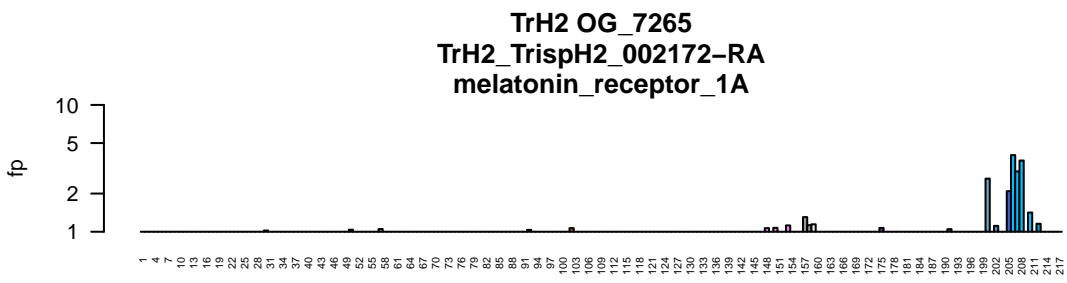
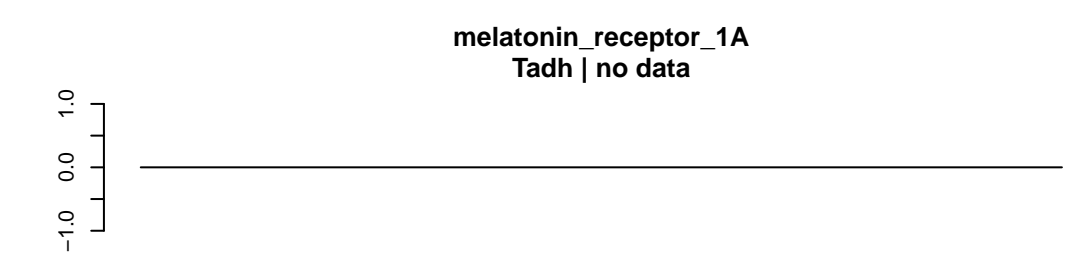


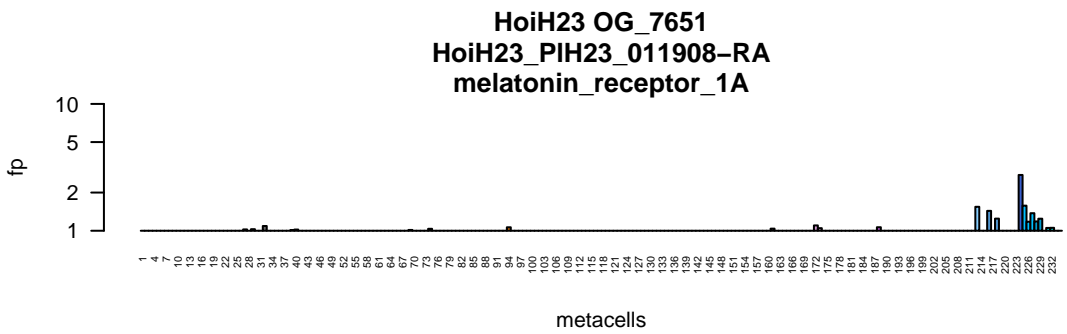
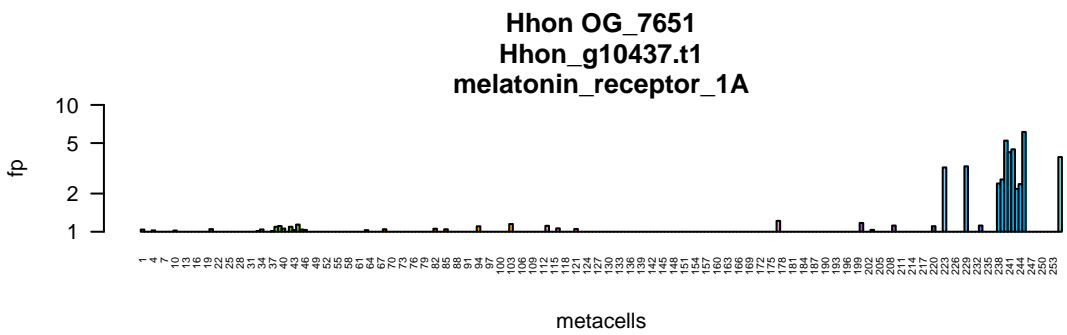
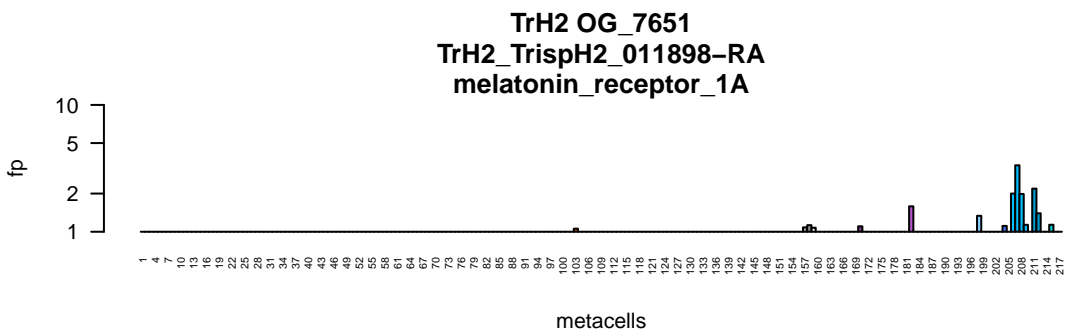
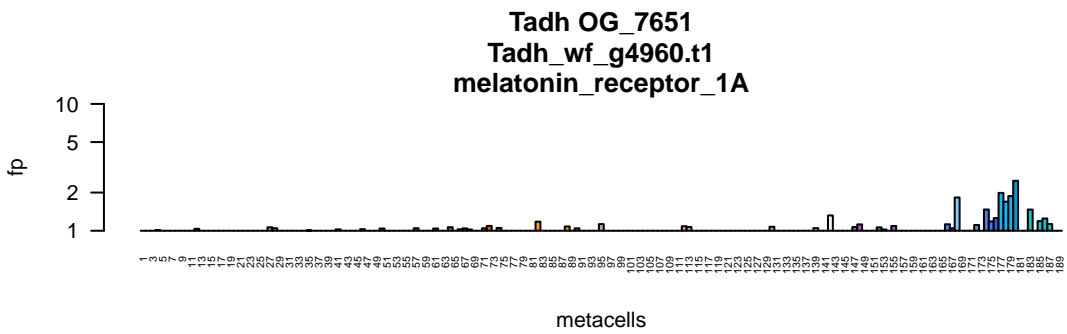


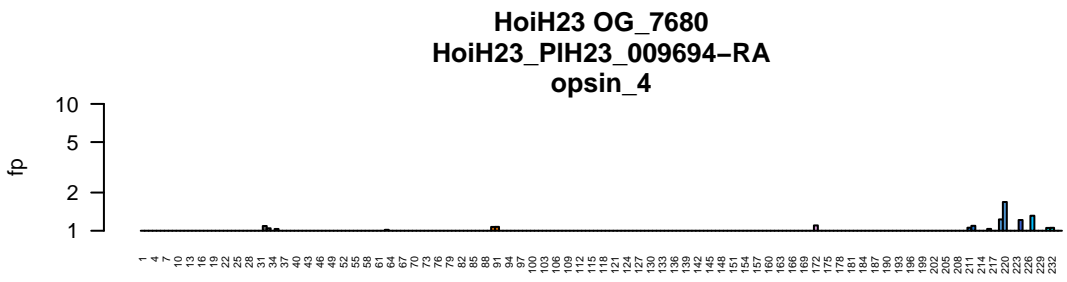
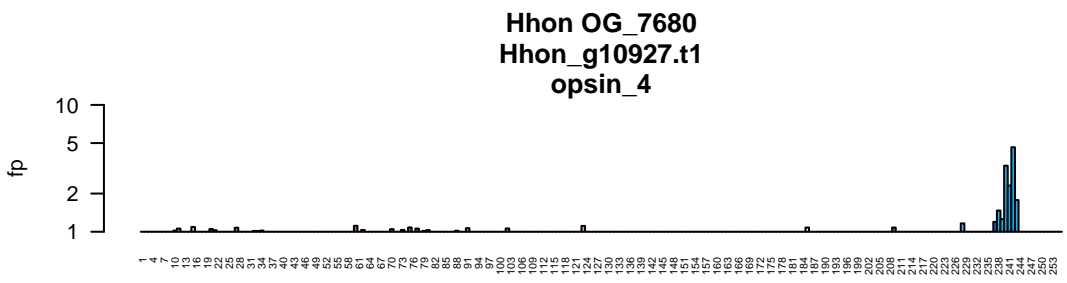
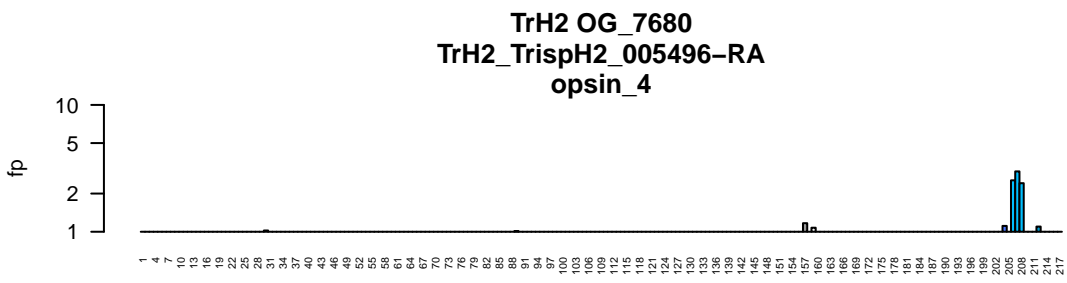
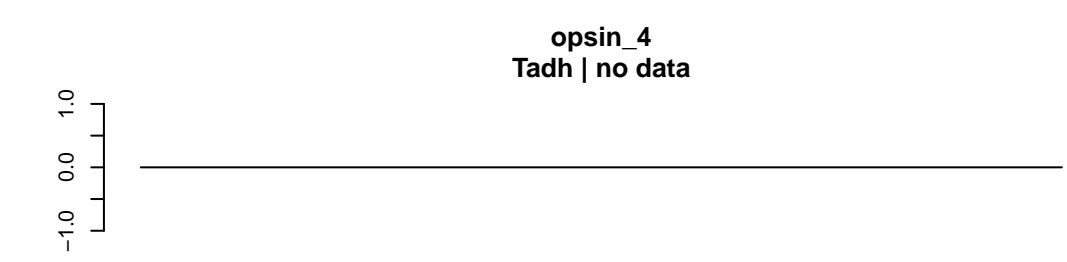


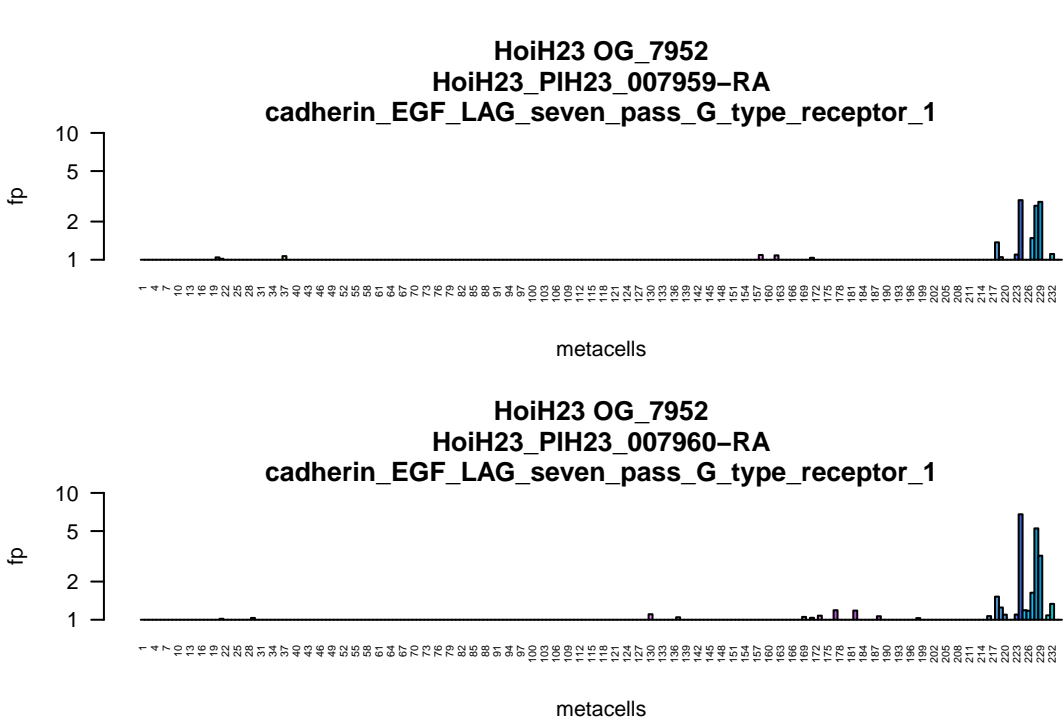
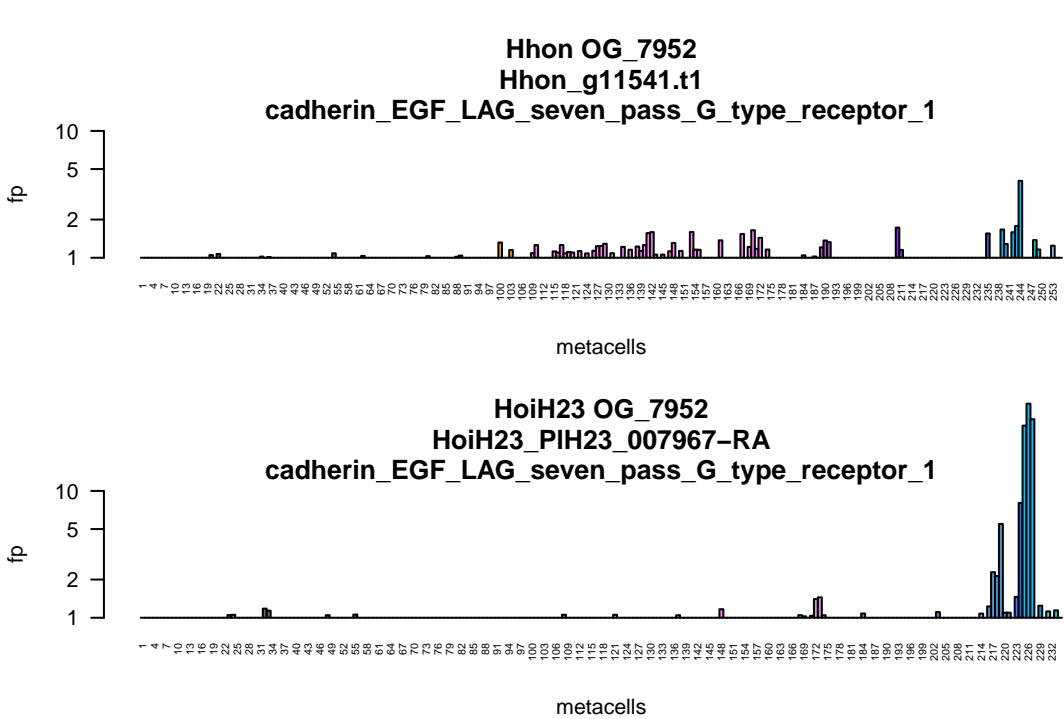
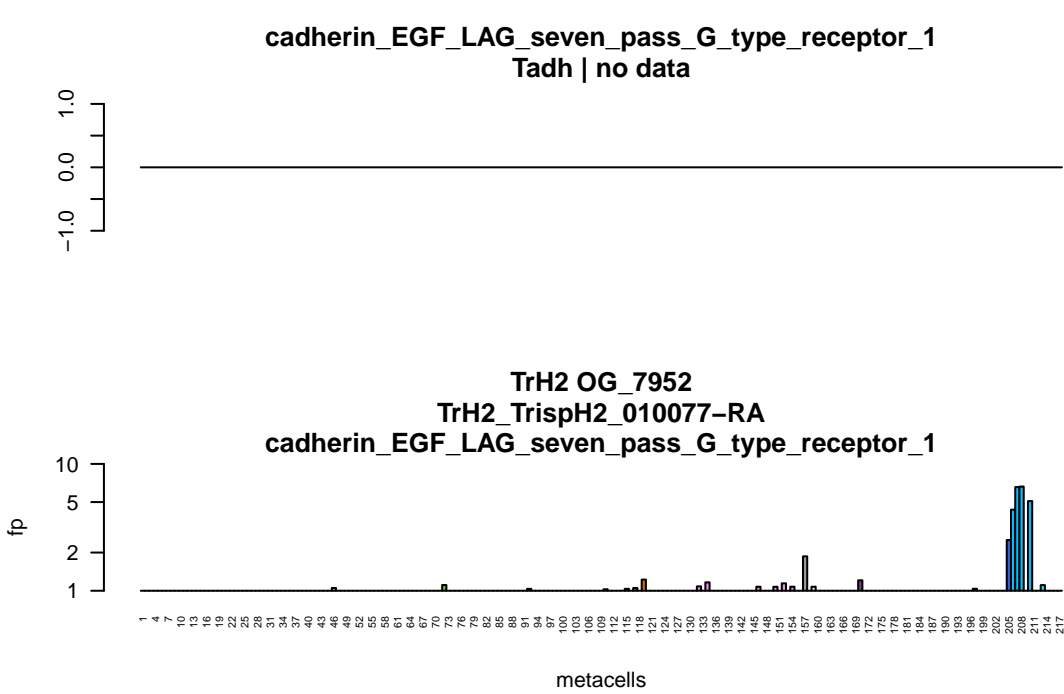




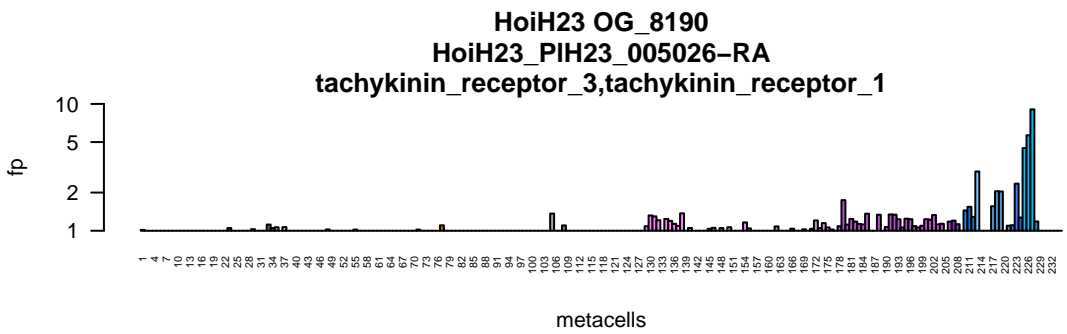
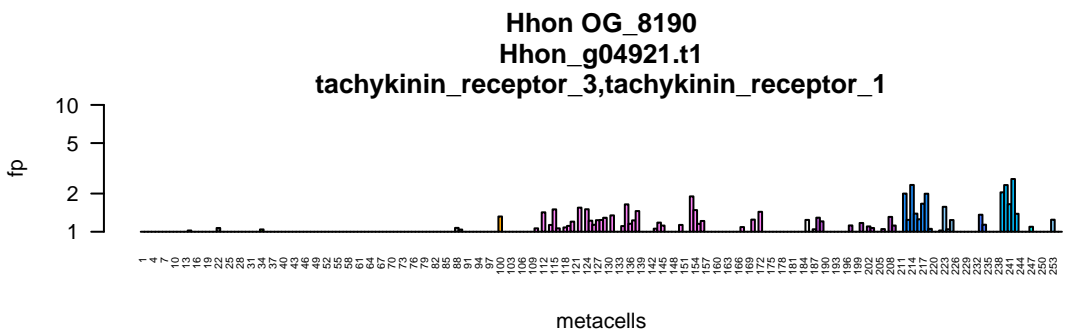
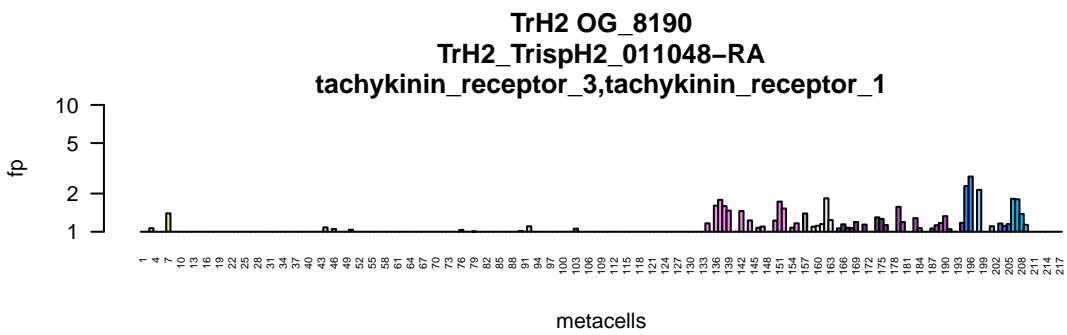
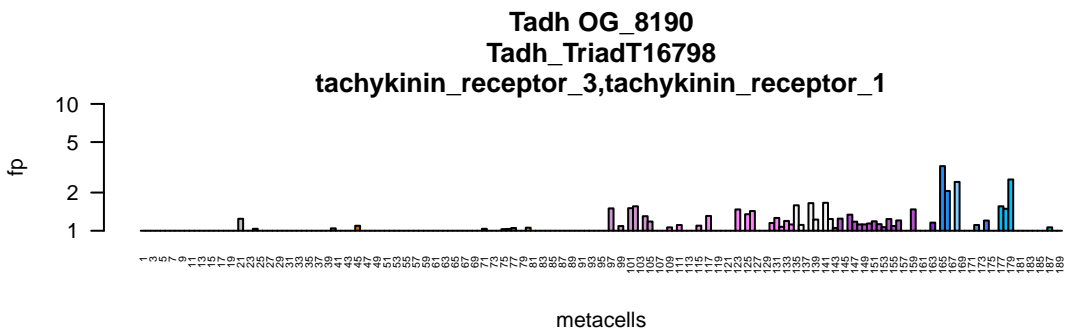


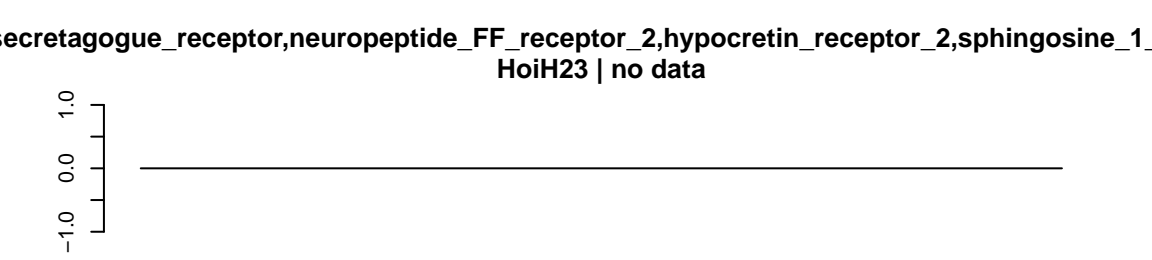
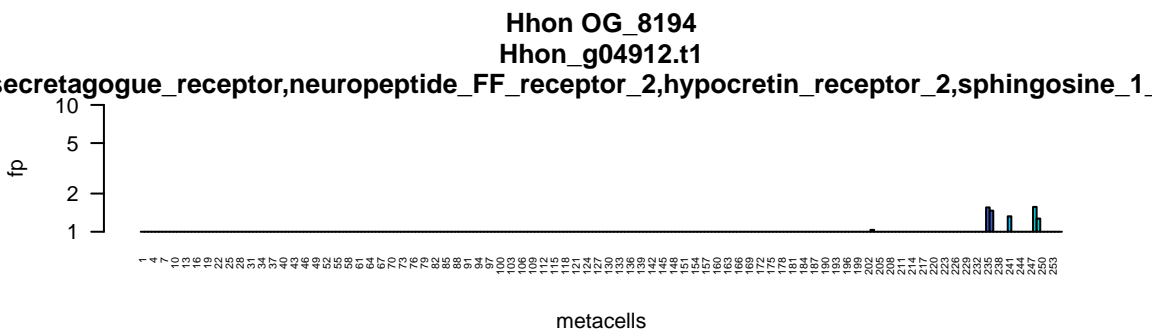
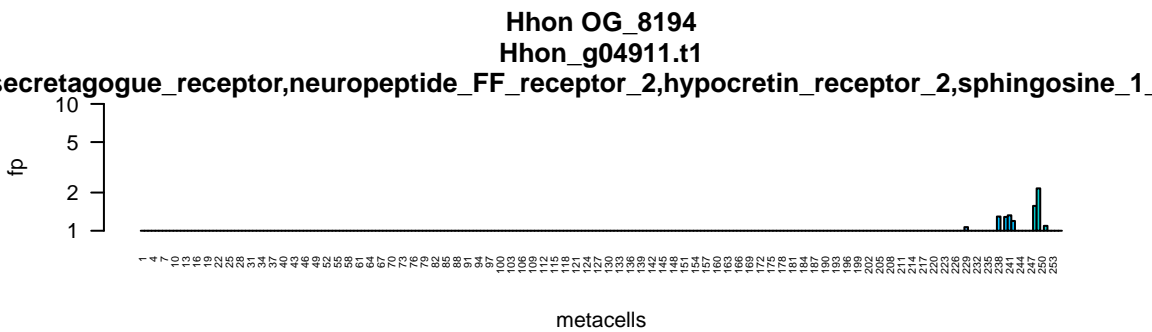
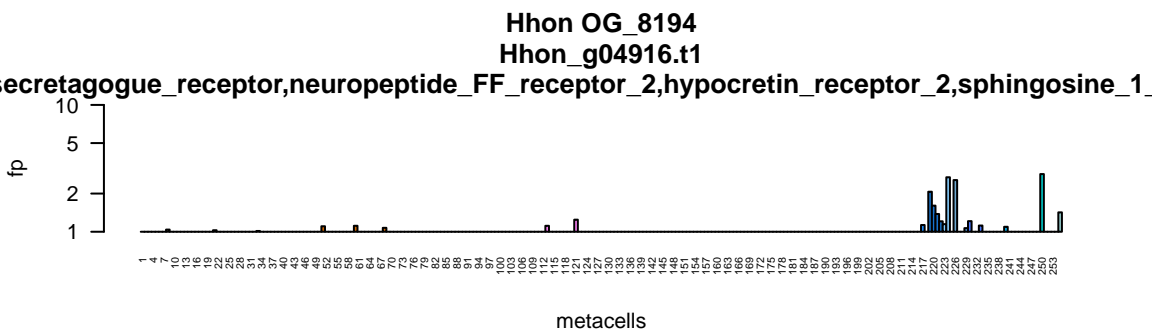
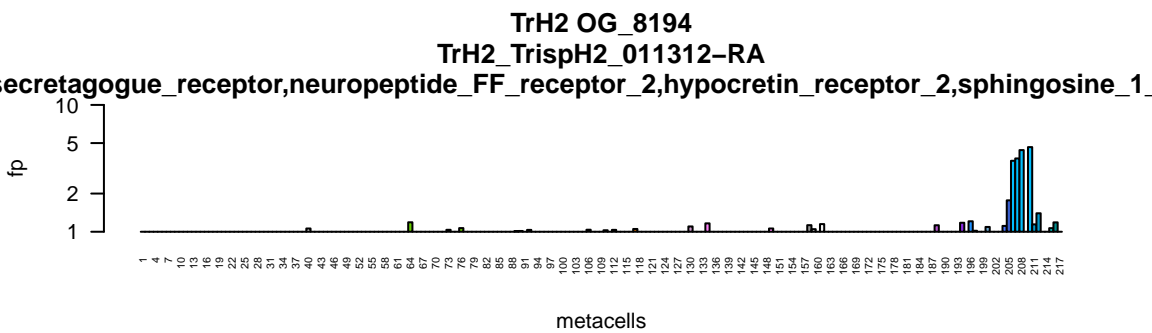
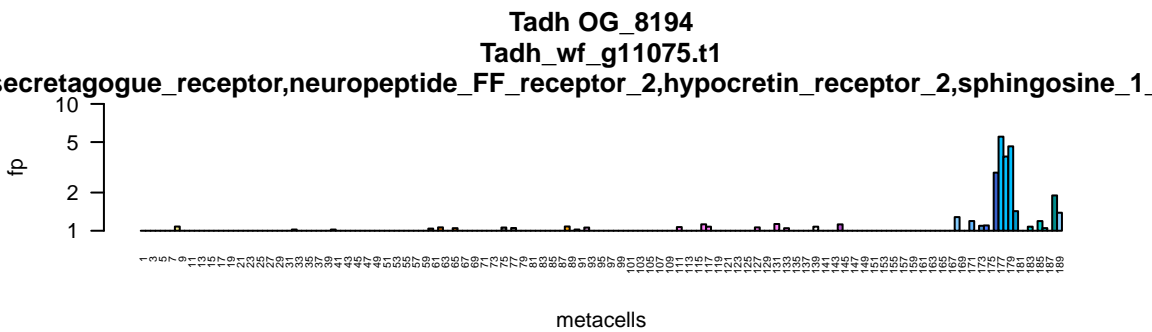






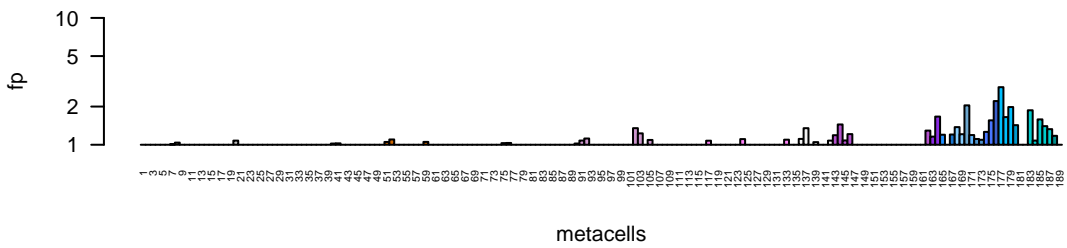






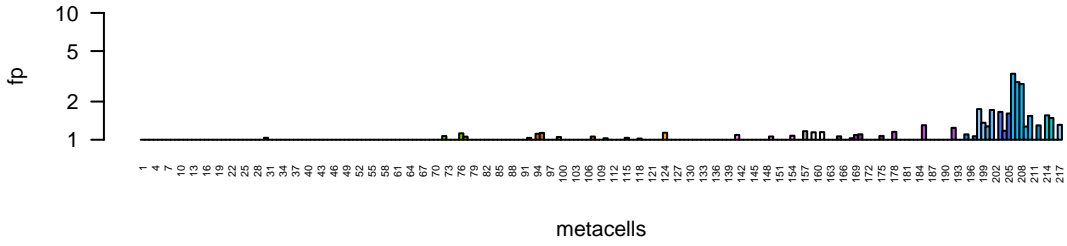
Tadh OG\_8295

Tadh\_TriadT54897



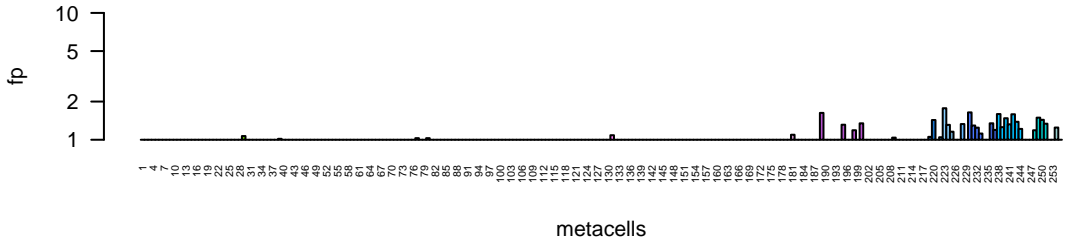
TrH2 OG\_8295

TrH2\_TrispH2\_007497-RA



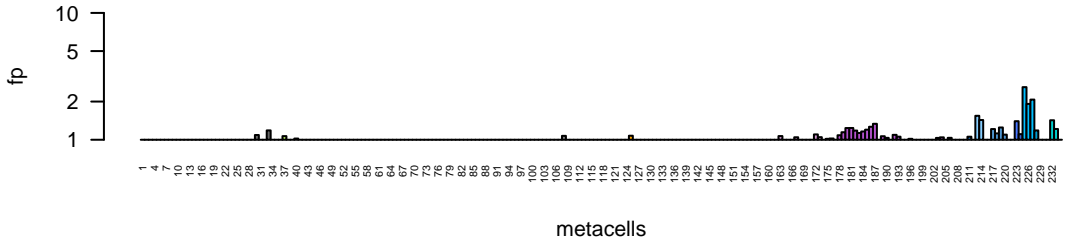
Hhon OG\_8295

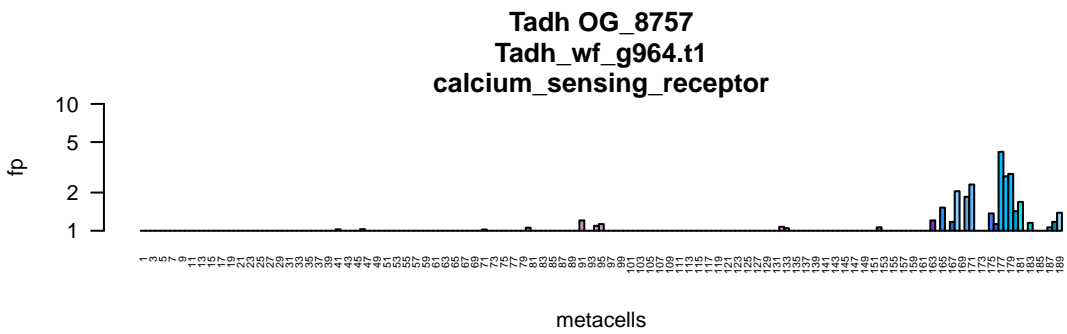
Hhon\_g07101.t1

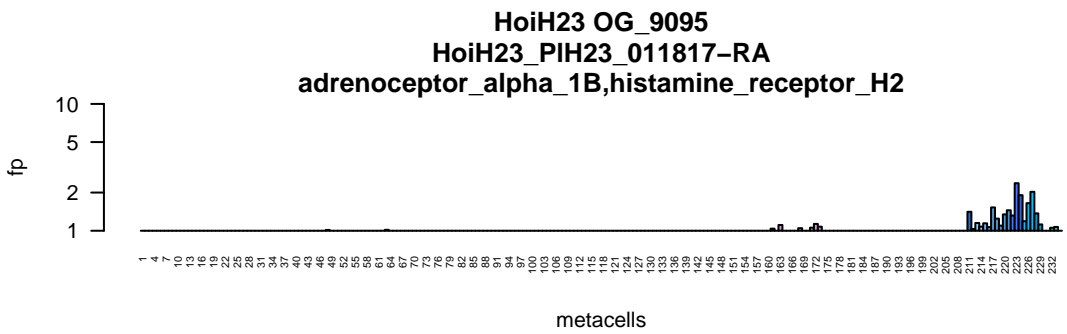
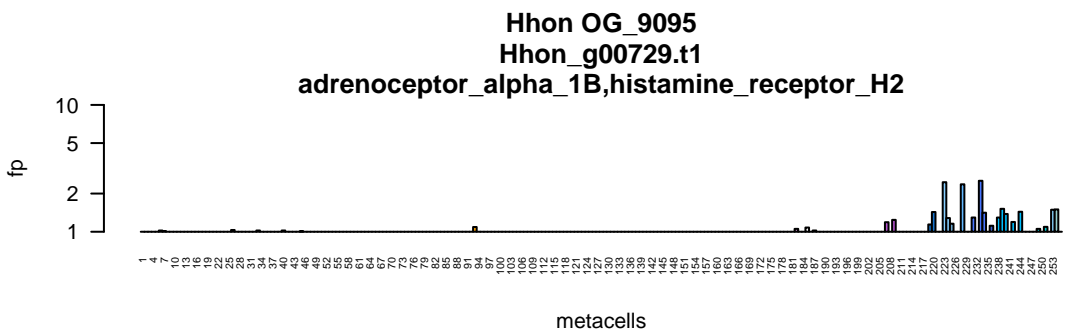
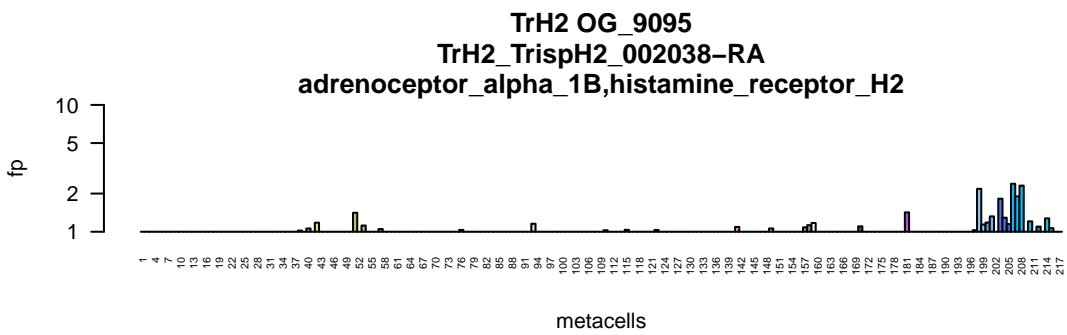
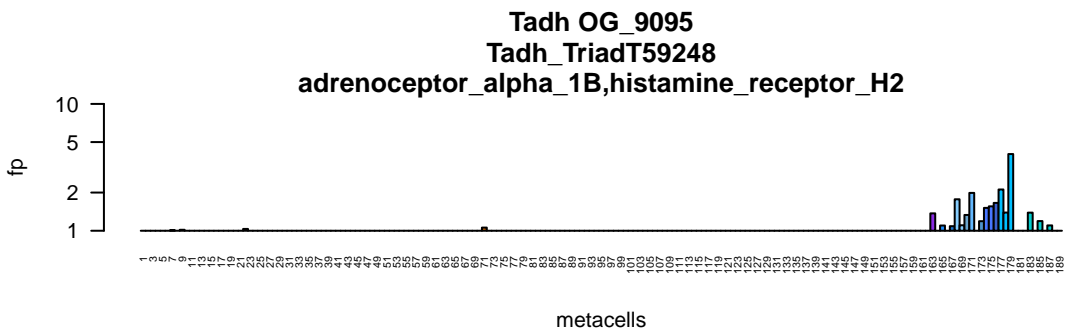


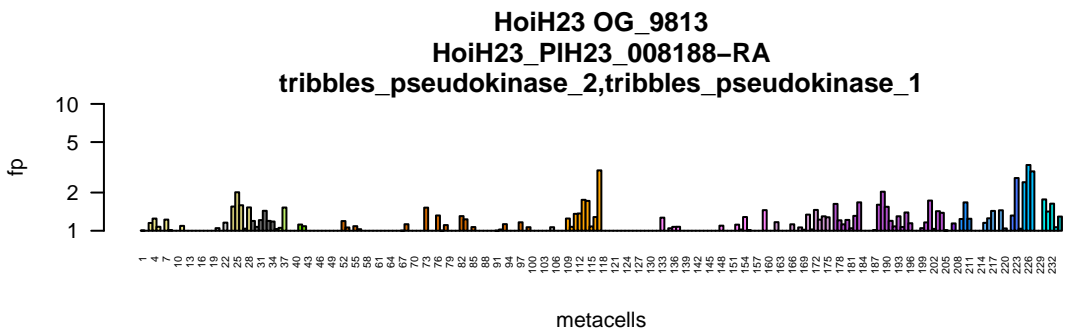
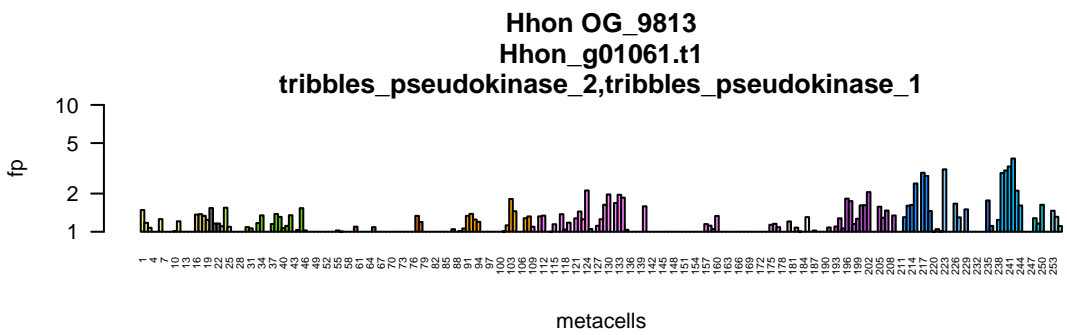
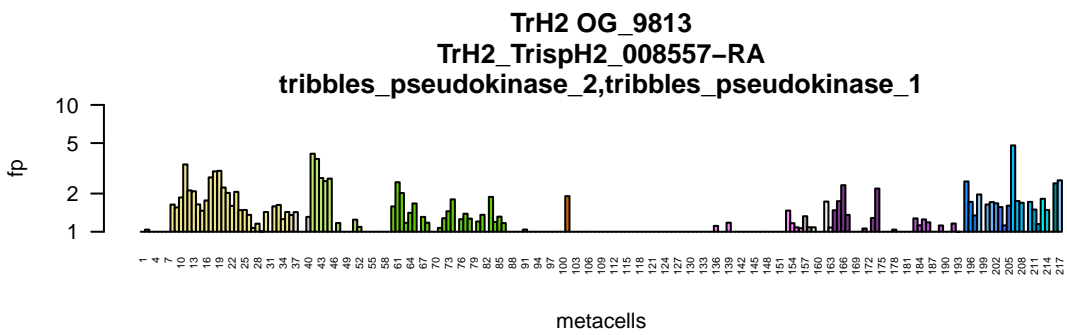
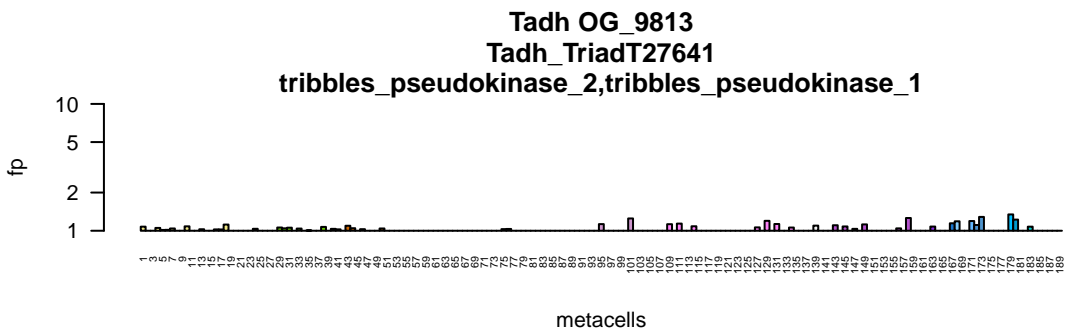
HoiH23 OG\_8295

HoiH23\_PIH23\_002701-RA

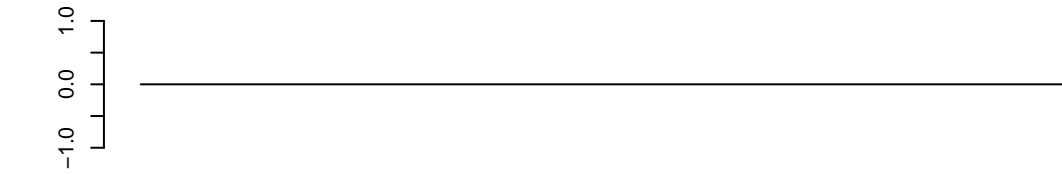




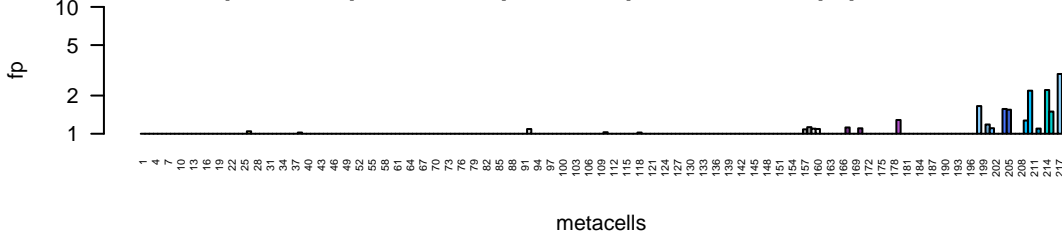




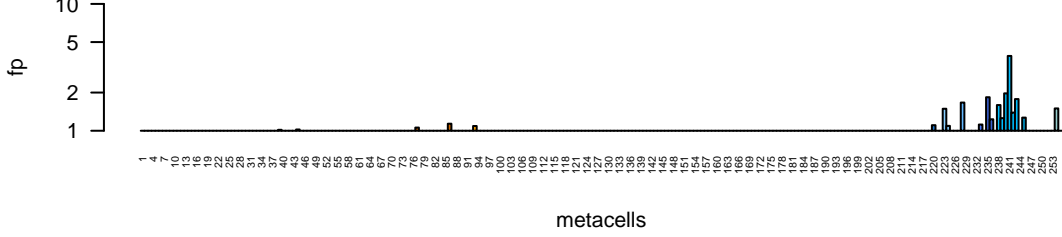
omatostatin\_receptor\_2,G\_protein\_coupled\_receptor\_183,neuropeptides\_B\_and\_W\_recept  
Tadh | no data



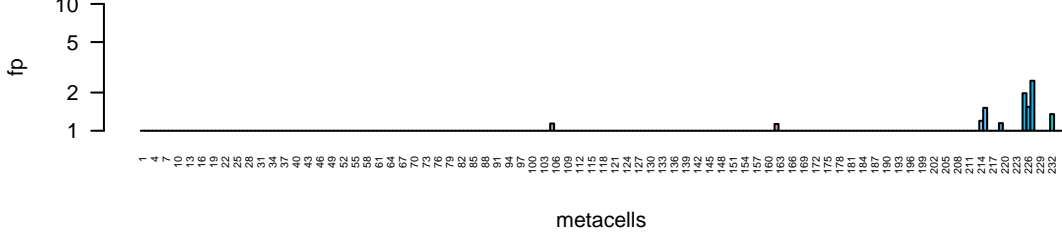
omatostatin\_receptor\_2,G\_protein\_coupled\_receptor\_183,neuropeptides\_B\_and\_W\_recept  
TrH2 OG\_10233  
TrH2\_TrispH2\_006452-RA



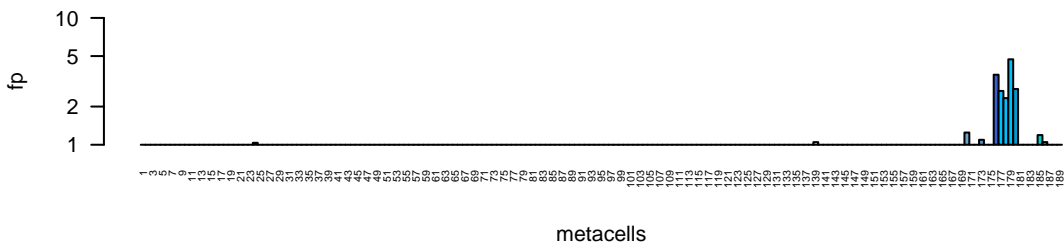
omatostatin\_receptor\_2,G\_protein\_coupled\_receptor\_183,neuropeptides\_B\_and\_W\_recept  
Hhon OG\_10233  
Hhon\_g05148.t1



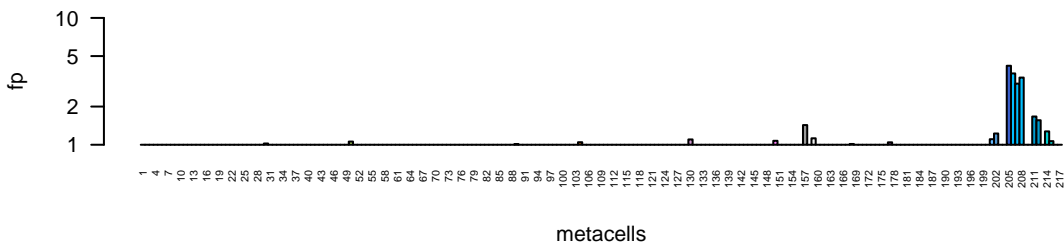
omatostatin\_receptor\_2,G\_protein\_coupled\_receptor\_183,neuropeptides\_B\_and\_W\_recept  
HoiH23 OG\_10233  
HoiH23\_PIH23\_001231-RA



Tadh OG\_10419  
Tadh\_wf\_g4956.t1



TrH2 OG\_10419  
TrH2\_TrispH2\_008975-RA



Hhon | no data

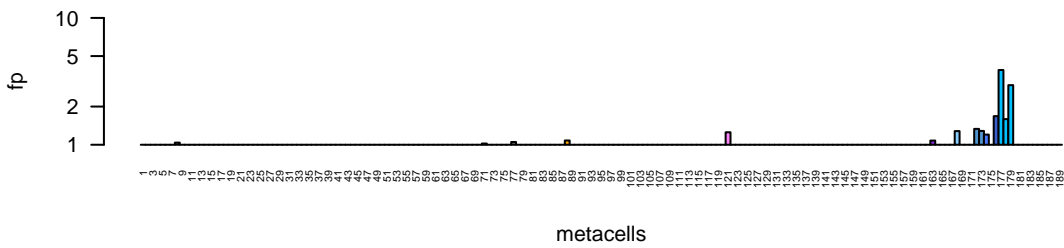


HoiH23 | no data

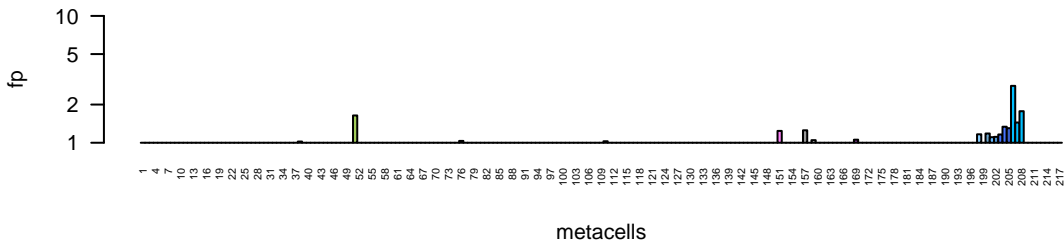




Tadh OG\_10549  
Tadh\_wf\_g9549.t1



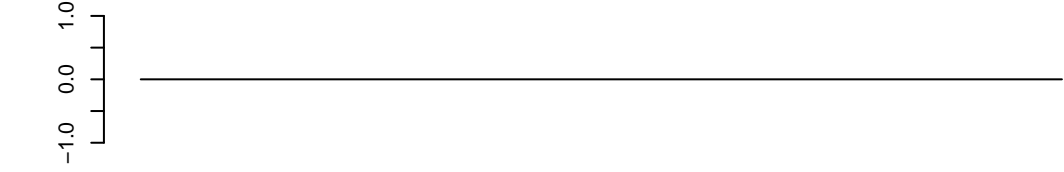
TrH2 OG\_10549  
TrH2\_TrispH2\_011188-RA

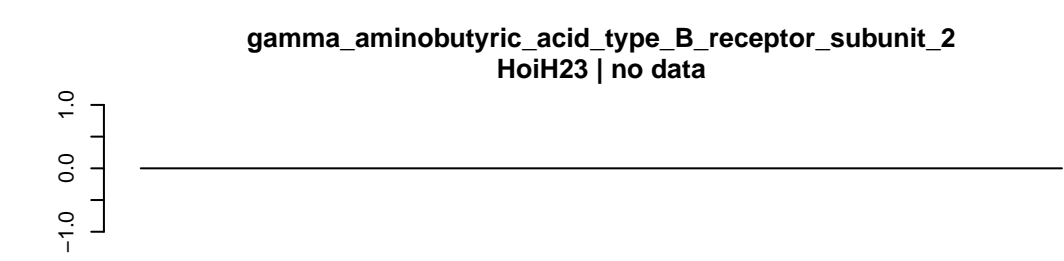
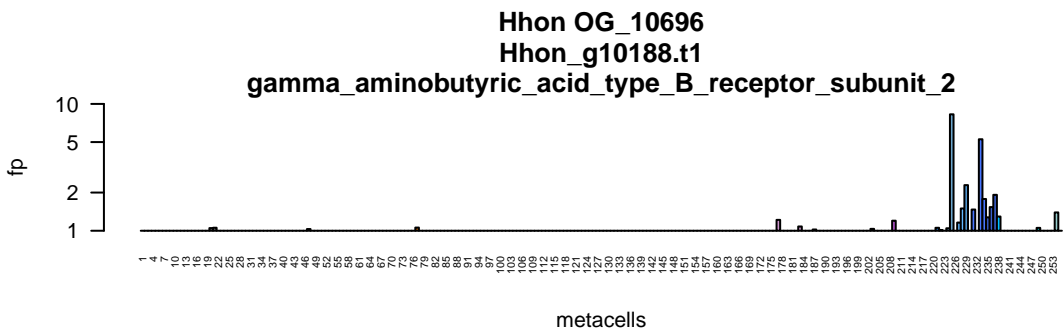
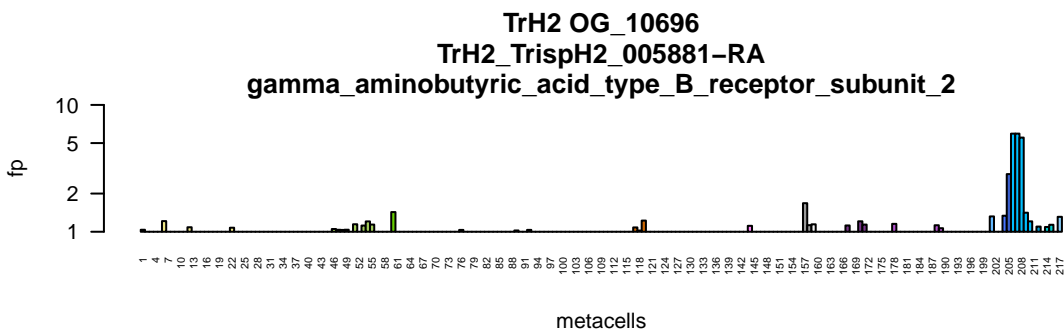
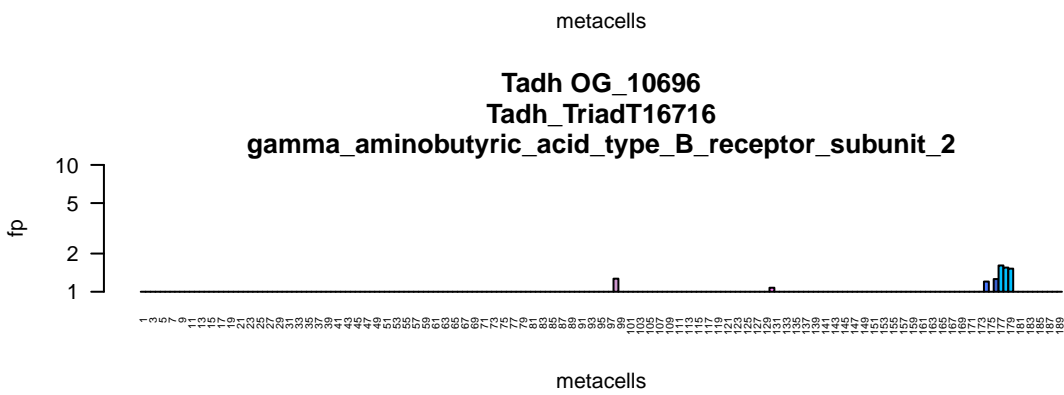
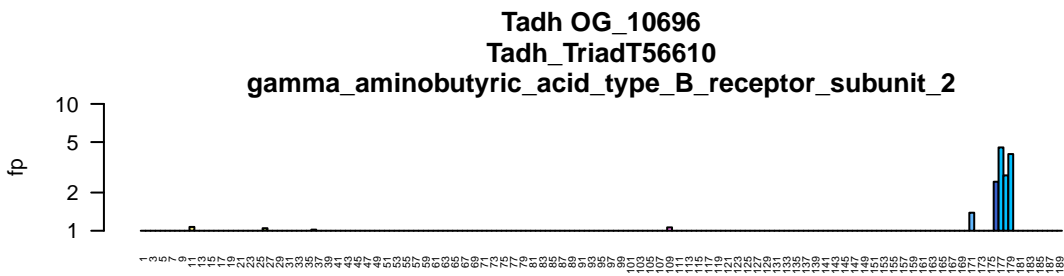


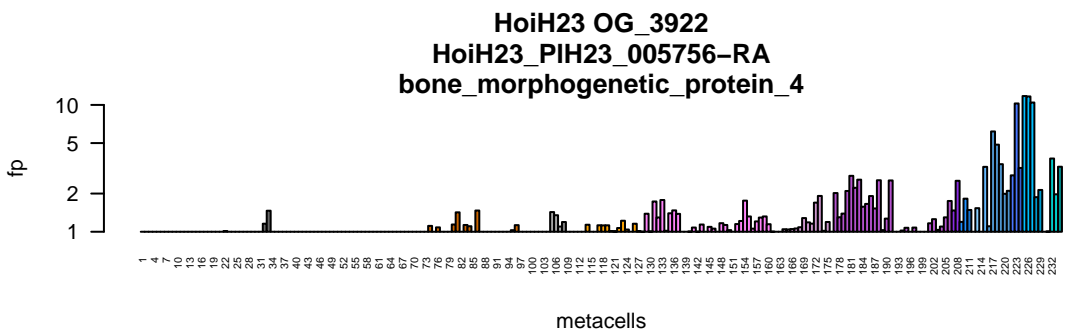
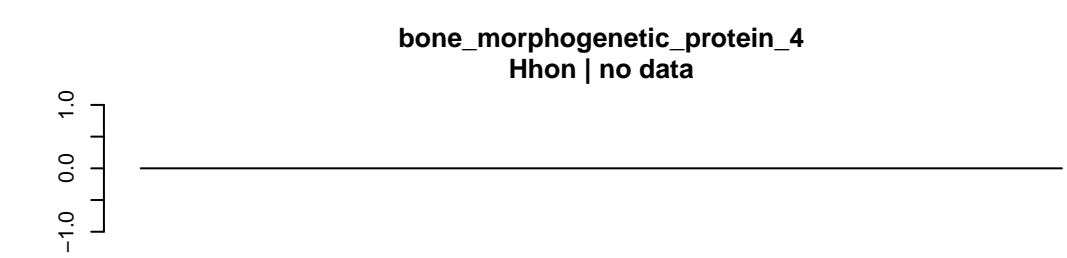
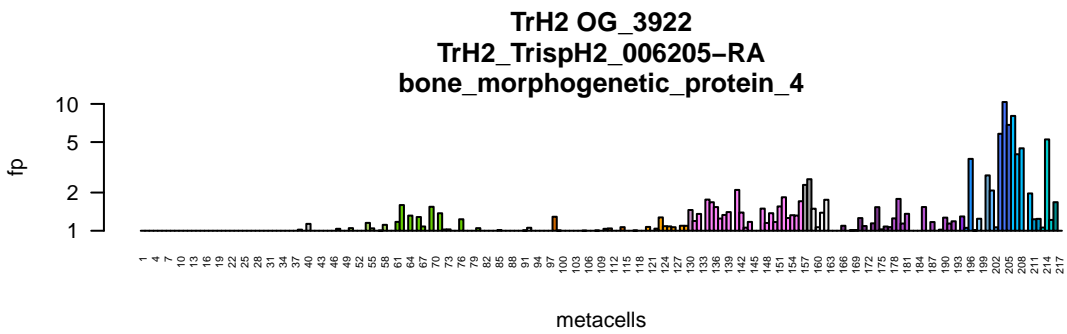
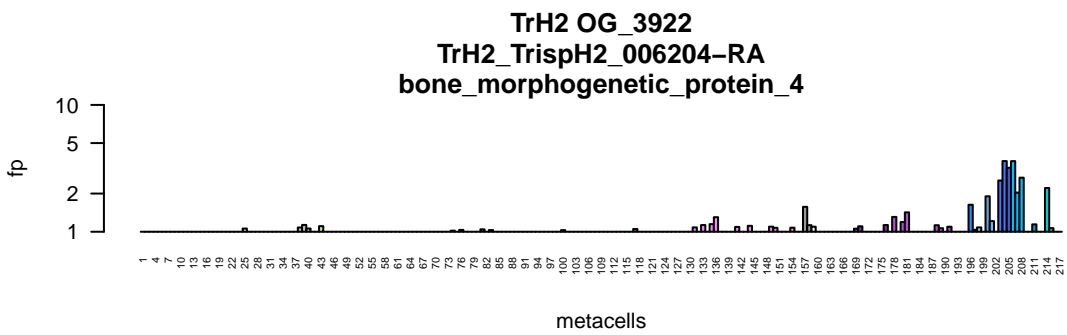
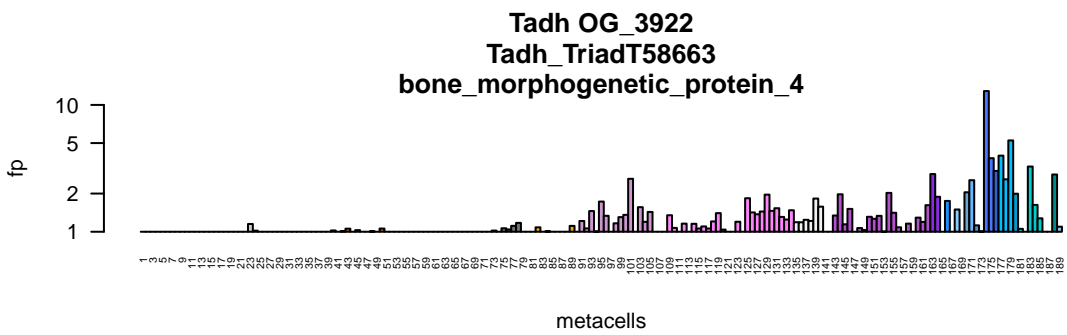
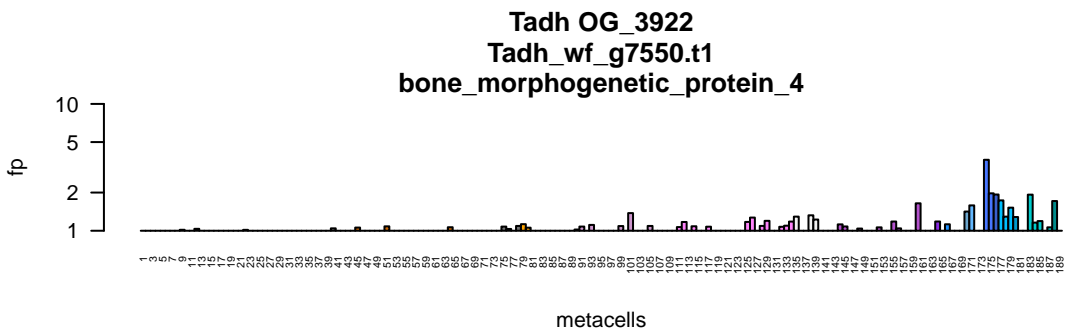
Hhon | no data

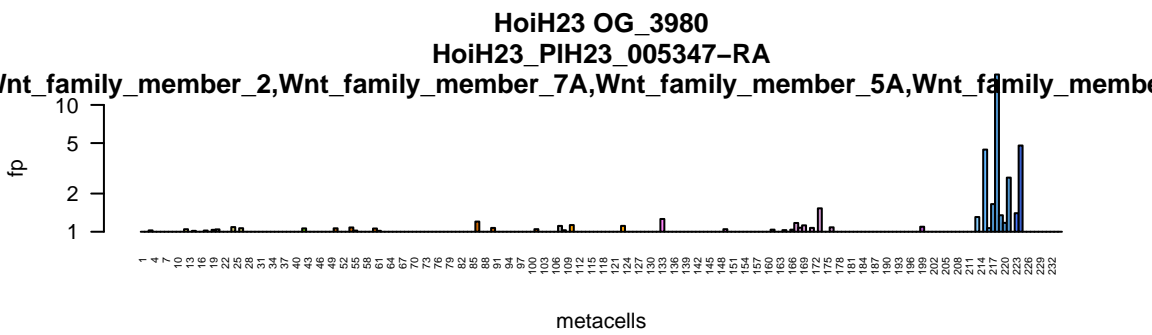
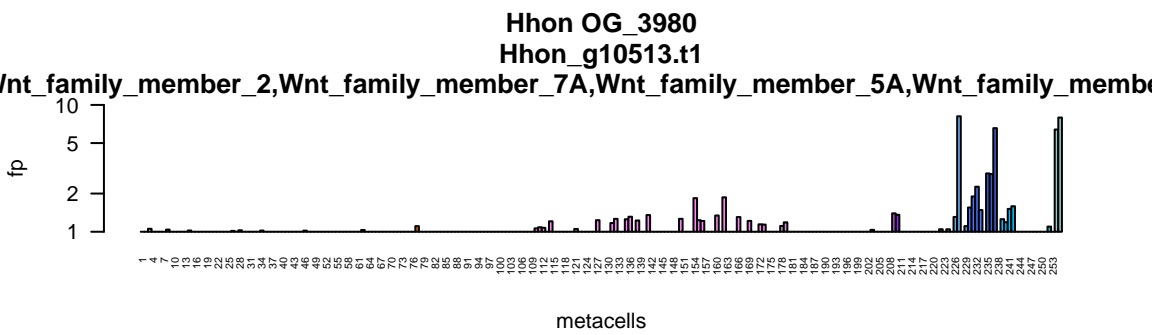
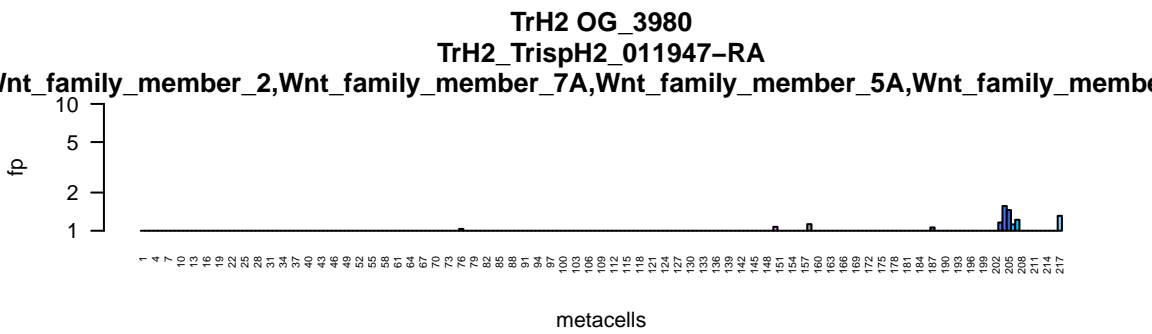
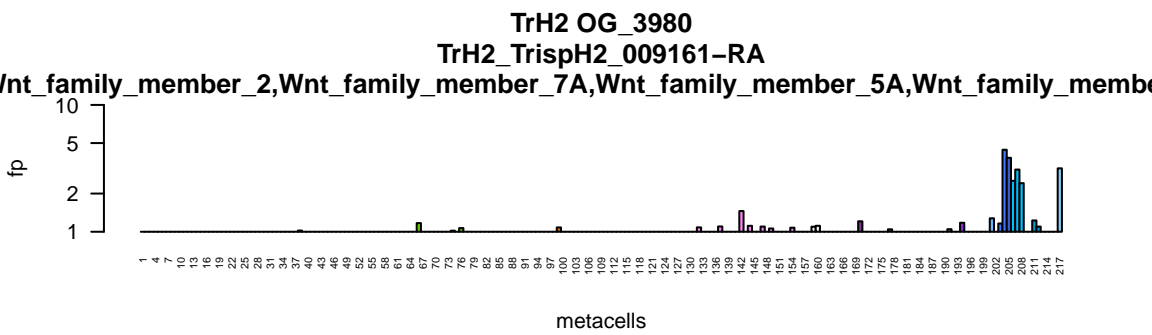
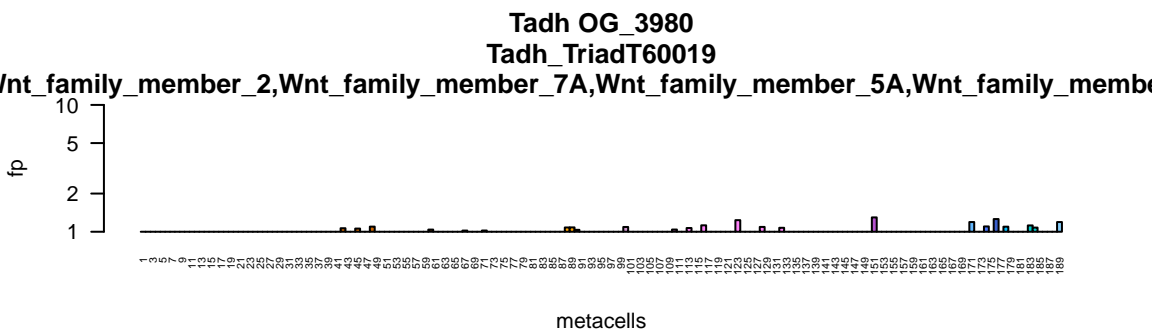
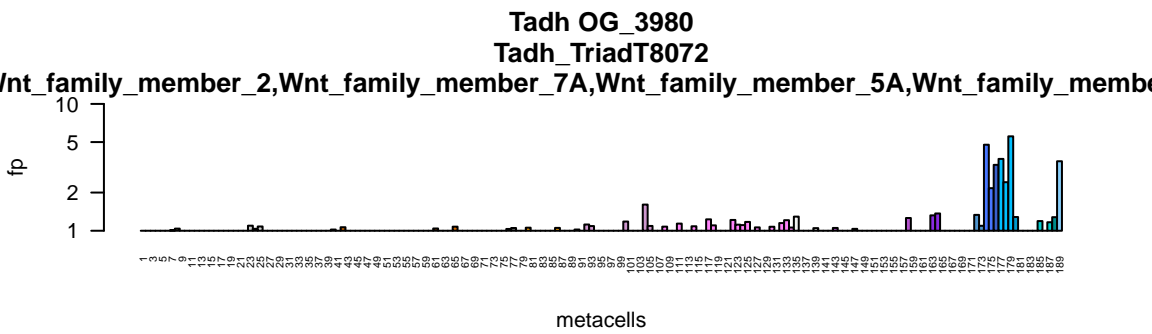


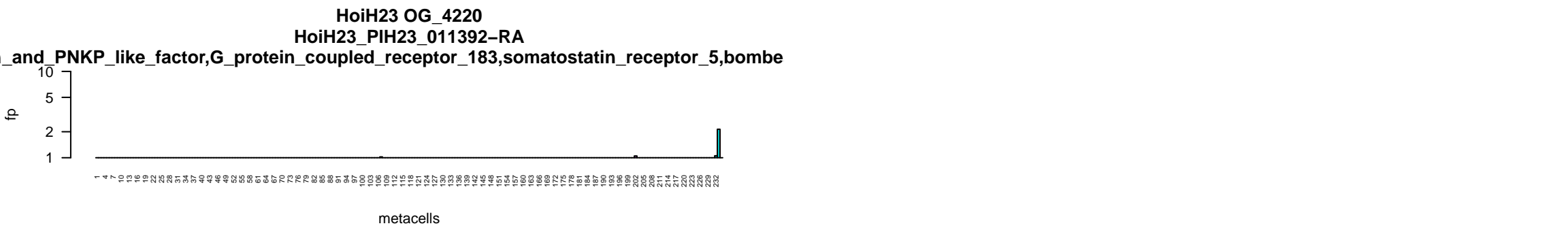
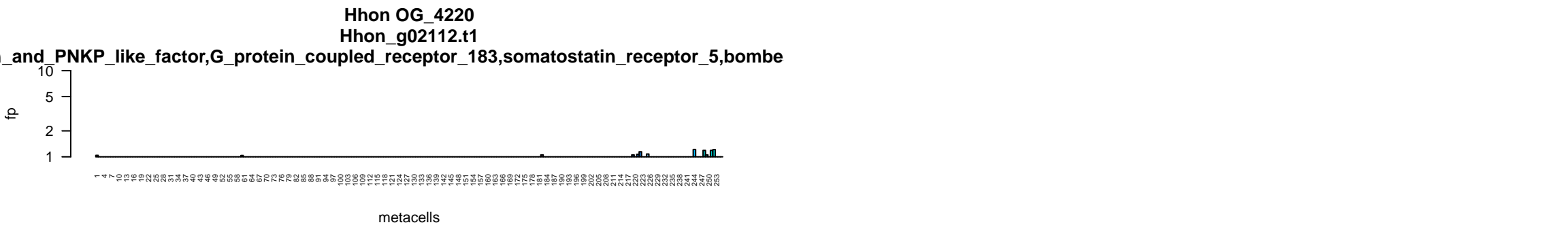
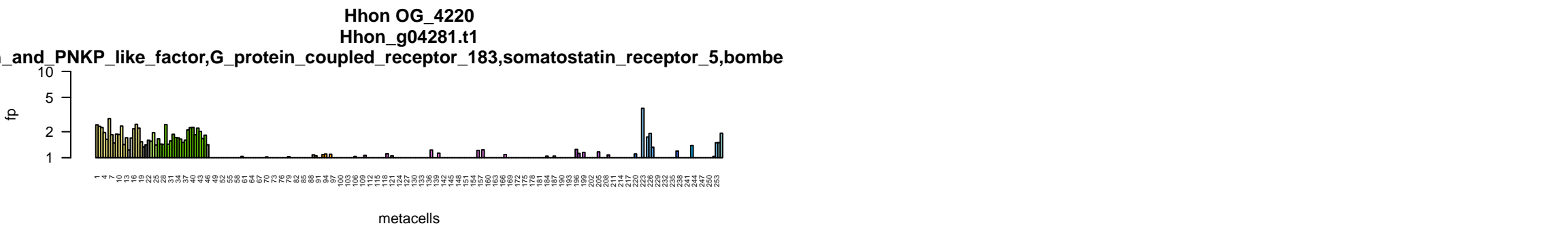
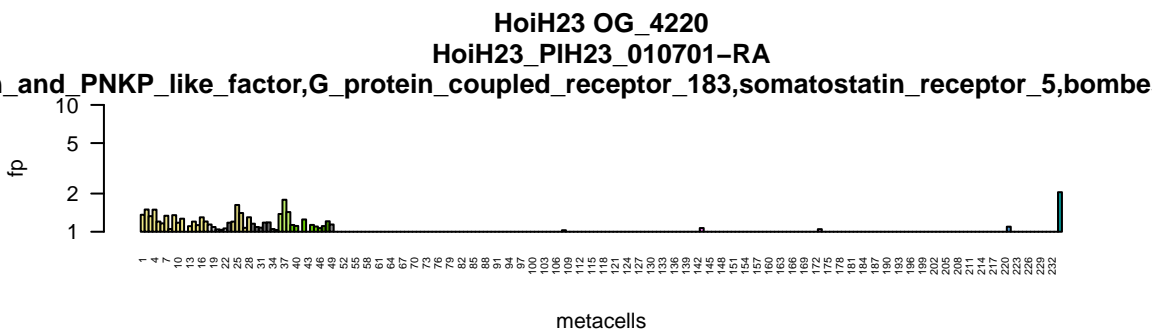
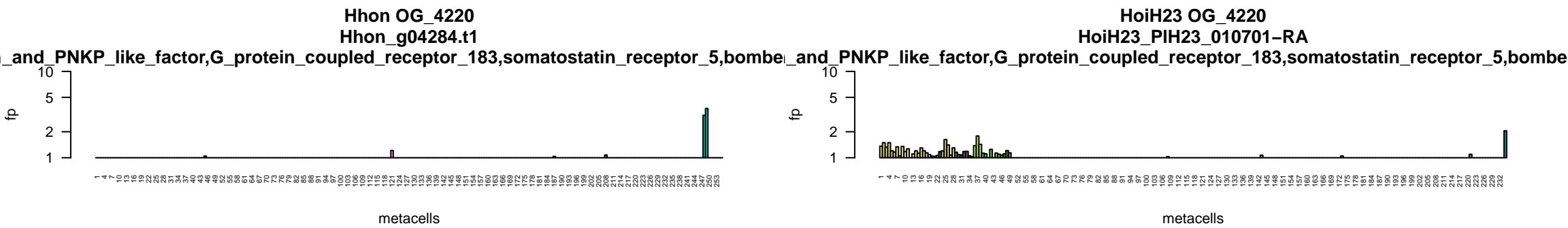
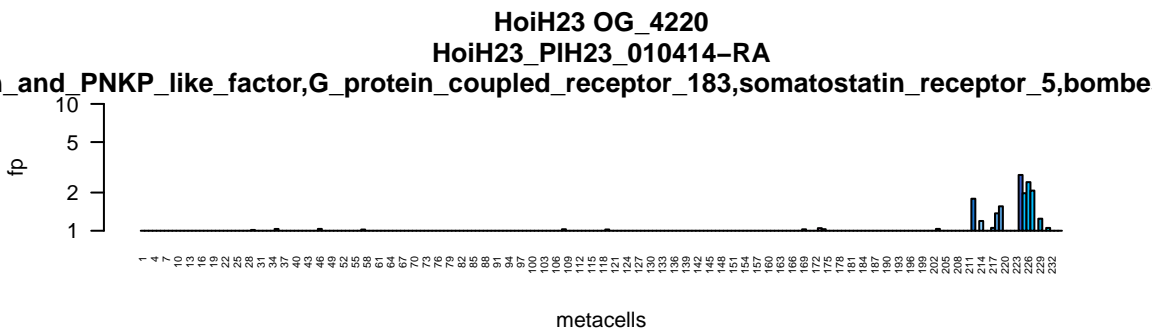
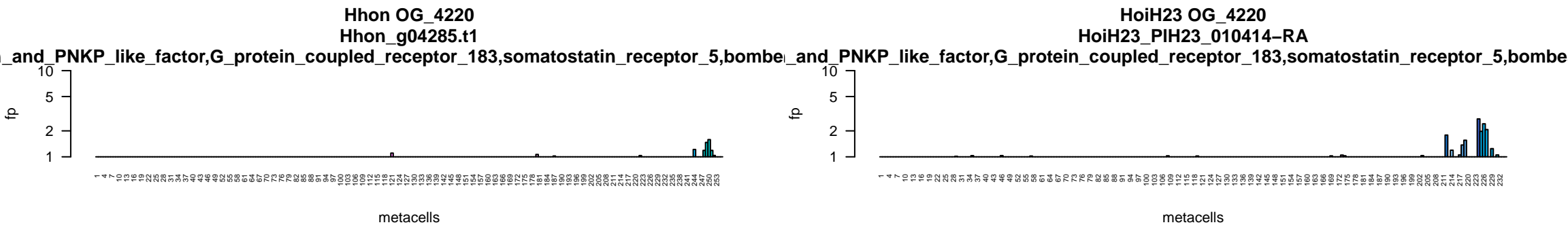
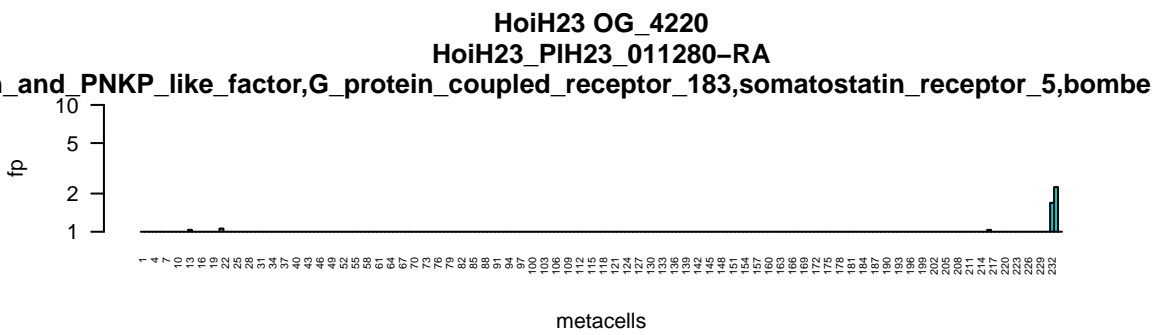
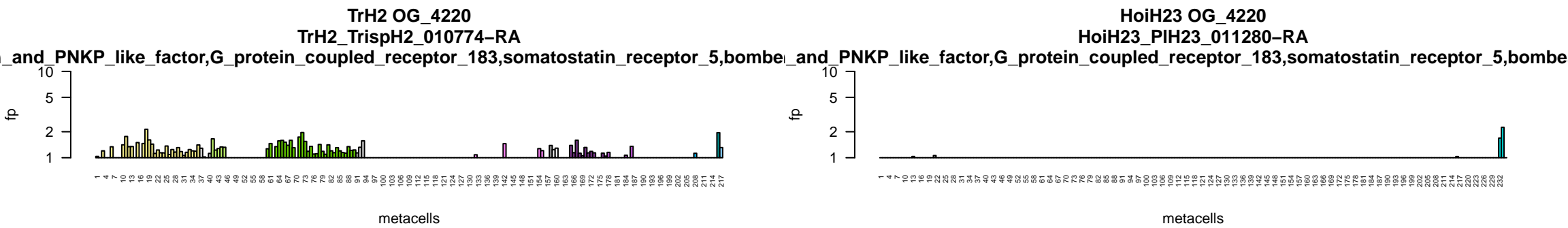
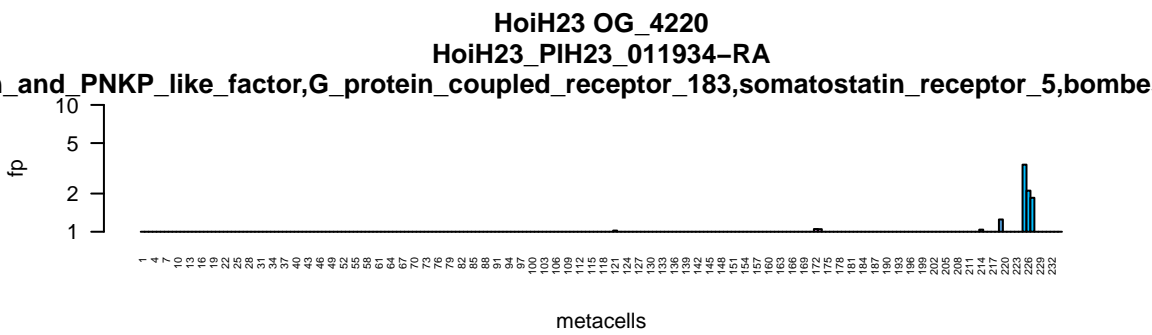
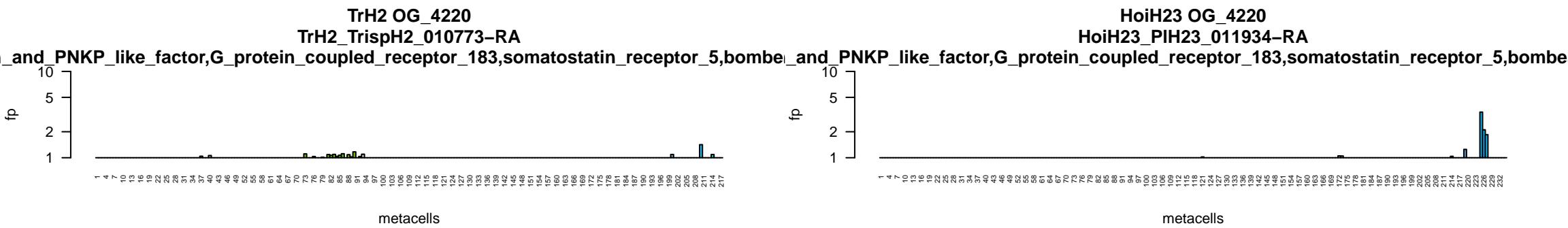
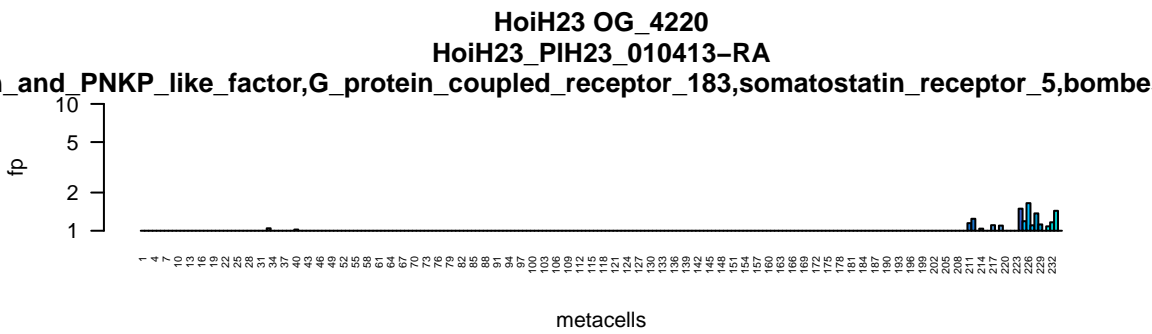
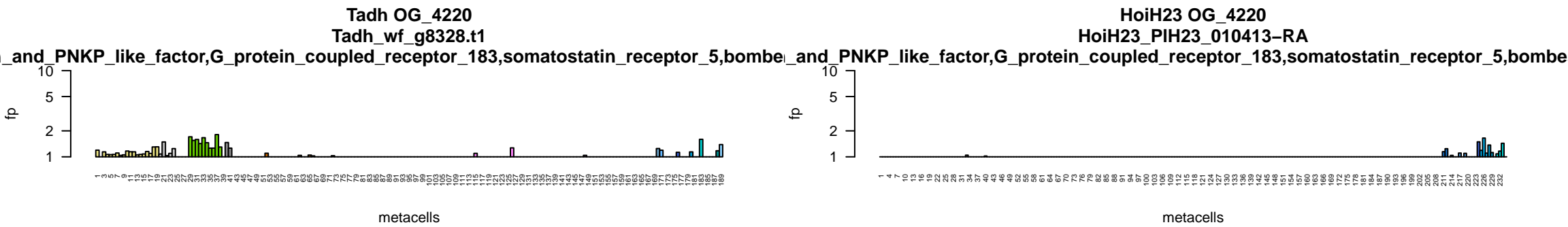
HoiH23 | no data

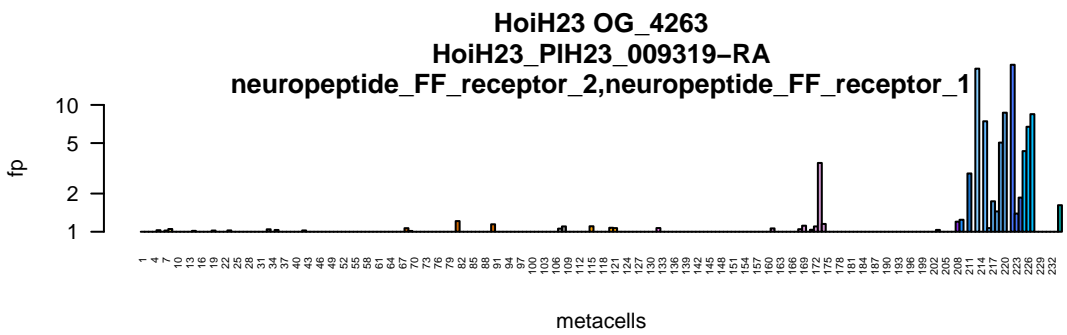
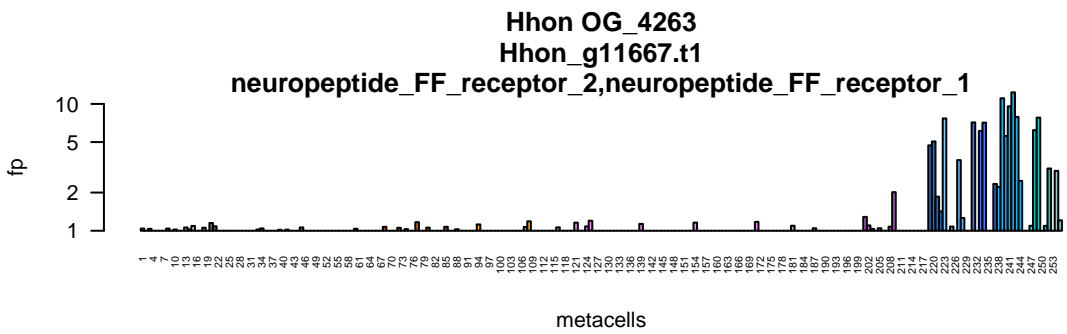
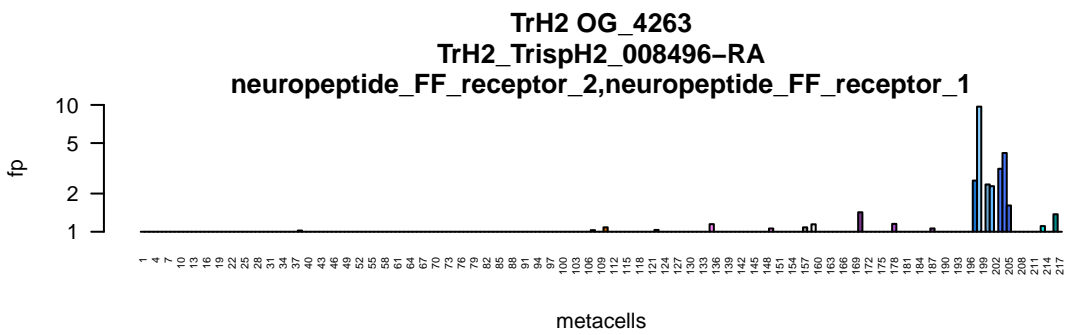
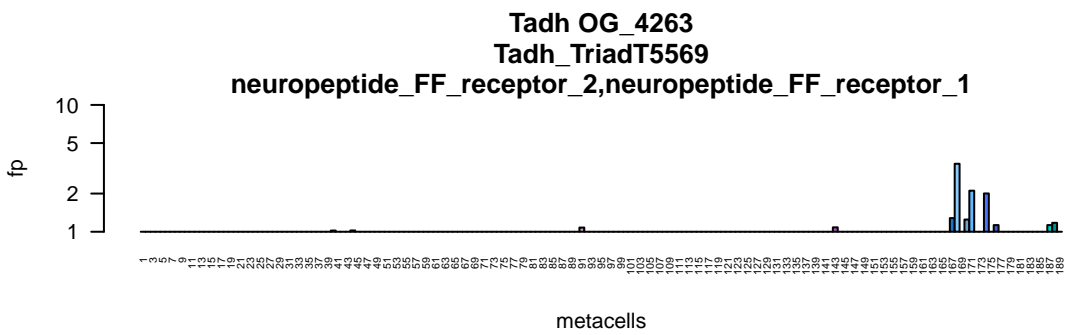
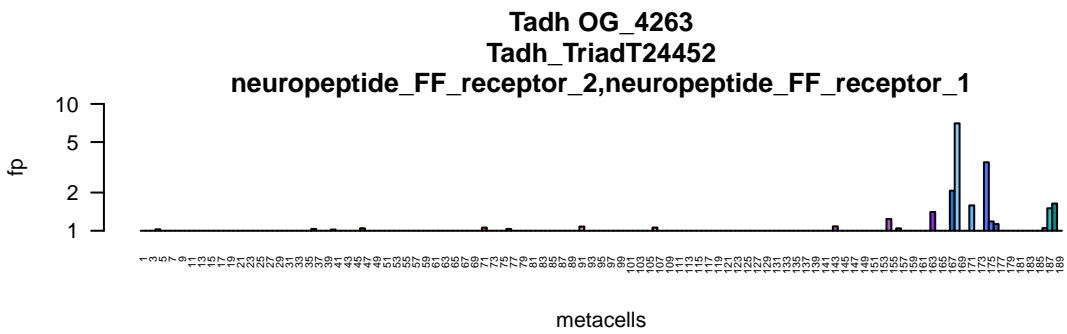


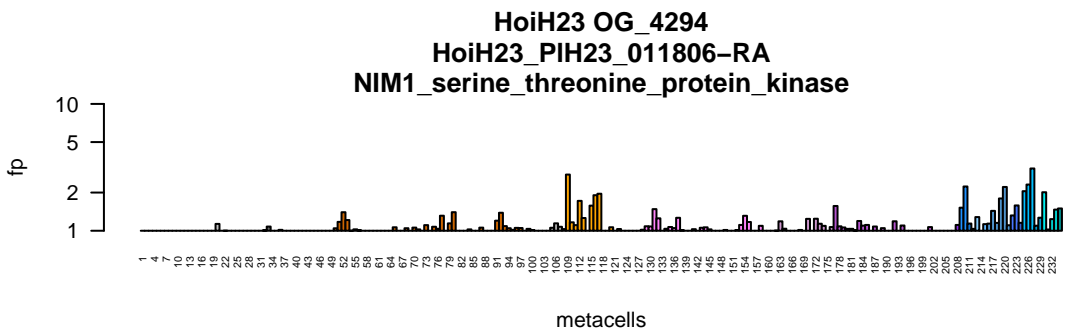
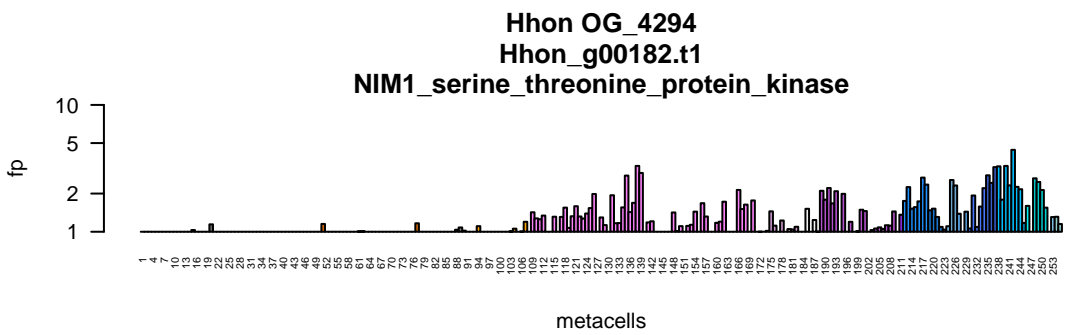
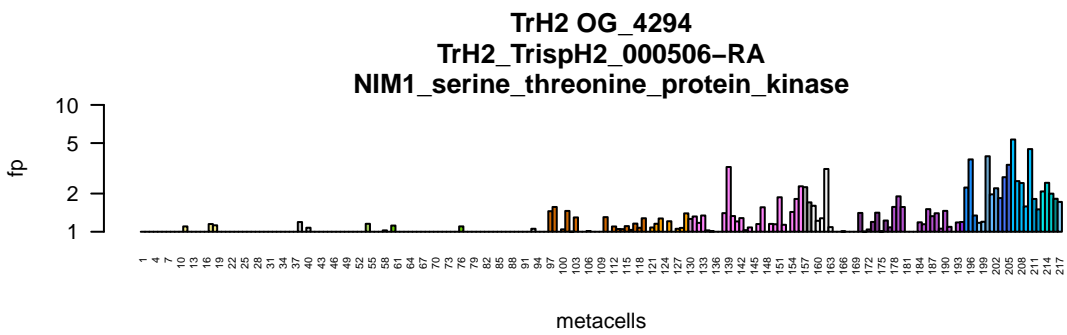
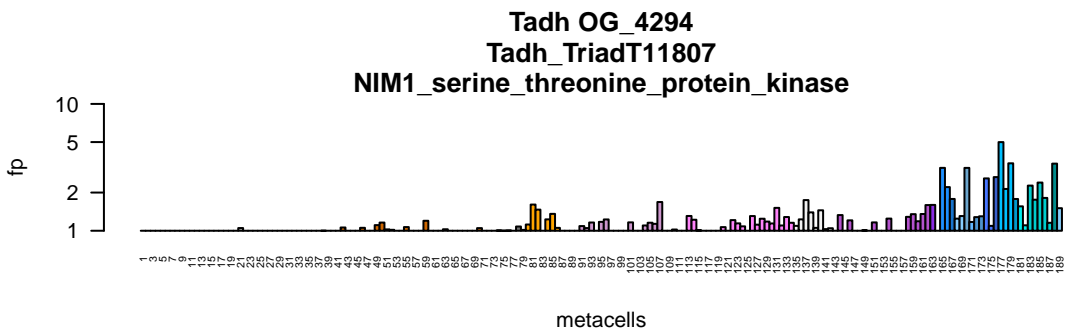








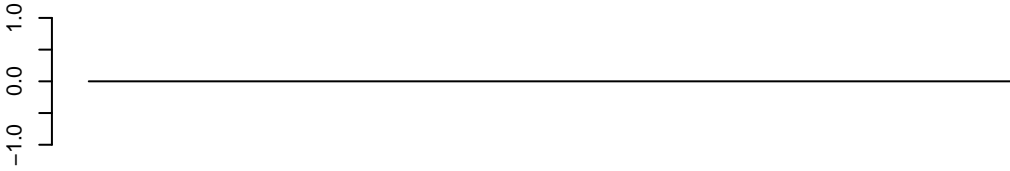




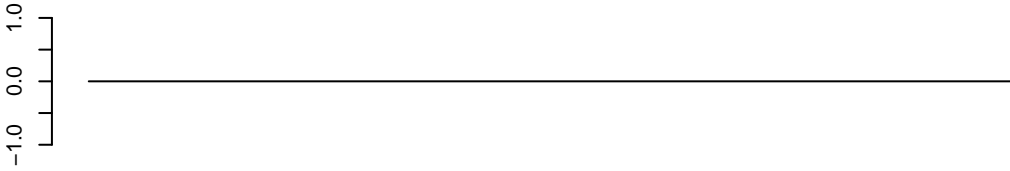
opsin\_4,opsin\_3  
Tadh | no data



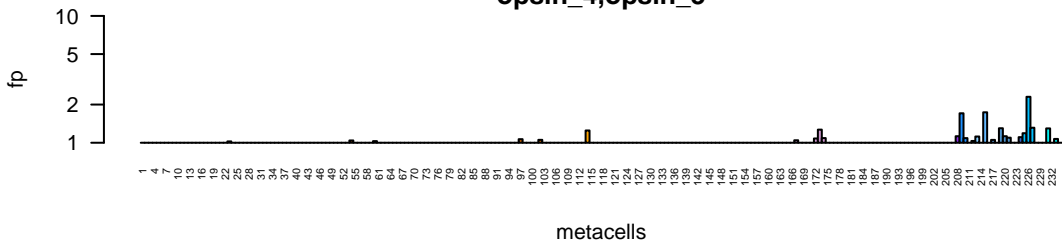
opsin\_4,opsin\_3  
TrH2 | no data



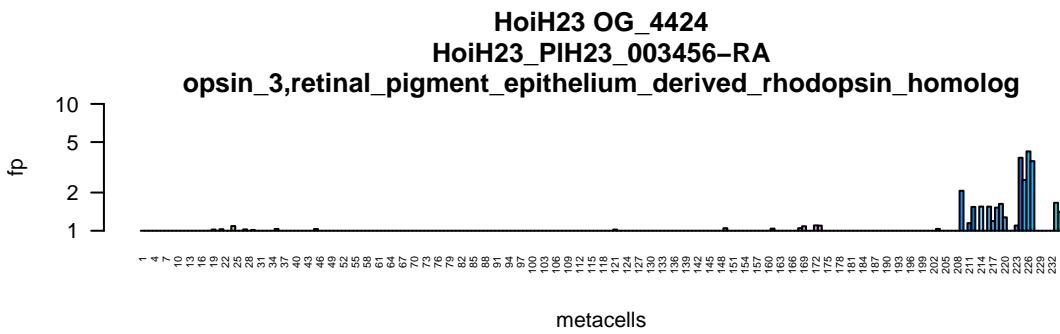
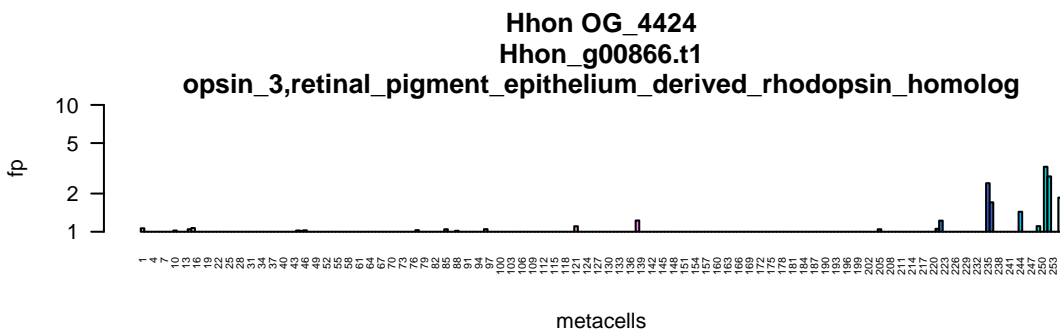
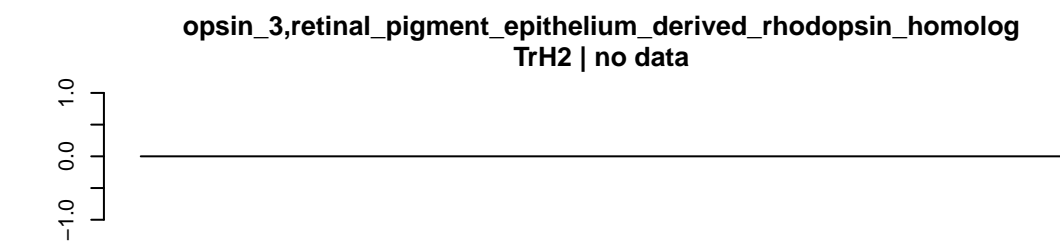
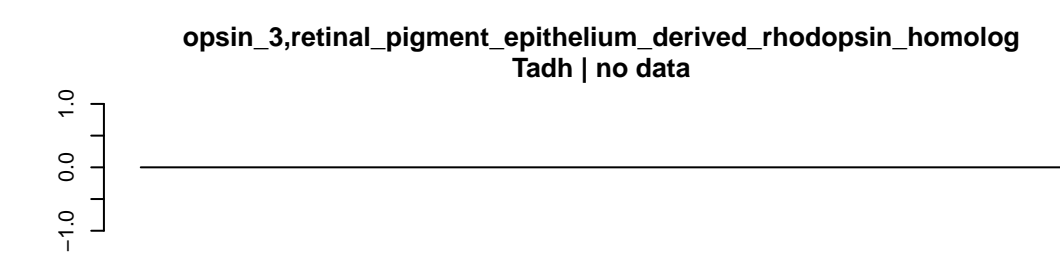
opsin\_4,opsin\_3  
Hhon | no data

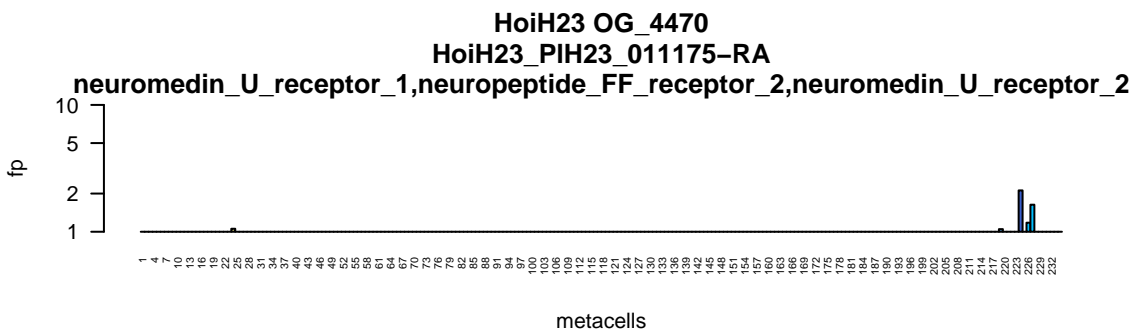
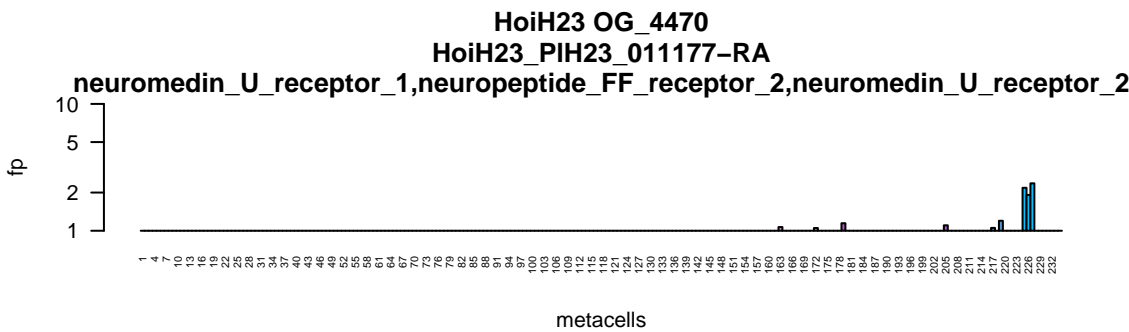
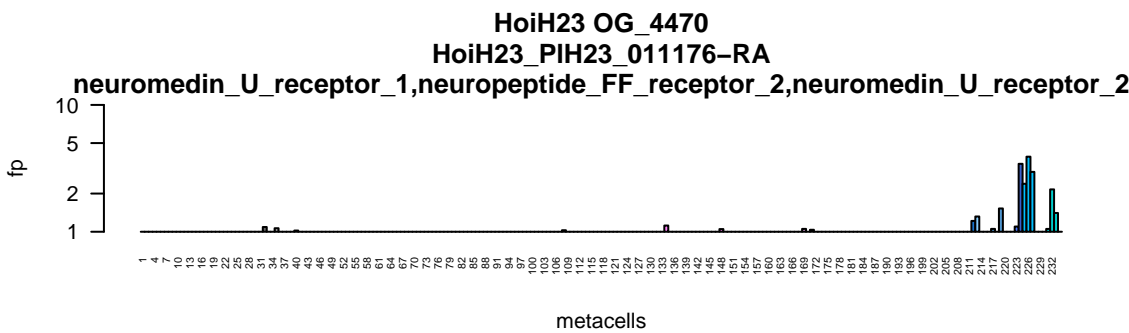
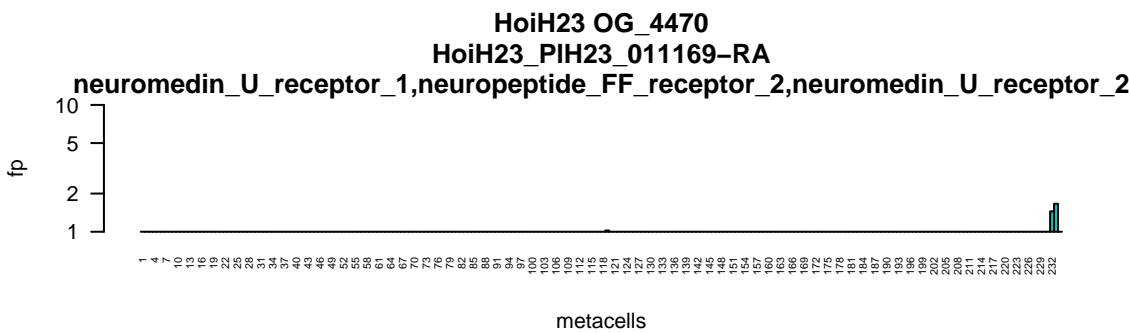
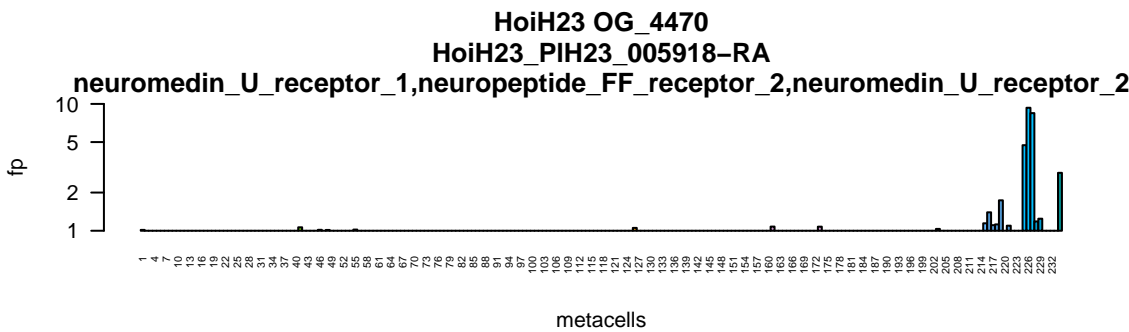
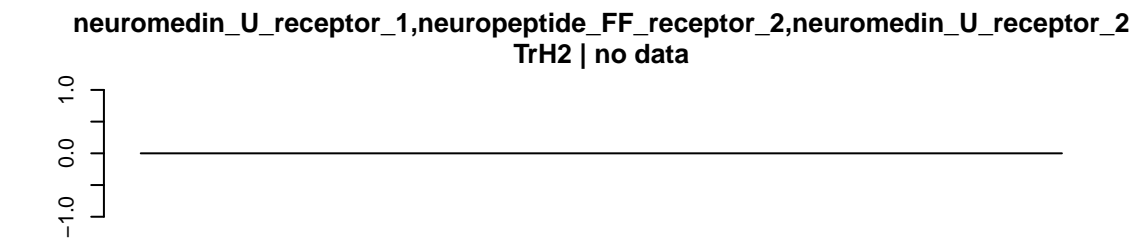
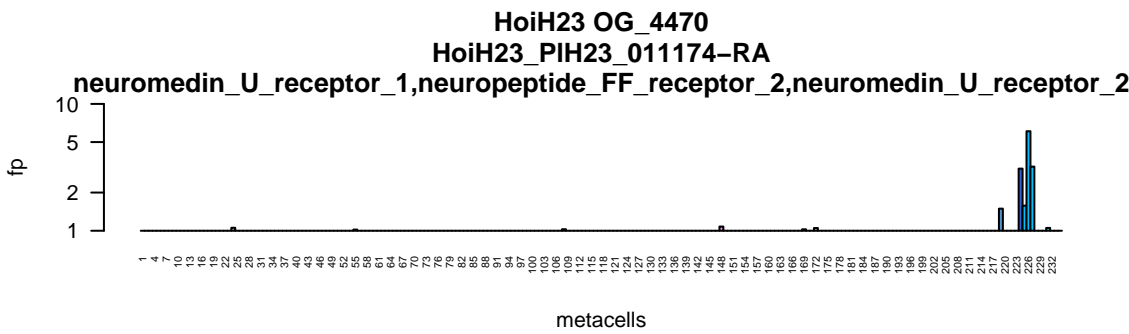
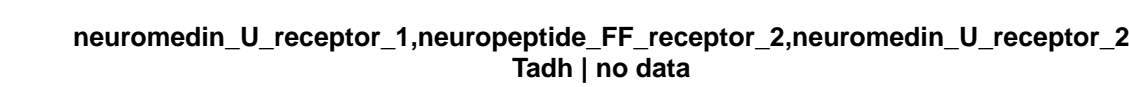


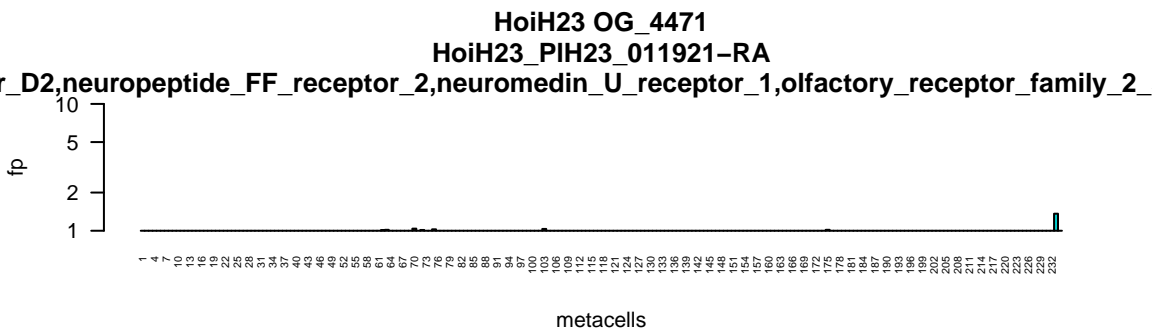
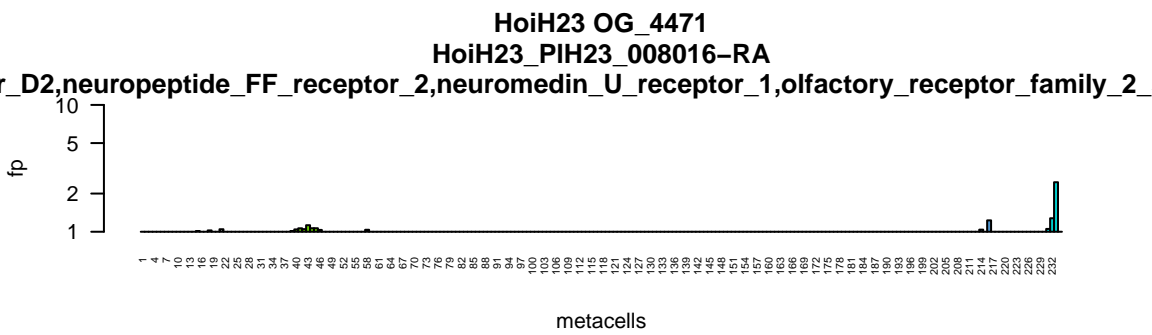
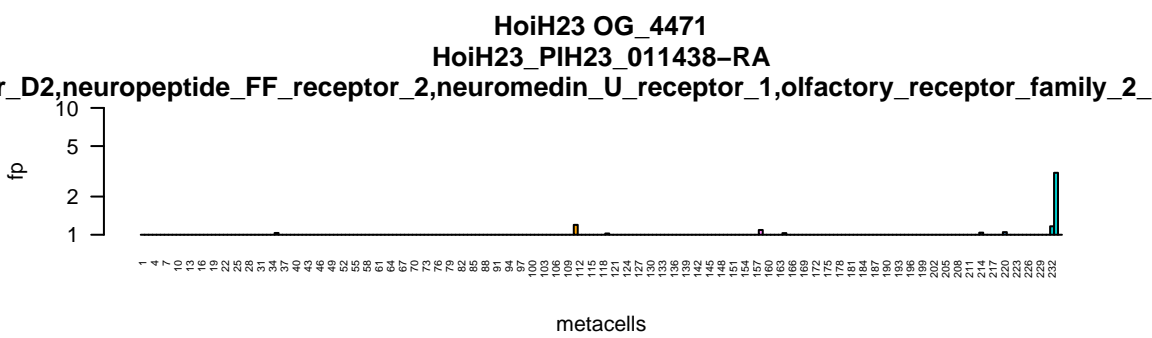
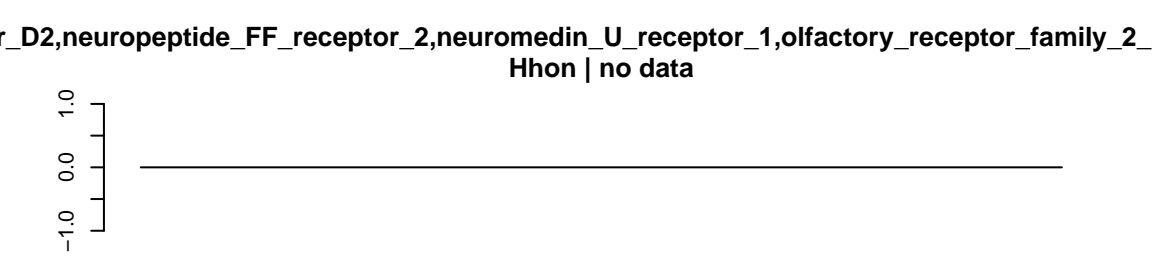
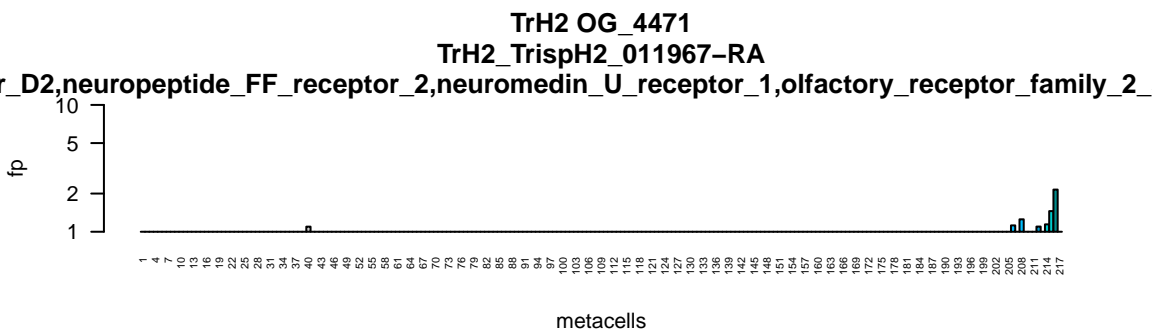
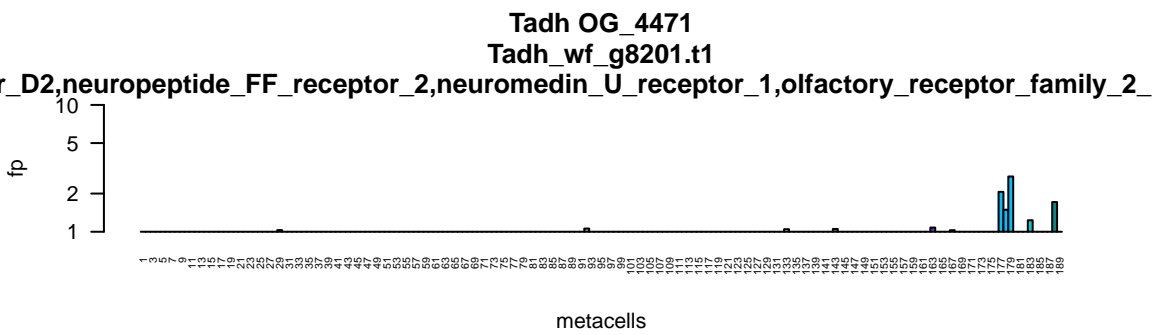
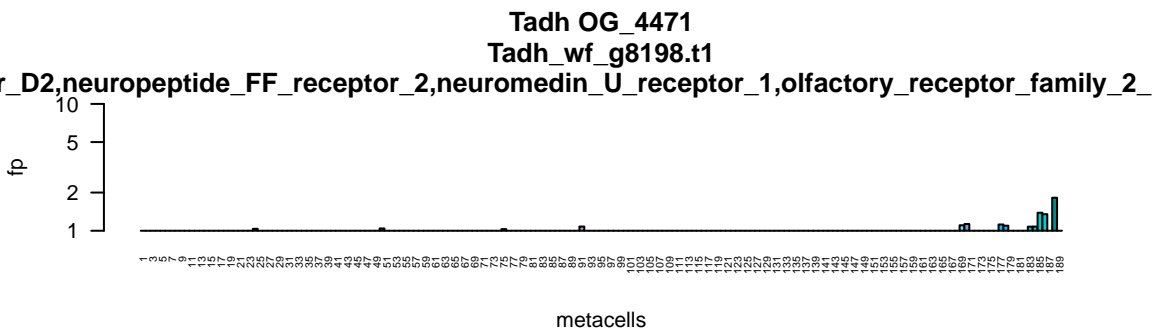
HoiH23 OG\_4423  
HoiH23\_PIH23\_003455--RA  
opsin\_4,opsin\_3

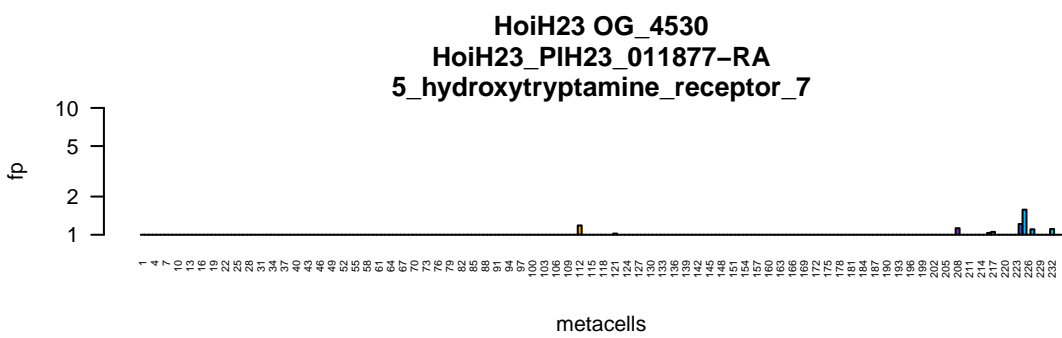
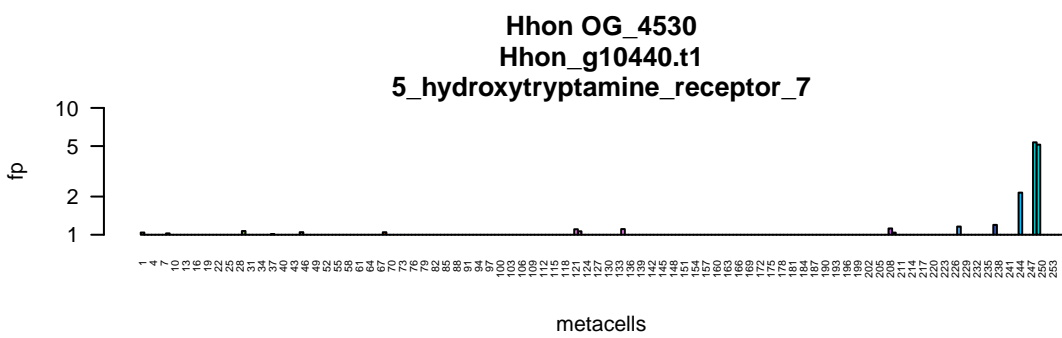
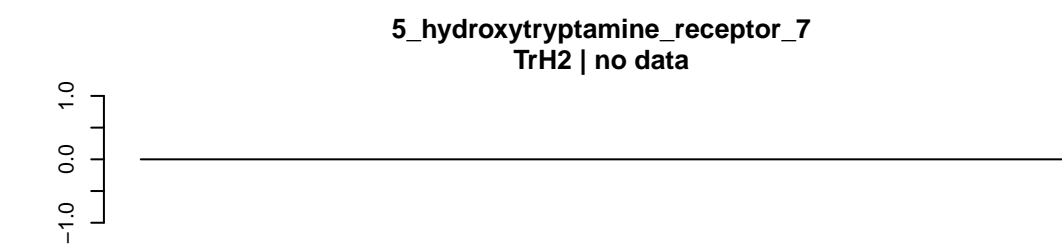
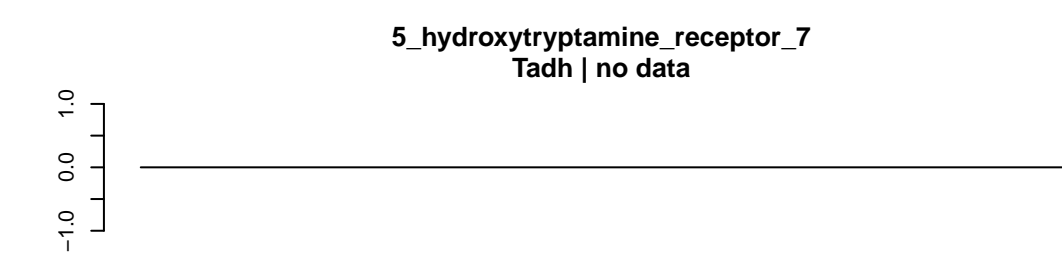


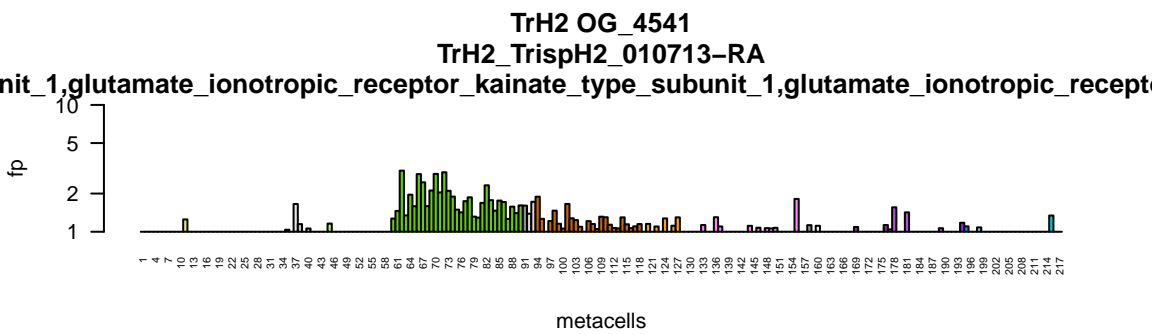
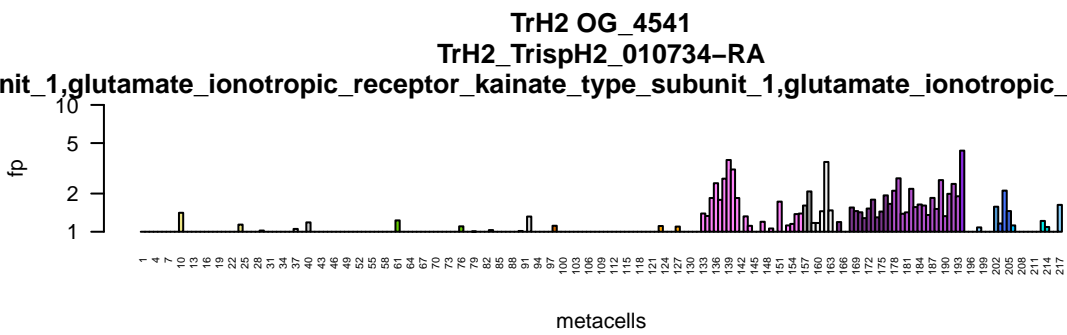
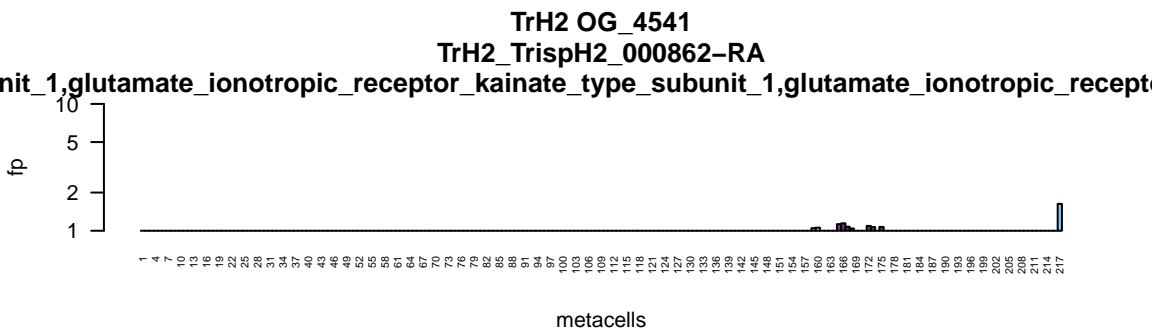
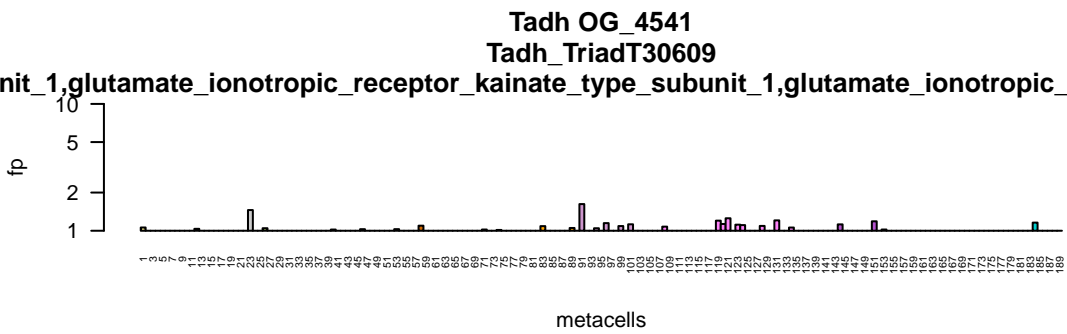
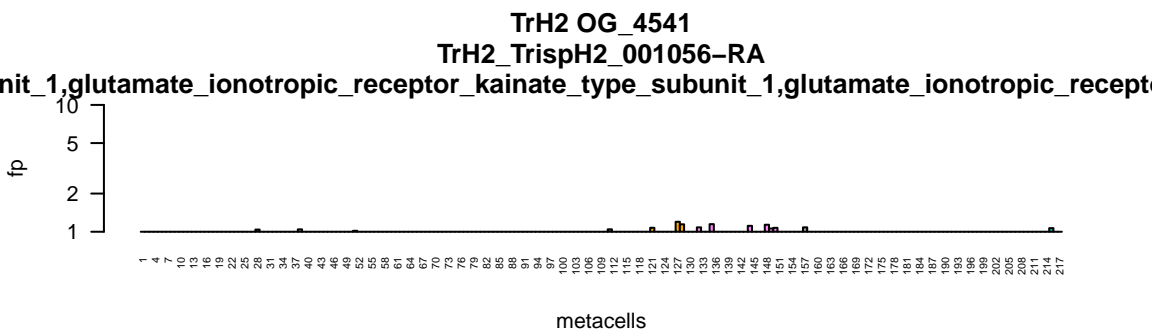
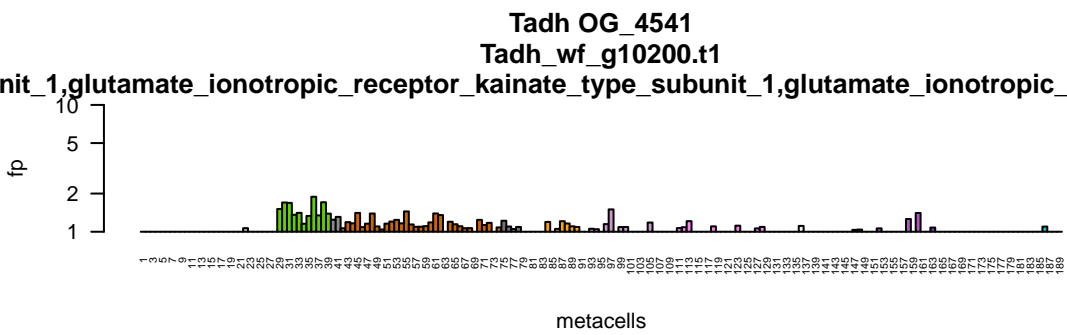
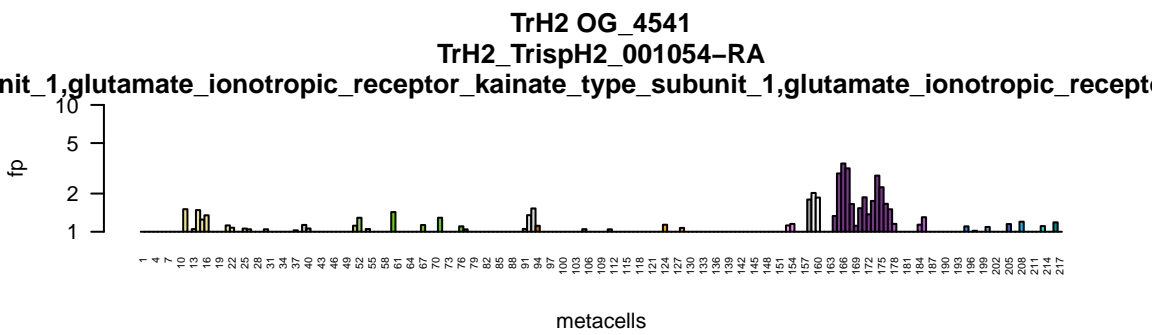
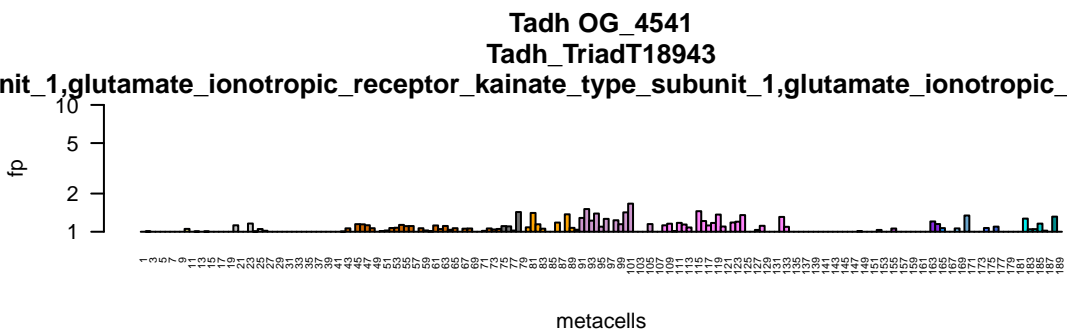
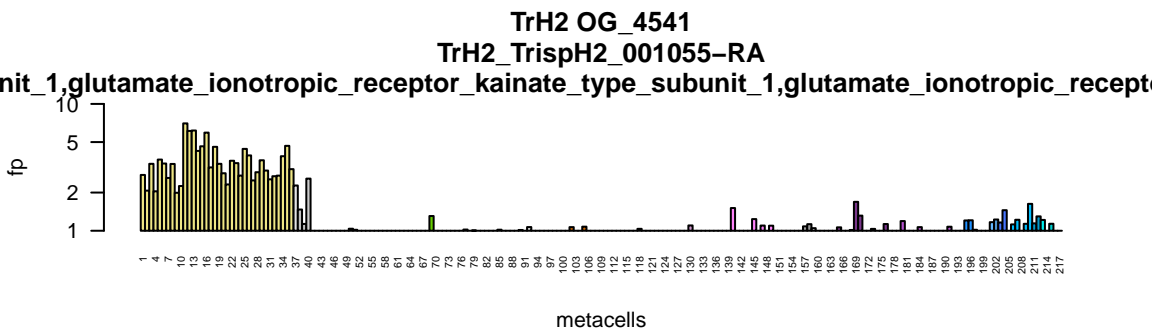
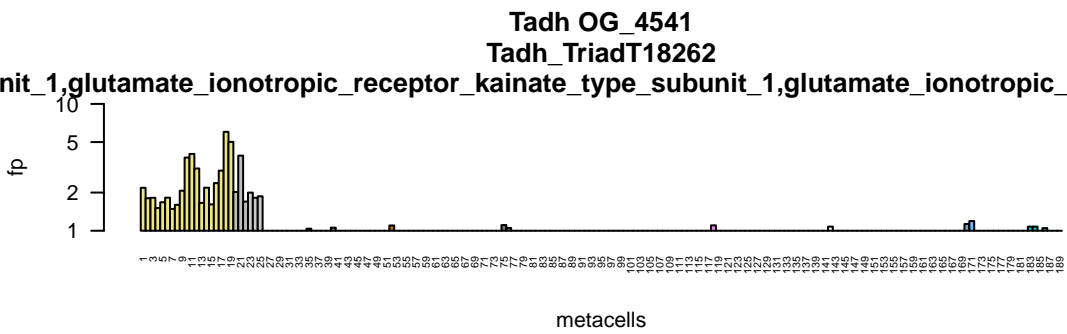
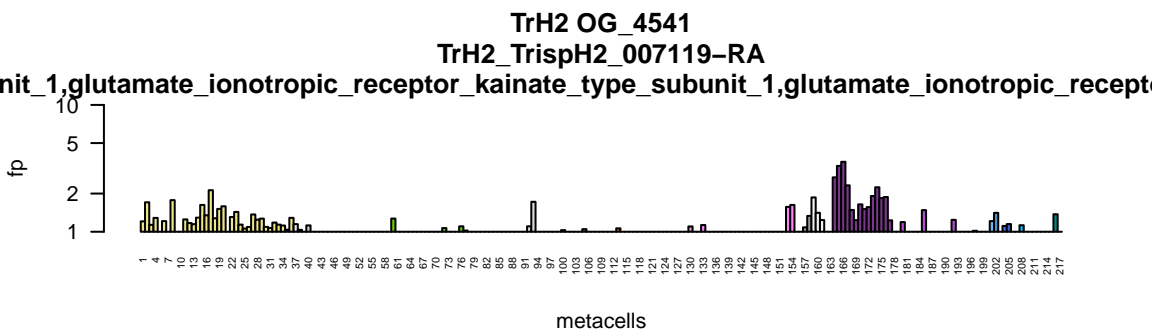
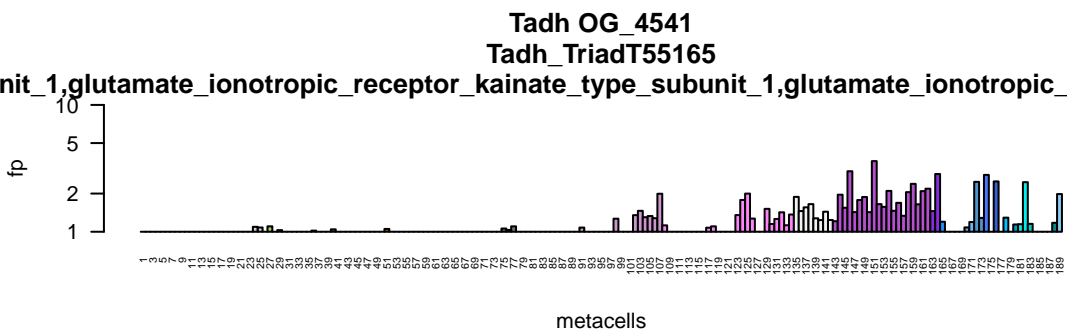
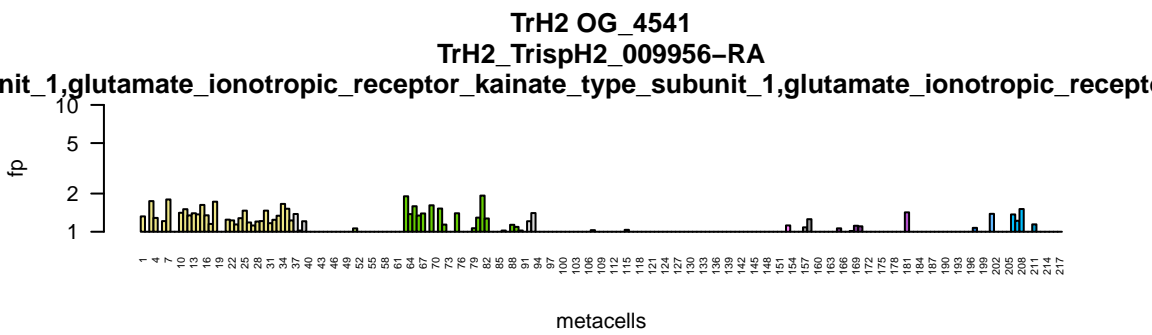
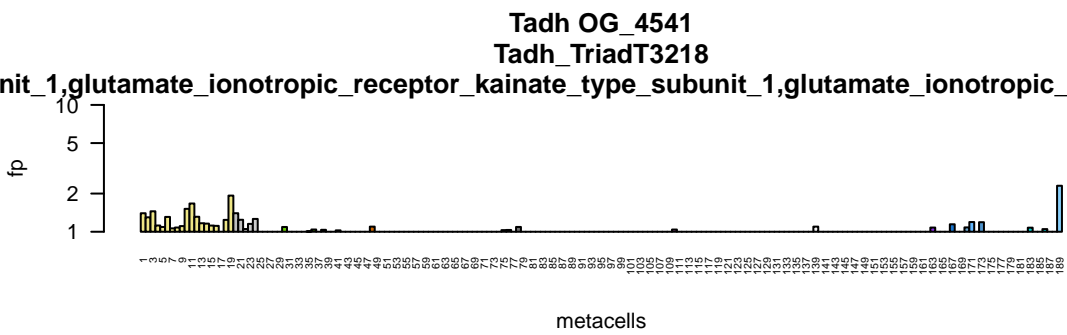
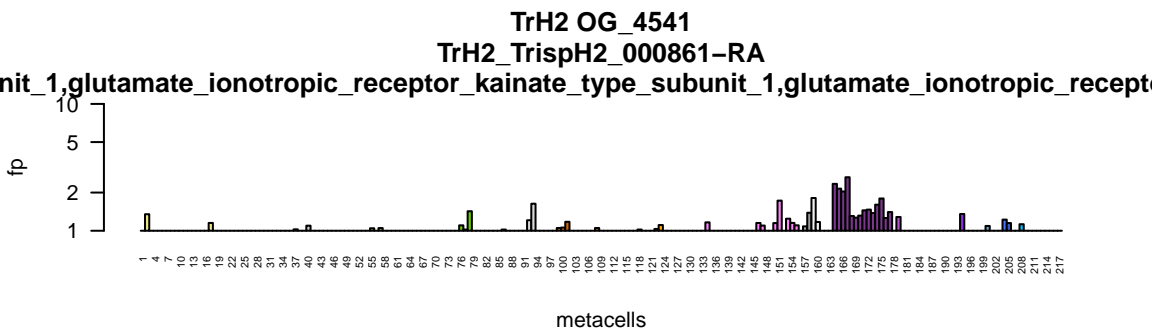
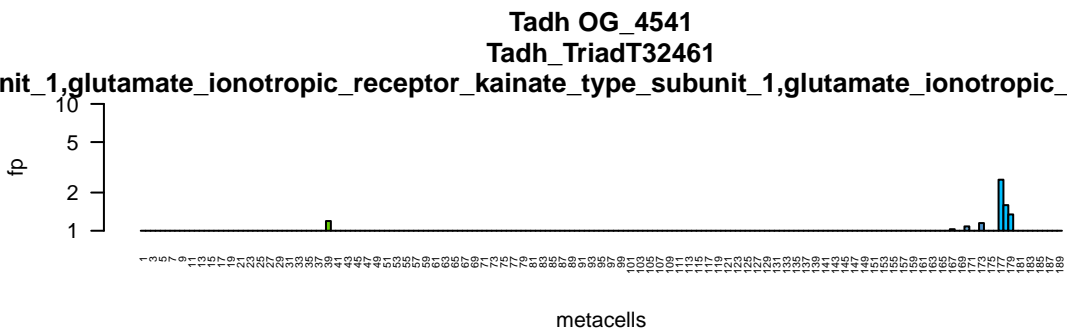


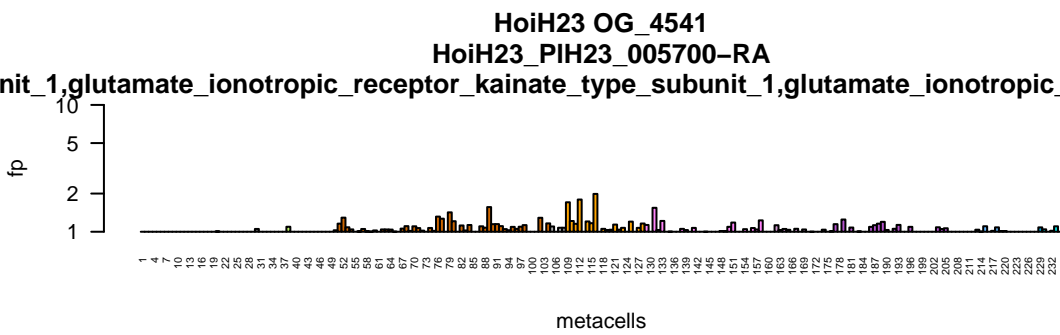
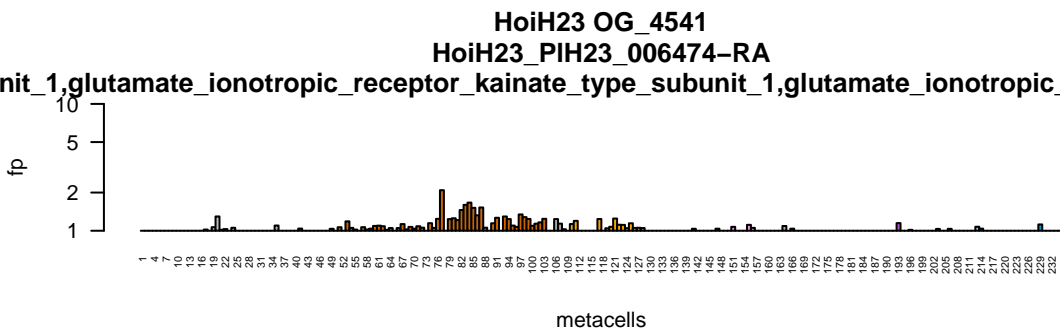
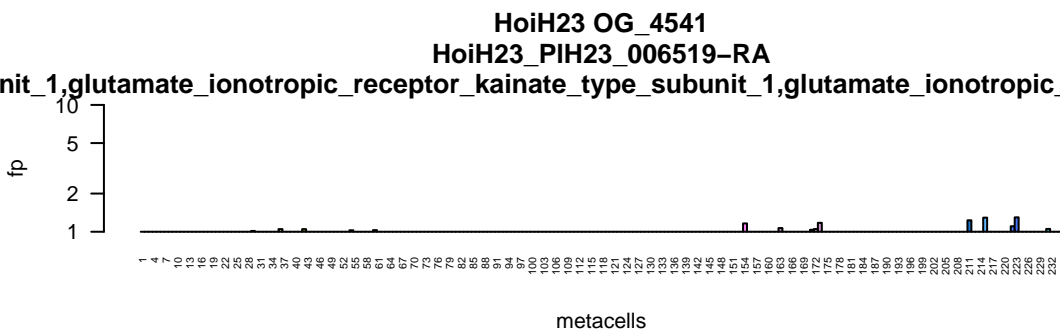
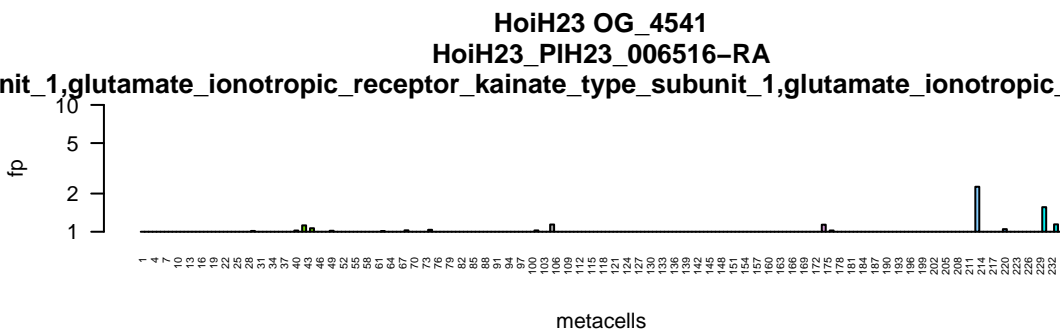
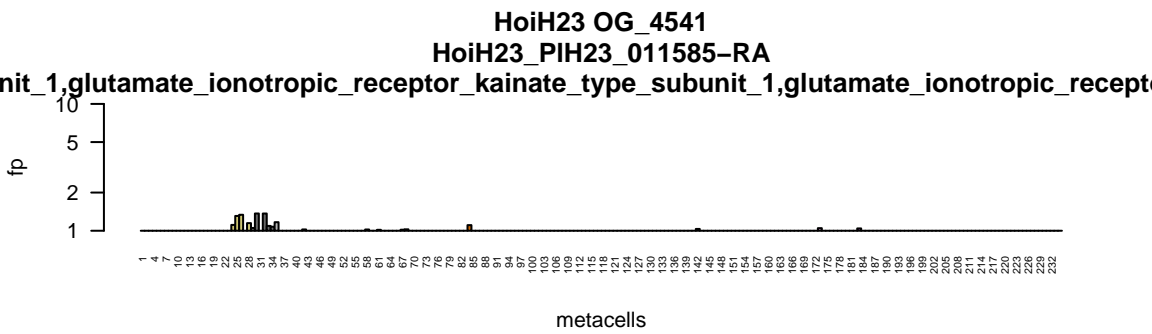
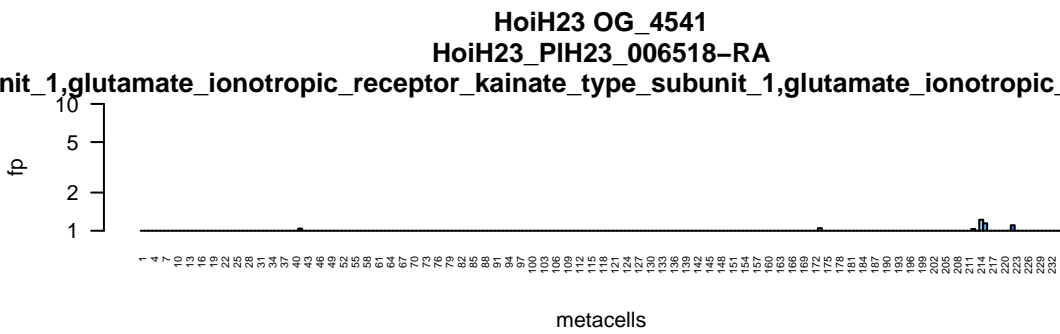
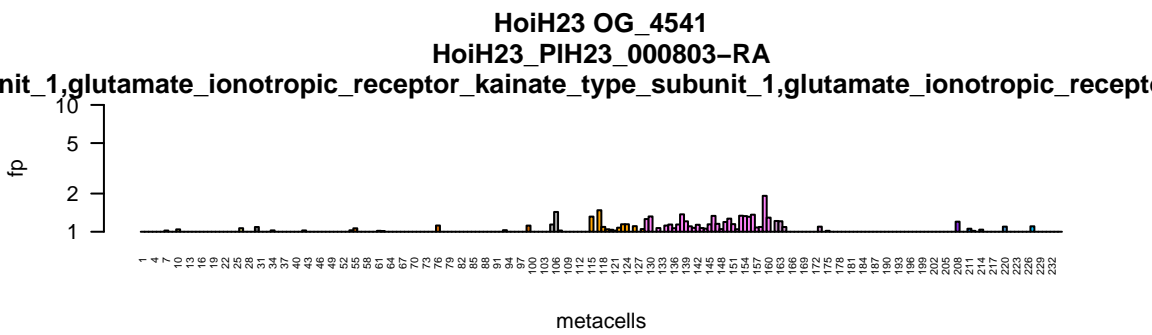
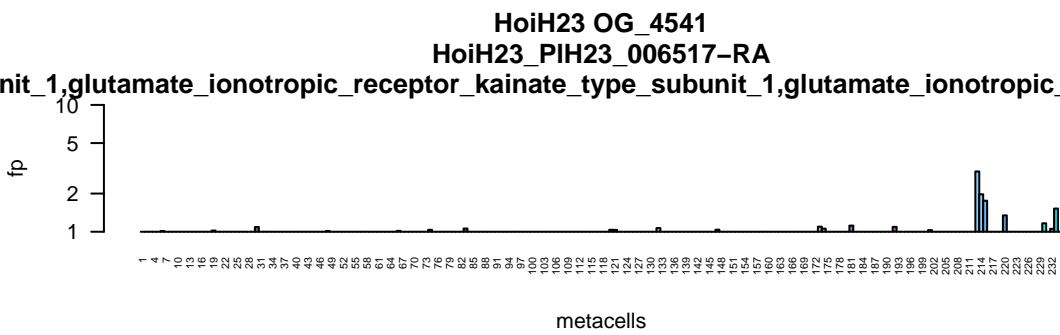
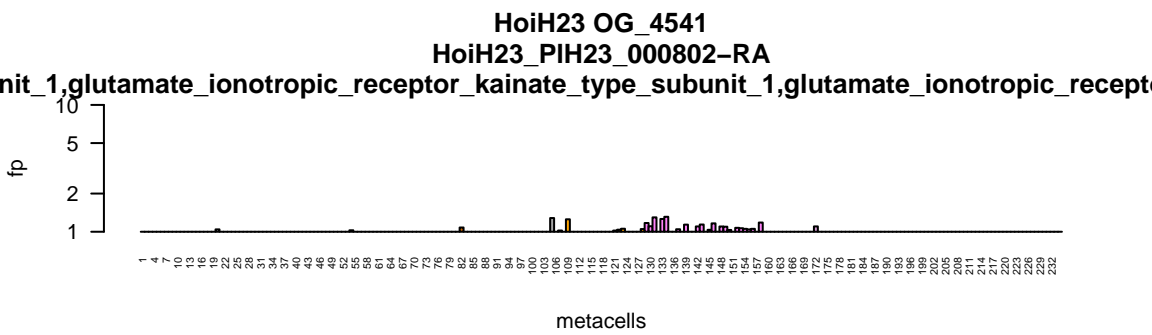
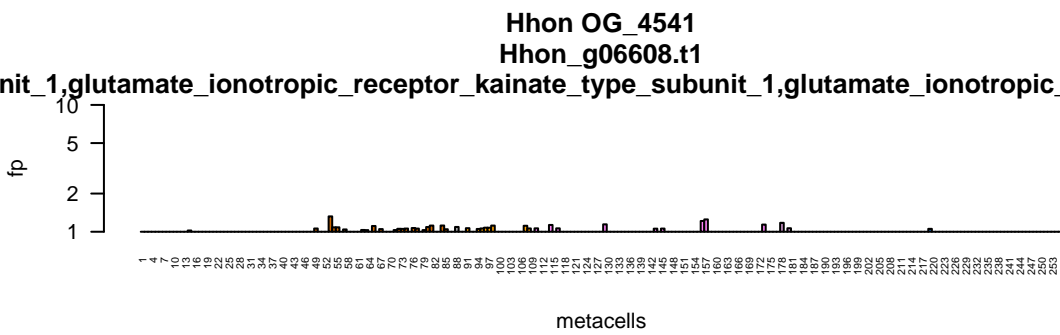
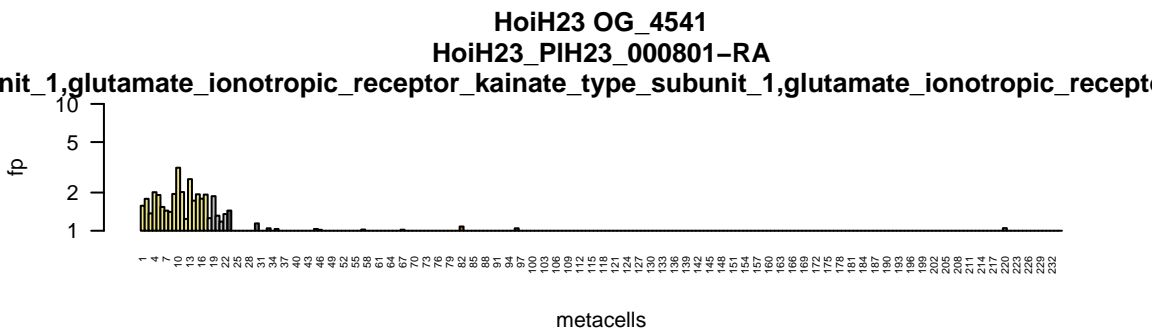
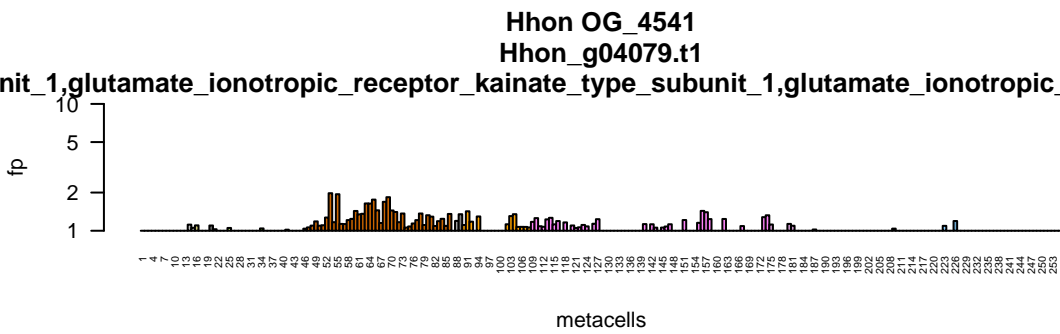


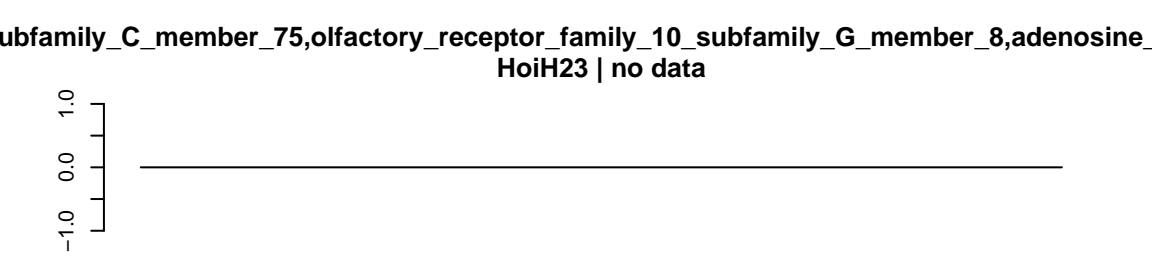
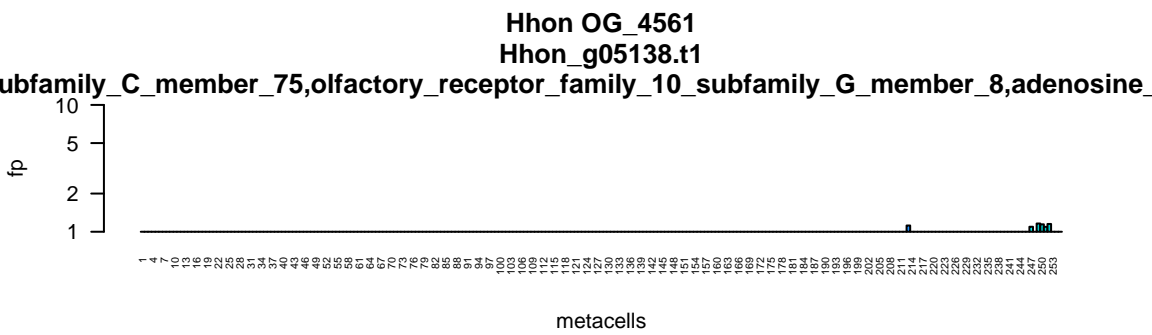
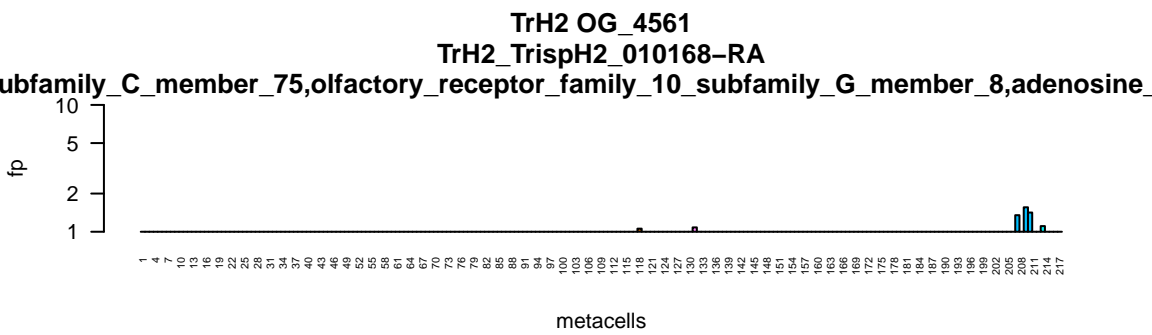
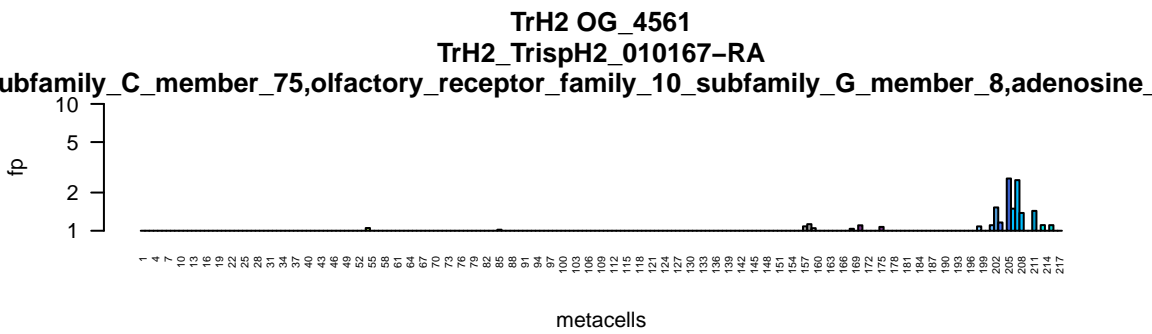
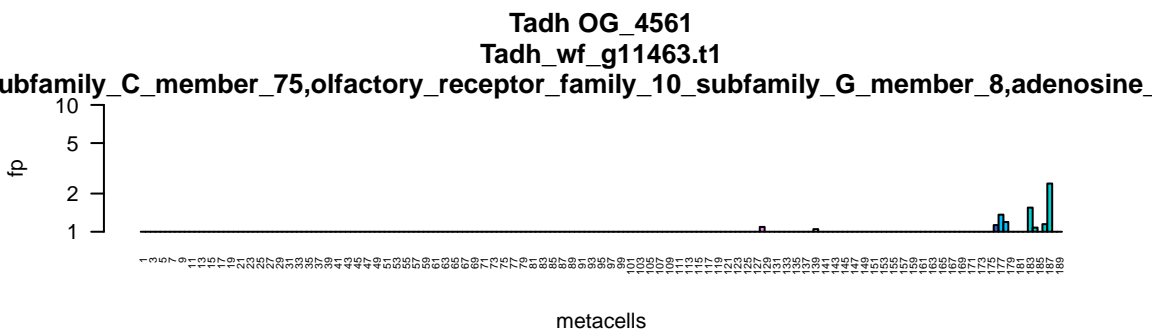
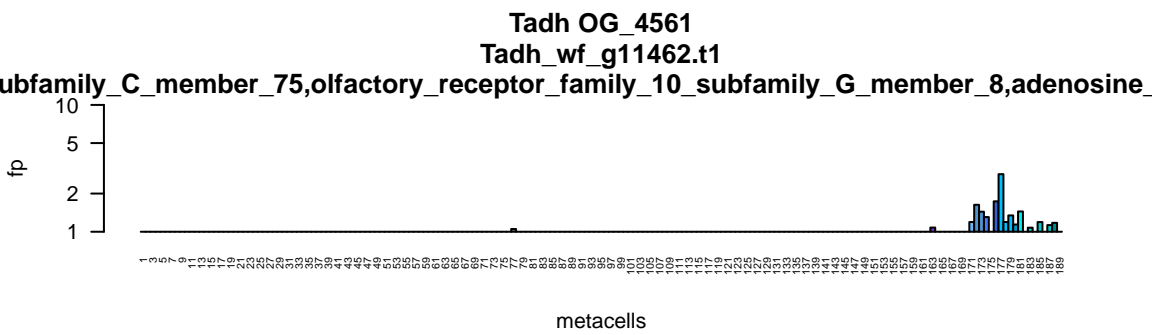
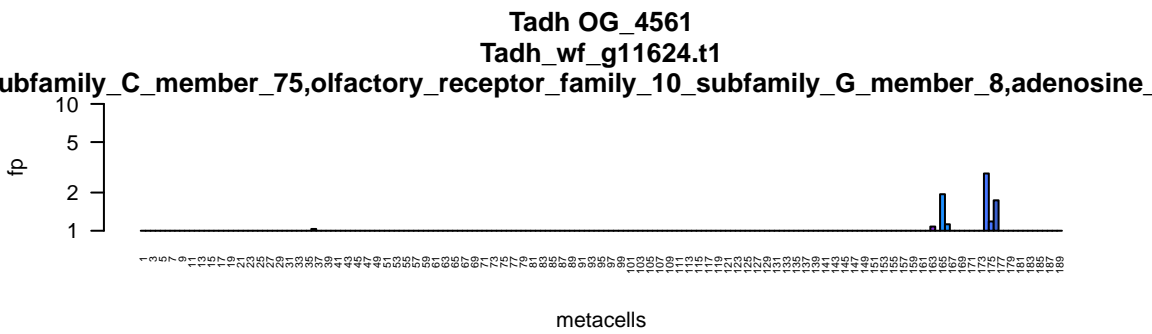


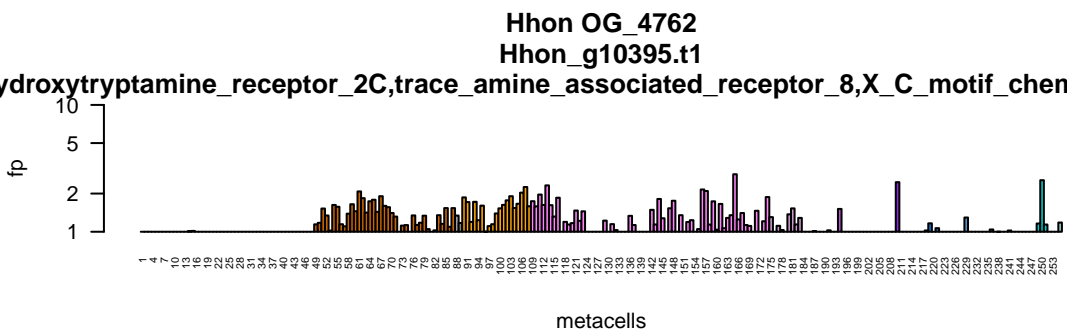
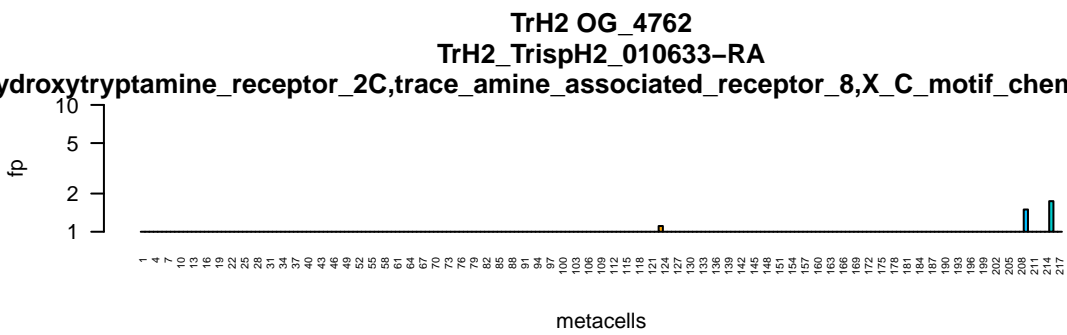
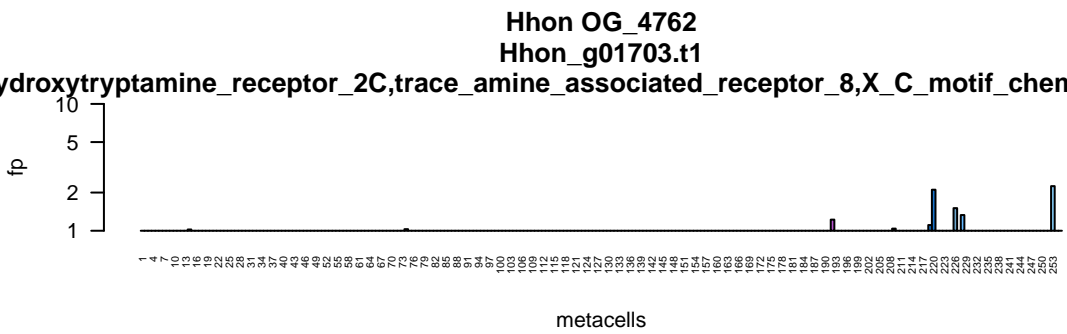
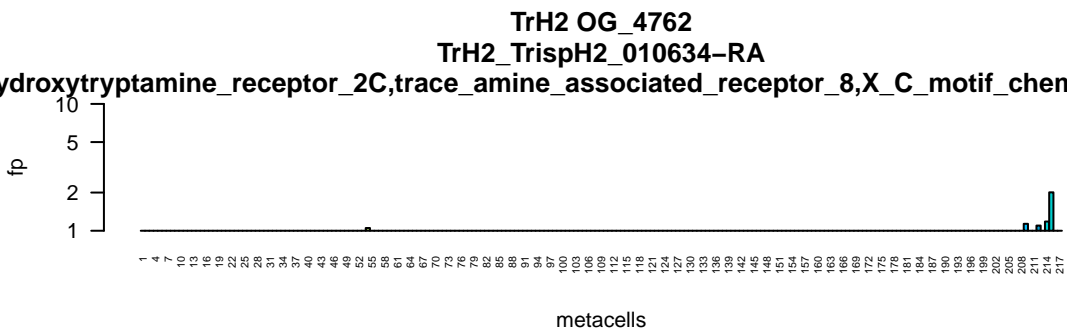
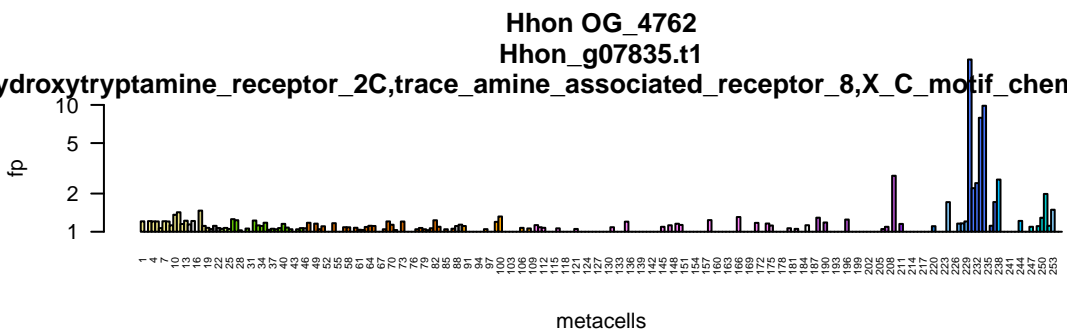
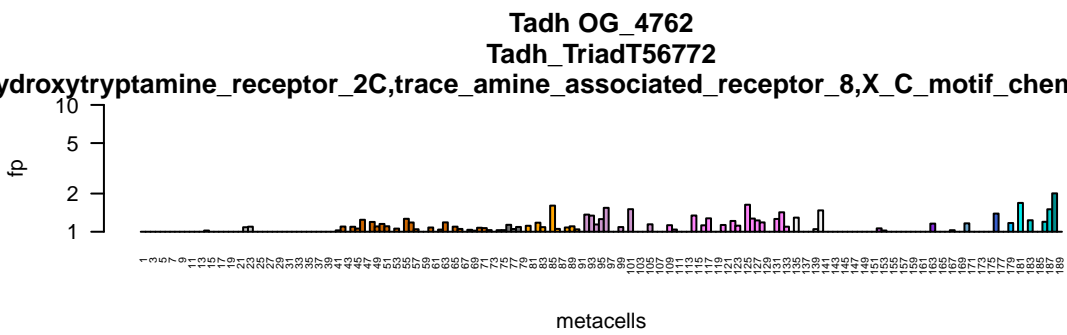
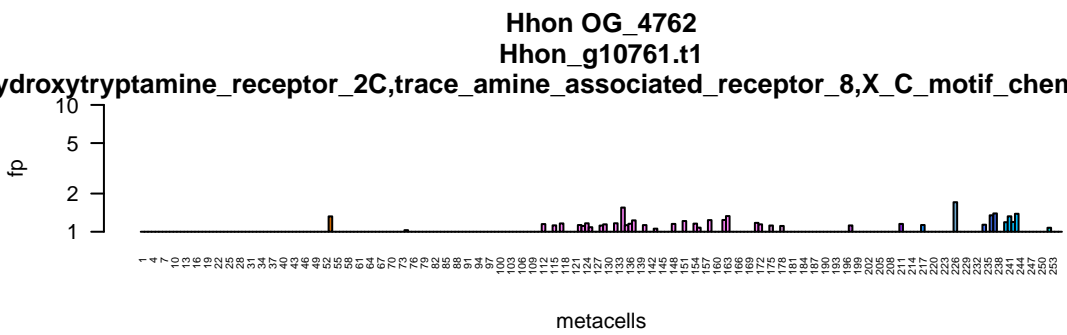
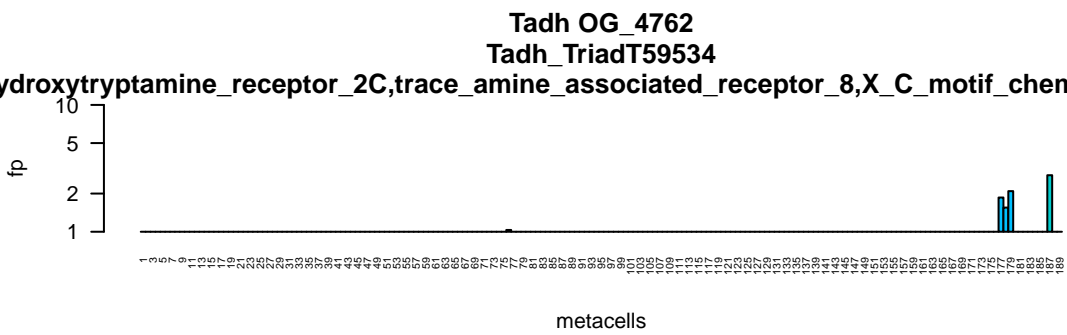
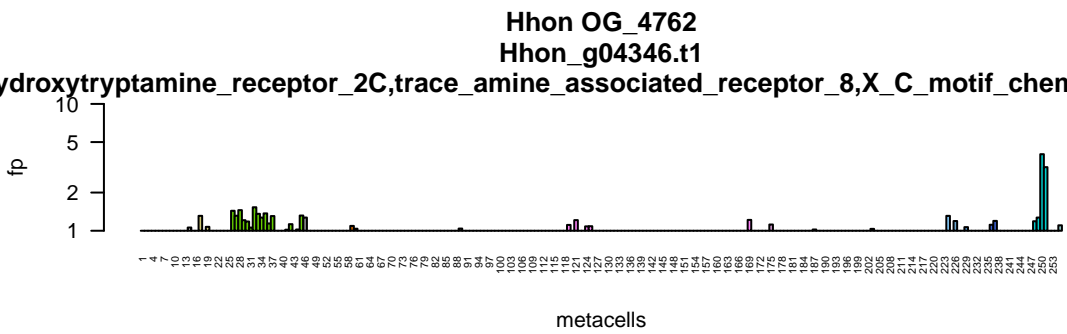
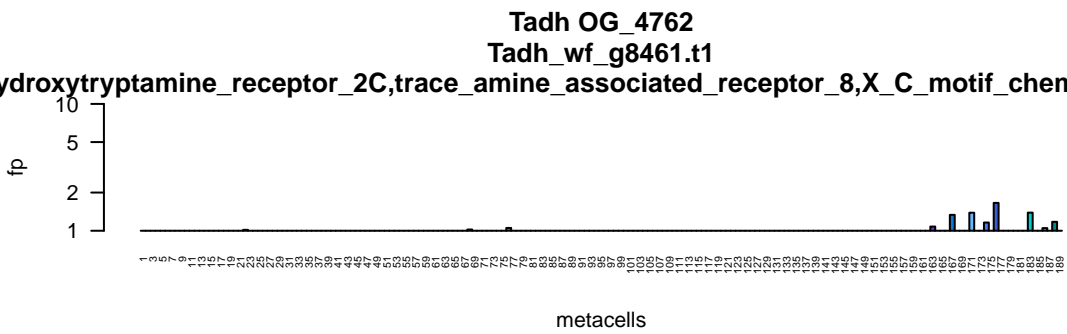
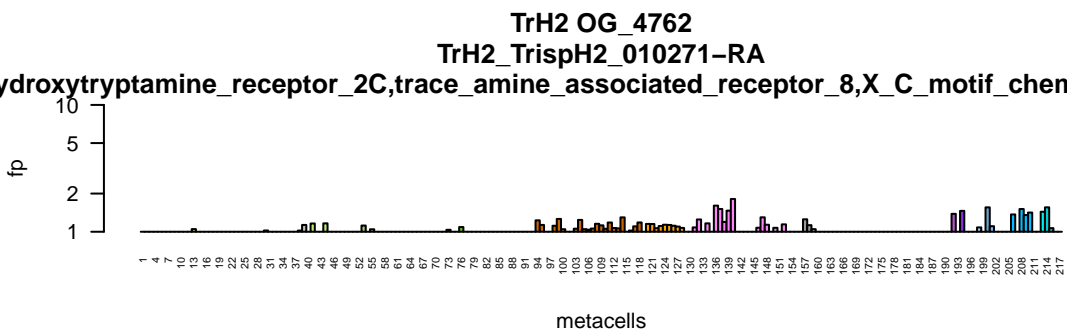
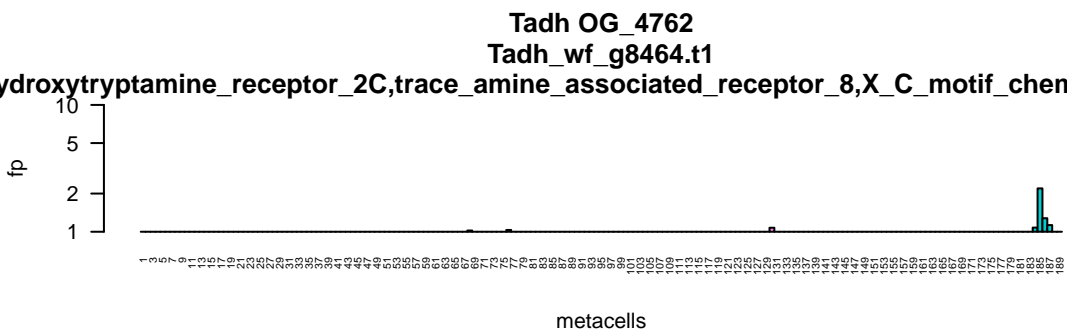
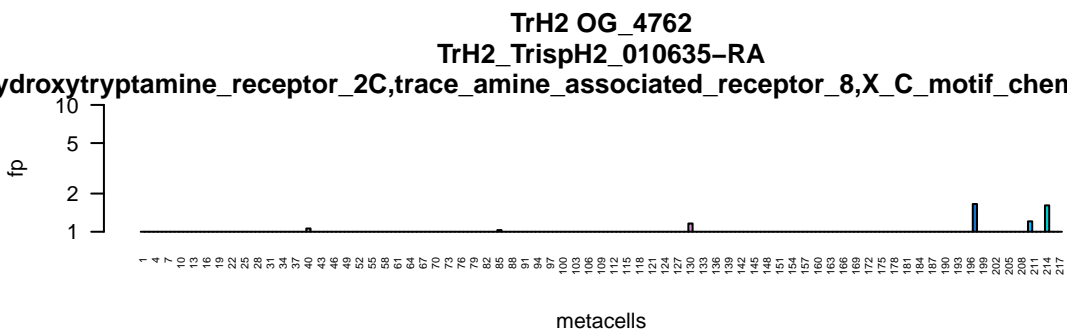
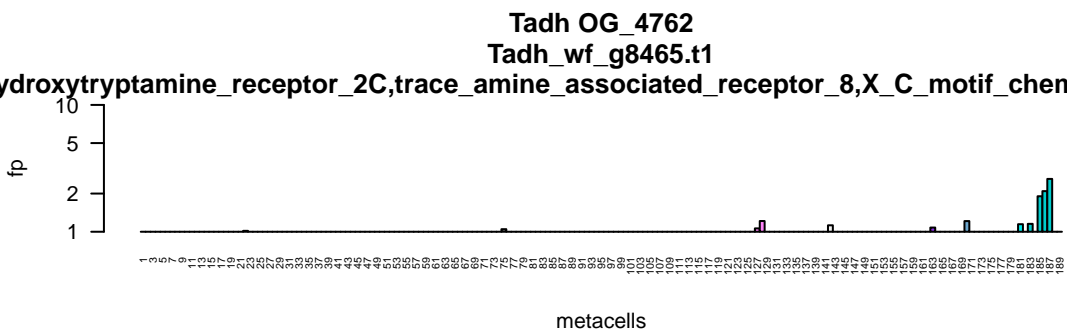
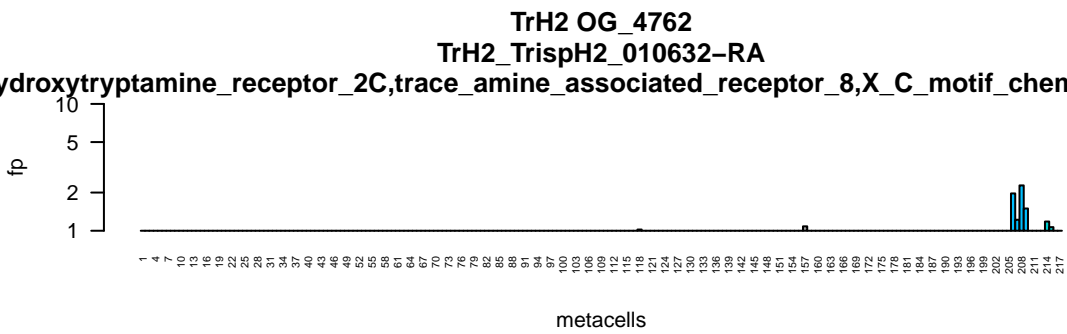
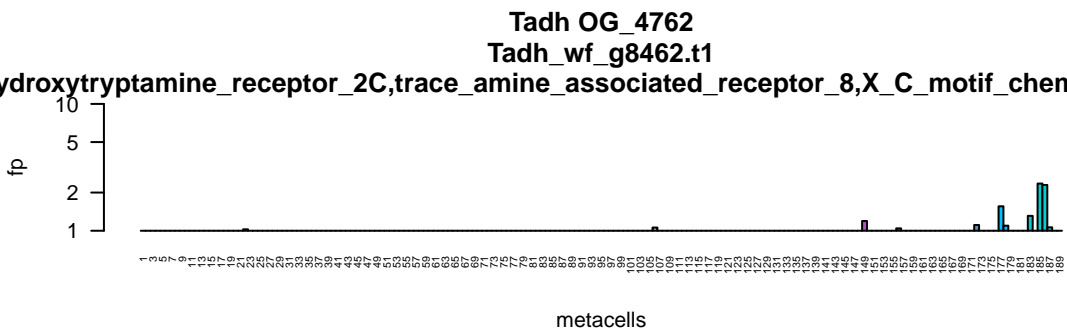




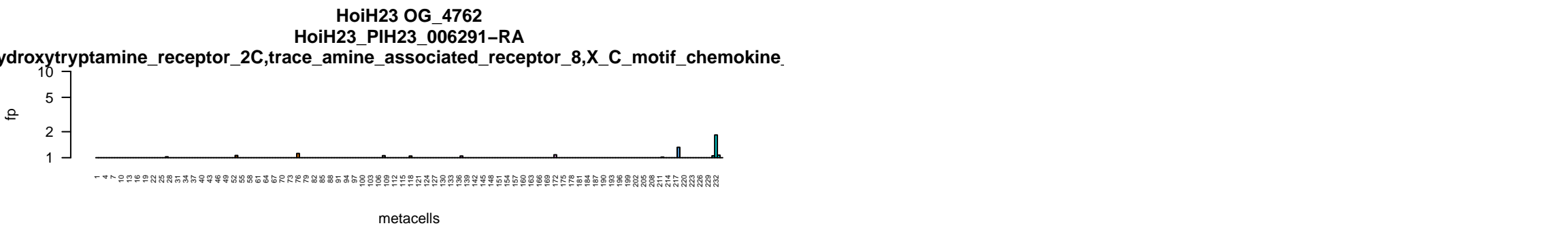
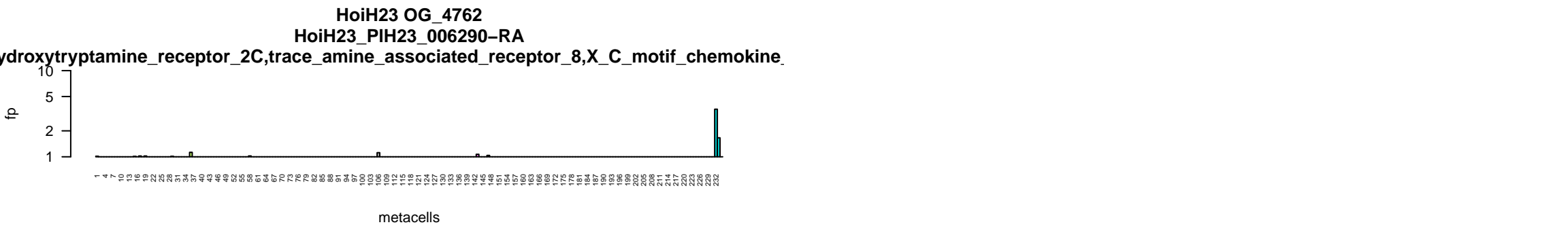
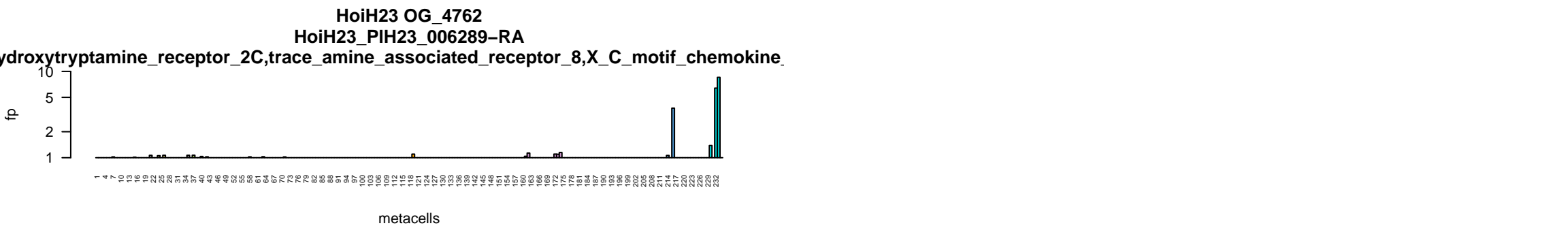
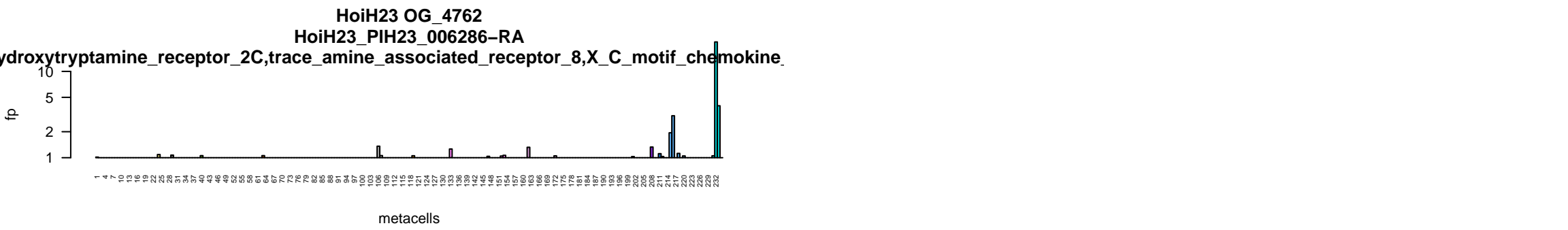
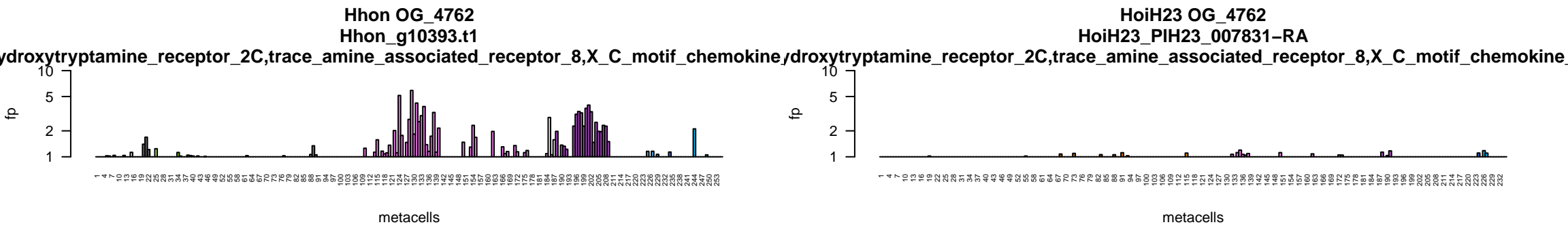


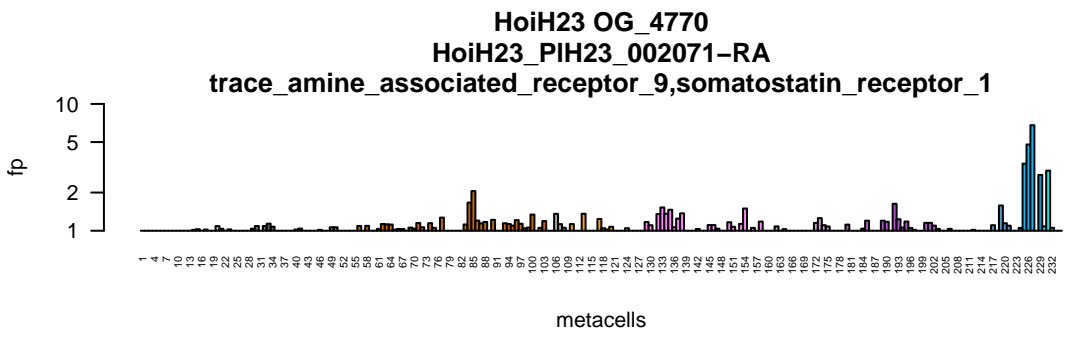
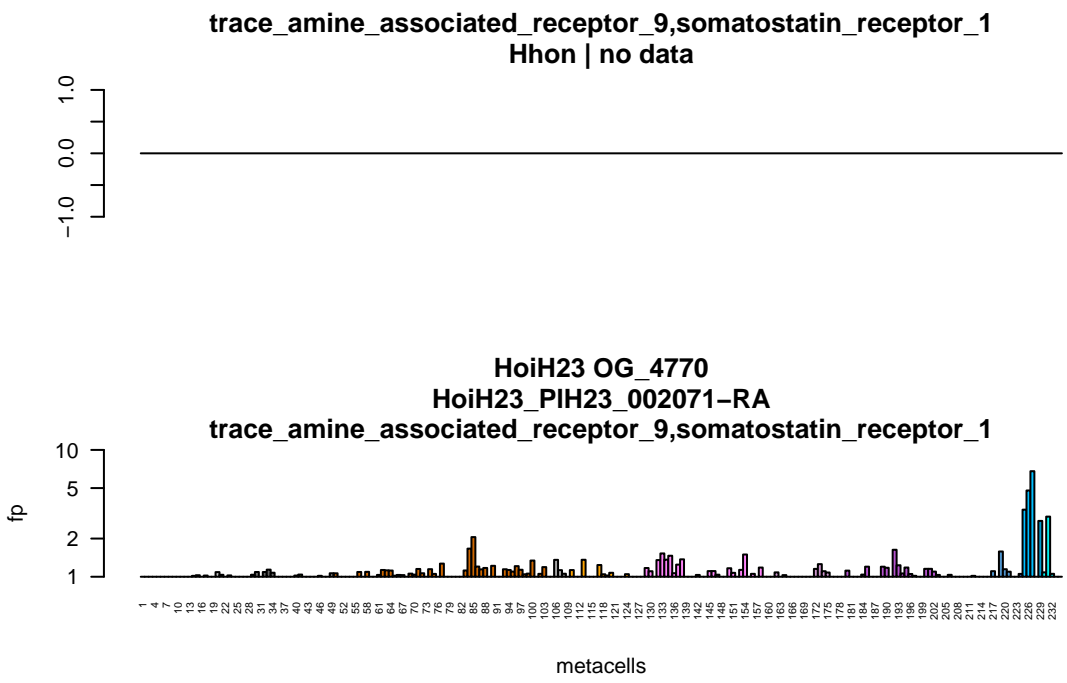
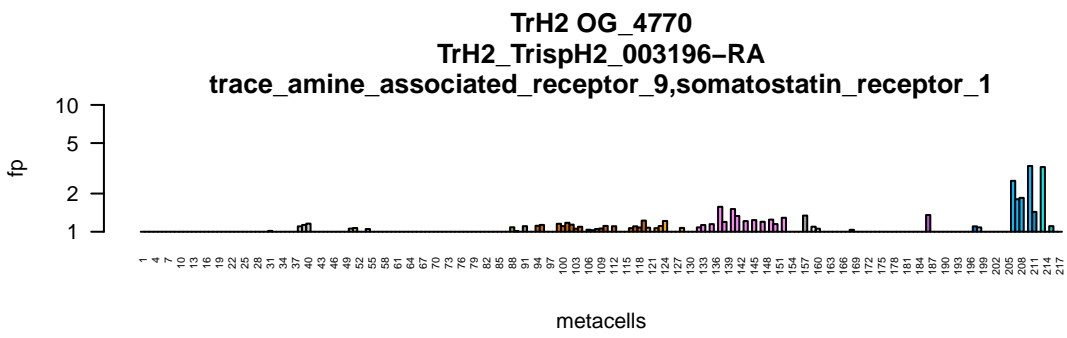
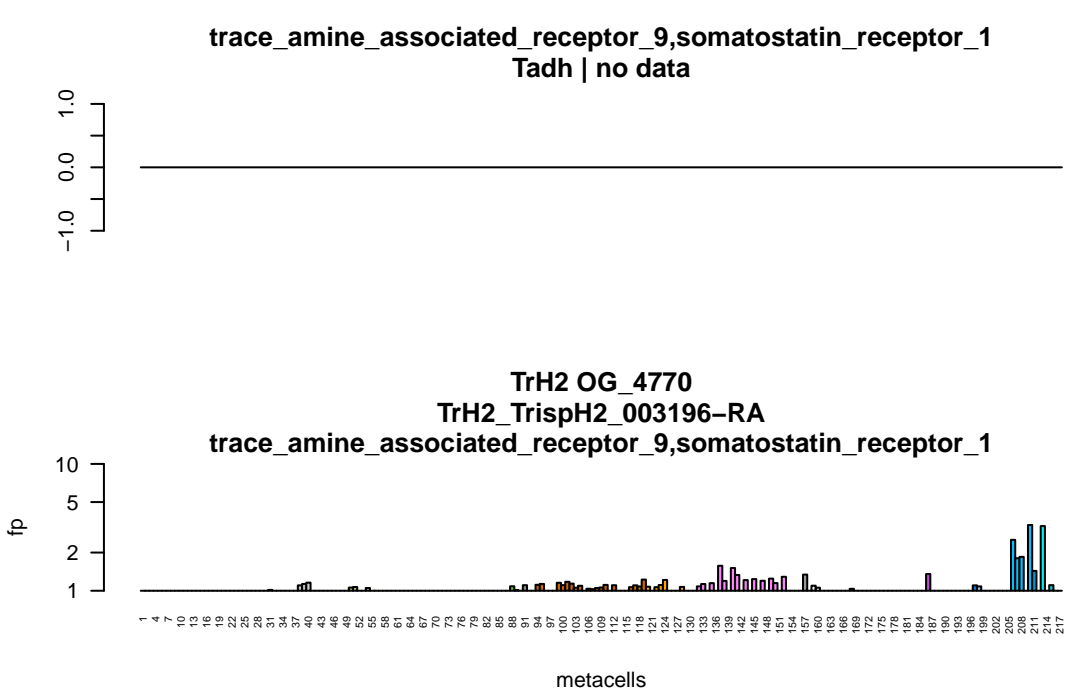


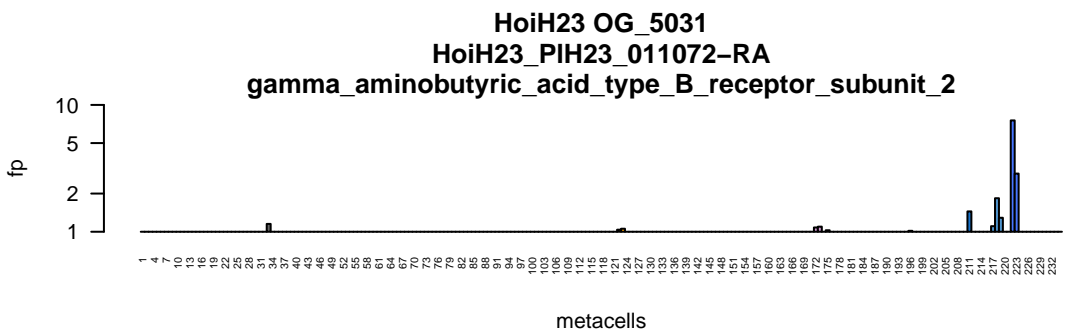
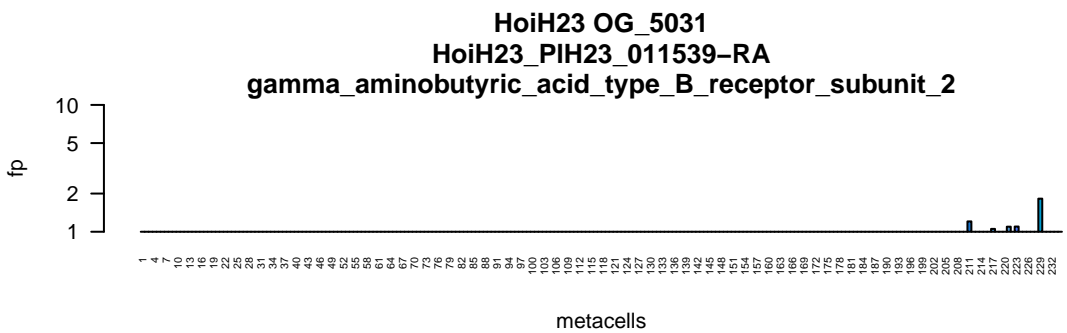
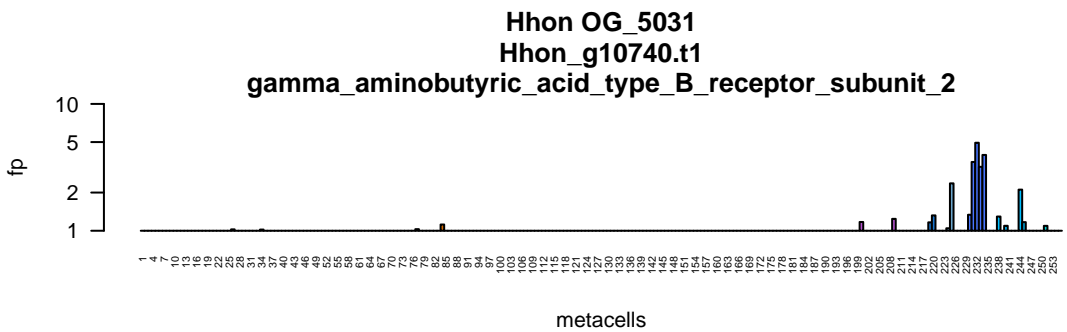
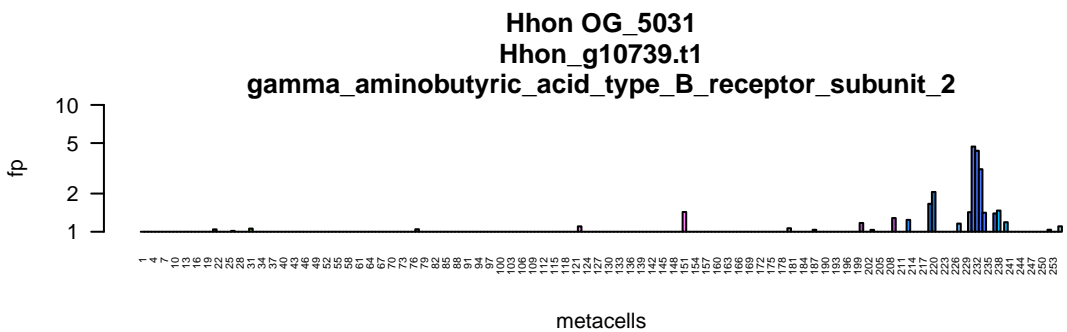
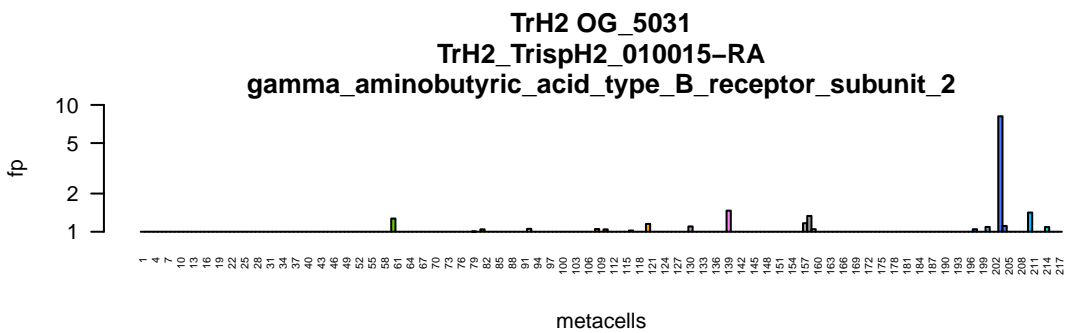
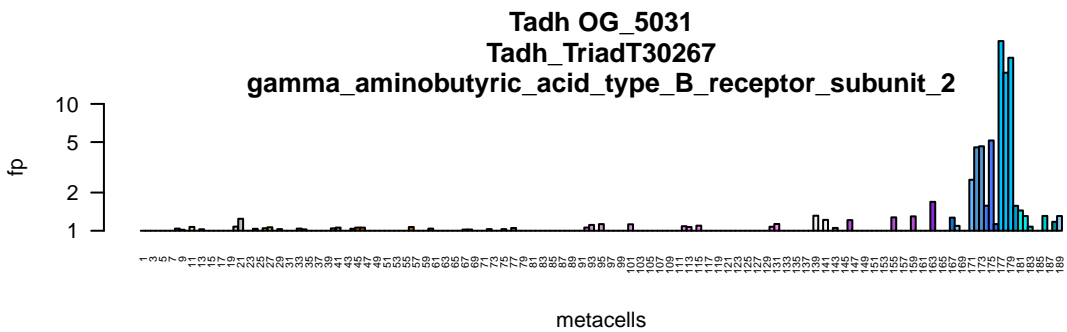


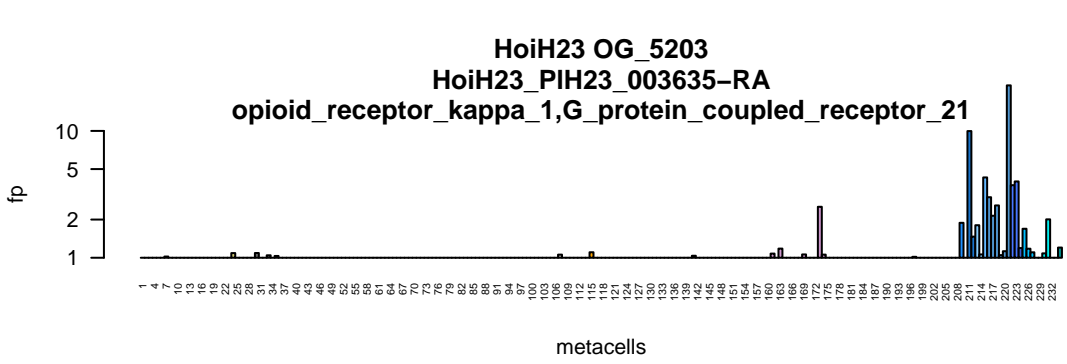
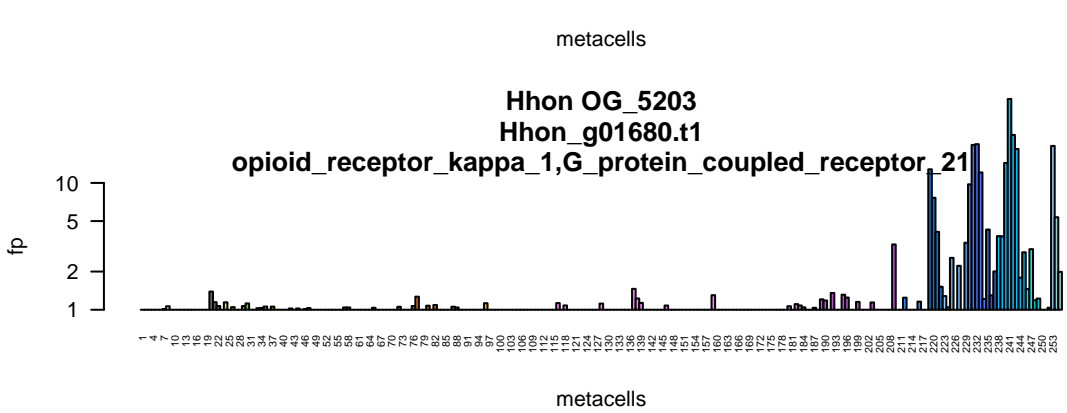
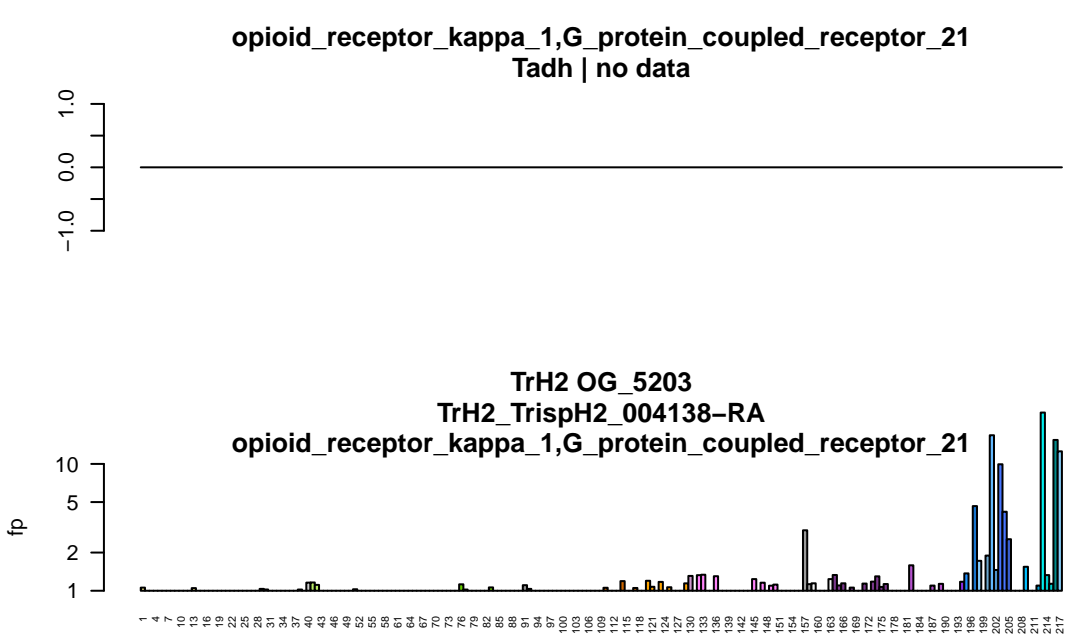


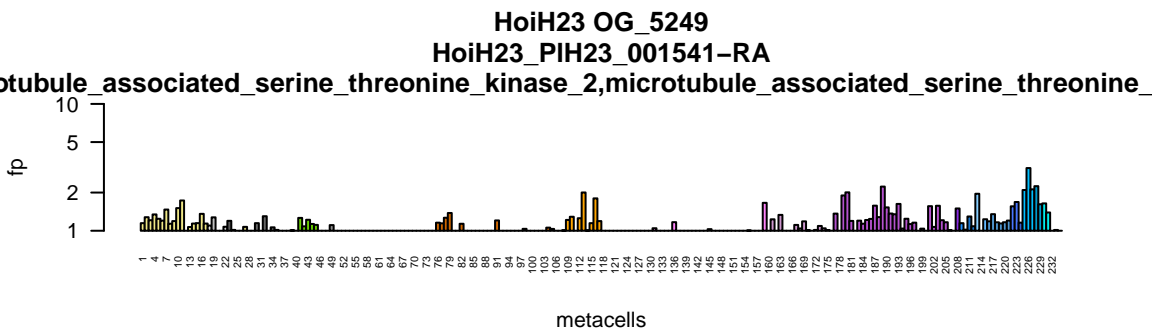
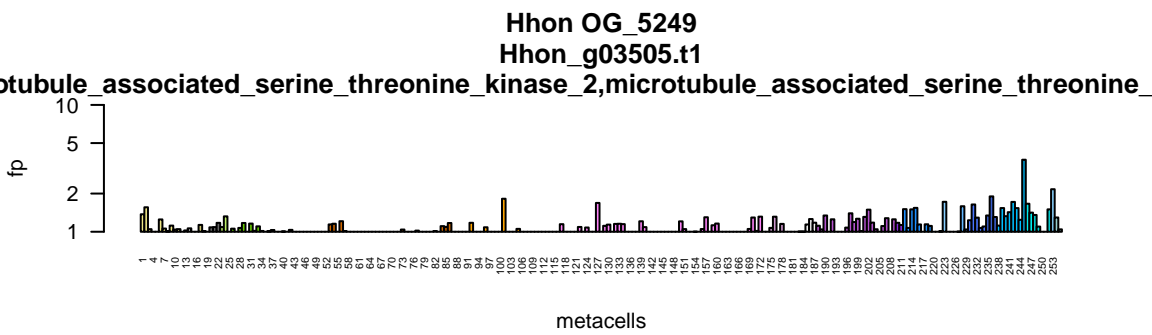
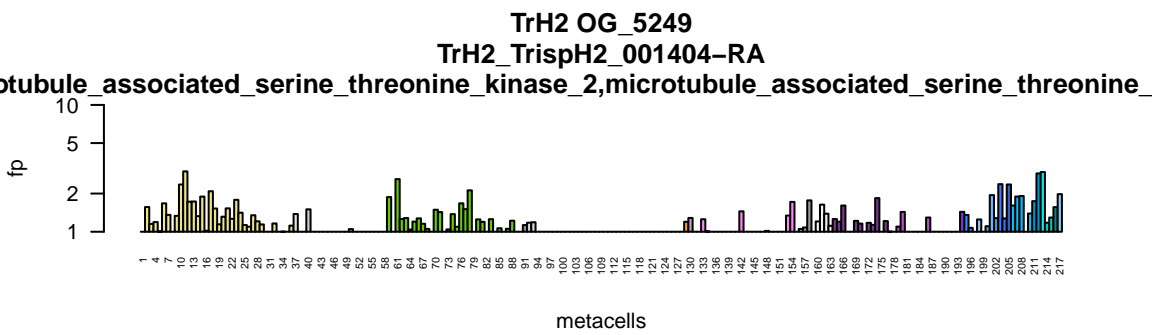
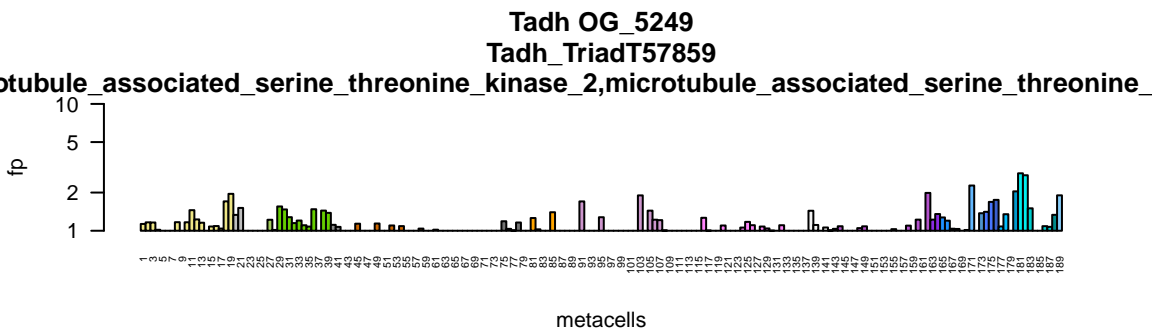


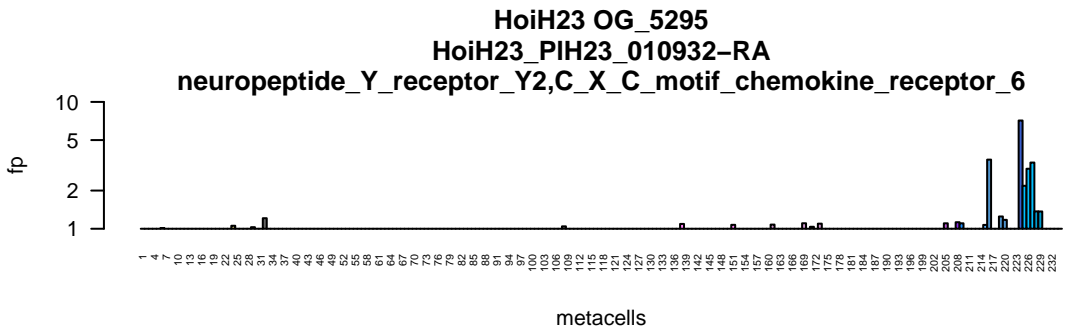
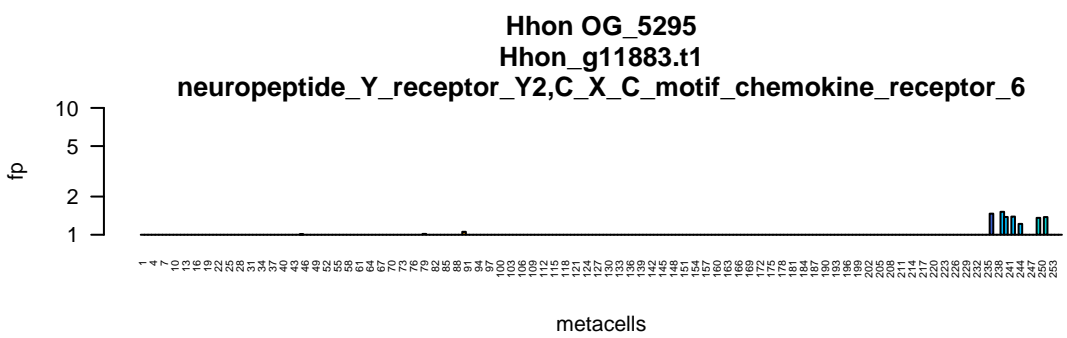
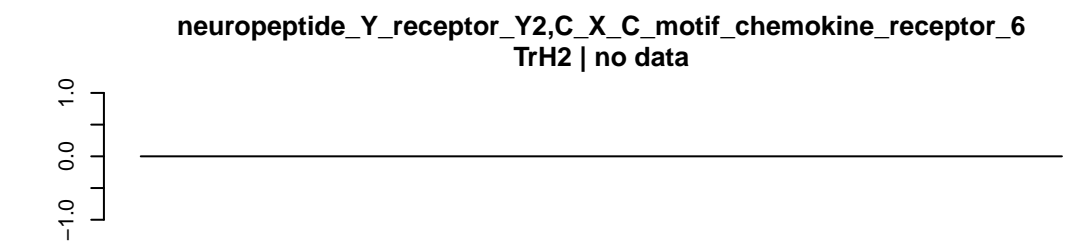
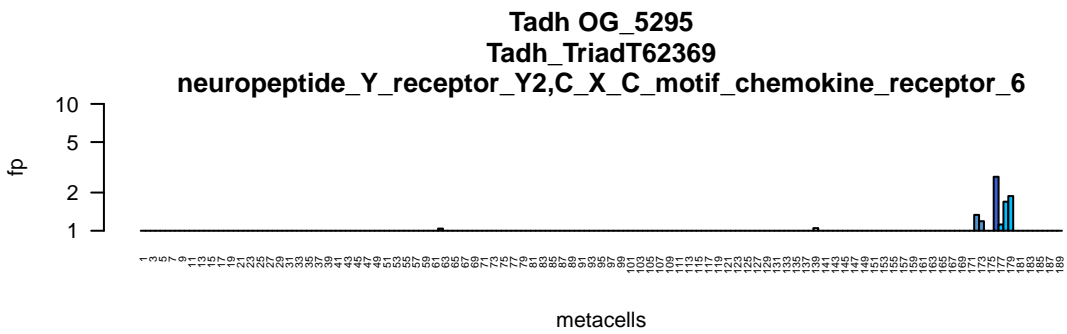


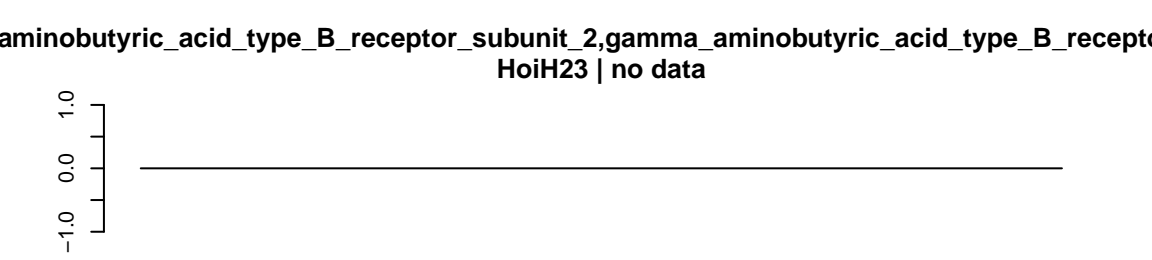
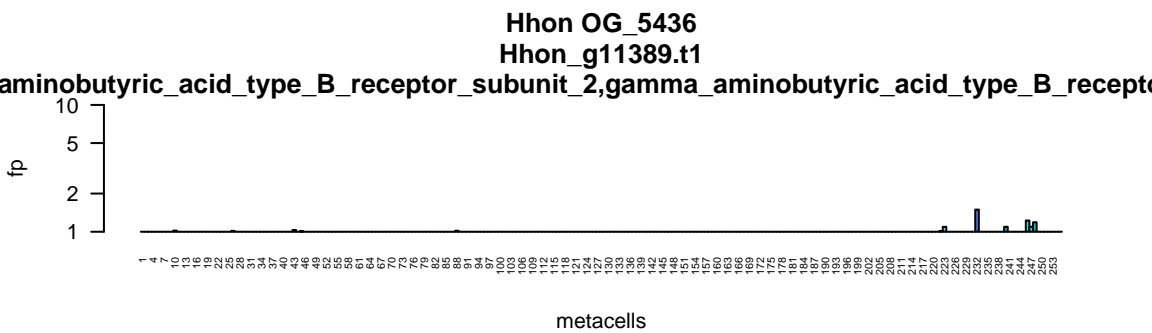
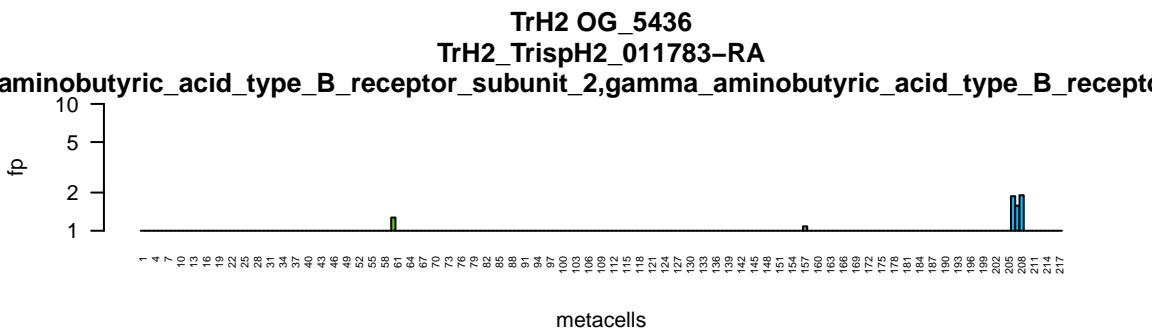
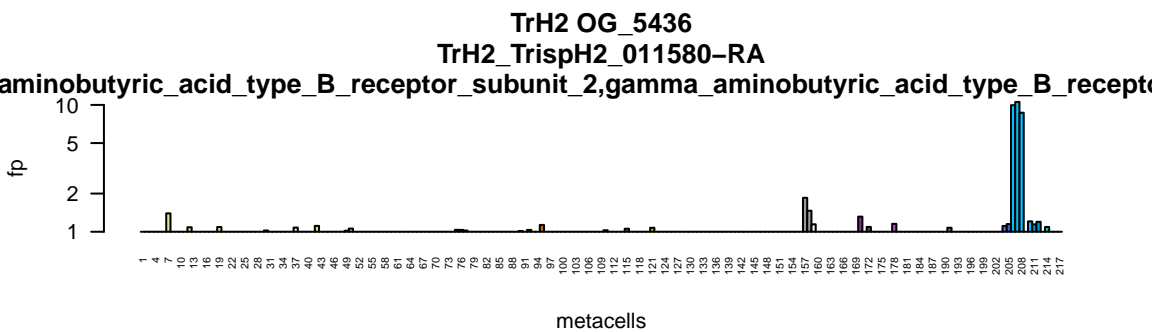
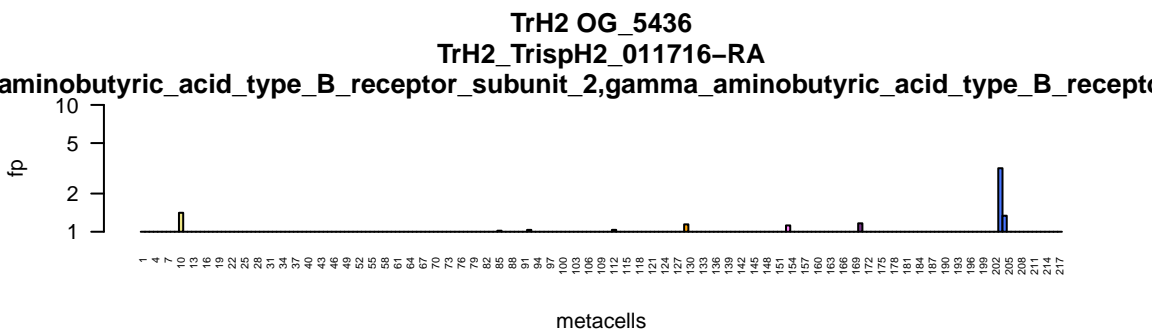
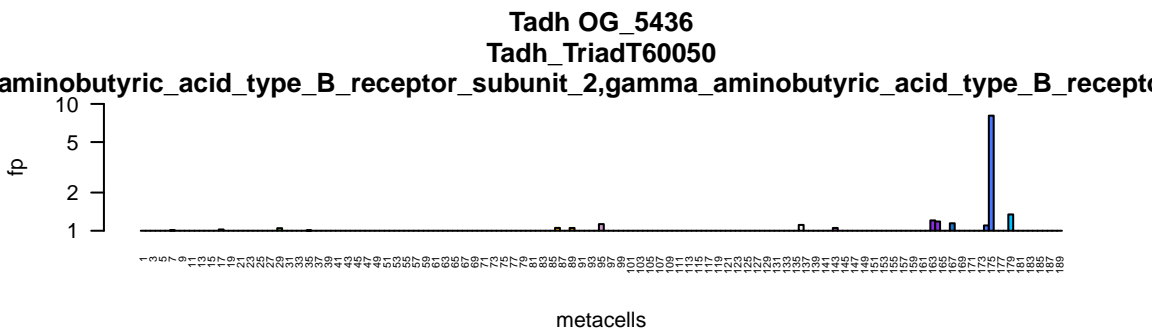


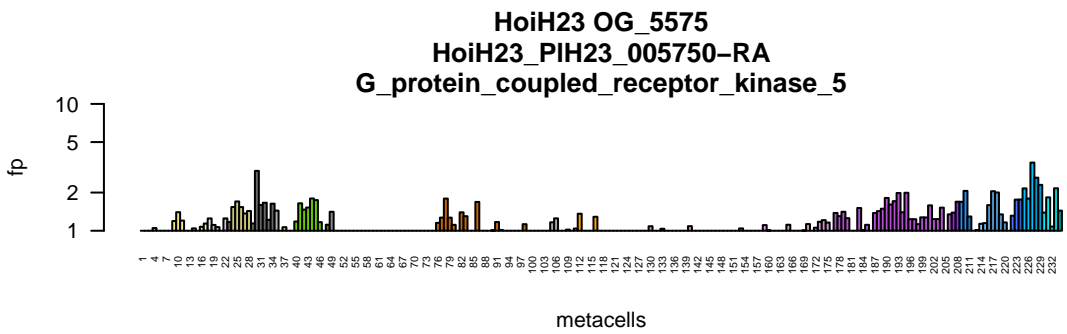
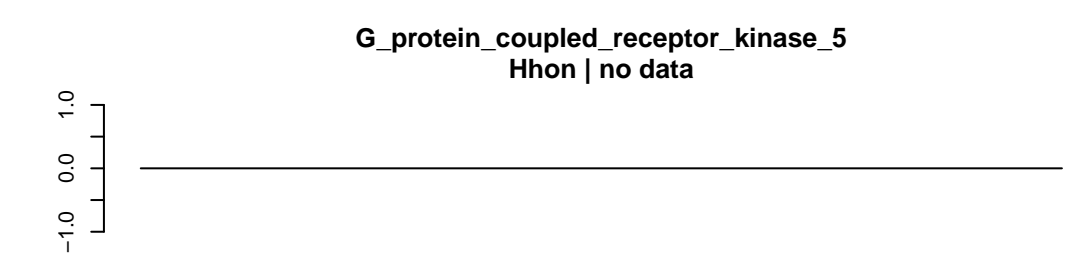
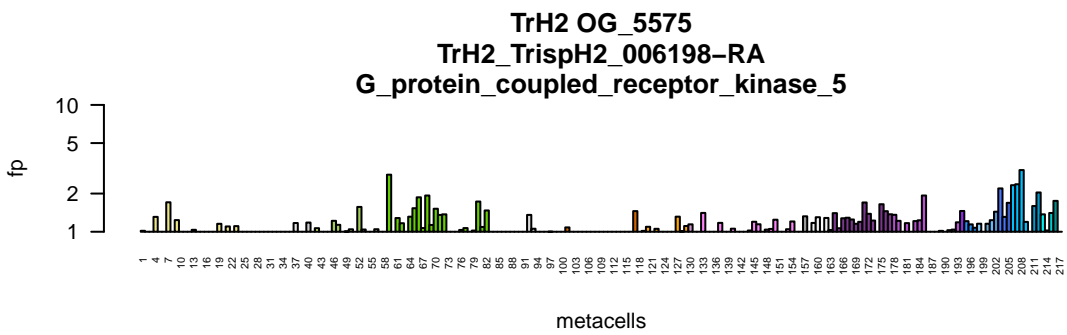
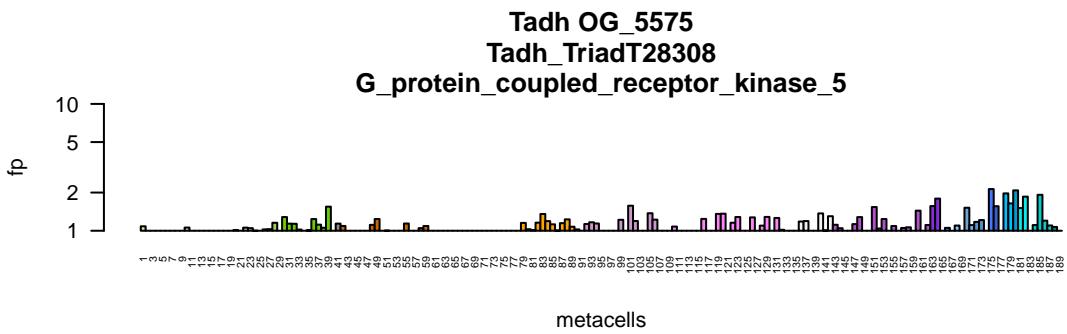




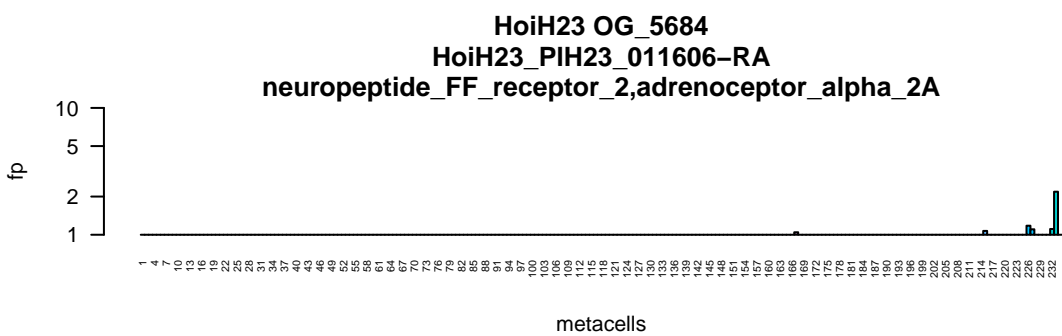
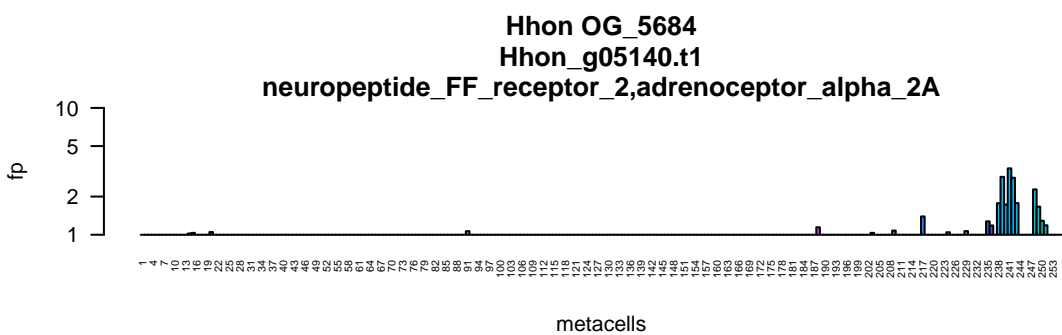
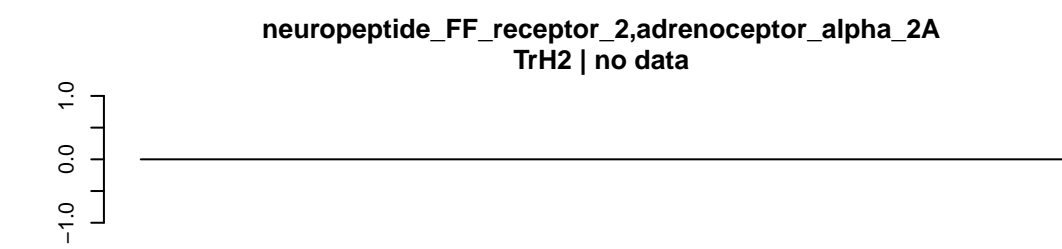
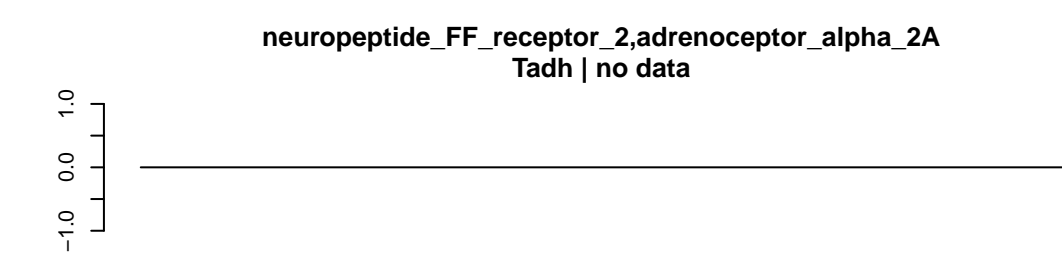


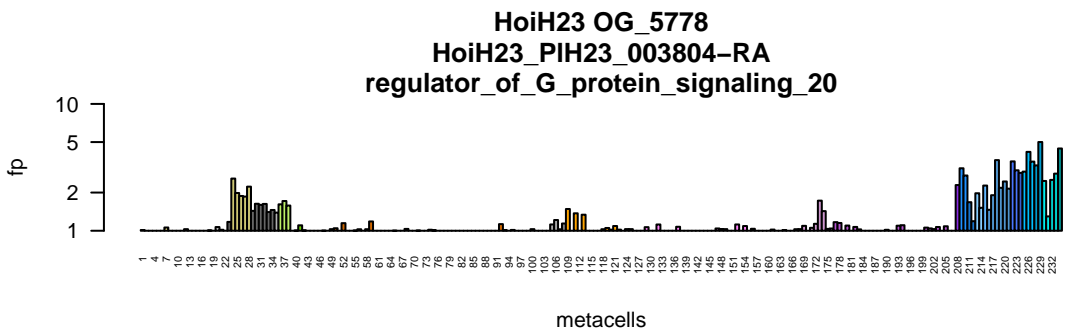
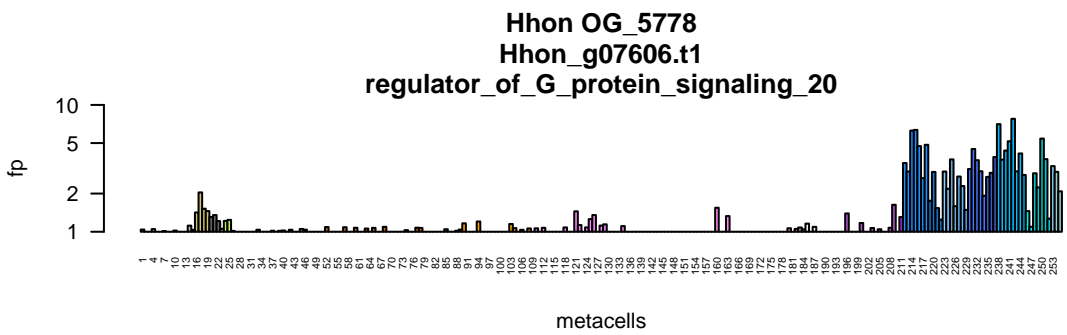
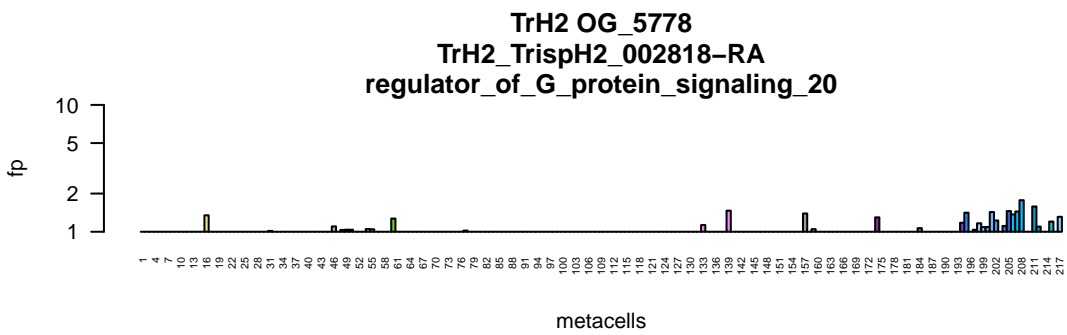
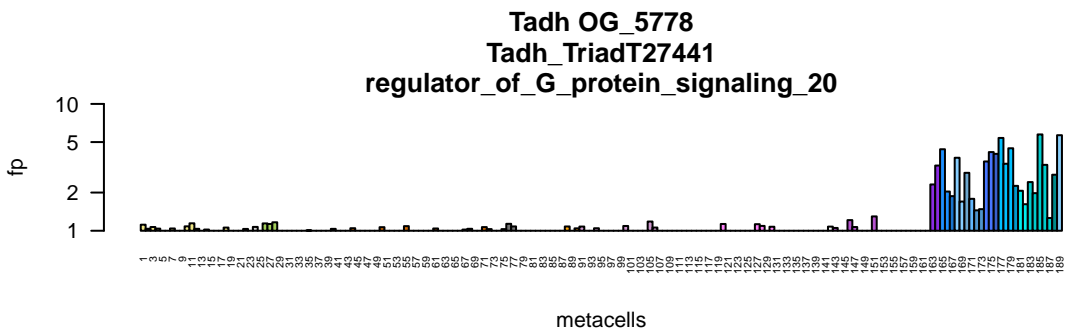


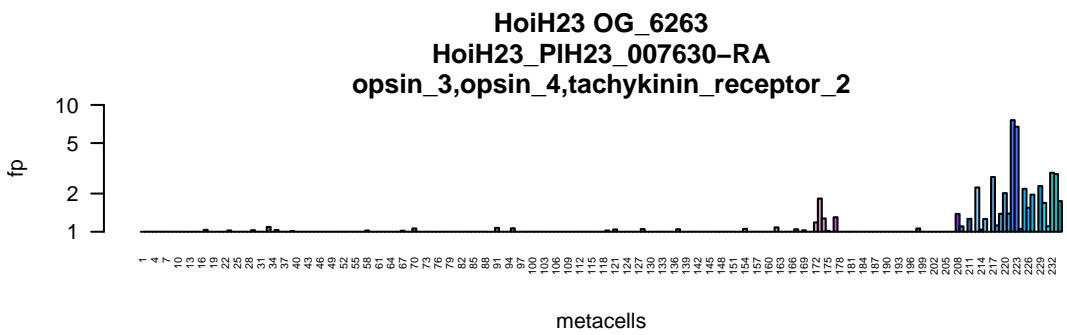
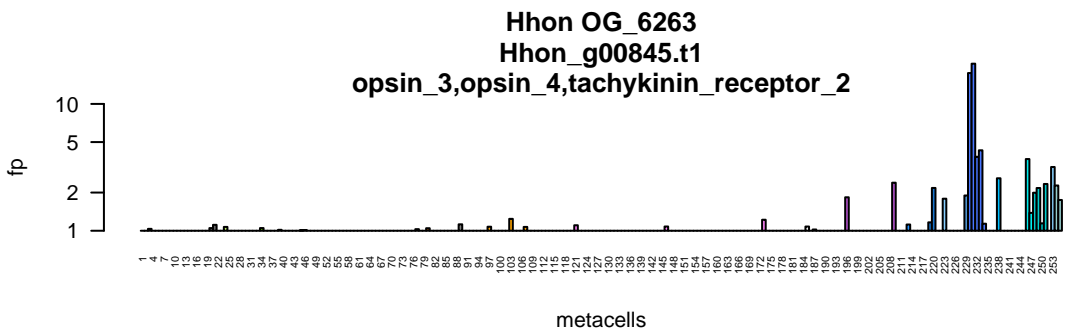
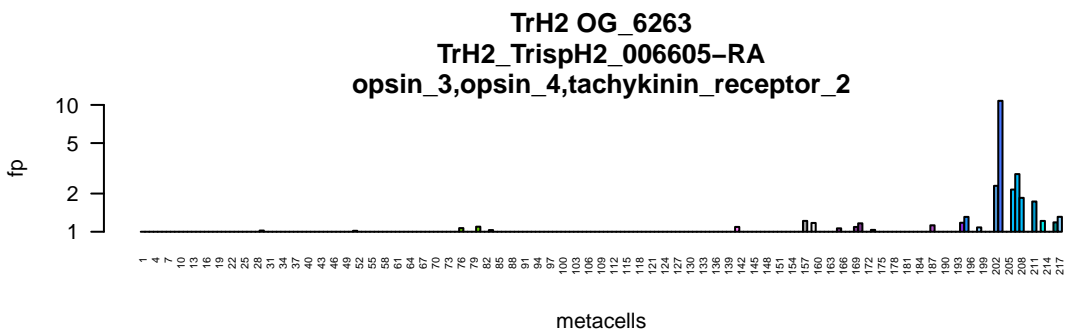
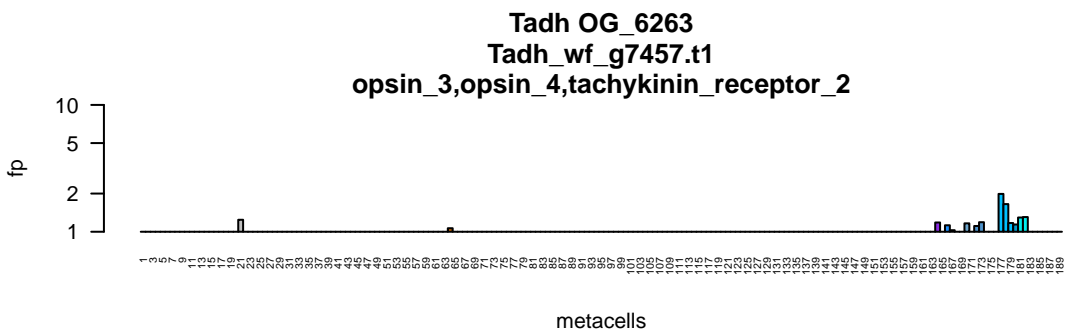
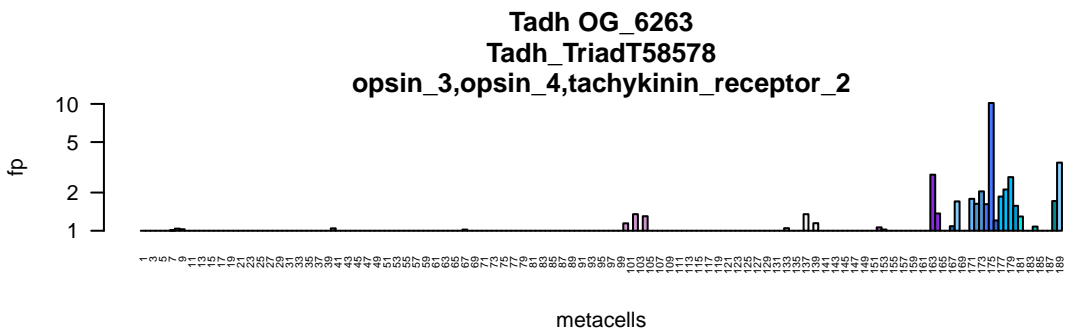


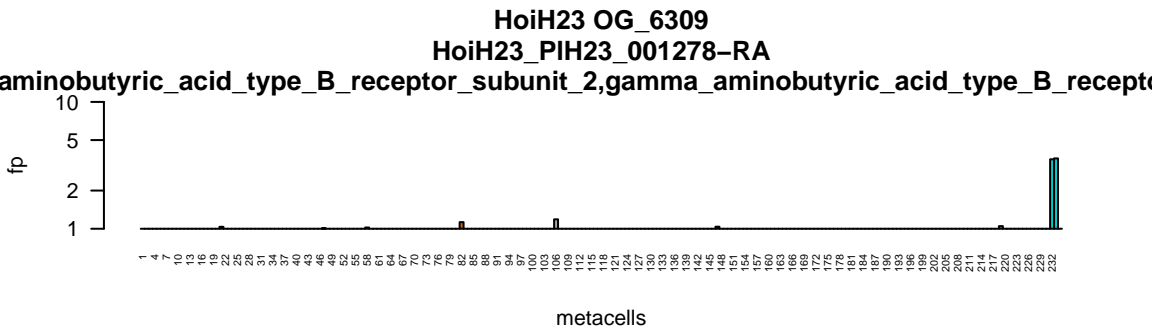
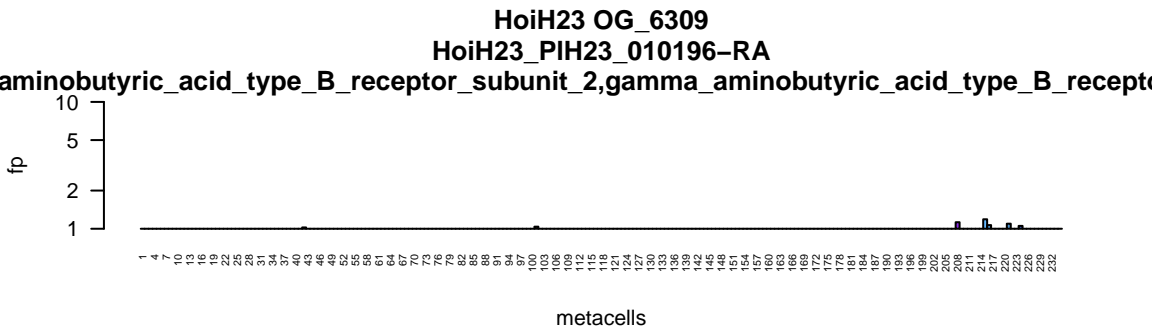
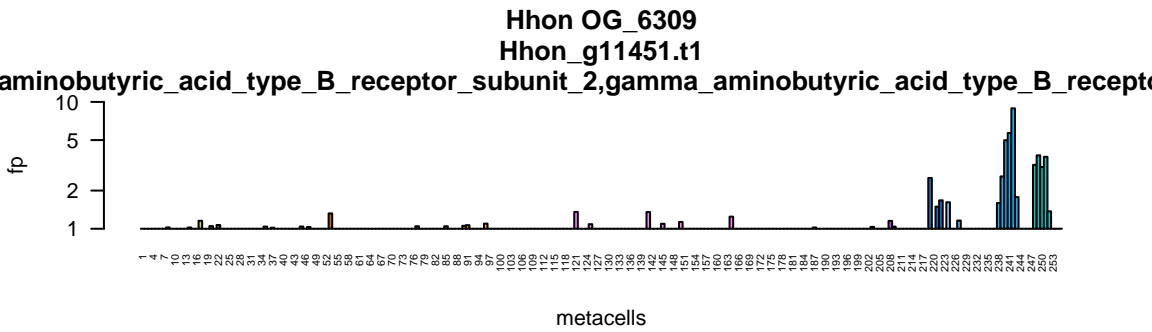
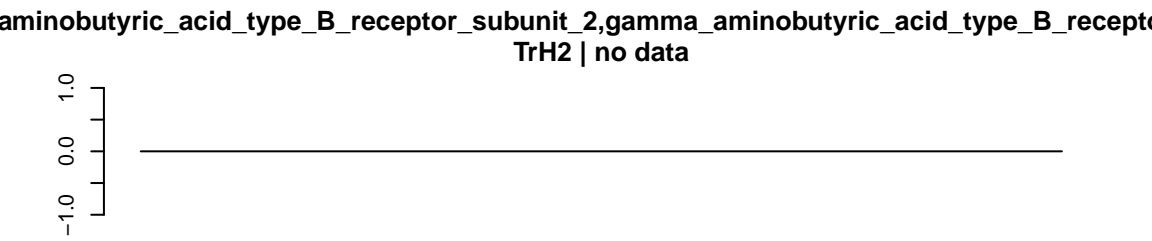
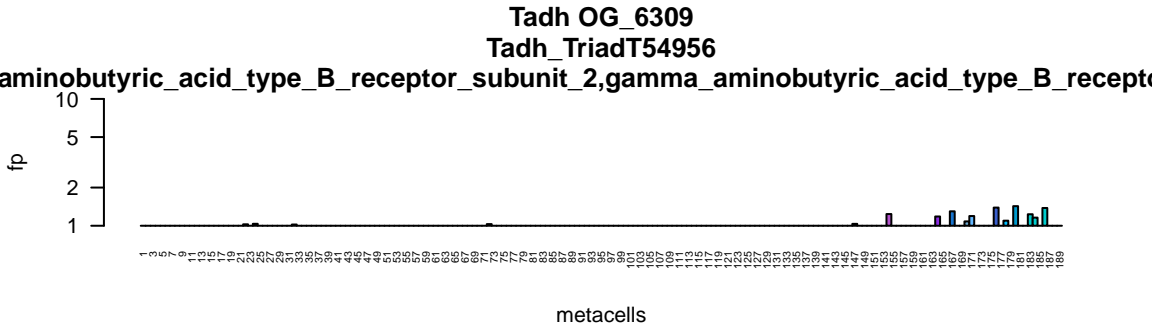
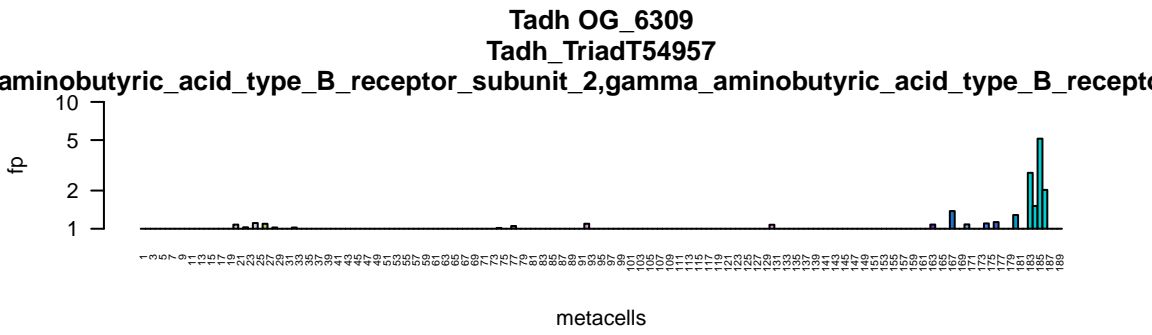
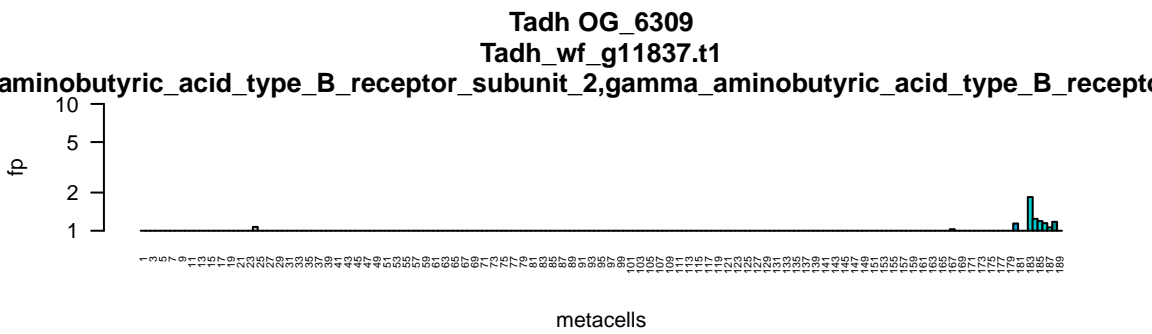


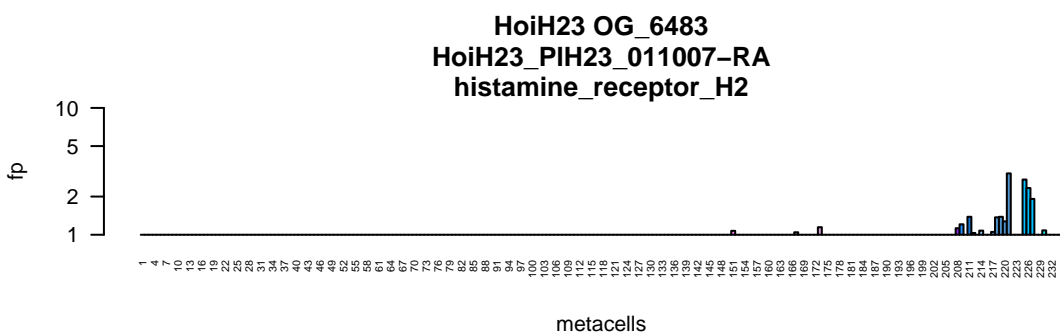
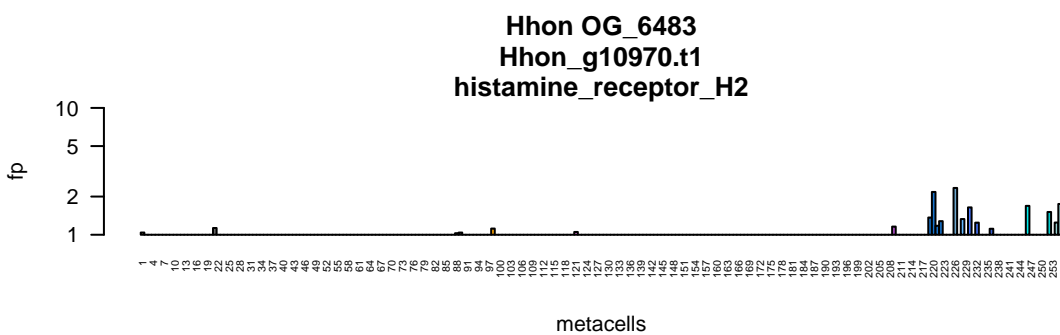
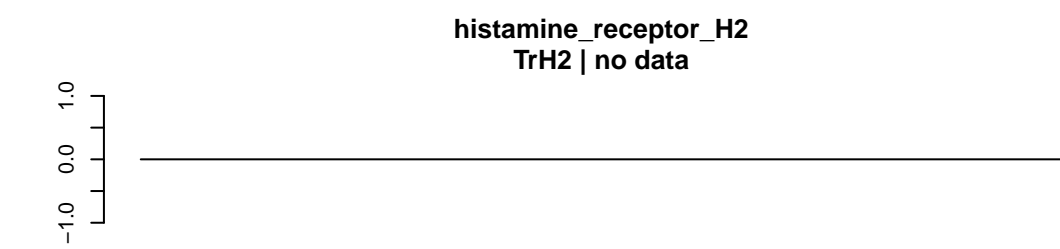
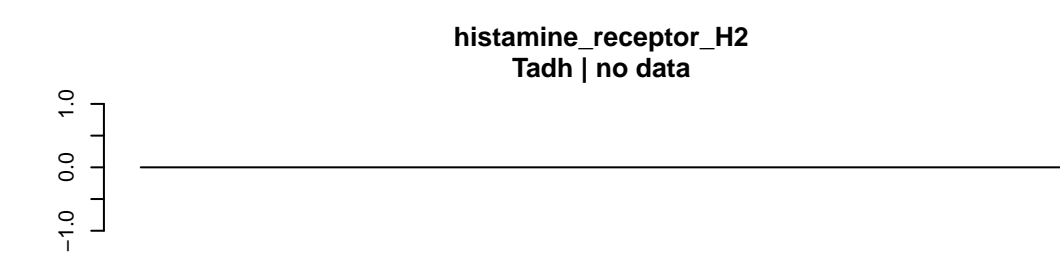












melanin\_concentrating\_hormone\_receptor\_1,C\_X\_C\_motif\_chemokine\_receptor\_1  
Tadh | no data

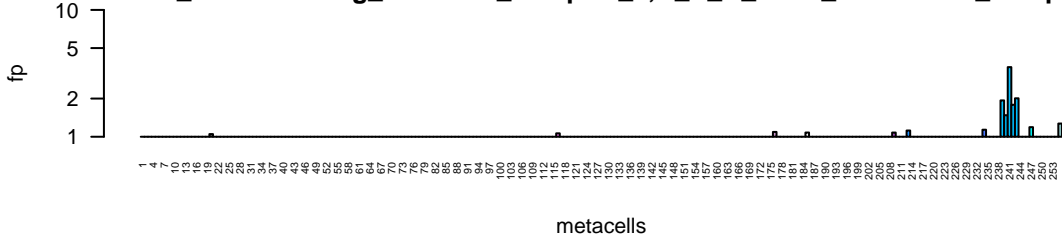


melanin\_concentrating\_hormone\_receptor\_1,C\_X\_C\_motif\_chemokine\_receptor\_1  
TrH2 | no data



Hhon OG\_6495  
Hhon\_g08607.t1

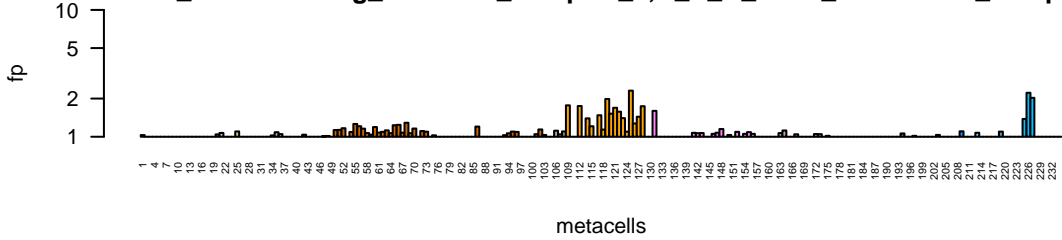
melanin\_concentrating\_hormone\_receptor\_1,C\_X\_C\_motif\_chemokine\_receptor\_1



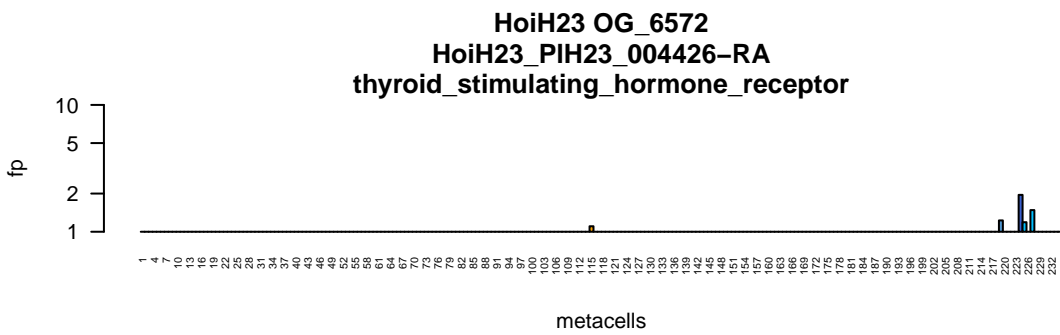
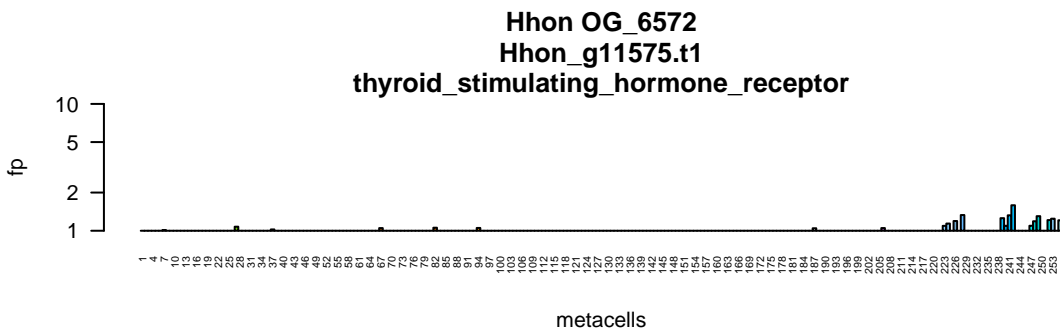
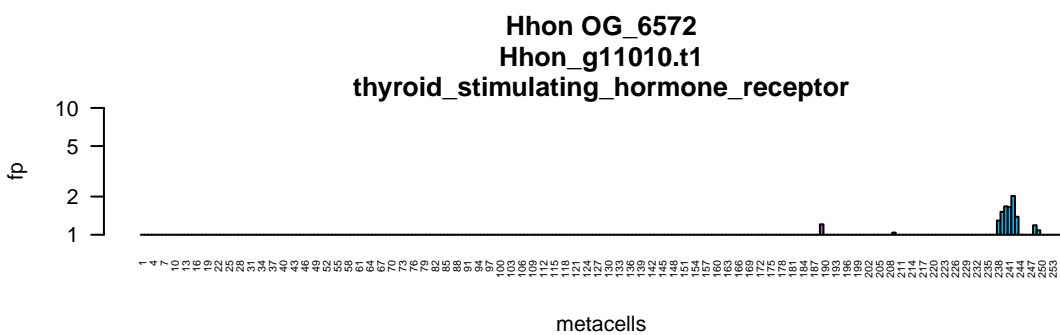
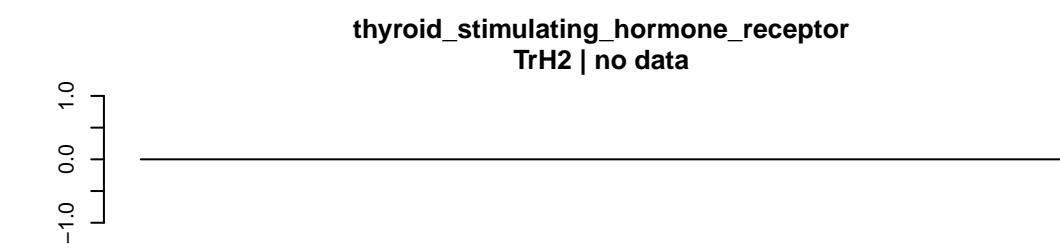
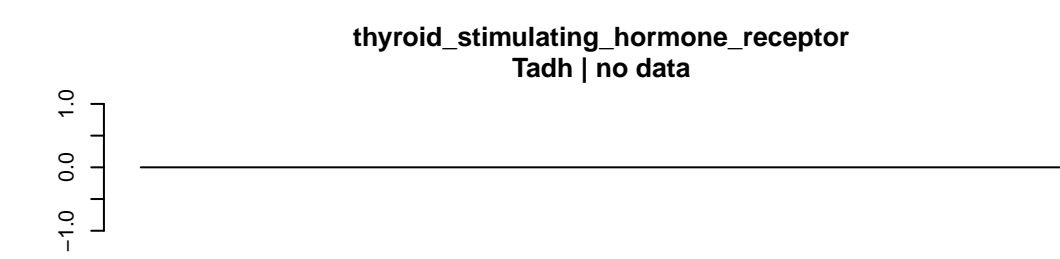
metacells

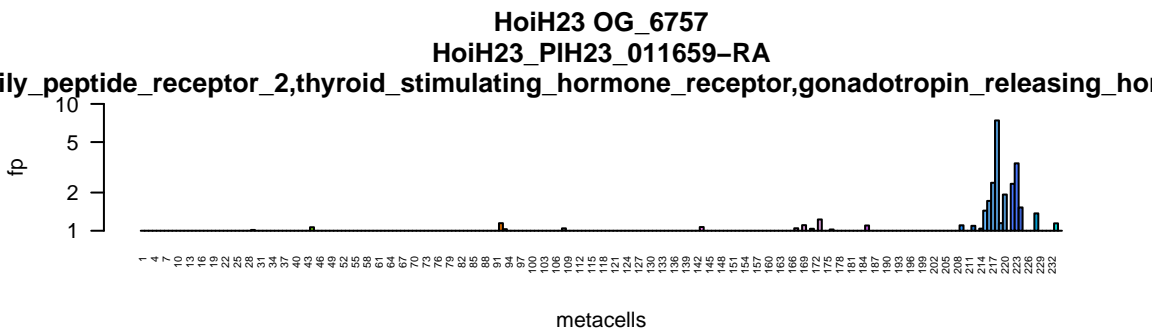
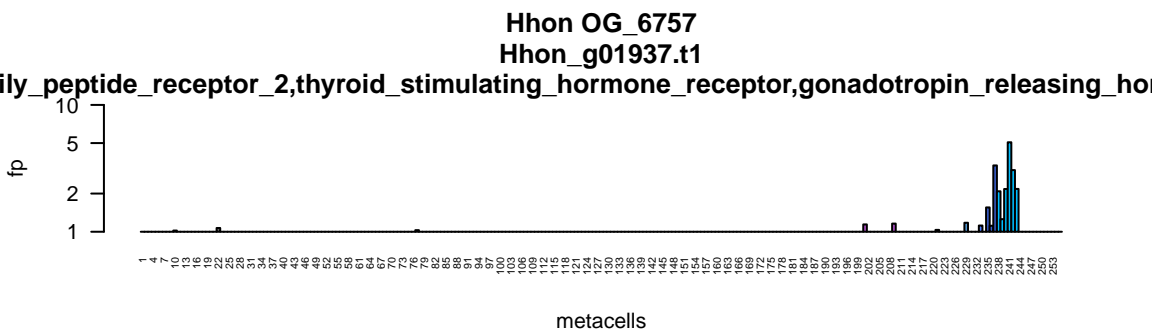
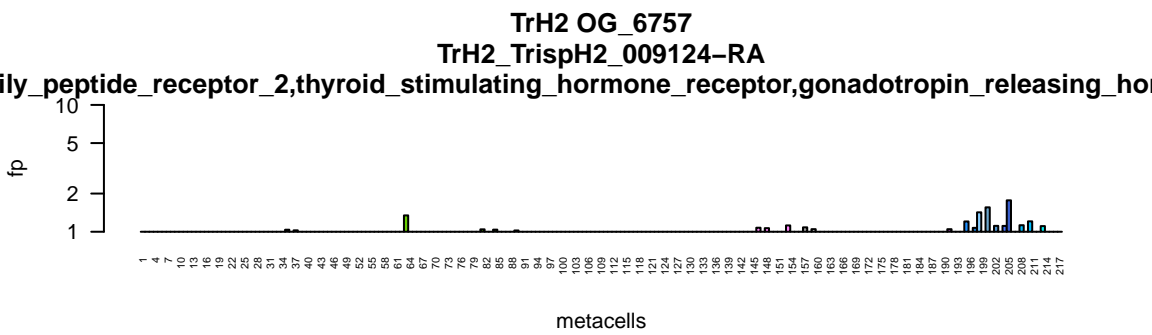
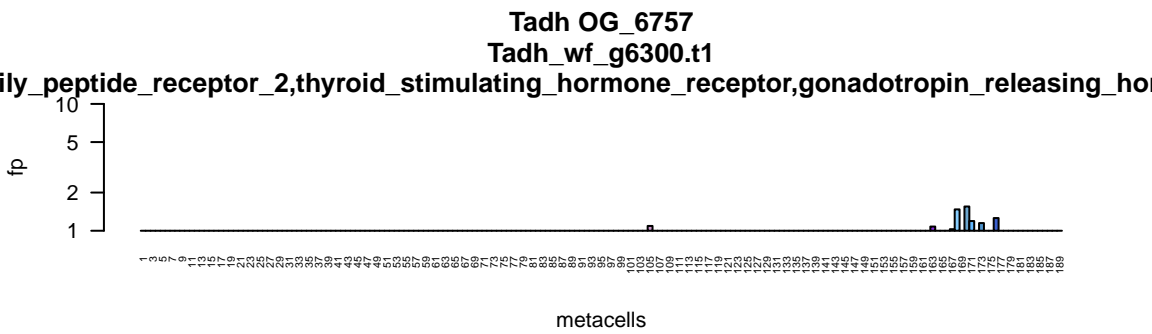
HoiH23 OG\_6495  
HoiH23\_PIH23\_004725-RA

melanin\_concentrating\_hormone\_receptor\_1,C\_X\_C\_motif\_chemokine\_receptor\_1



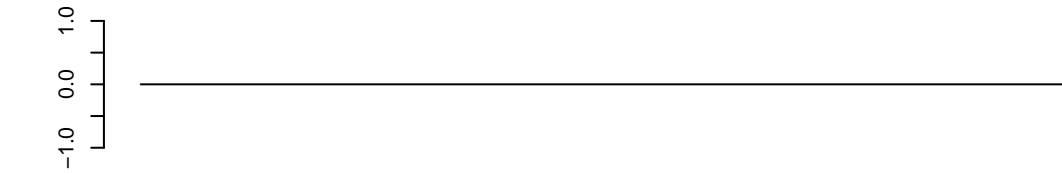
metacells



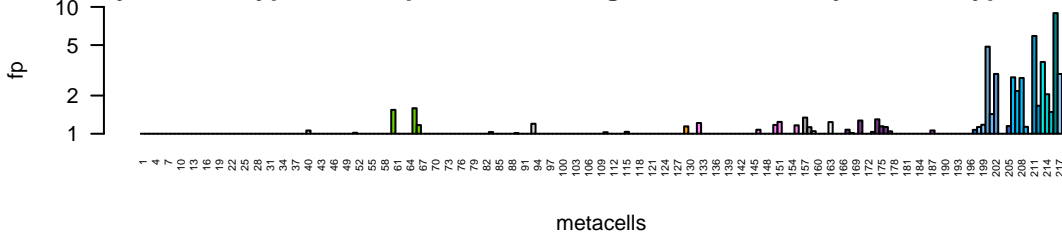




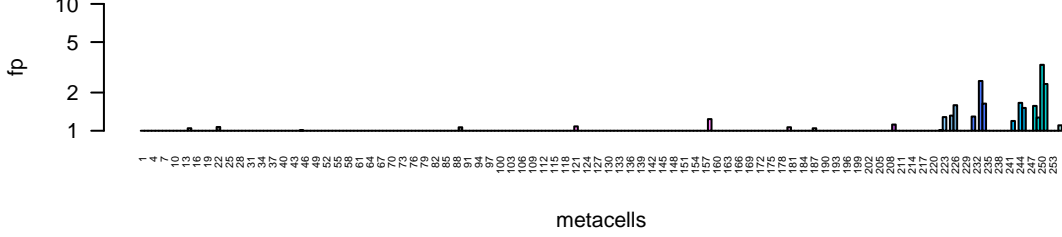
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recept  
Tadh | no data



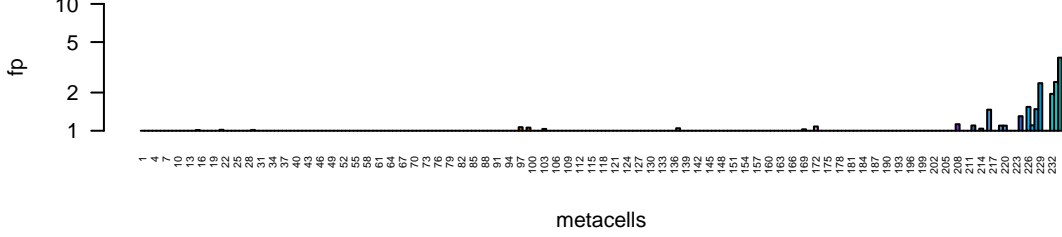
TrH2 OG\_7186  
TrH2\_TrispH2\_010138-RA  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recept



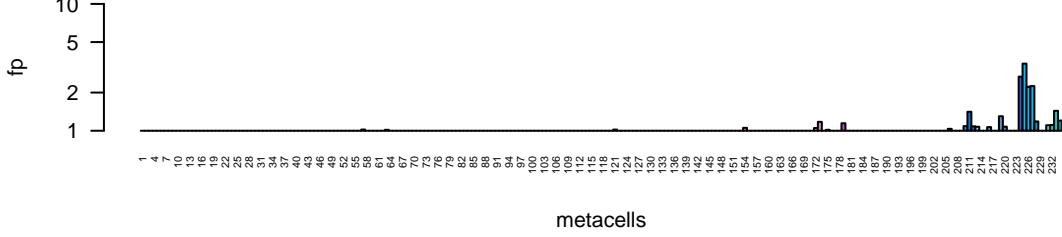
Hhon OG\_7186  
Hhon\_g01977.t1  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recept



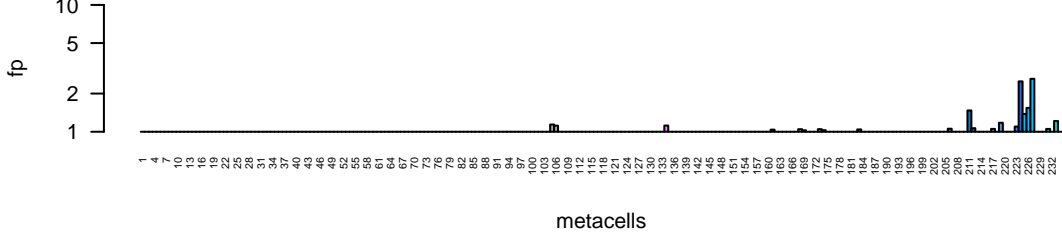
HoiH23 OG\_7186  
HoiH23\_PIH23\_011144-RA  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recept



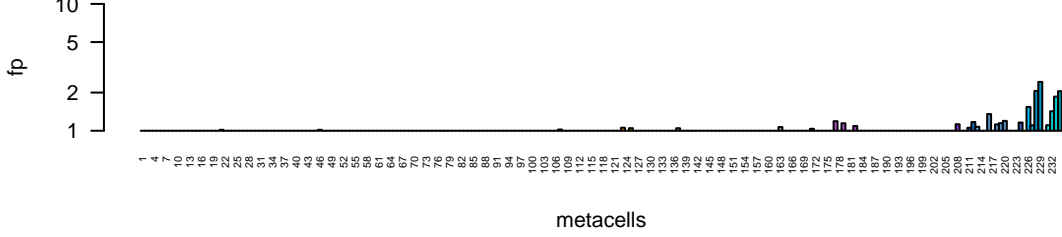
HoiH23 OG\_7186  
HoiH23\_PIH23\_011313-RA  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recept

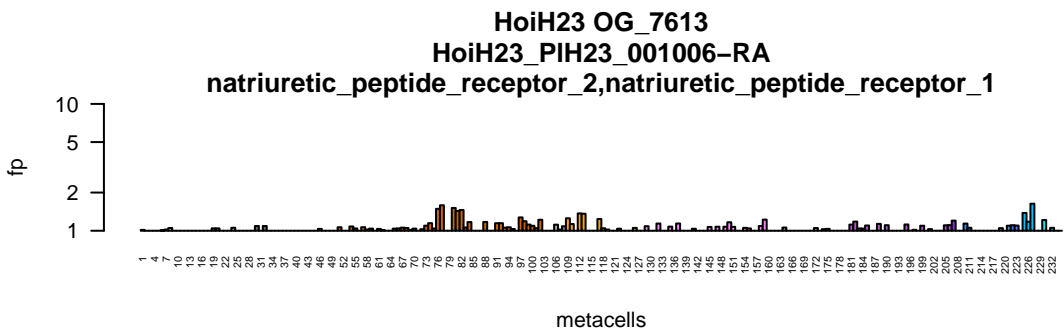
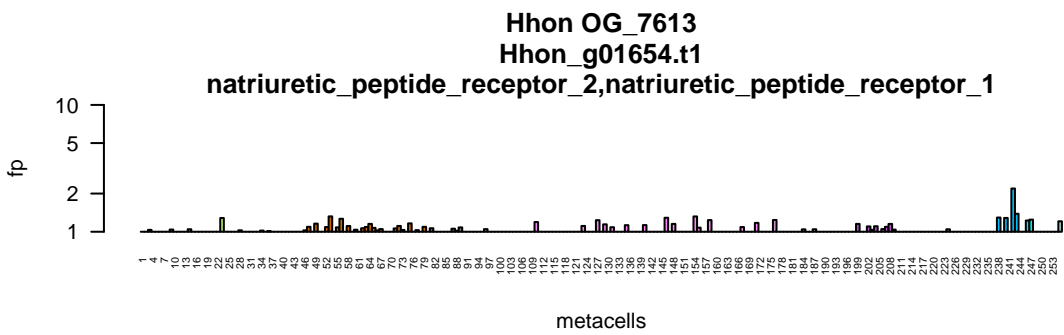
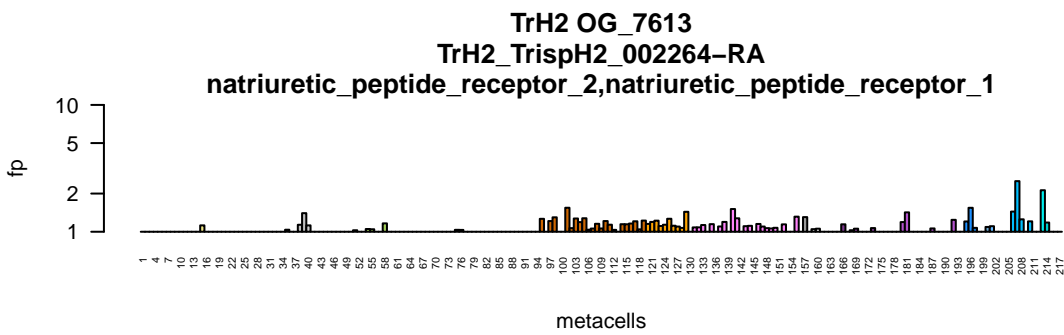
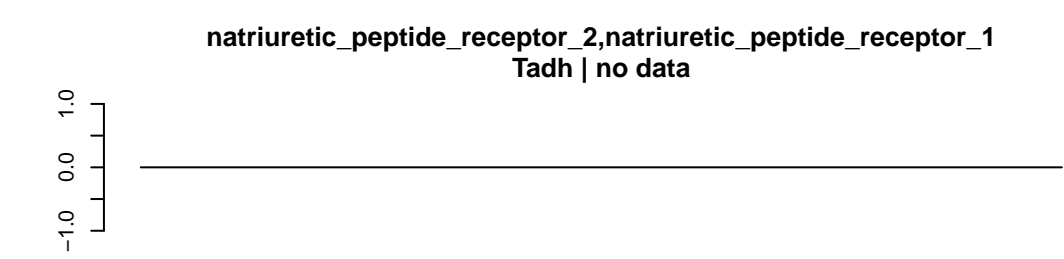


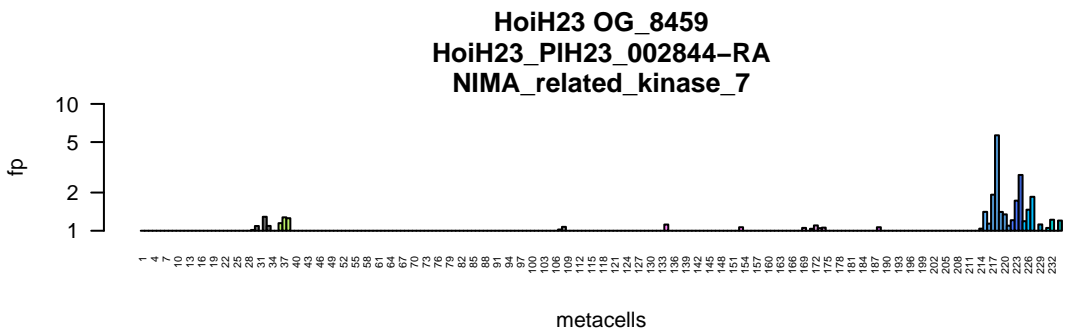
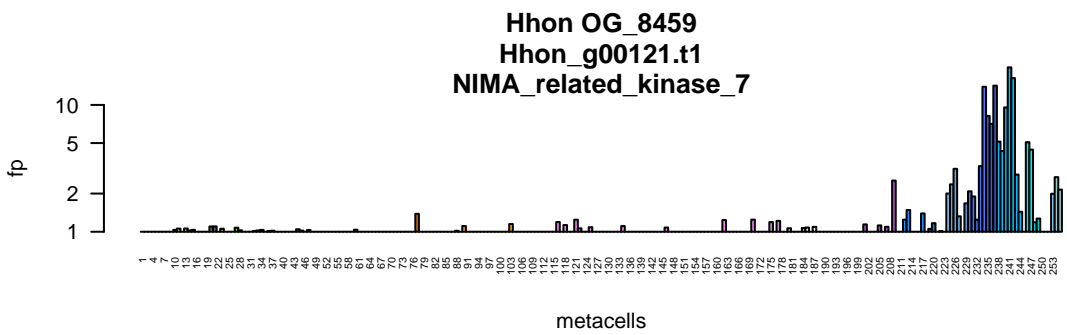
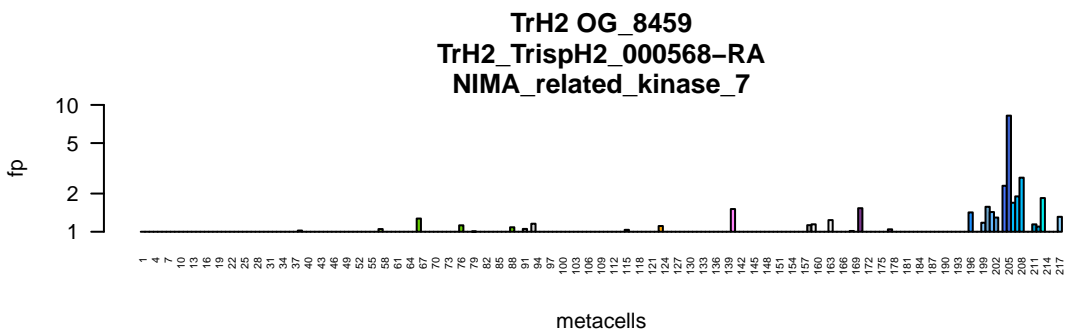
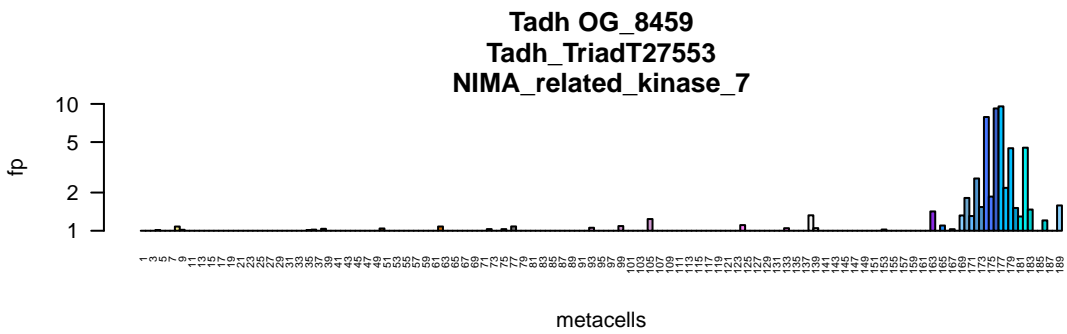
HoiH23 OG\_7186  
HoiH23\_PIH23\_004027-RA  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recept

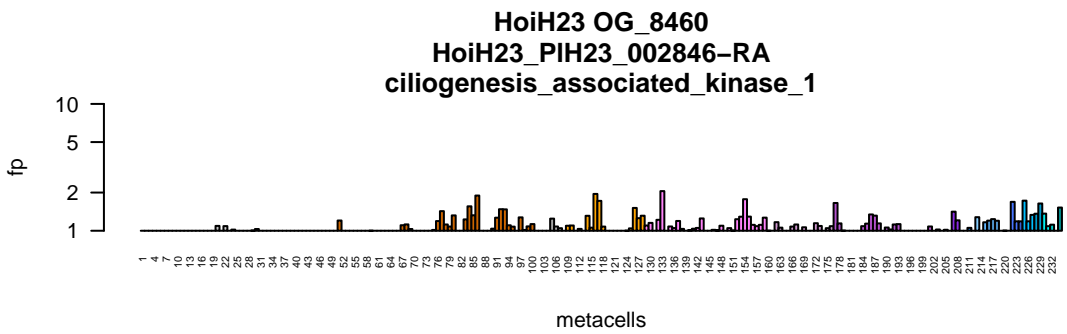
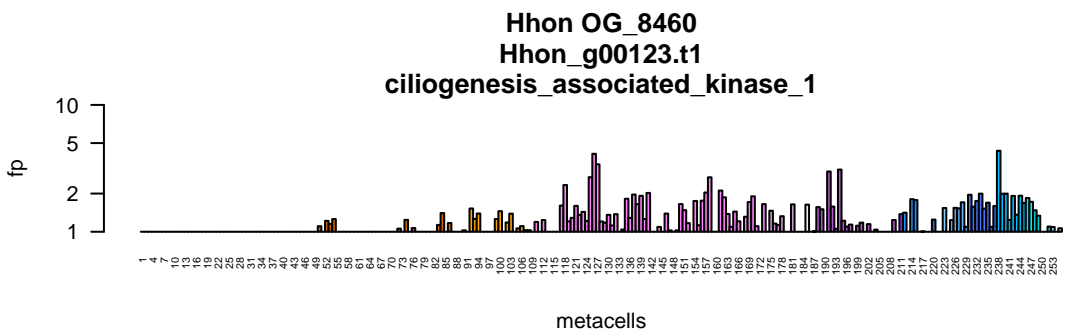
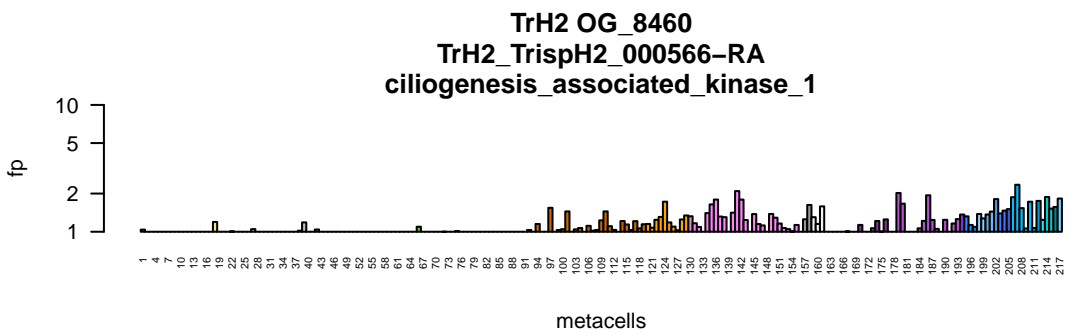
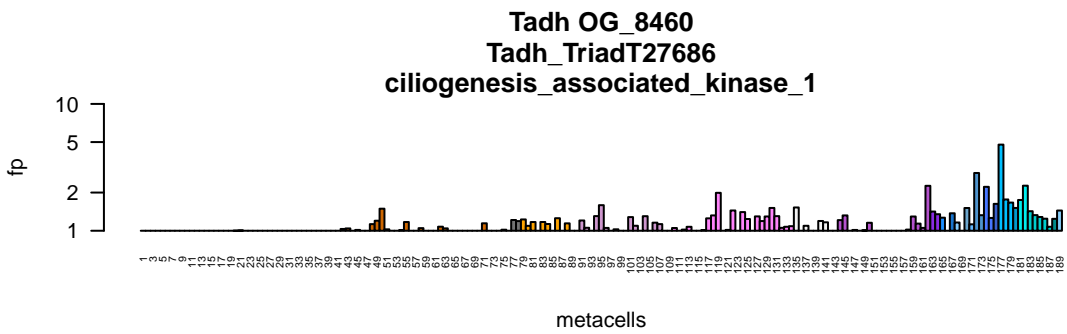


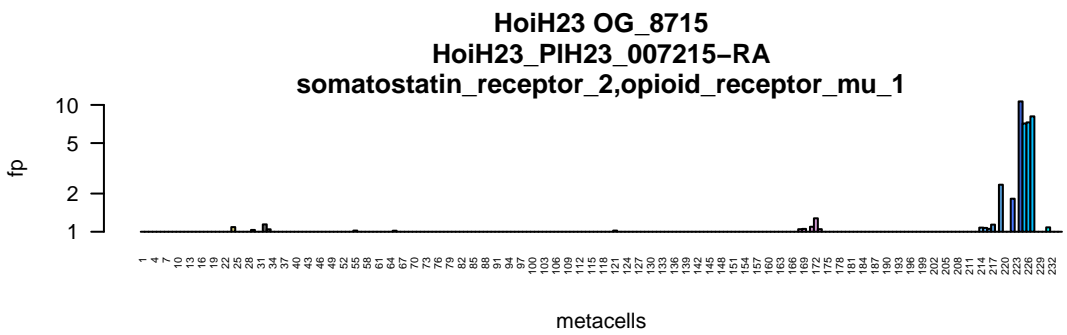
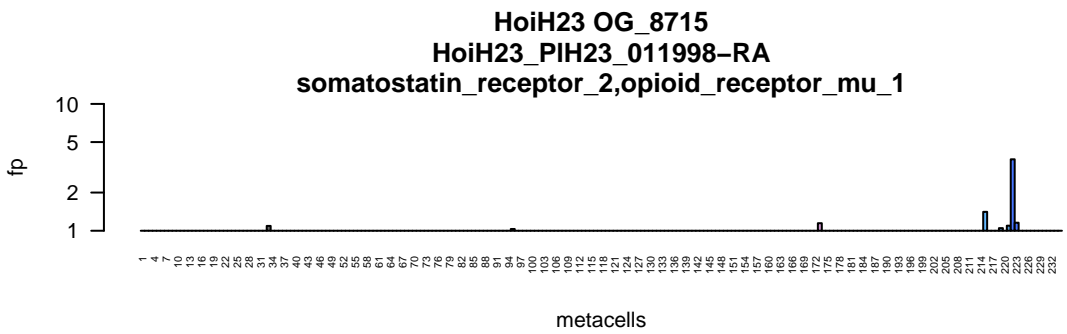
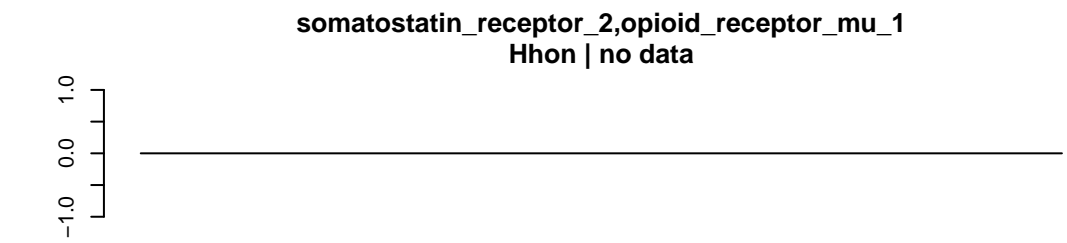
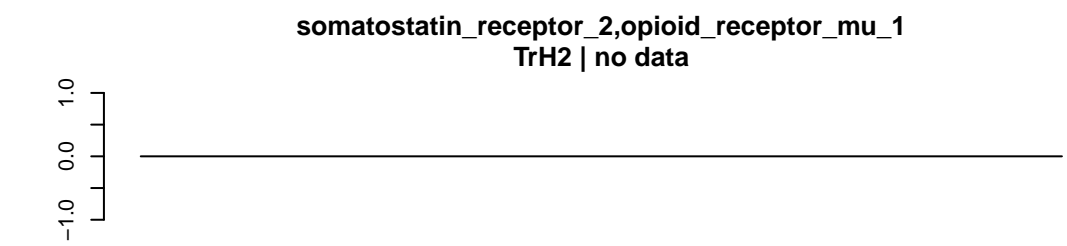
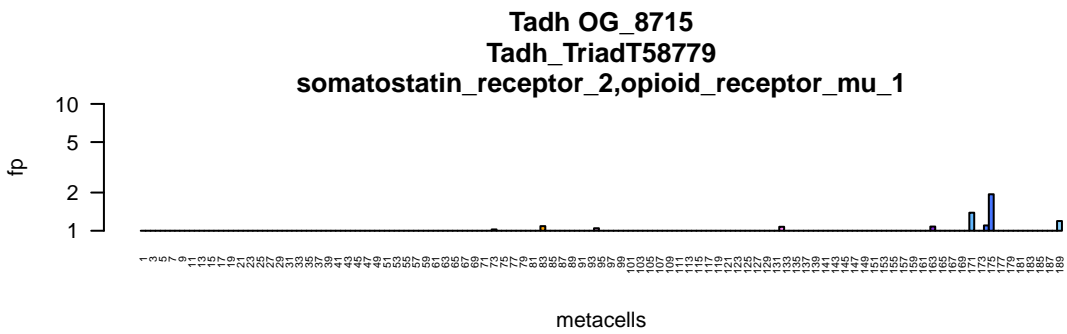
HoiH23 OG\_7186  
HoiH23\_PIH23\_011312-RA  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recept

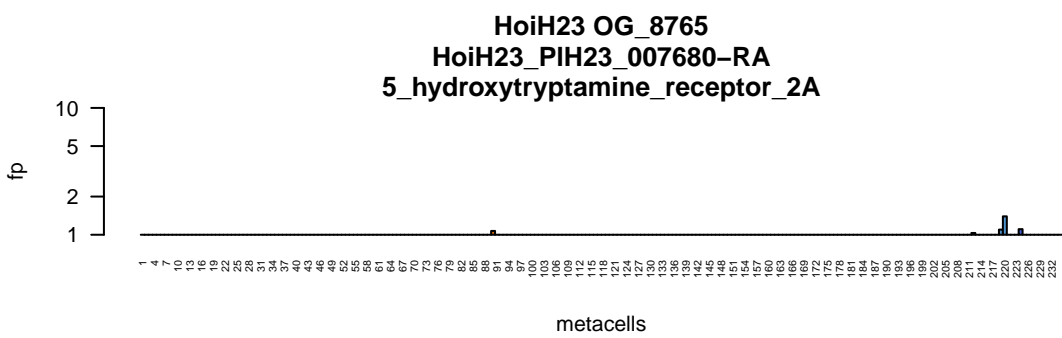
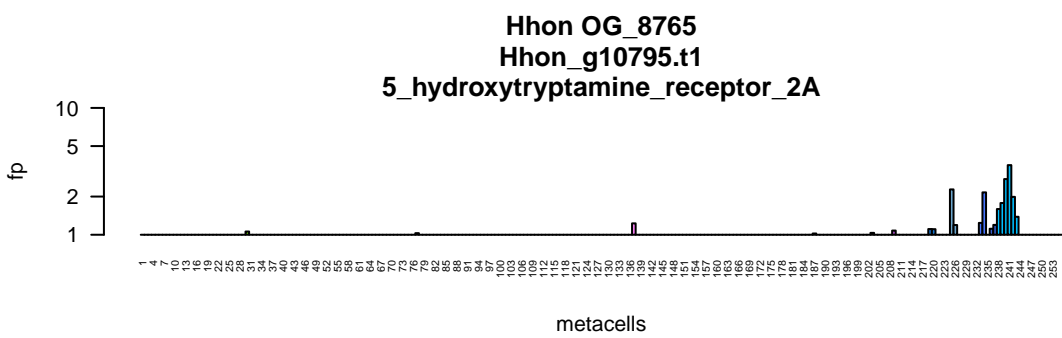
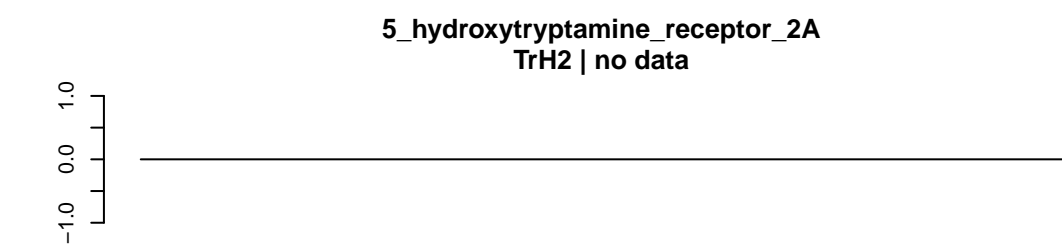
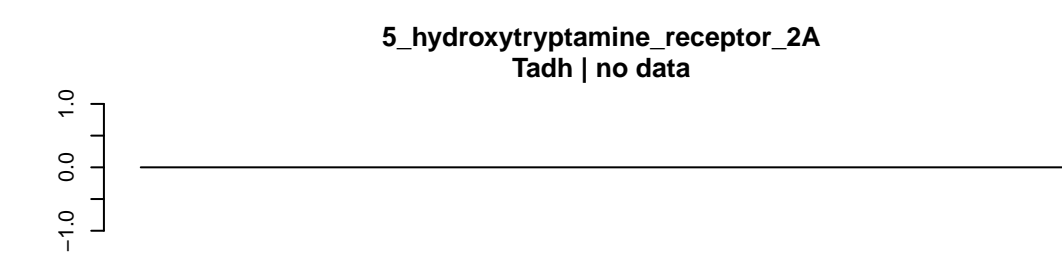


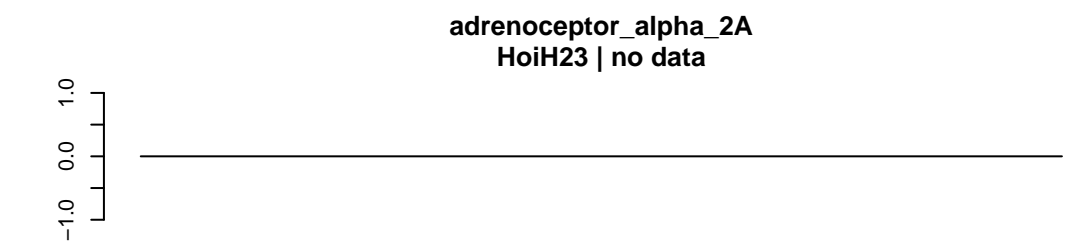
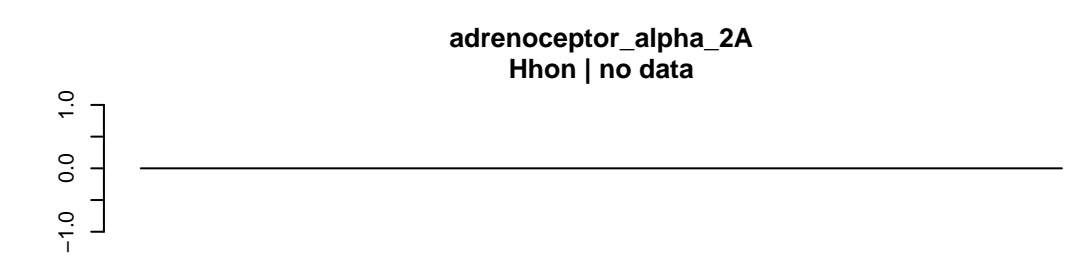
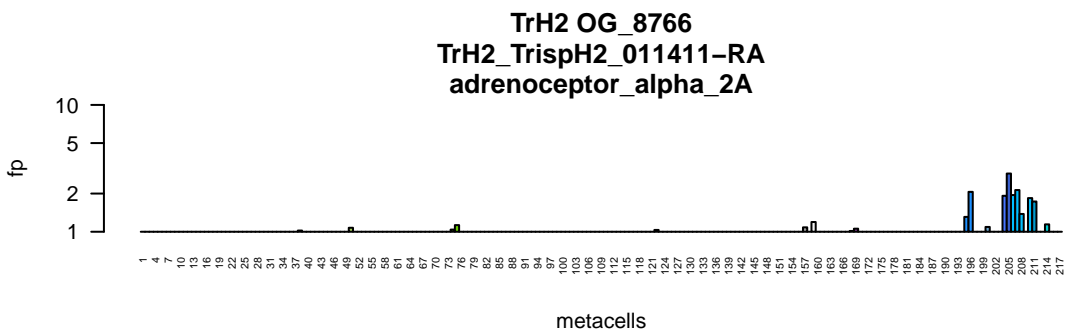
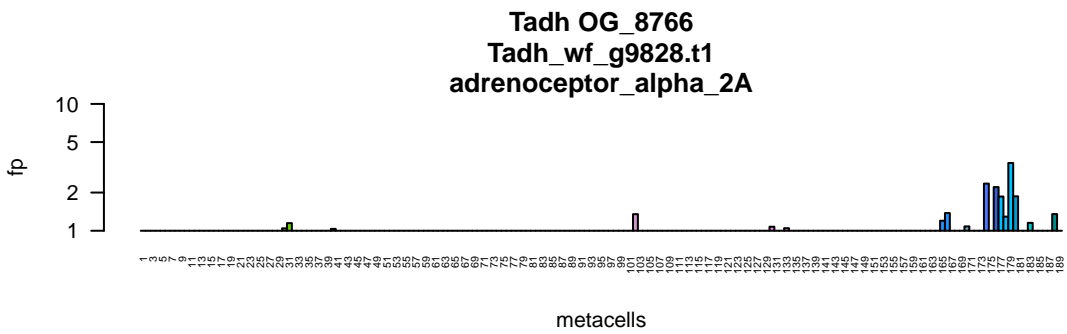


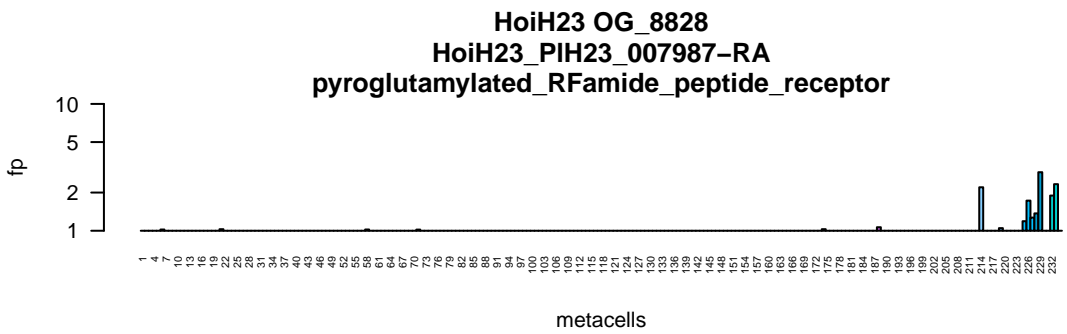
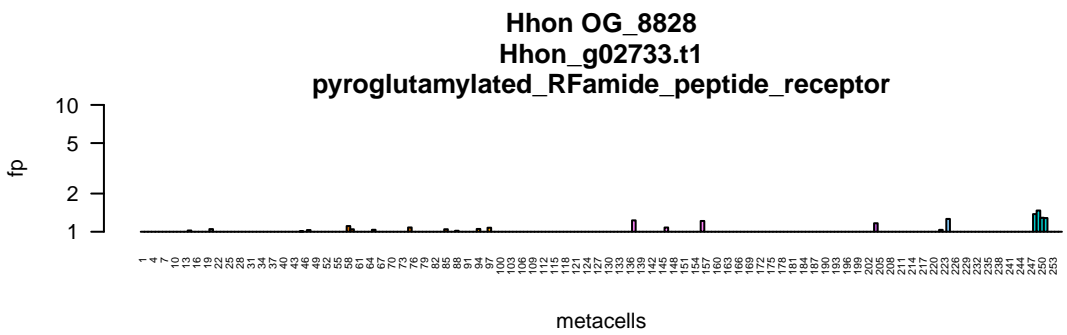
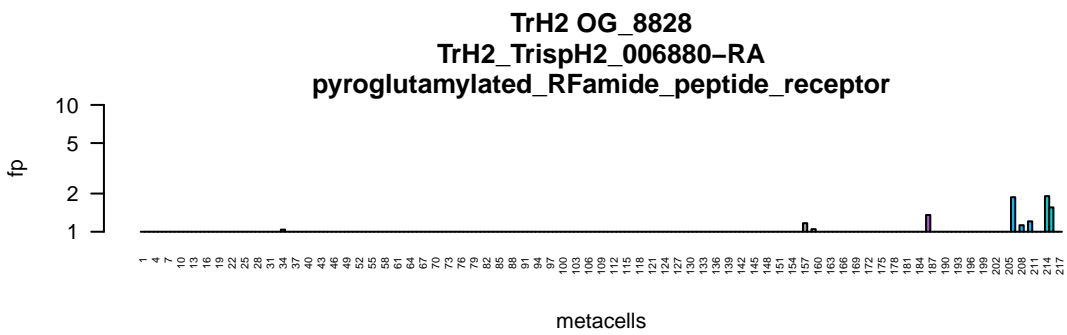
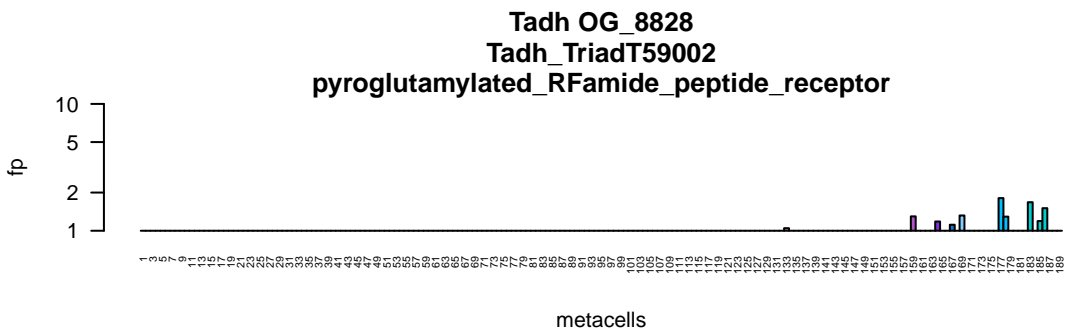




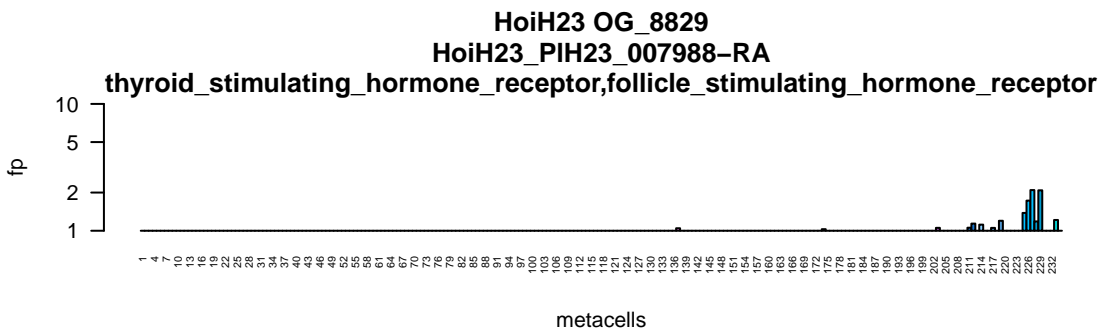
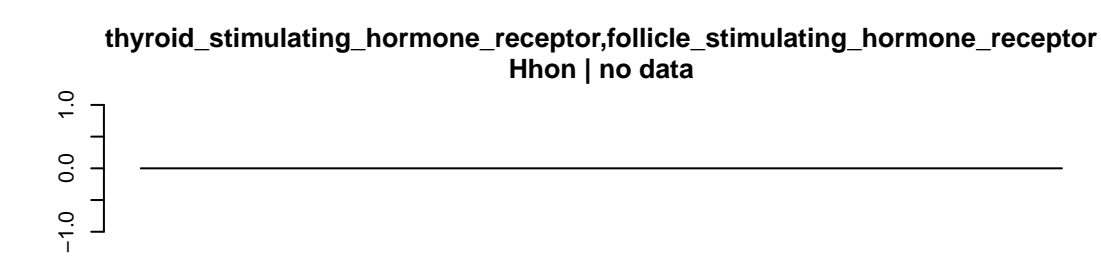
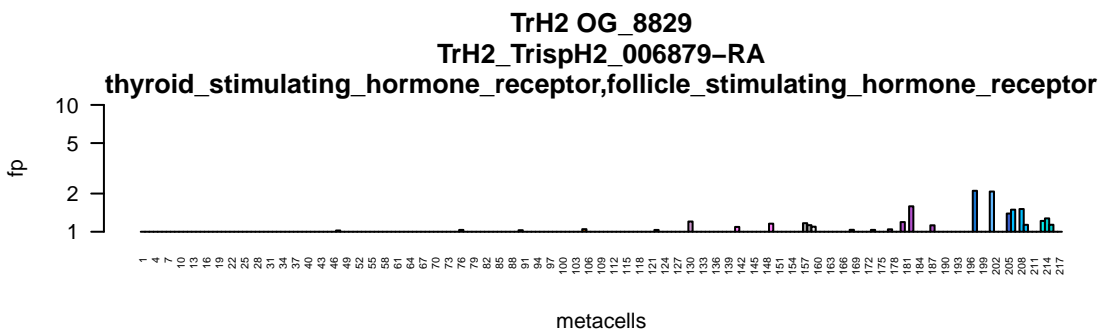
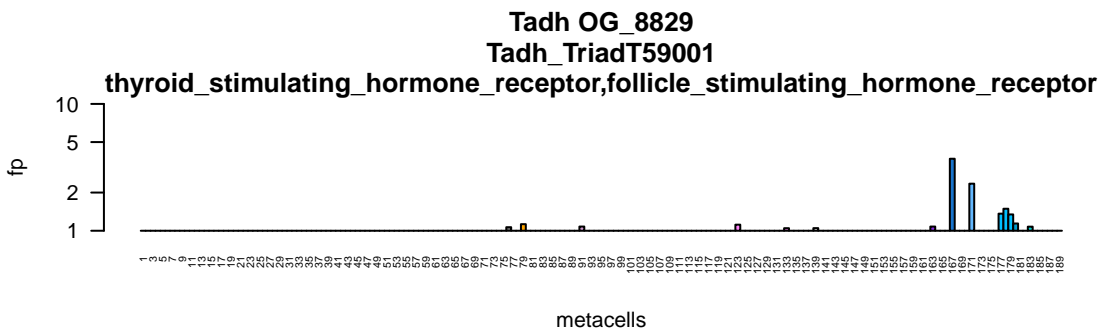


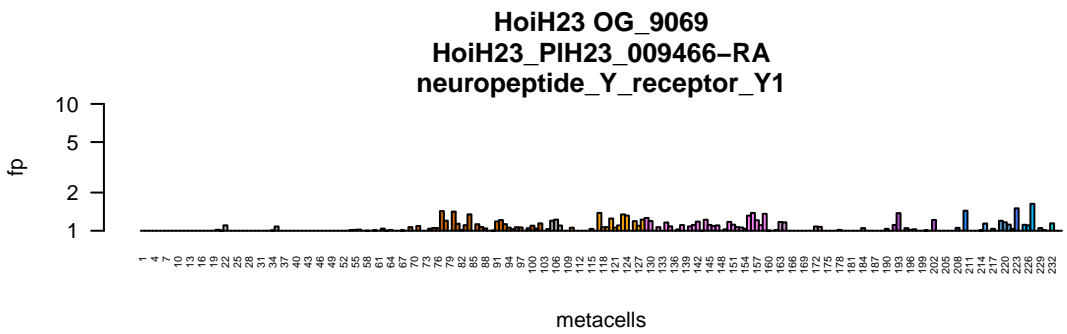
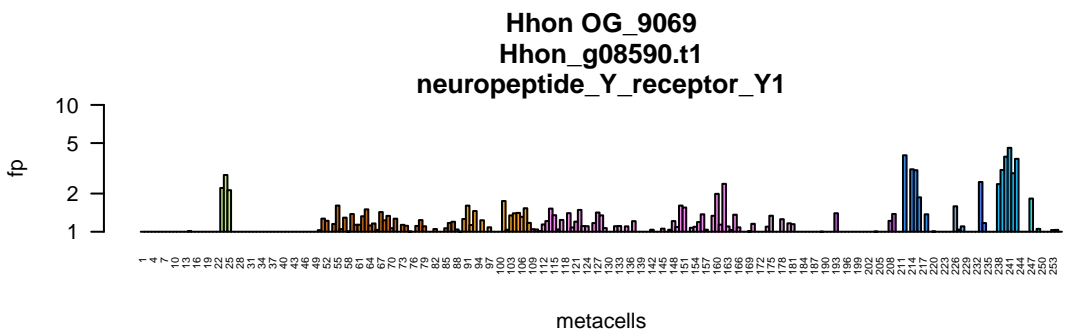
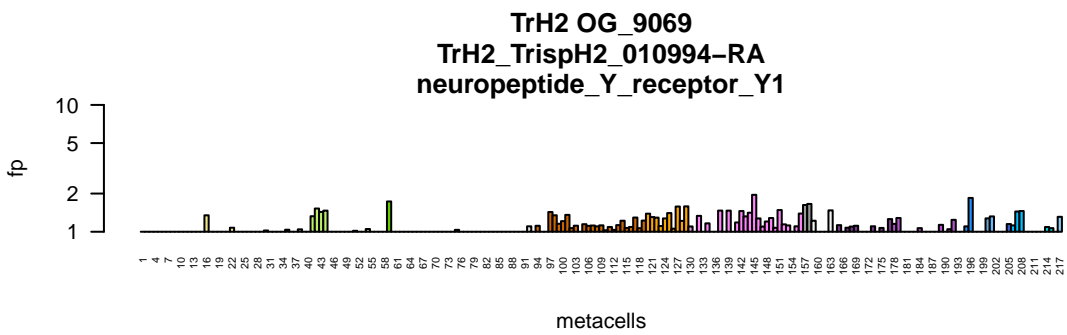
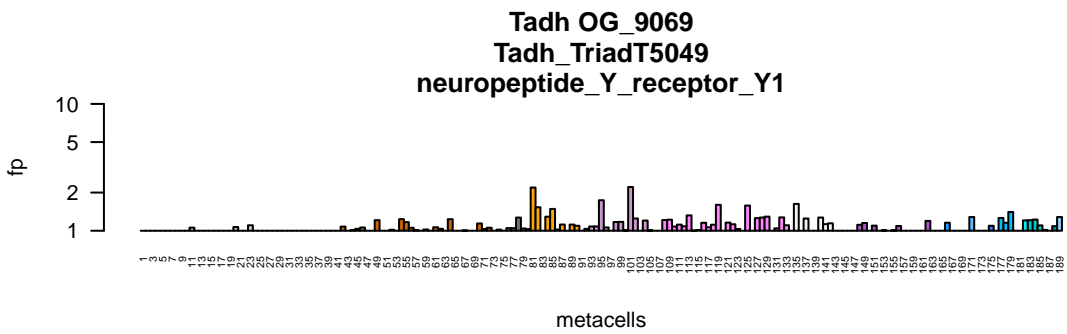


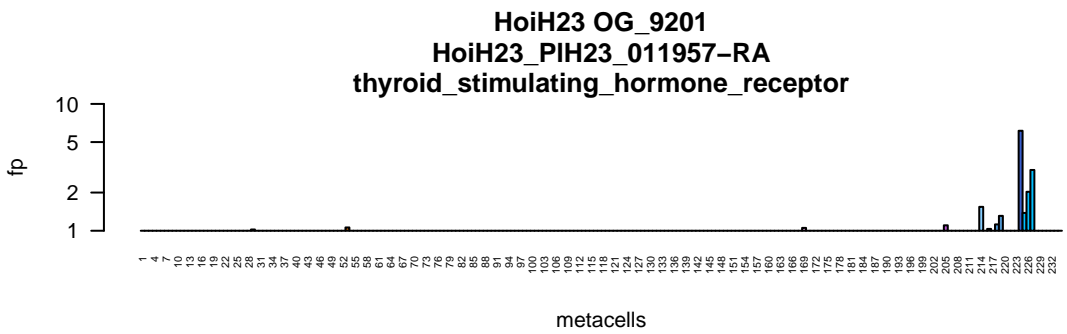
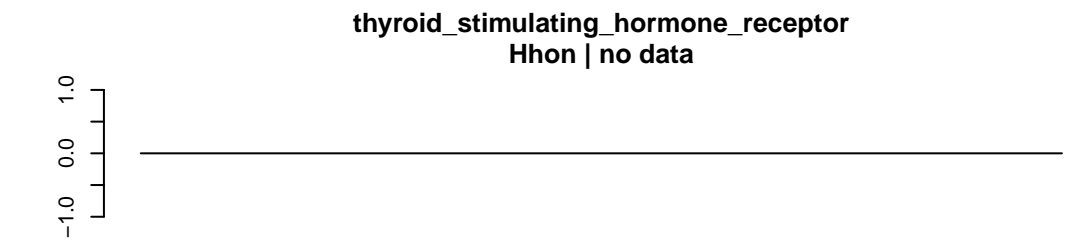
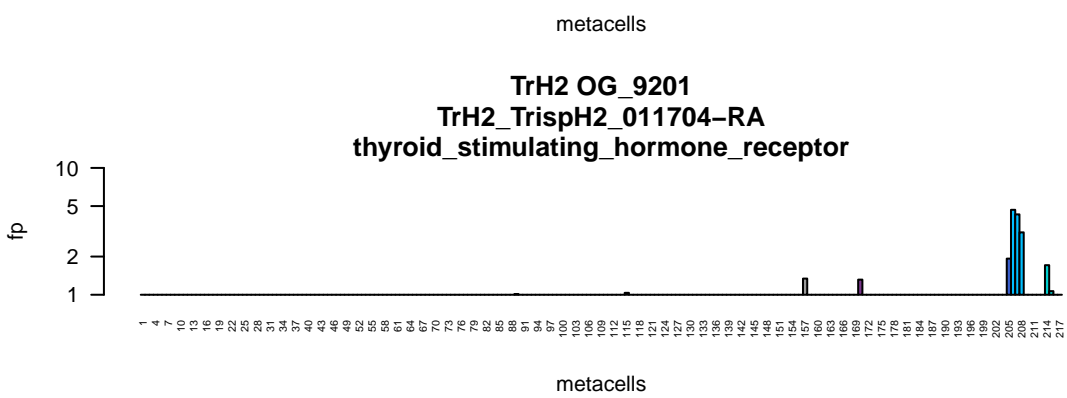
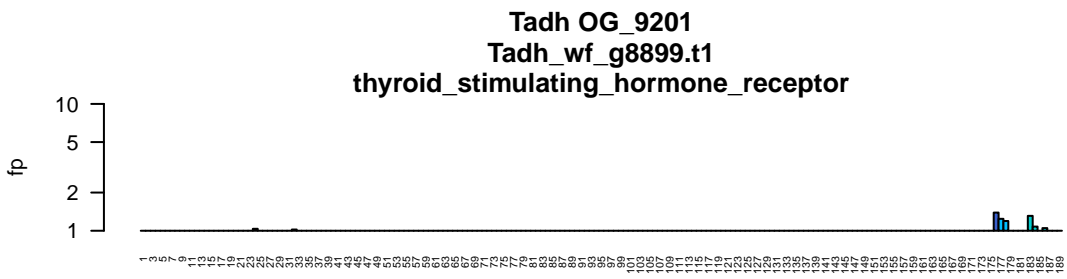


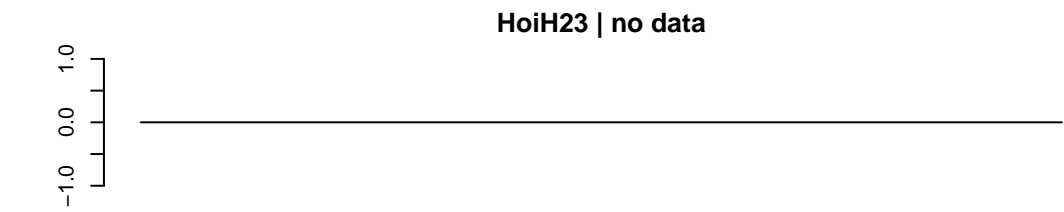
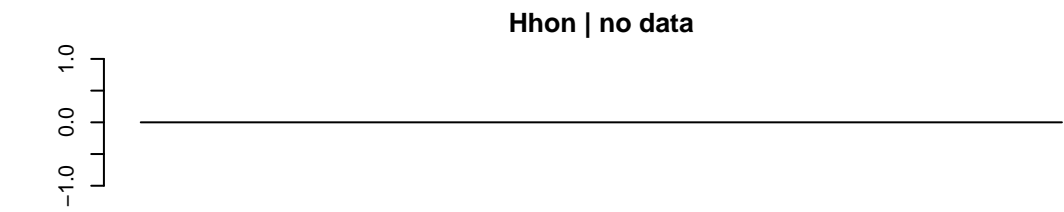
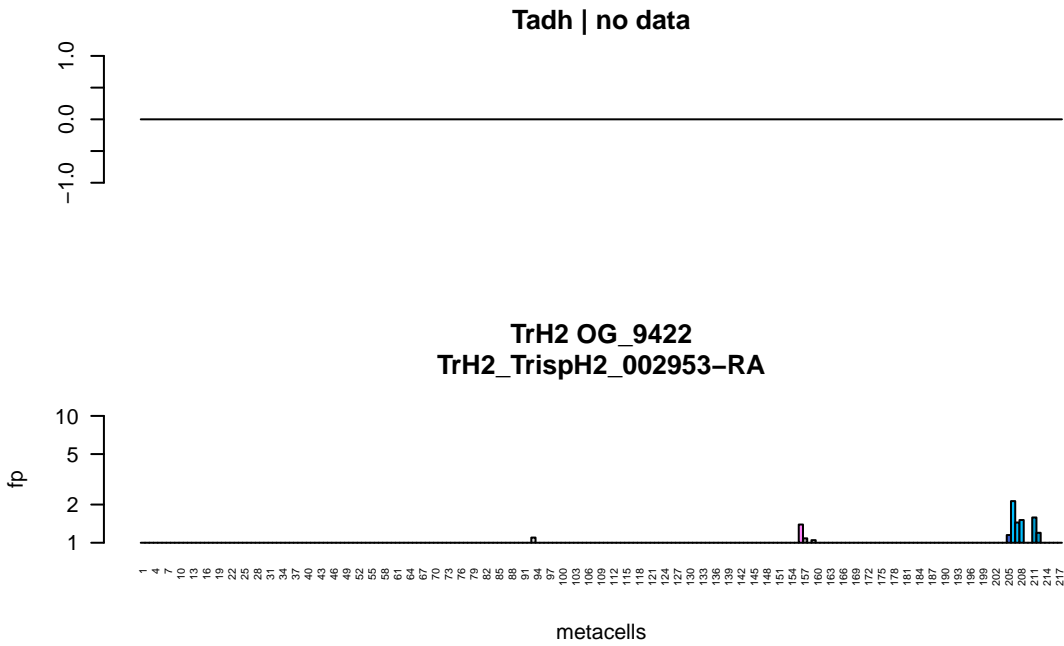


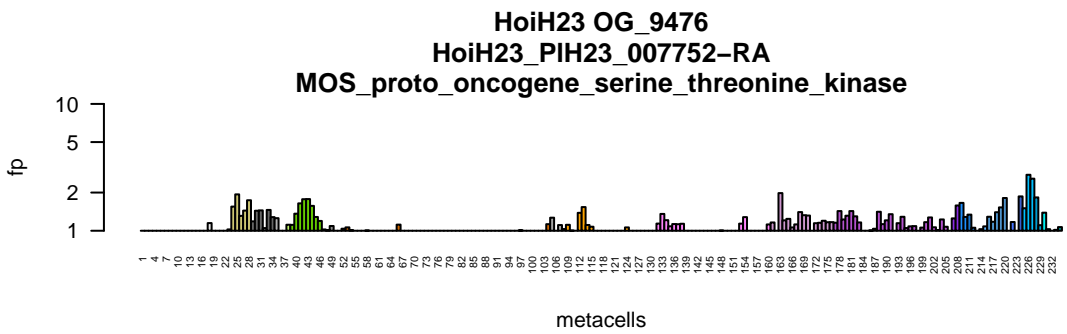
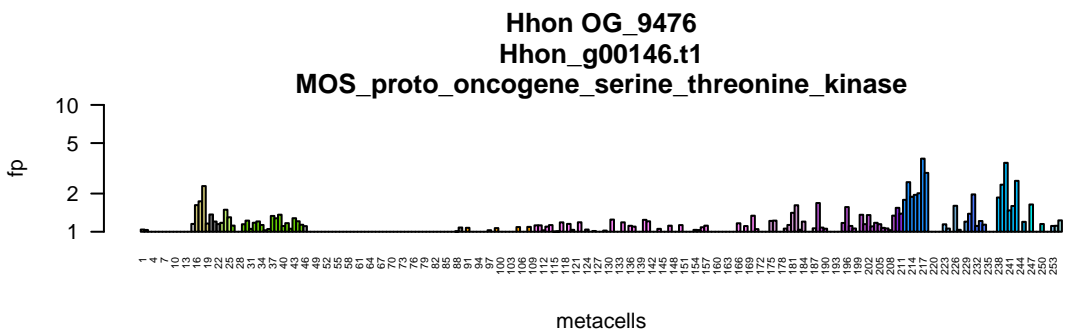
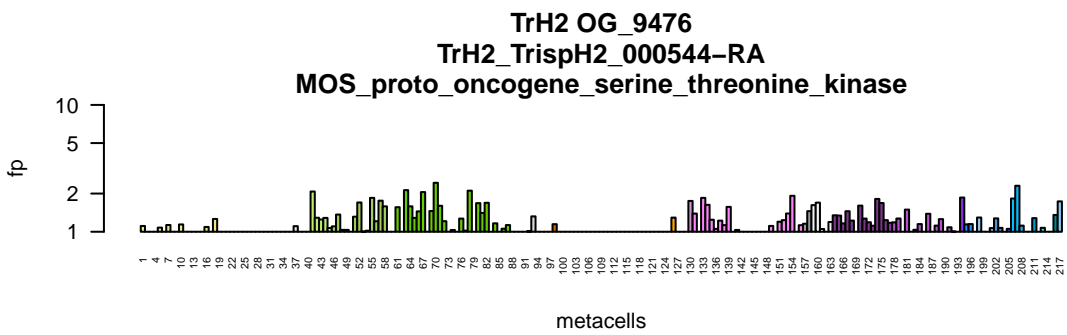
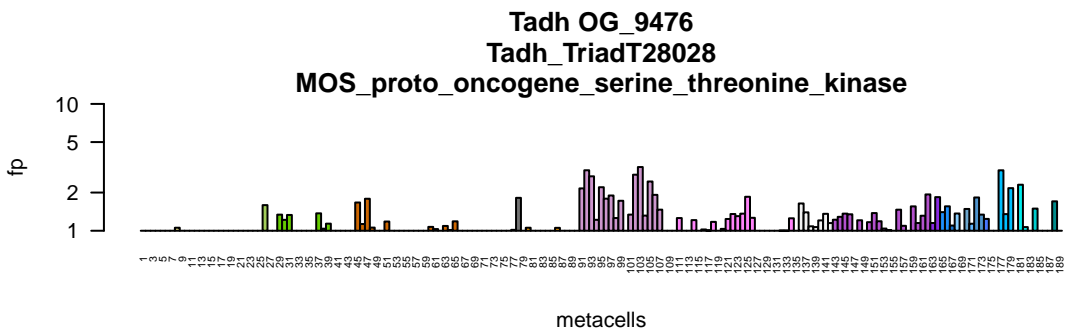


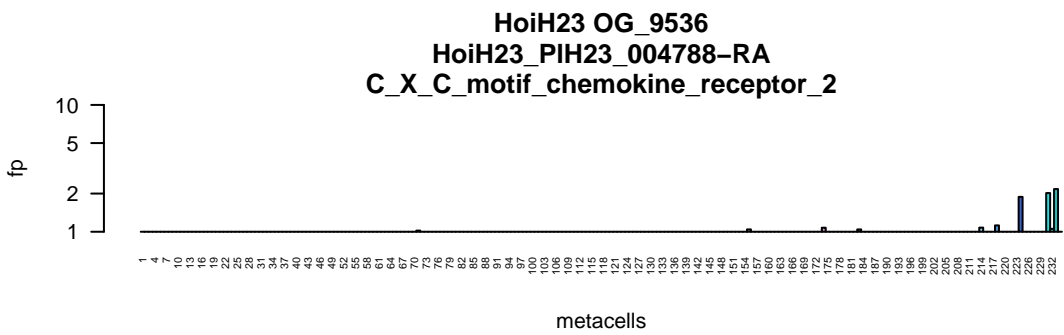
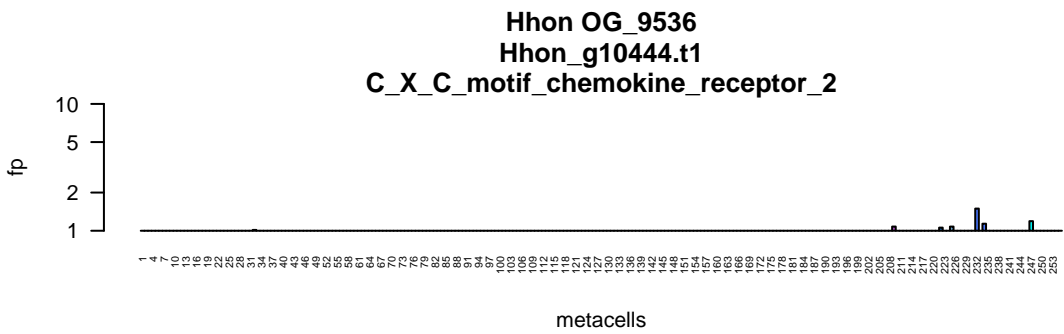
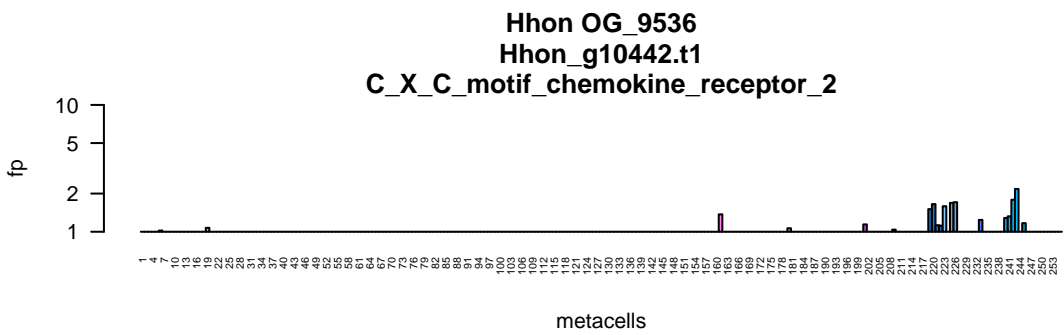
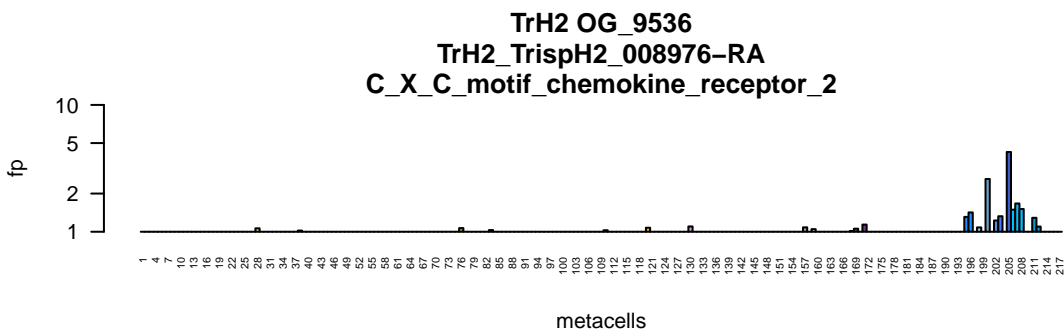
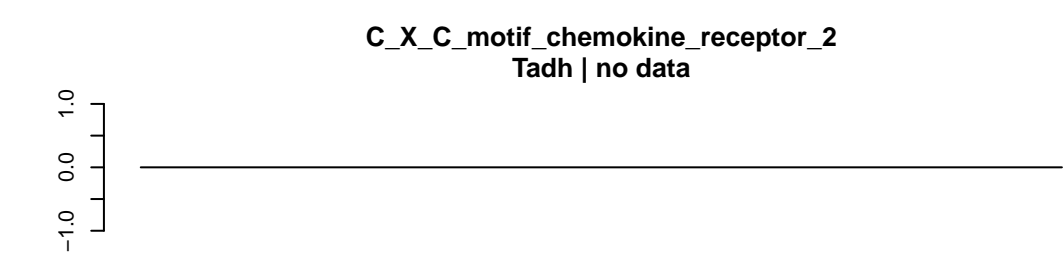


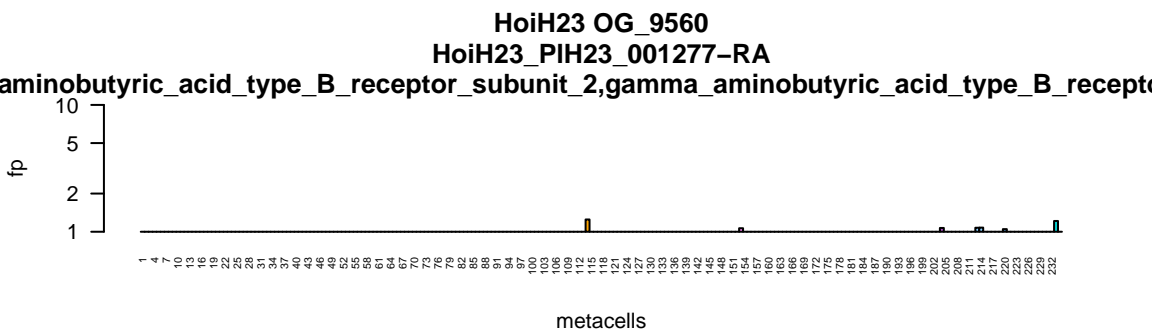
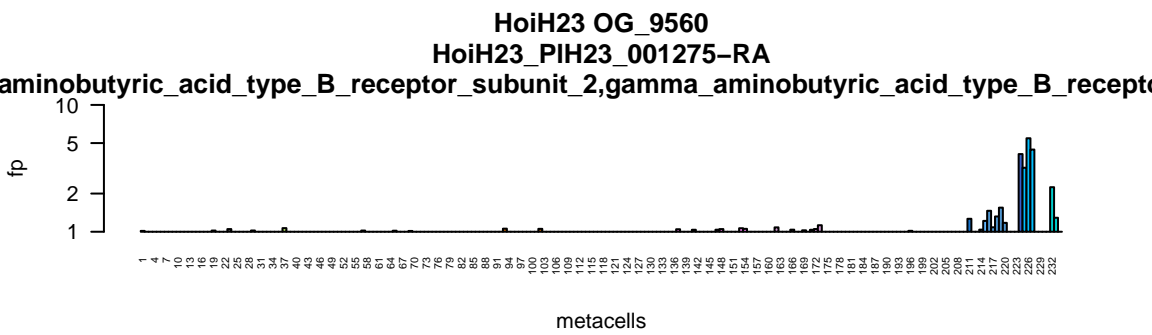
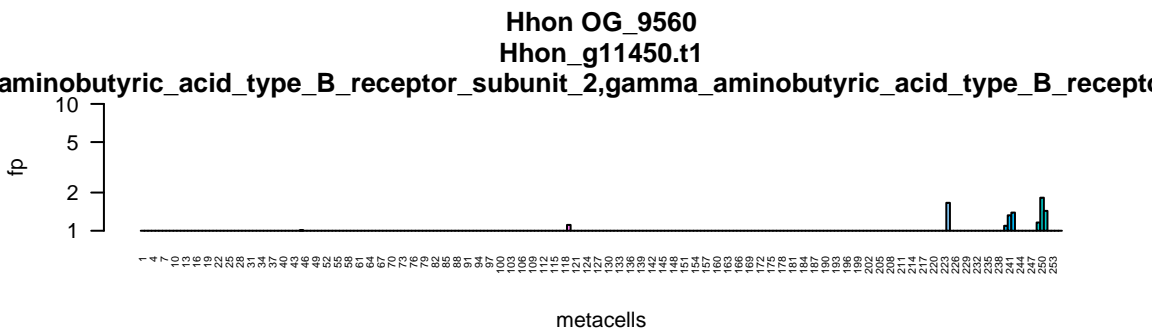
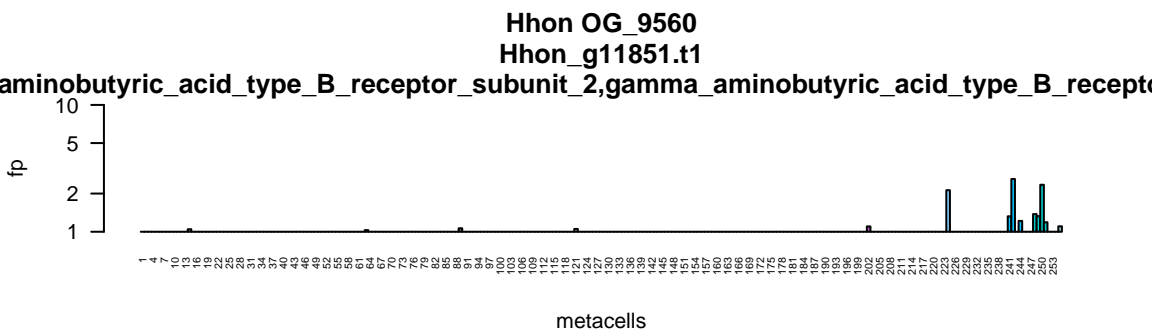
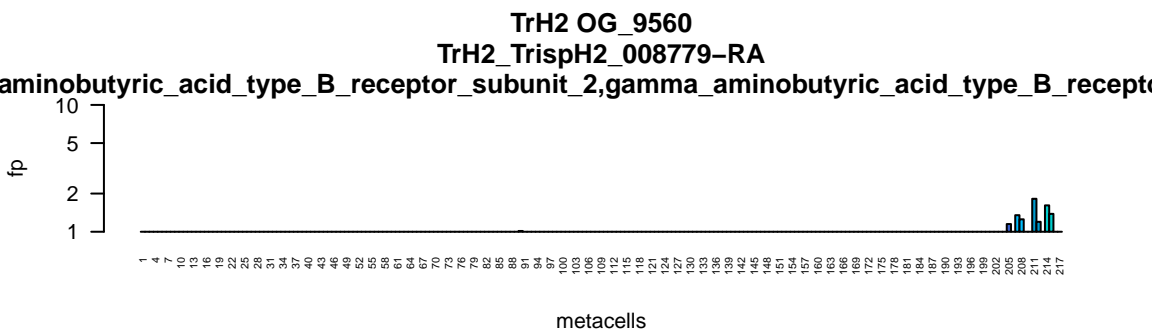
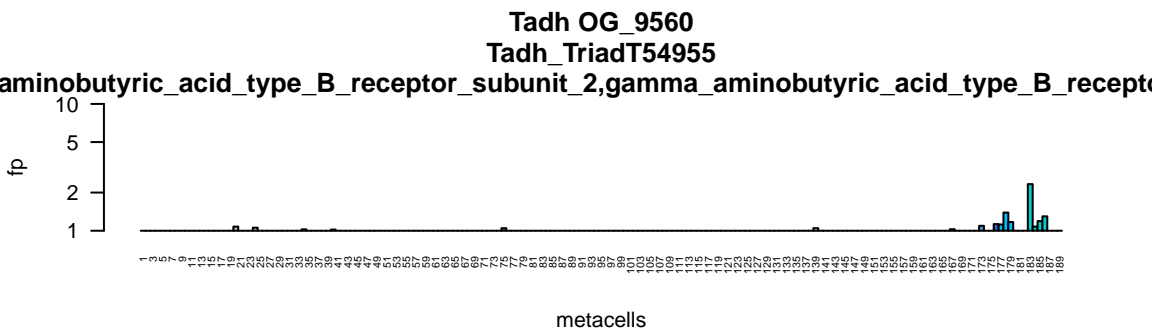


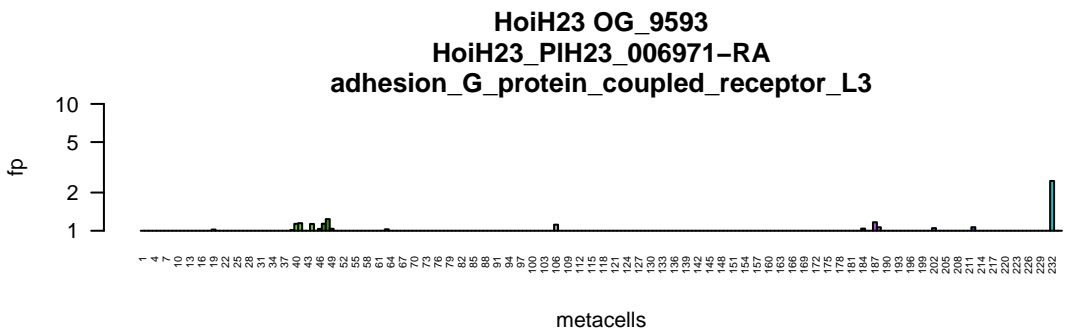
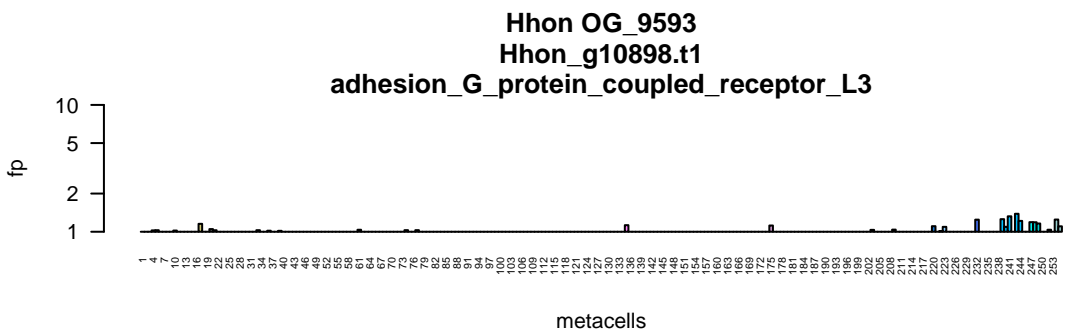
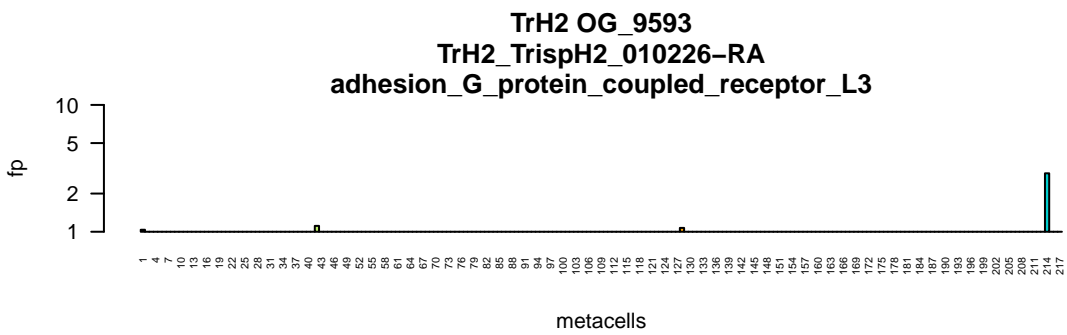
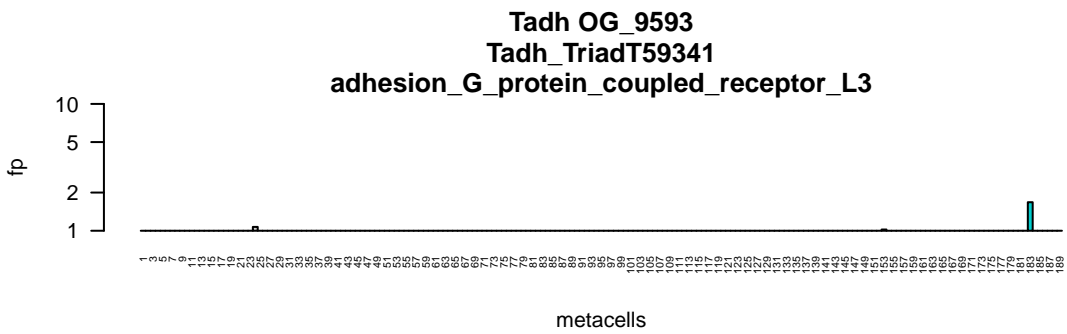






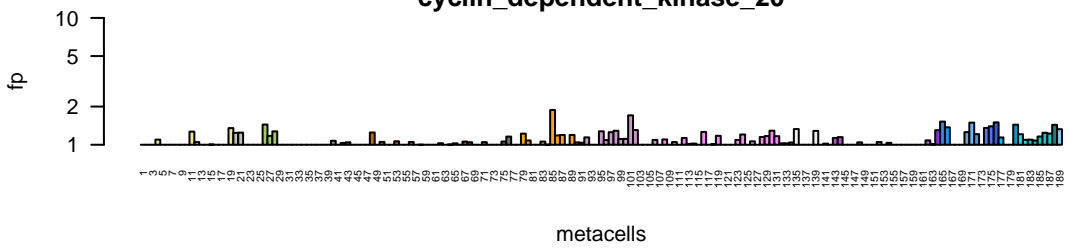




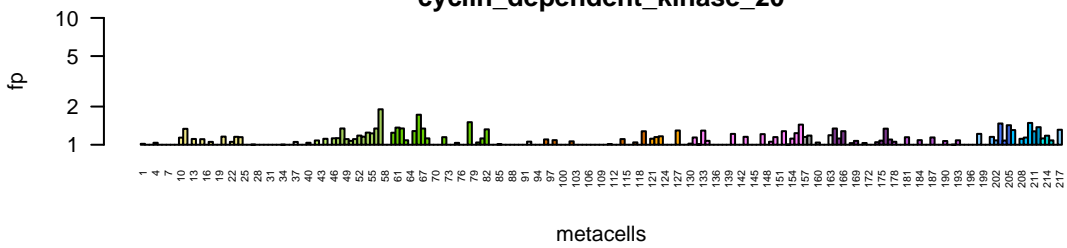




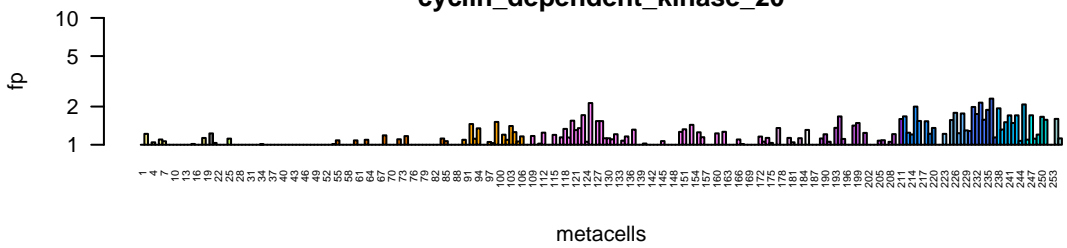
Tadh OG\_9697  
Tadh\_TriadT20272  
cyclin\_dependent\_kinase\_20



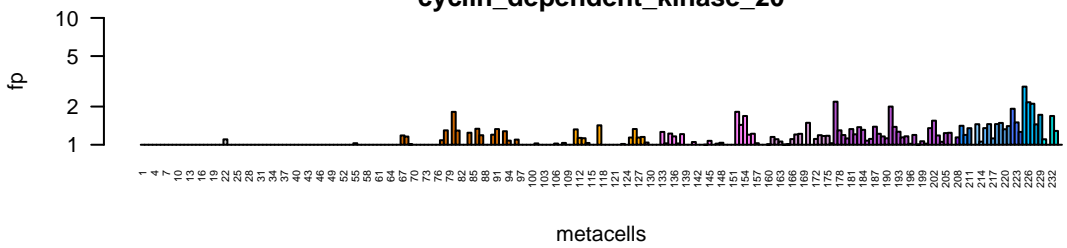
TrH2 OG\_9697  
TrH2\_TrispH2\_000088-RA  
cyclin\_dependent\_kinase\_20



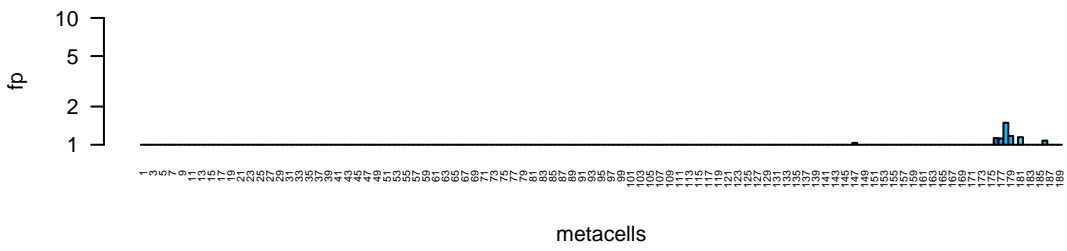
Hhon OG\_9697  
Hhon\_g09947.t1  
cyclin\_dependent\_kinase\_20



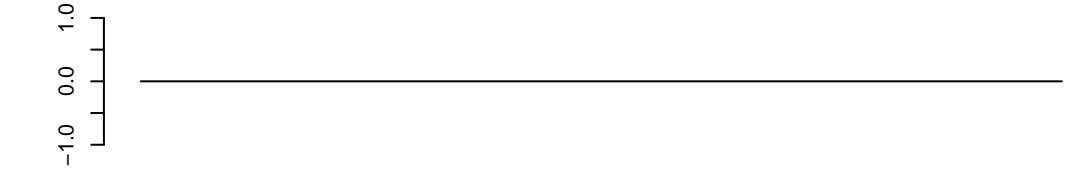
HoiH23 OG\_9697  
HoiH23\_PIH23\_001922-RA  
cyclin\_dependent\_kinase\_20



Tadh OG\_9754  
Tadh\_TriadT58809



TrH2 | no data



Hhon | no data



HoiH23 | no data



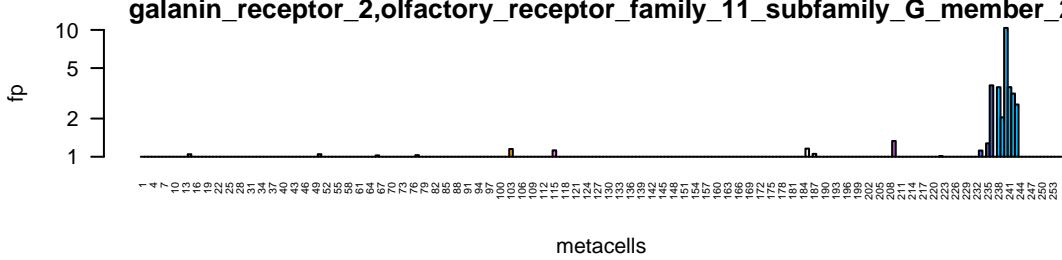
galanin\_receptor\_2,olfactory\_receptor\_family\_11\_subfamily\_G\_member\_2  
Tadh | no data



galanin\_receptor\_2,olfactory\_receptor\_family\_11\_subfamily\_G\_member\_2  
TrH2 | no data



Hhon OG\_9909  
Hhon\_g08355.t1  
galanin\_receptor\_2,olfactory\_receptor\_family\_11\_subfamily\_G\_member\_2



galanin\_receptor\_2,olfactory\_receptor\_family\_11\_subfamily\_G\_member\_2  
HoiH23 | no data



