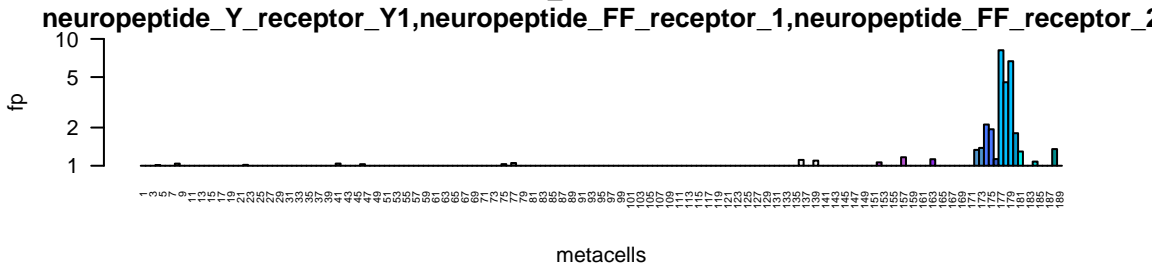
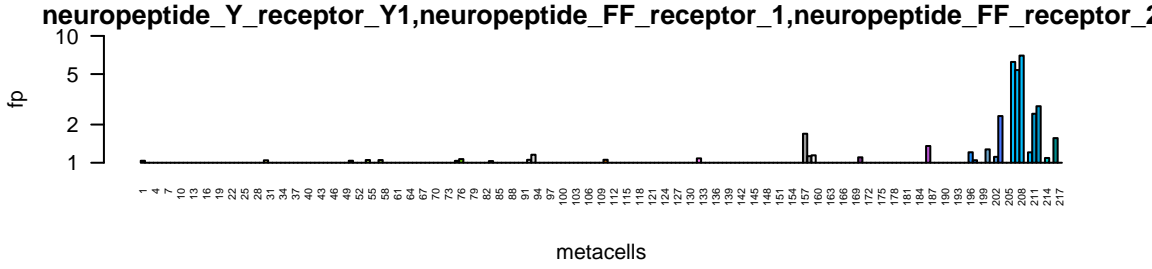


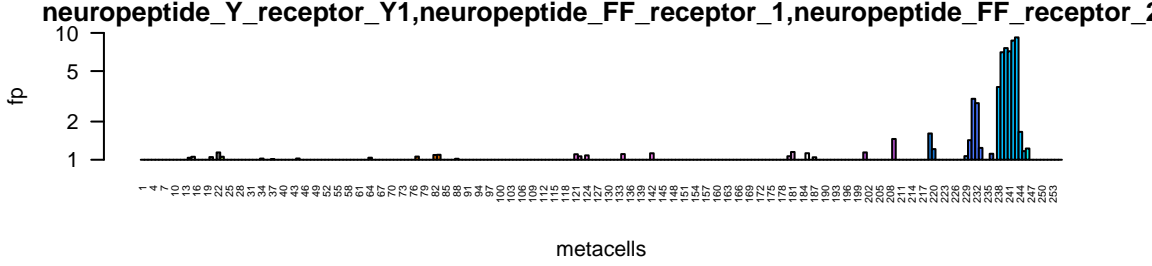
Tadh OG\_6978  
Tadh\_TriadT32926



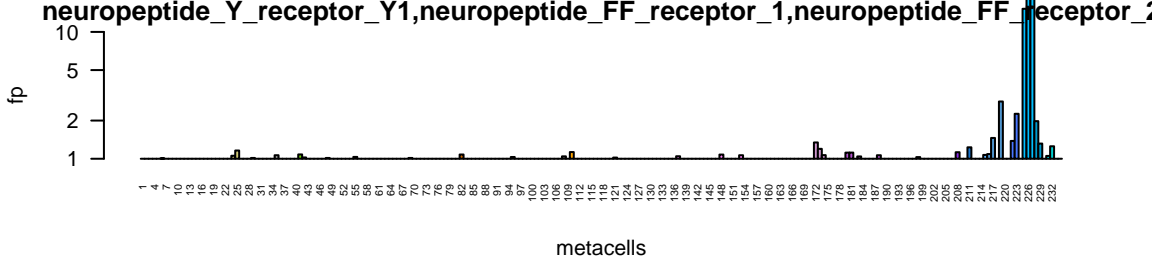
TrH2 OG\_6978  
TrH2\_TrispH2\_009841-RA

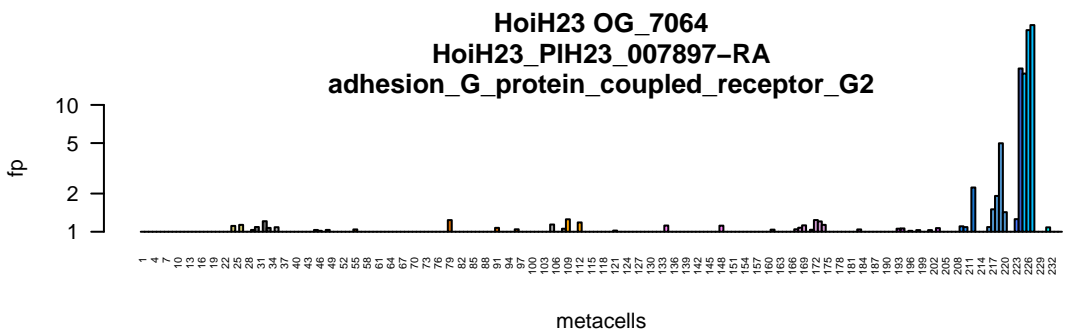
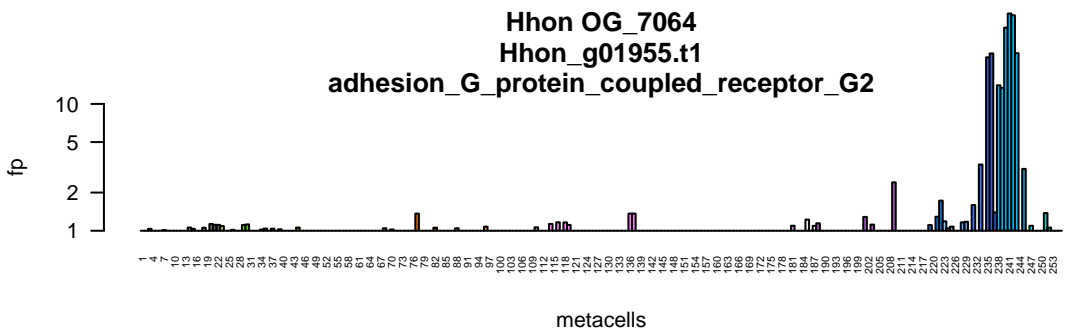
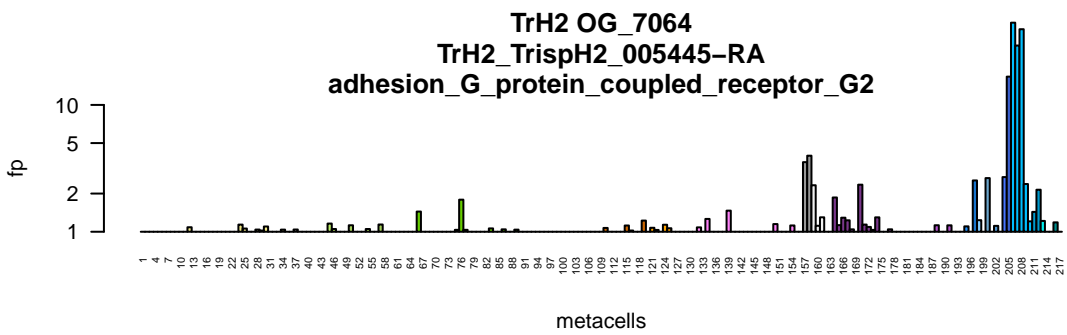
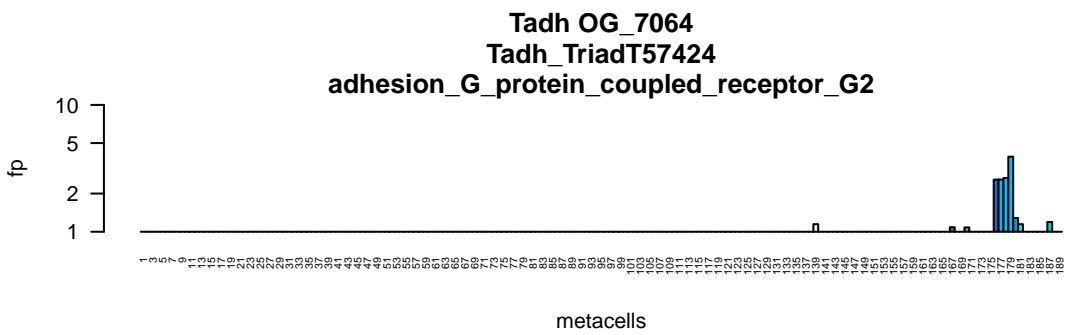
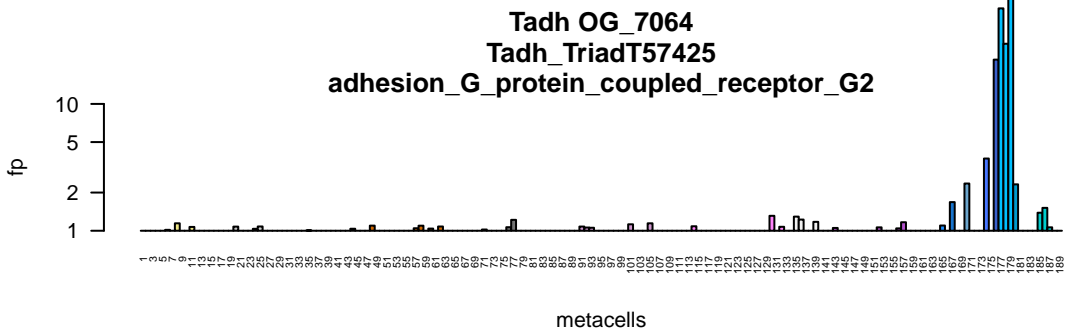


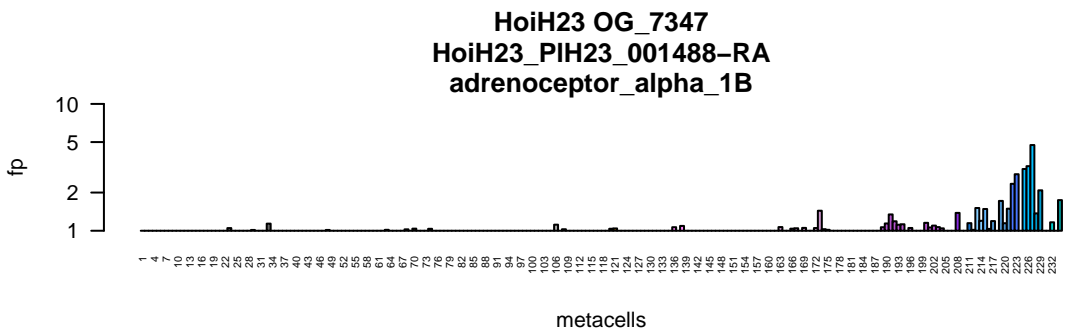
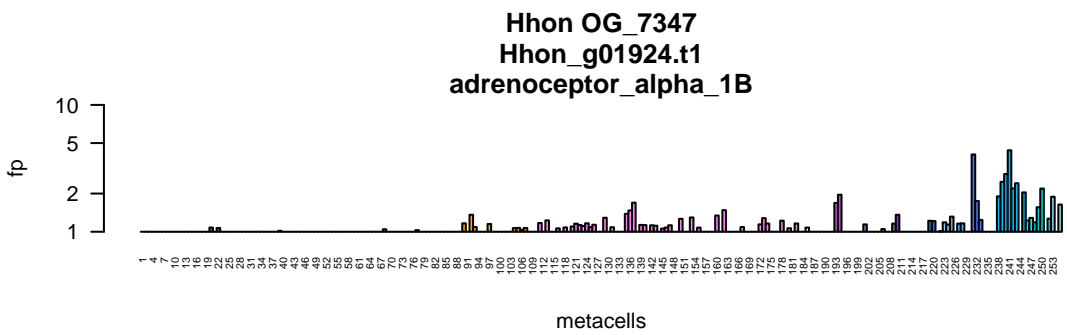
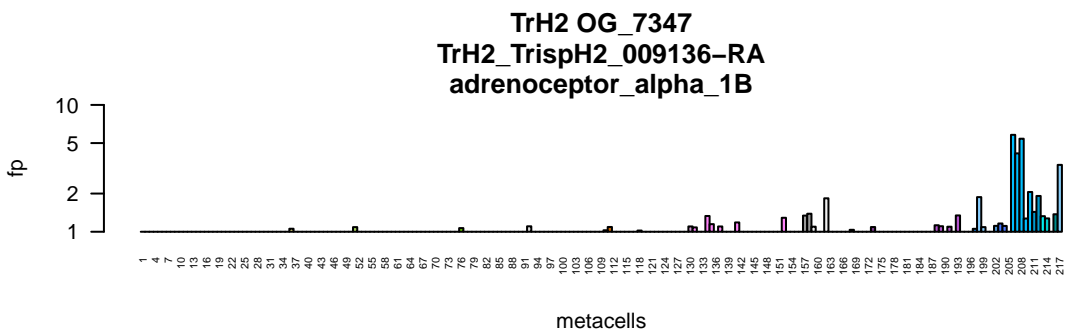
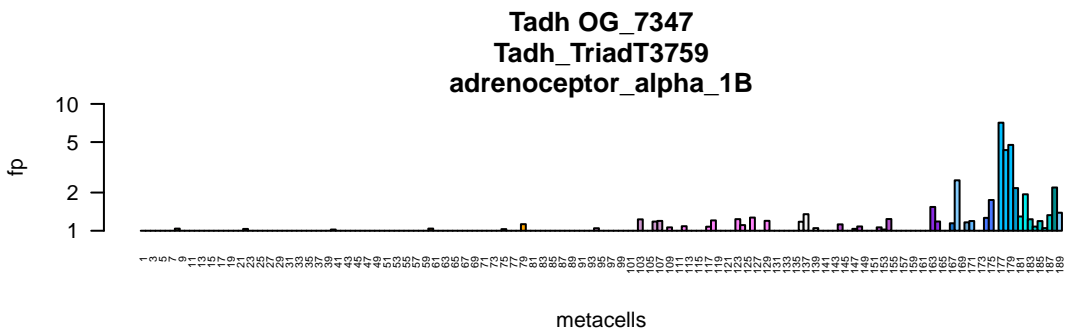
Hhon OG\_6978  
Hhon\_g04425.t1



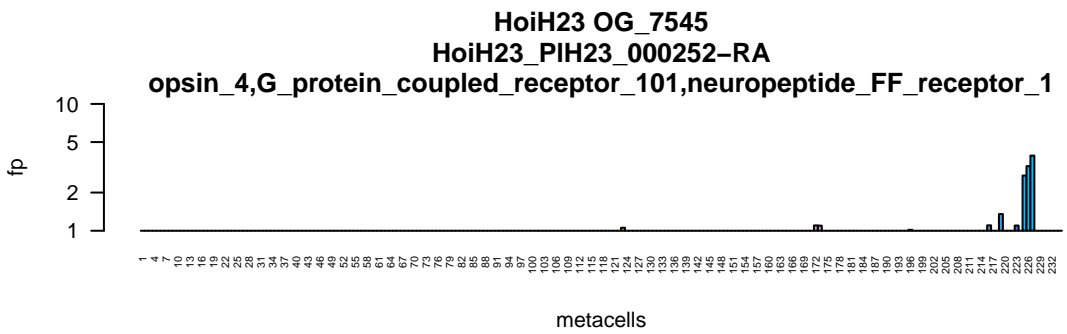
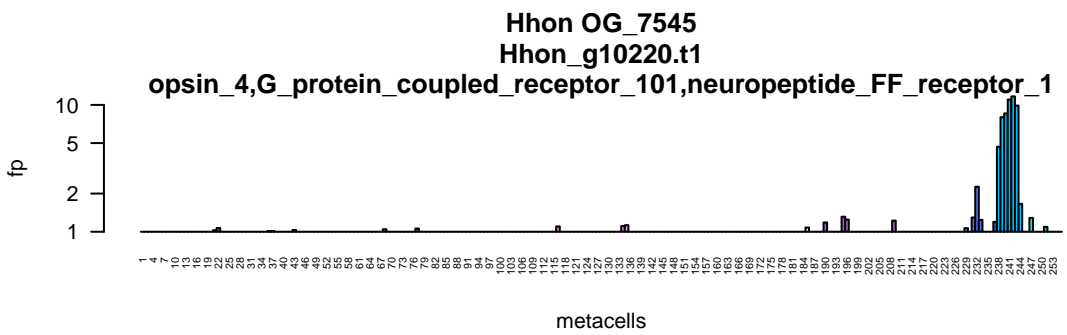
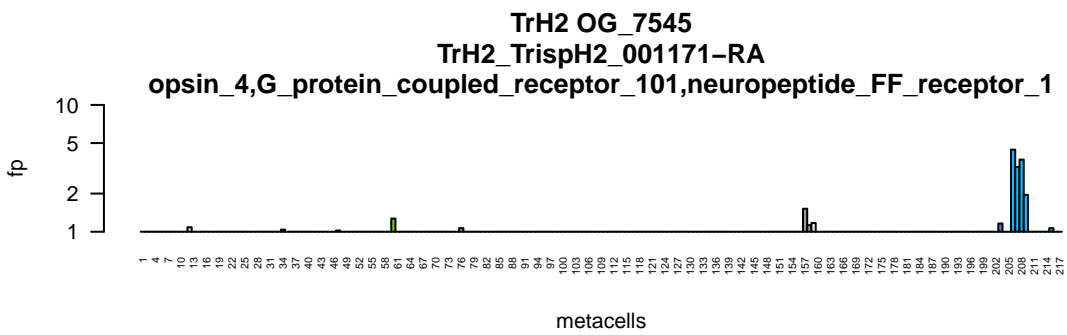
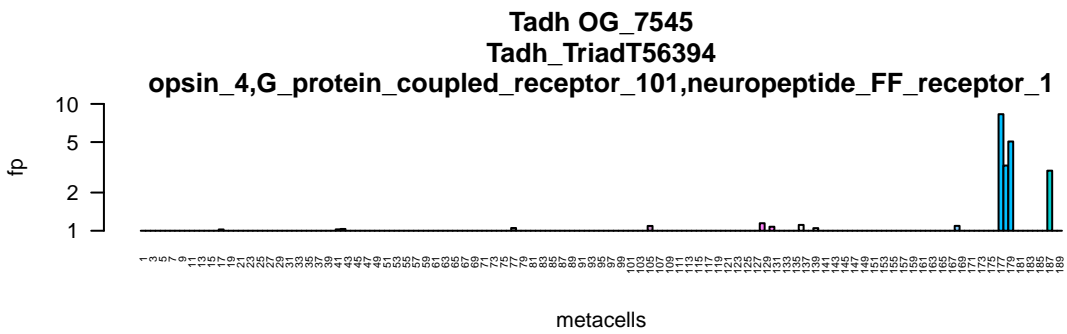
HoiH23 OG\_6978  
HoiH23\_PIH23\_011166-RA







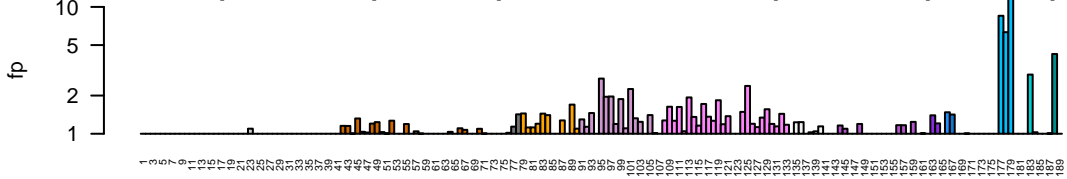




Tadh OG\_7669

Tadh\_TriadT57516

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

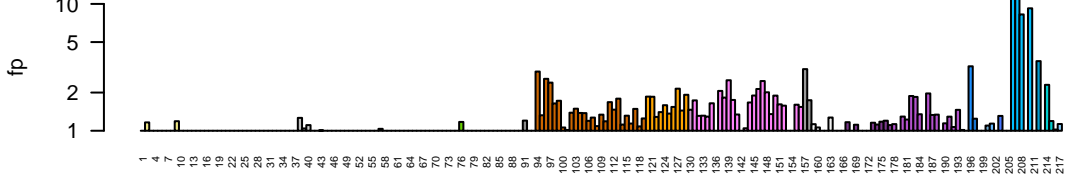


metacells

TrH2 OG\_7669

TrH2\_TrispH2\_004529-RA

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

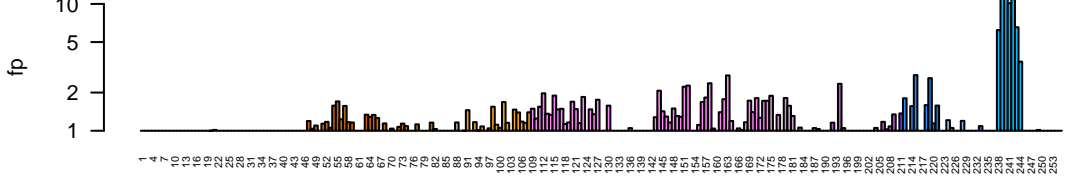


metacells

Hhon OG\_7669

Hhon\_g09503.t1

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

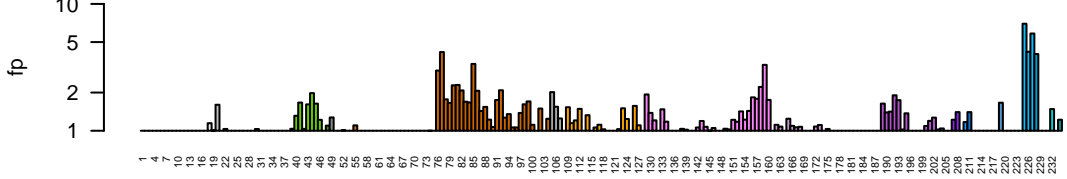


metacells

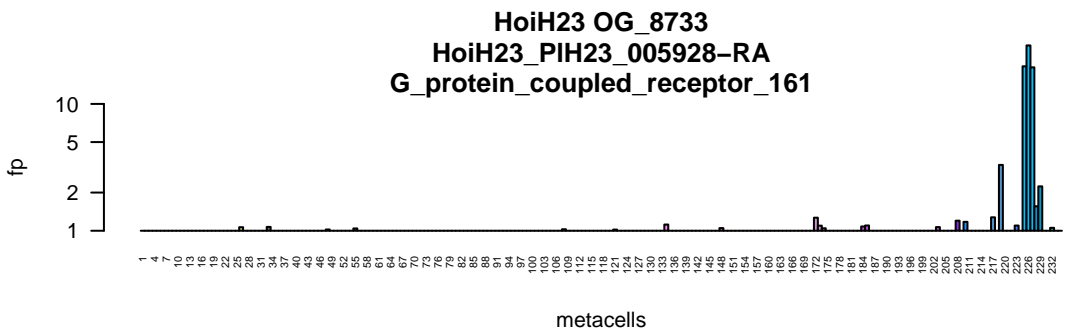
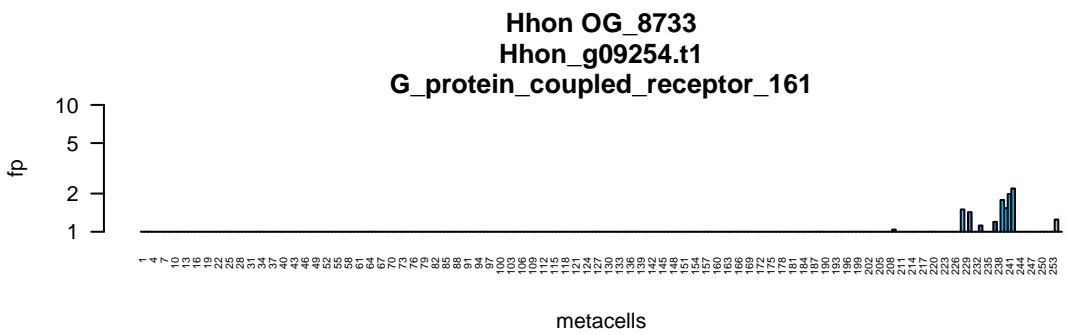
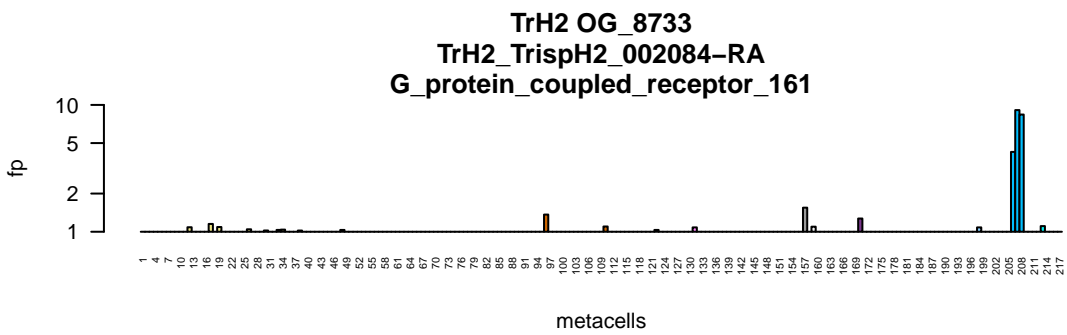
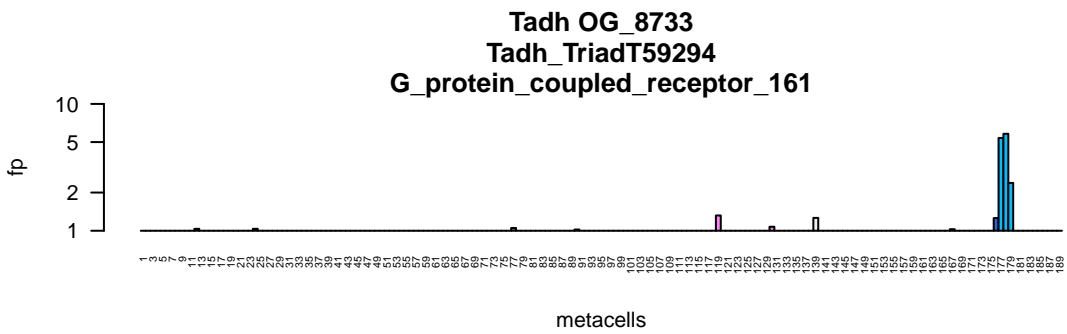
HoiH23 OG\_7669

HoiH23\_PIH23\_005798-RA

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



metacells



## Tadh OG 8863



**TrH2 OG 8863**

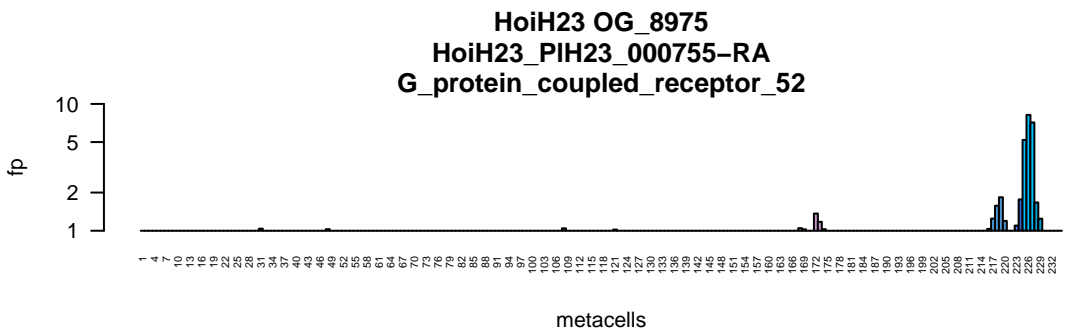
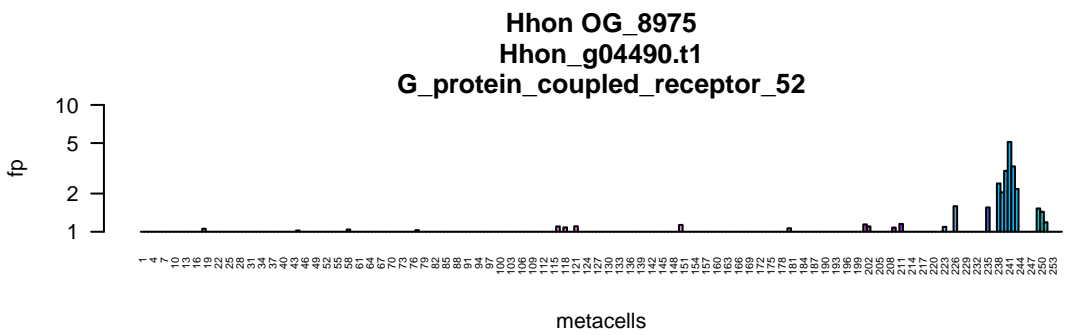
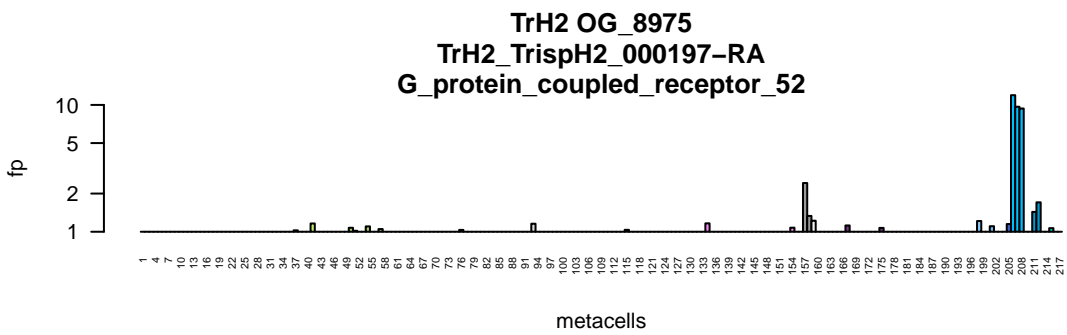
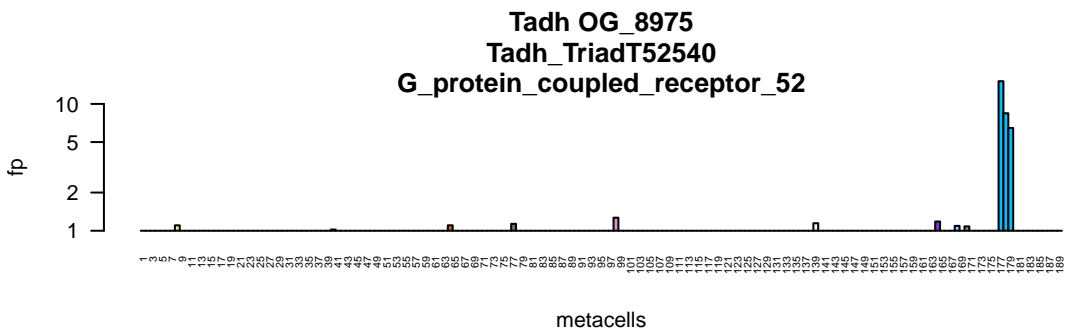


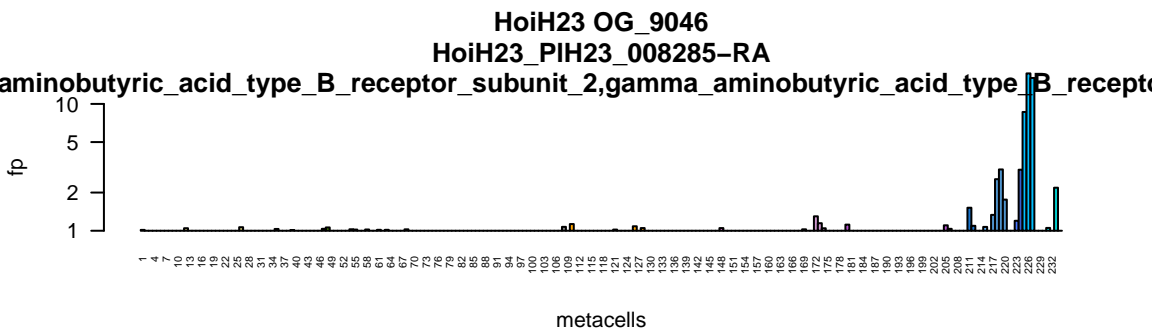
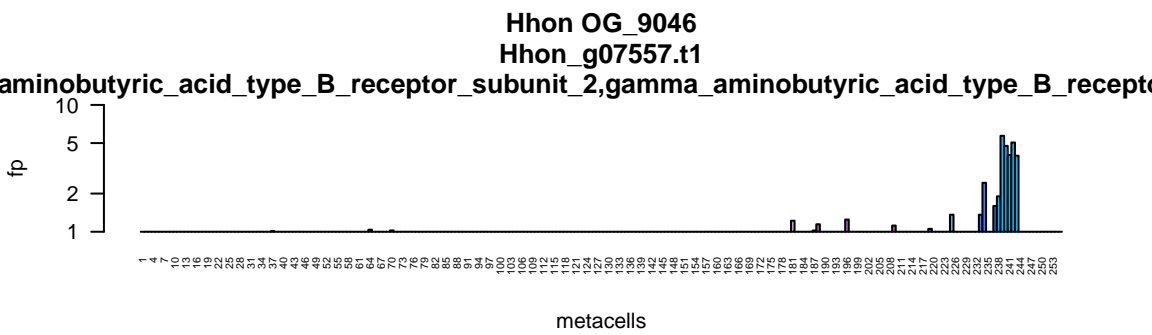
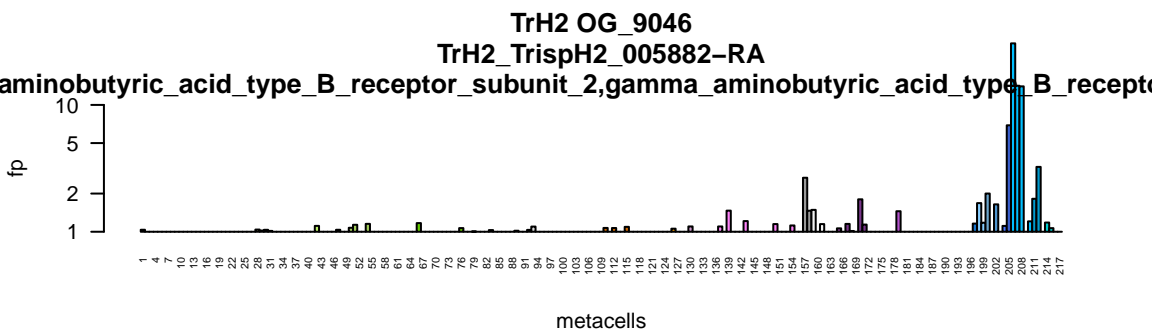
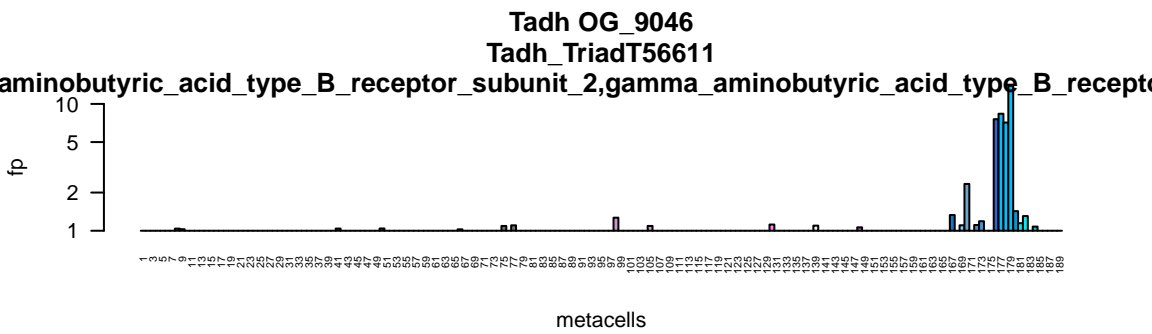
**Hhon OG 8863**

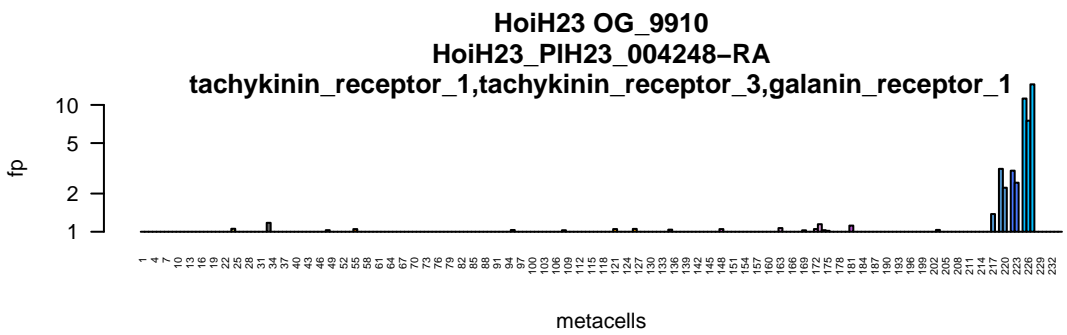
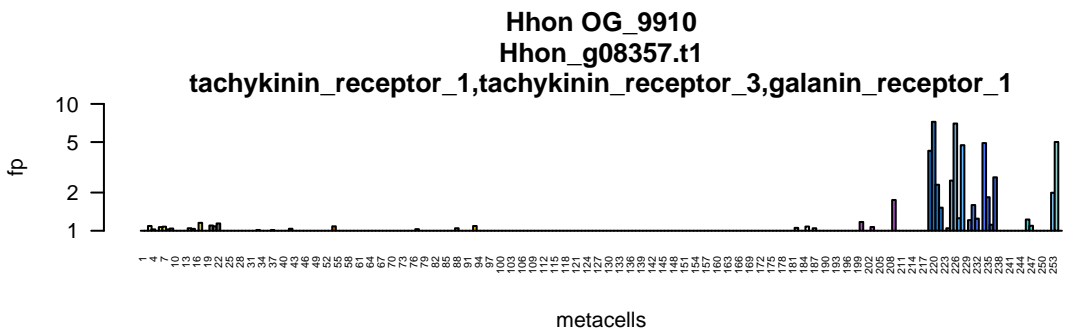
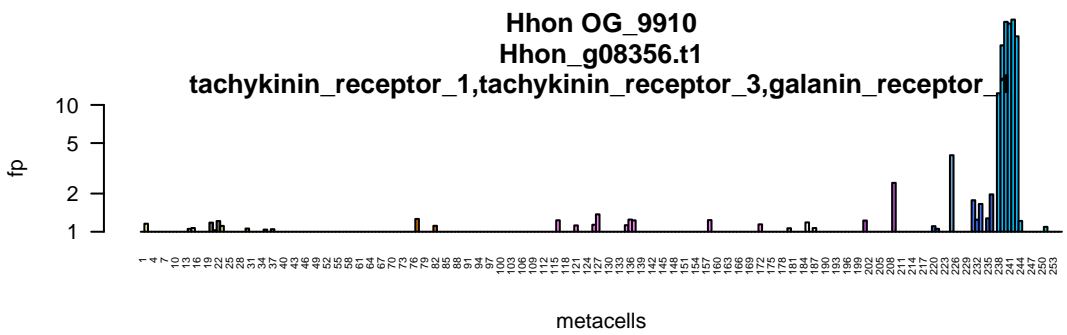
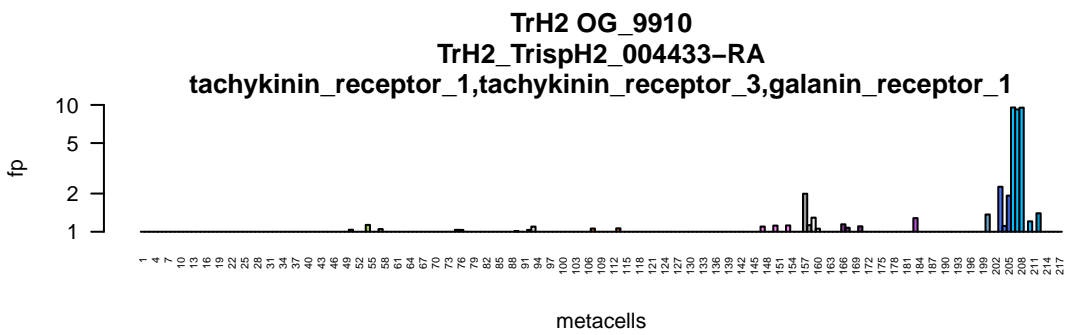
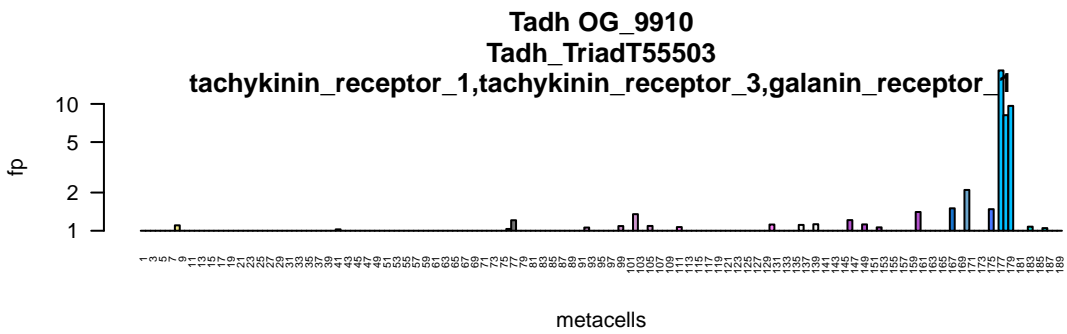


## HoiH23 OG 8863



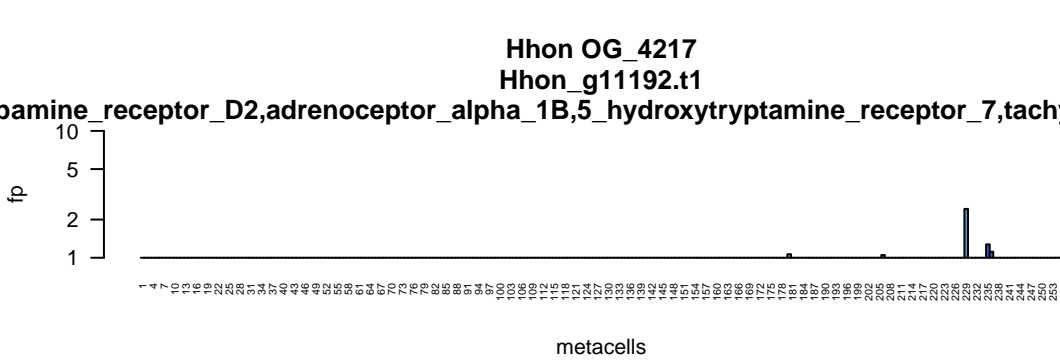
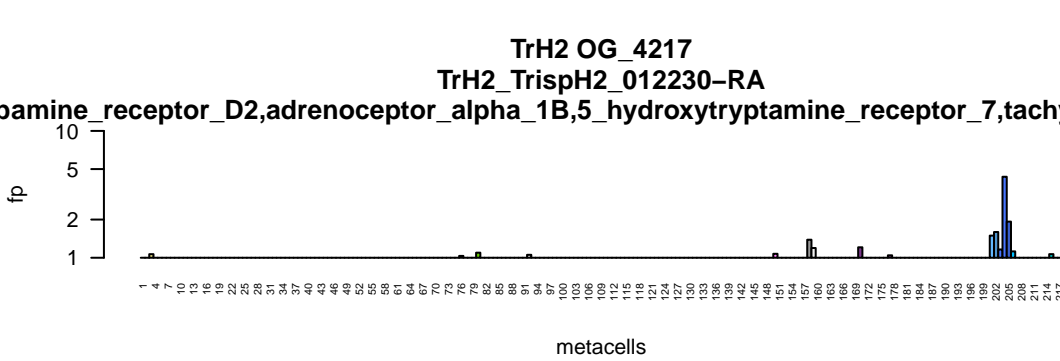
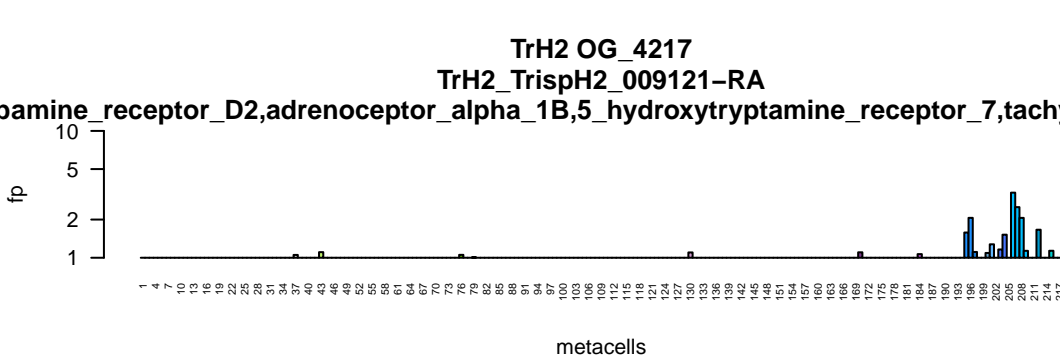
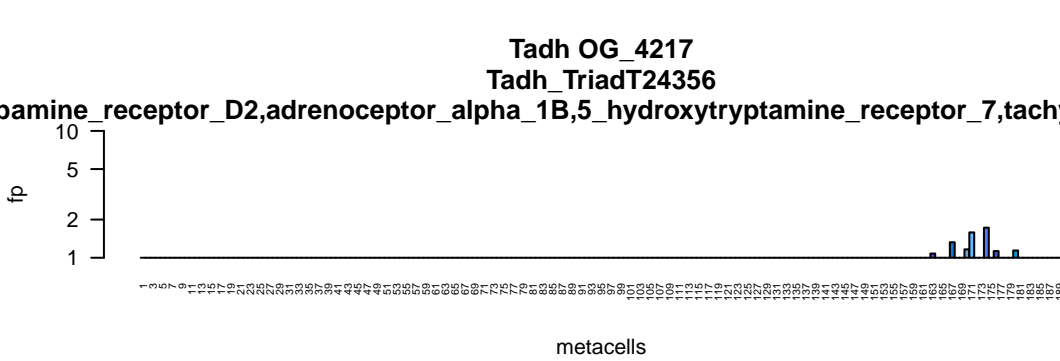




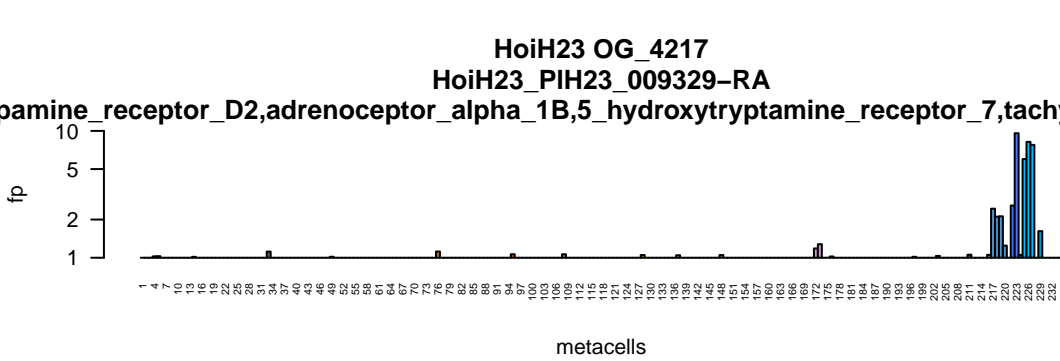


Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists 180 metacells. Most metacells have 0 false positives, but a few have 1, and a small cluster of metacells (173-187) has 2, 3, or 4 false 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	2
174	2
175	3
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
191	1
192	1
193	1
194	1
195	1
196	1
197	1
198	1
199	1
200	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 metacell IDs from 1 to 232. Most metacells have a false positive count of 1. Metacells 169, 172, and 178 have a false positive count of 2. Metacells 214, 215, 216, 217, 218, 219, 220, 221, 222, 223, 224, 225, 226, 227, 228, 229, 230, 231, 232 have false positive counts of 3 or more.





**Tadh OG\_4391**  
**Tadh\_wf\_g11074.t1**

or\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KISS1\_receptor

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

TrH2 OG\_4391  
TrH2\_TripH2\_009542-RA

or\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KISS1\_receptor

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	1.2
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	1.5
160	1
163	1
166	1
169	1
172	1.3
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
202	1
205	2.0
206	2.0
209	1
211	2.0
214	1
217	1

TrH2 OG\_4391  
TrH2\_TrispH2\_009543-RA

or\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KISS1\_receptor

TrH2 OG\_4391  
TrH2\_TripH2\_011814-RA

or\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KISS1\_receptor

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
202	1
205	4
208	3
211	4
214	1
217	1

TrH2 OG\_4391  
TrH2\_TripH2\_012158-RA

or\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KISS1\_receptor

fp

metacells

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

TrH2 OG\_4391  
TrH2\_TripH2\_011813-RA

or\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KISS1\_receptor

metacells

TrH2 OG\_4391  
TrH2\_TrispH2\_012225-RA

for\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KISS1\_receptor

fp

metacells

TrH2 OG\_4391  
TrH2\_TrispH2\_011811-RA

or\_2,hypocretin\_receptor\_2,hypocretin\_receptor\_1,neuropeptide\_Y\_receptor\_Y1,KISS1\_receptor

metacells	fp
1	1
7	1
13	1
16	1
19	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	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	10
214	2
217	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 small cluster of higher values (2, 3, 5, 6) for metacells 228 through 232.

metacell	fp
1	1
4	1
10	1
13	1
19	1
22	1
28	1
31	1
34	1
38	1
40	1
43	1
46	1
52	1
55	1
61	1
64	1
73	1
76	1
82	1
85	1
88	1
91	1
94	1
103	1
108	1
112	1
115	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
178	1
184	1
187	1
190	1
196	1
199	1
205	1
208	1
211	1
214	1
217	2
220	3
223	5
228	6
229	1
232	1

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.

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 48 metacells by index. Most metacells have a false positive count of 1, with a few having 2 or 3. Metacell 228 has the highest count at 6.

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	1
223	1
228	6
232	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.

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 at the end of the list.

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	1
85	1
88	1
91	1
94	1
97	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
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	5
228	6
229	5
232	3

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
61	1
64	1
70	1
73	1
76	1
82	1
85	1
88	1
91	1
94	1
102	1
103	1
106	1
112	1
115	1
118	1
121	1
124	1
127	1
133	1
136	1
145	1
148	1
154	1
157	1
163	1
166	1
169	1
175	1
178	1
184	1
187	1
190	1
199	1
186	1
205	1
208	1
211	1
214	1
217	1
220	1
223	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 show significantly higher values, around 8 and 6 respectively.

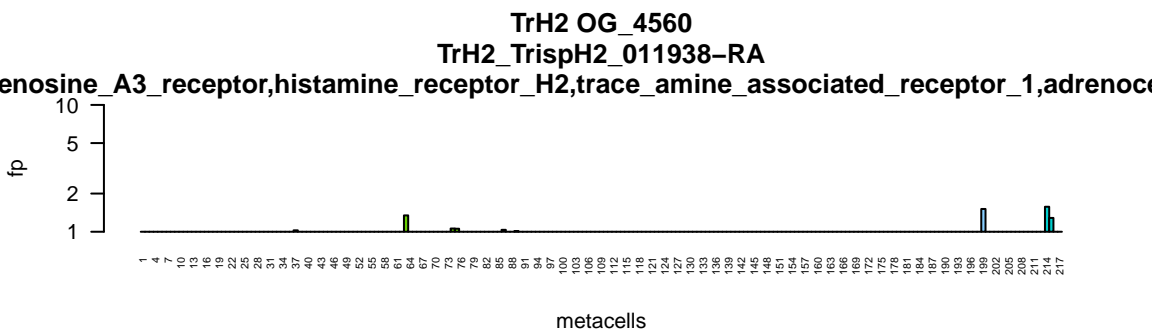
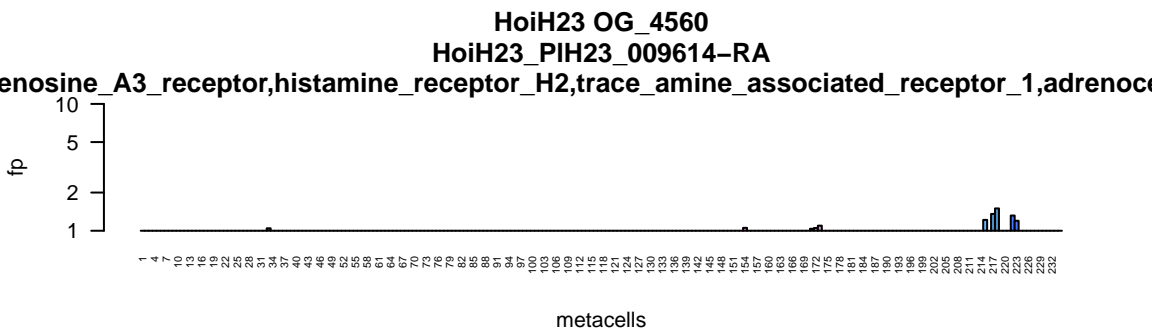
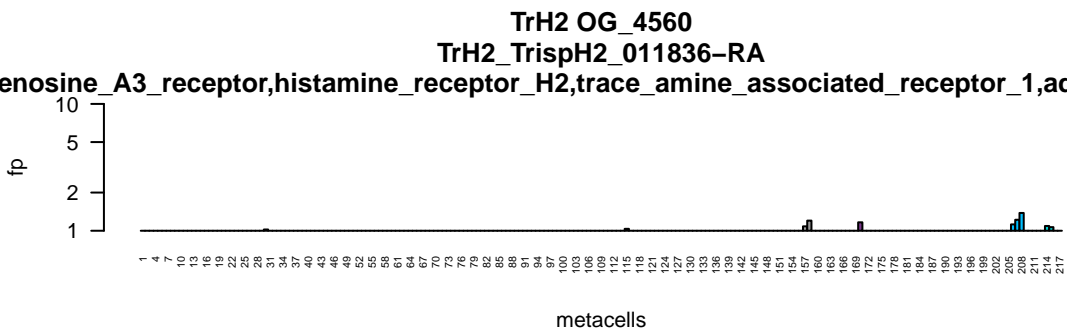
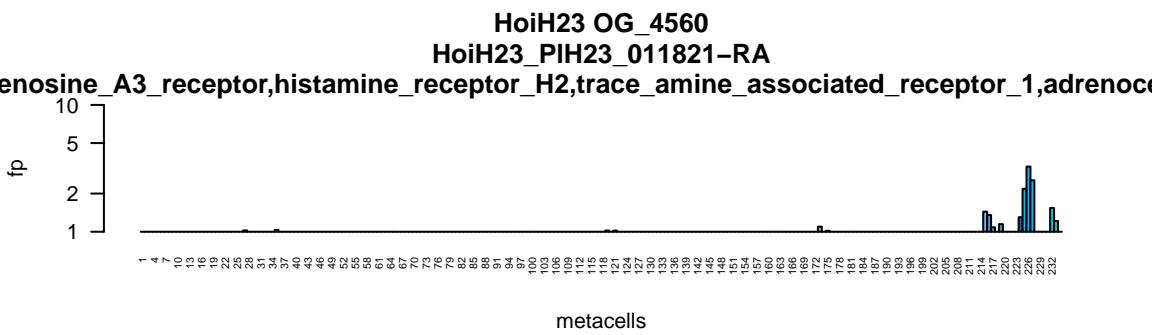
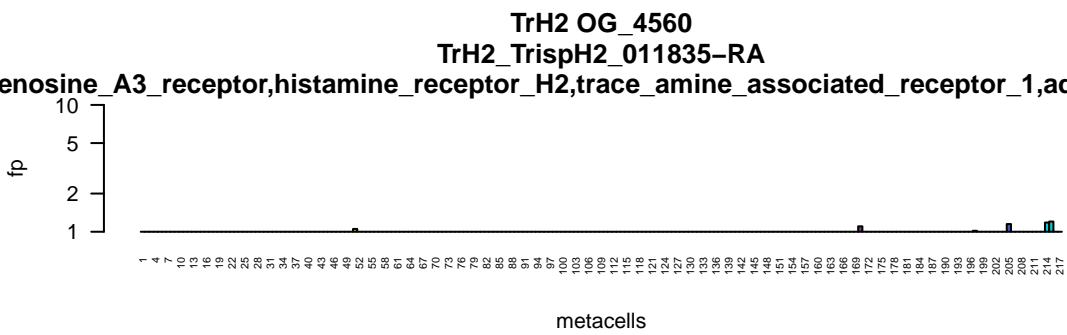
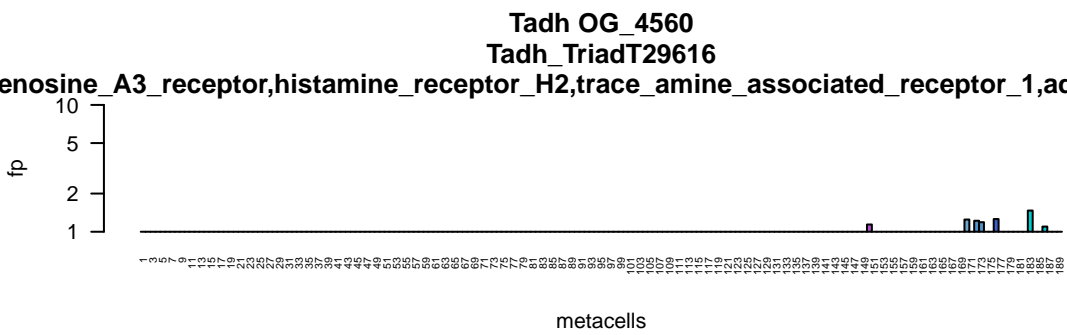
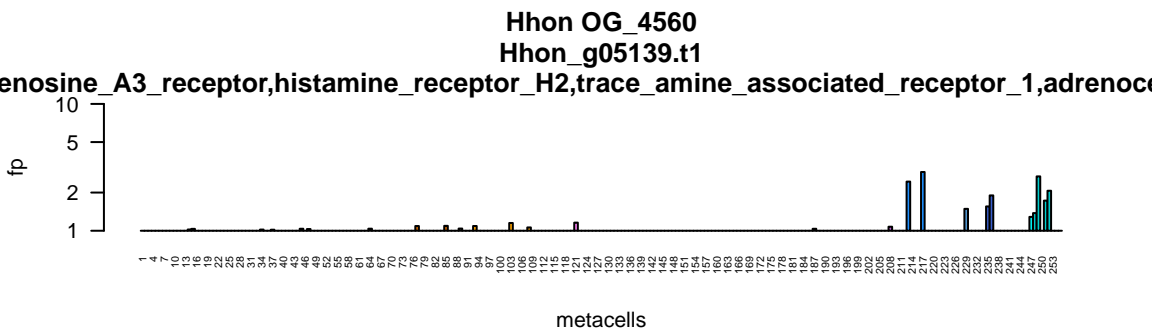
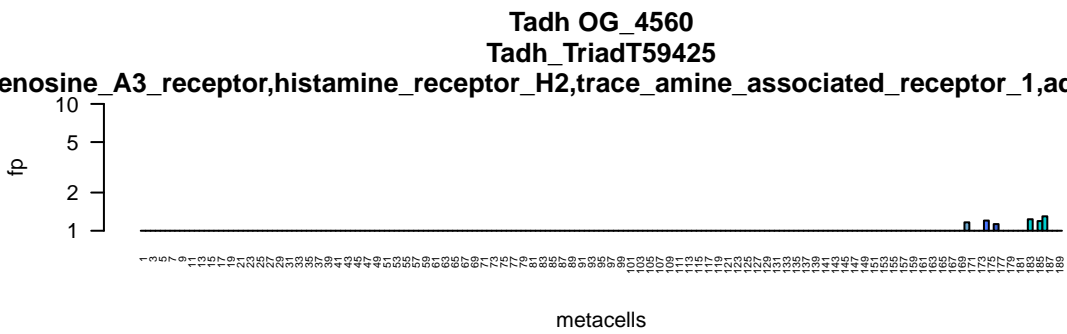
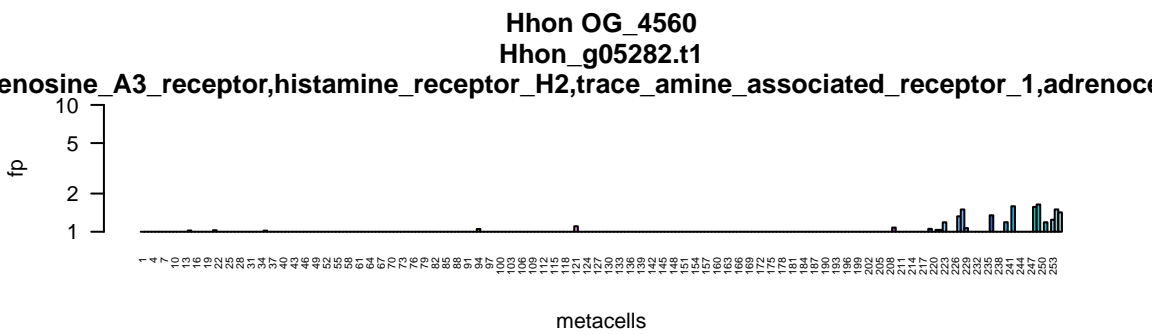
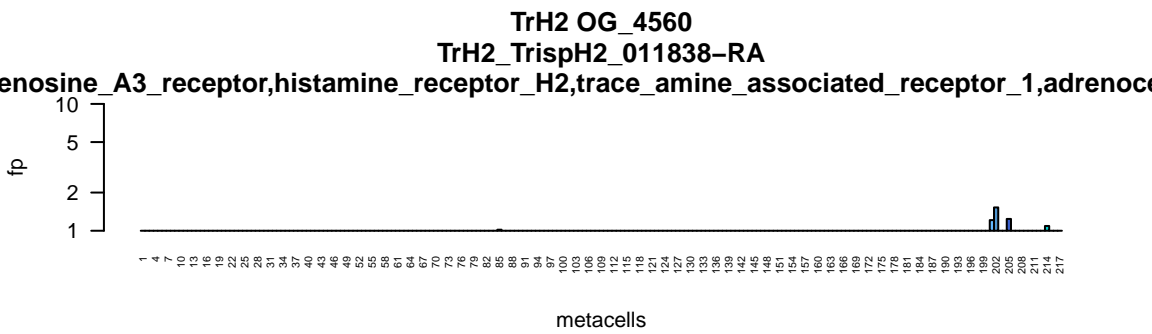
metacell	fp
1	1
4	1
10	1
13	1
19	1
22	1
25	1
28	1
31	1
34	1
40	1
43	1
46	1
55	1
58	1
61	1
64	1
73	1
76	1
82	2
85	2
88	1
91	1
94	1
97	1
102	1
103	1
106	1
112	1
115	1
118	1
121	1
124	1
127	1
133	1
136	1
142	1
145	1
148	1
154	1
157	1
163	1
166	1
169	1
175	1
178	1
181	1
184	1
187	1
190	1
196	1
199	1
205	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 239. 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
38	1
40	1
43	1
46	1
52	1
55	1
58	1
61	1
64	1
73	1
76	1
82	1
85	1
88	1
91	1
94	1
97	1
102	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 a false positive count of 1, with a small cluster of higher counts (2, 3, 4) for metacells 228, 229, and 232.

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
110	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
178	1
184	1
187	1
190	1
196	1
199	1
205	1
208	1
211	1
214	1
217	1
220	1
222	1
223	1
228	2
229	3
232	4



Bar chart showing the frequency of metacells (x-axis) versus the frequency of paths (fp, y-axis). The x-axis lists metacells from 1 to 189. The y-axis ranges from 1 to 10. The chart shows a distribution of path frequencies across metacells, with most metacells having a frequency of 1, and a few having higher frequencies up to 6.

Bar chart showing the frequency of metacells (x-axis) versus the frequency of pairs (fp, y-axis). The x-axis lists metacells from 1 to 189. The y-axis ranges from 1 to 10. The chart shows a distribution of frequencies, with most metacells having a frequency of 1 or 2. A notable peak is at metacell 175, which has a frequency of 3.

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 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 2 or 4. Metacells 175, 176, and 179 have the highest counts at 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. The bars are colored in a repeating pattern of light blue, dark blue, and light green. Most metacells have 0 false positives, with some having 1, 2, or 3. Metacells 178 and 181 have 5 false positives each.

metacell	fp
1	0
4	1
10	0
13	0
16	1
19	0
22	0
25	1
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	0
67	0
70	0
73	0
76	1
79	0
82	0
85	0
88	0
91	1
94	2
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	2
124	1
127	2
130	2
133	3
136	3
139	4
142	3
145	3
148	2
151	3
154	2
157	3
160	2
163	1
166	1
169	1
172	3
175	1
178	5
181	4
184	1
187	1
190	1
193	2
196	1
199	1
202	1
205	1
208	1
211	3
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 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	1.5
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	1.5
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
202	2.5
205	1.5
208	4.5
211	4
214	1
217	1

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

metacells	fp
1	1
4	1
10	1
13	1
16	1
22	1
23	1
25	1
28	1
31	1
32	1
37	1
40	1
43	2
44	1
48	1
49	1
52	1
55	1
59	1
61	1
64	1
67	1
73	1
76	1
79	1
82	1
83	1
88	1
91	1
94	1
97	1
100	1
103	1
108	1
110	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	2
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
200	1
202	2
205	1
208	1
211	1
214	1
217	1

A bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 217. Most metacells have a false positive count of 1, with a few outliers reaching up to 5.

metacell	fp
1	1
4	1
10	1
13	1
16	1
22	1
23	1
25	1
28	1
31	1
32	1
33	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

[illegible]

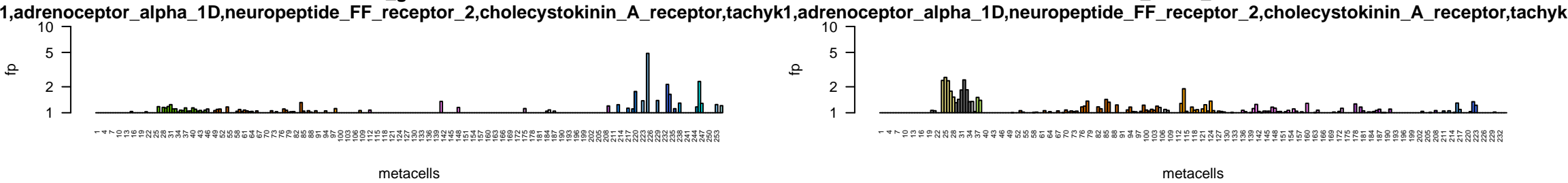
A bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 217. The chart shows a distribution of false positives across the metacells, with most having 1 or 2 false positives, and a few having up to 4.

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

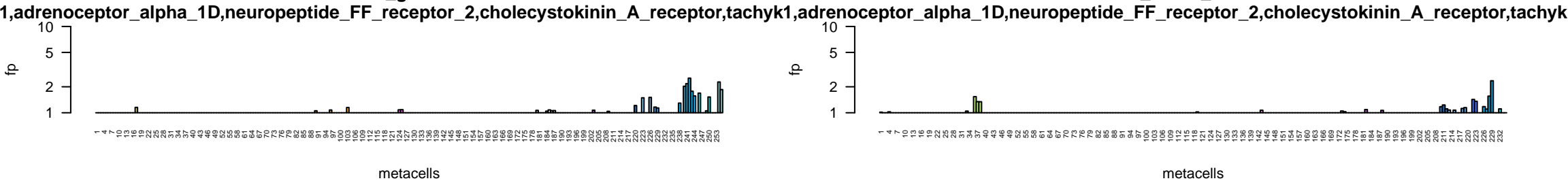
A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 250. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that for most metacells, the frequency is 1. There are a few metacells with a frequency of 2 (metacells 19, 89, 94), and a small cluster of metacells with frequencies of 3 and 4 (metacells 241, 242, 243, 244, 245, 246, 247). The highest frequency is 4, which occurs for metacells 241, 242, 243, and 244.

metacells	fp
1	1
7	1
10	1
16	1
18	1
19	2
25	1
28	1
34	1
37	1
43	1
46	1
52	1
55	1
61	1
64	1
67	1
70	1
76	1
79	1
85	1
89	2
94	2
103	1
106	1
112	1
115	1
121	1
122	1
127	1
130	1
133	1
136	1
140	1
145	1
148	1
154	1
157	1
163	1
166	1
172	1
175	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
205	1
208	1
214	1
217	1
223	1
226	1
232	1
235	1
236	1
241	4
242	3
243	3
244	3
245	3
246	3
247	3
250	1
253	1
254	1

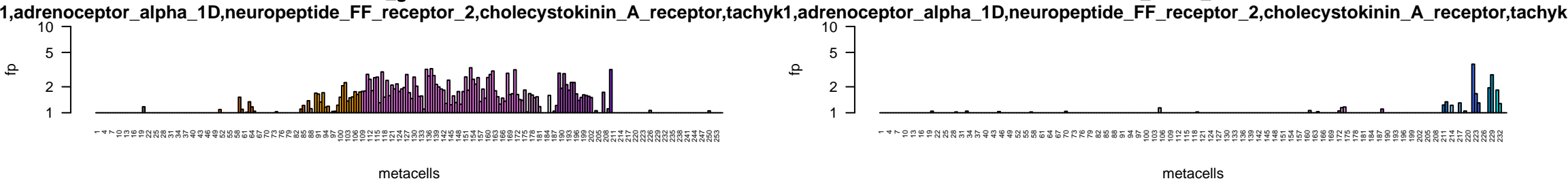
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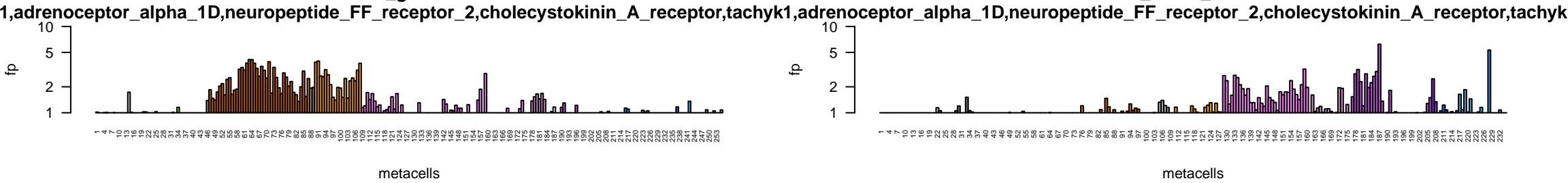
**Hhon q06114.t1**



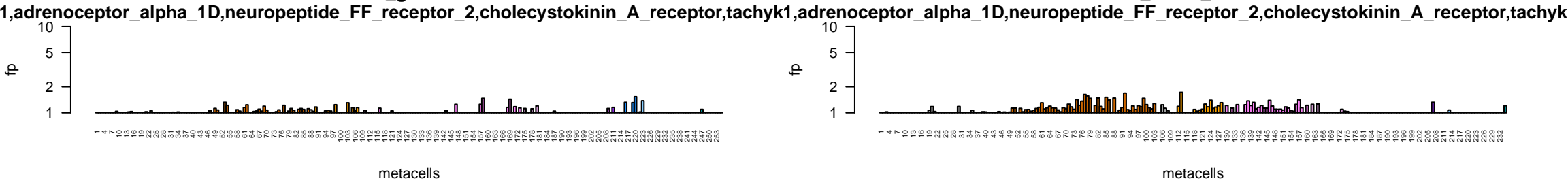
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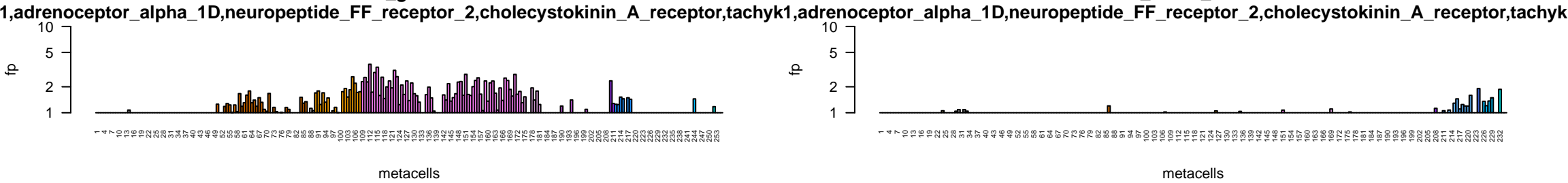
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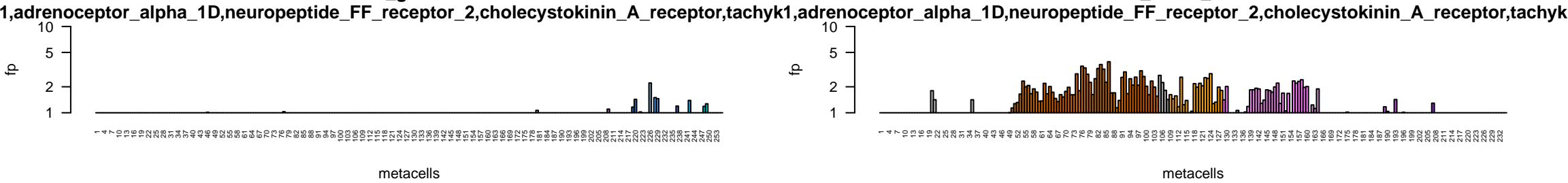
**Hhon q07529.t1**



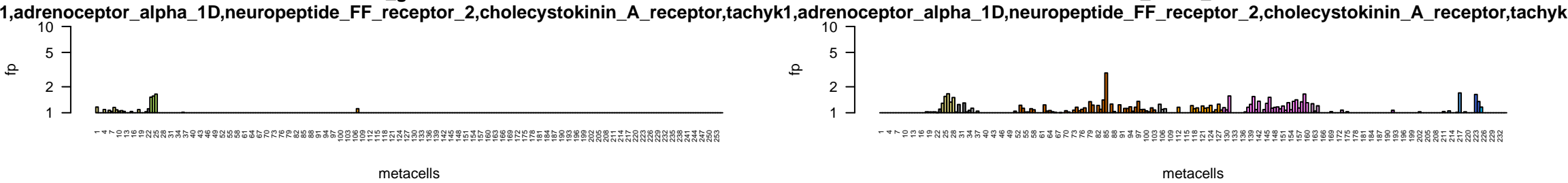
**Hhon q07527.t1**

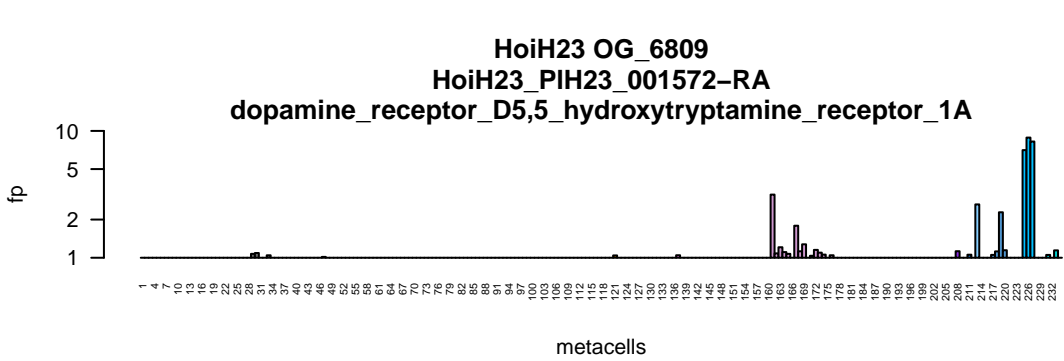
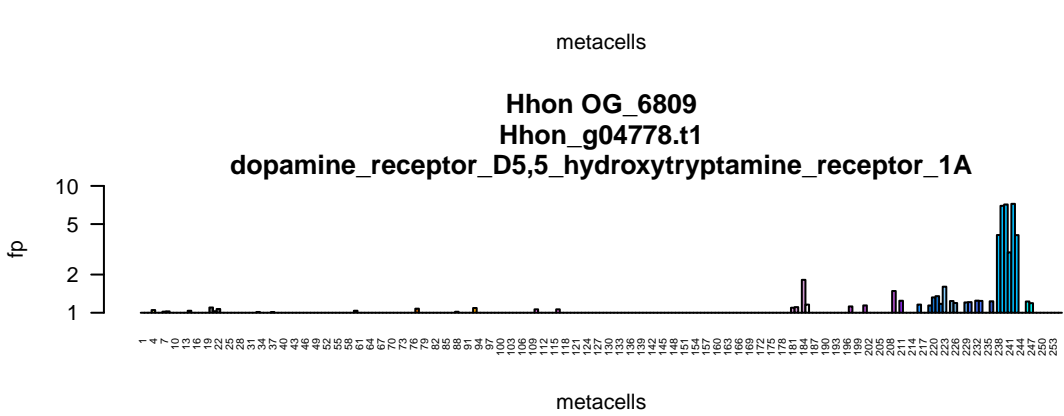
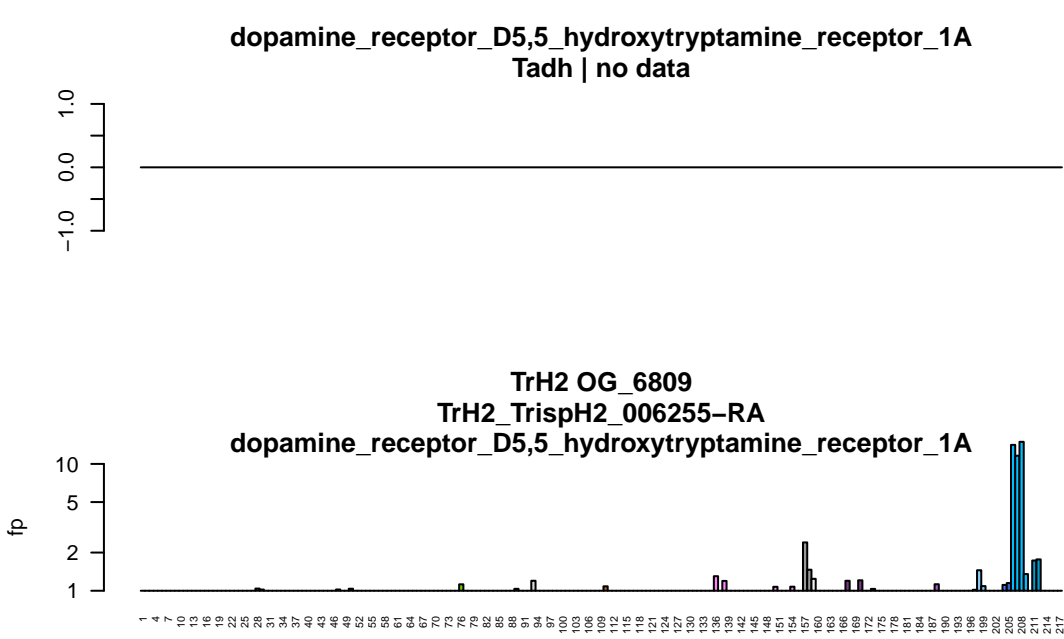


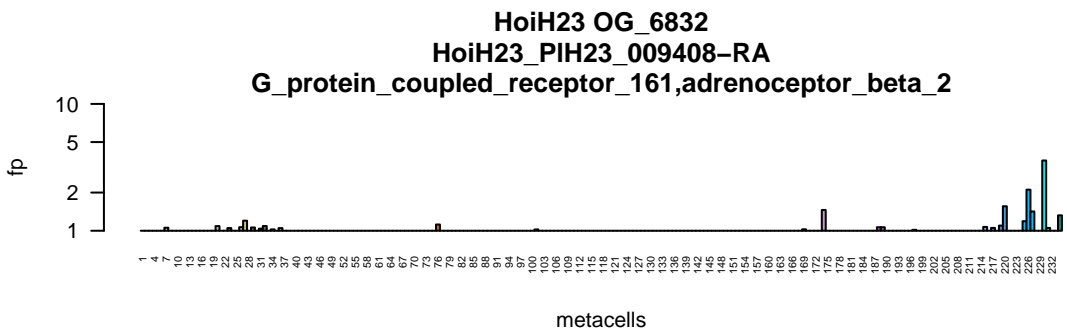
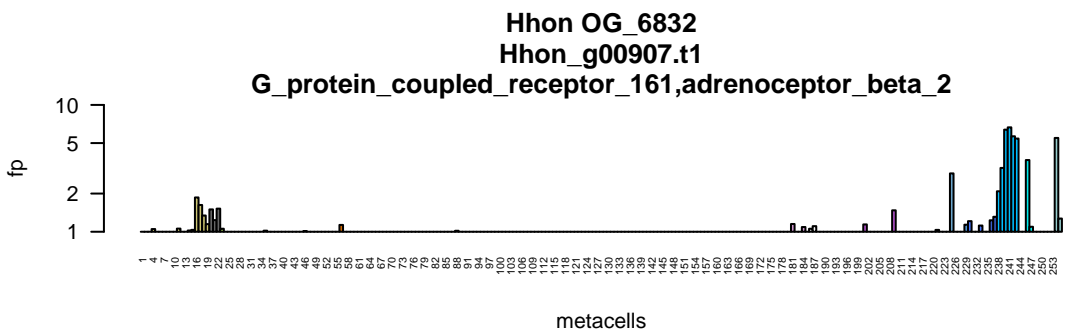
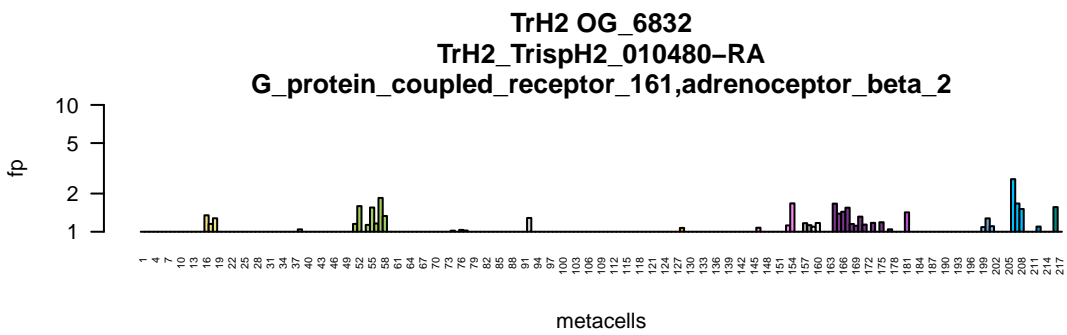
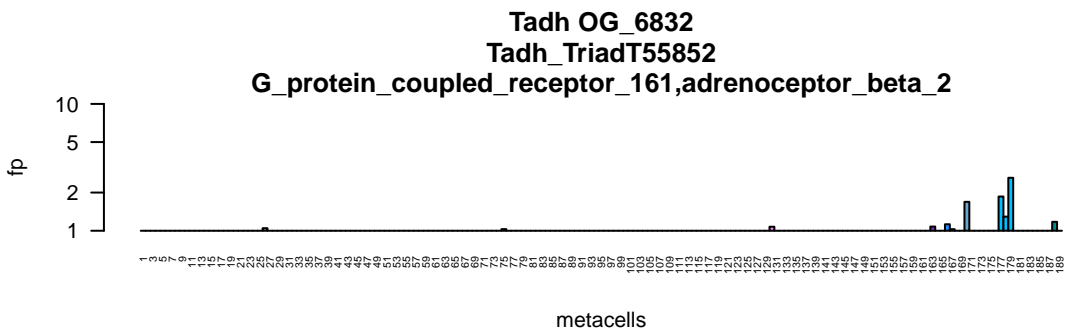
**Hhon q10923.t1**

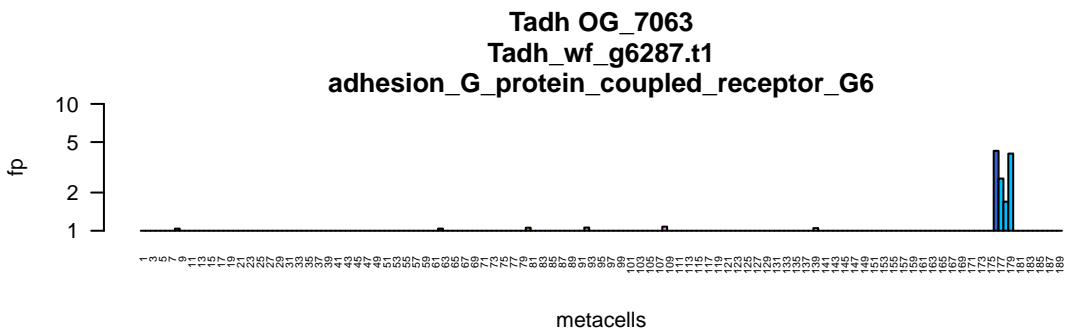


**Hhon q08418.t1**

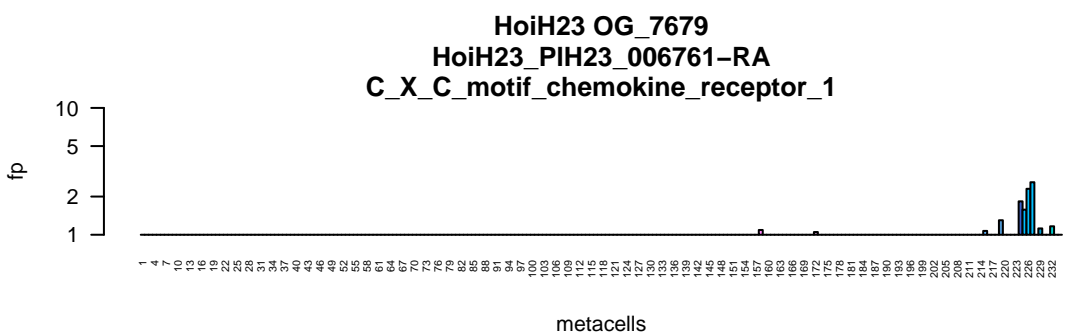
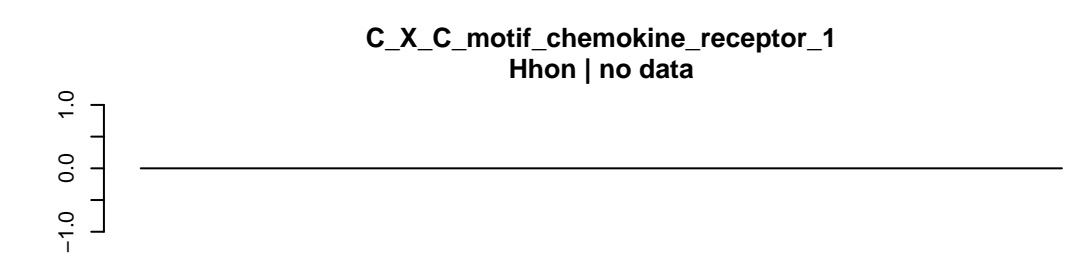
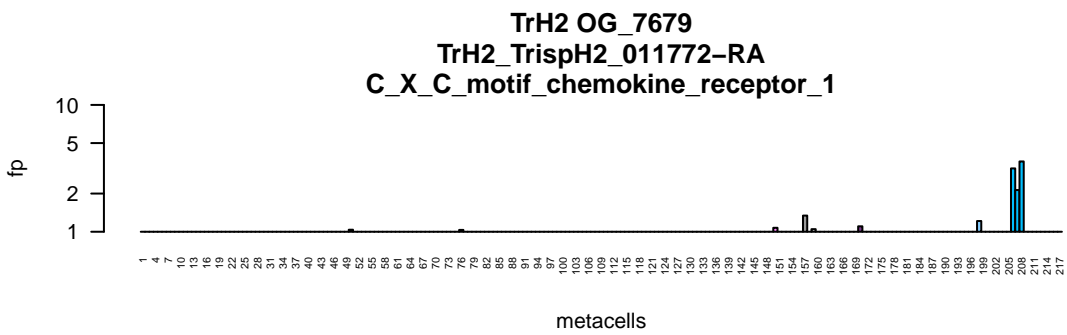
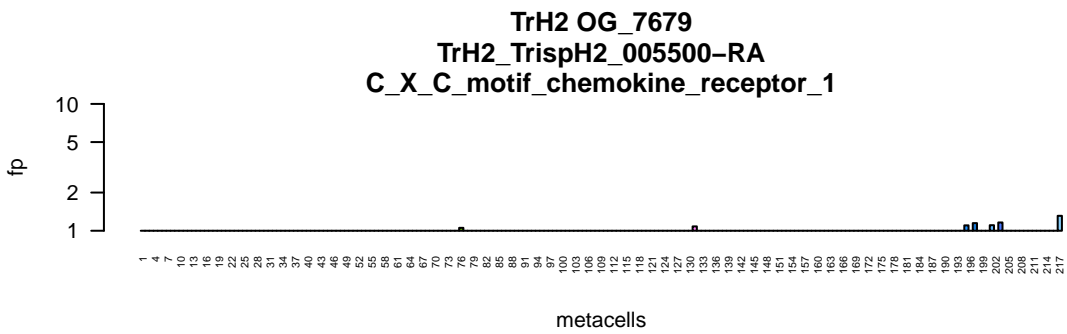
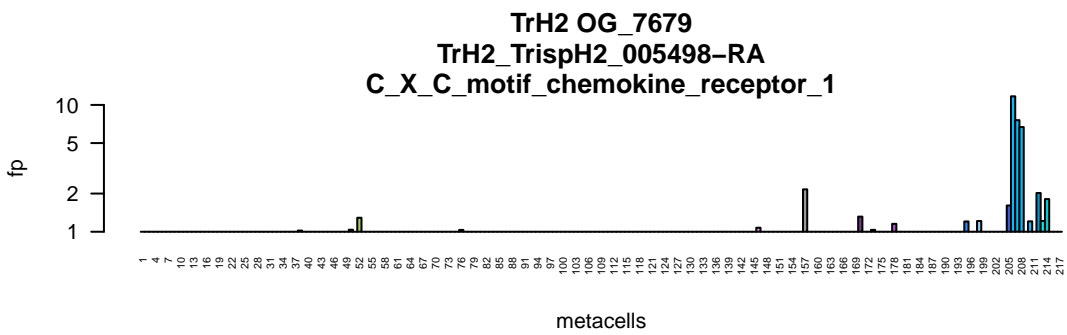
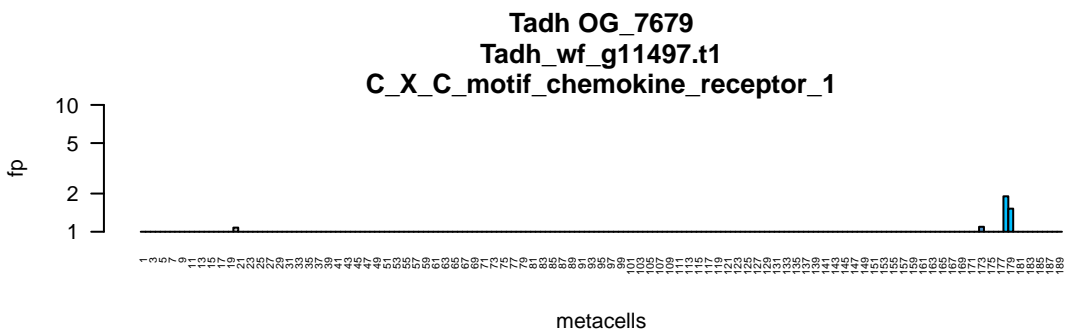
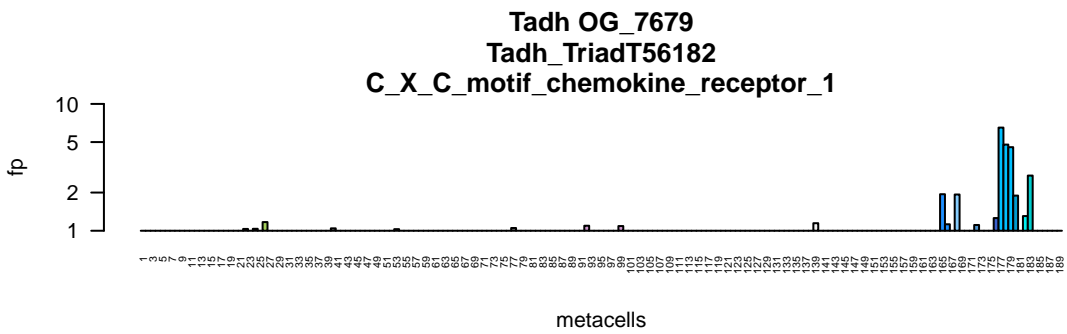


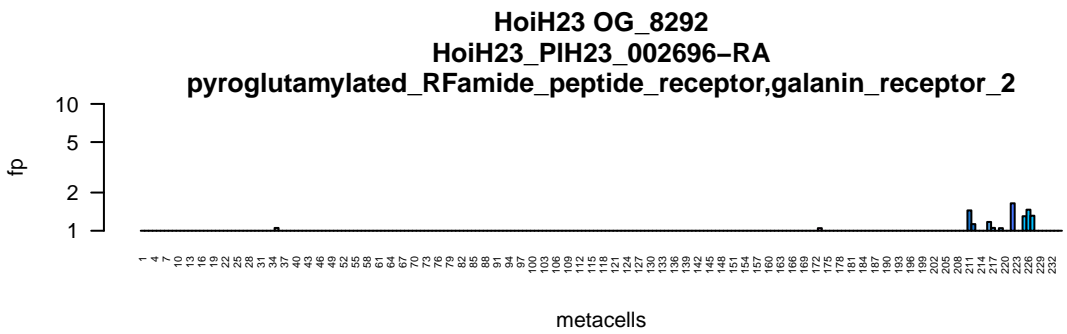
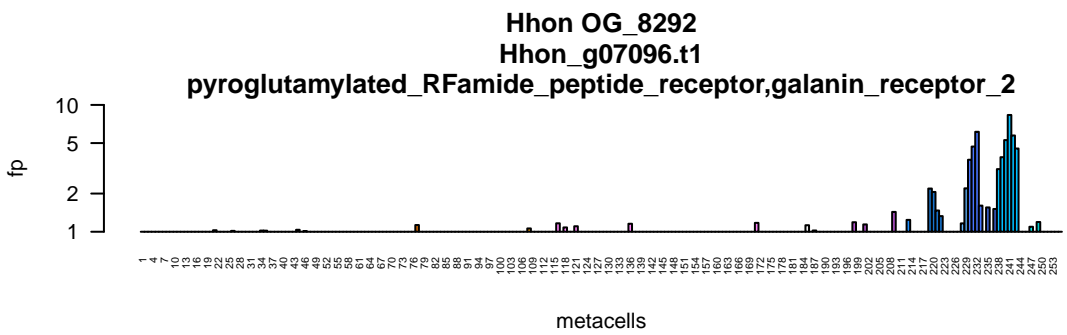
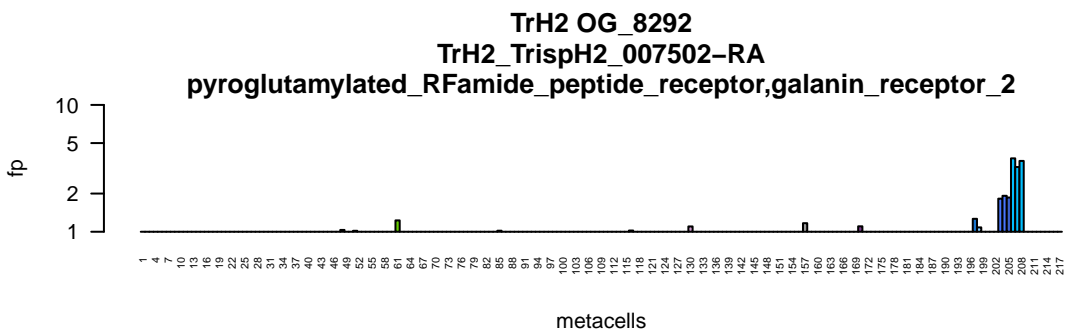
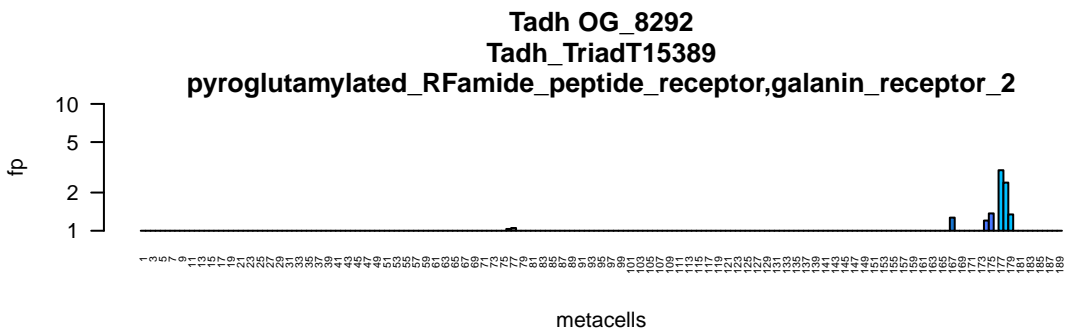


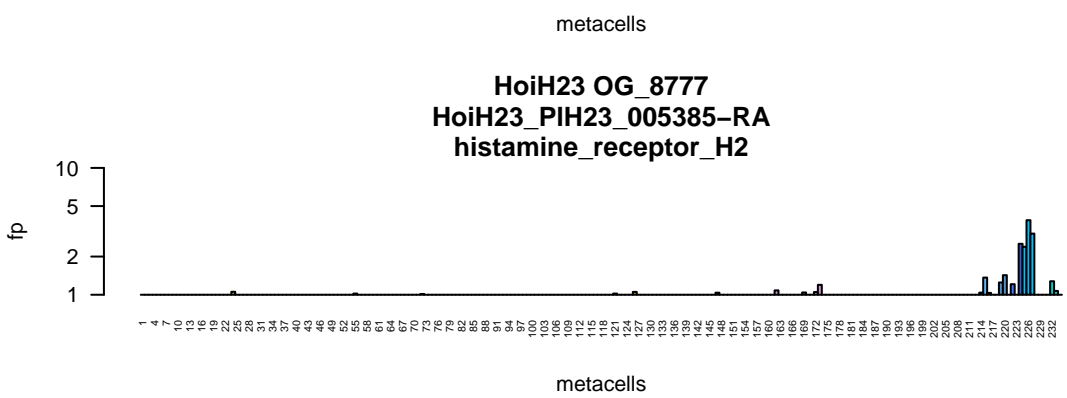
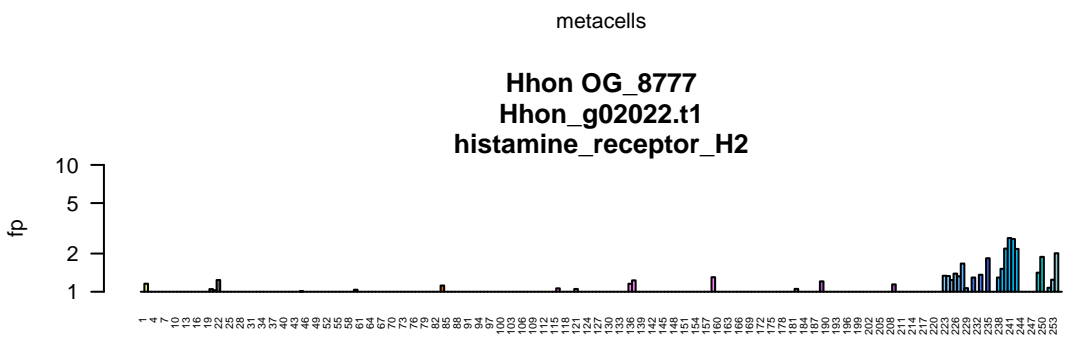
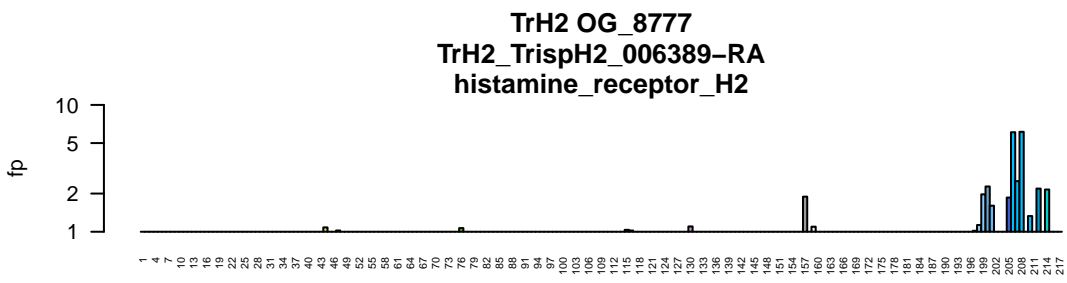
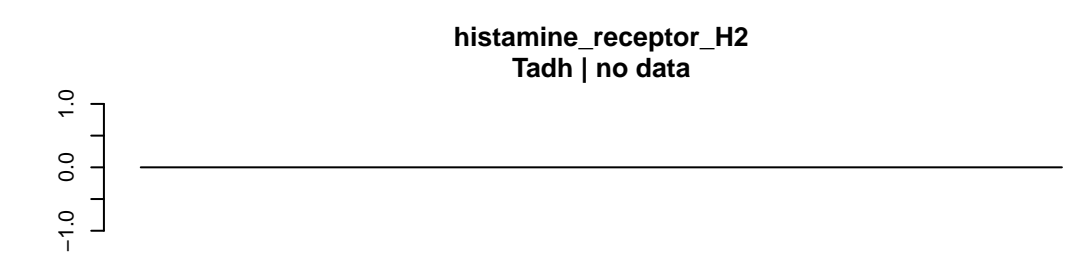




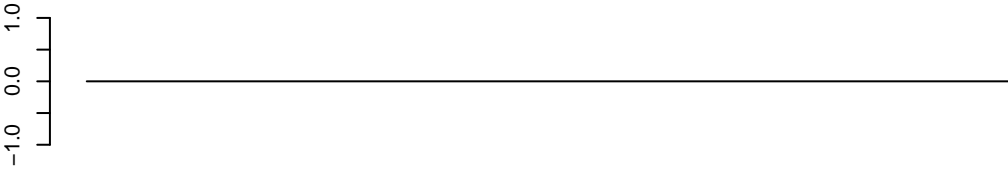




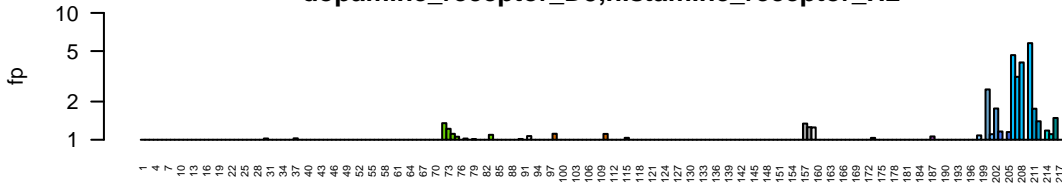




dopamine\_receptor\_D5,histamine\_receptor\_H2  
Tadh | no data

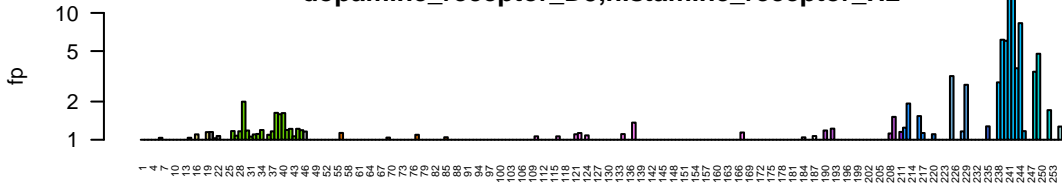


TrH2 OG\_8890  
TrH2\_TrispH2\_010269-RA  
dopamine\_receptor\_D5,histamine\_receptor\_H2



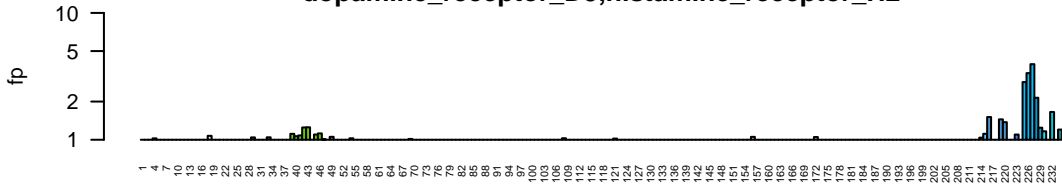
metacells

Hhon OG\_8890  
Hhon\_g10763.t1  
dopamine\_receptor\_D5,histamine\_receptor\_H2

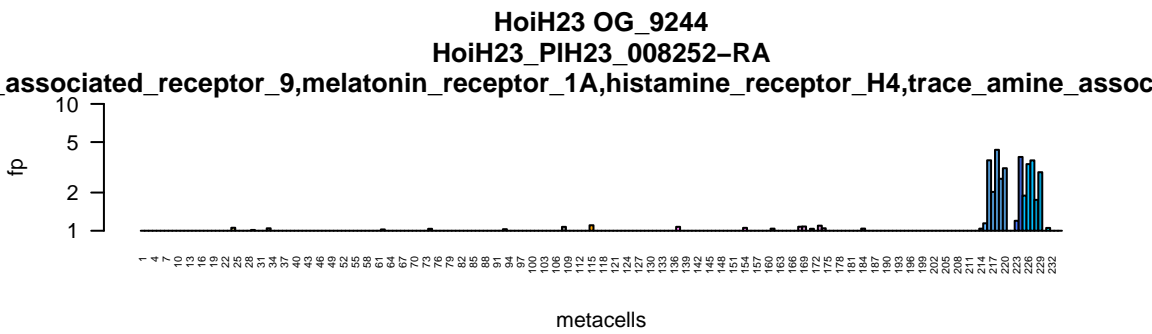
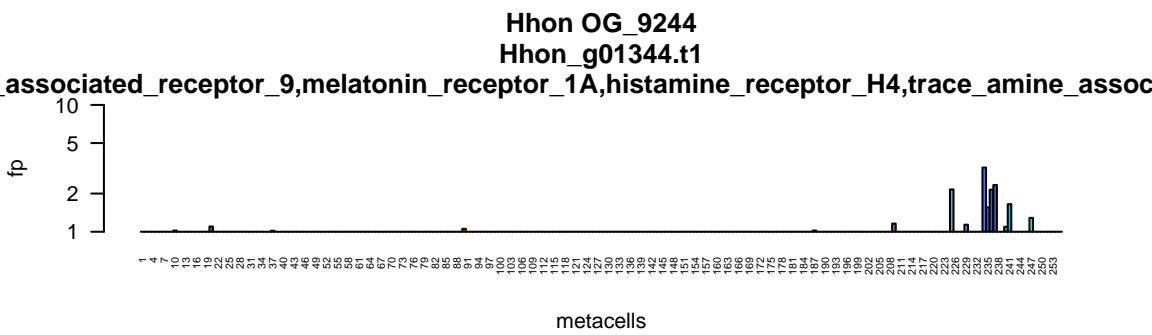
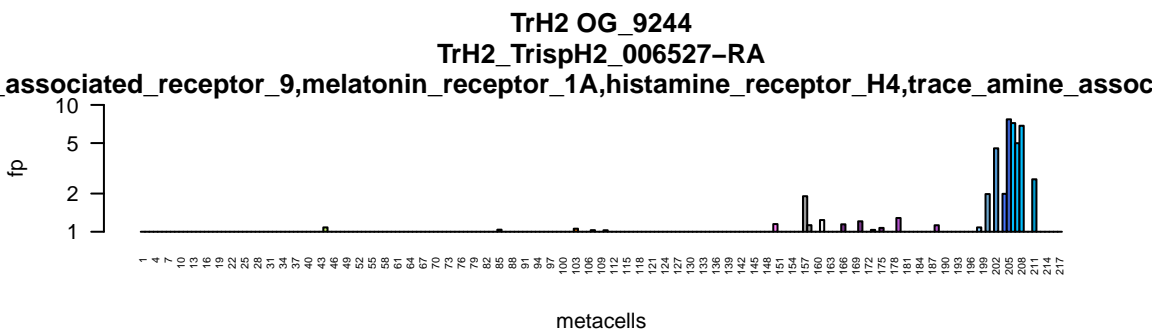
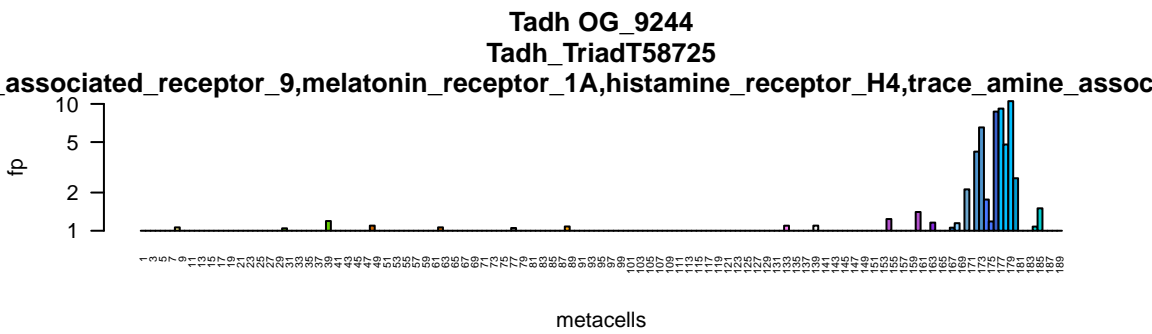


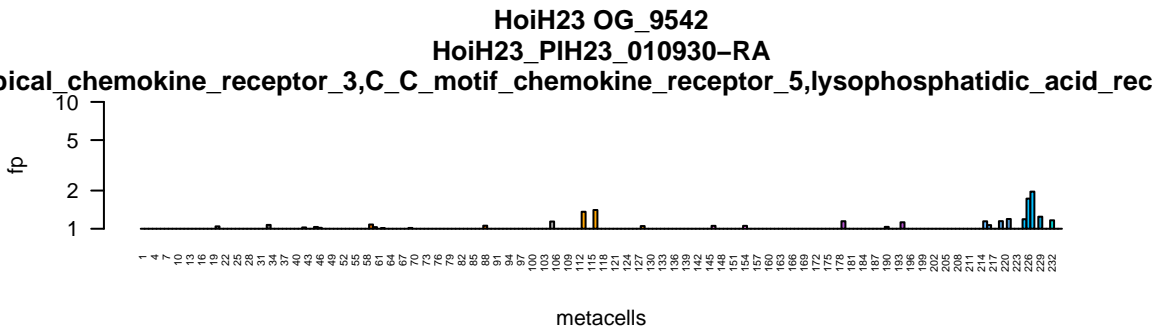
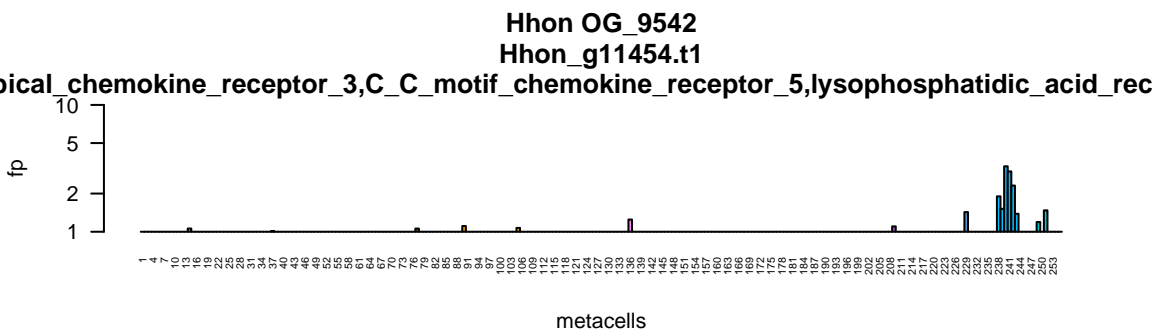
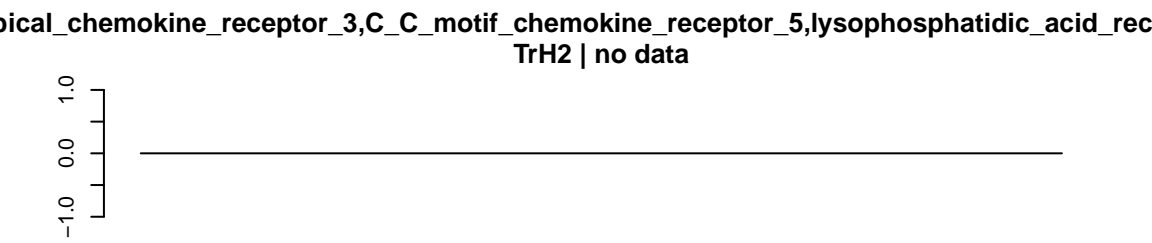
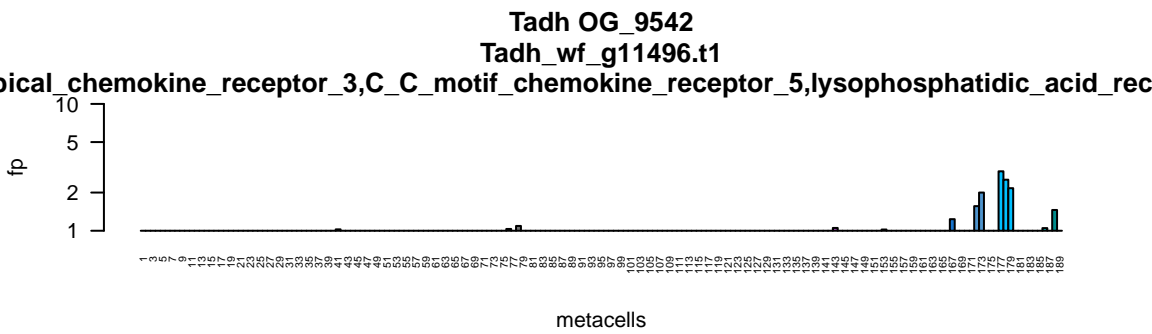
metacells

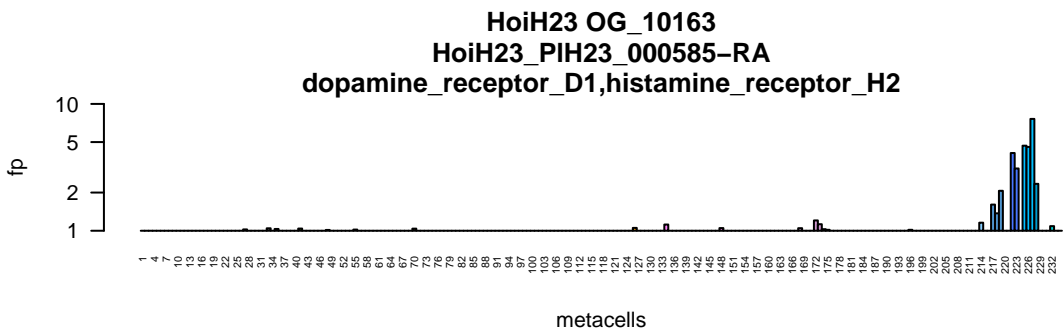
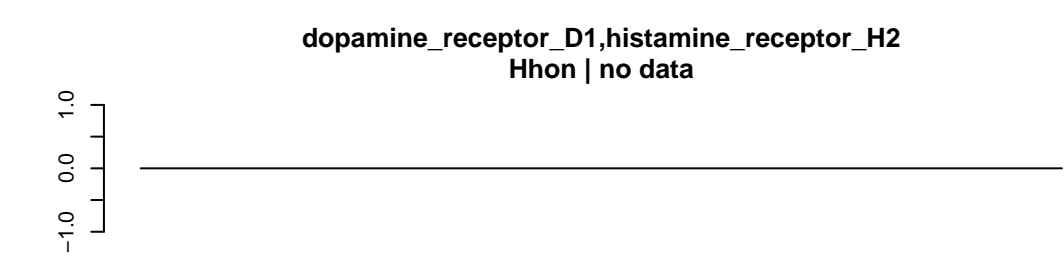
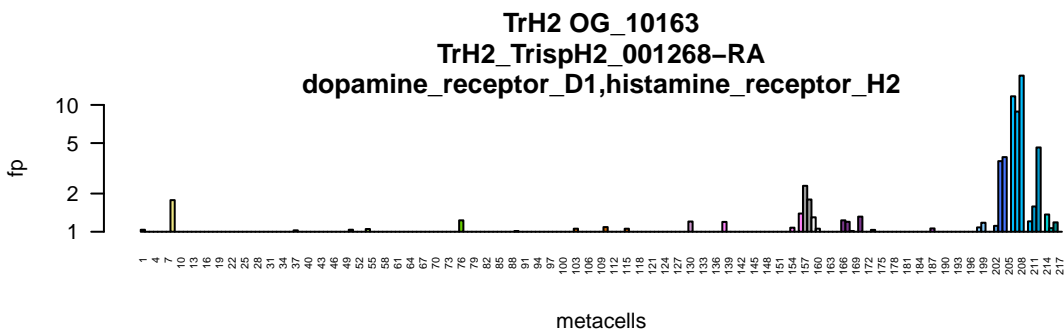
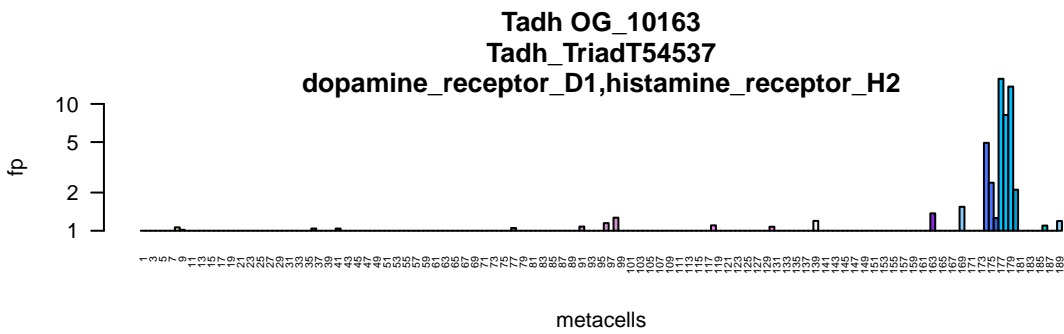
HoiH23 OG\_8890  
HoiH23\_PIH23\_008421-RA  
dopamine\_receptor\_D5,histamine\_receptor\_H2



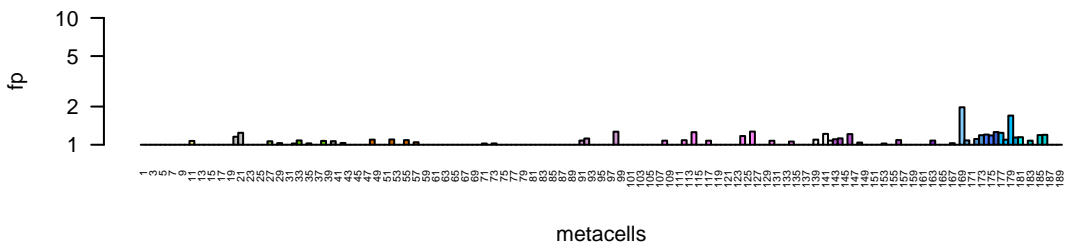
metacells



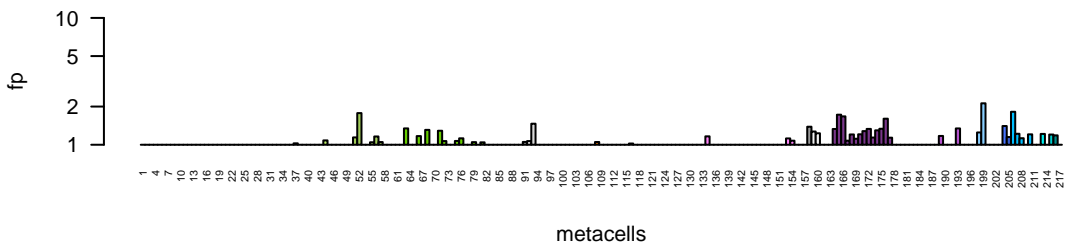




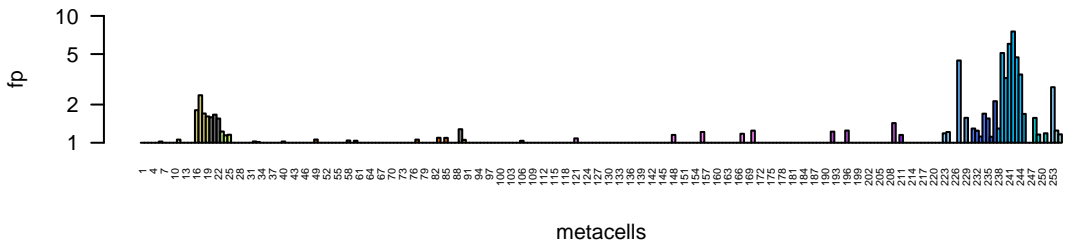
Tadh OG\_1881  
Tadh\_wf\_g2794.t1



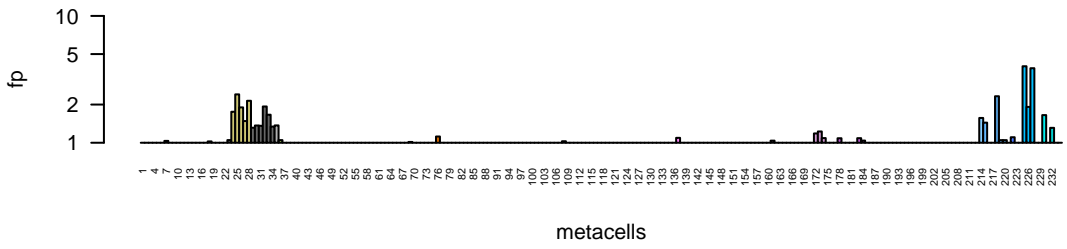
TrH2 OG\_1881  
TrH2\_TrispH2\_007804-RA



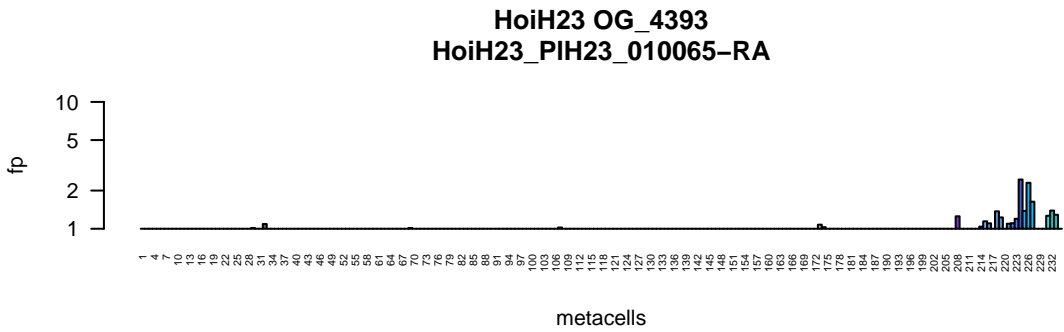
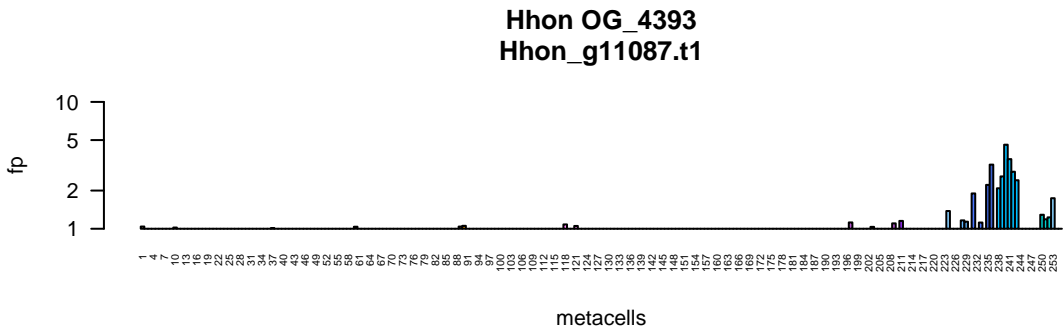
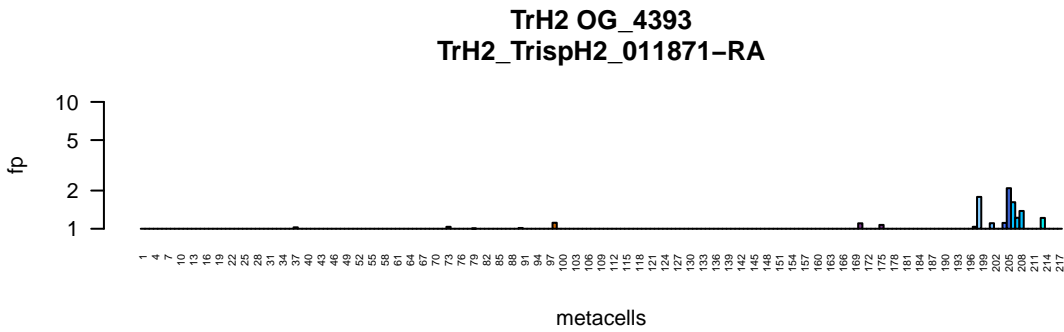
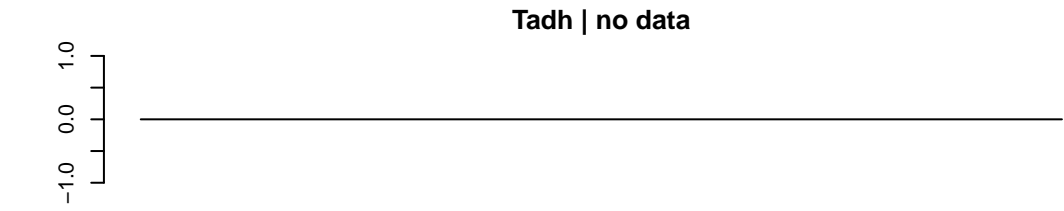
Hhon OG\_1881  
Hhon\_g06778.t1

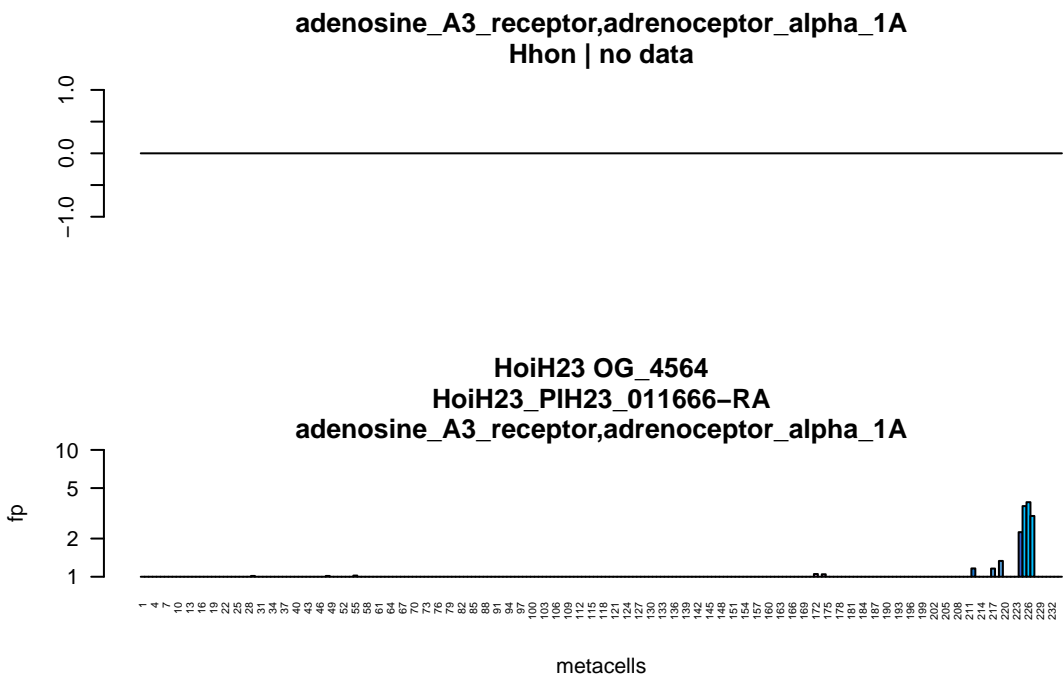
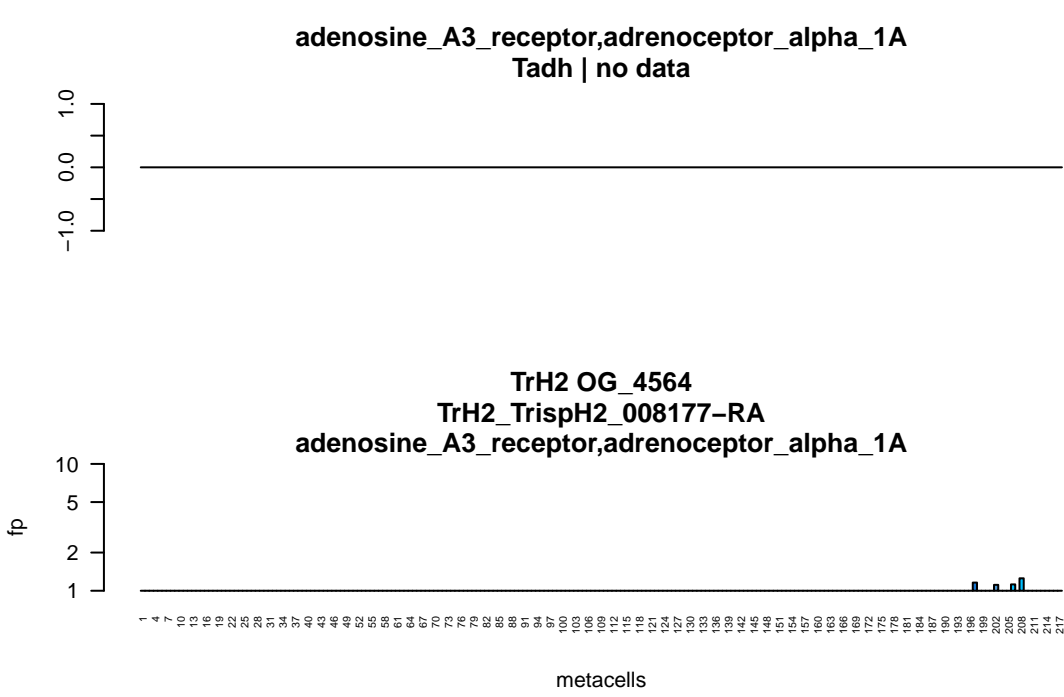


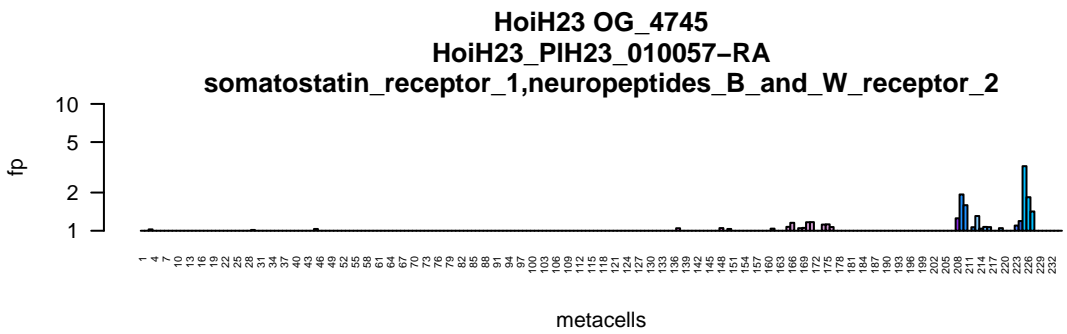
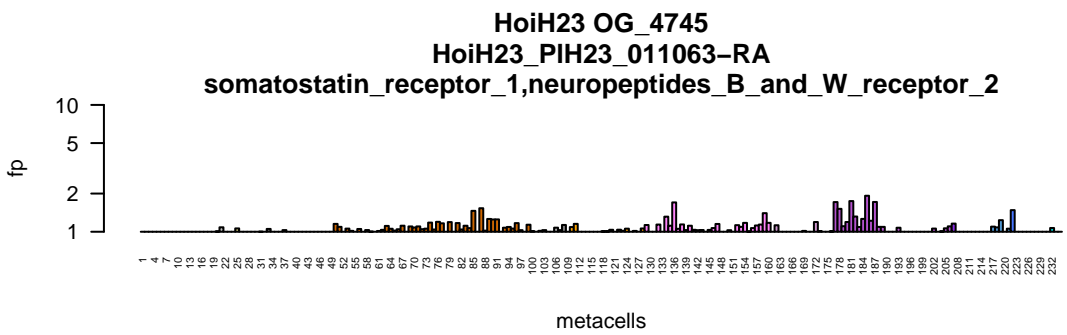
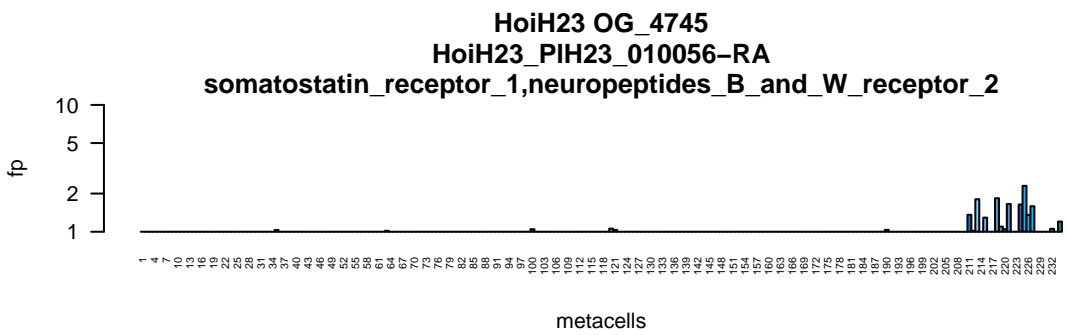
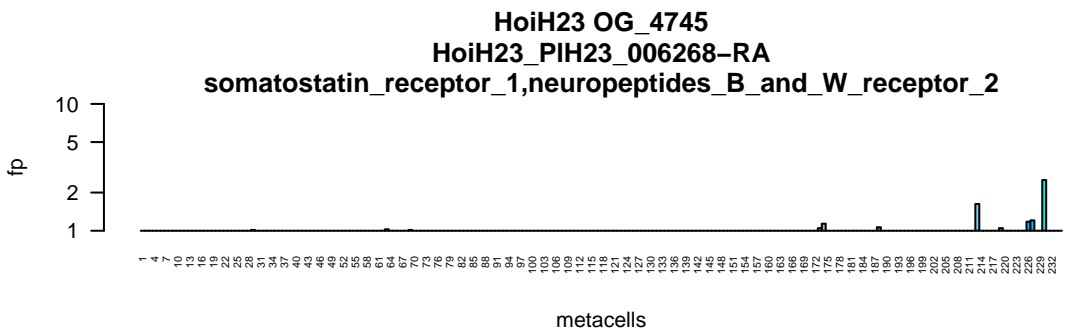
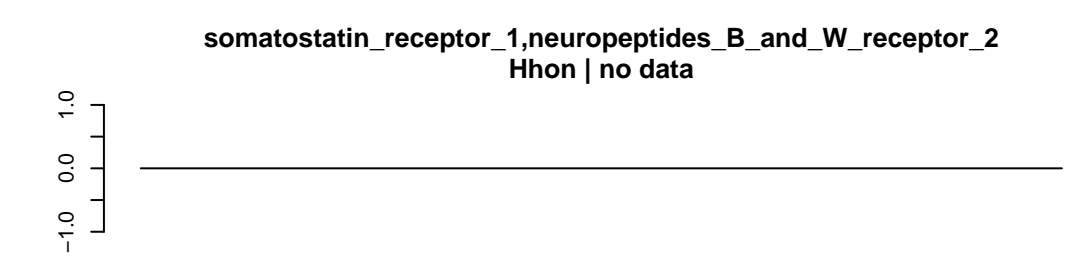
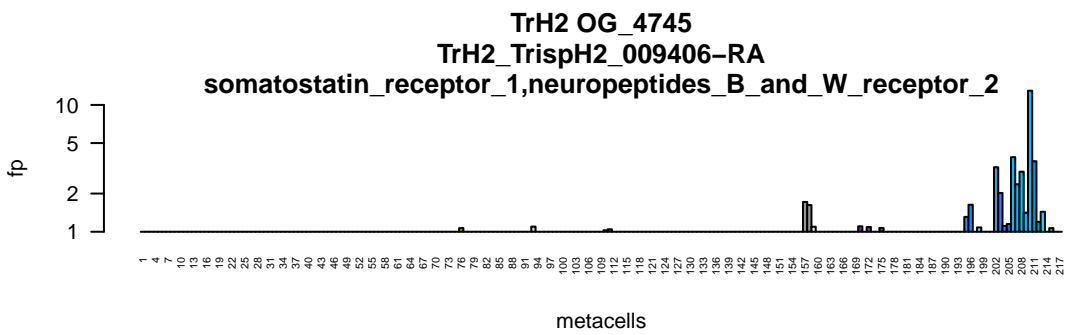
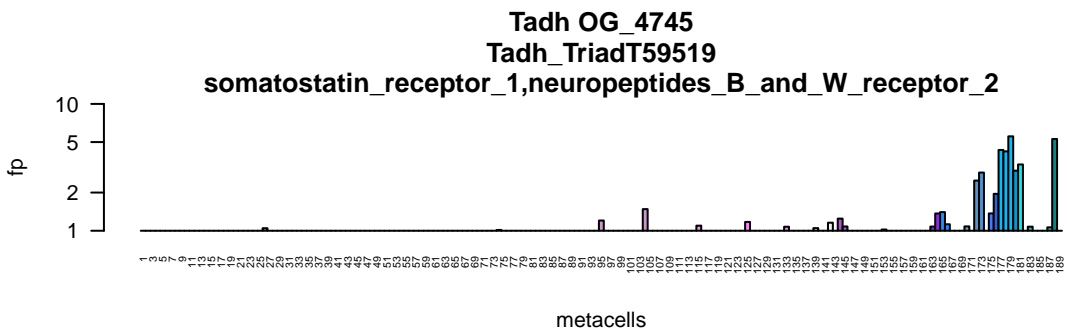
HoiH23 OG\_1881  
HoiH23\_PIH23\_005068-RA

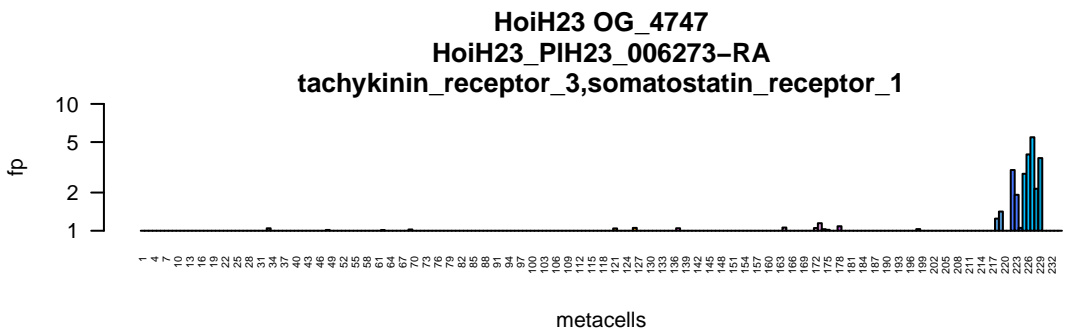
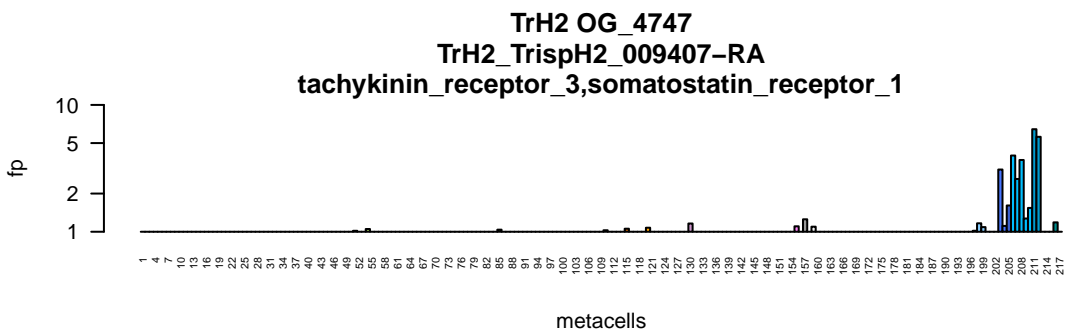
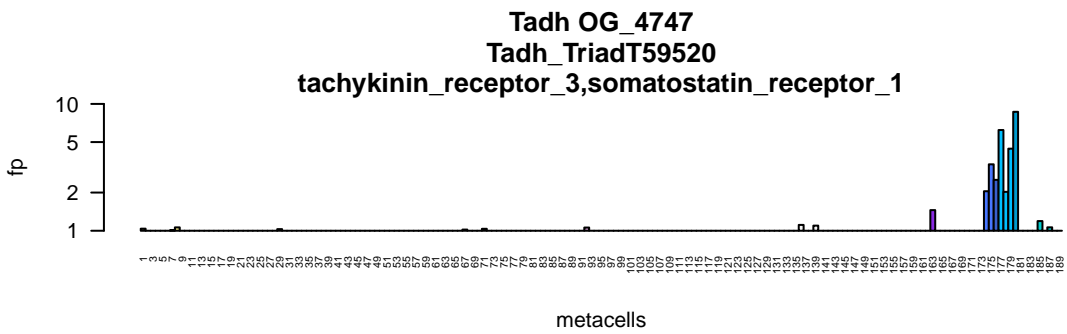


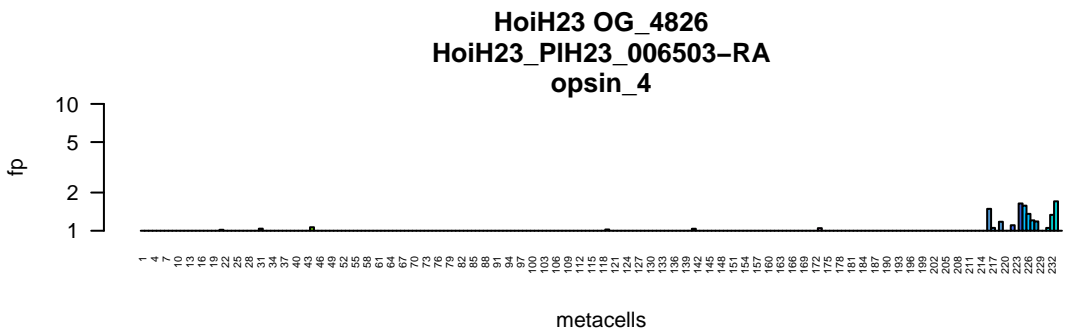
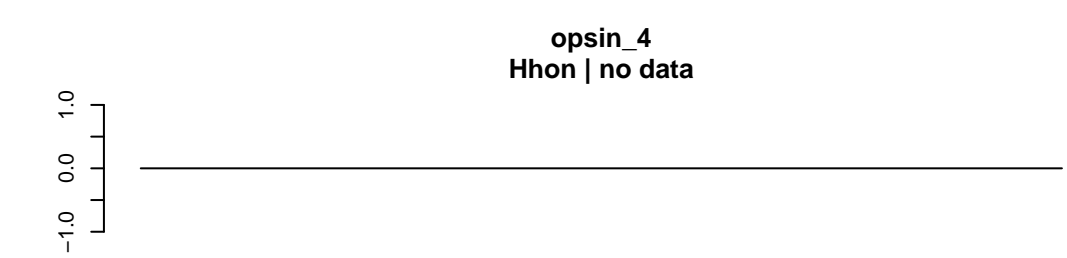
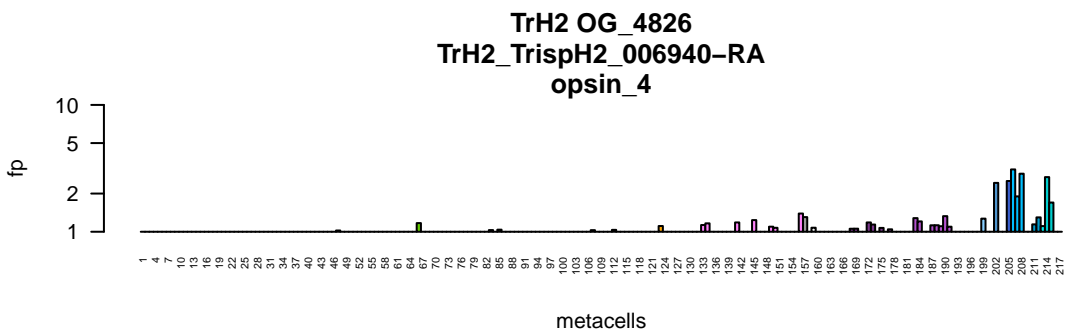
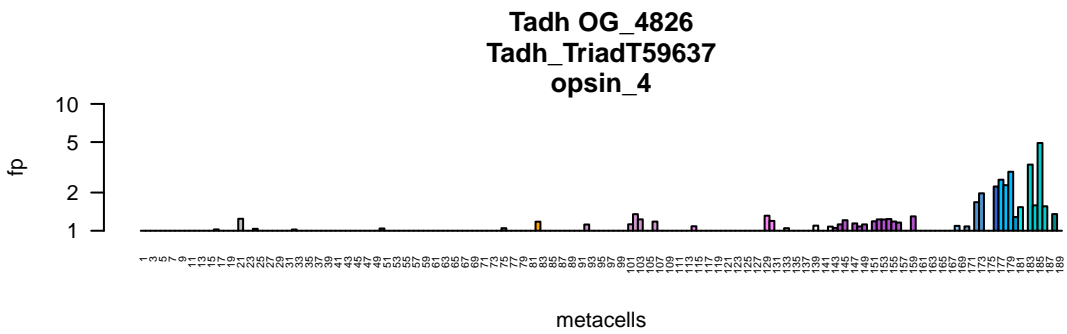


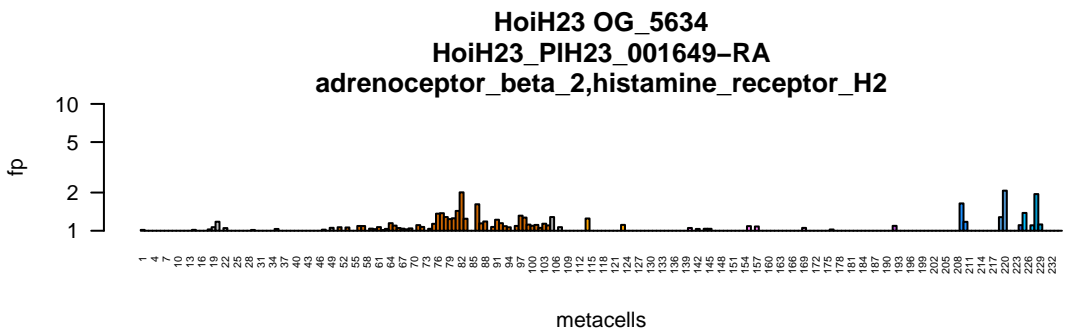
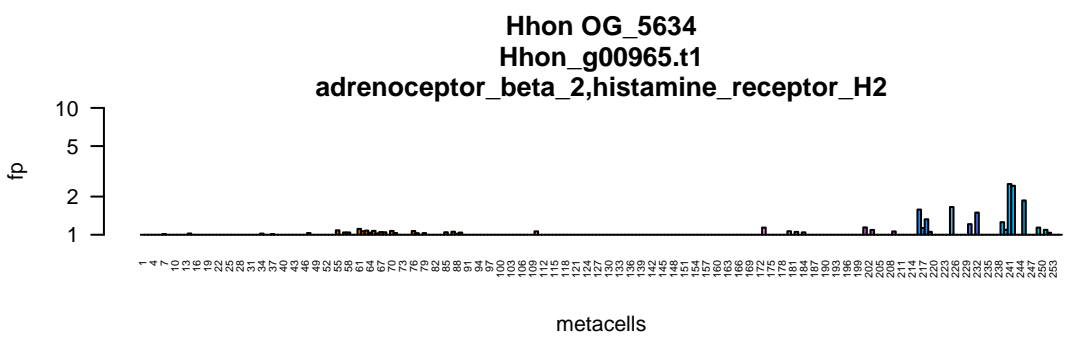
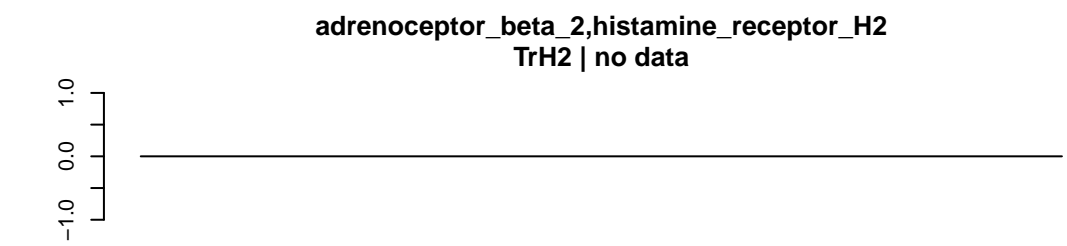
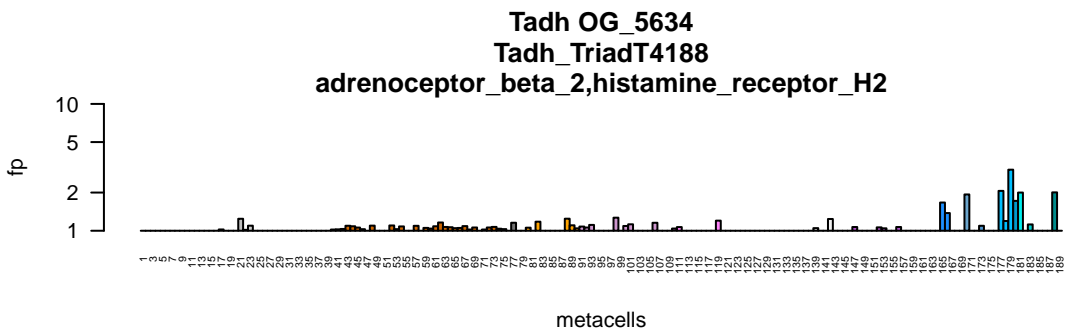


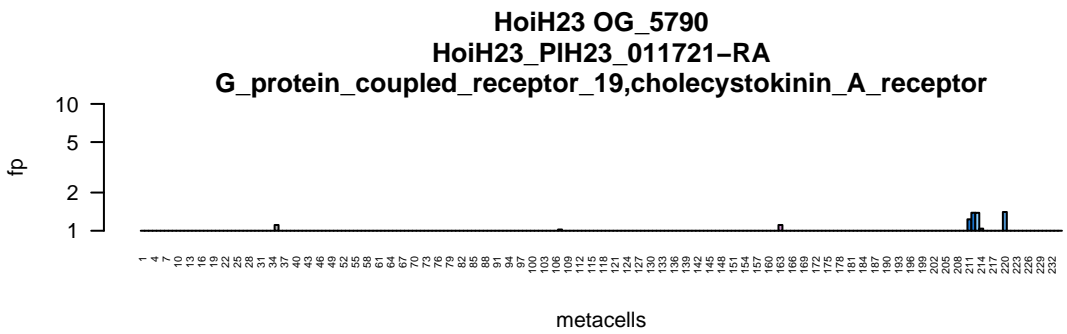
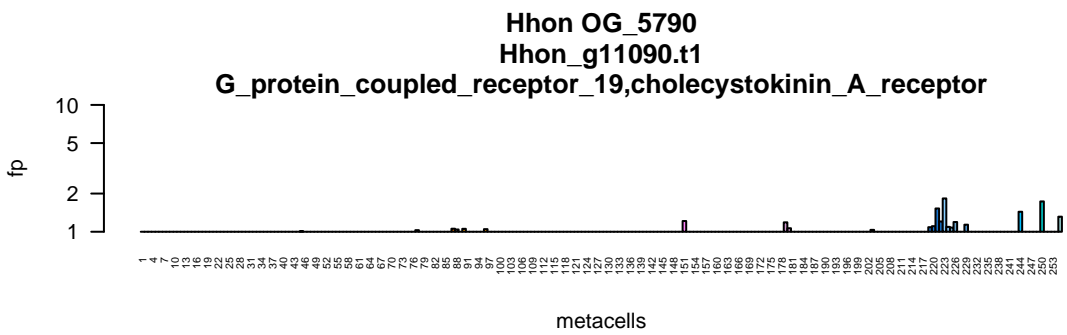
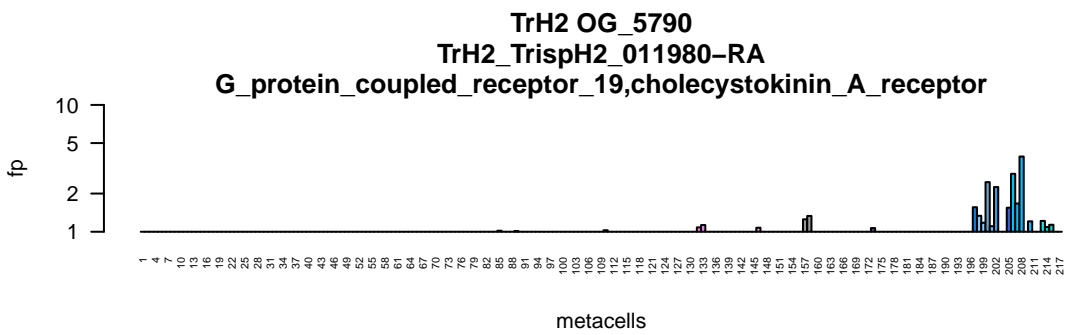
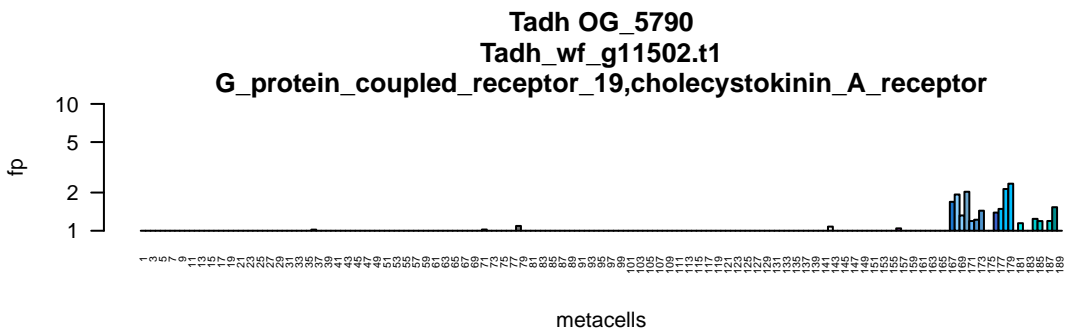




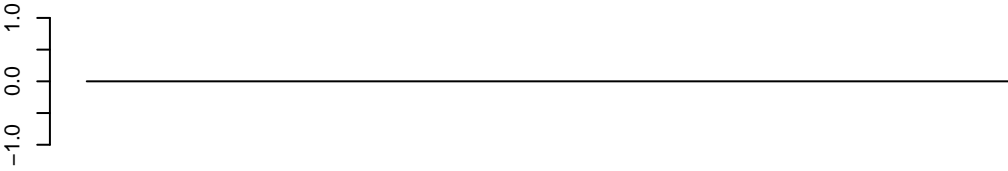




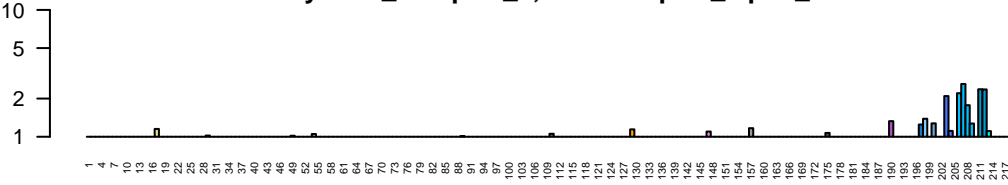




tachykinin\_receptor\_1,adrenoceptor\_alpha\_1B  
Tadh | no data

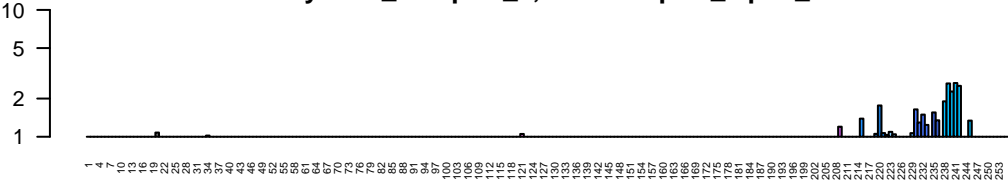


TrH2 OG\_5841  
TrH2\_TrispH2\_002792-RA  
tachykinin\_receptor\_1,adrenoceptor\_alpha\_1B



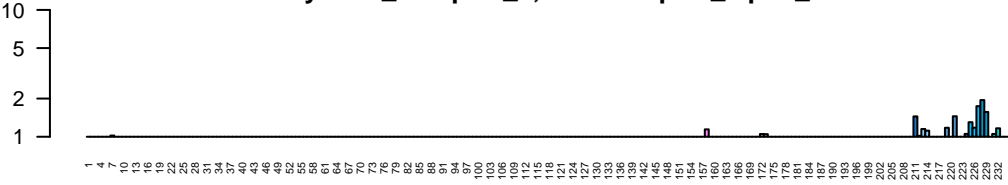
metacells

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



metacells

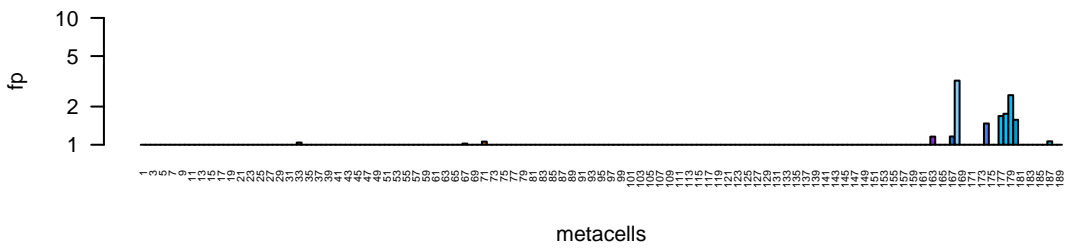
HoiH23 OG\_5841  
HoiH23\_PIH23\_001771-RA  
tachykinin\_receptor\_1,adrenoceptor\_alpha\_1B



metacells

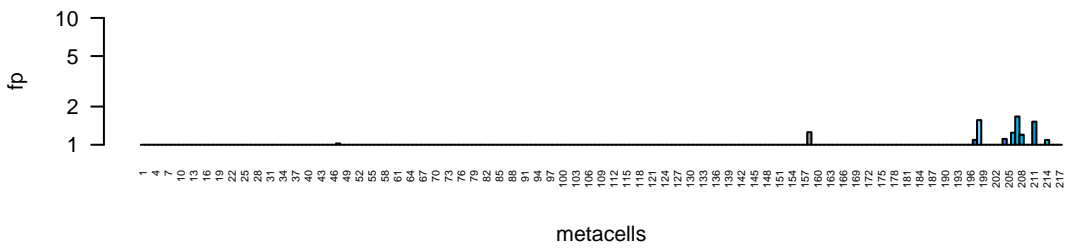


Tadh OG\_6120  
Tadh\_TriadT60881



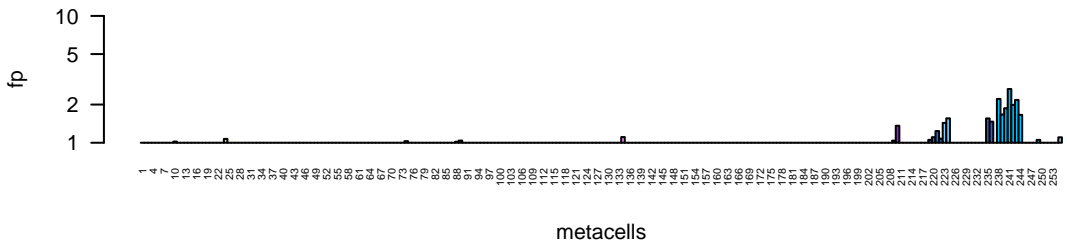
metacells

TrH2 OG\_6120  
TrH2\_TrispH2\_008845-RA



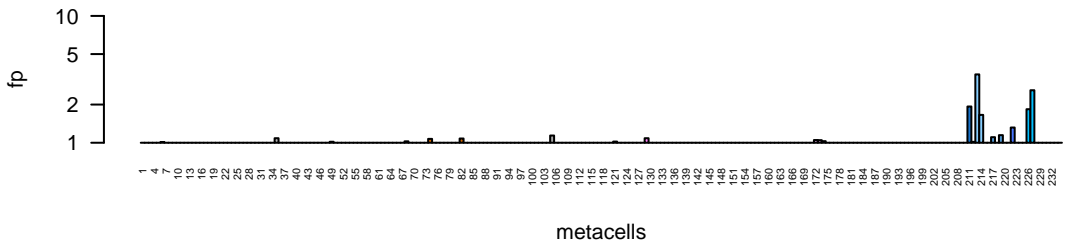
metacells

Hhon OG\_6120  
Hhon\_g10766.t1

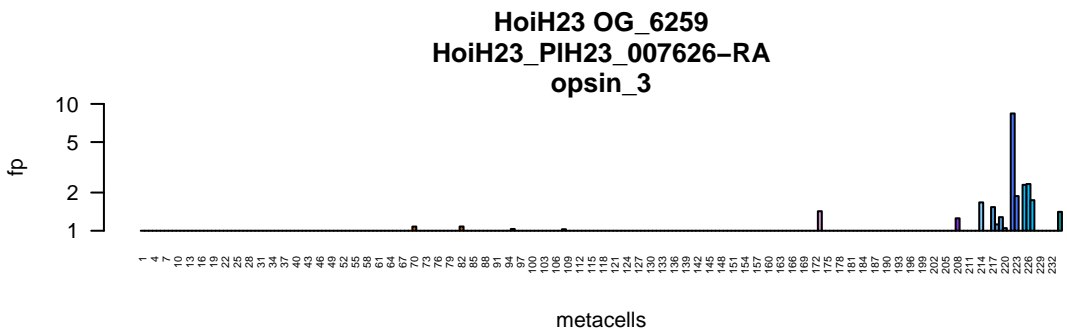
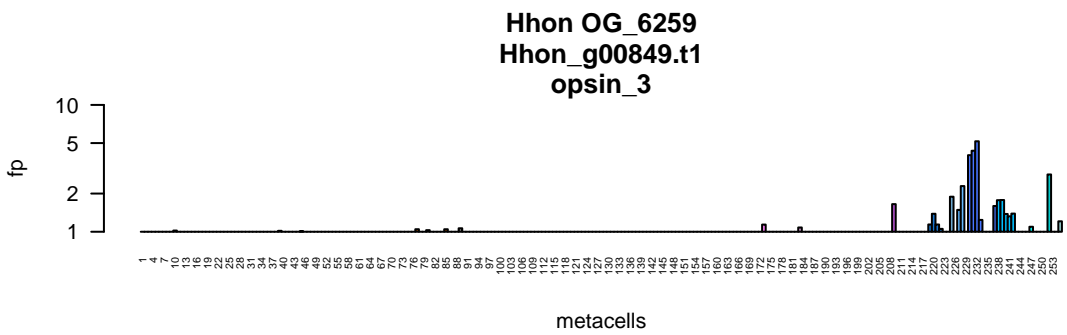
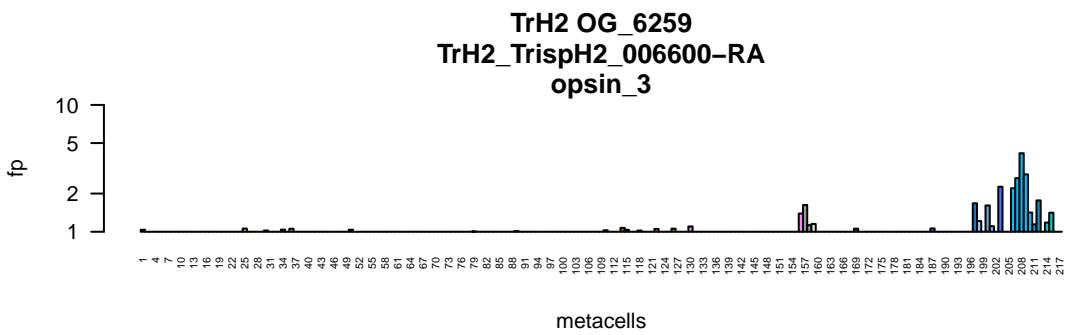
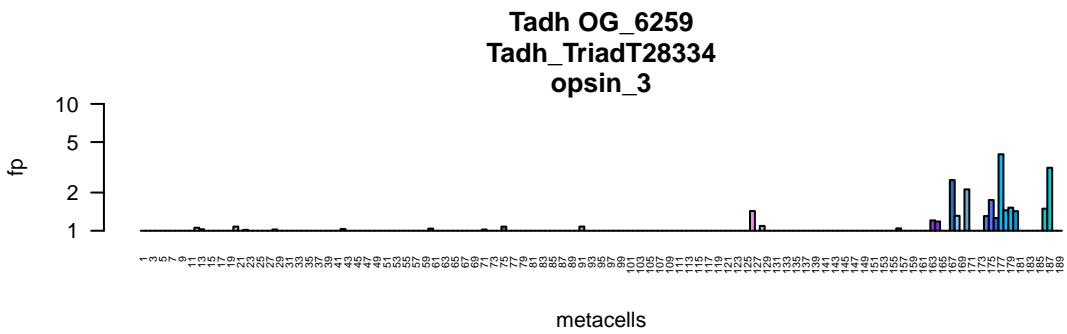


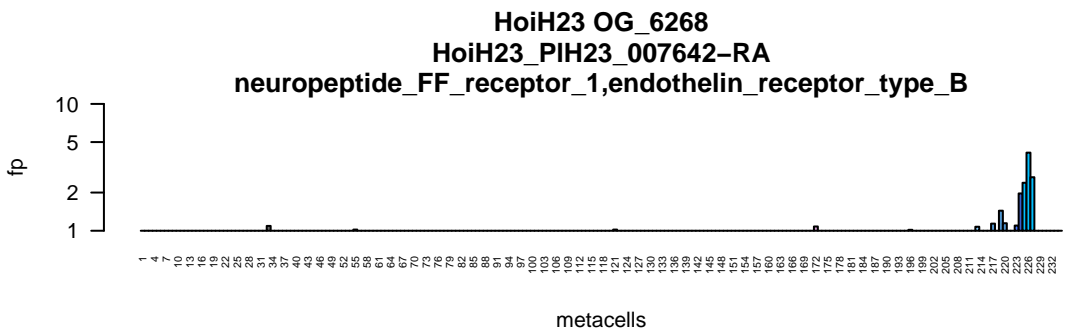
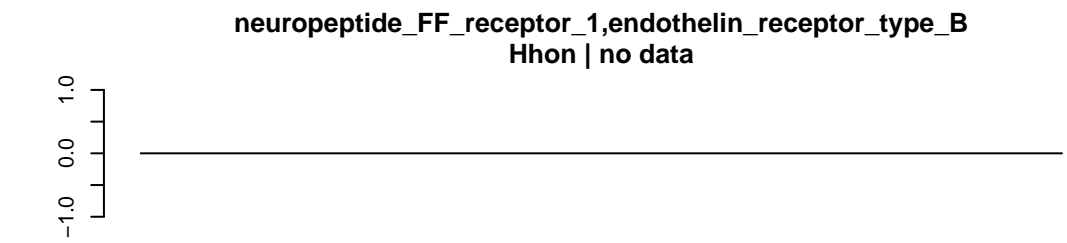
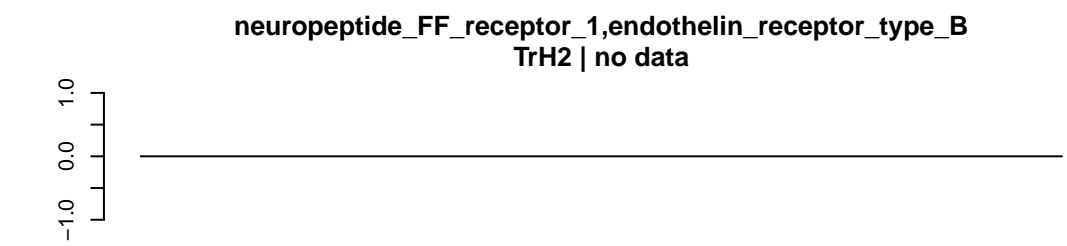
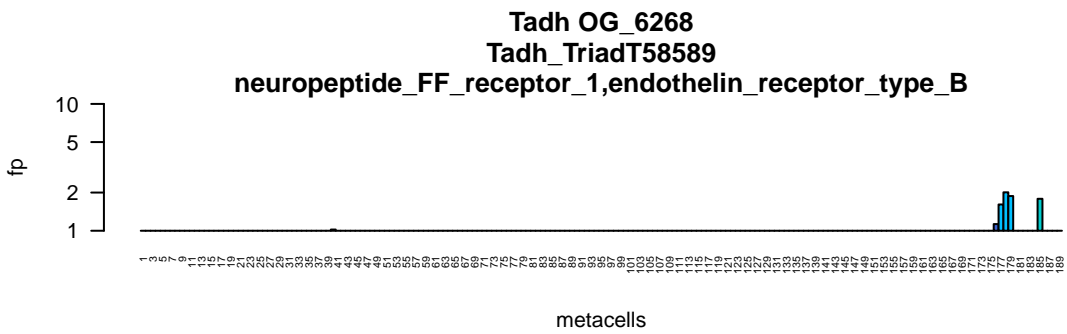
metacells

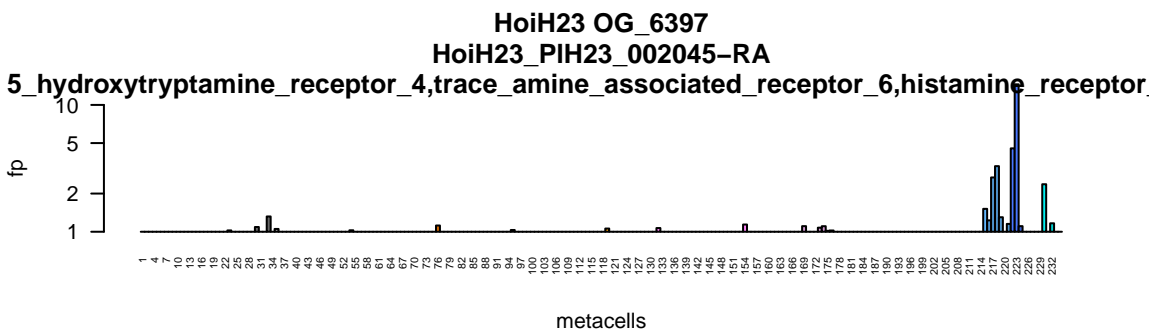
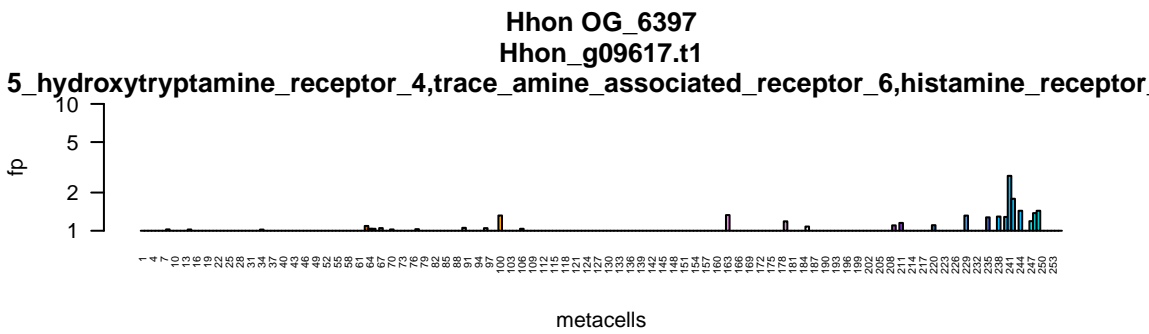
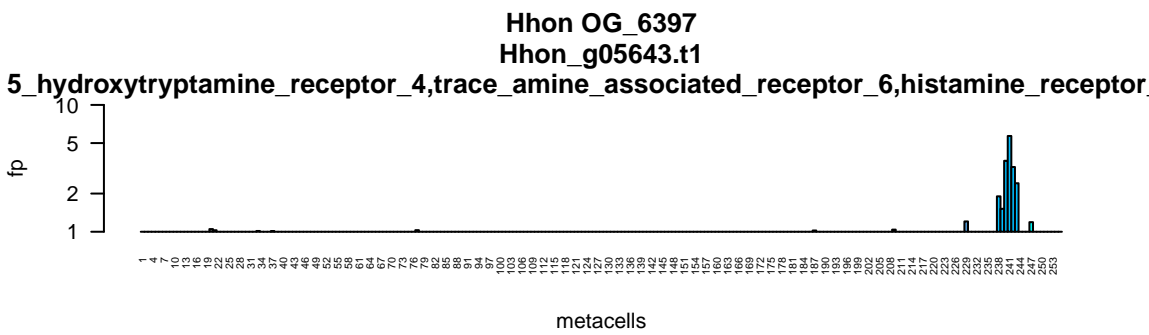
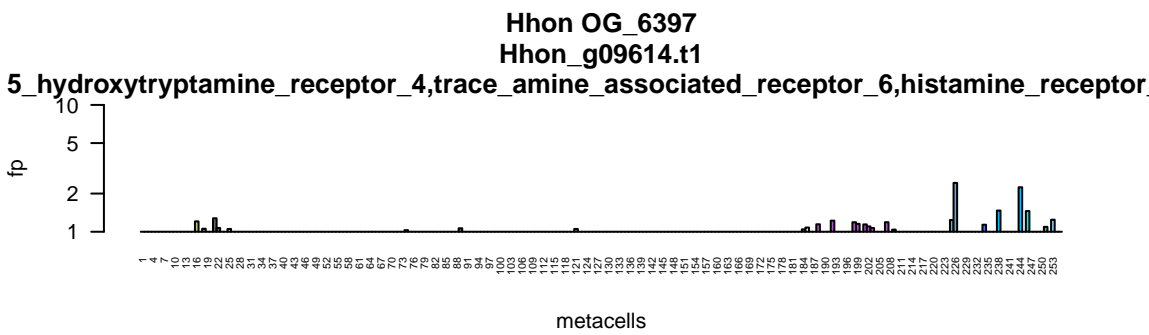
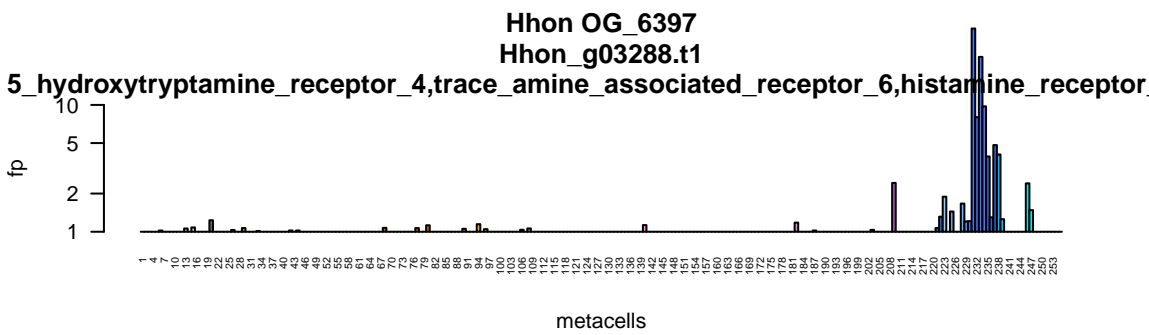
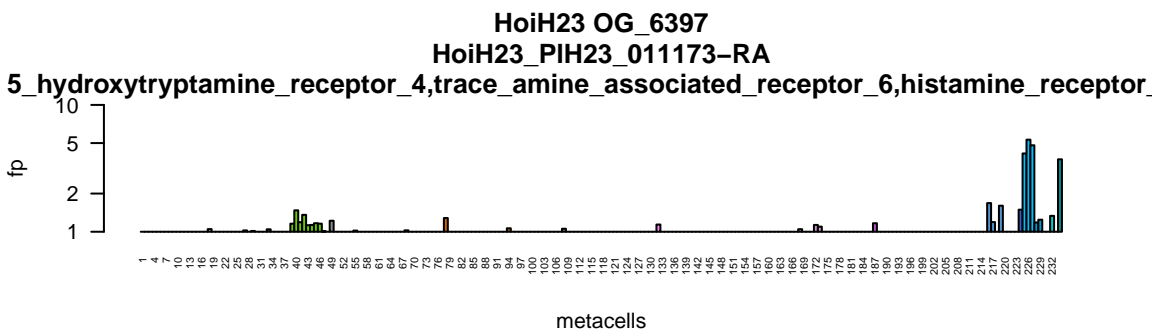
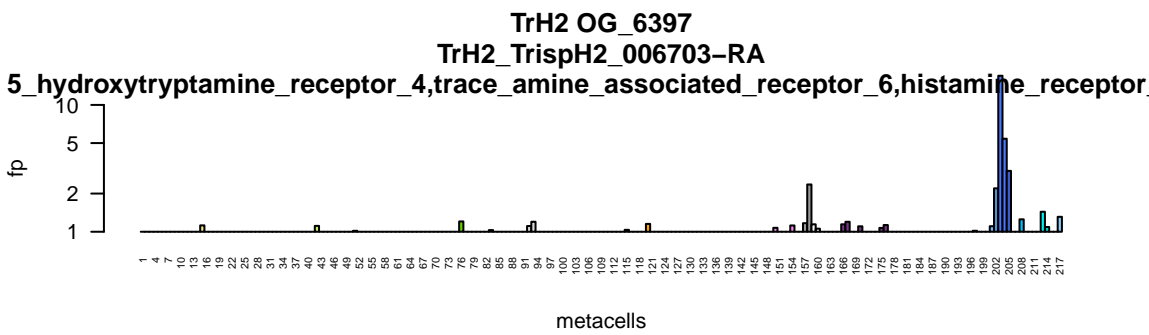
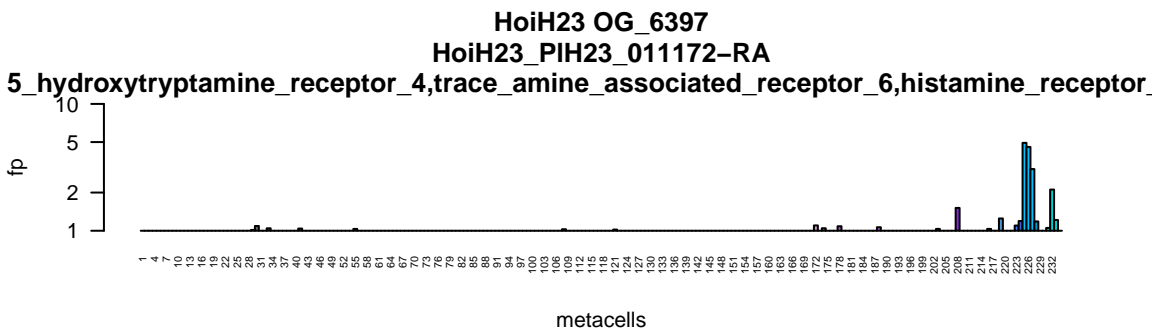
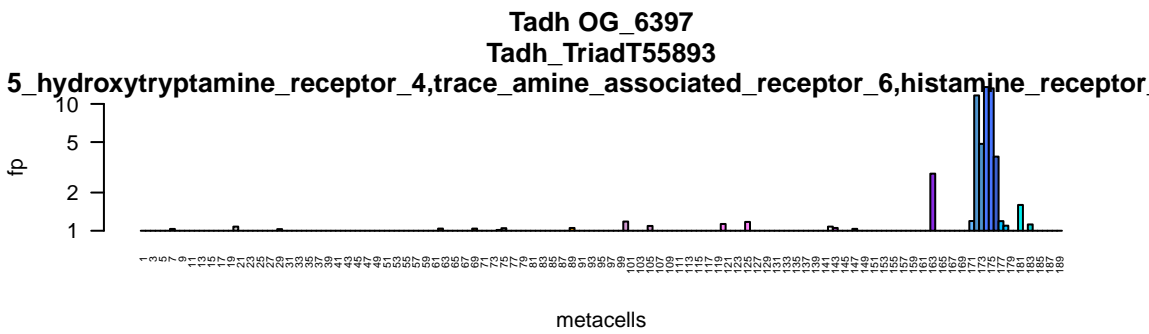
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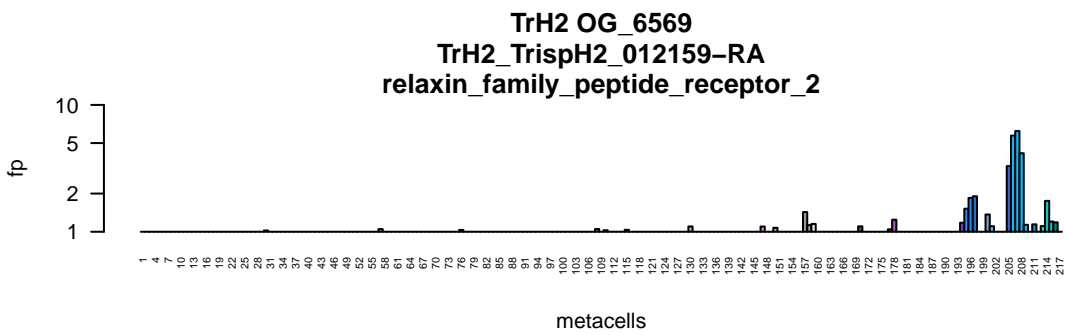
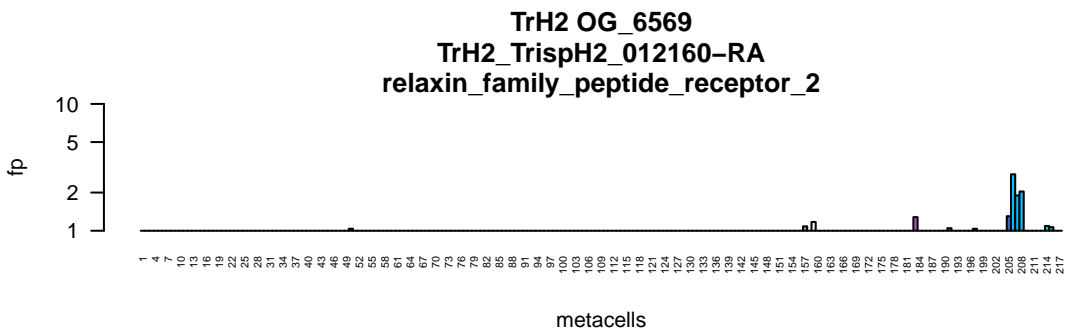
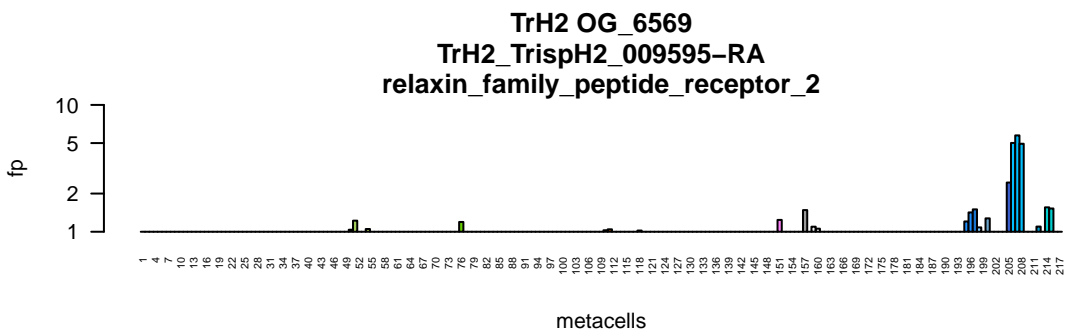
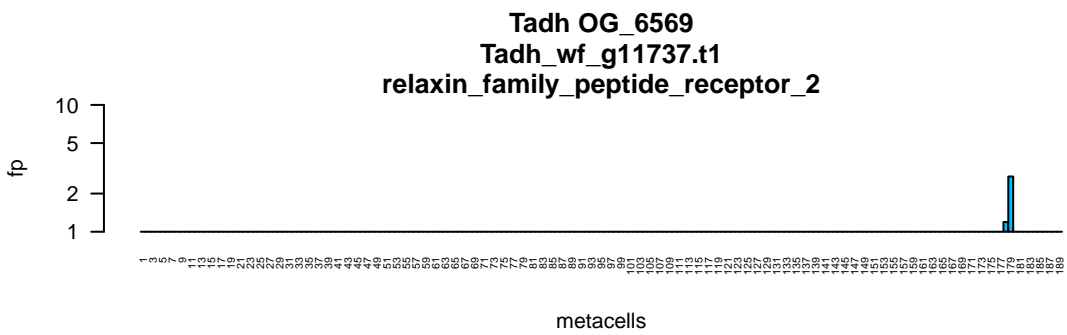
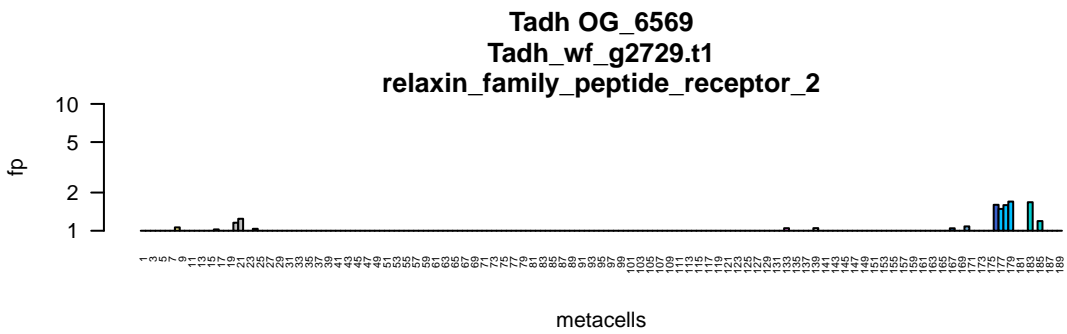
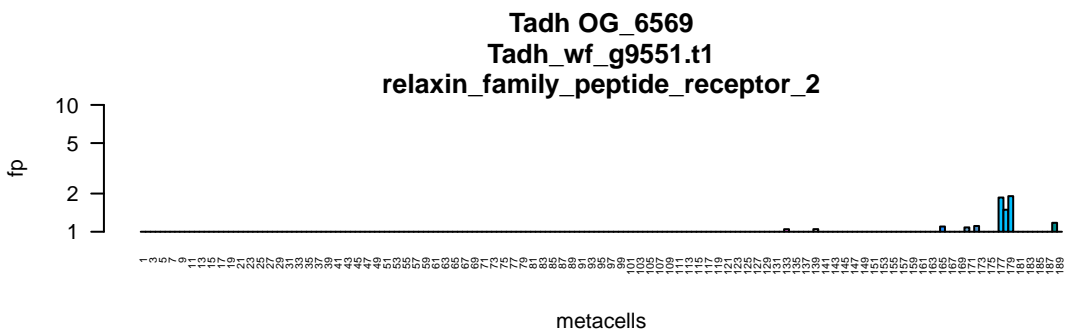
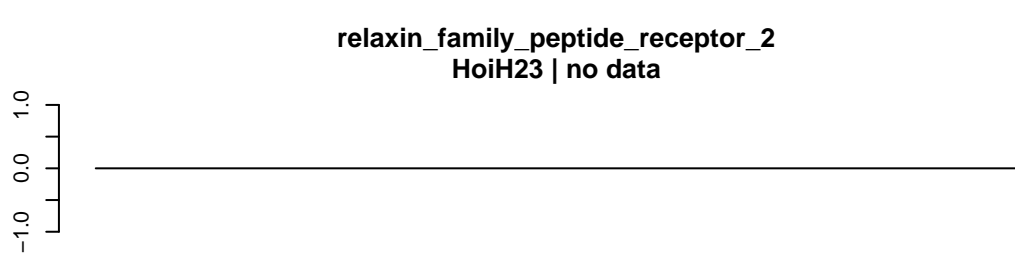
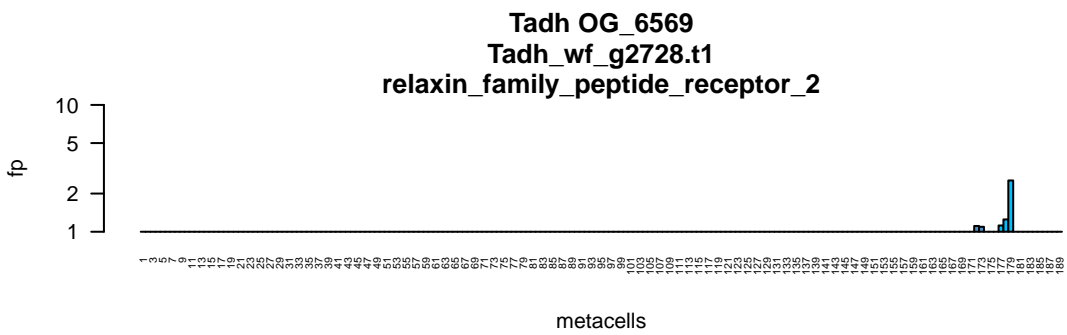
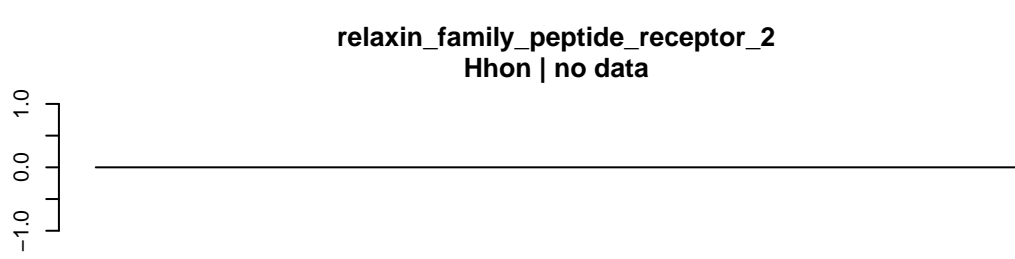
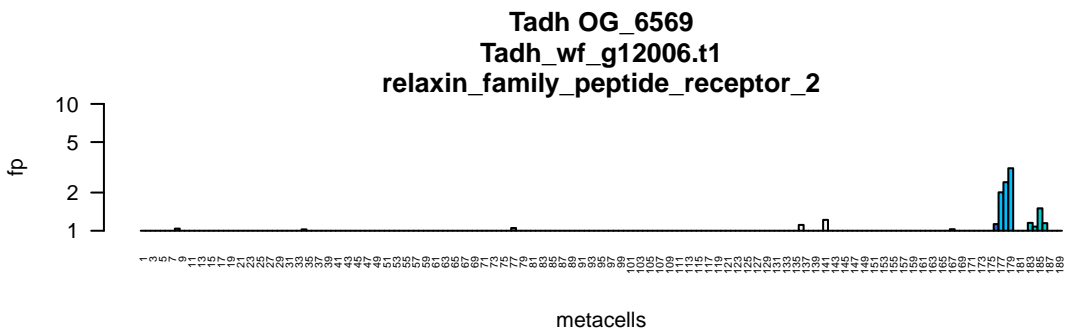


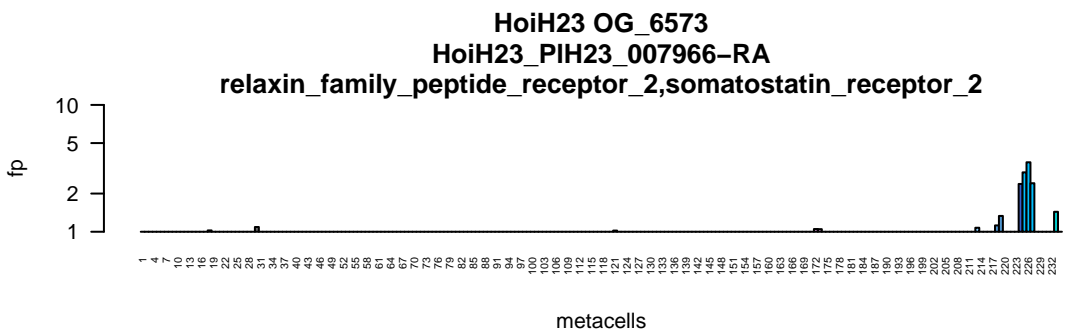
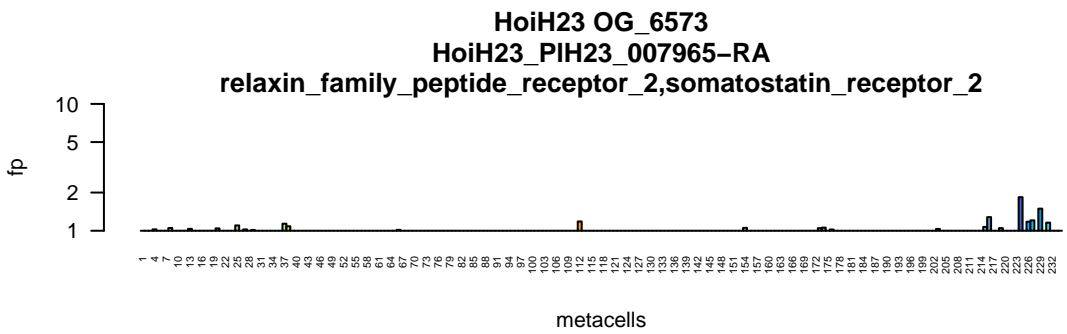
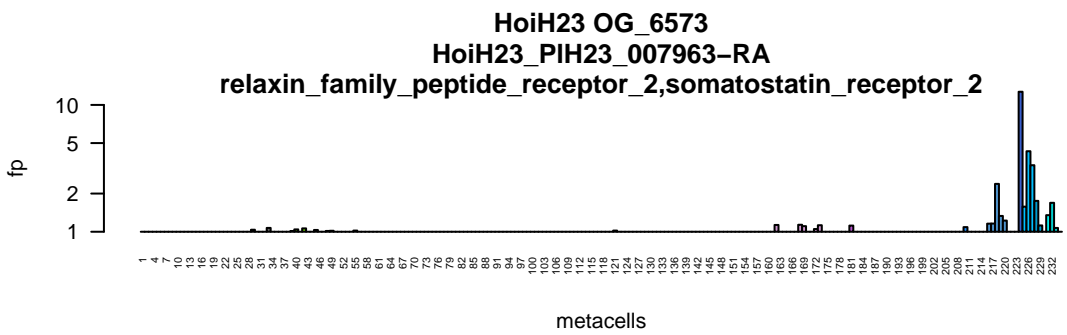
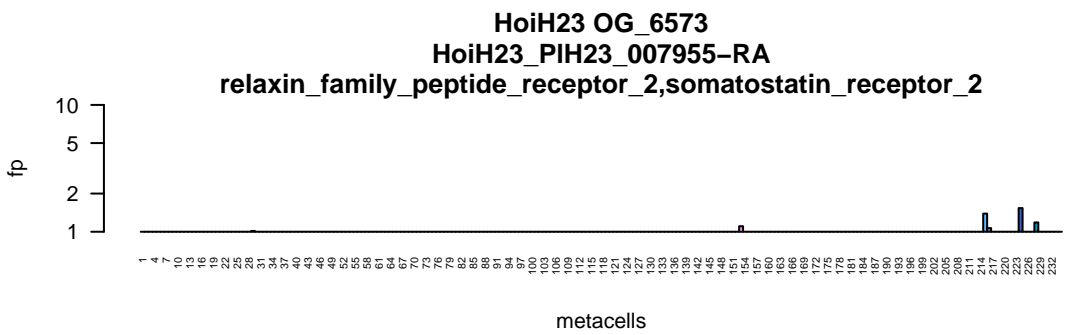
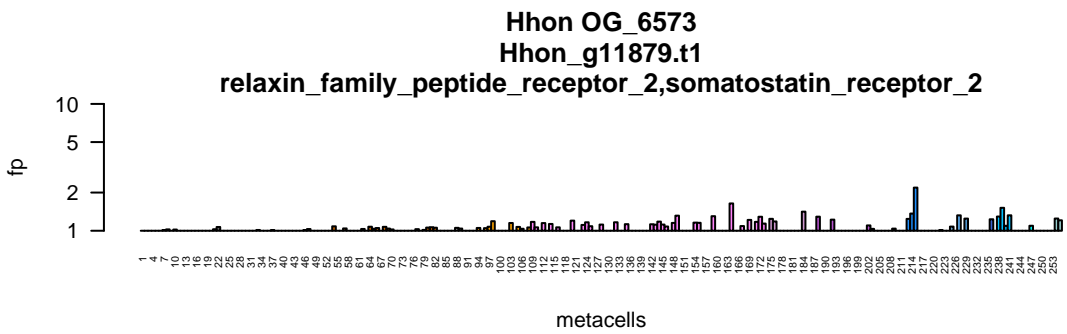
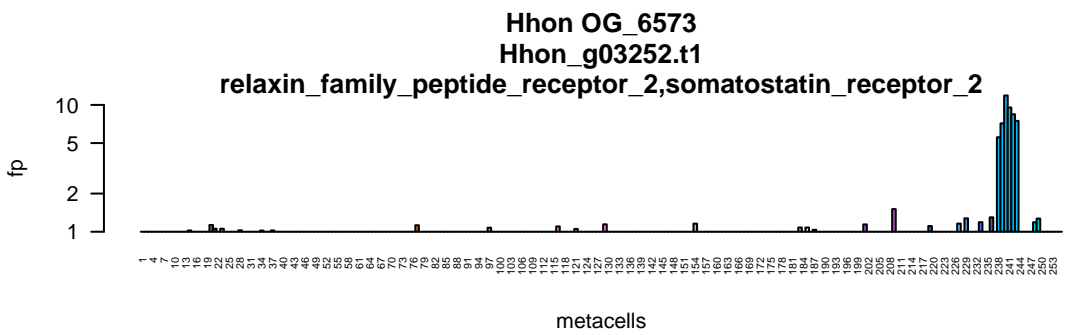
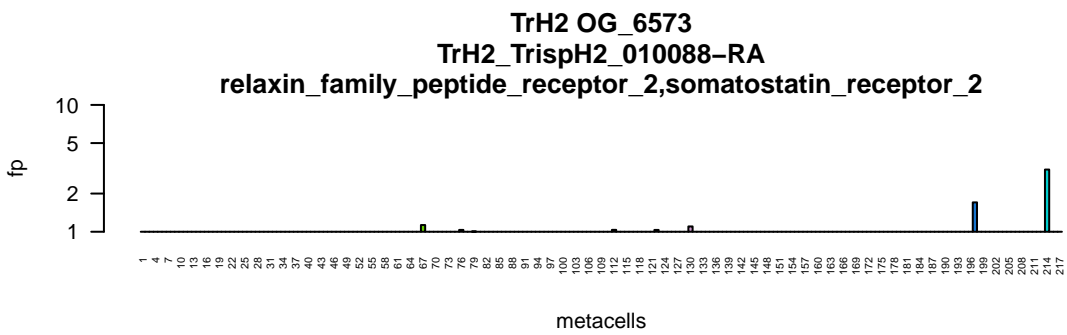
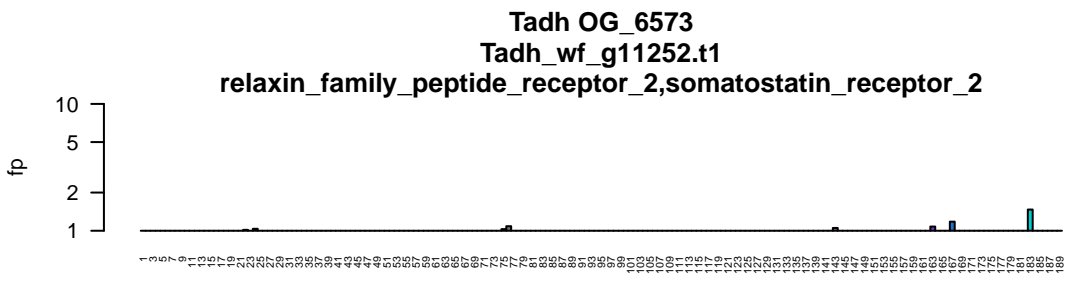
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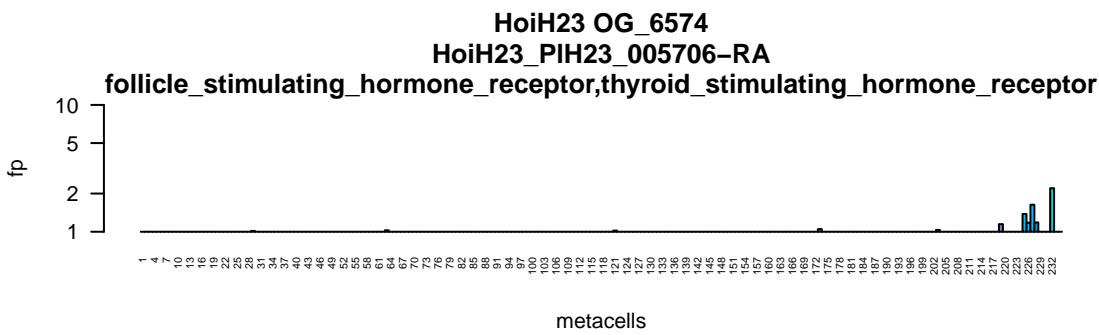
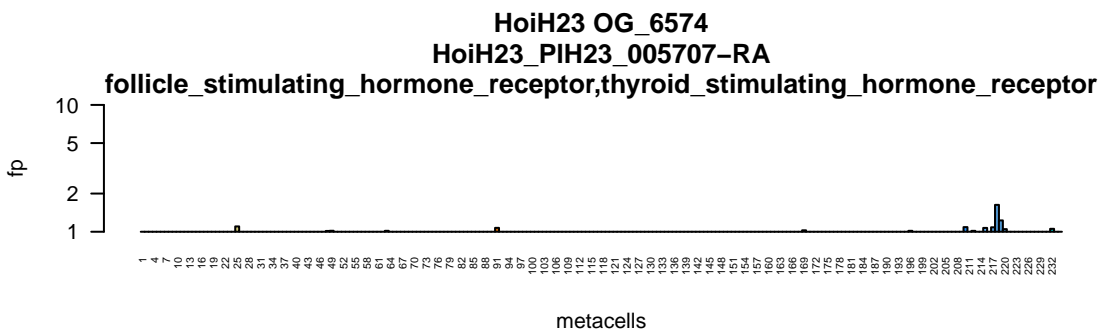
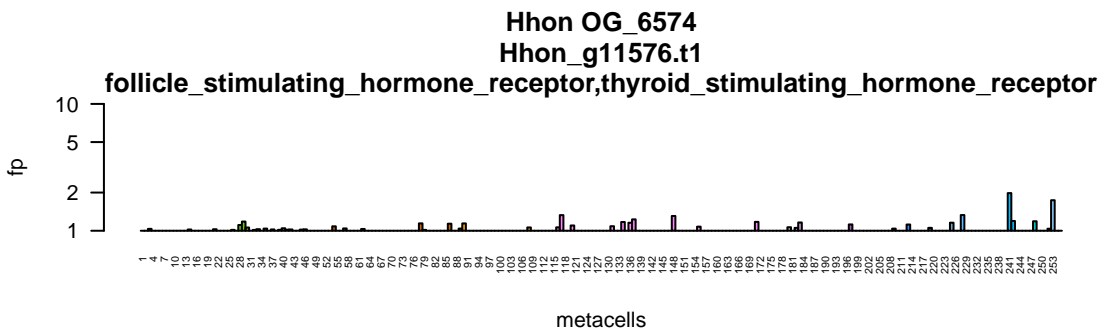
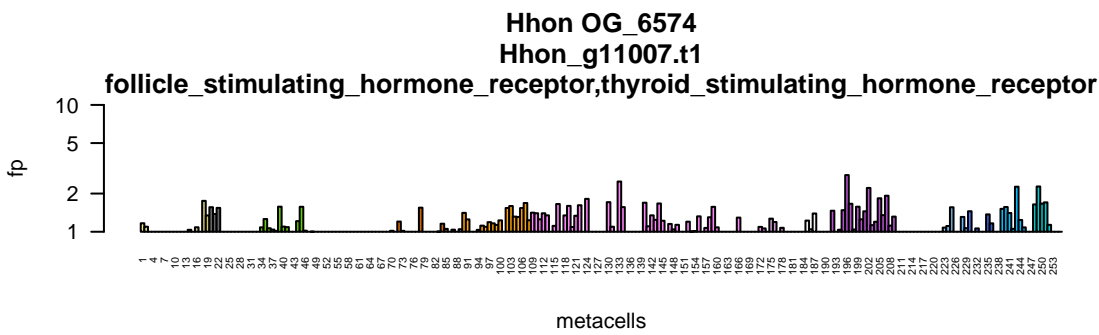
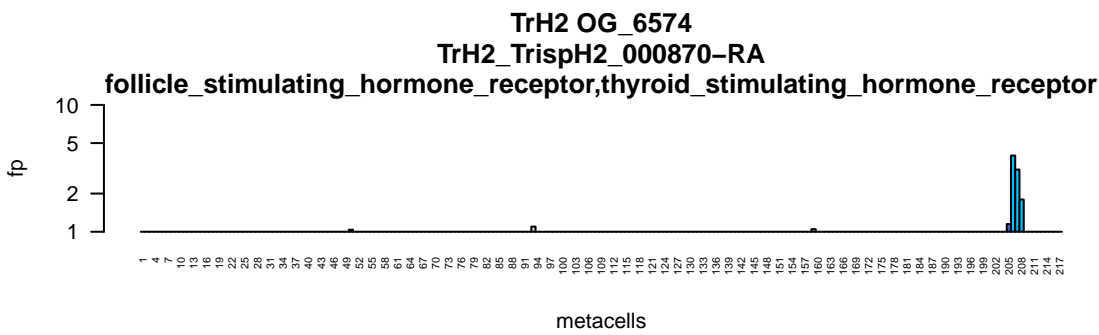
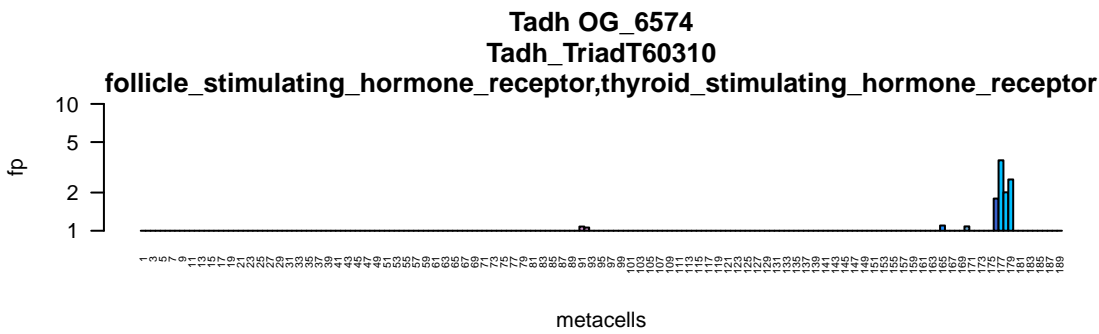


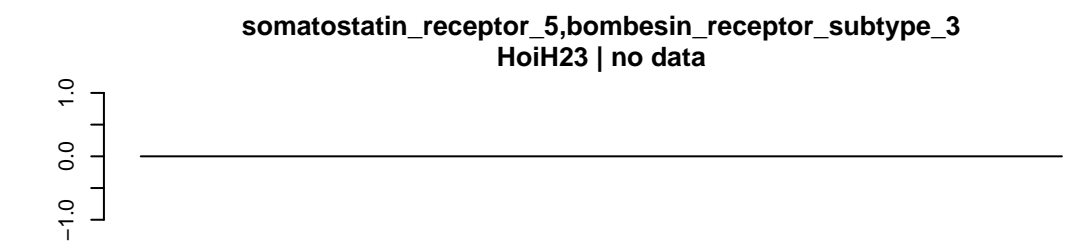
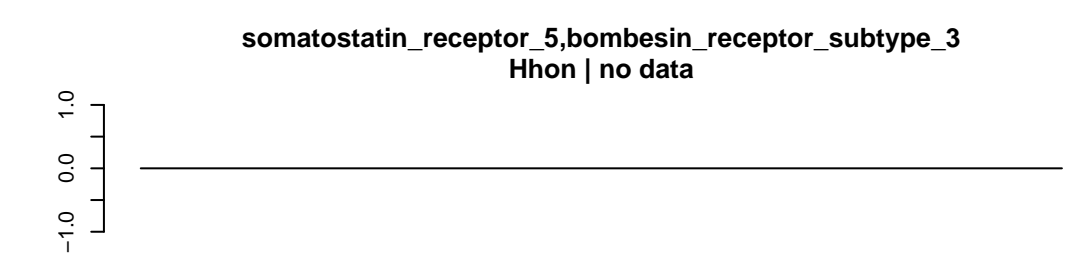
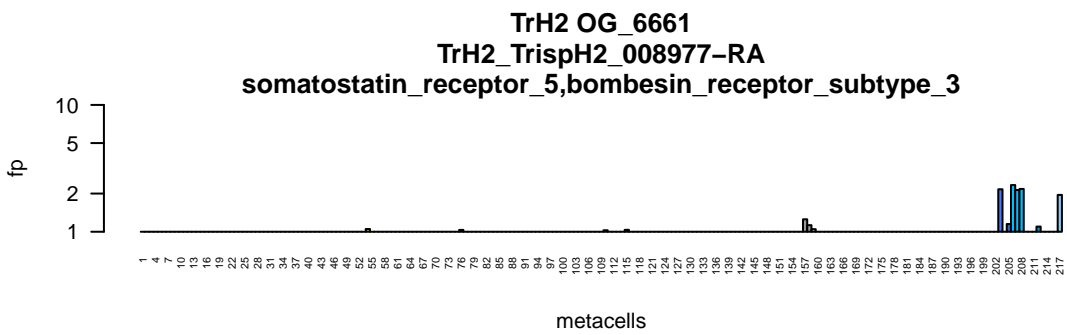
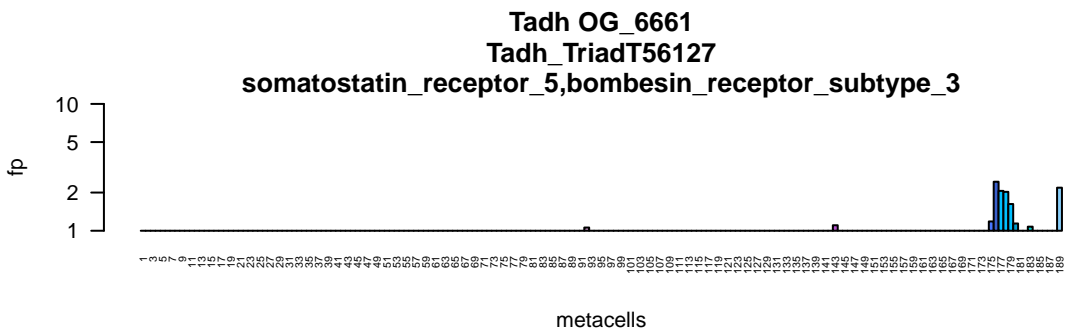




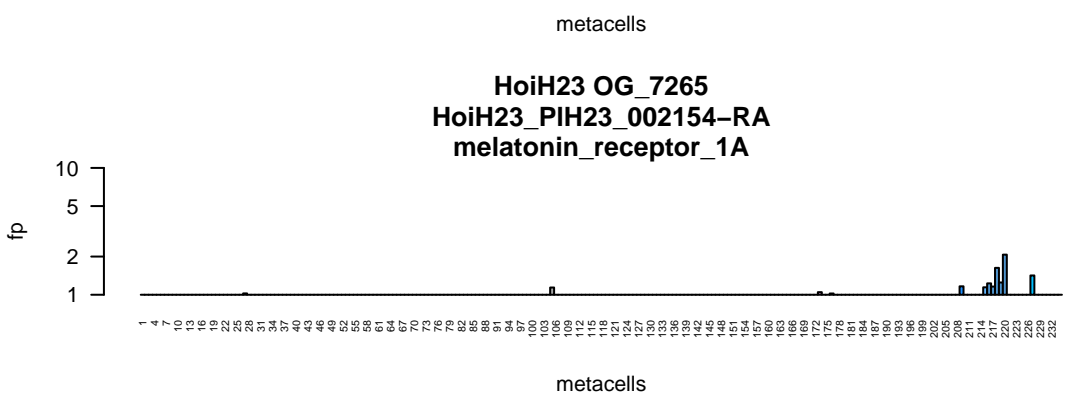
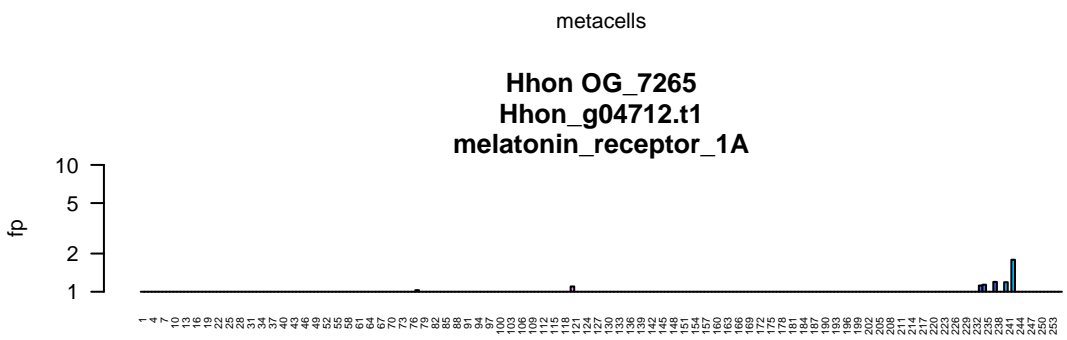
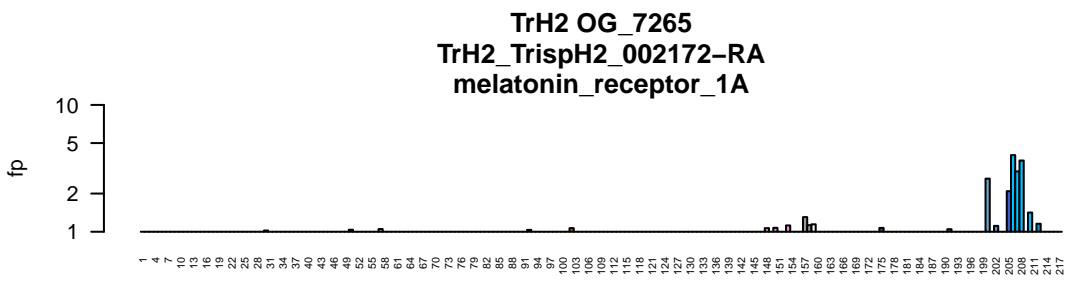
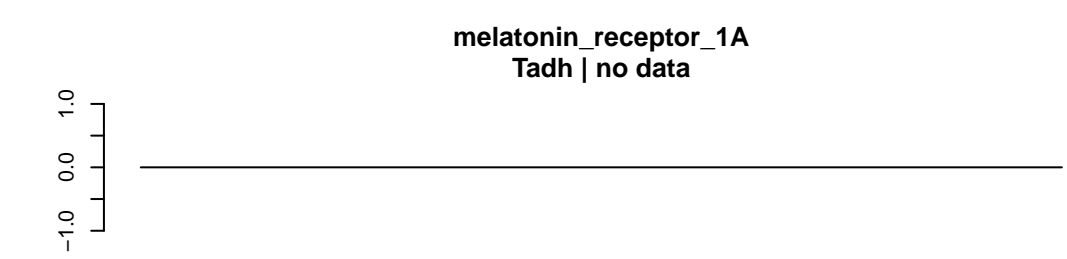


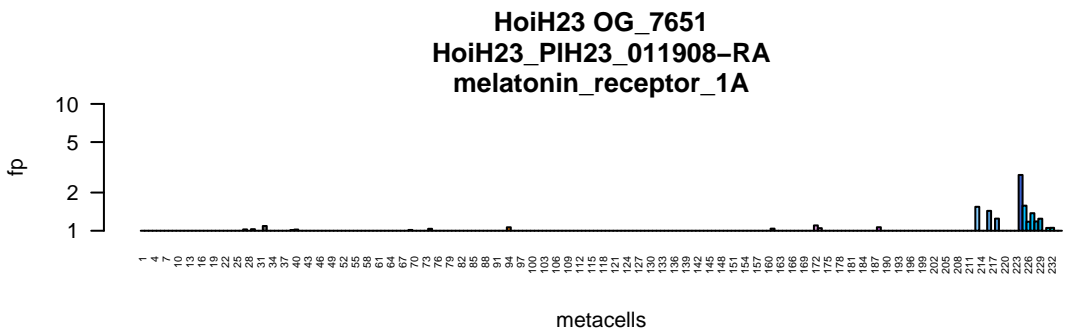
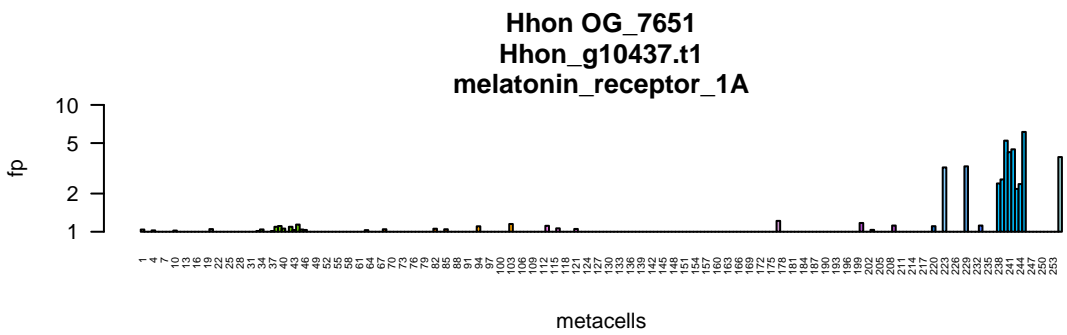
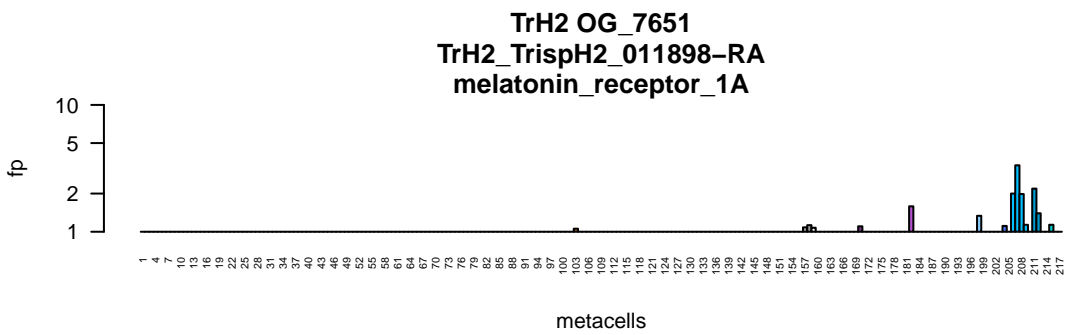
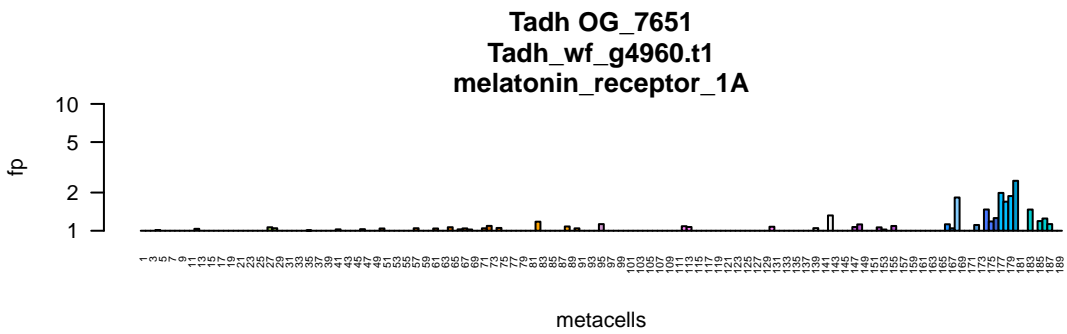


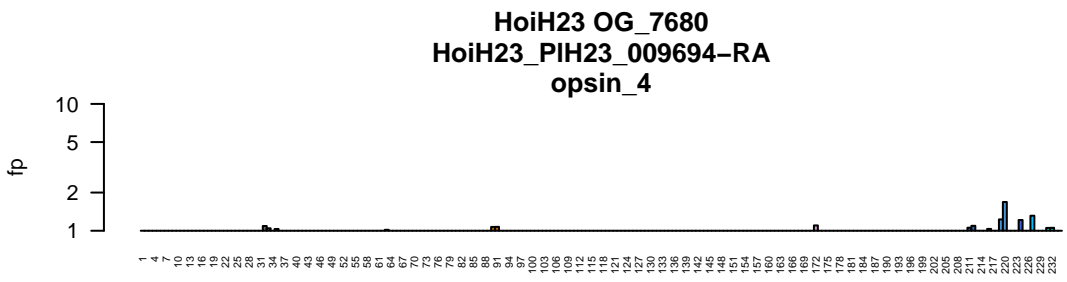
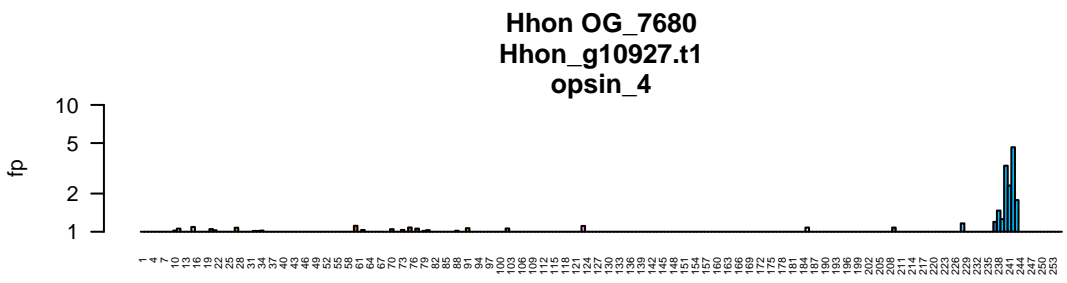
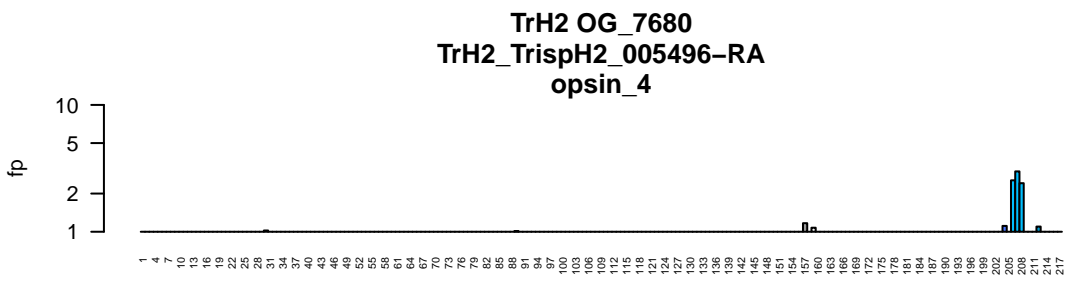
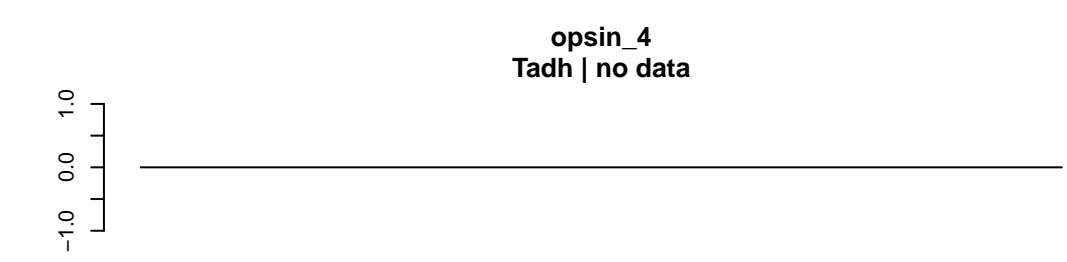


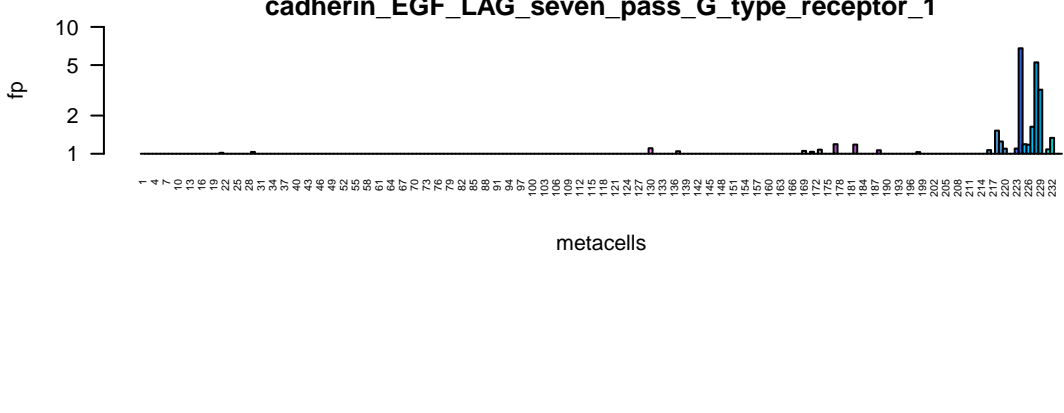
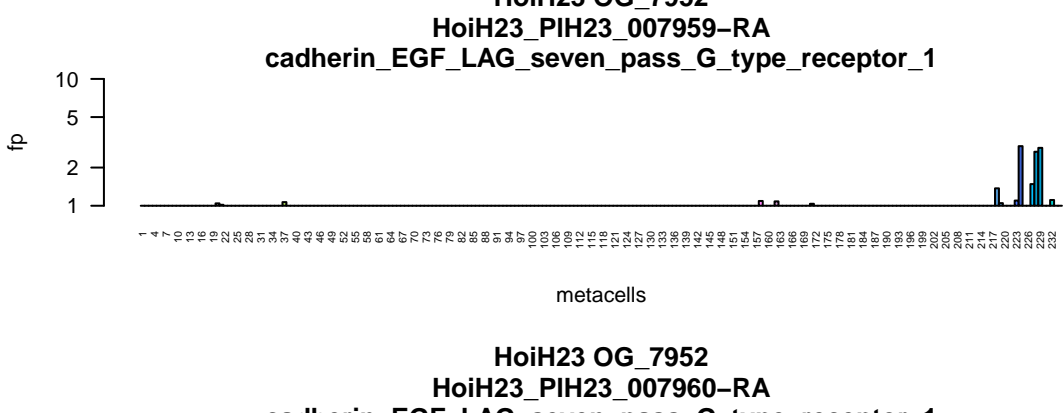
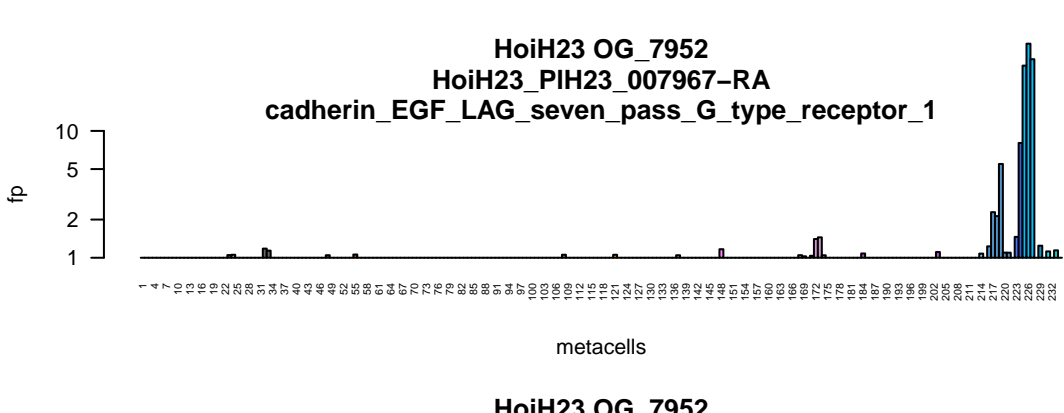
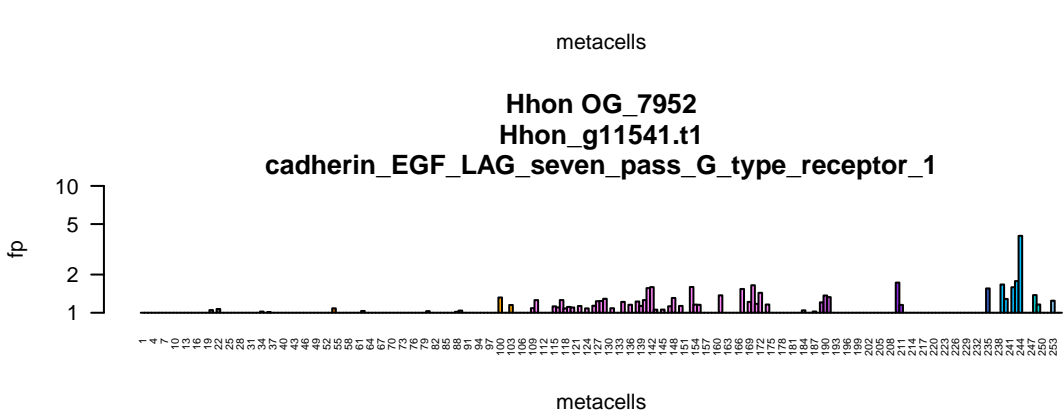
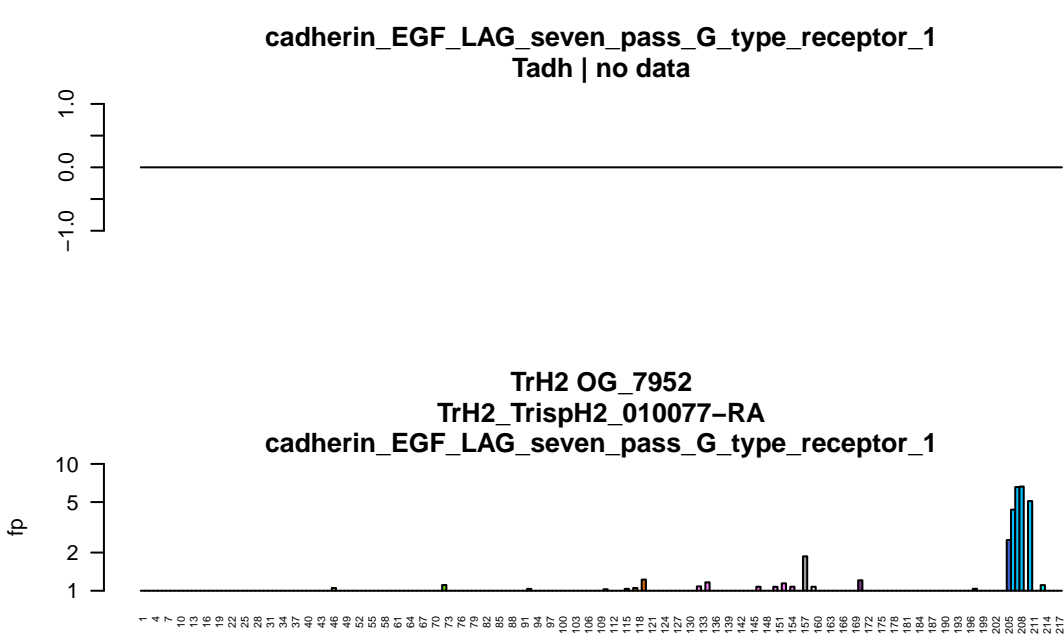


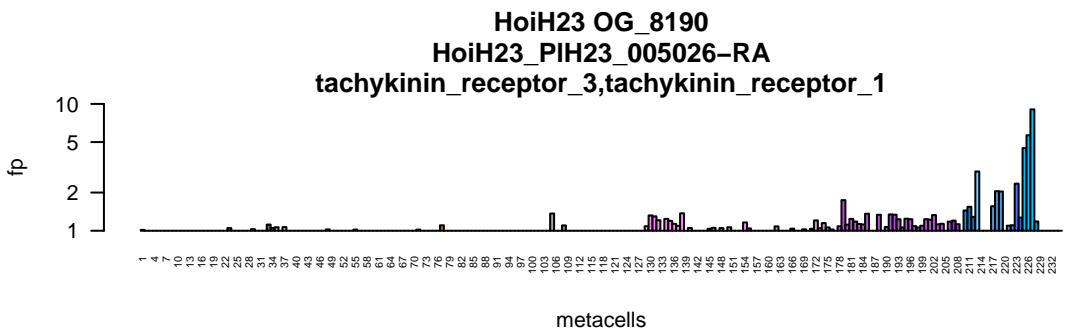
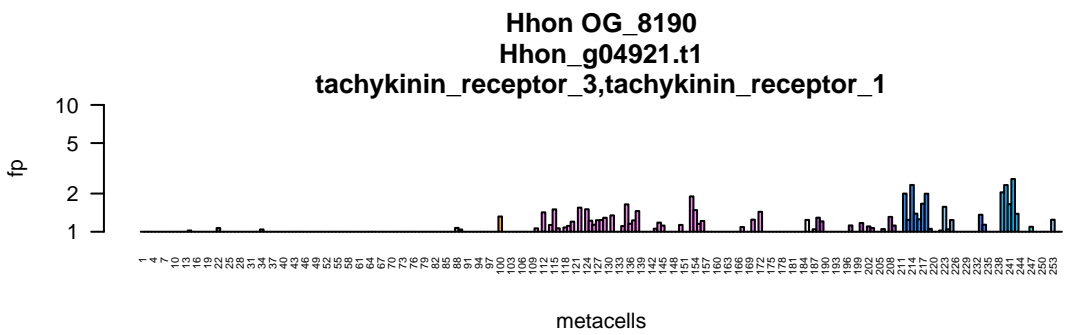
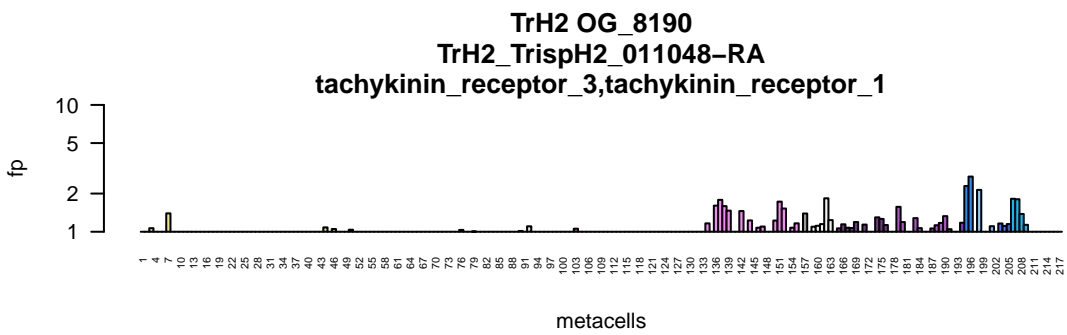
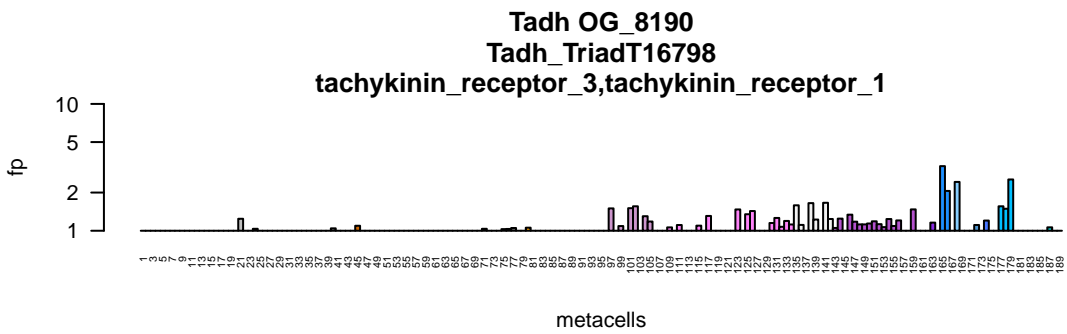


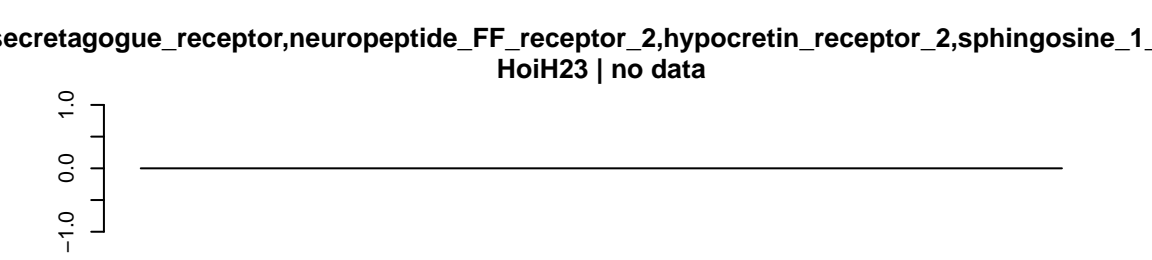
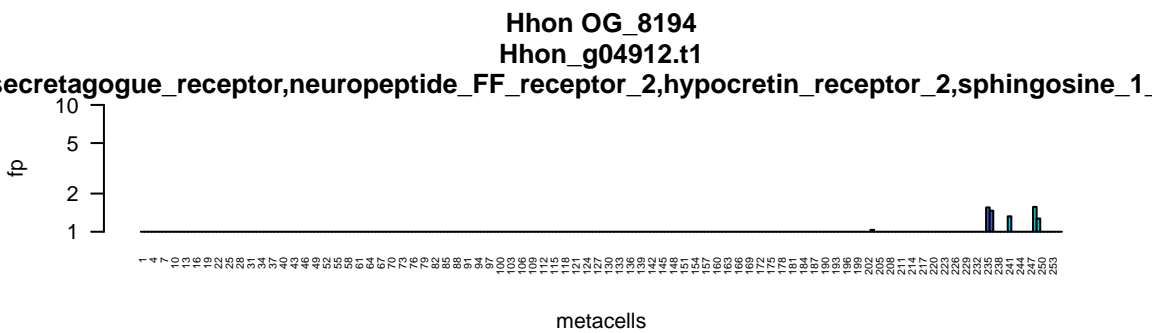
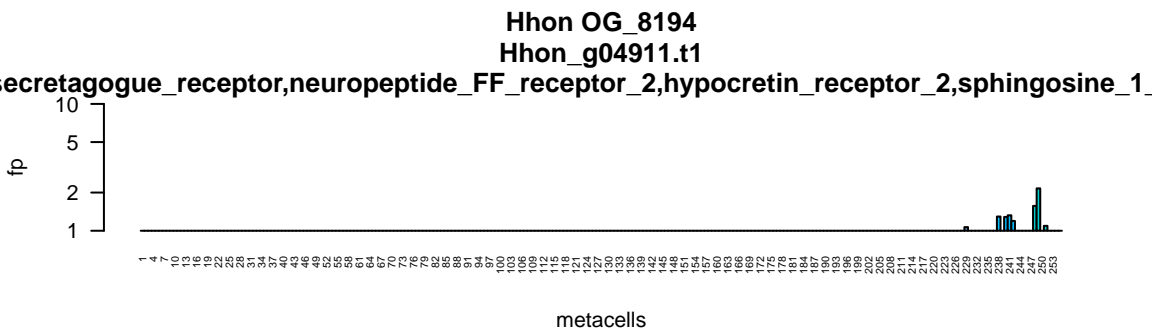
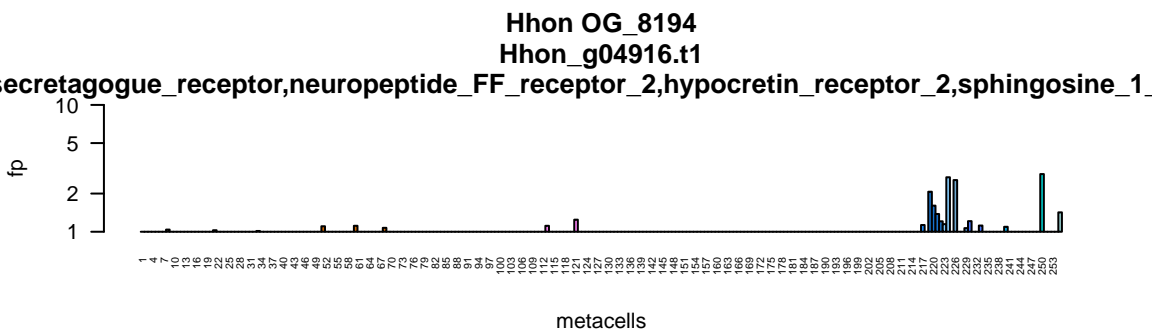
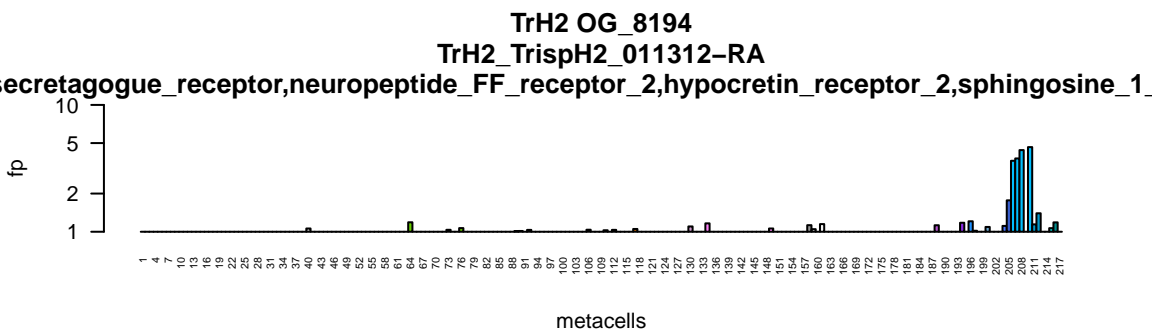
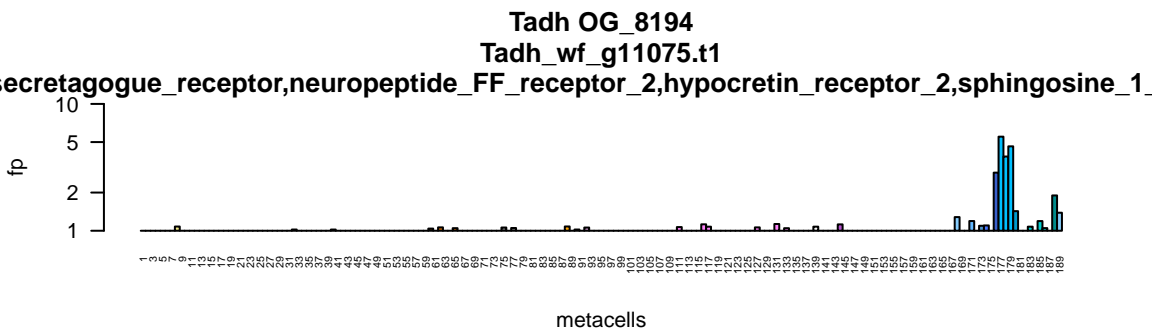






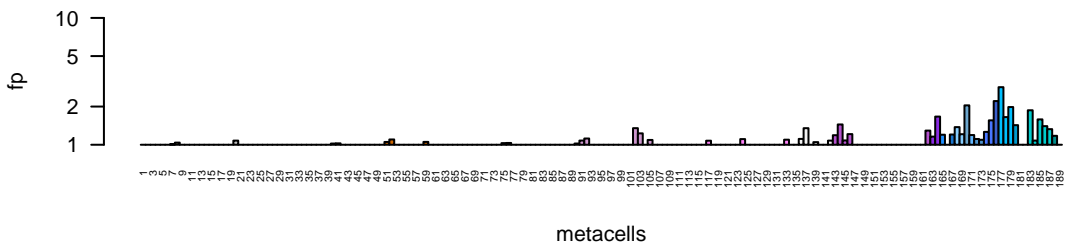






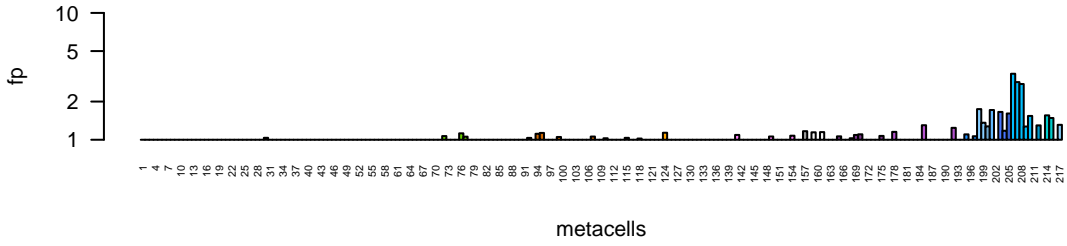
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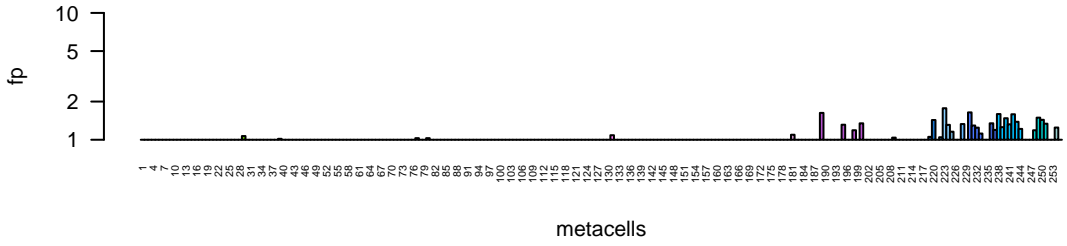
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TrH2\_TrispH2\_007497-RA



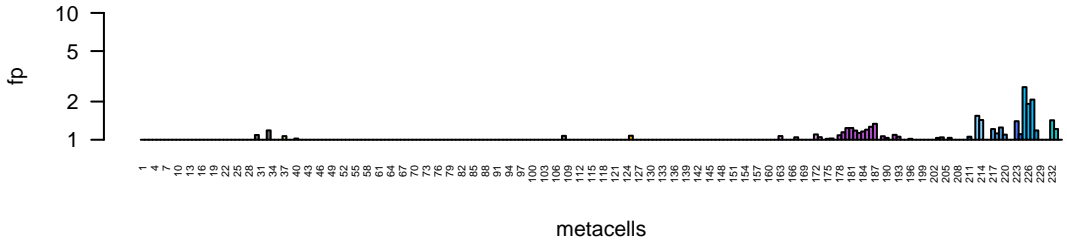
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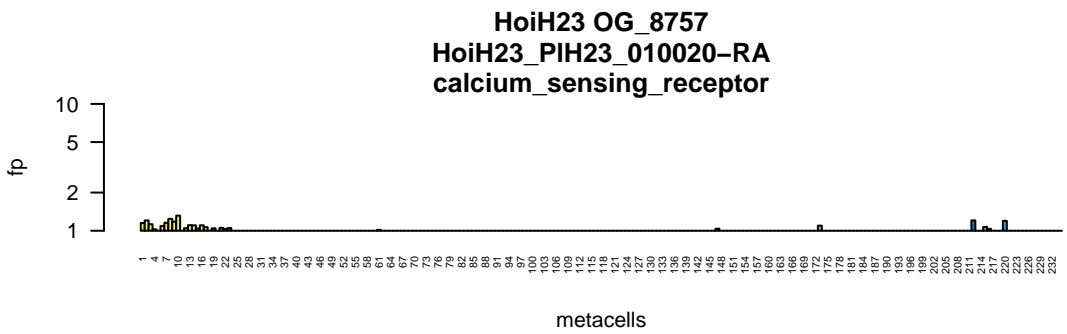
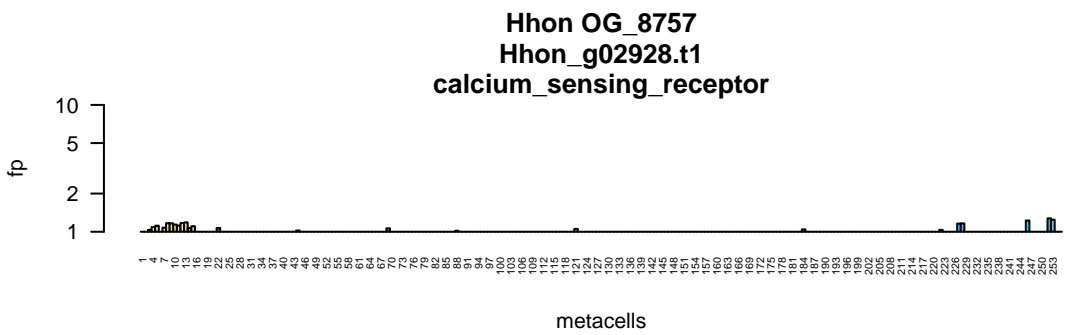
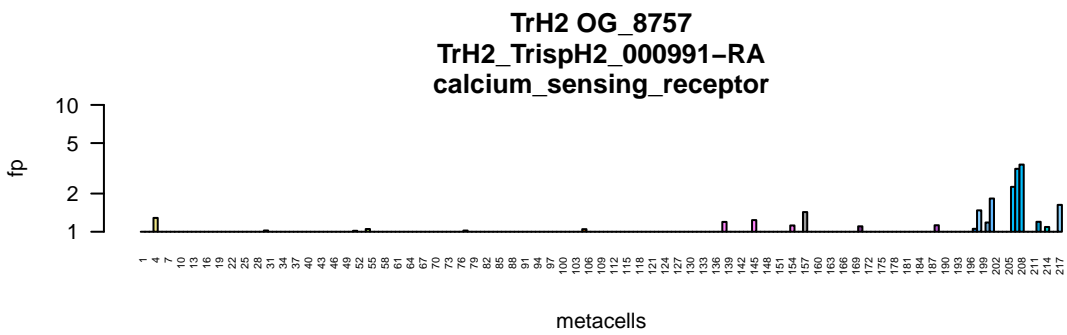
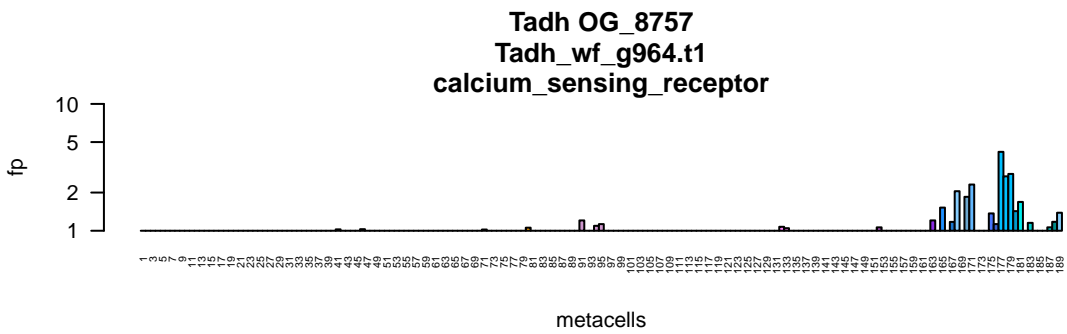
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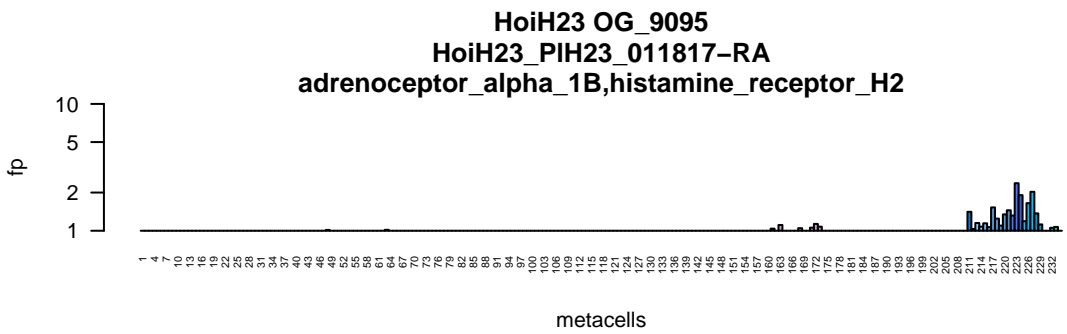
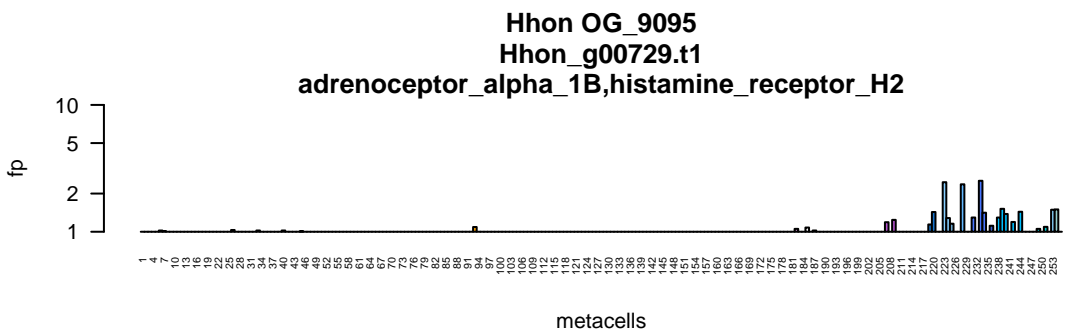
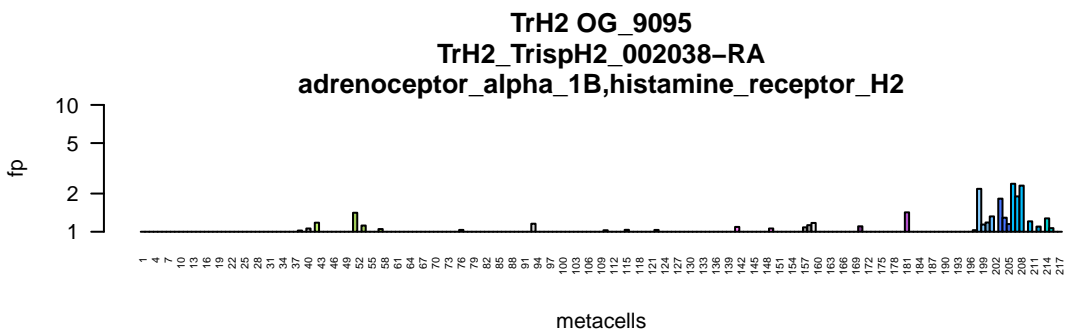
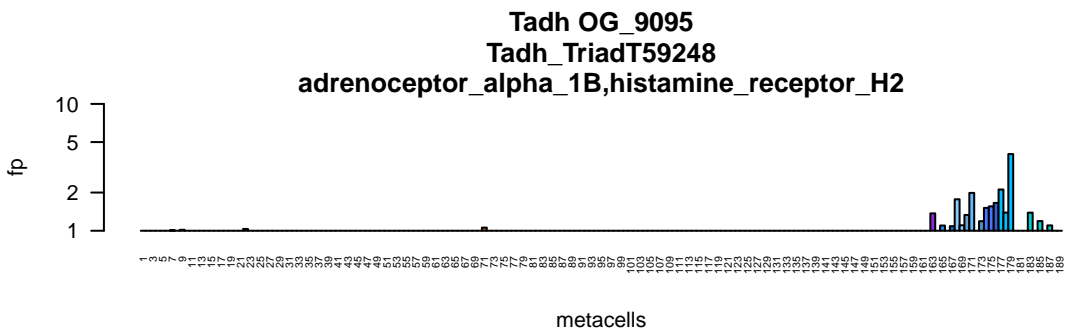
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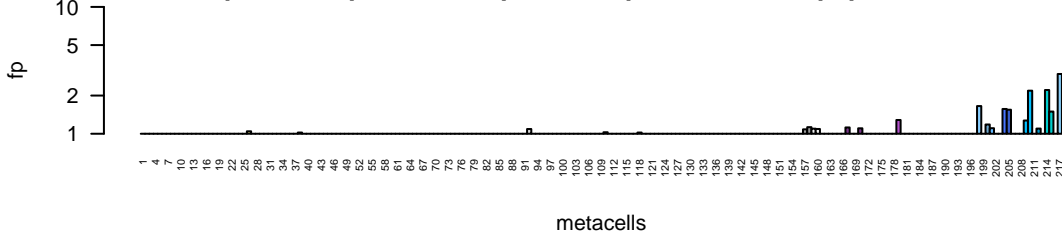




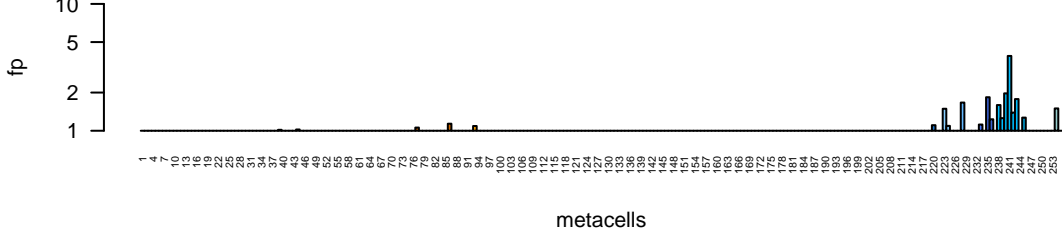
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Tadh | no data



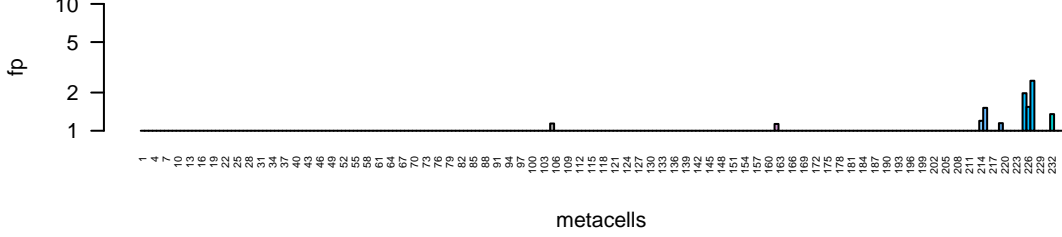
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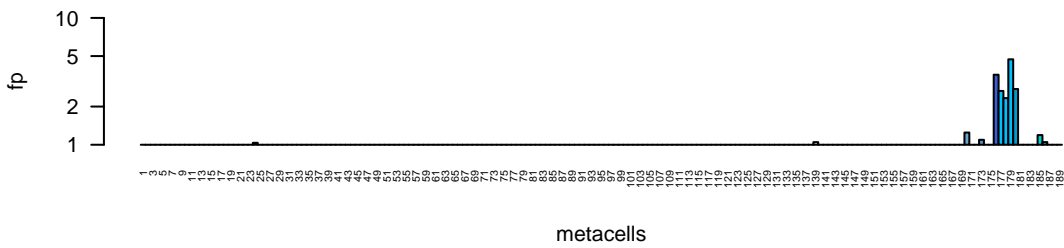
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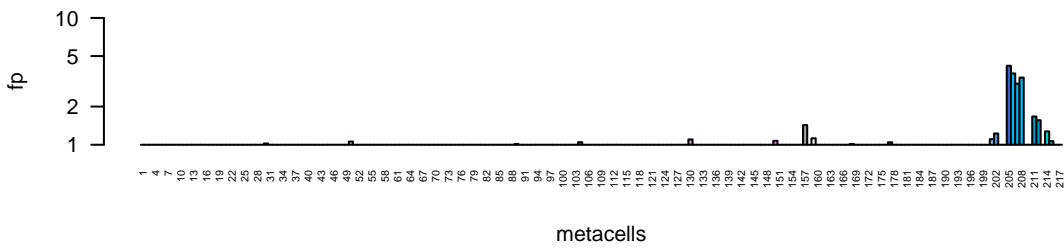
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HoiH23\_PIH23\_001231-RA



Tadh OG\_10419  
Tadh\_wf\_g4956.t1



TrH2 OG\_10419  
TrH2\_TrispH2\_008975-RA



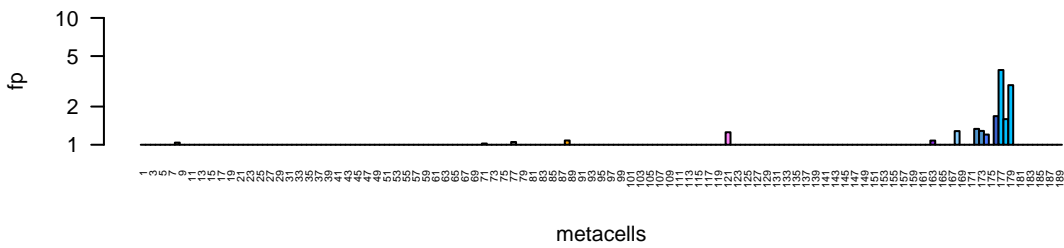
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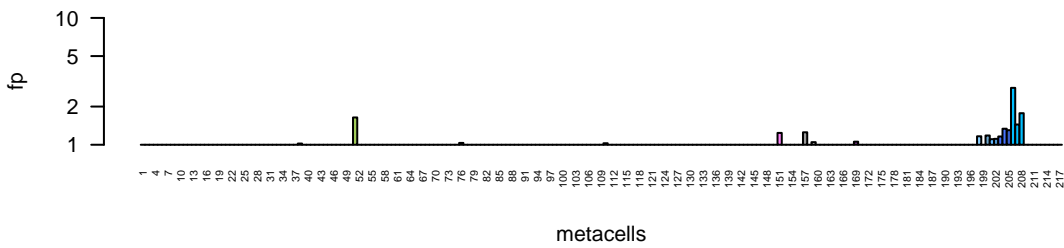
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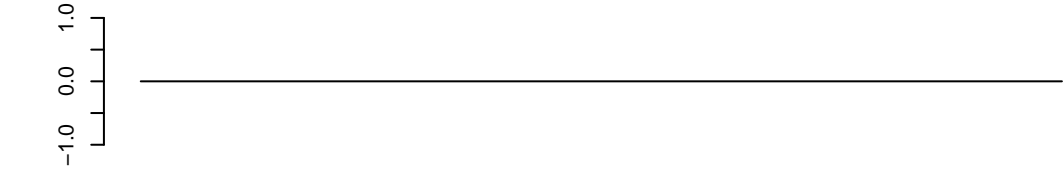
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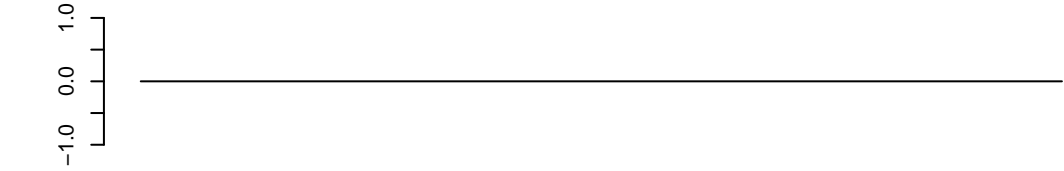
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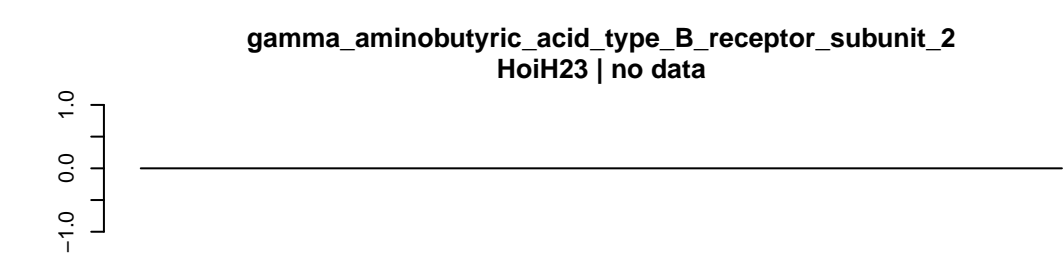
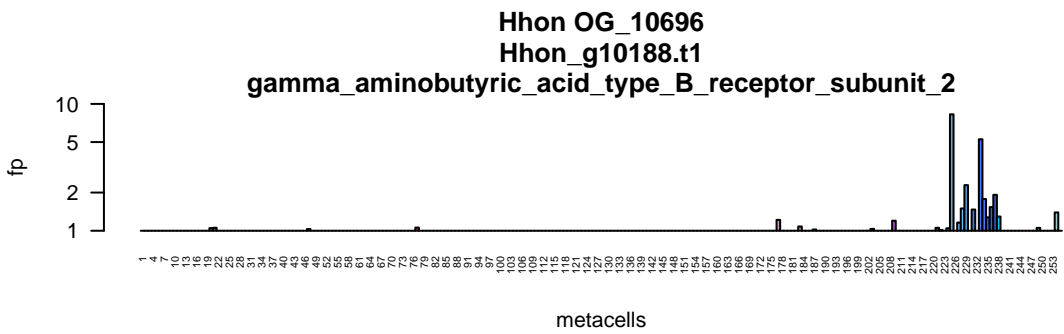
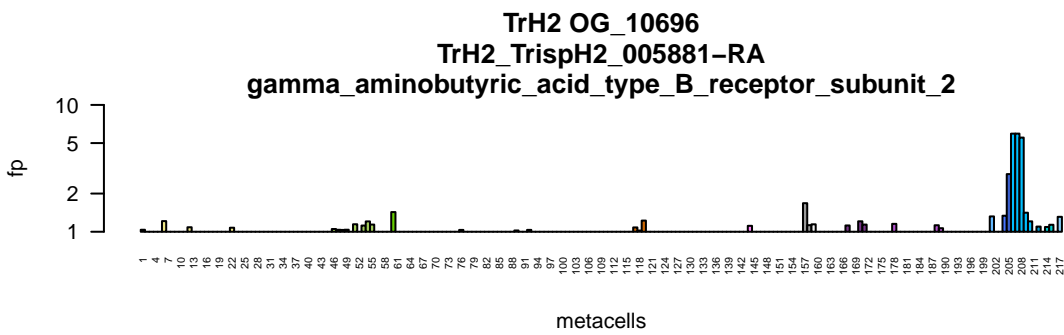
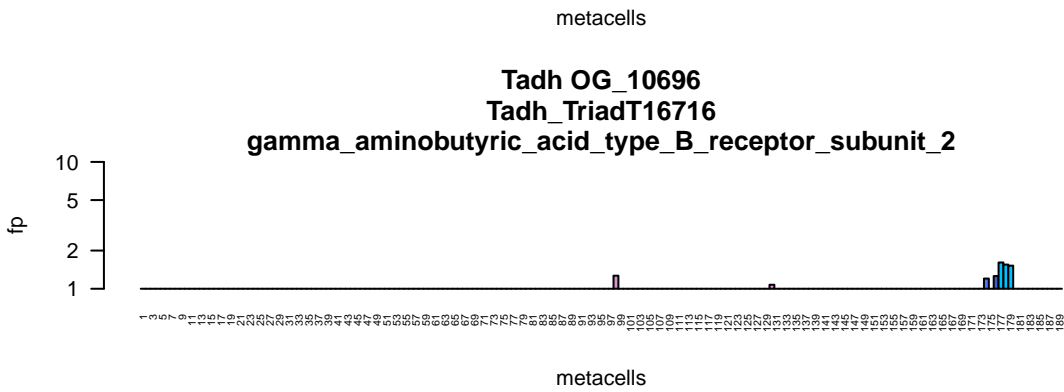
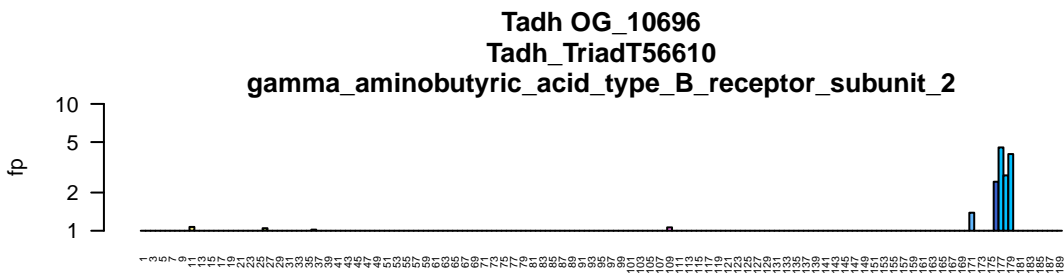


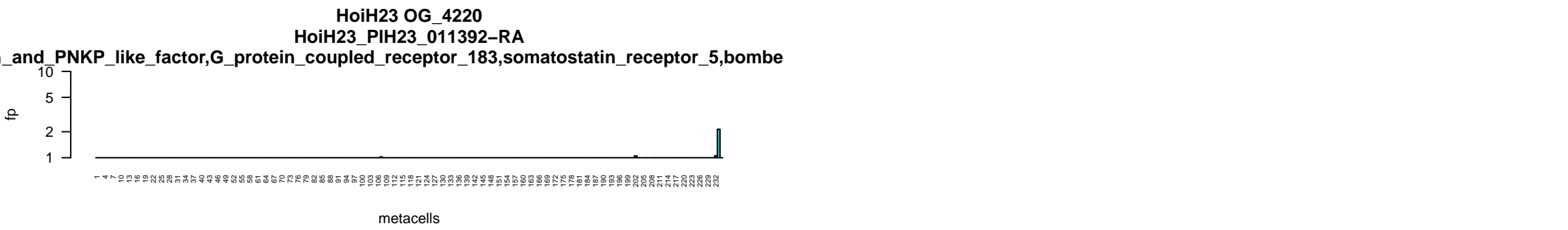
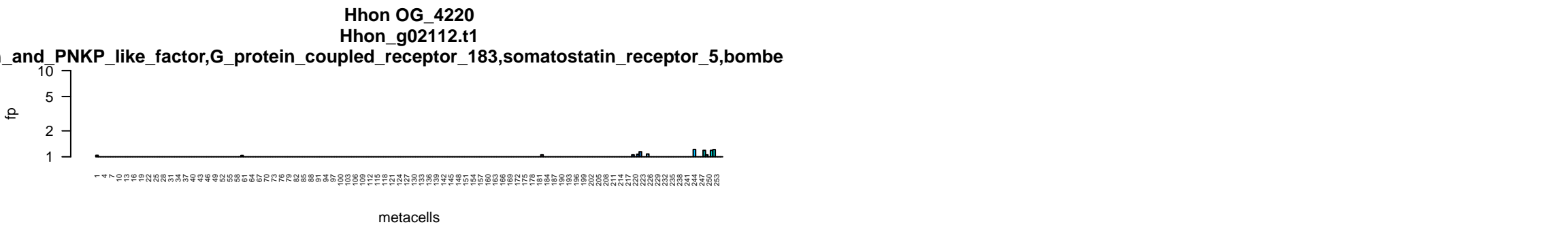
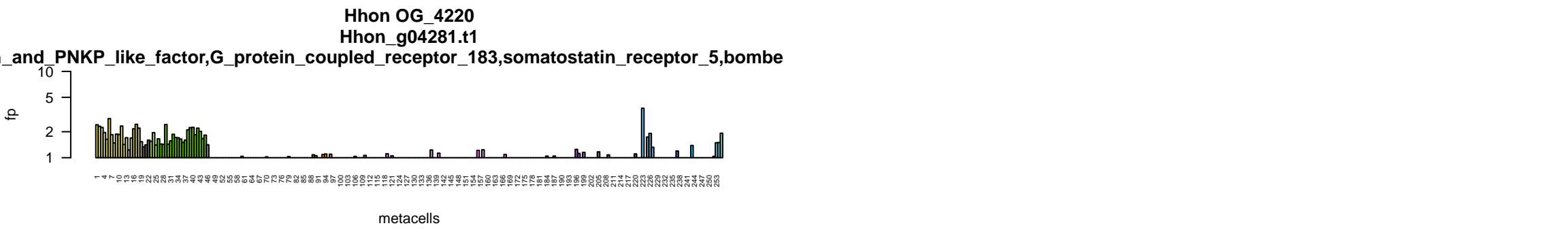
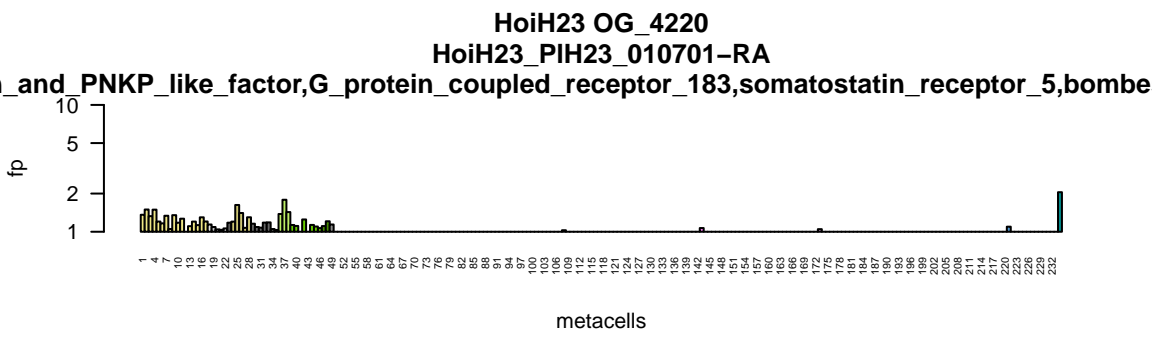
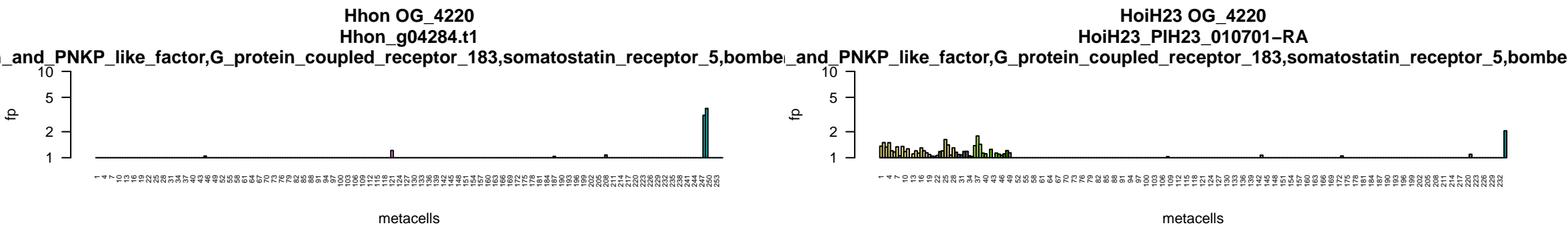
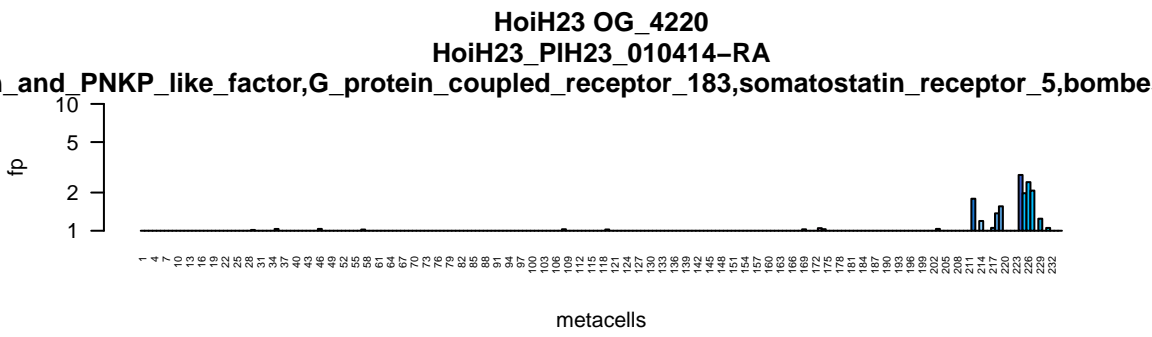
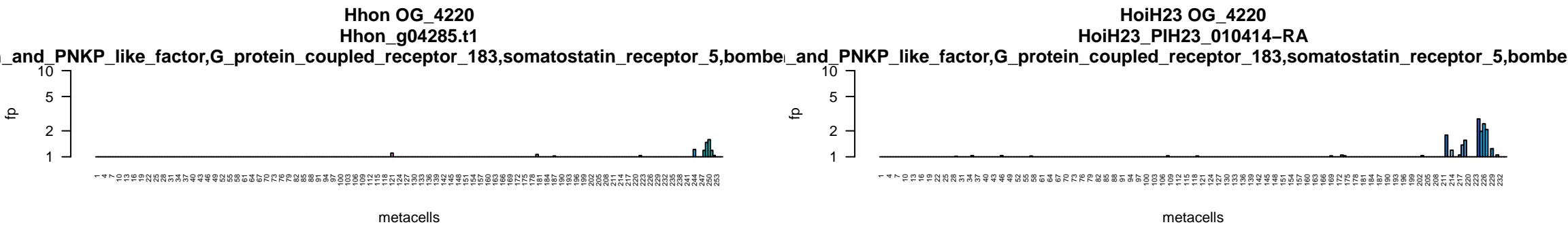
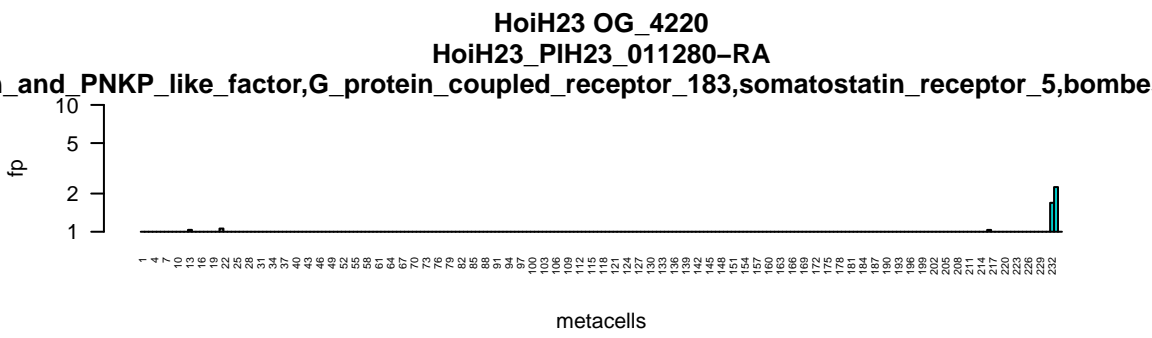
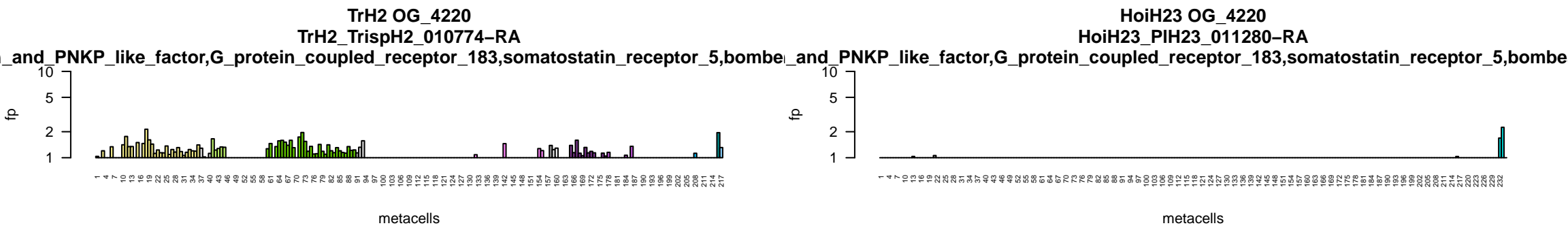
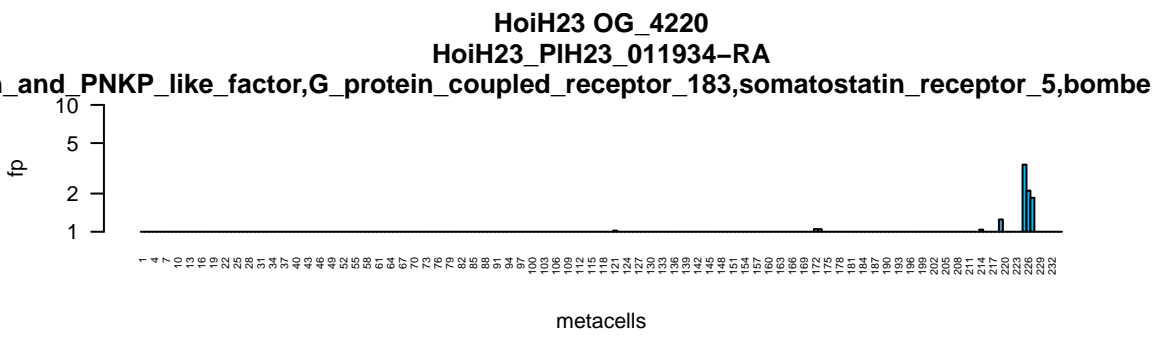
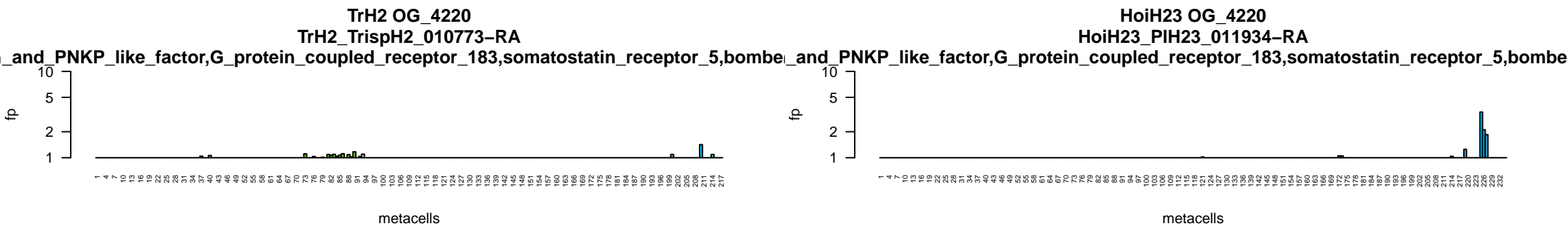
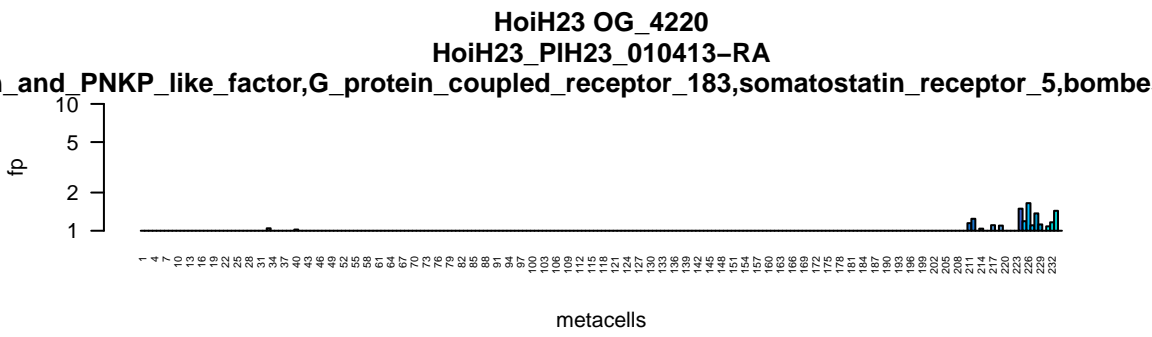
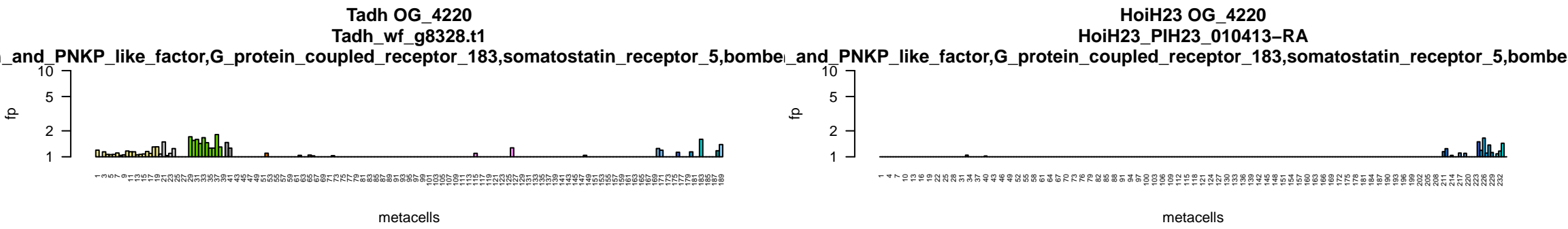
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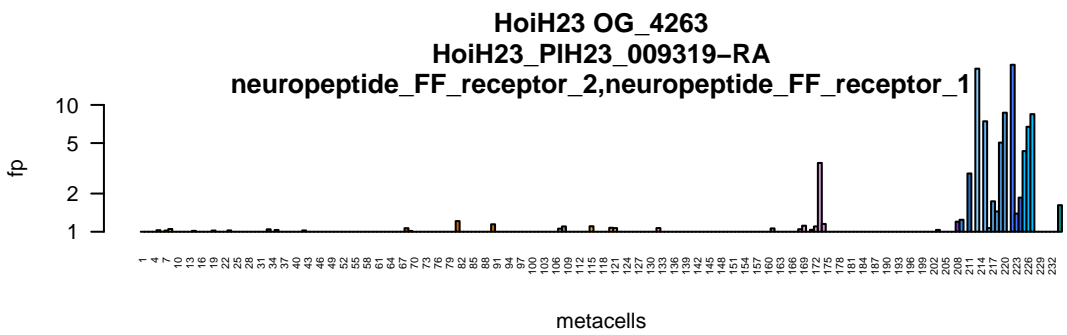
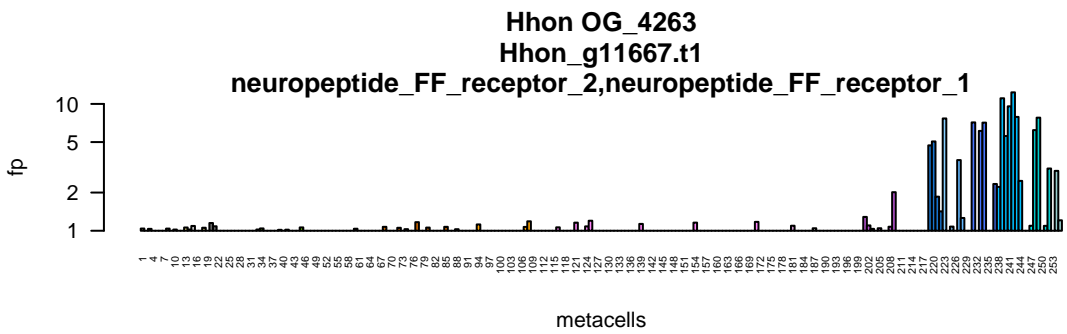
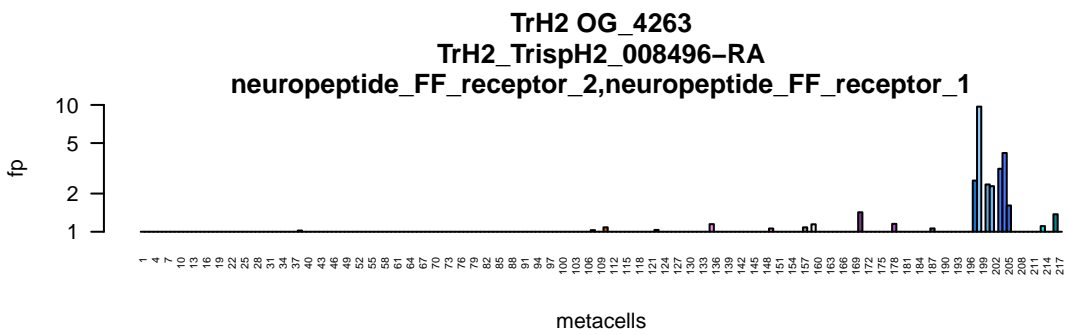
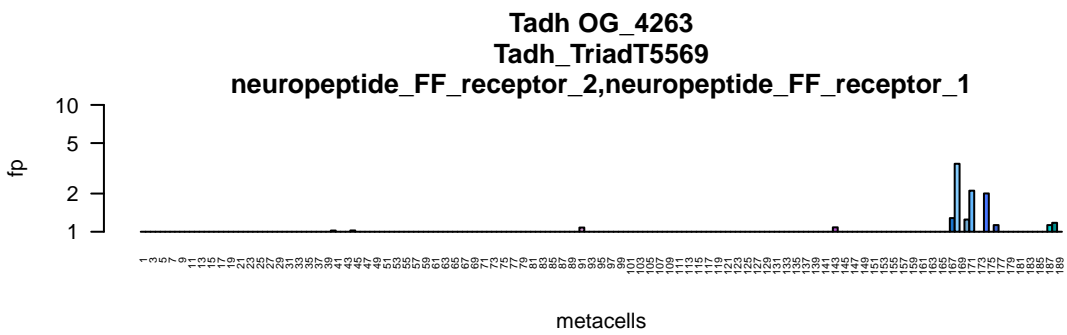
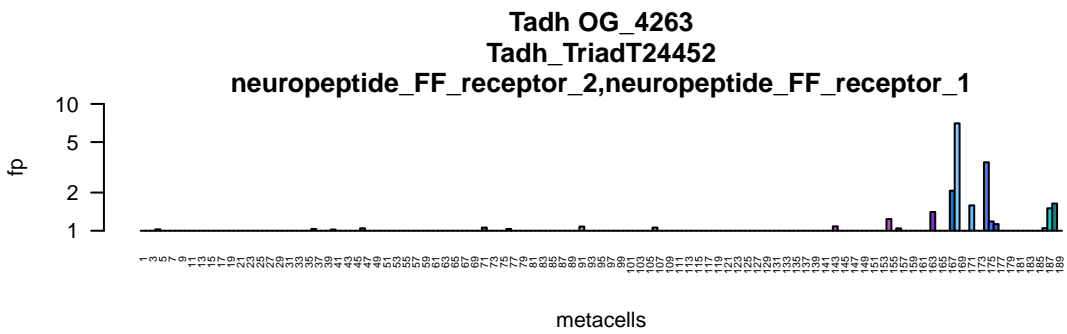


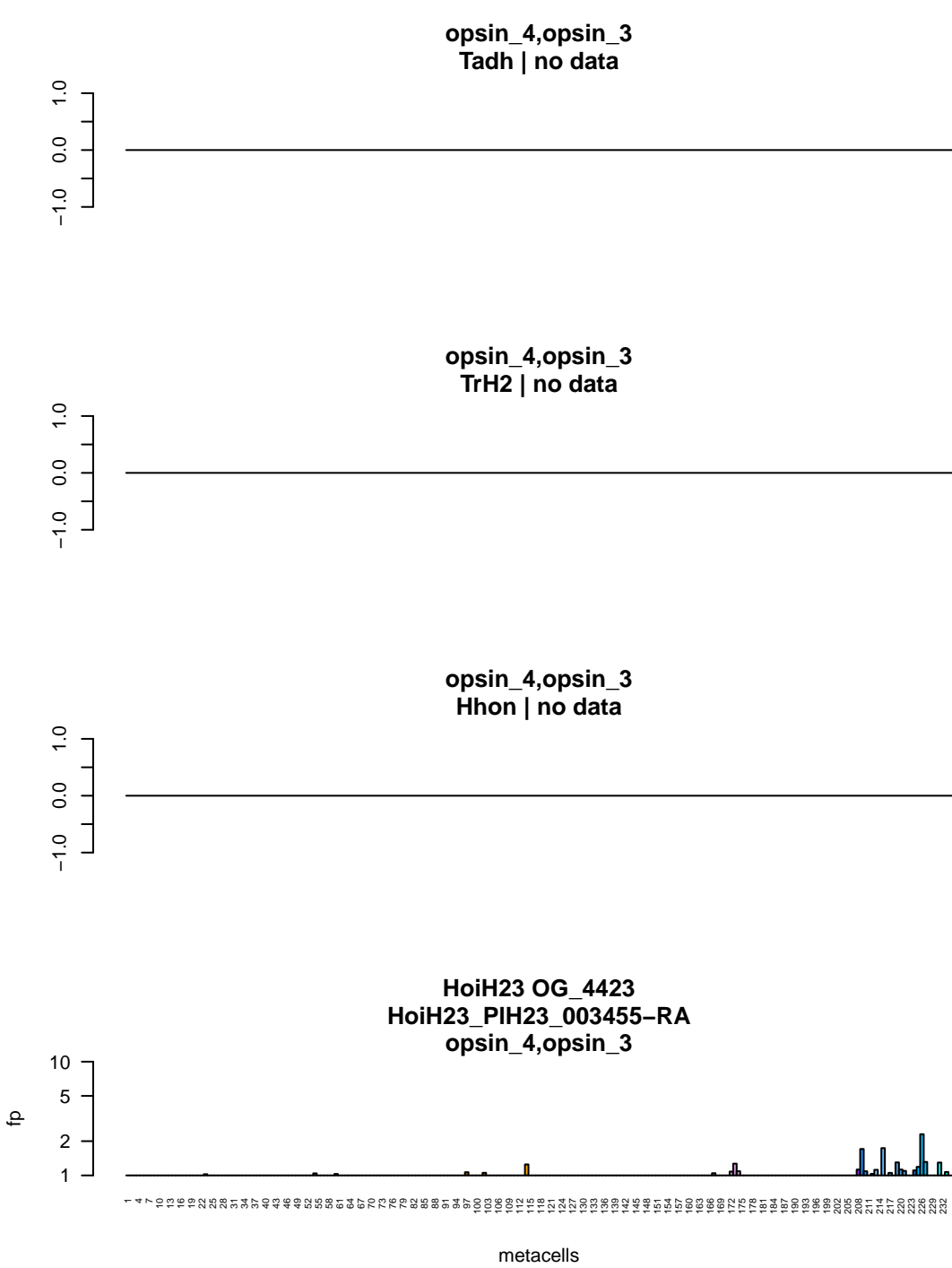
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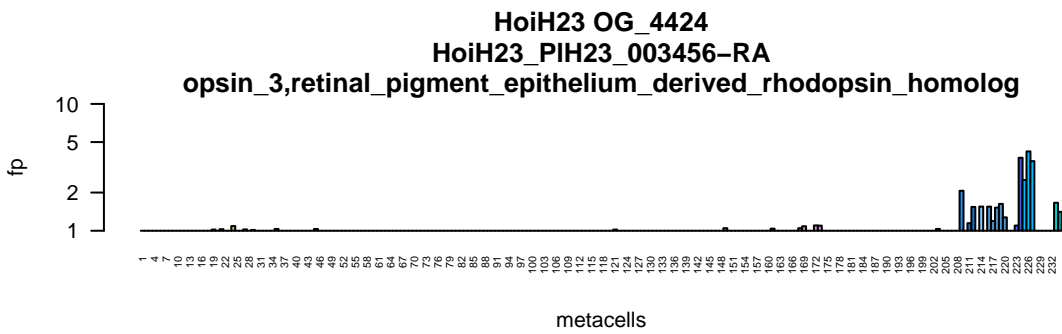
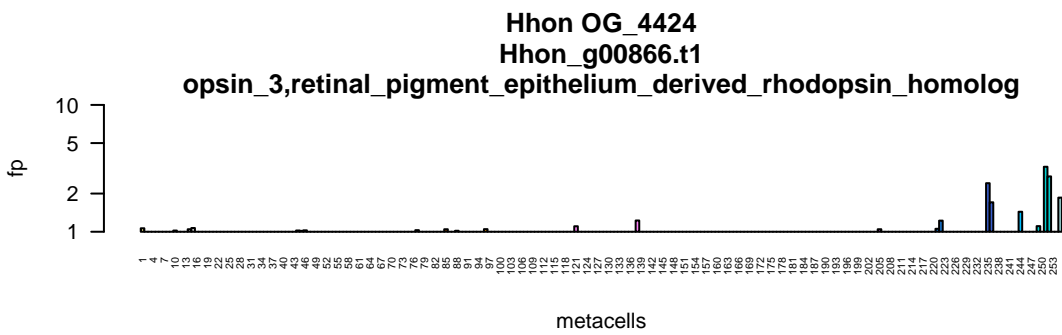
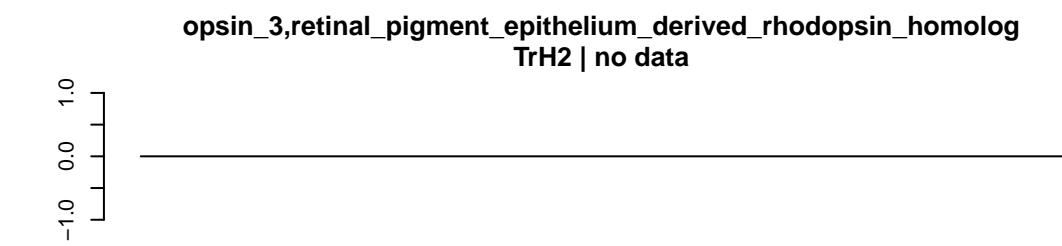
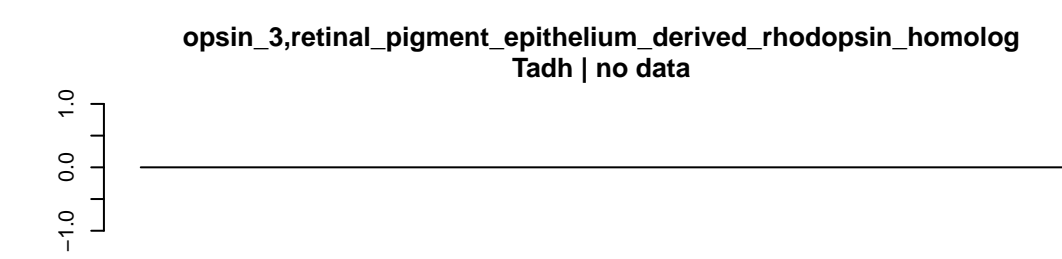


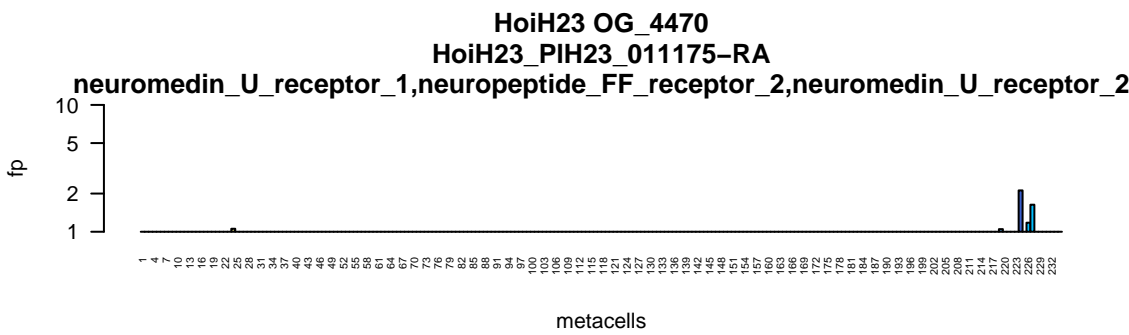
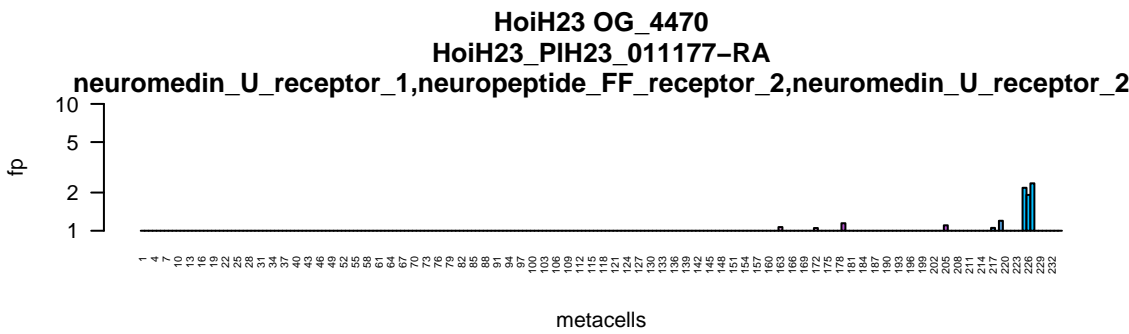
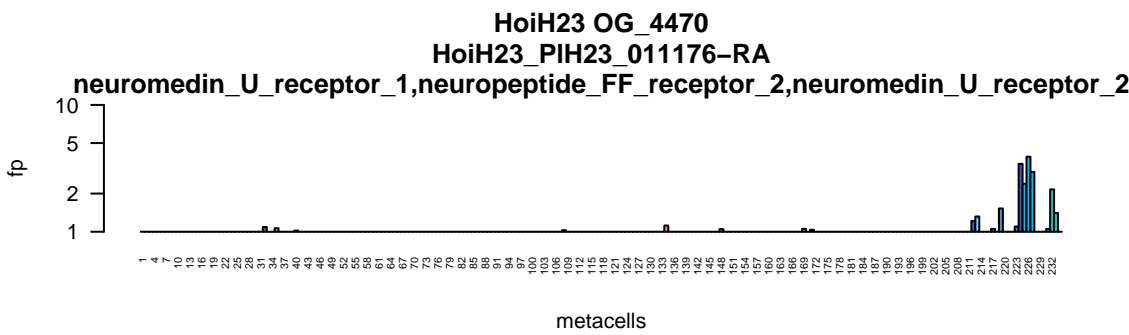
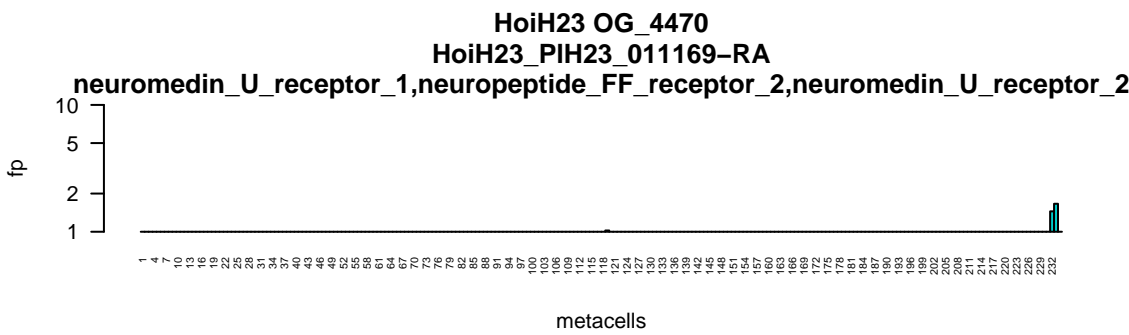
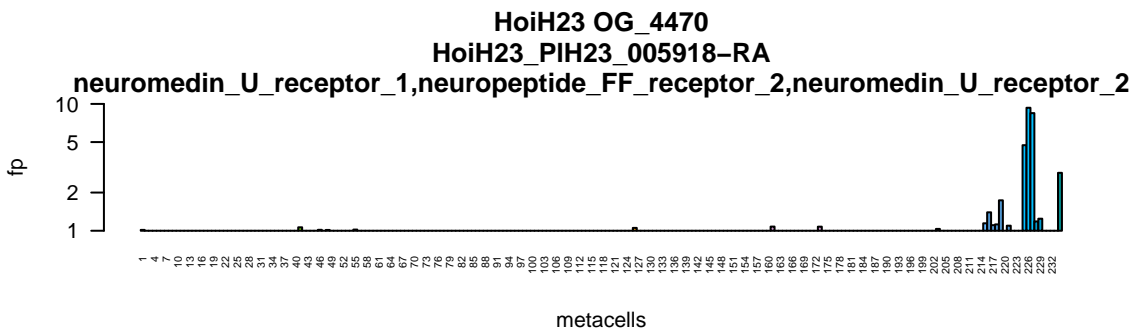
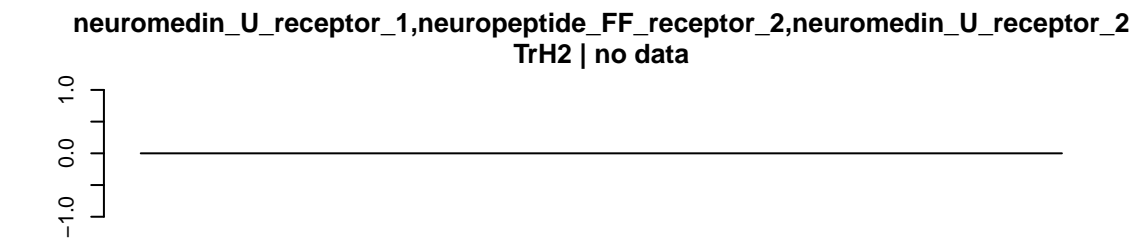
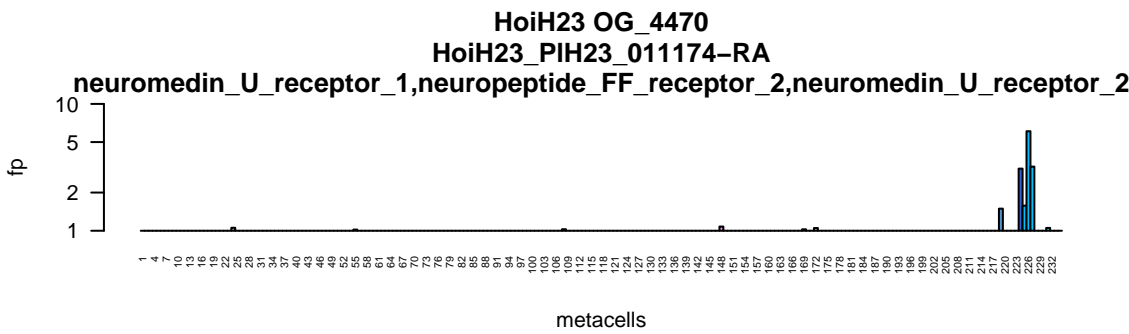
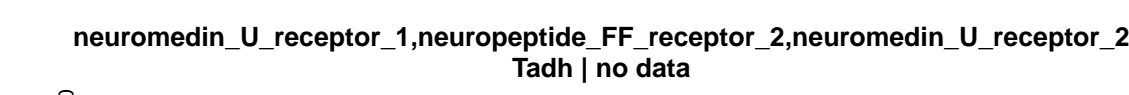


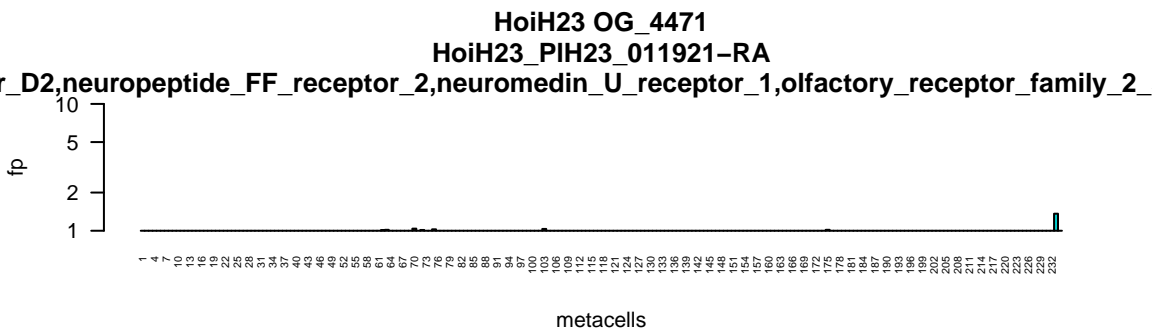
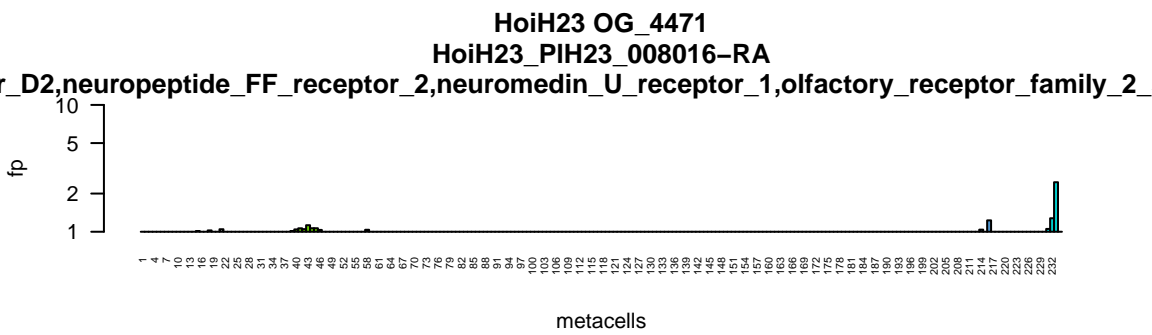
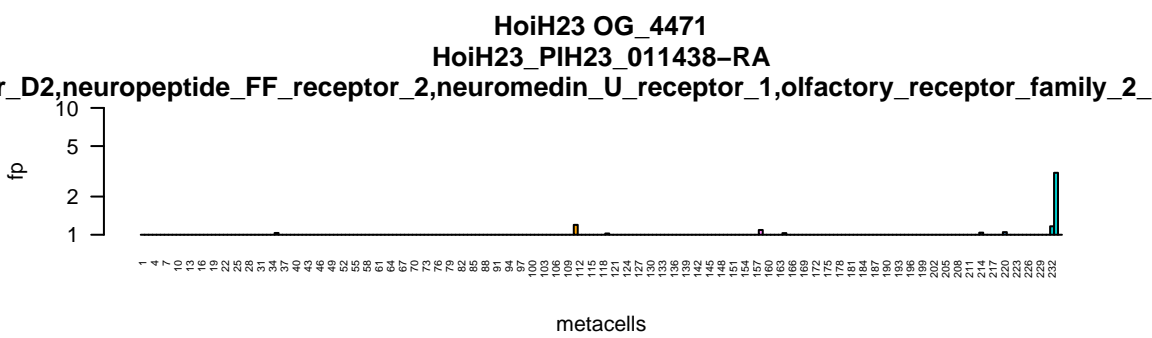
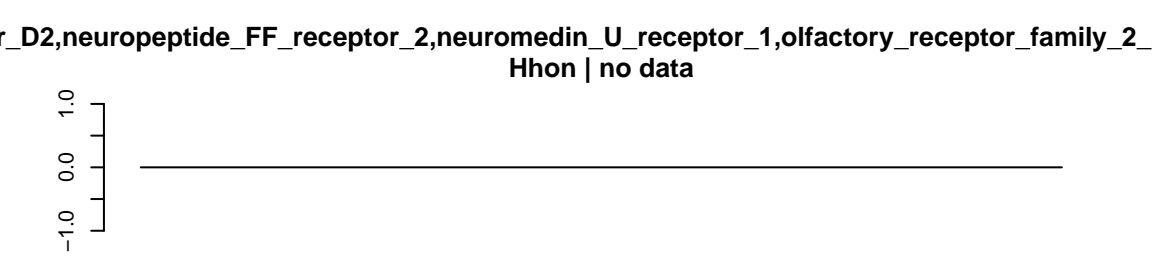
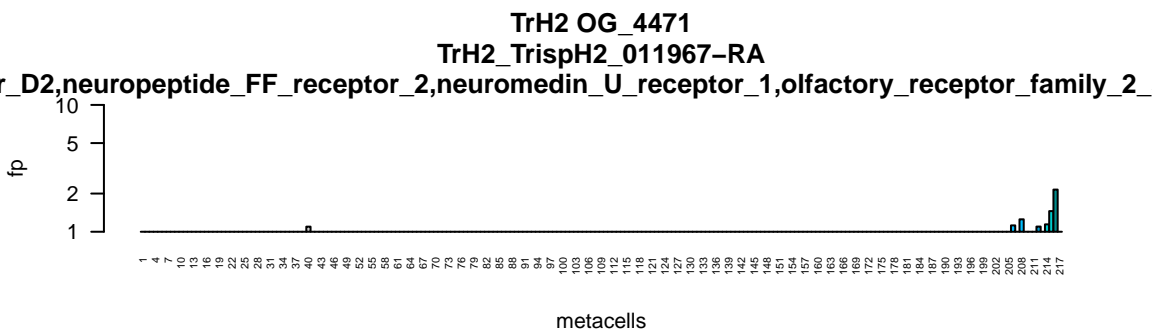
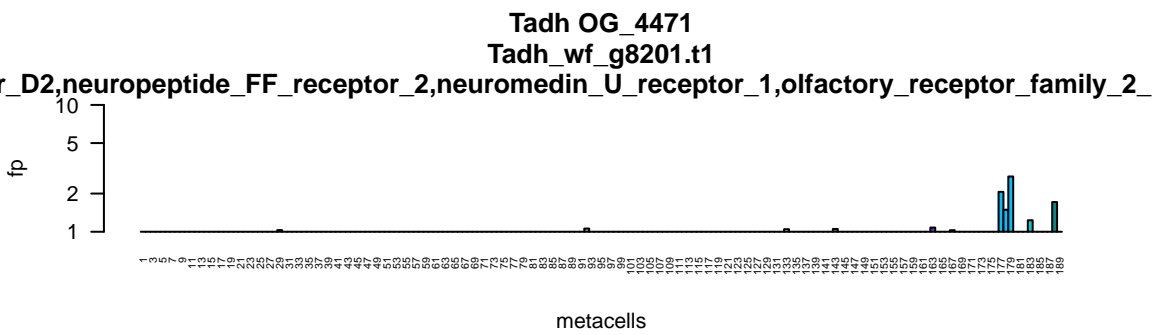
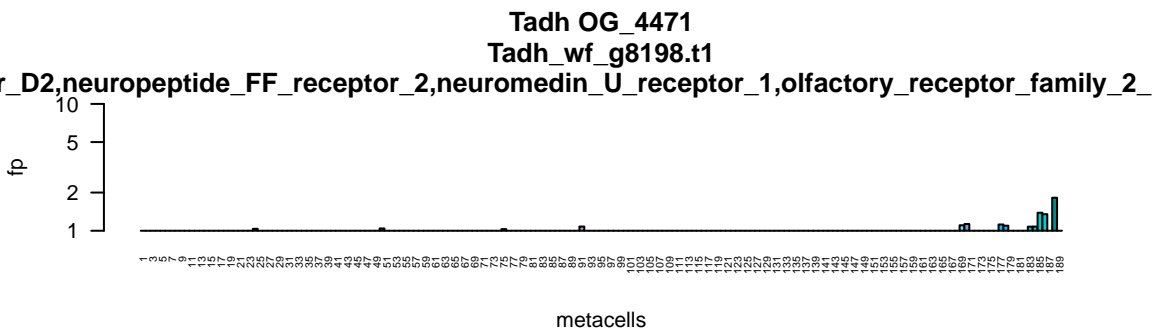




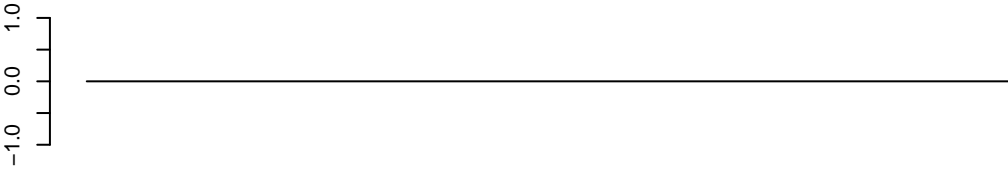








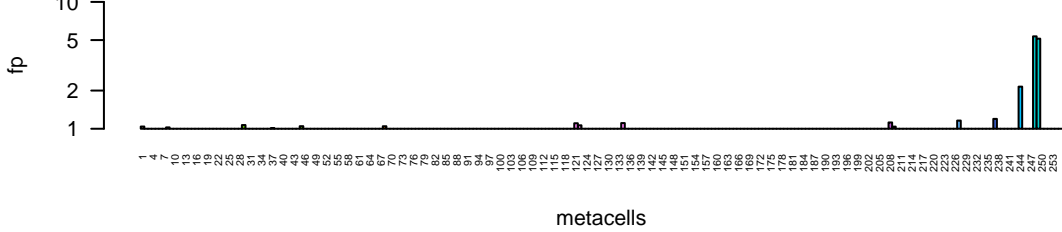
5\_hydroxytryptamine\_receptor\_7  
Tadh | no data



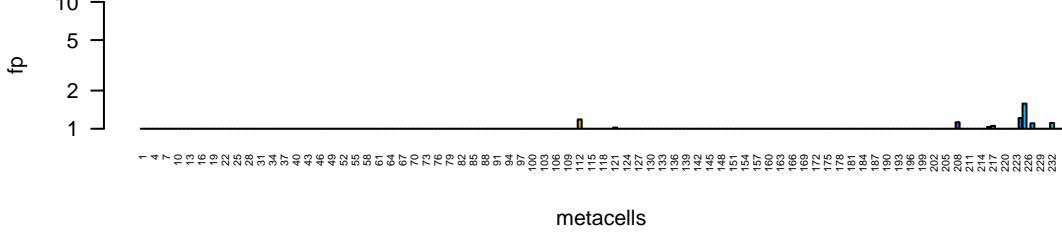
5\_hydroxytryptamine\_receptor\_7  
TrH2 | no data

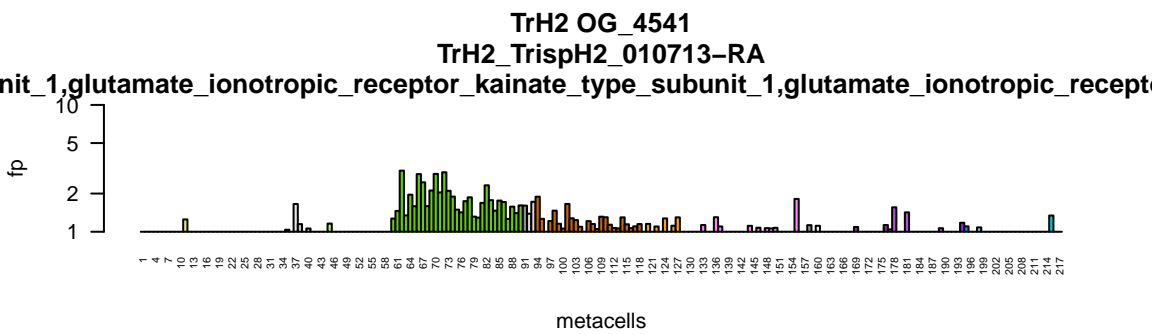
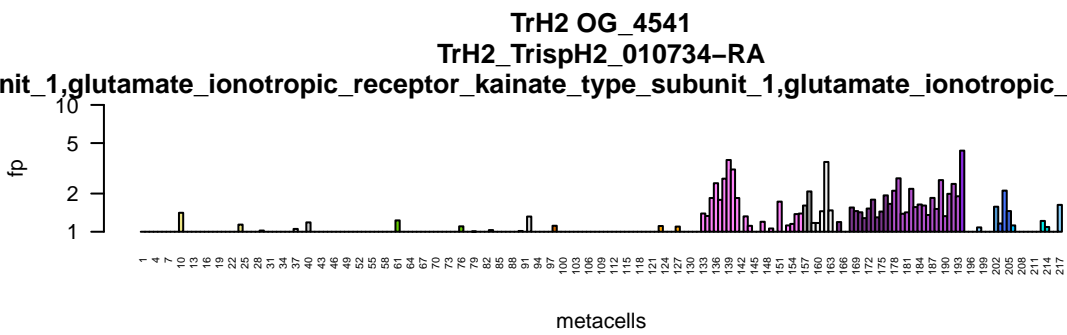
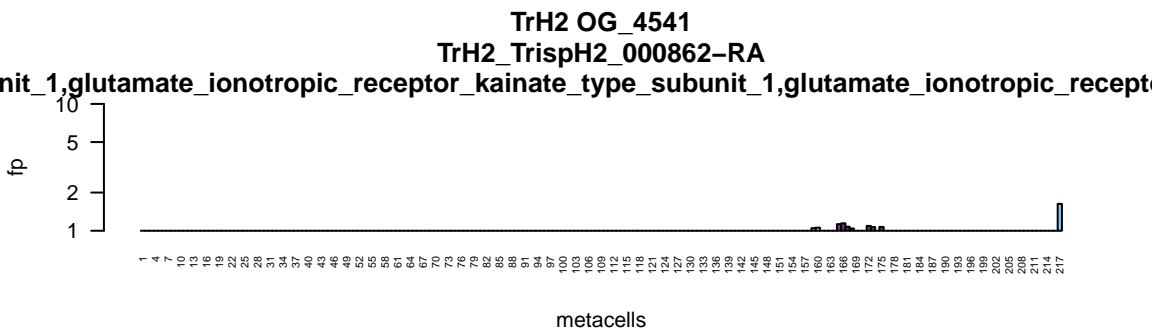
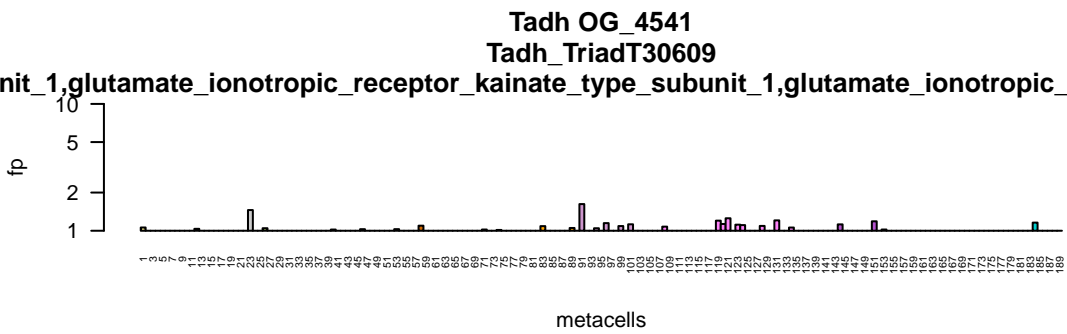
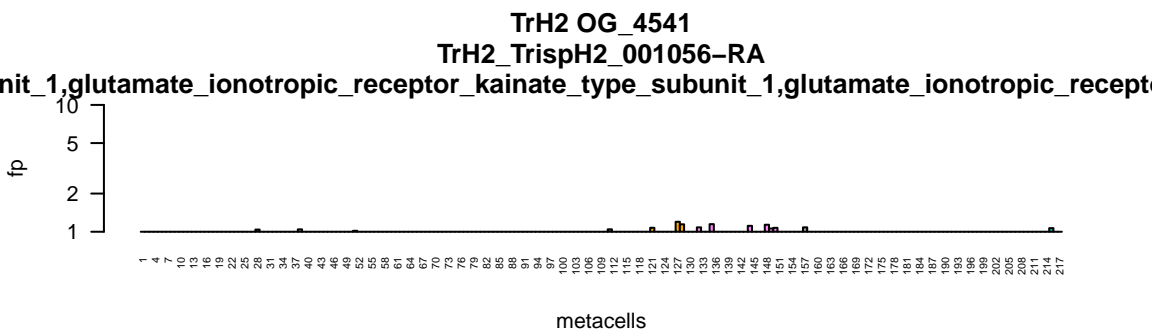
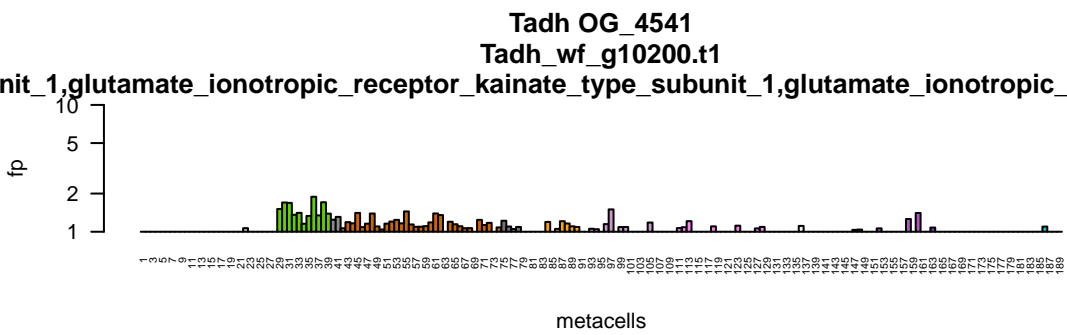
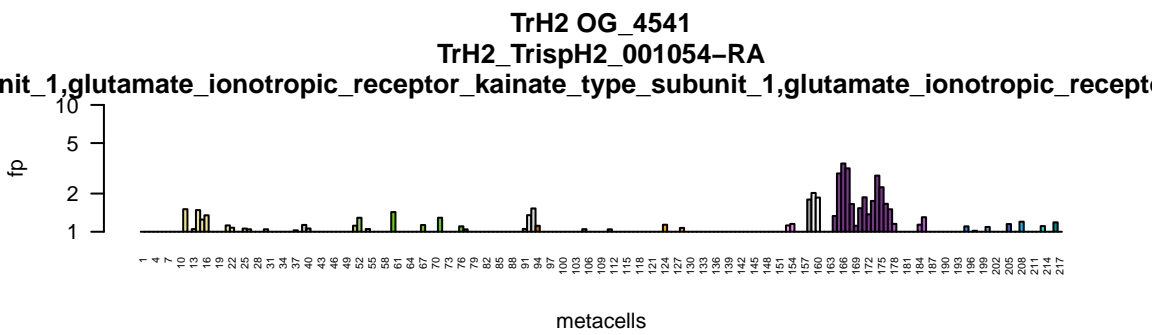
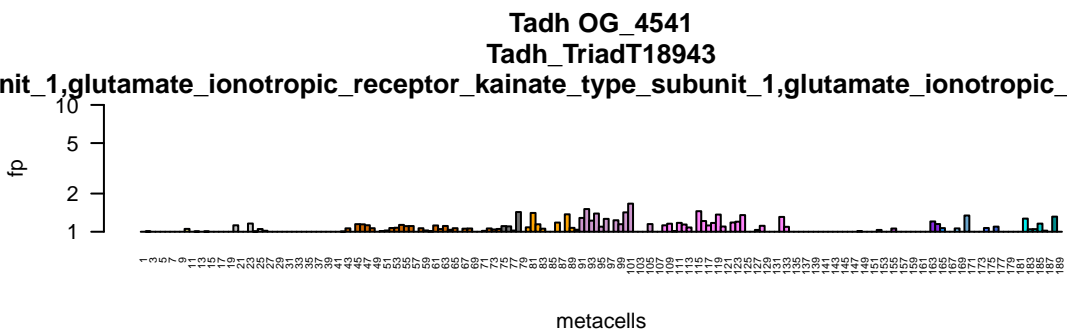
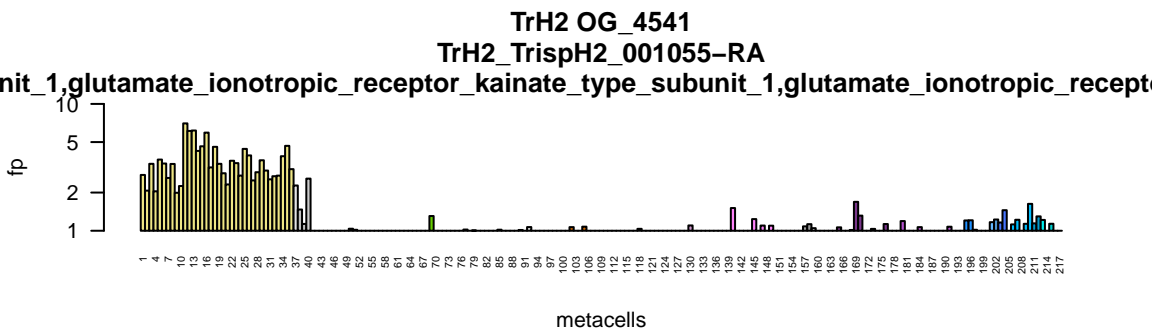
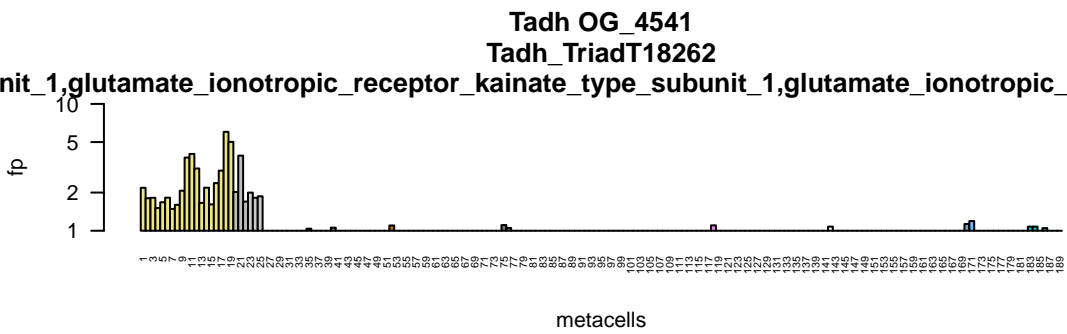
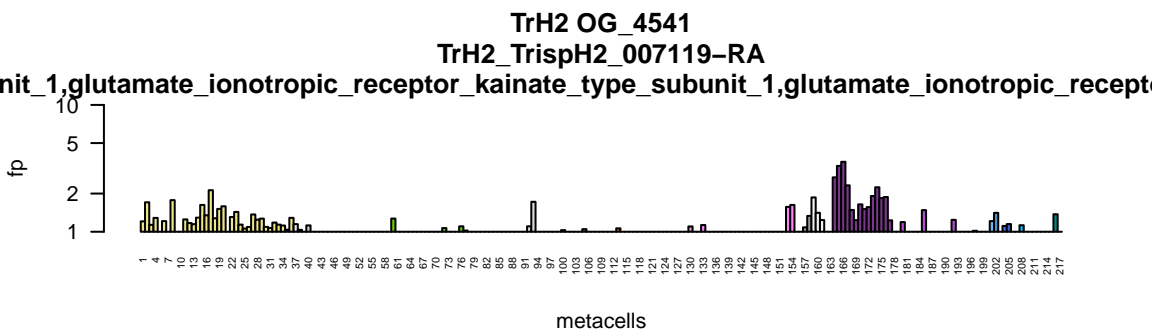
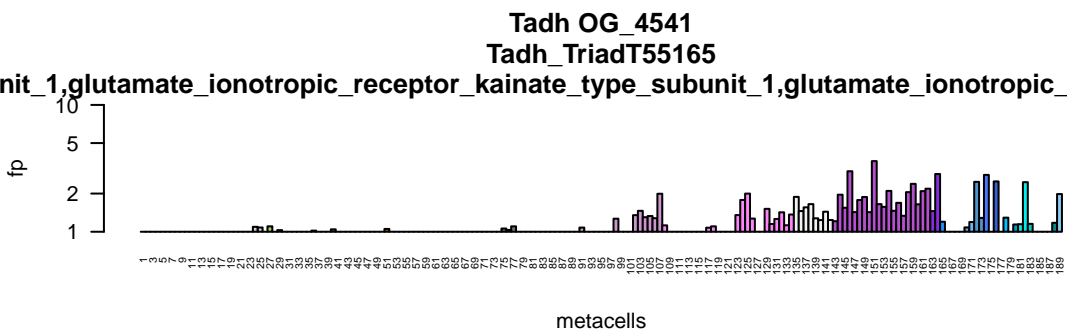
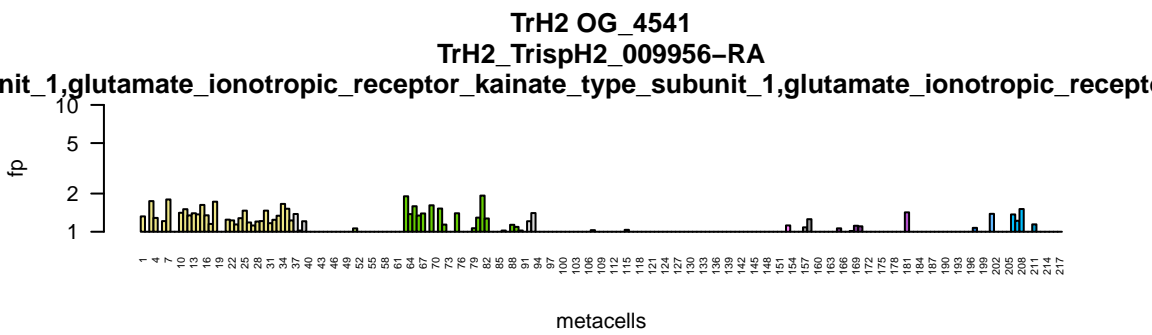
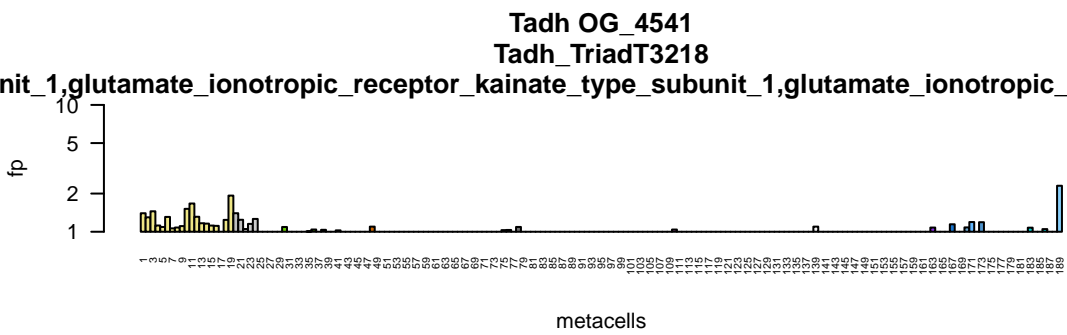
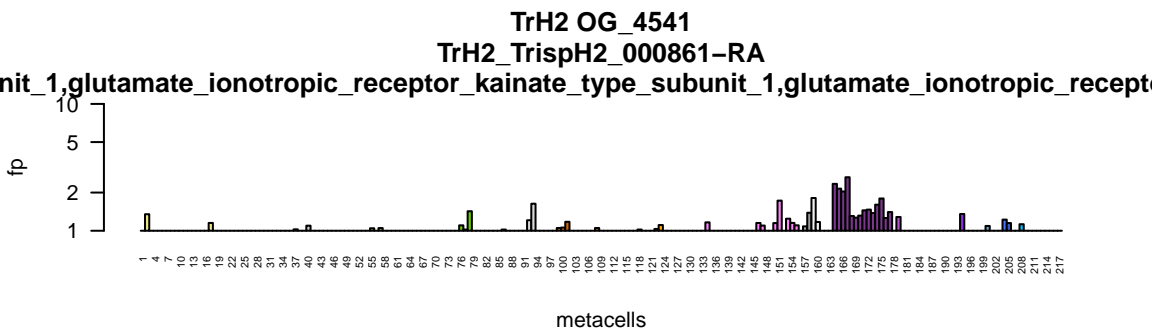
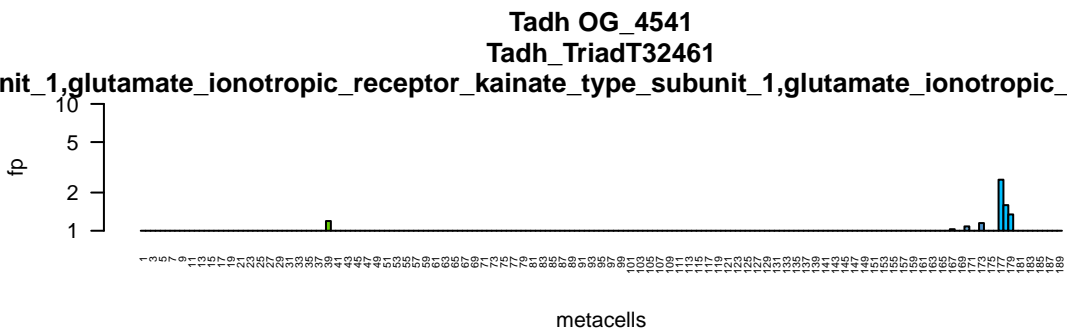


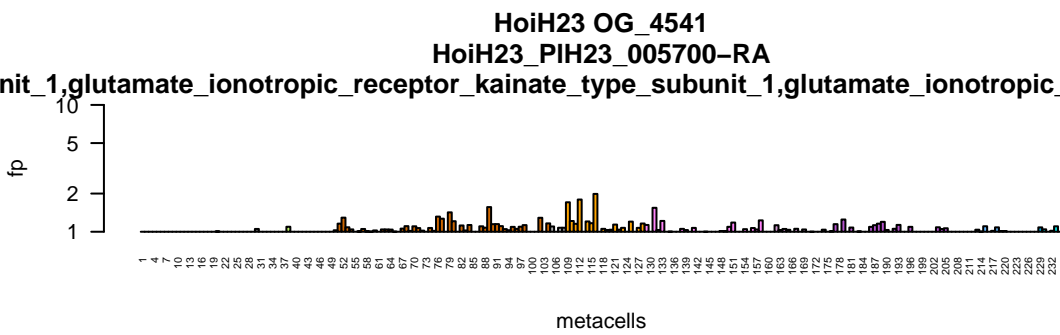
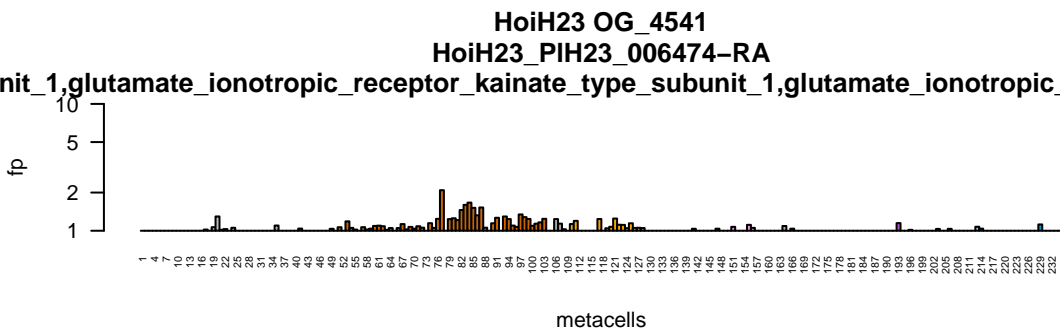
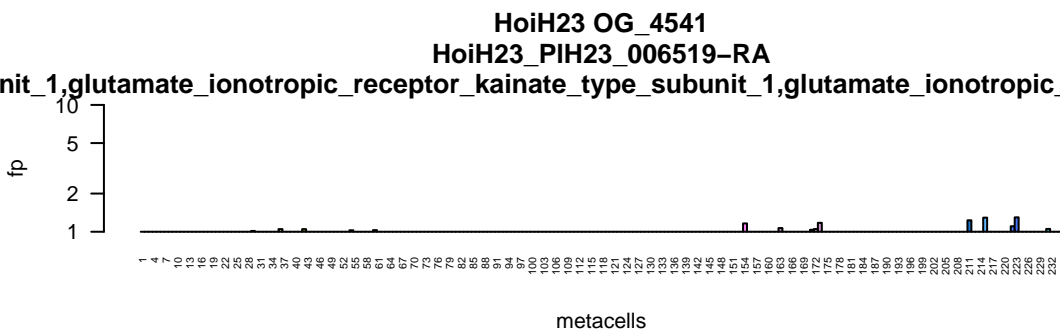
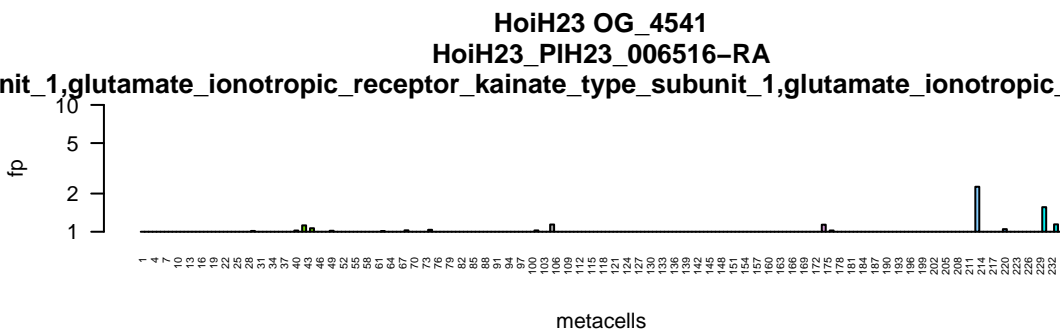
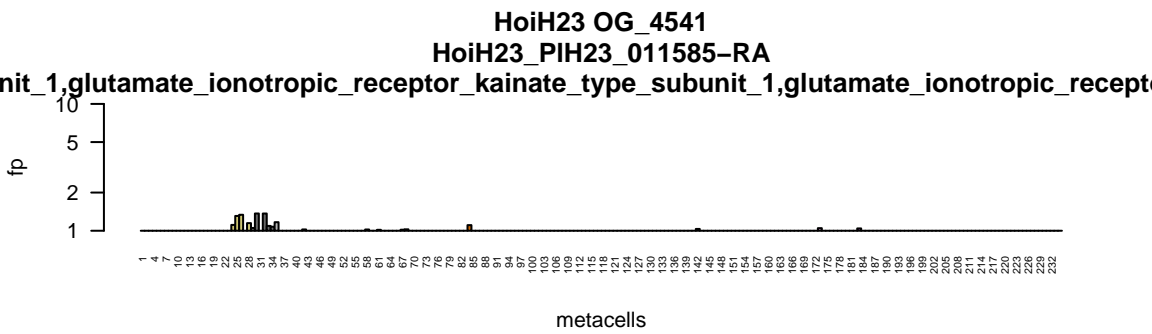
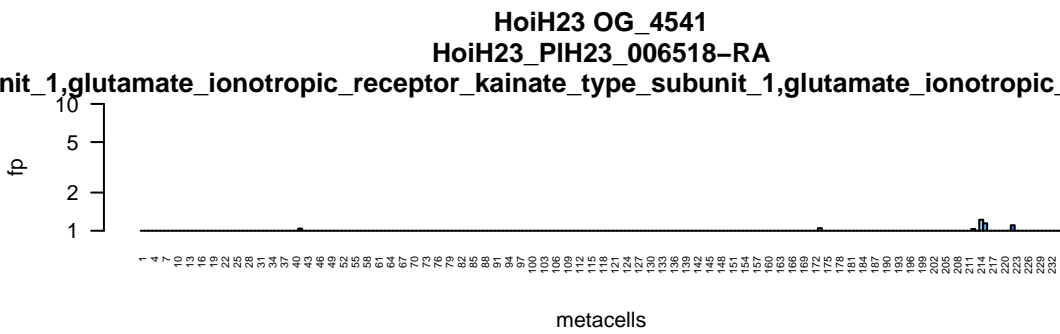
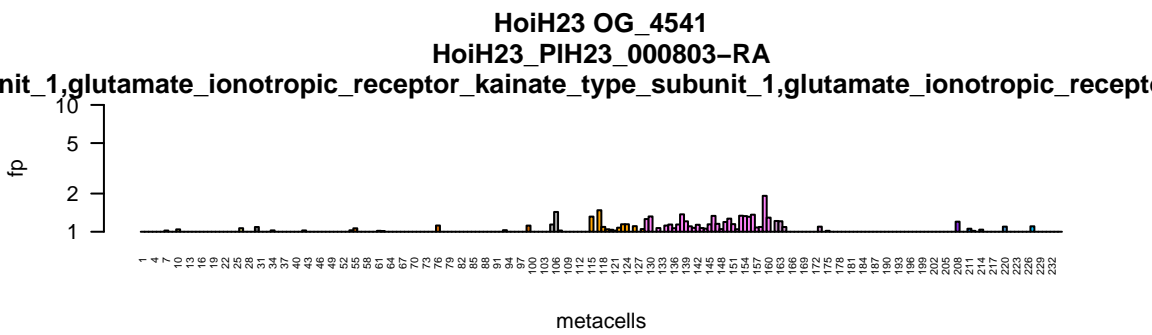
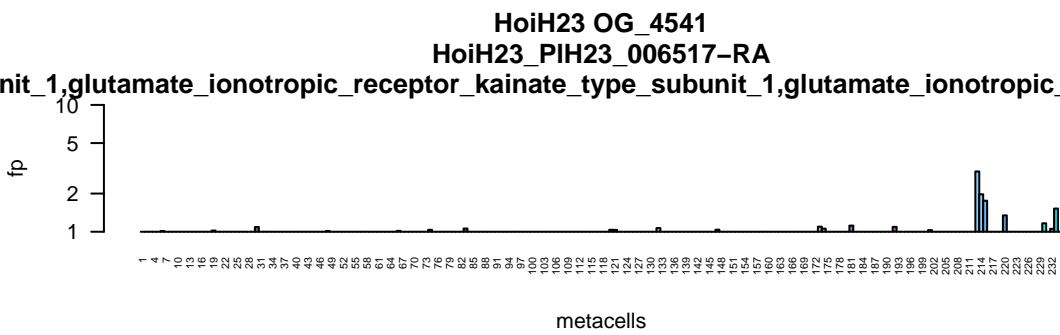
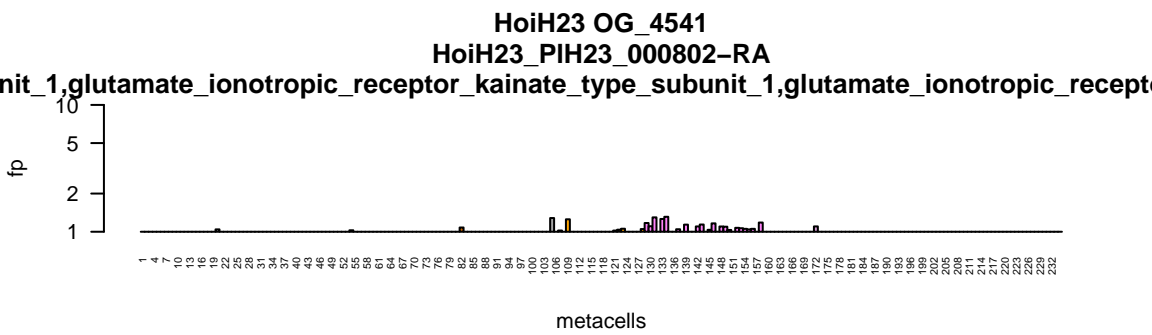
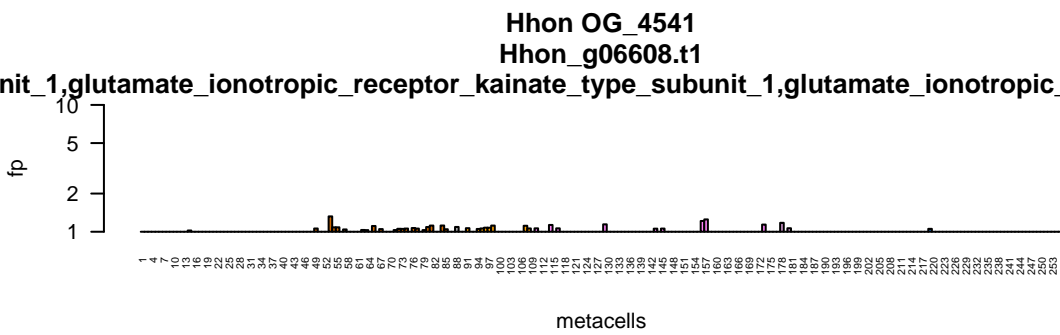
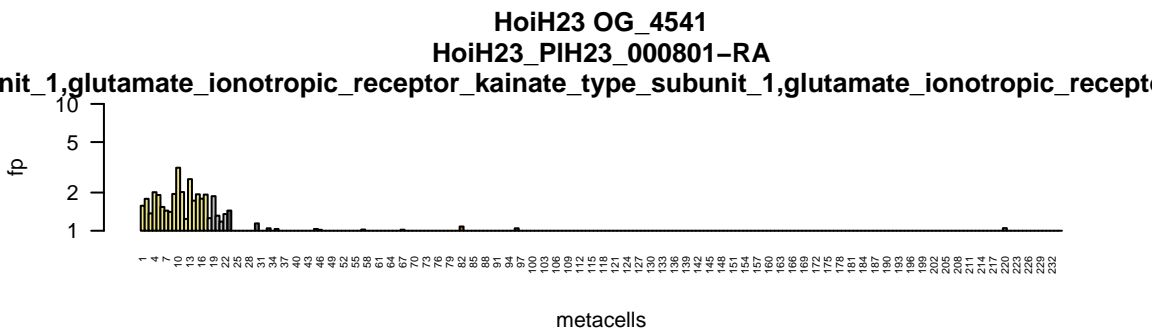
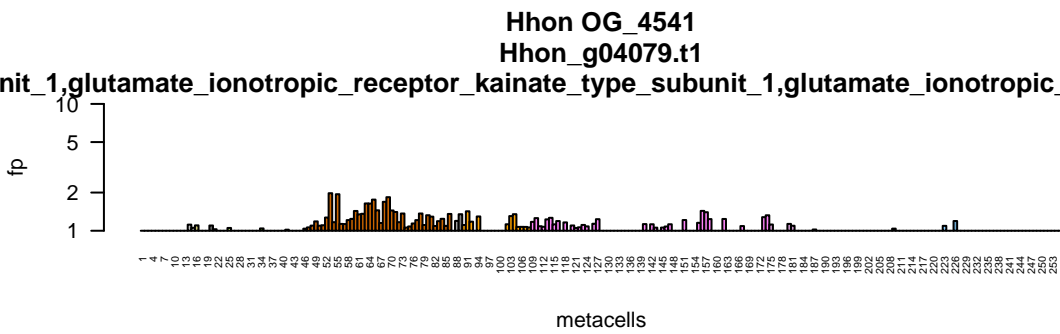
Hhon OG\_4530  
Hhon\_g10440.t1  
5\_hydroxytryptamine\_receptor\_7

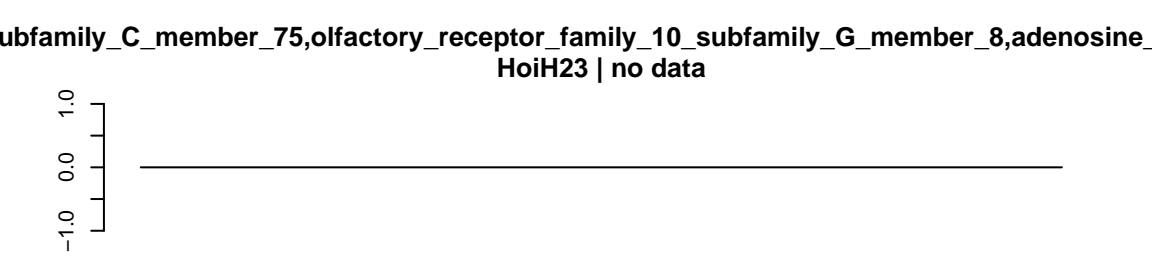
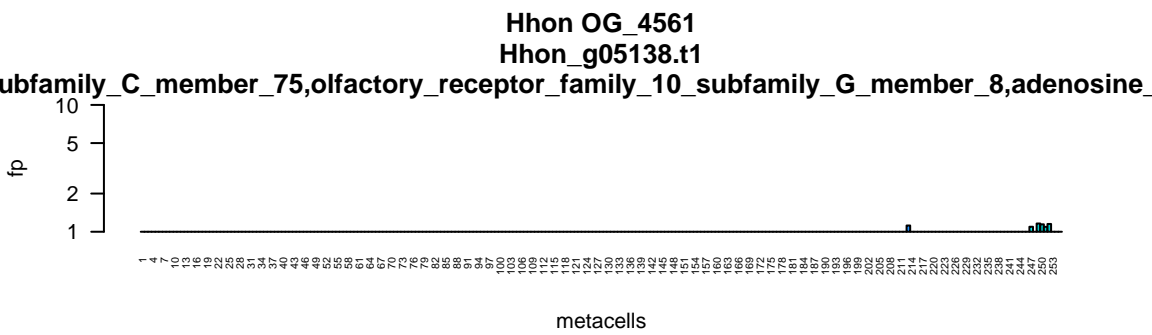
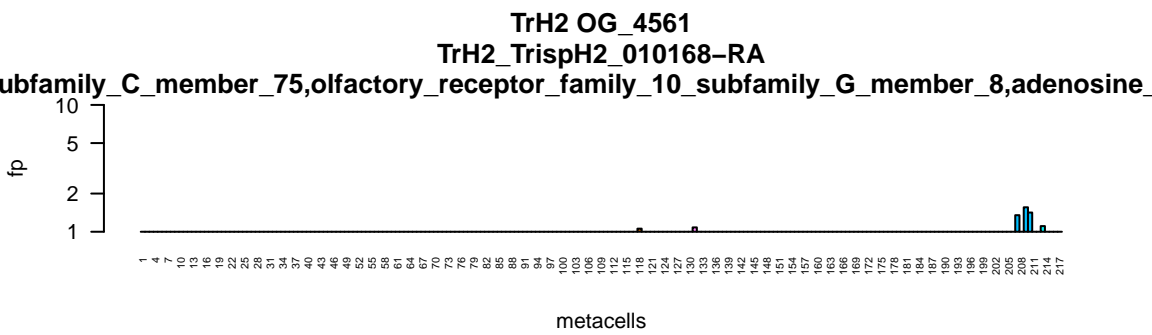
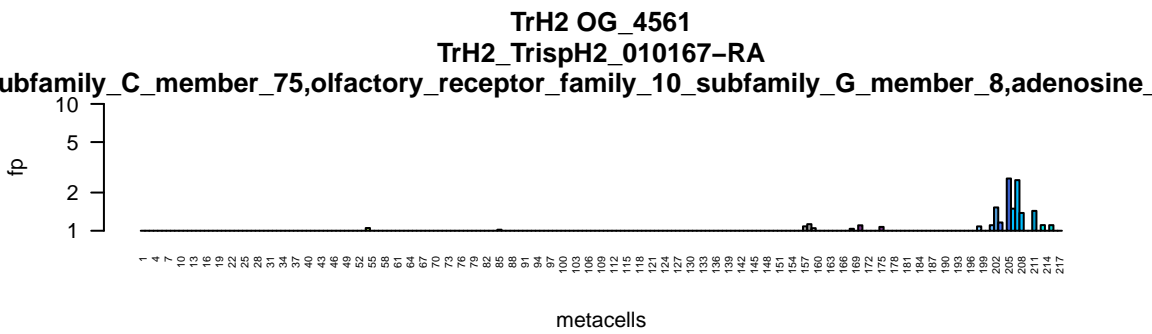
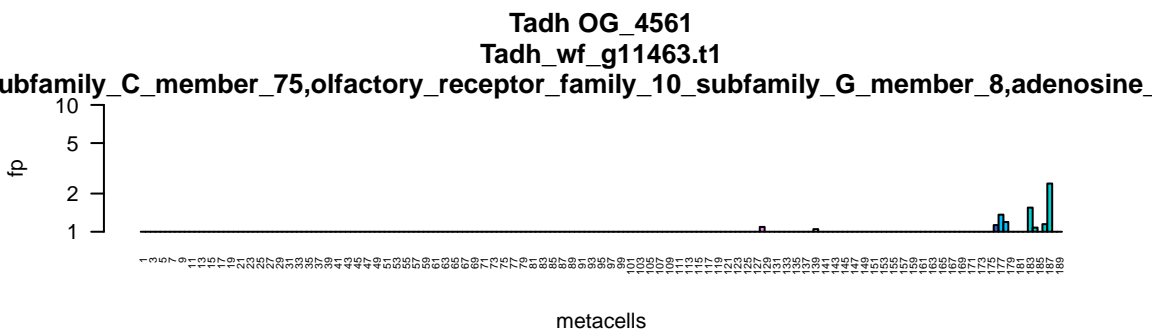
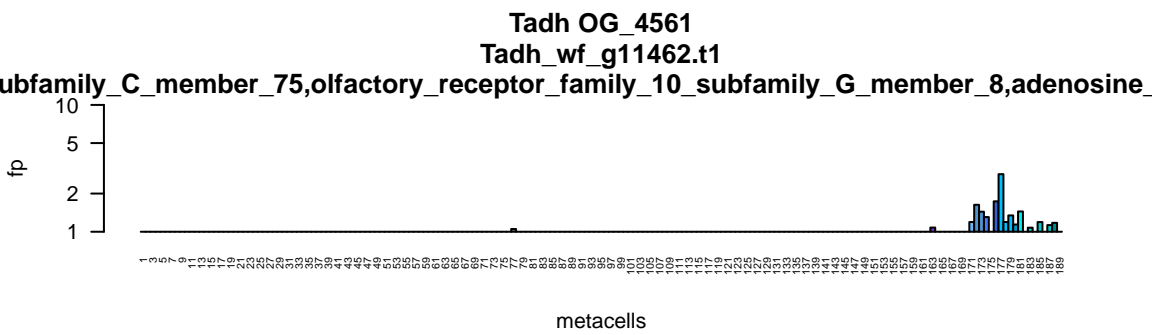
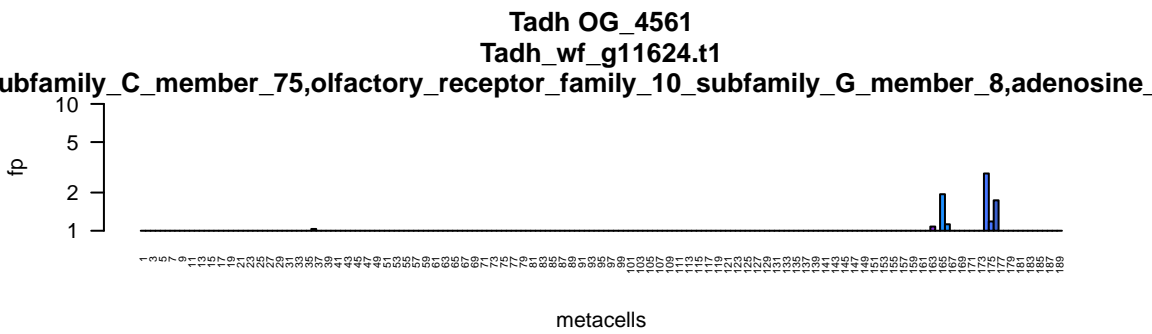


HoiH23 OG\_4530  
HoiH23\_PIH23\_011877-RA  
5\_hydroxytryptamine\_receptor\_7









TrH2 OG\_4762  
TrH2\_TrispH2\_010632-RA  
hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chemokine

fp

metacells

TrH2 OG\_4762  
TrH2\_TrispH2\_010635-RA  
hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chemokine

metacells	fp
1	1
199	2
214	2

TrH2 OG\_4762  
TrH2\_TrispH2\_010271-RA

hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chemokine

fp

metacells

hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chemokine

fp

metacells

Hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chemokine

metacell	fp
1	1
4	1
10	1
13	1
19	1
22	1
28	1
31	1
37	1
43	1
46	1
52	1
55	2
61	1
64	1
70	1
73	1
79	1
82	1
88	1
91	1
97	1
100	1
106	1
108	1
109	1
115	1
118	1
127	1
133	1
136	1
145	1
146	1
151	1
154	1
160	1
163	1
169	1
175	1
178	1
181	1
184	1
193	1
196	1
202	1
205	1
211	1
214	1
220	1
223	1
229	2
232	1
238	1
241	1.5
247	1
250	1
253	1

Hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chemokine

metacells

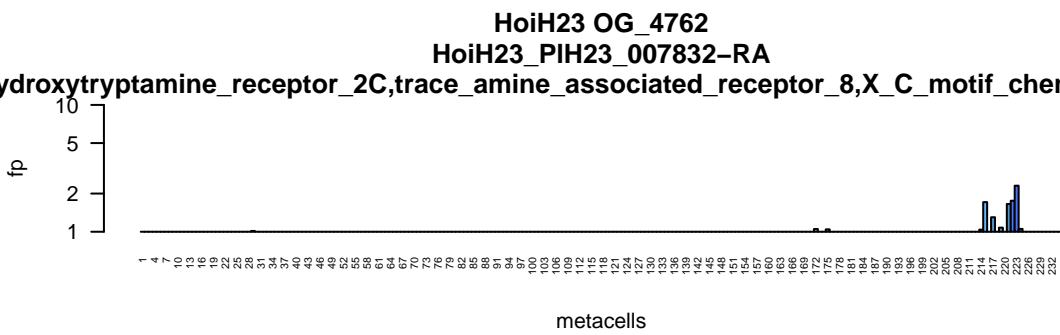
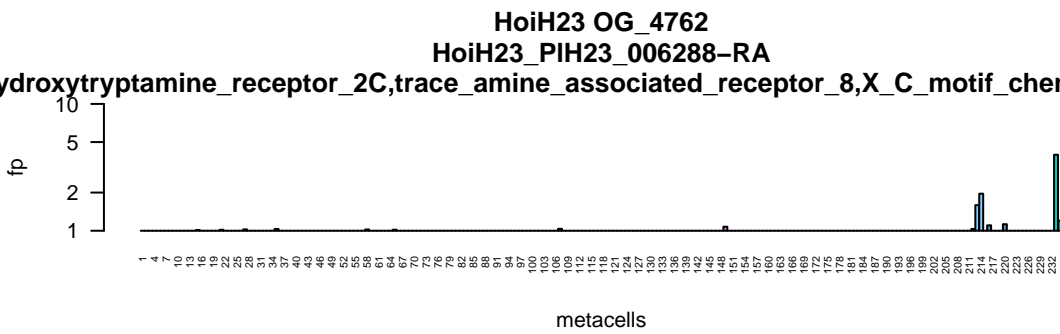
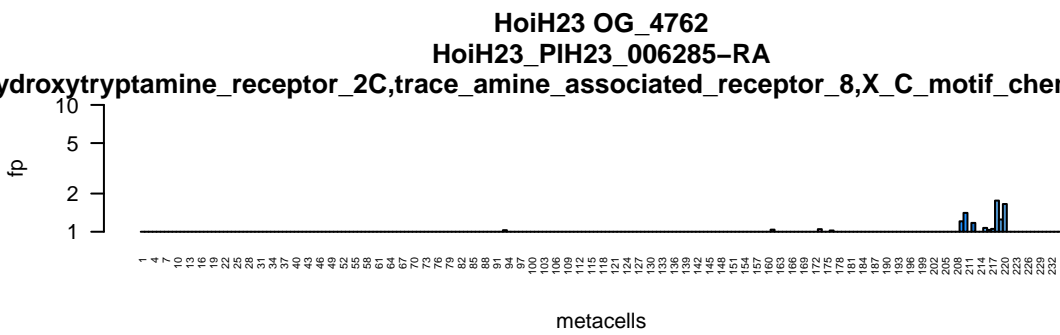
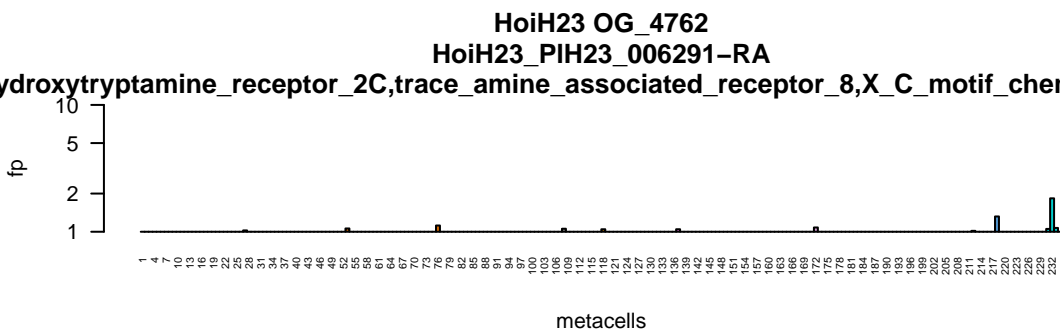
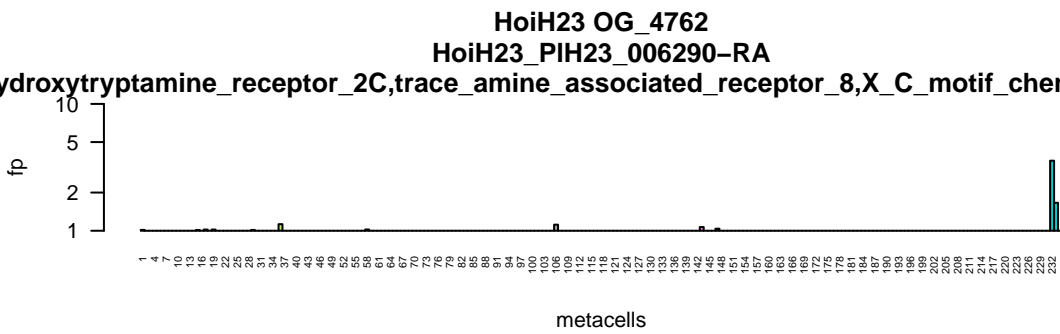
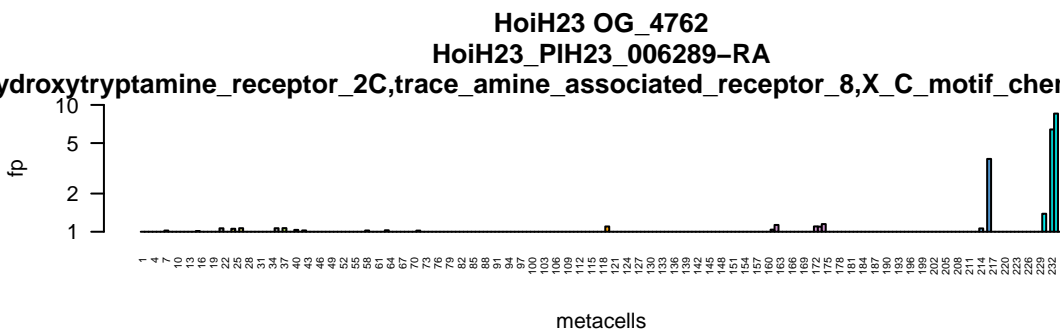
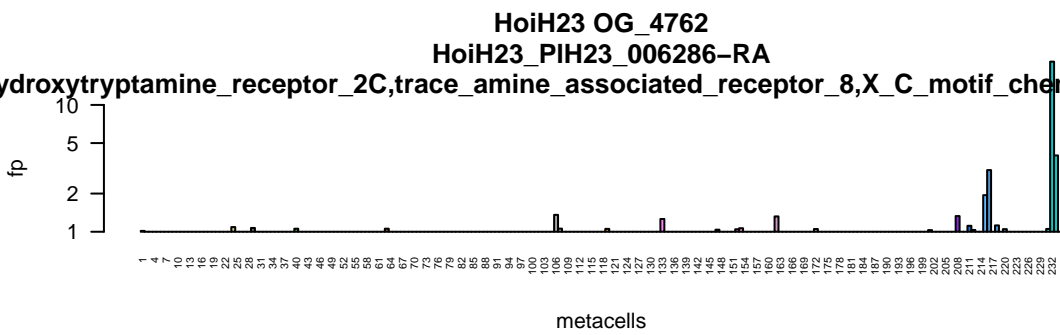
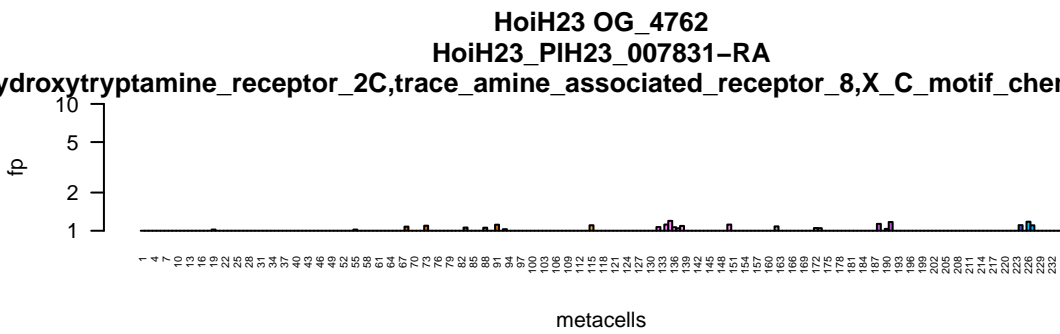
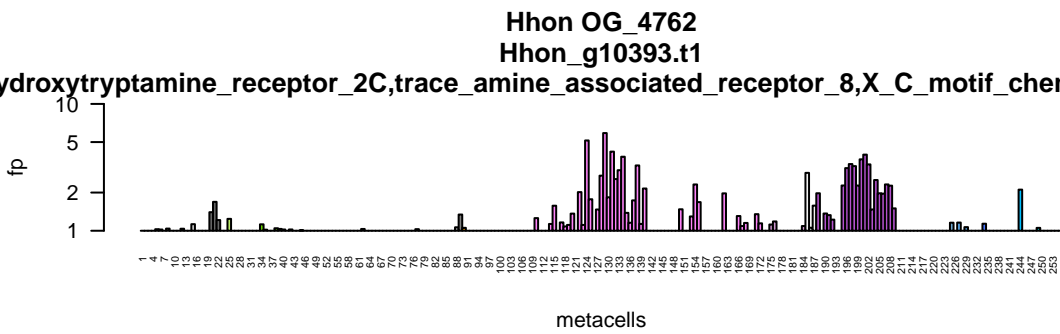
Hhon OG\_4762  
Hhon\_g10395.t1

hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chemokine

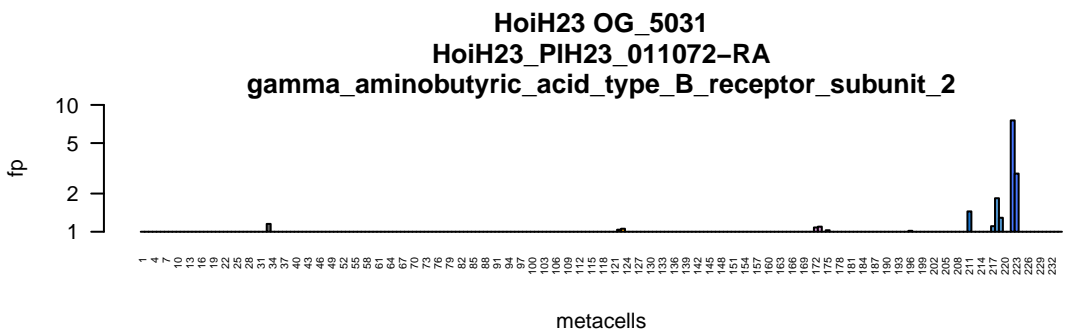
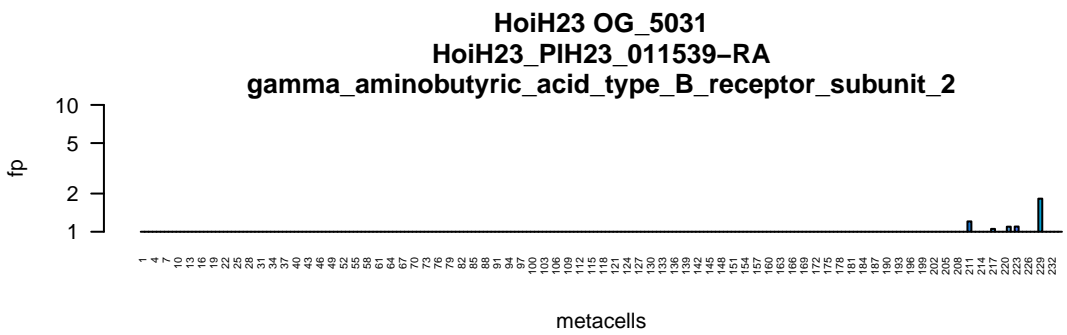
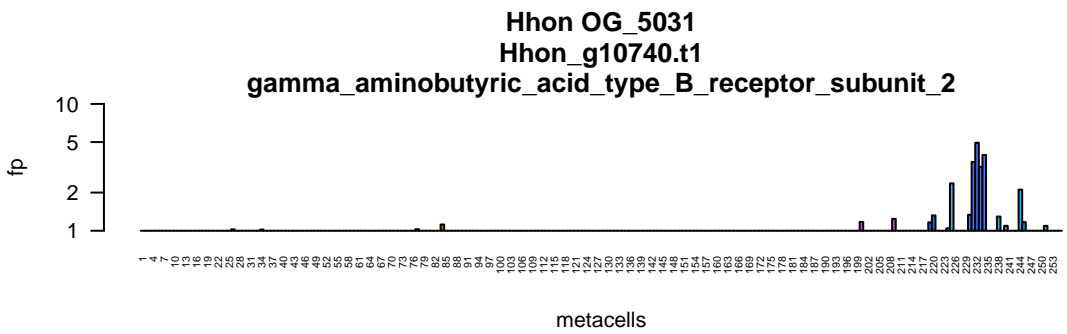
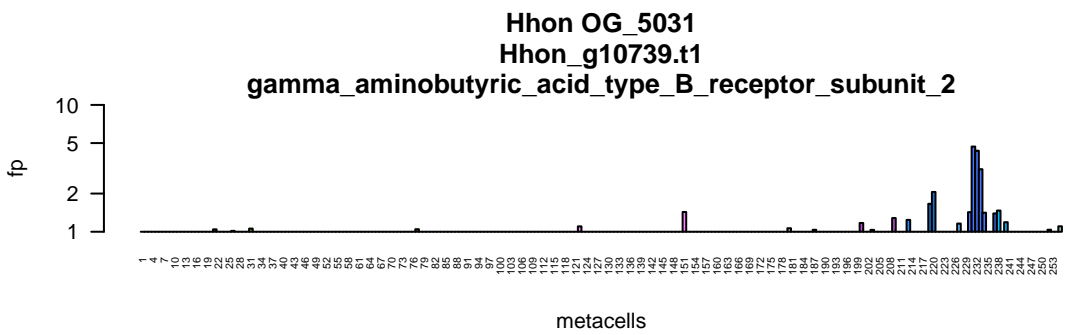
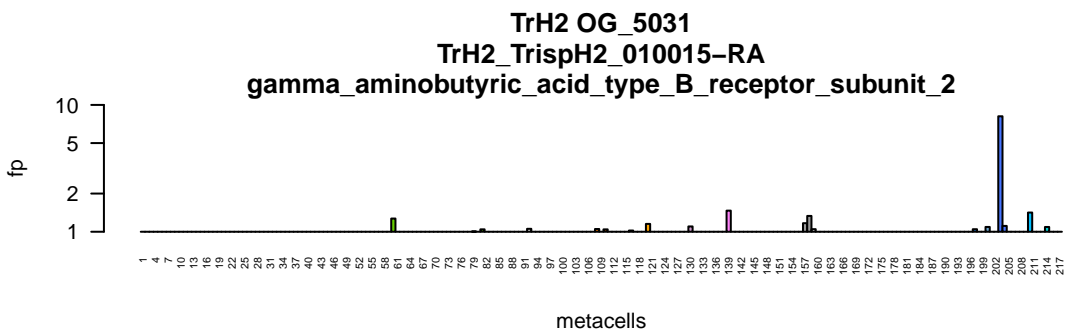
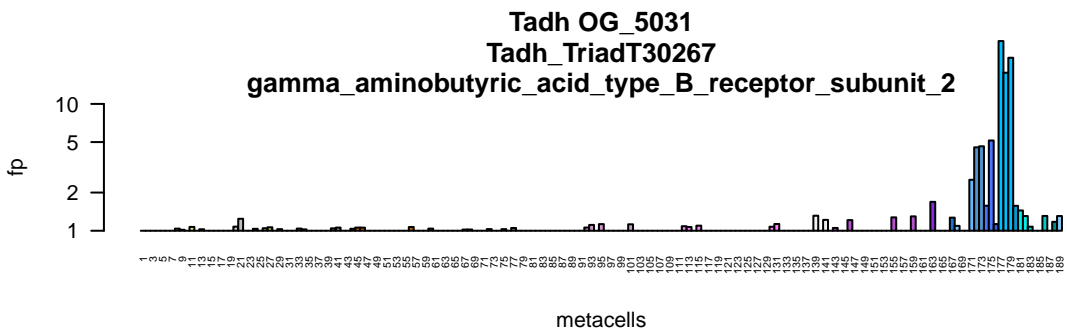
fp

metacells

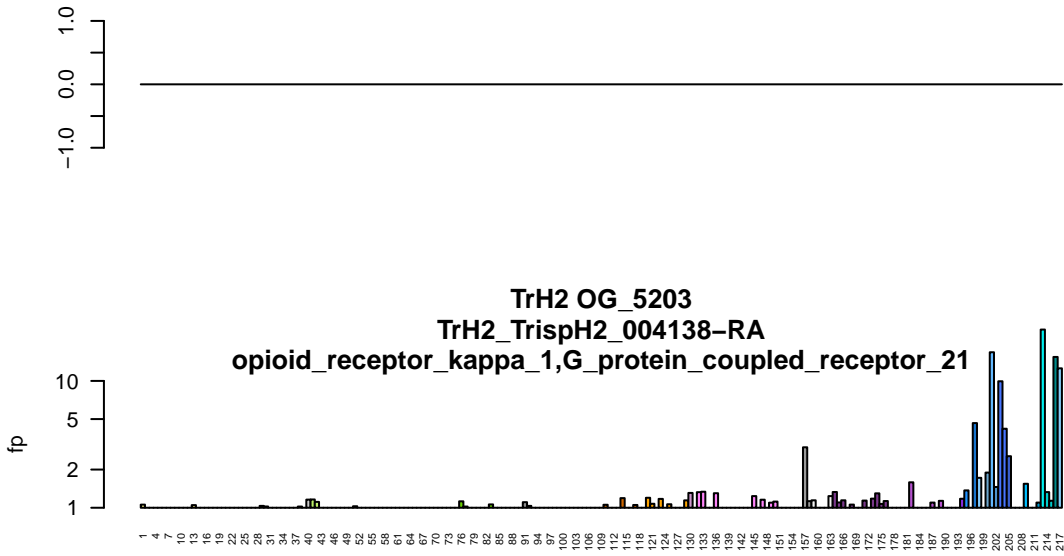




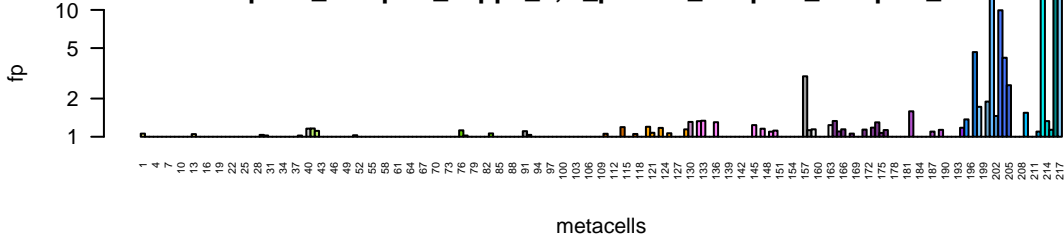




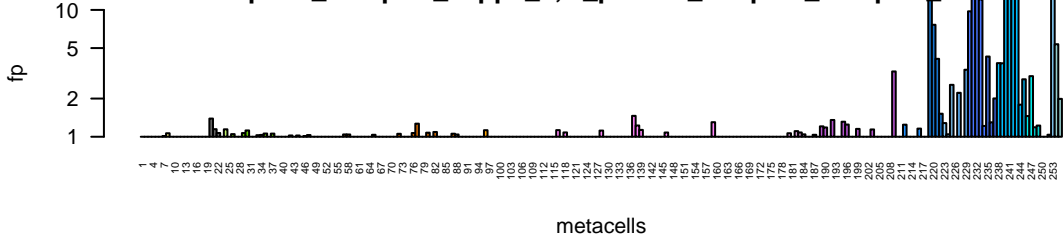
opioid\_receptor\_kappa\_1,G\_protein\_coupled\_receptor\_21  
Tadh | no data



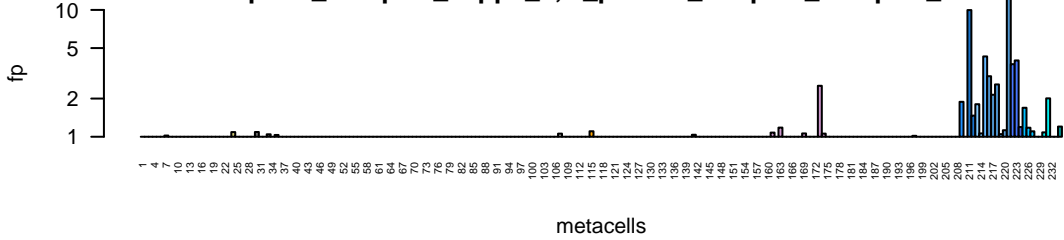
TrH2 OG\_5203  
TrH2\_TrispH2\_004138-RA  
opioid\_receptor\_kappa\_1,G\_protein\_coupled\_receptor\_21

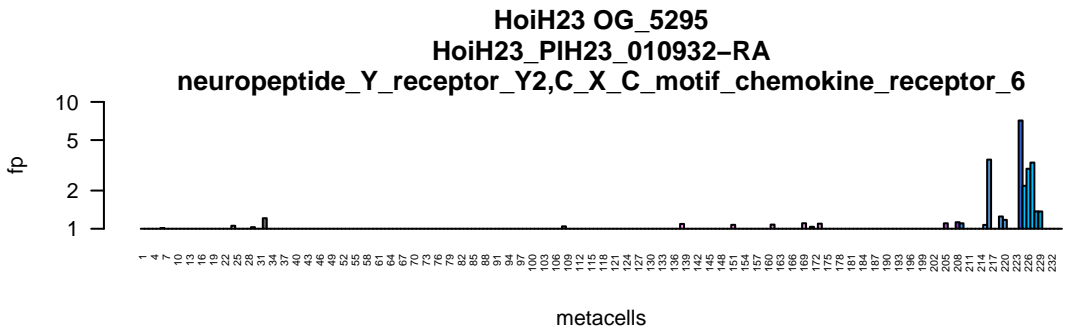
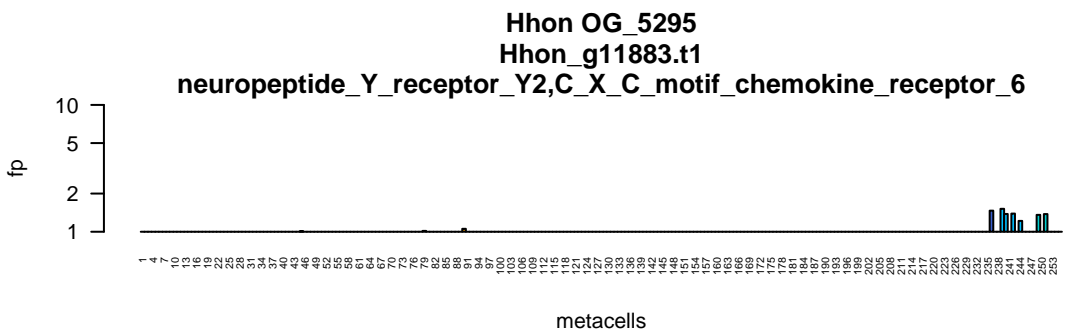
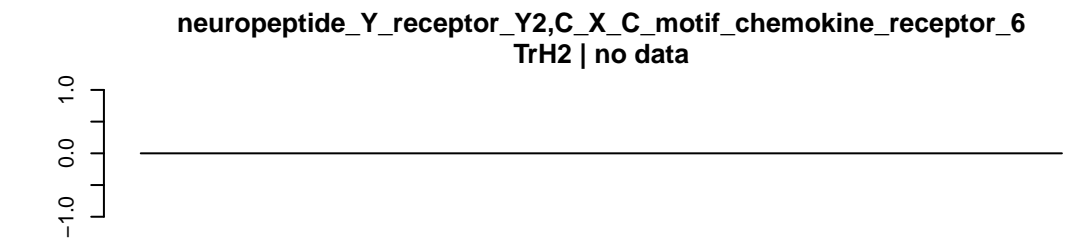
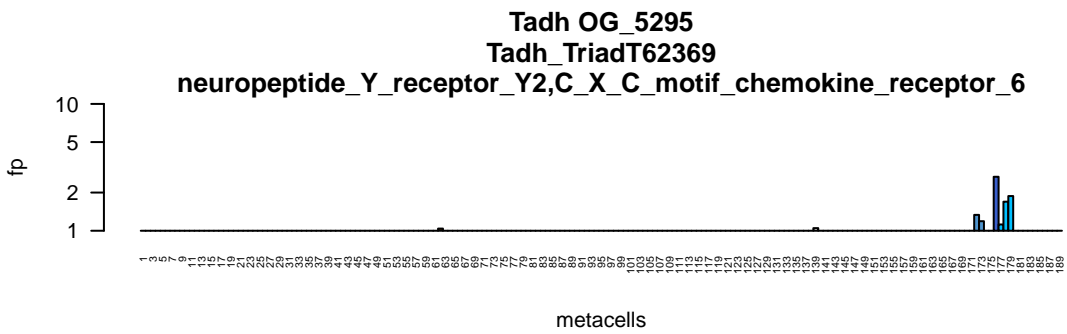


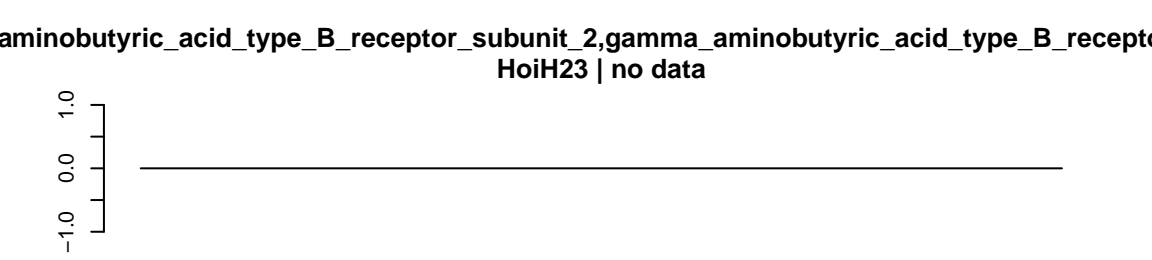
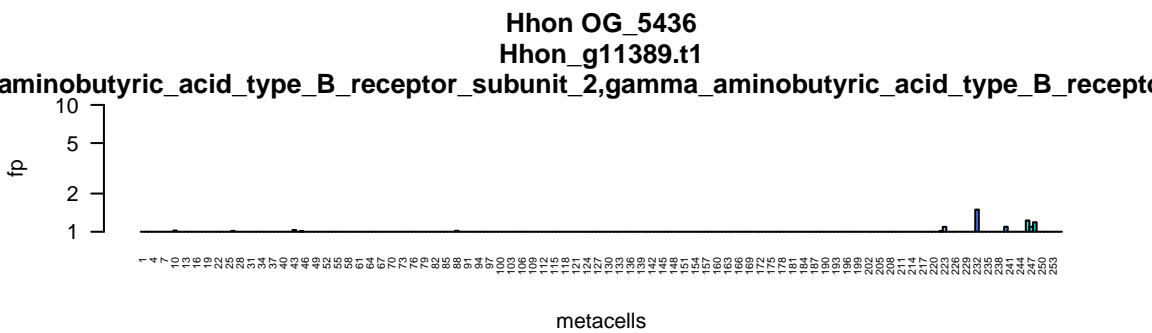
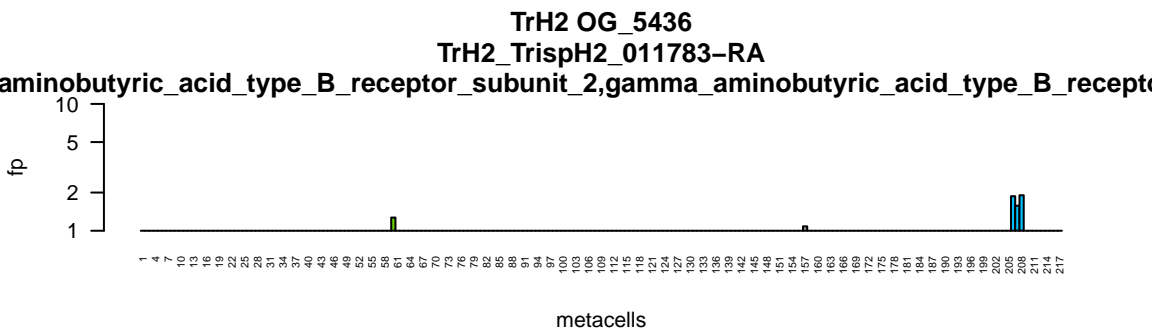
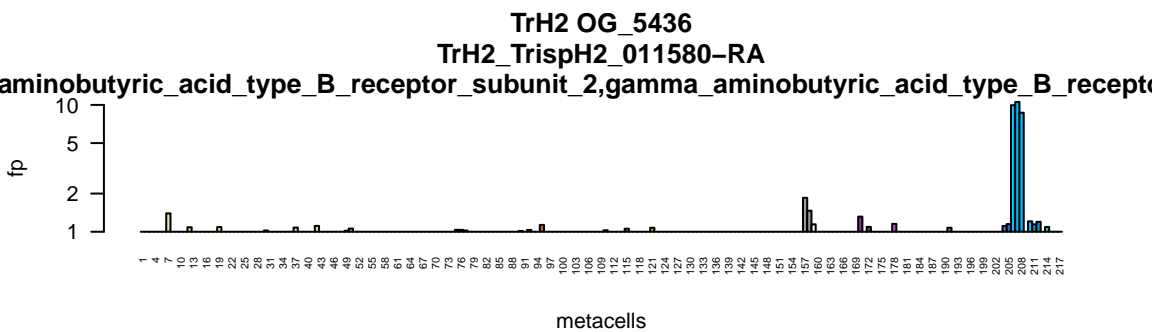
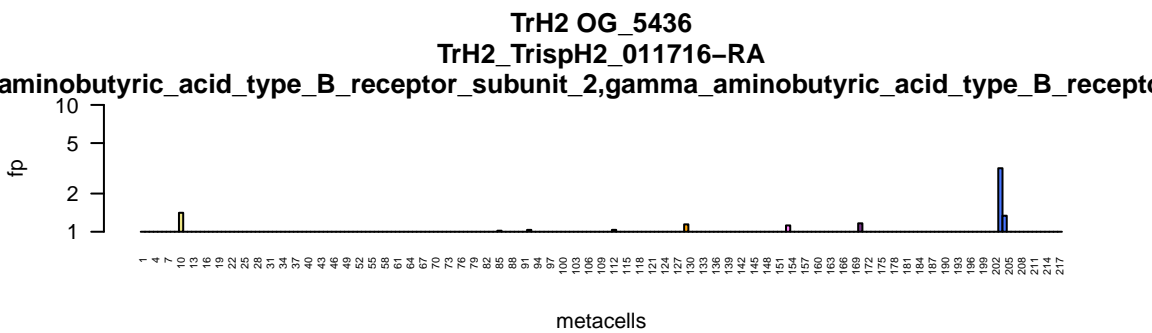
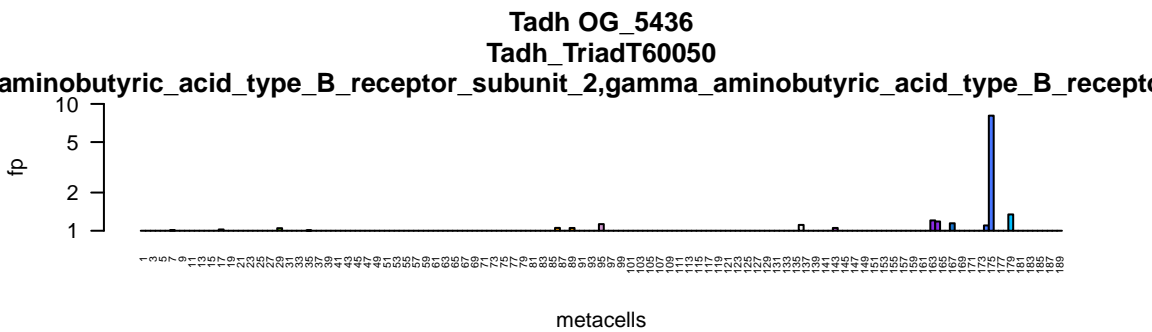
Hhon OG\_5203  
Hhon\_g01680.t1  
opioid\_receptor\_kappa\_1,G\_protein\_coupled\_receptor\_21

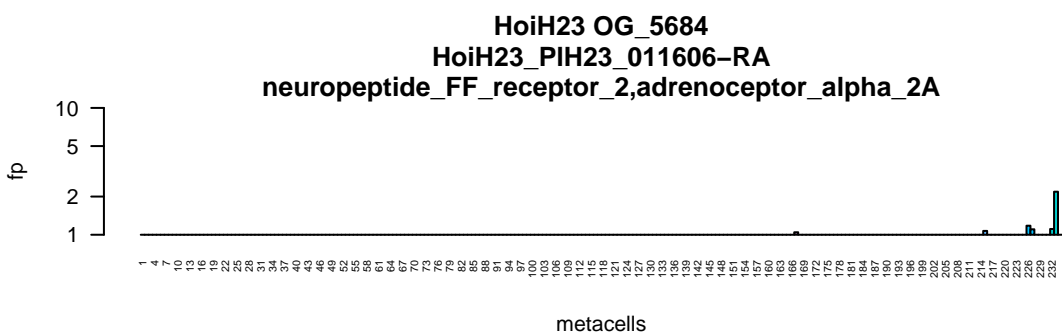
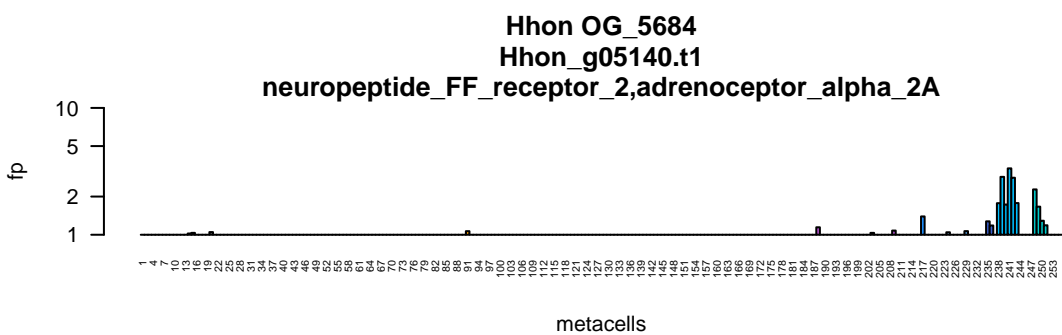
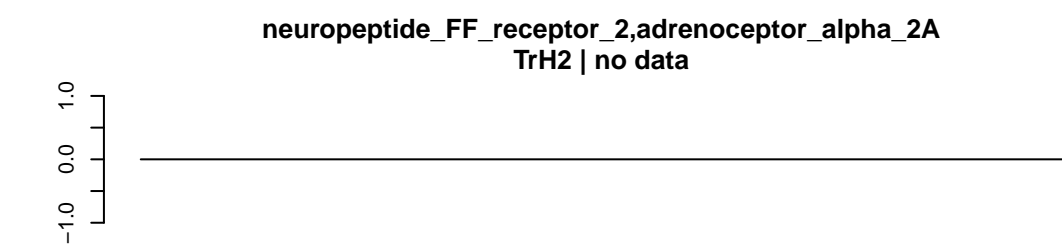
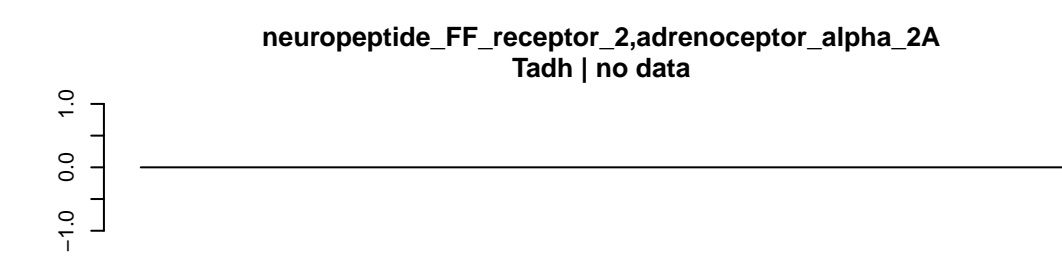


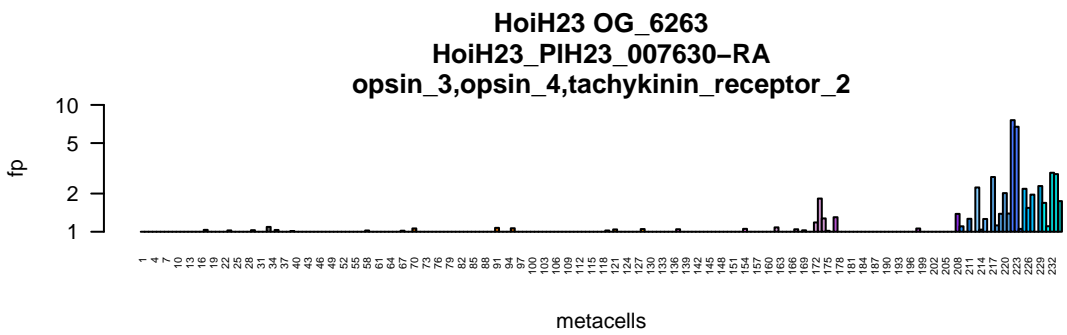
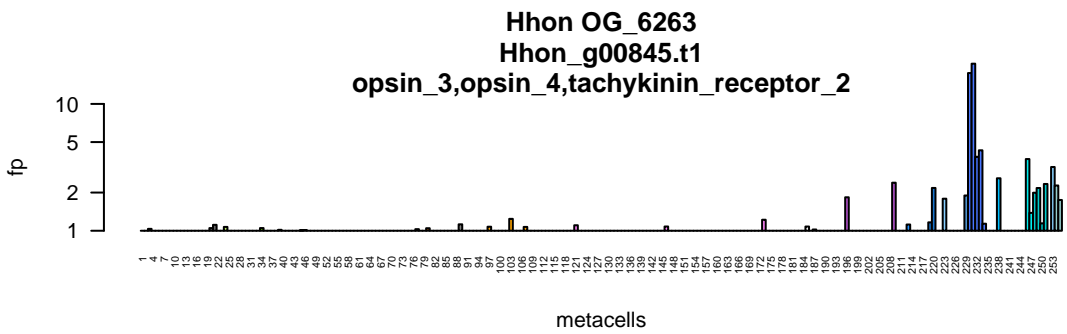
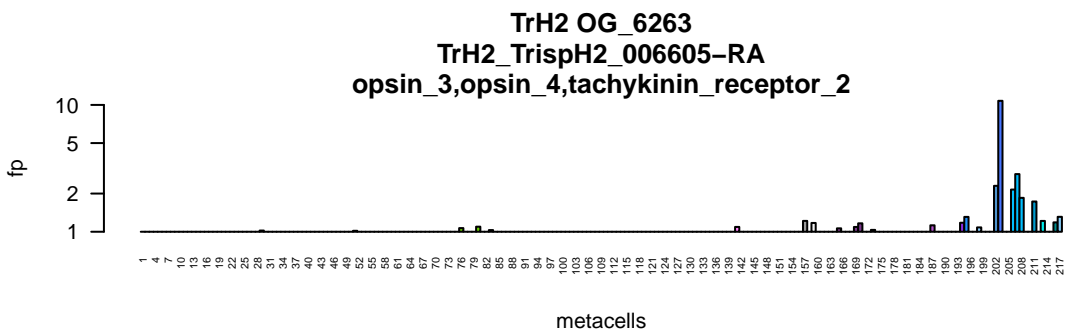
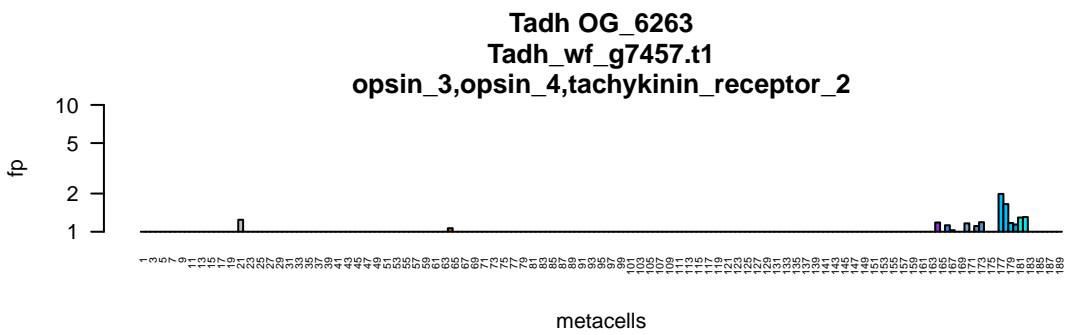
HoiH23 OG\_5203  
HoiH23\_PIH23\_003635-RA  
opioid\_receptor\_kappa\_1,G\_protein\_coupled\_receptor\_21



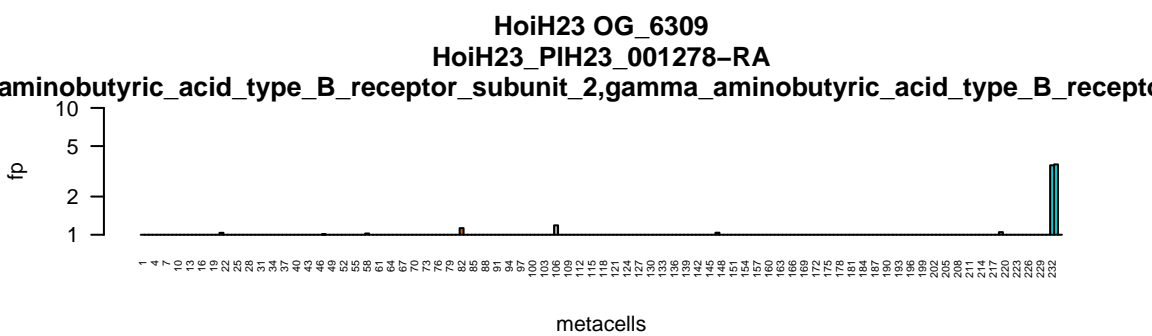
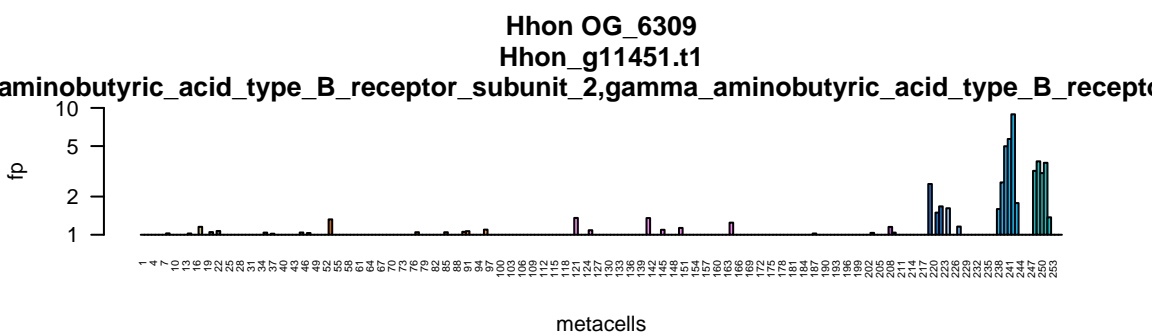
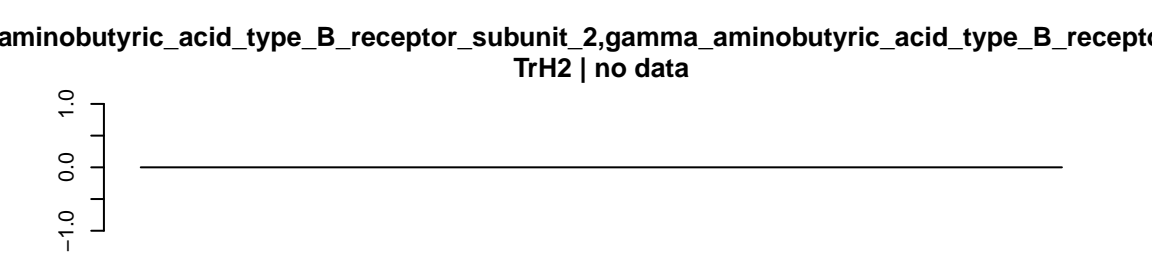
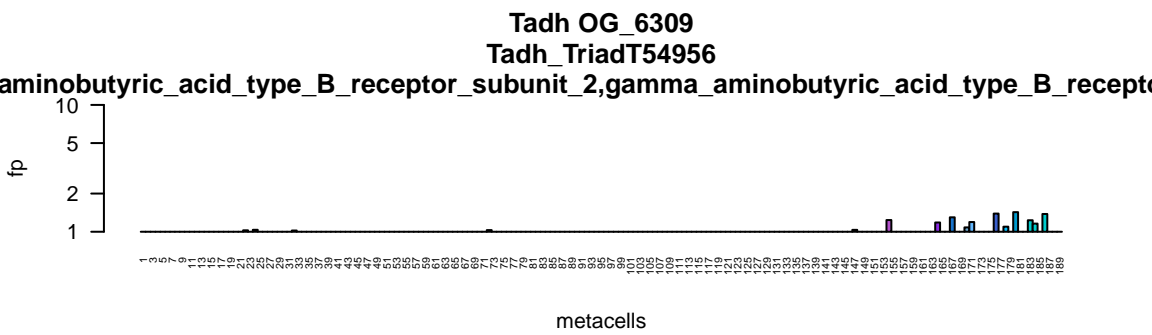
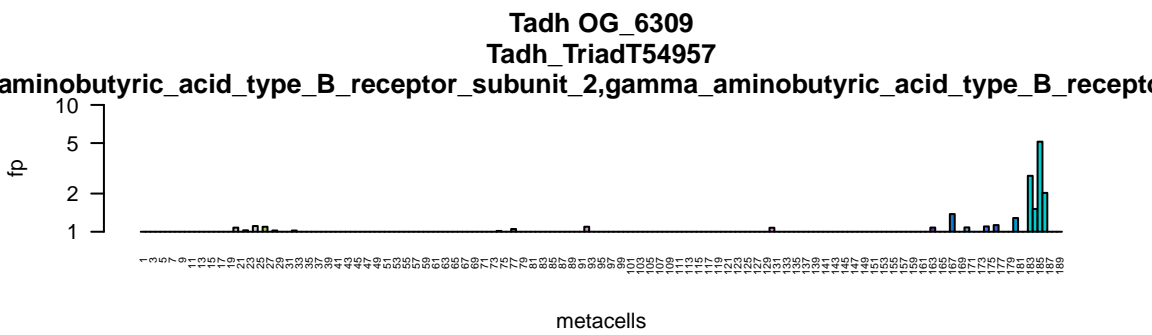
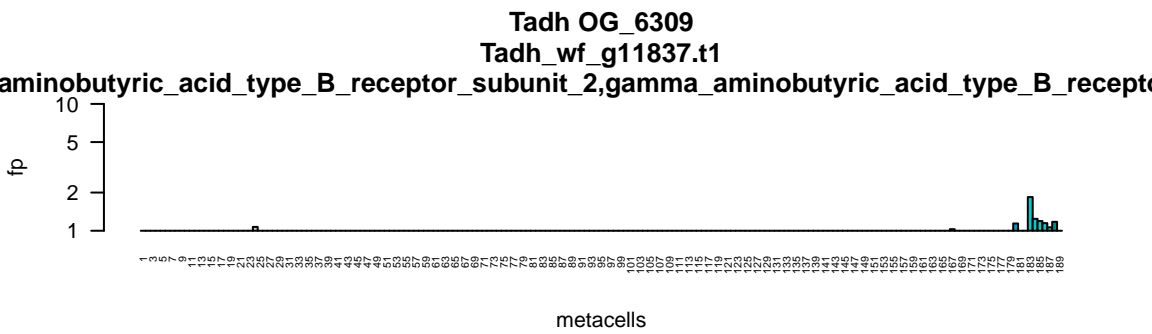


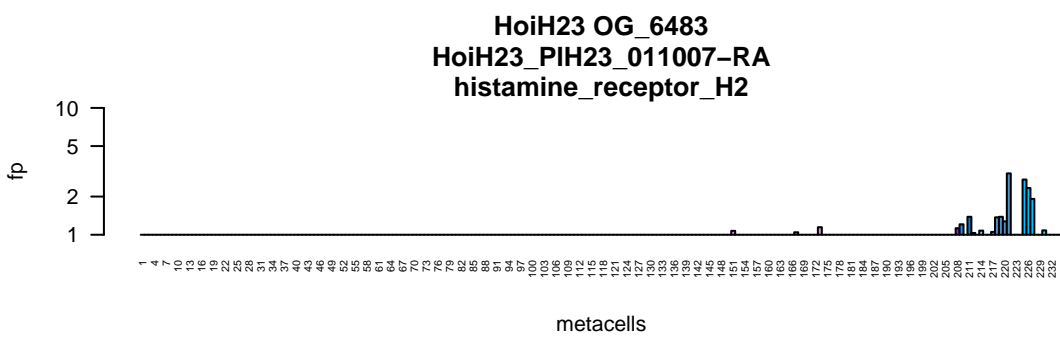
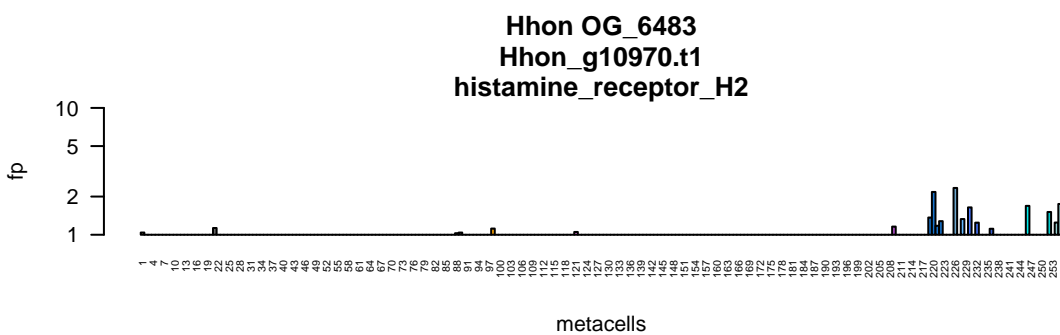
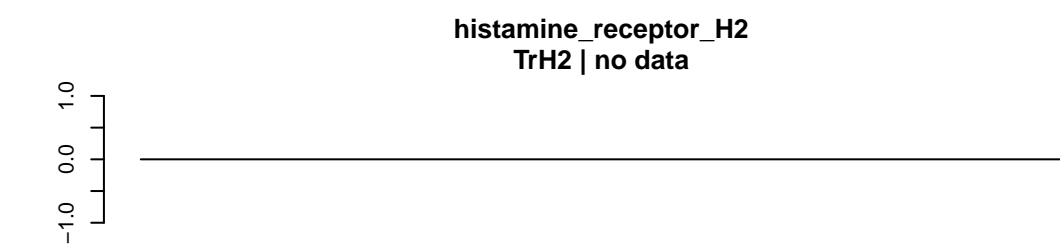
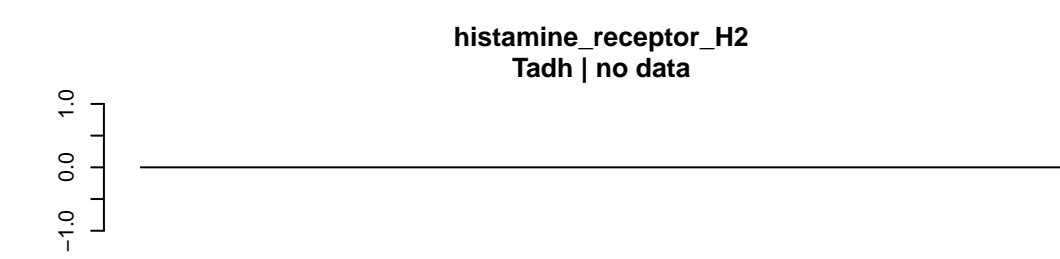












melanin\_concentrating\_hormone\_receptor\_1,C\_X\_C\_motif\_chemokine\_receptor\_1  
Tadh | no data

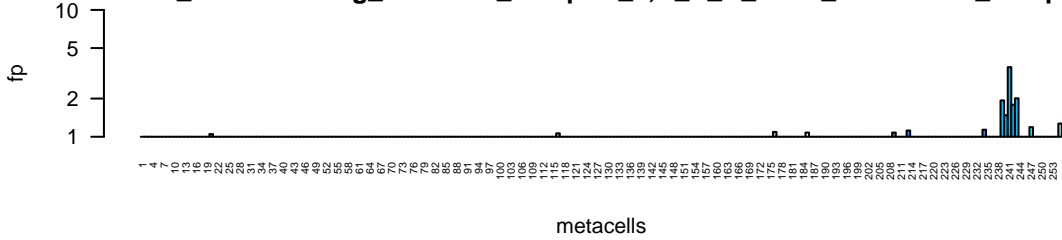


melanin\_concentrating\_hormone\_receptor\_1,C\_X\_C\_motif\_chemokine\_receptor\_1  
TrH2 | no data



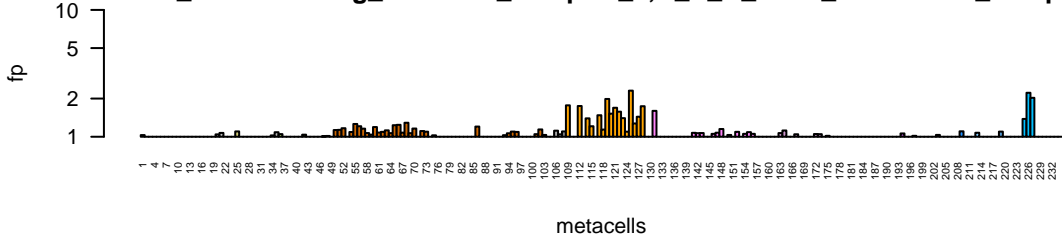
Hhon OG\_6495  
Hhon\_g08607.t1

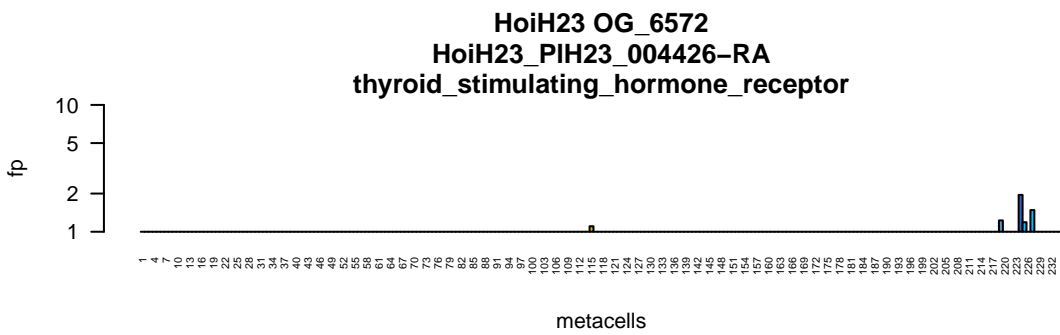
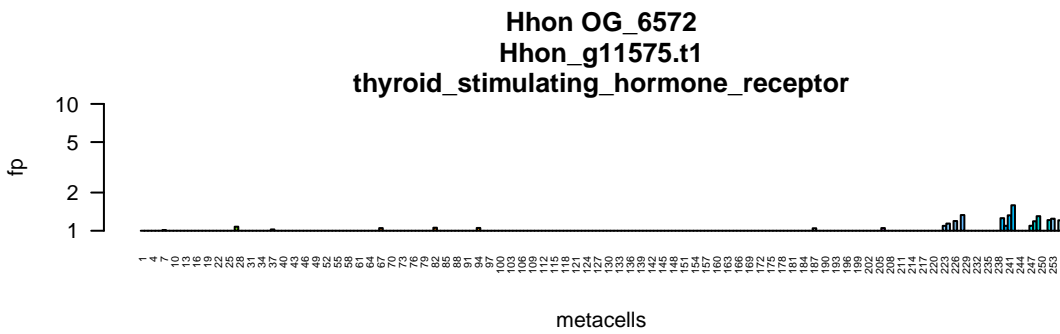
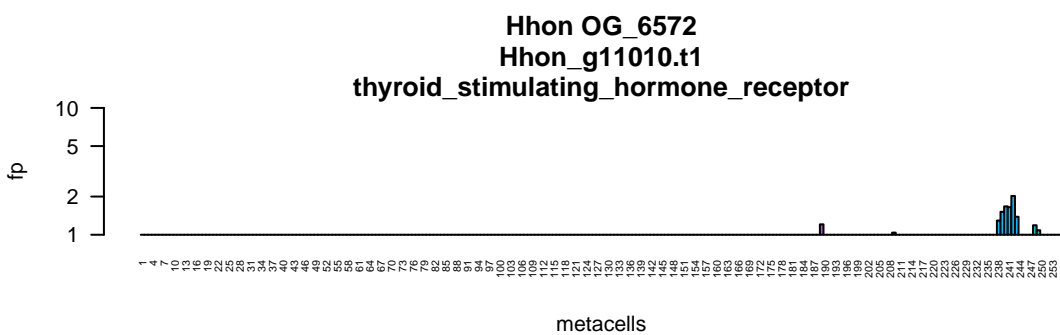
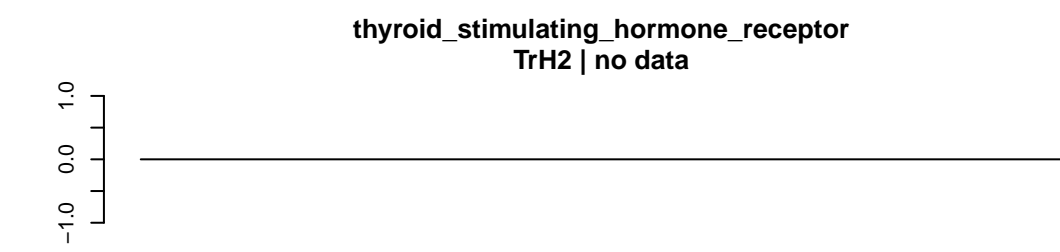
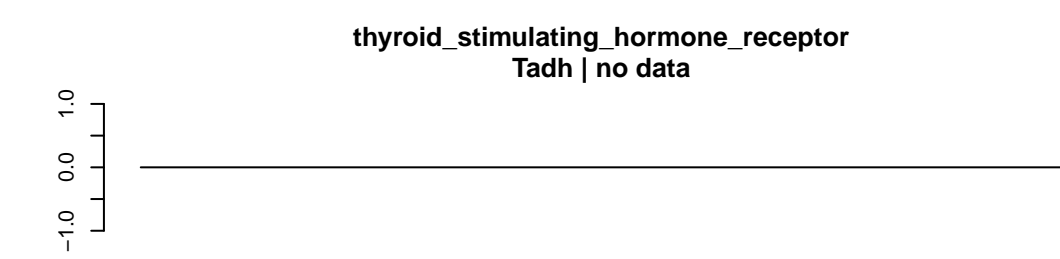
melanin\_concentrating\_hormone\_receptor\_1,C\_X\_C\_motif\_chemokine\_receptor\_1

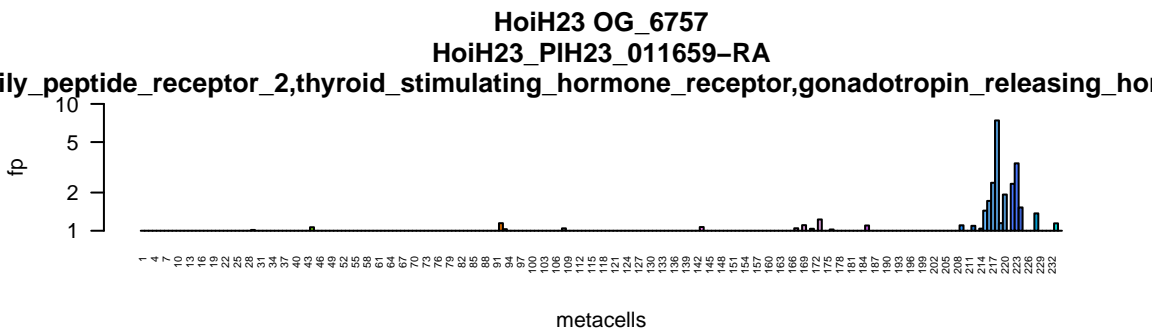
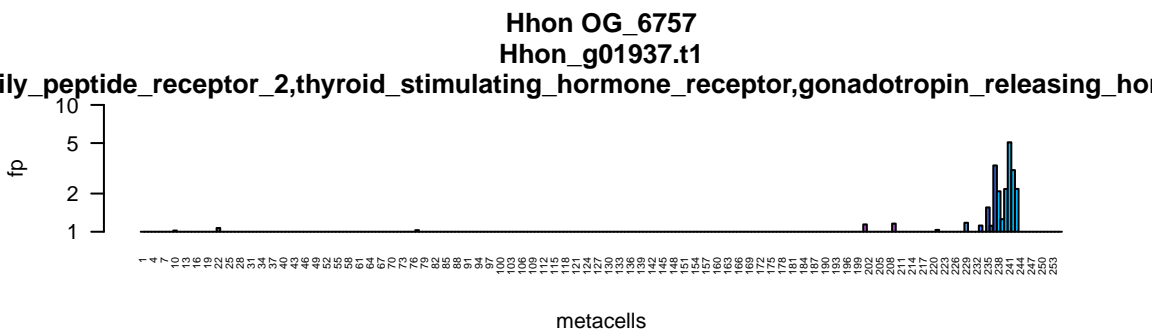
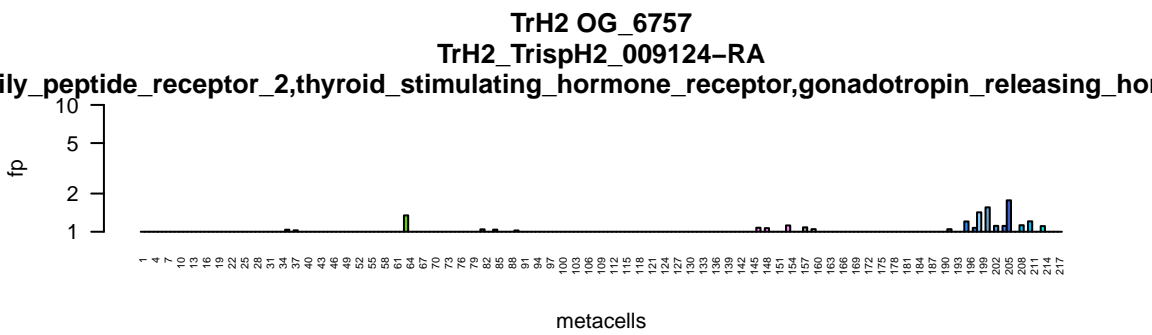
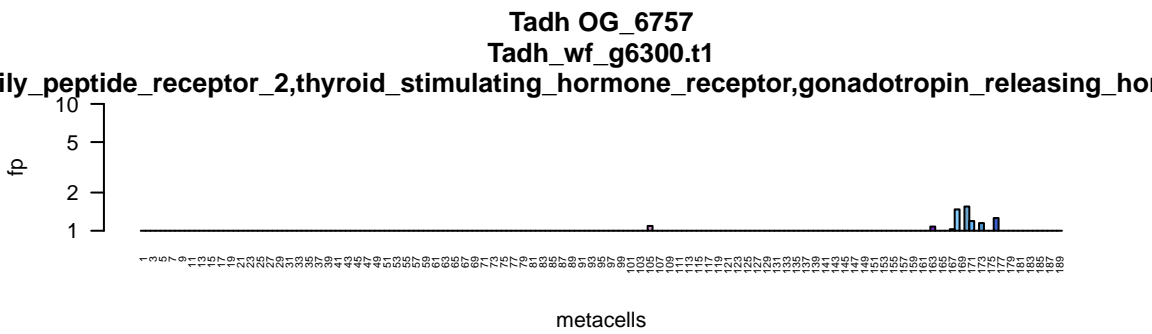


HoiH23 OG\_6495  
HoiH23\_PIH23\_004725-RA

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



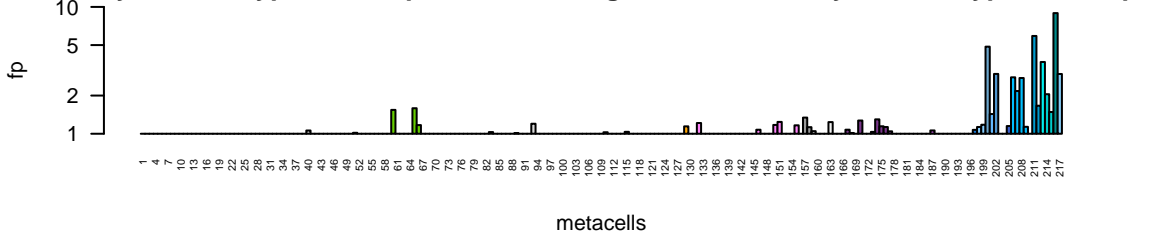




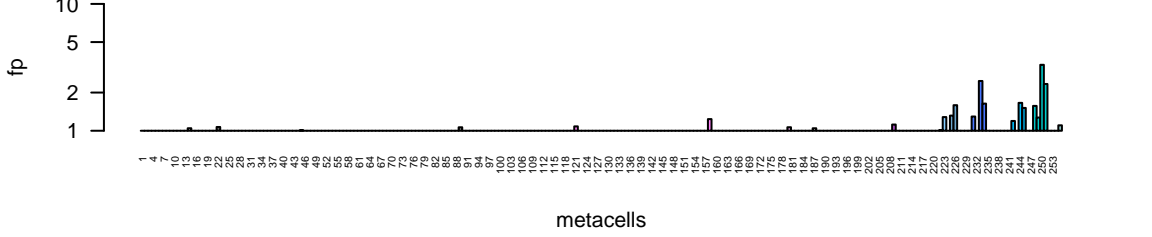
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recepto  
Tadh | no data



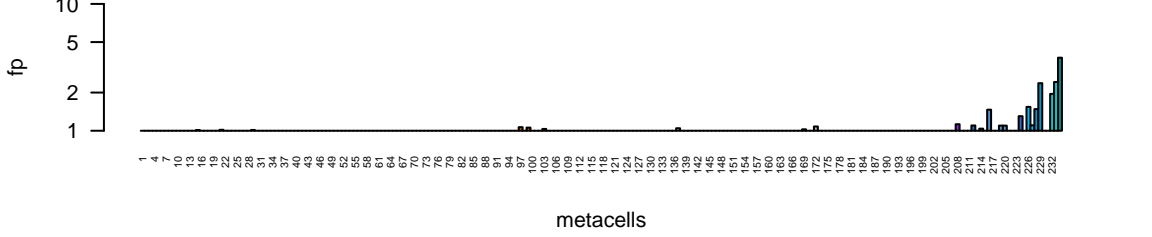
TrH2 OG\_7186  
TrH2\_TrispH2\_010138-RA  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recepto



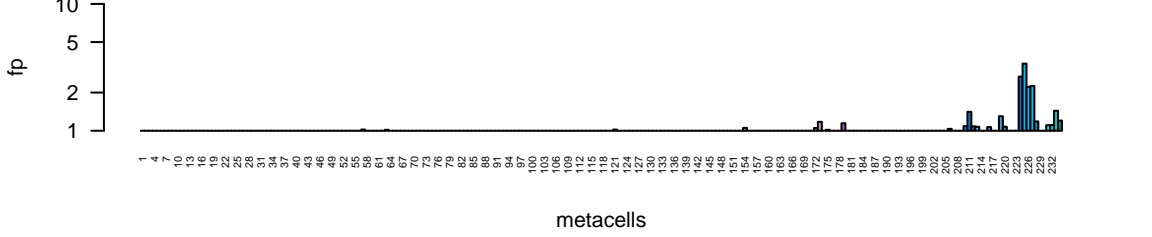
Hhon OG\_7186  
Hhon\_g01977.t1  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recepto



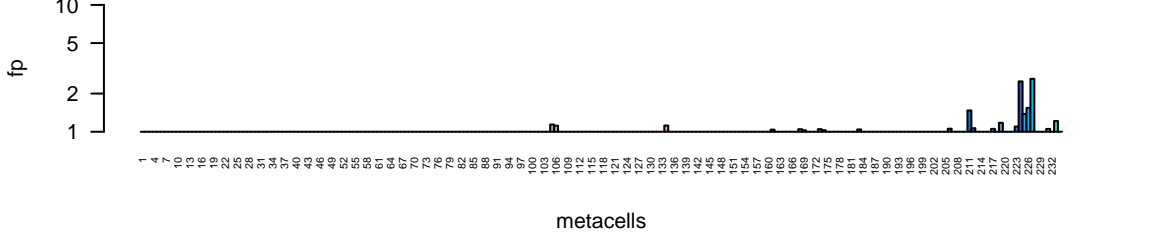
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HoiH23\_PIH23\_011144-RA  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recepto



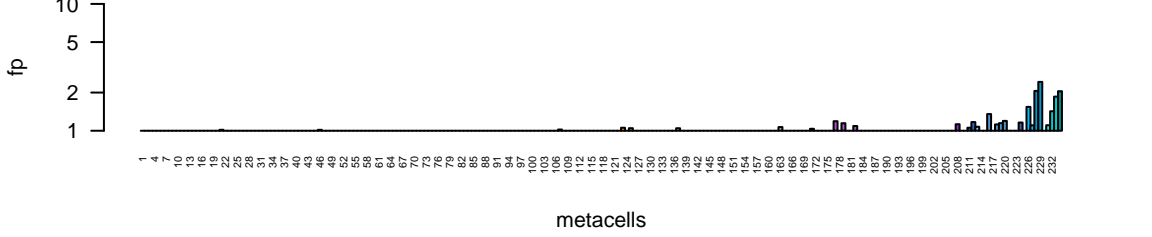
HoiH23 OG\_7186  
HoiH23\_PIH23\_011313-RA  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recepto

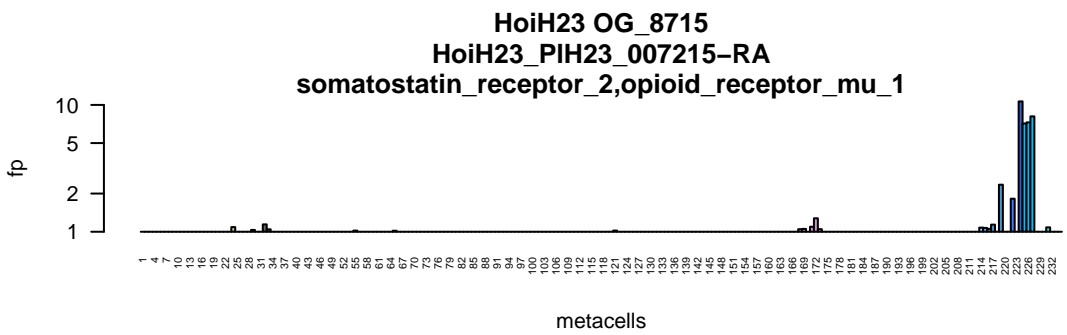
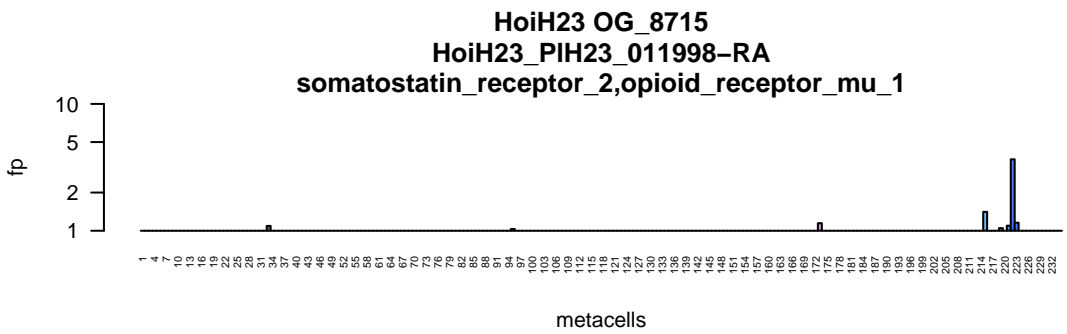
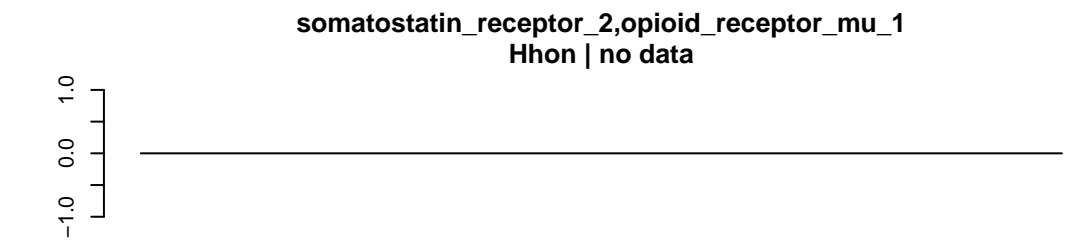
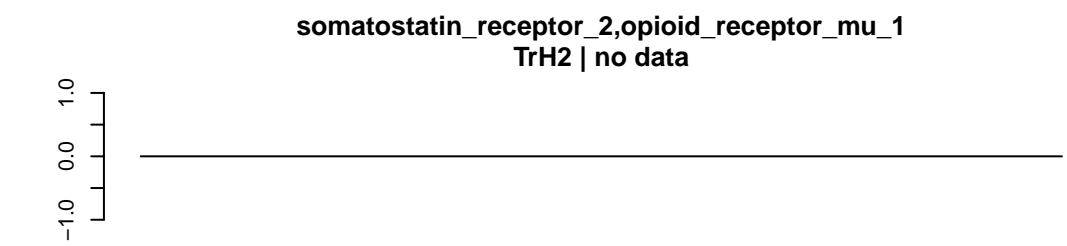
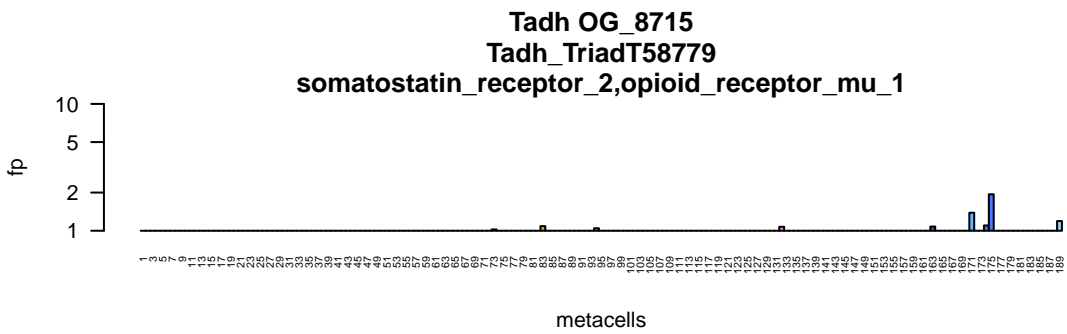


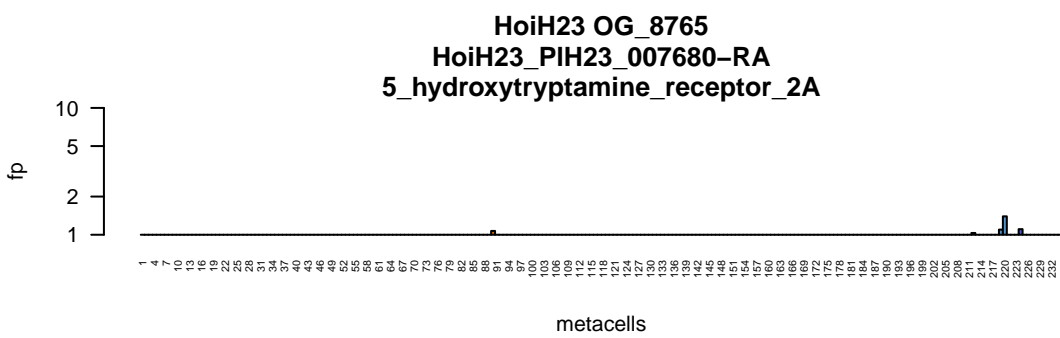
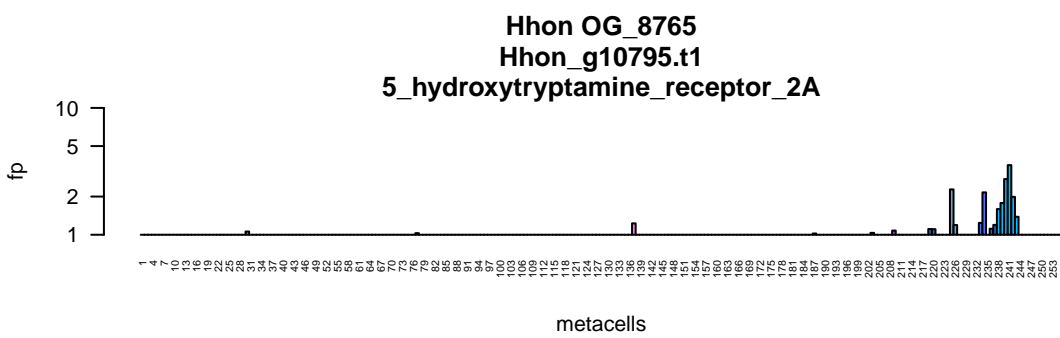
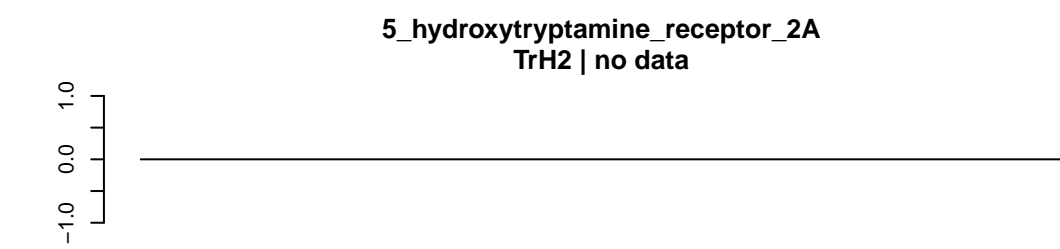
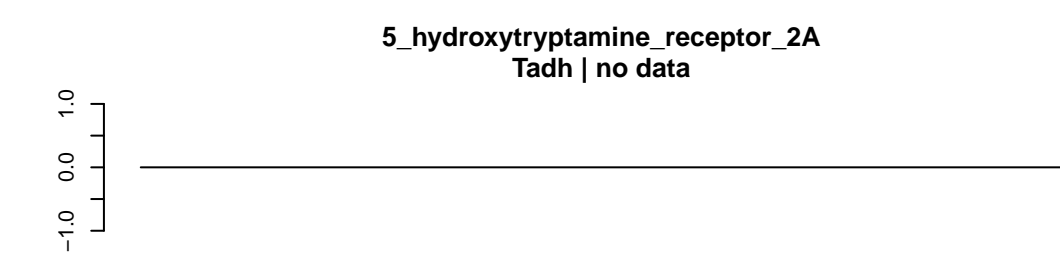
HoiH23 OG\_7186  
HoiH23\_PIH23\_004027-RA  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recepto



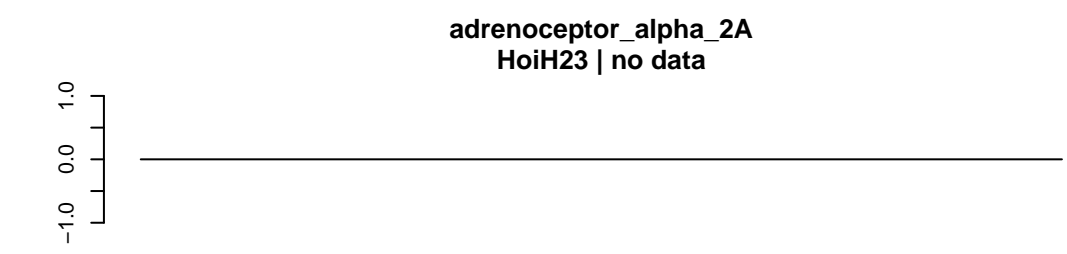
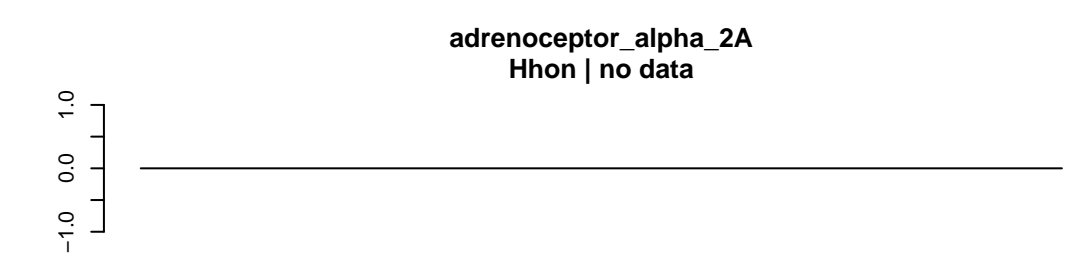
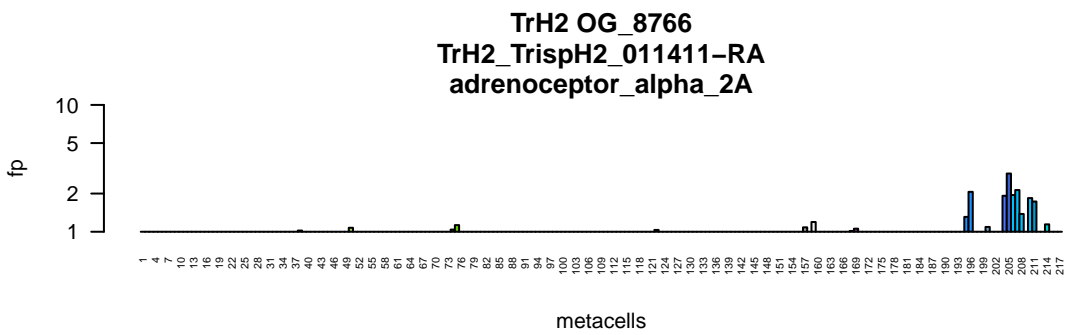
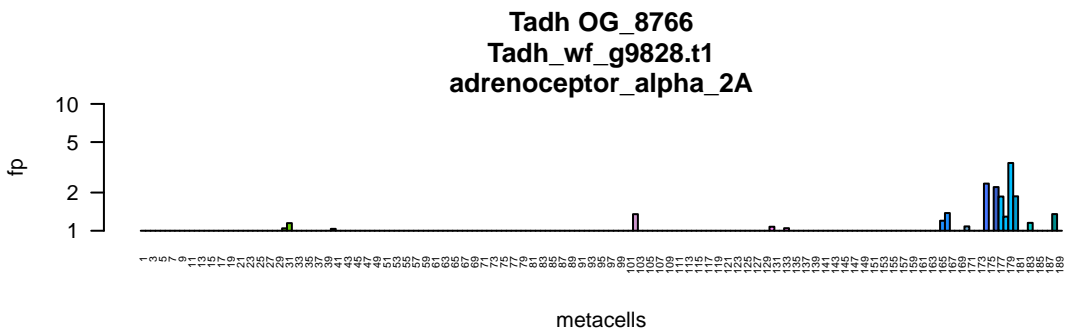
HoiH23 OG\_7186  
HoiH23\_PIH23\_011312-RA  
aminobutyric\_acid\_type\_B\_receptor\_subunit\_2,gamma\_aminobutyric\_acid\_type\_B\_recepto

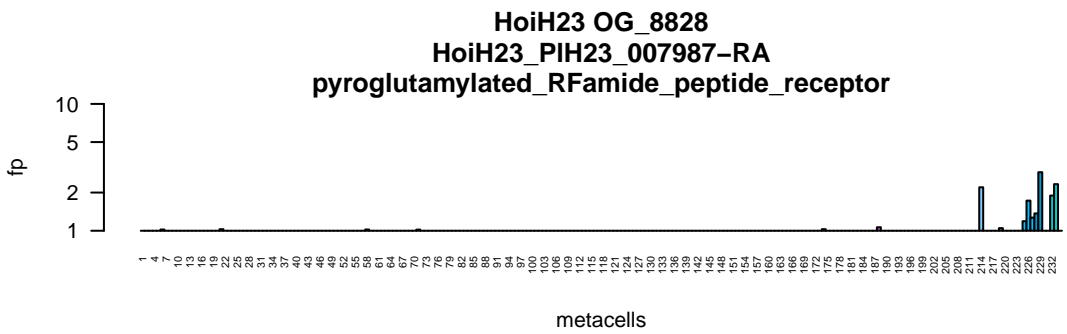
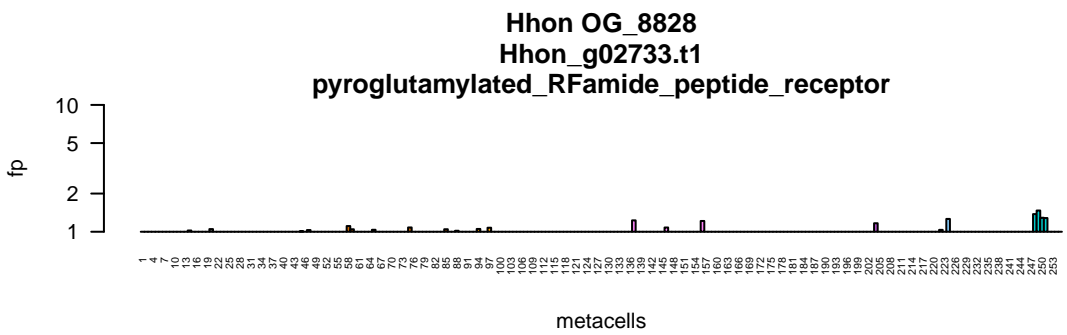
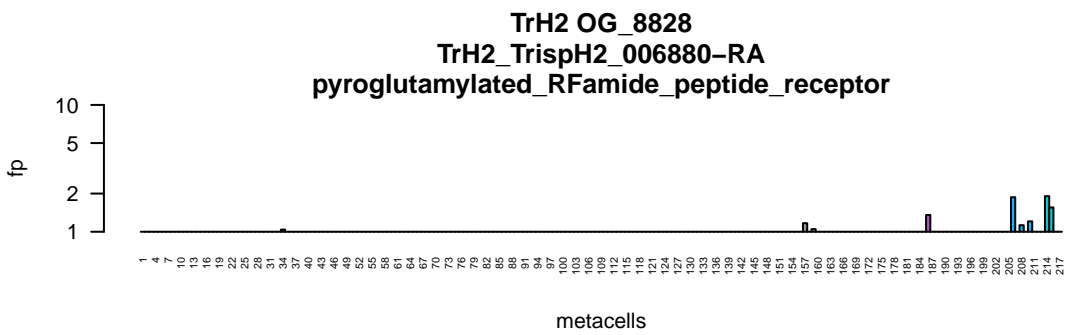
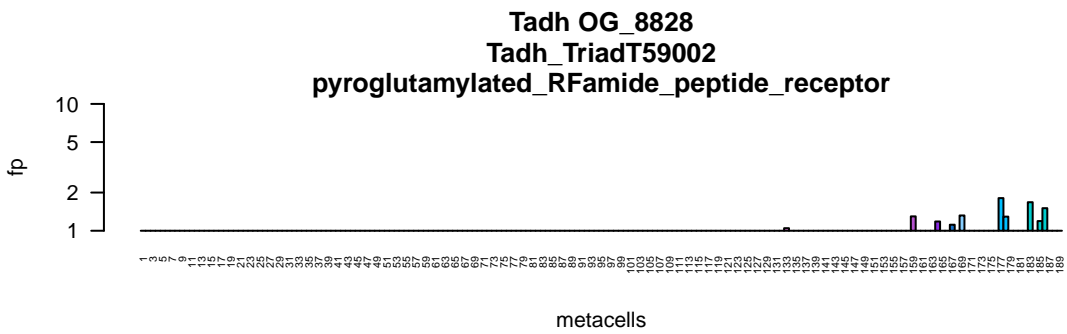


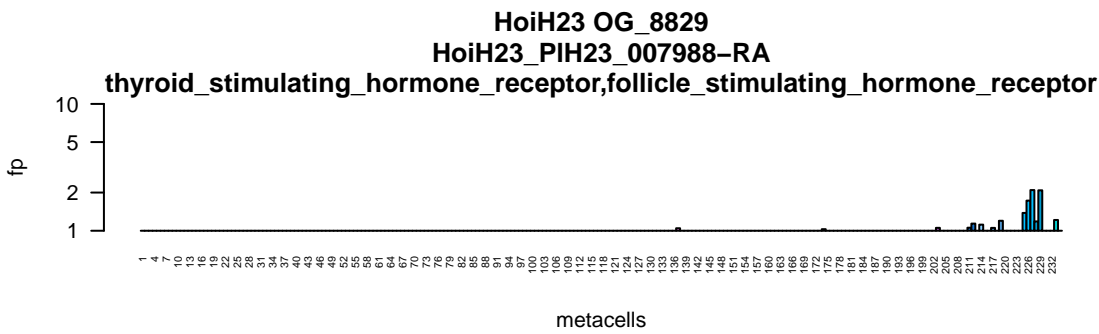
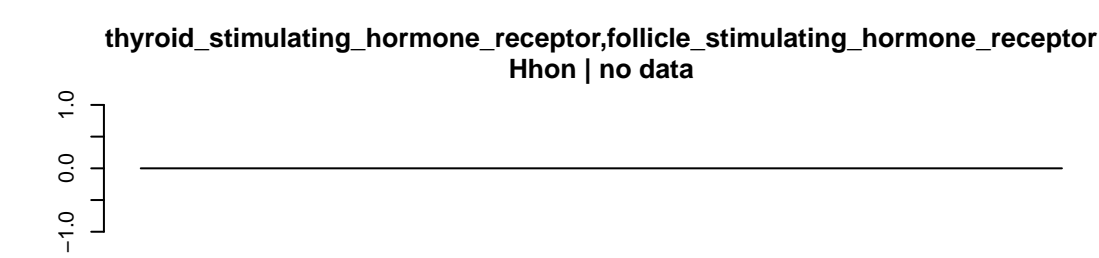
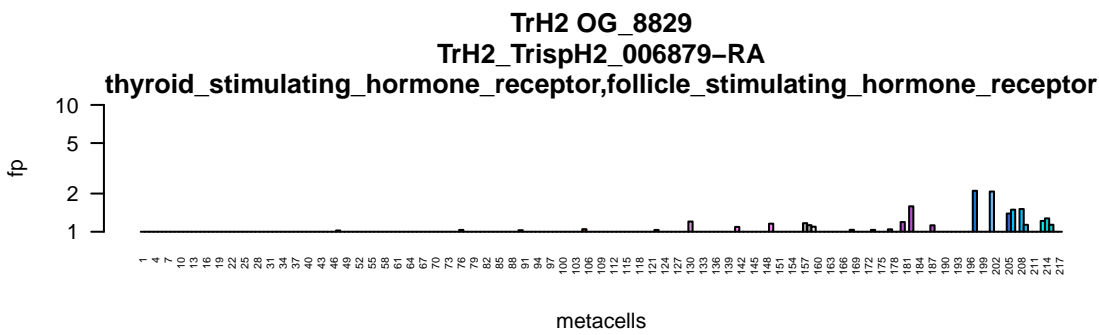
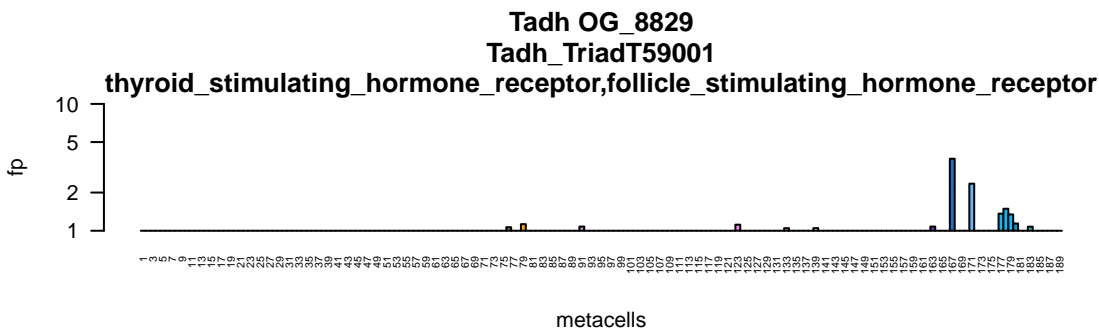


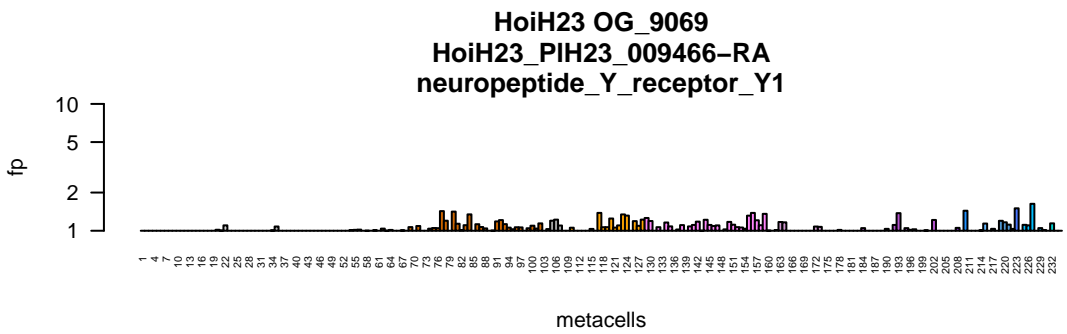
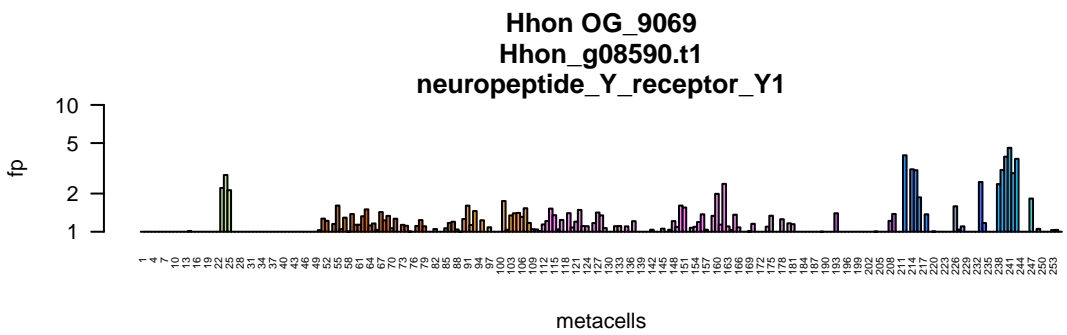
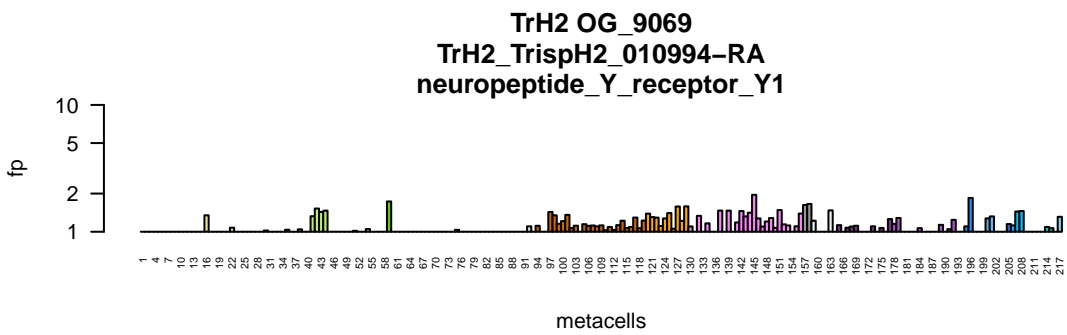
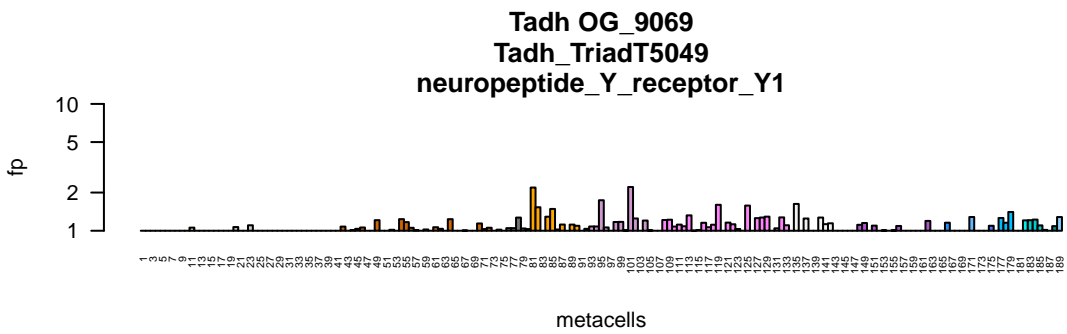


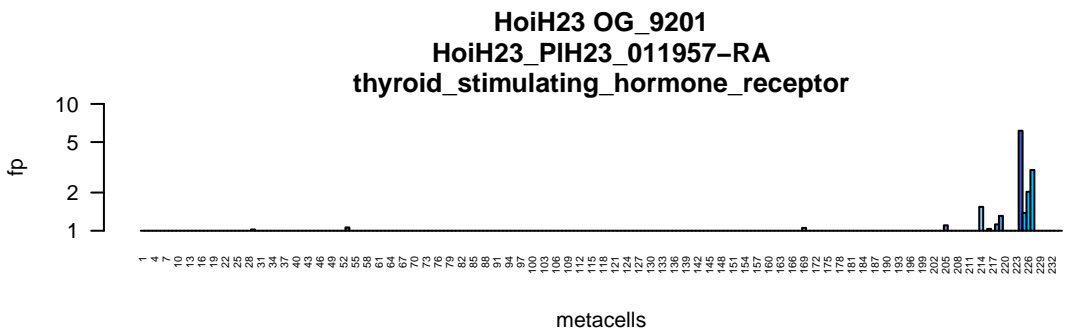
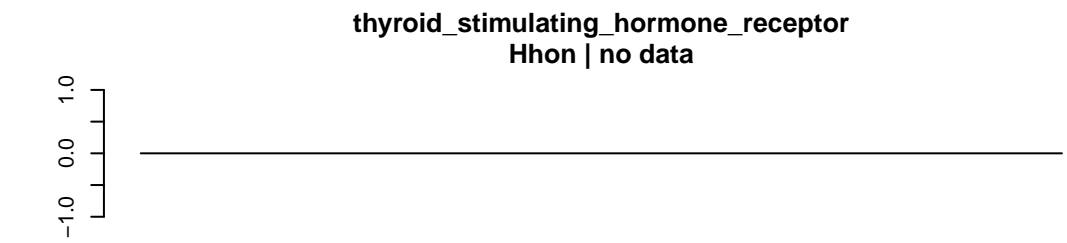
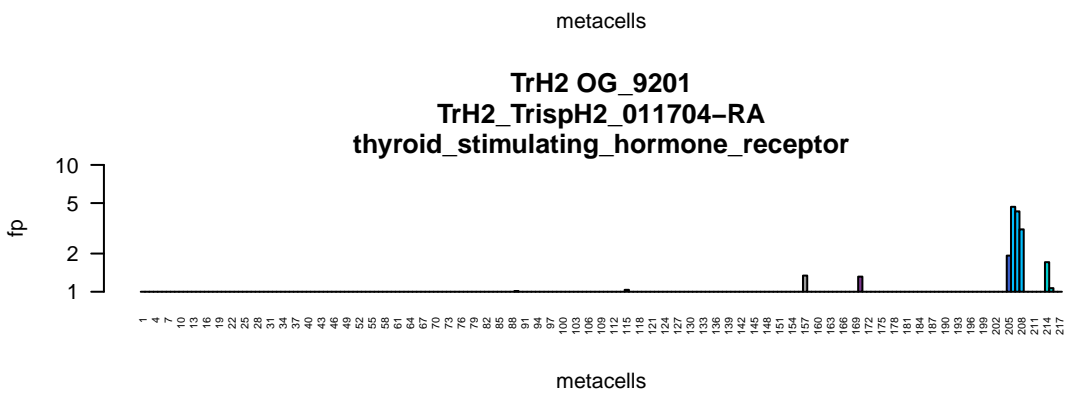
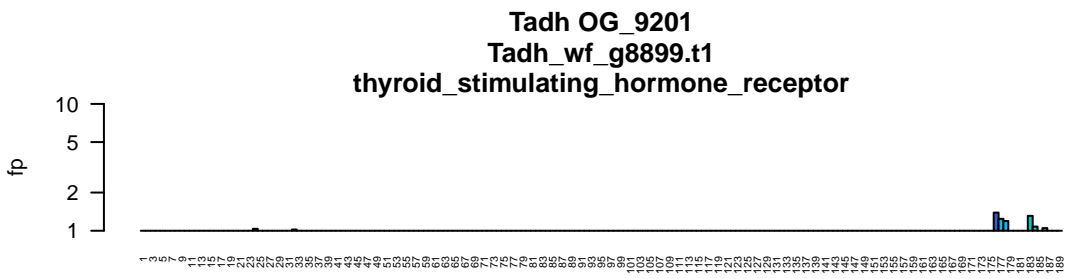


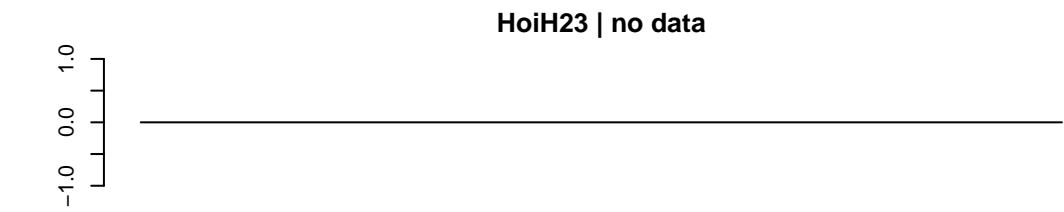
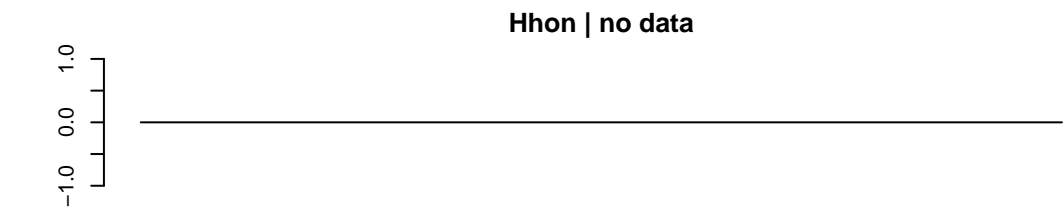
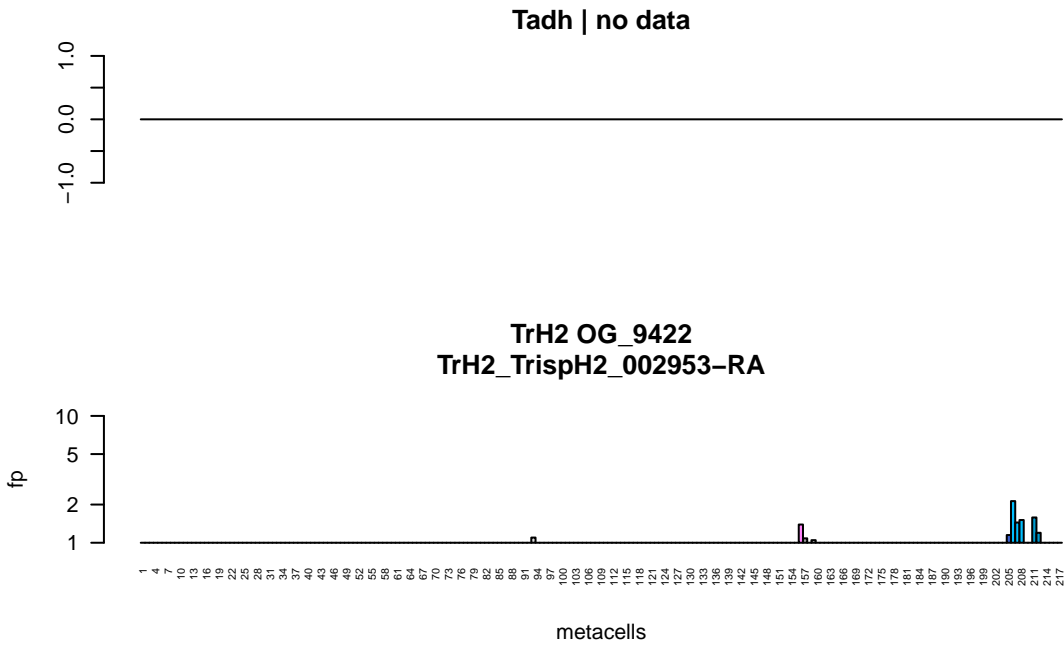


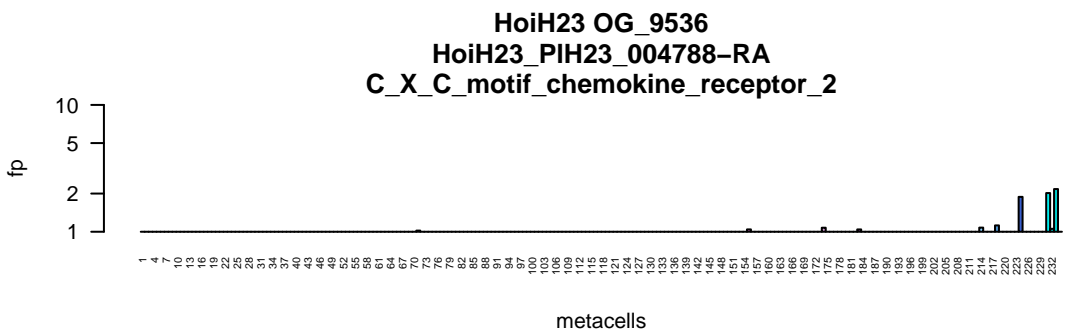
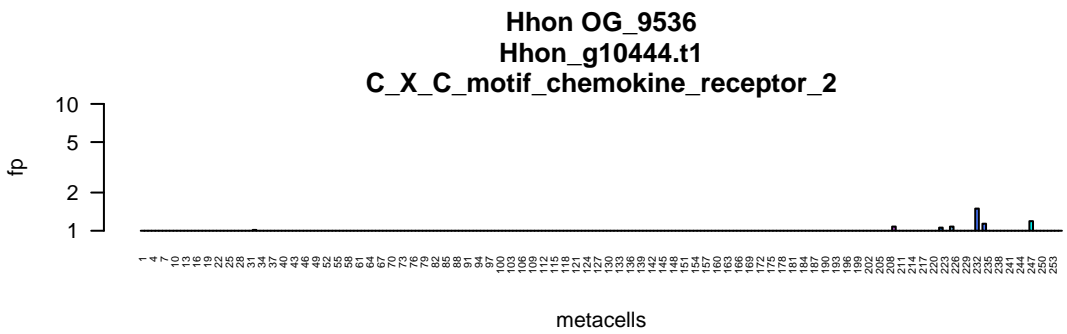
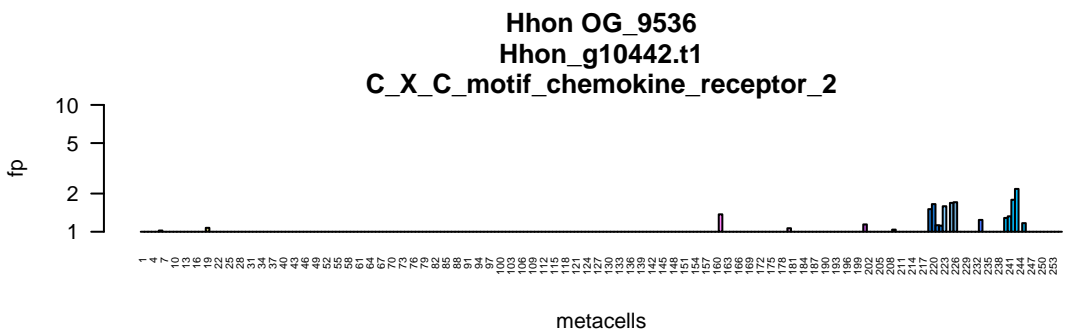
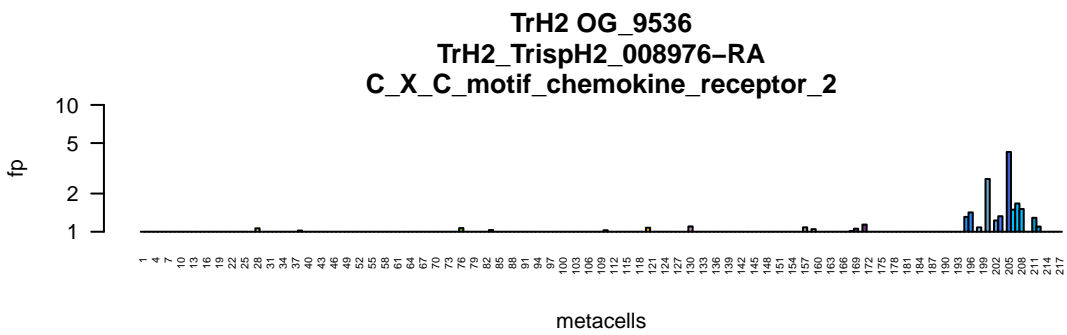
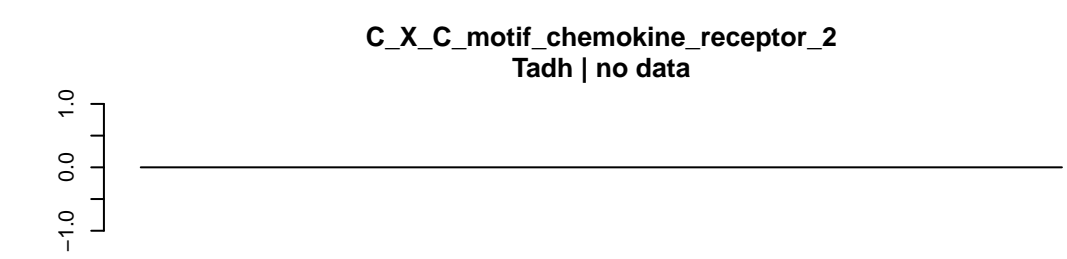


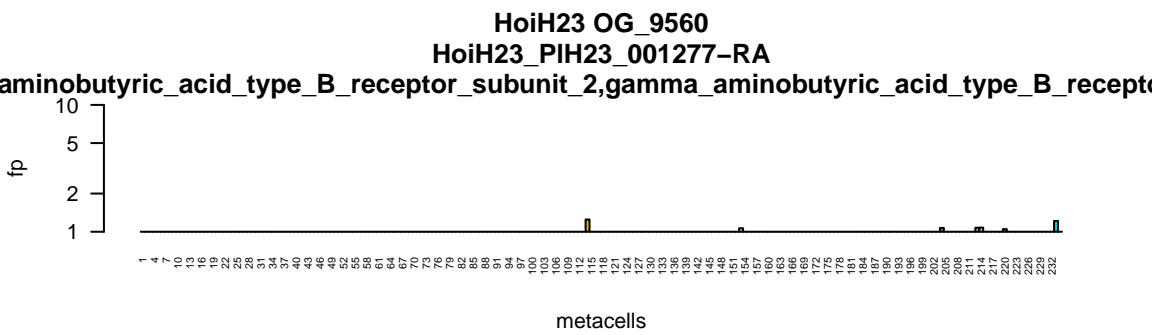
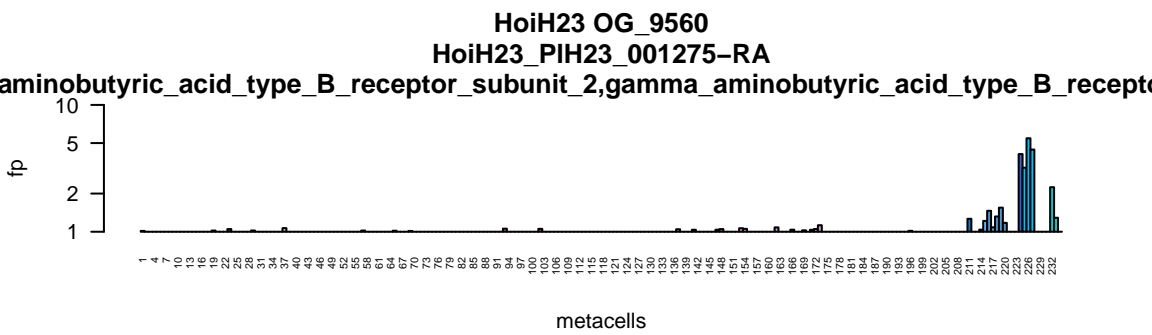
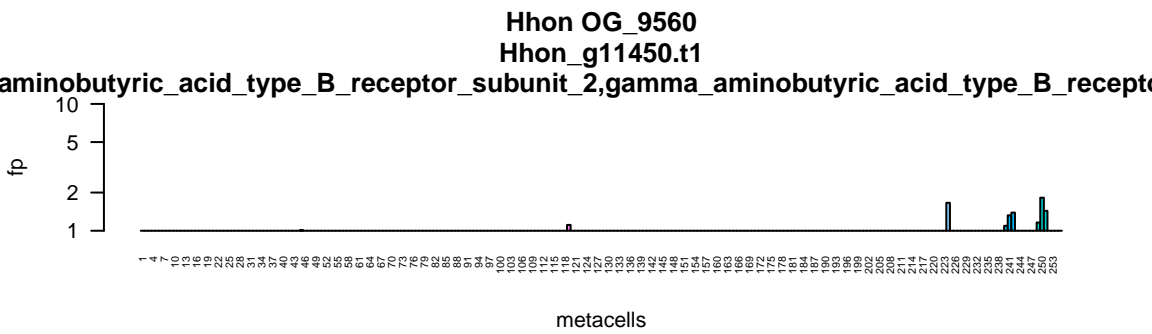
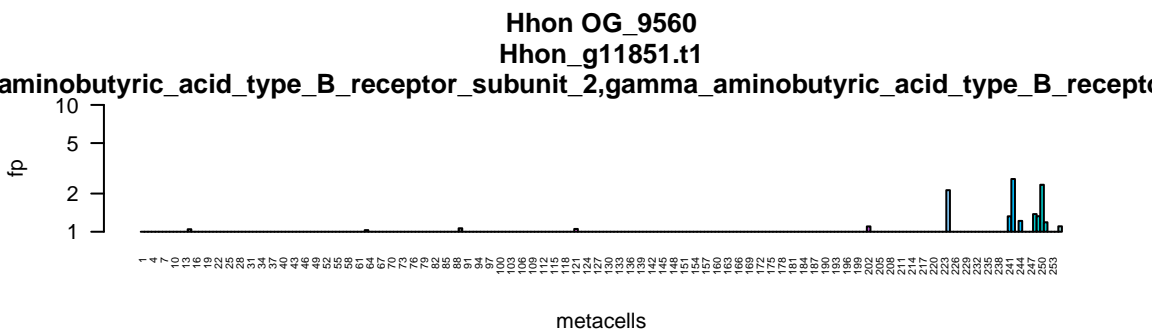
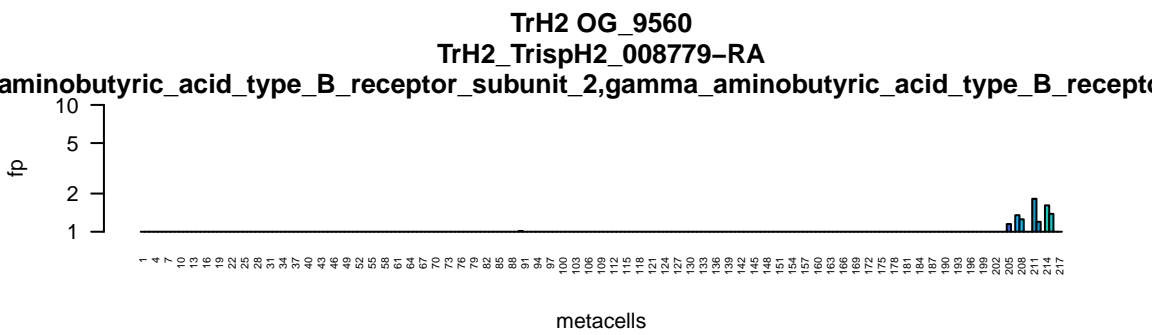
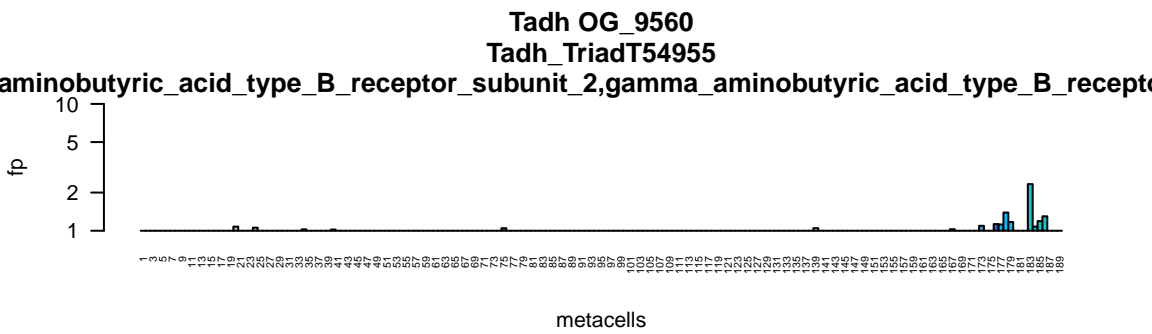




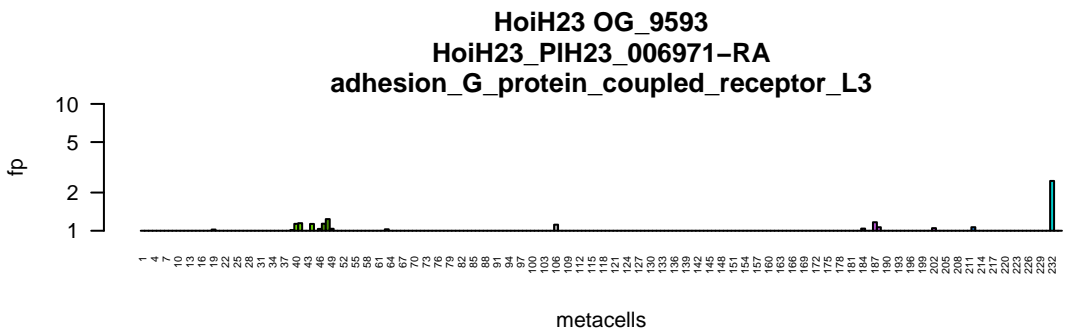
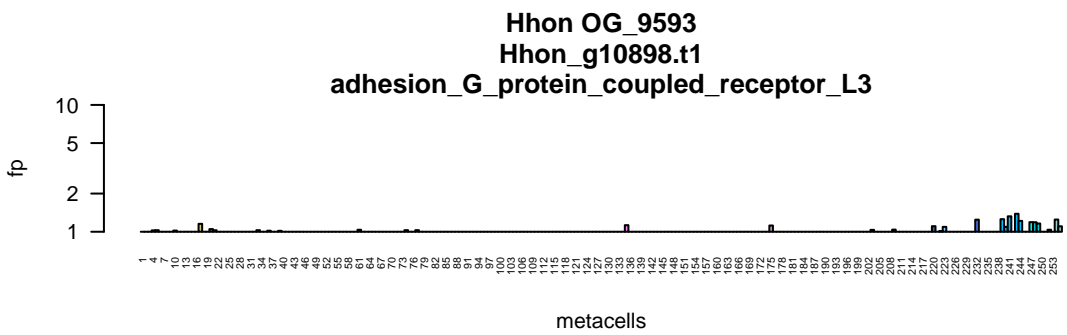
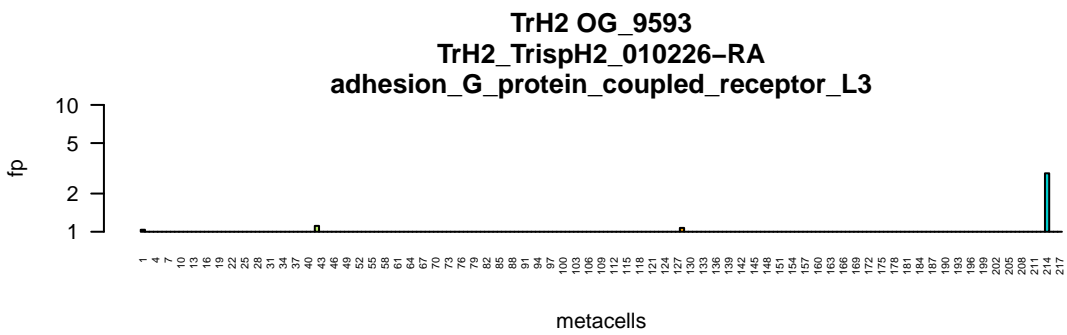
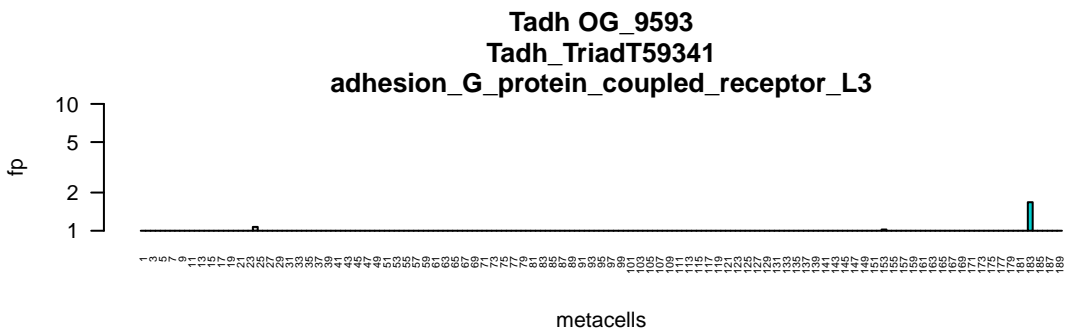




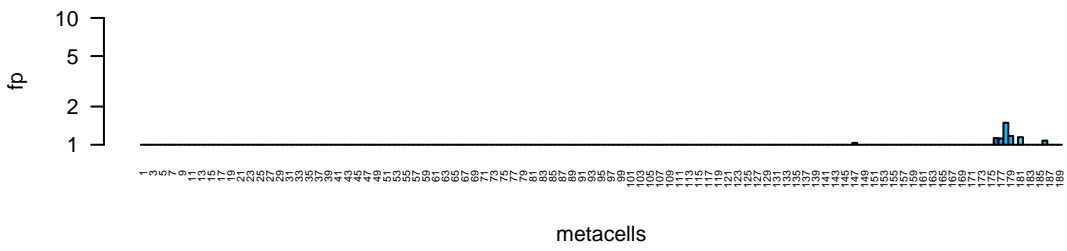




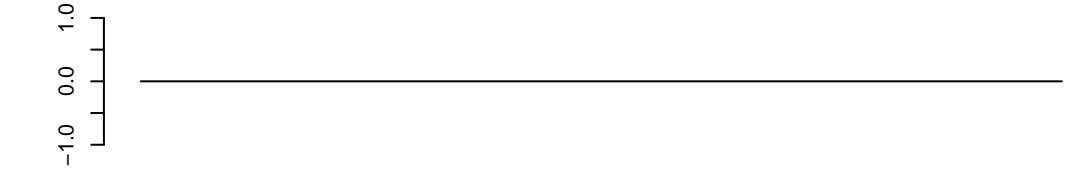




Tadh OG\_9754  
Tadh\_TriadT58809



TrH2 | no data



Hhon | no data



HoiH23 | no data



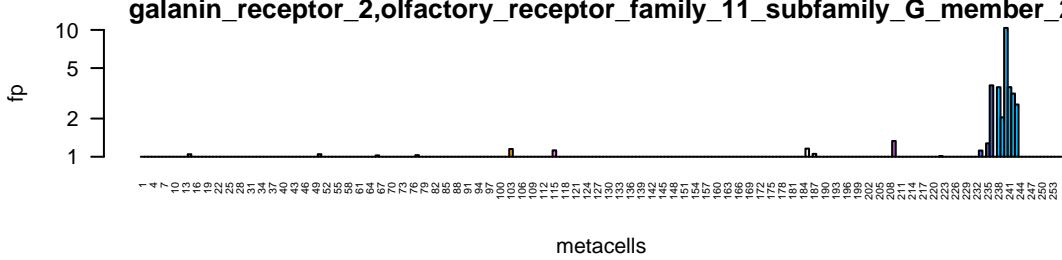
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Tadh | no data



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



