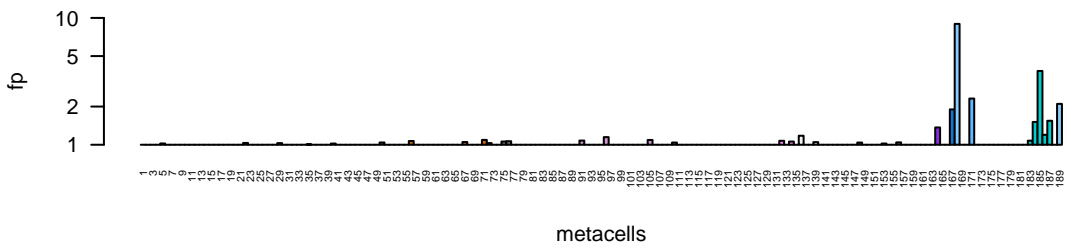
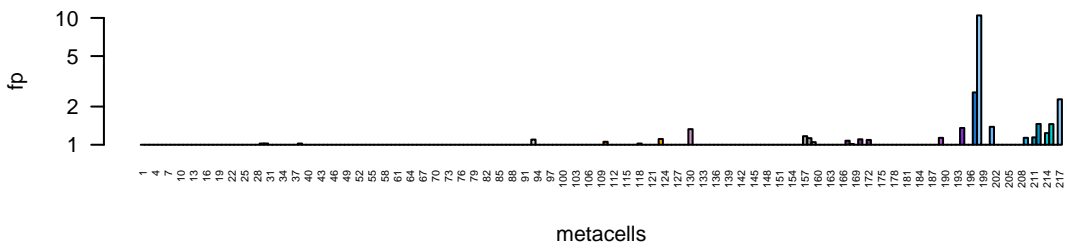


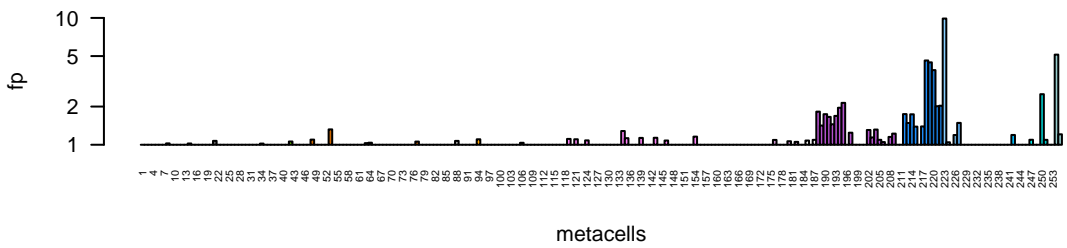
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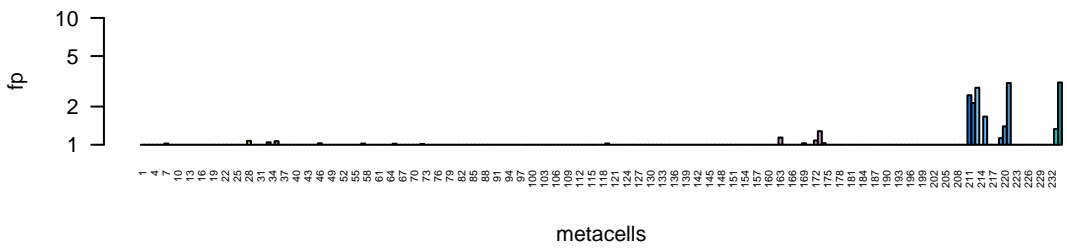
TrH2 OG_7537
TrH2_TrispH2_001160-RA

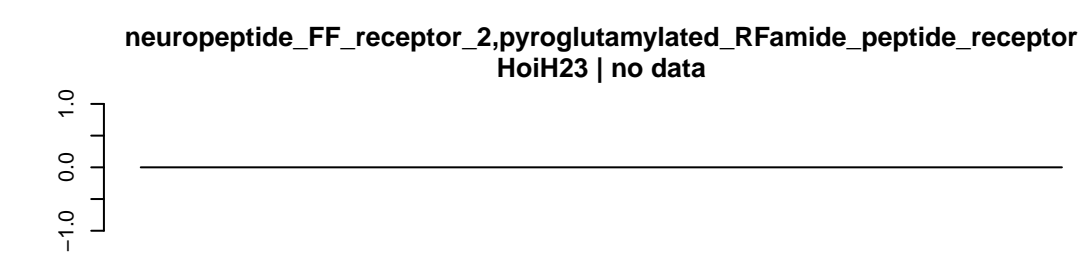
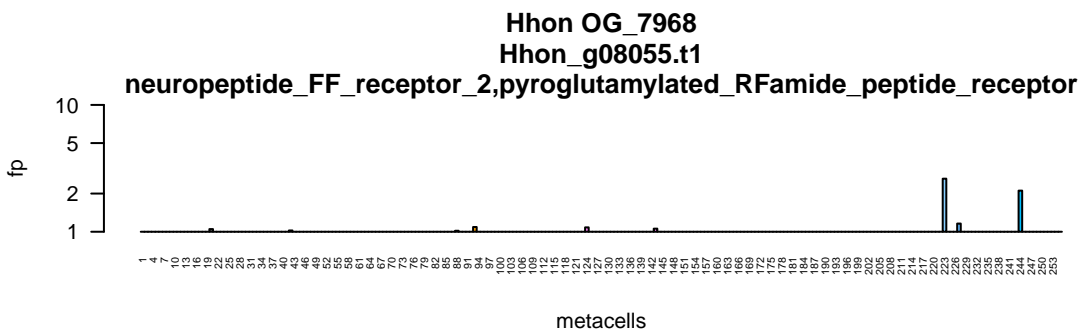
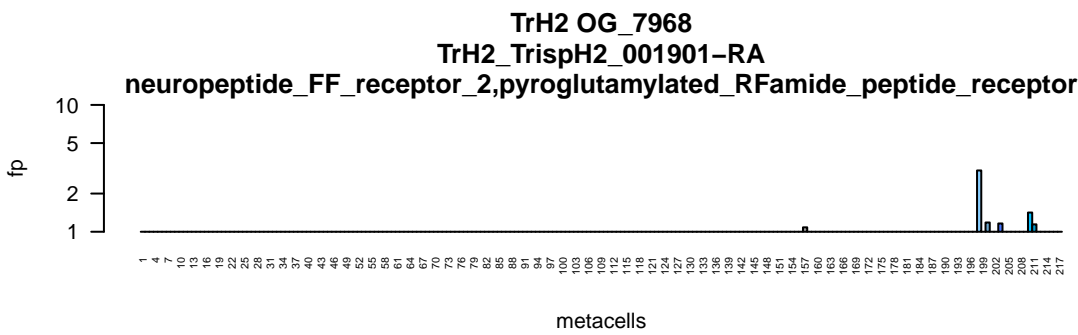
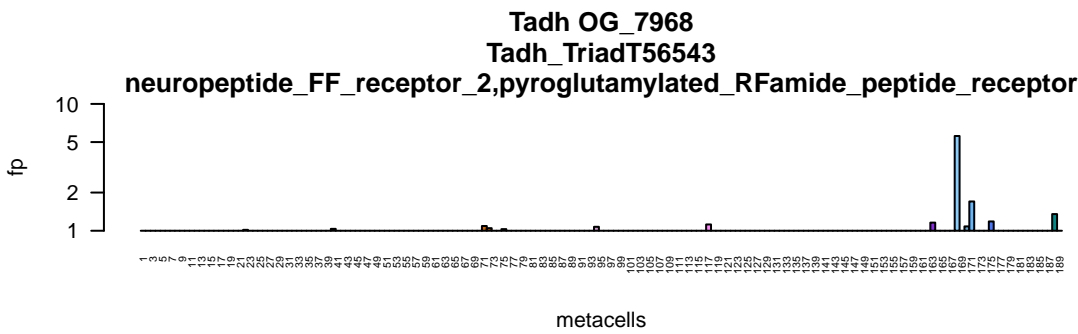


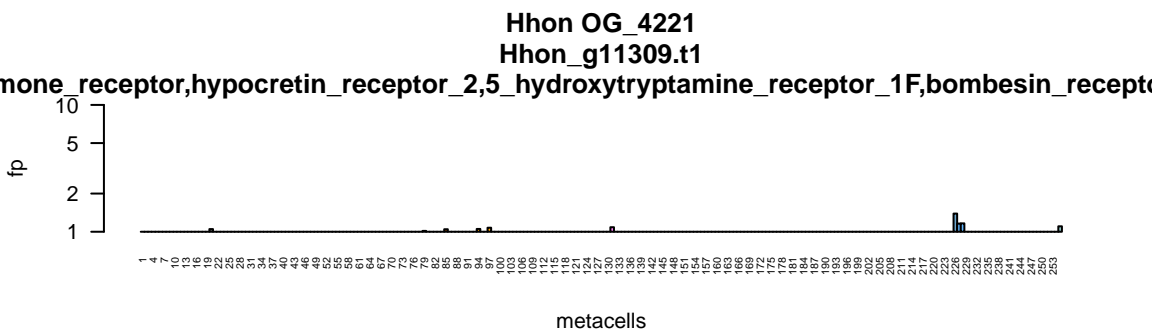
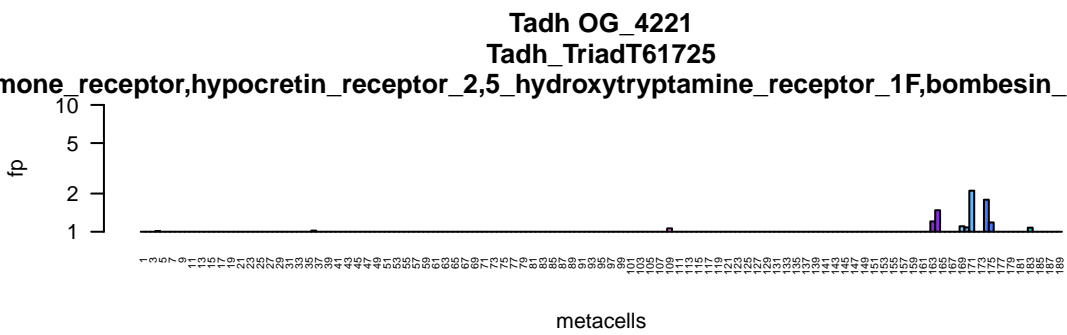
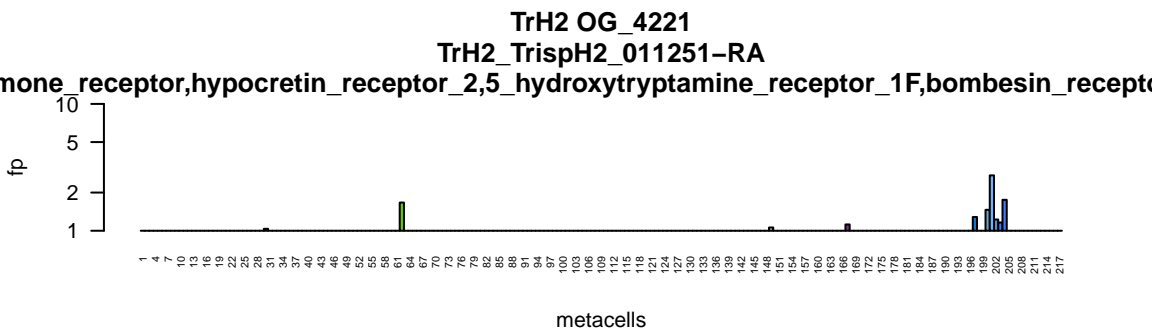
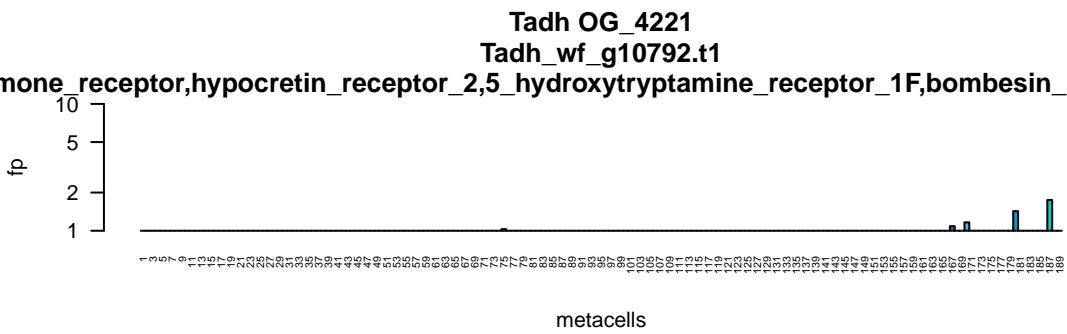
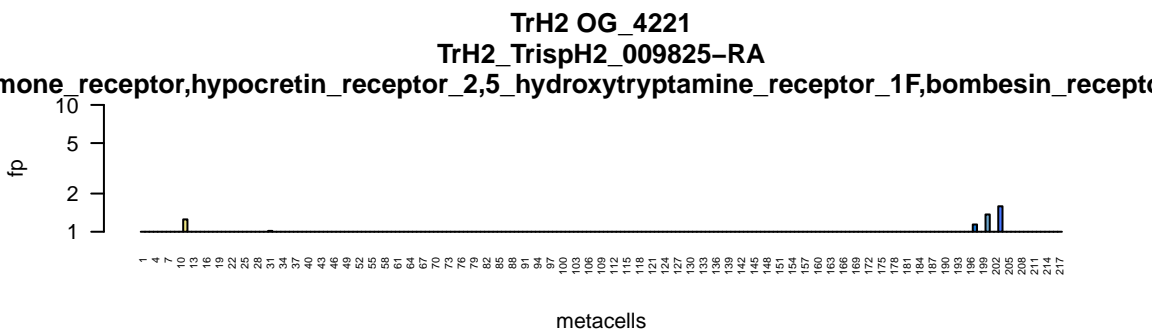
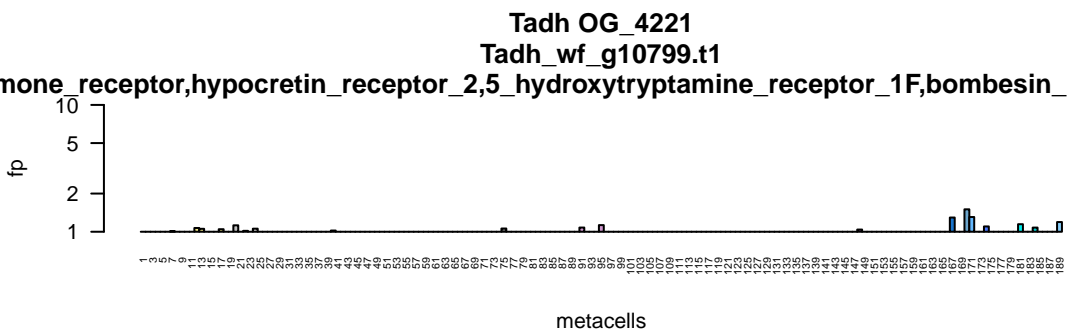
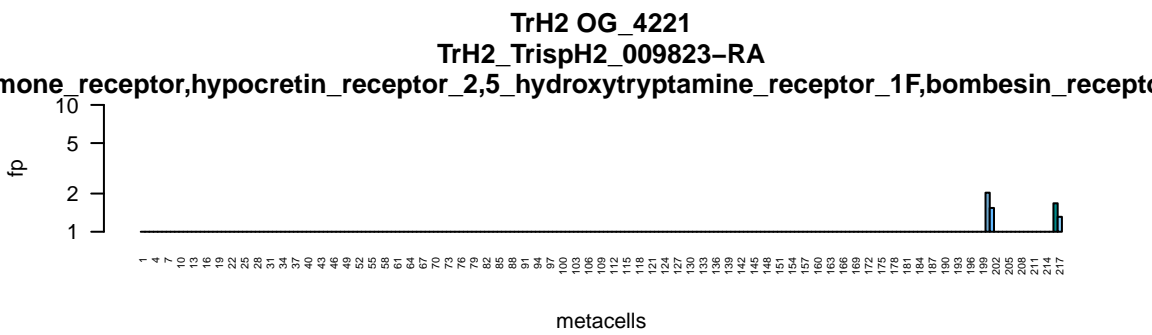
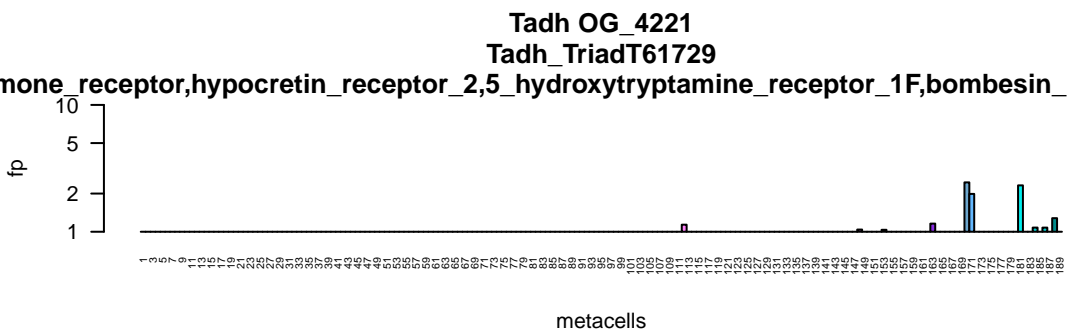
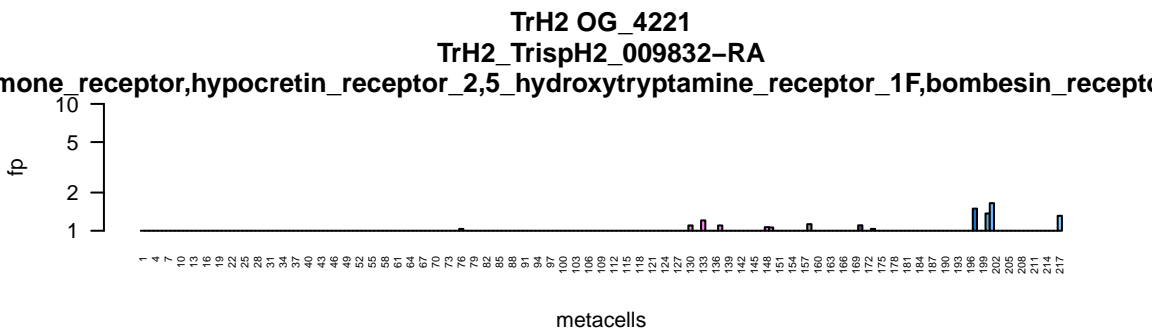
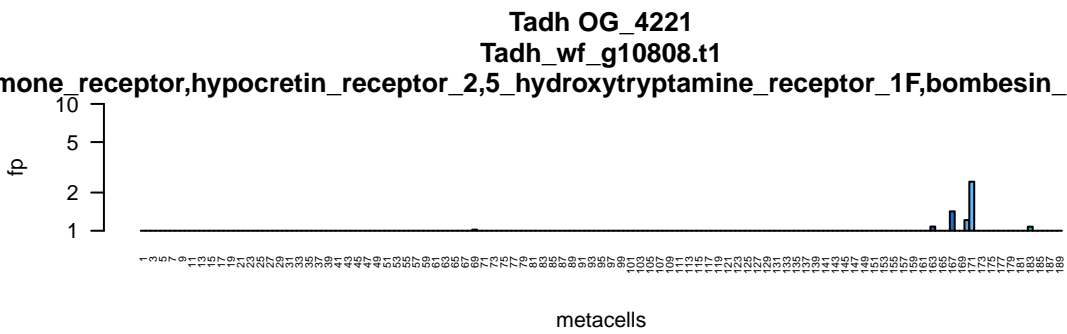
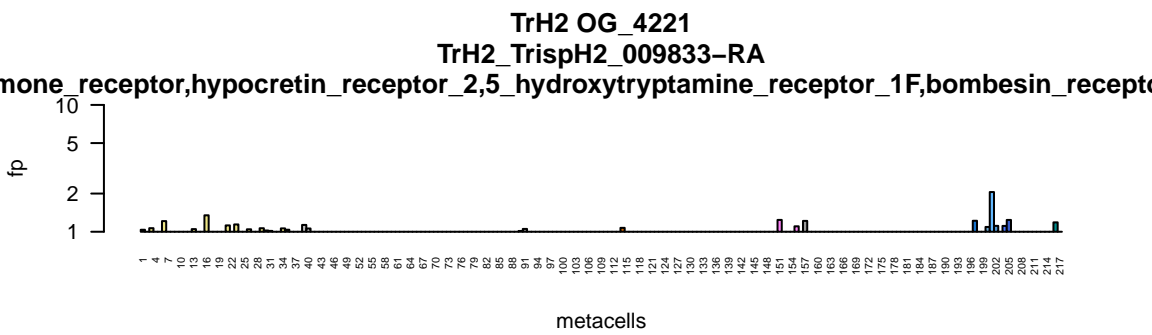
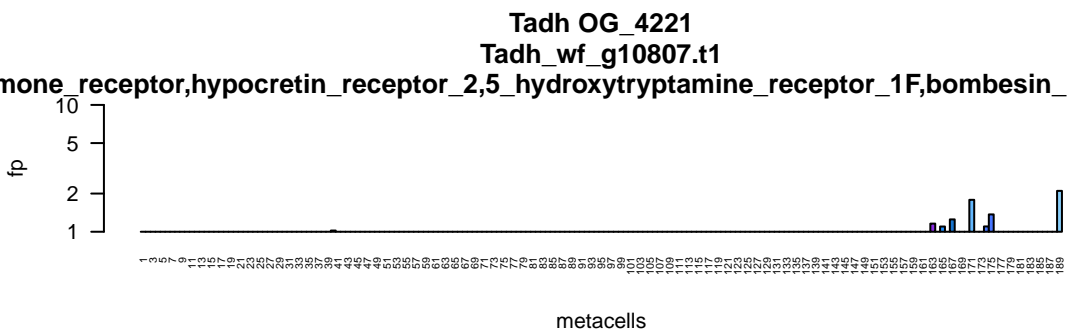
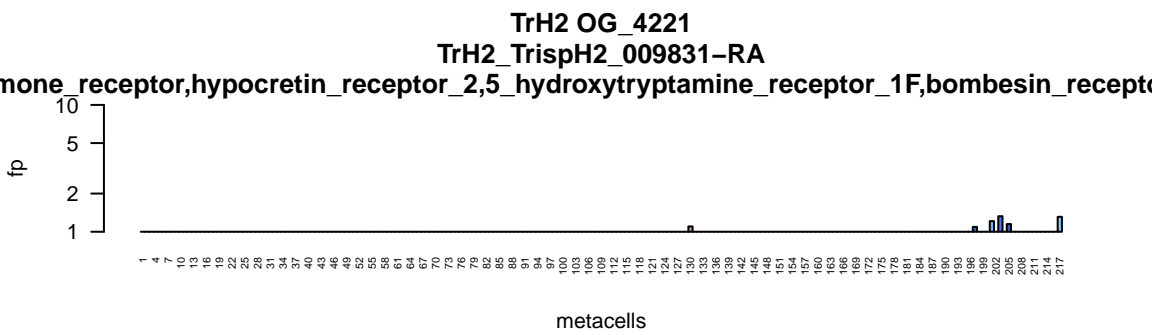
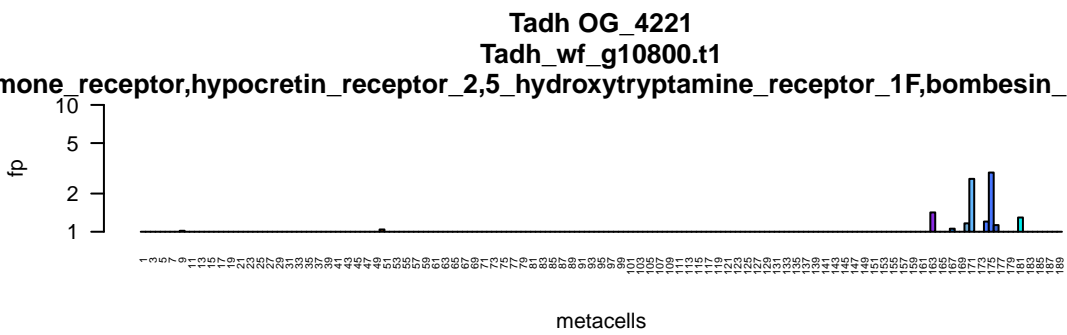
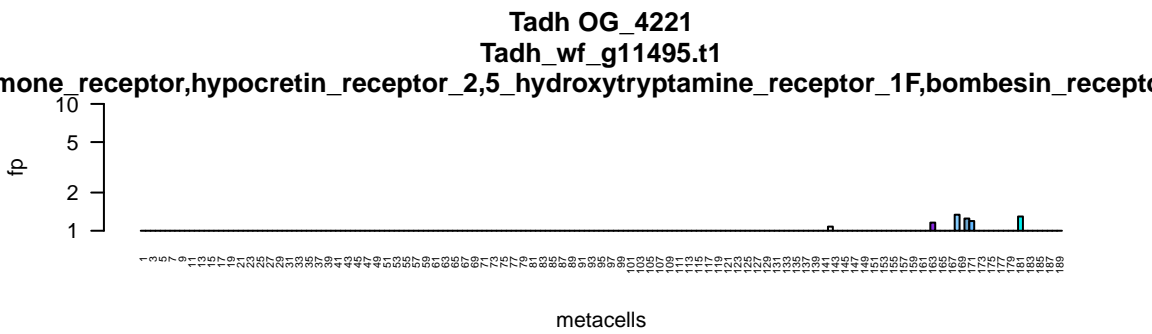
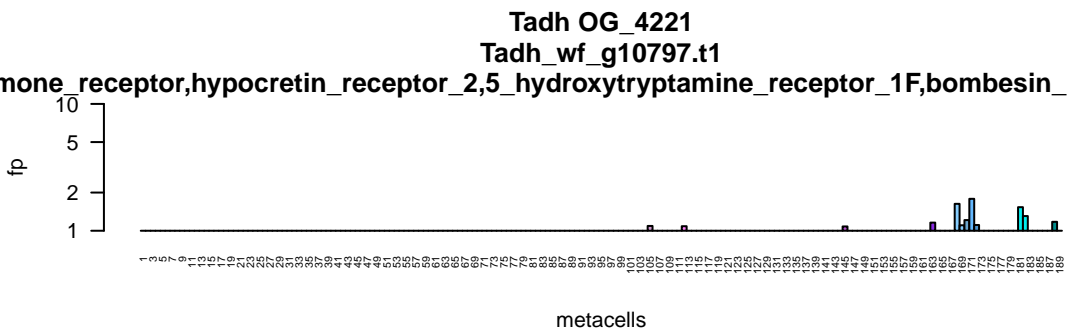
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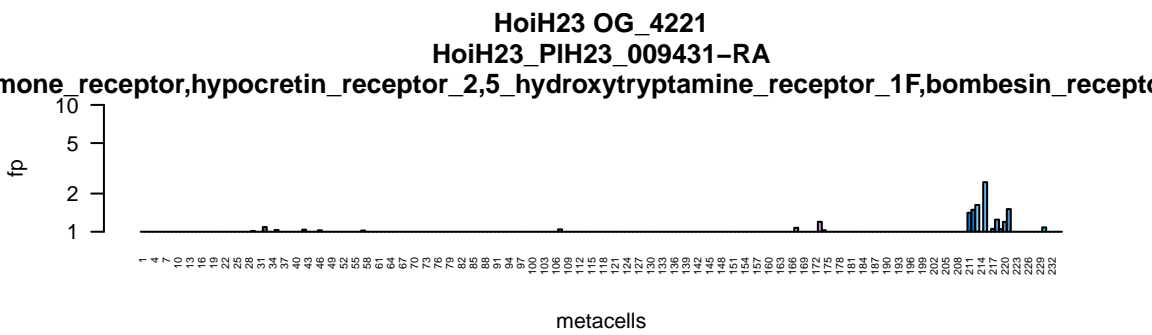
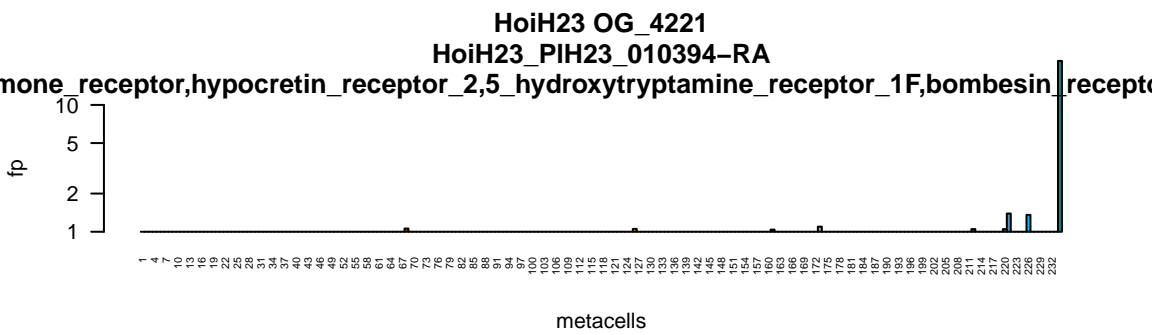
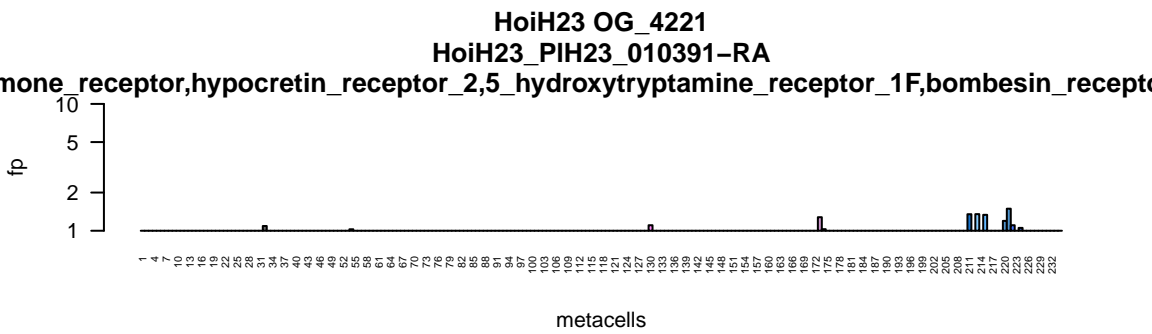
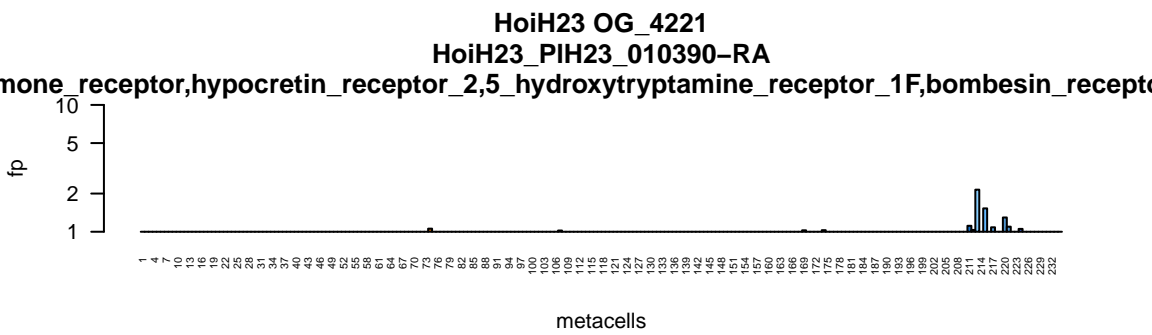
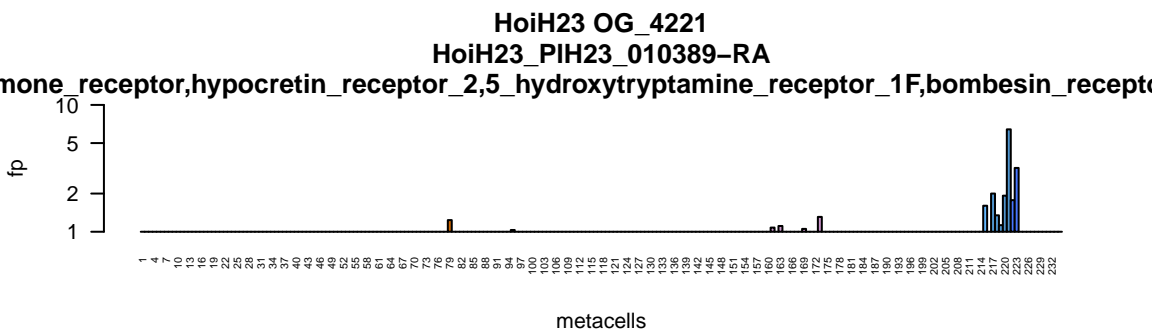
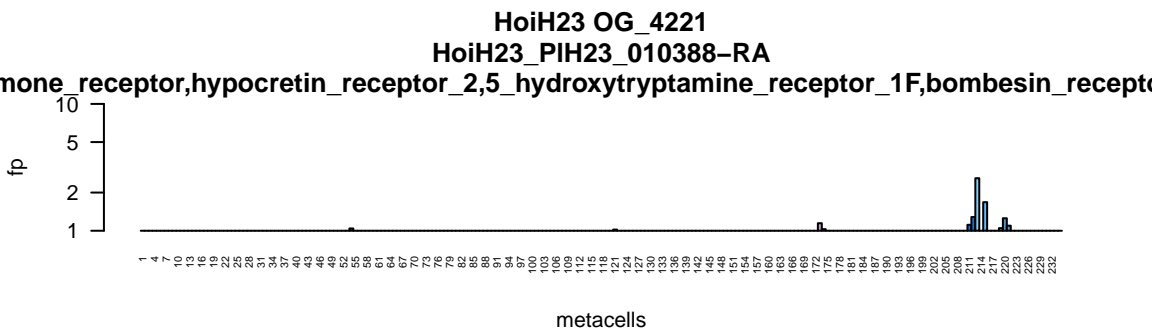


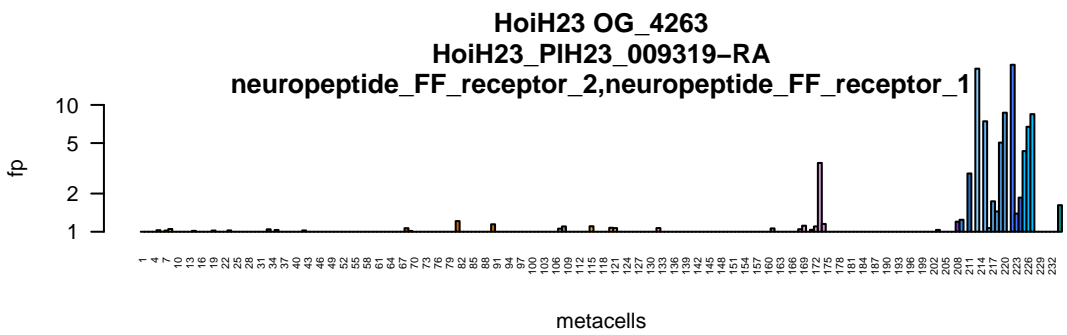
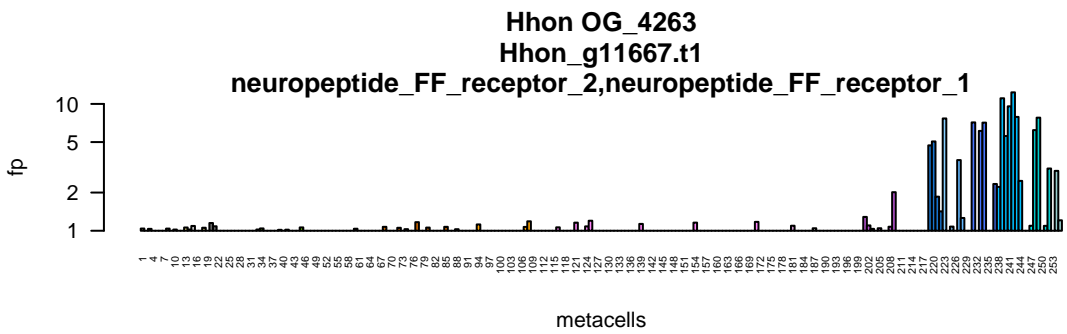
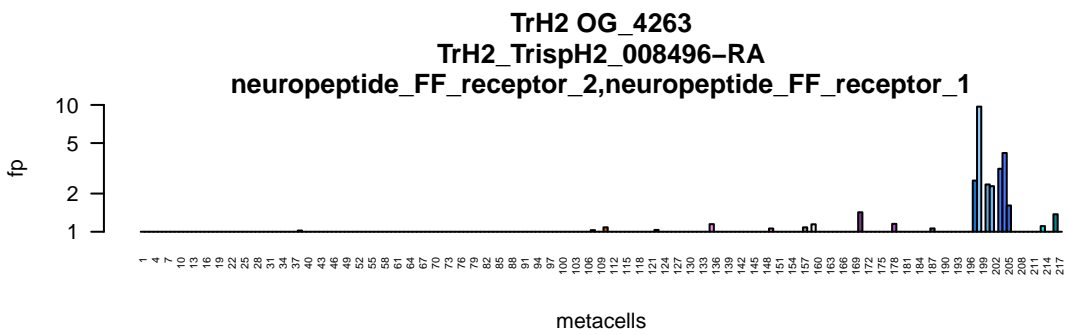
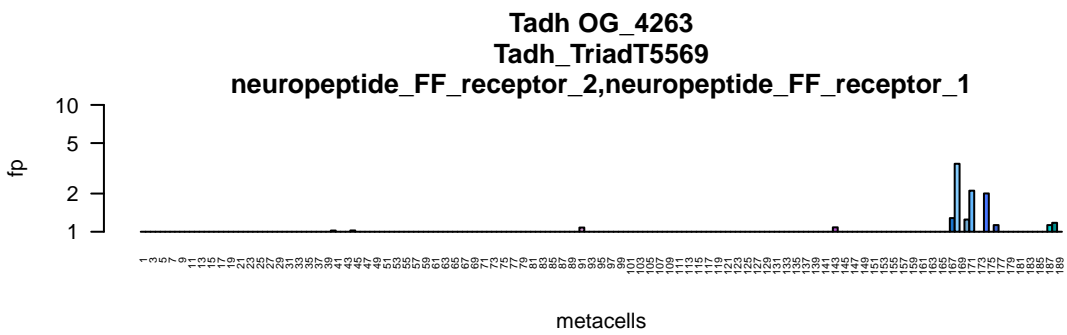
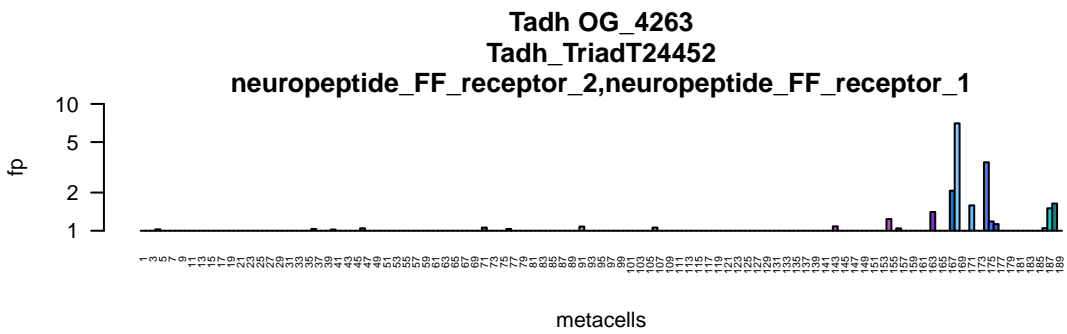
HoiH23 OG_7537
HoiH23_PIH23_000241-RA



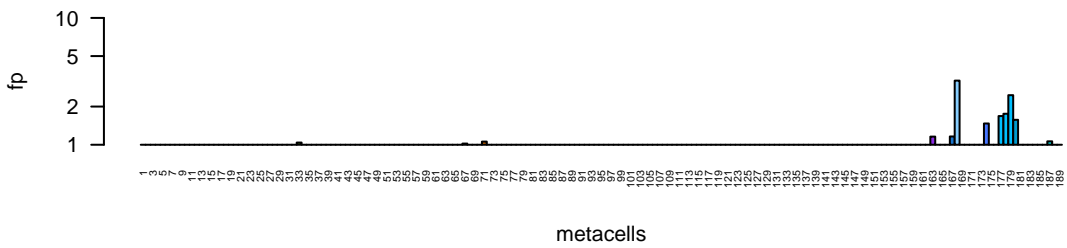






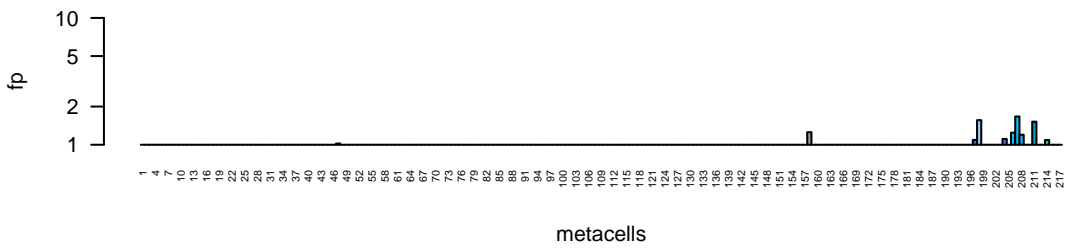


Tadh OG_6120
Tadh_TriadT60881



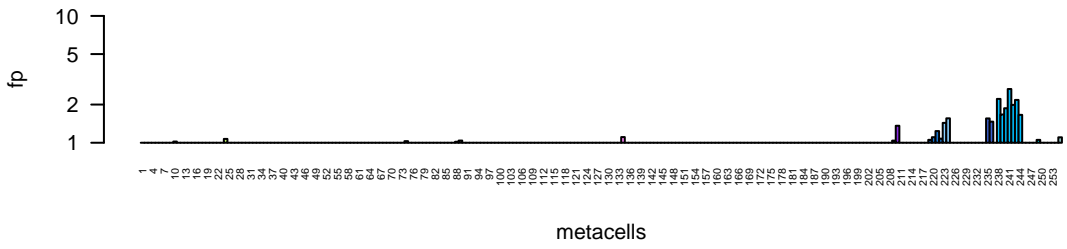
metacells

TrH2 OG_6120
TrH2_TrispH2_008845-RA



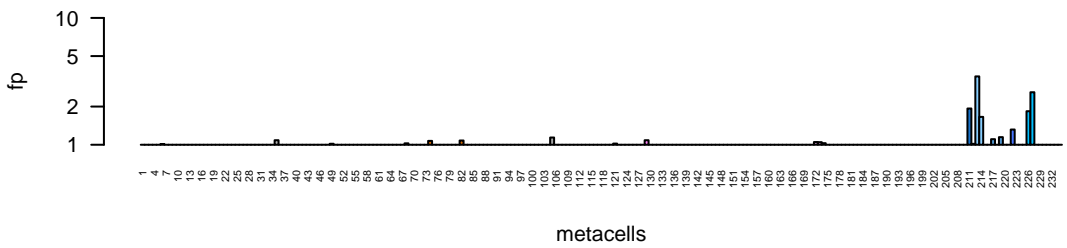
metacells

Hhon OG_6120
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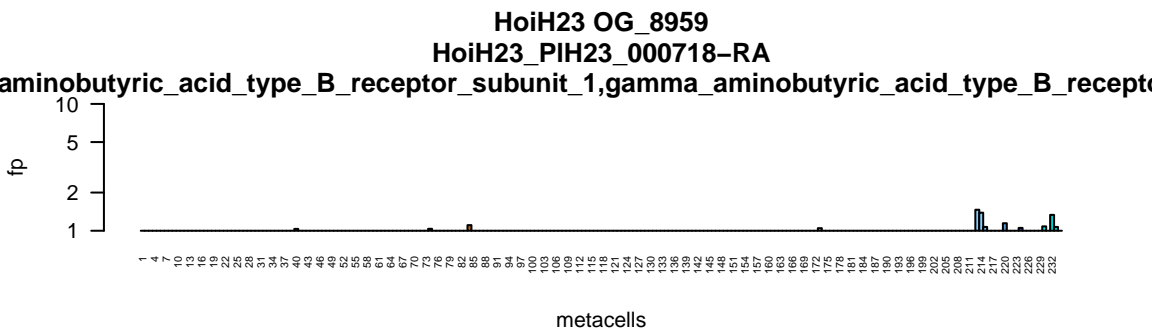
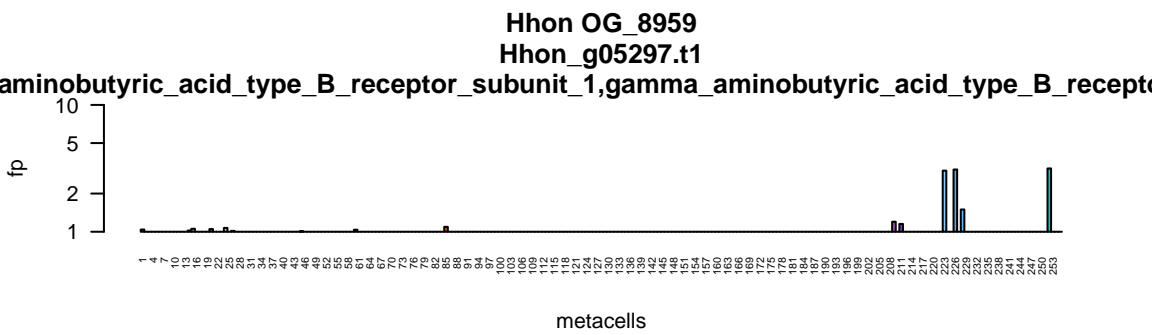
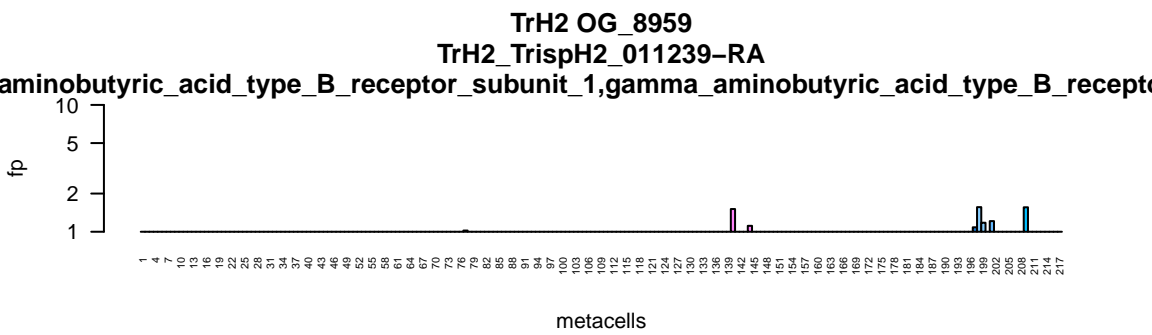
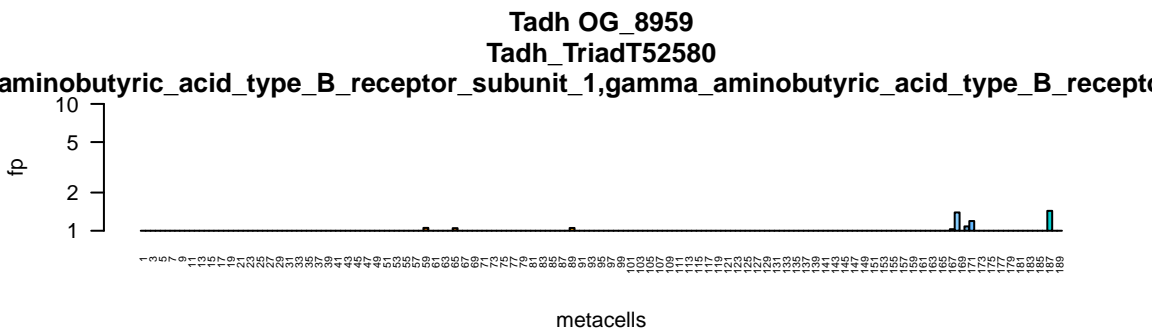


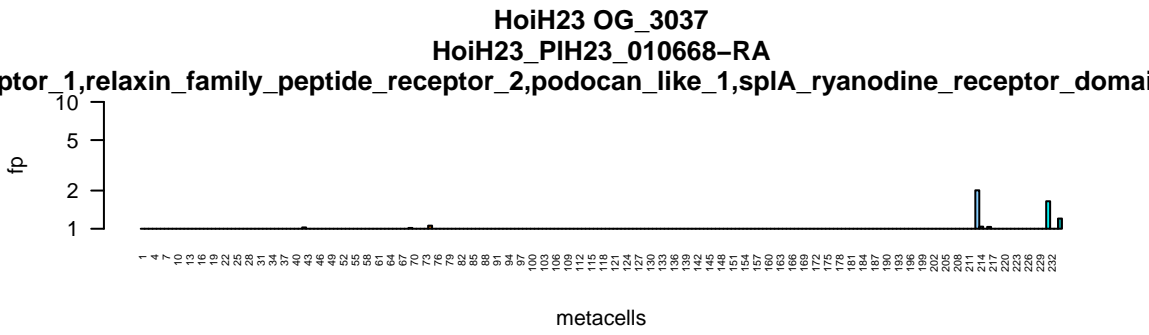
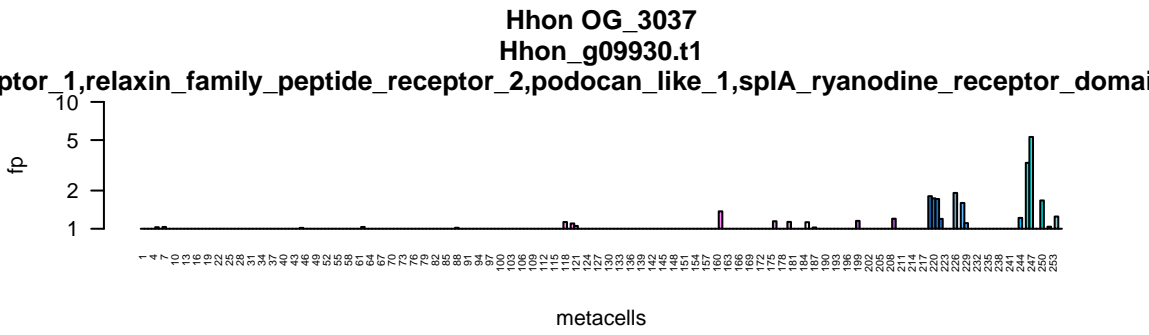
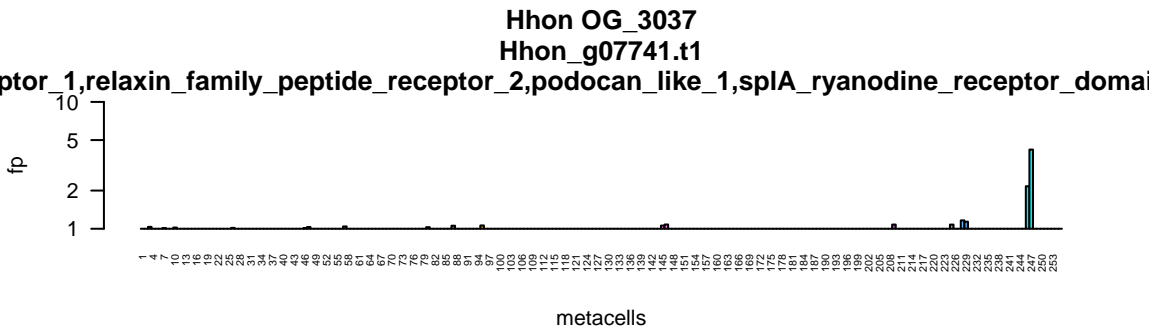
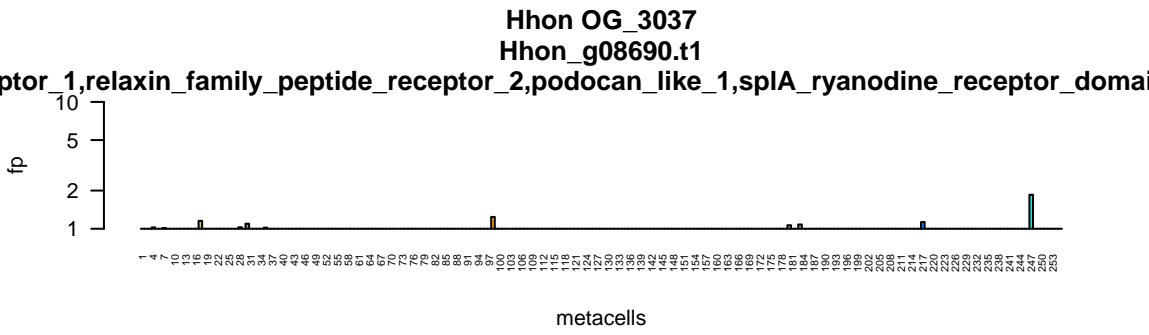
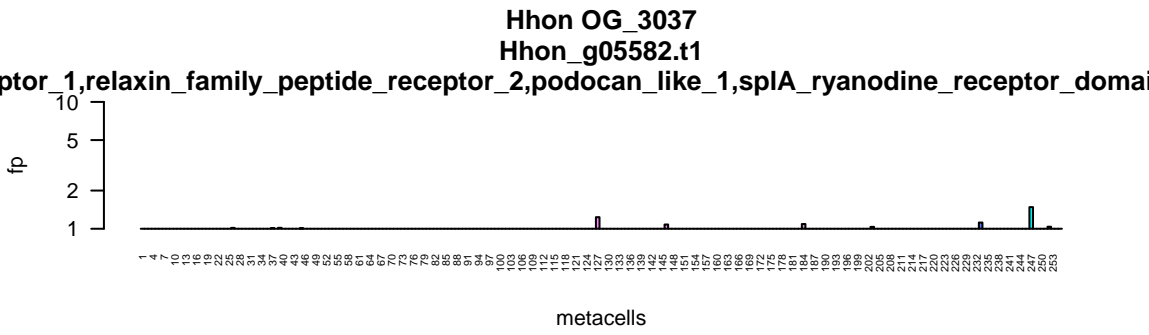
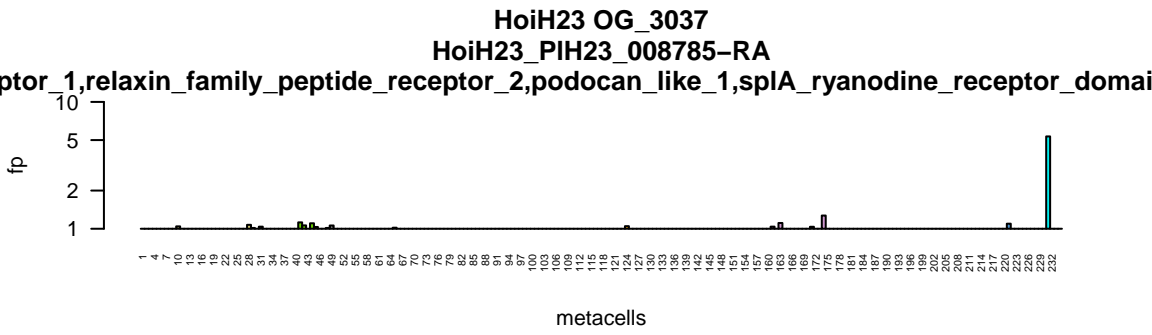
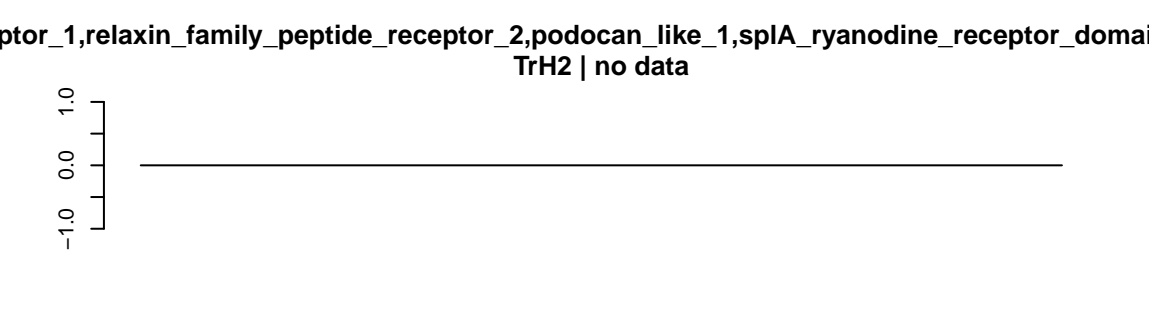
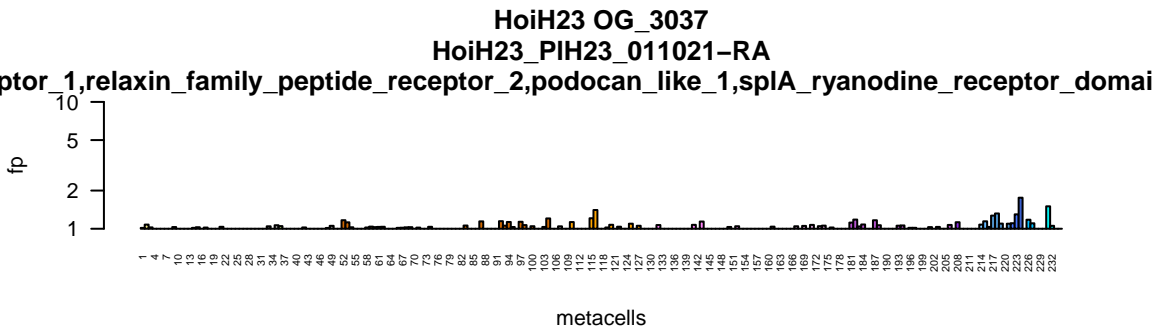
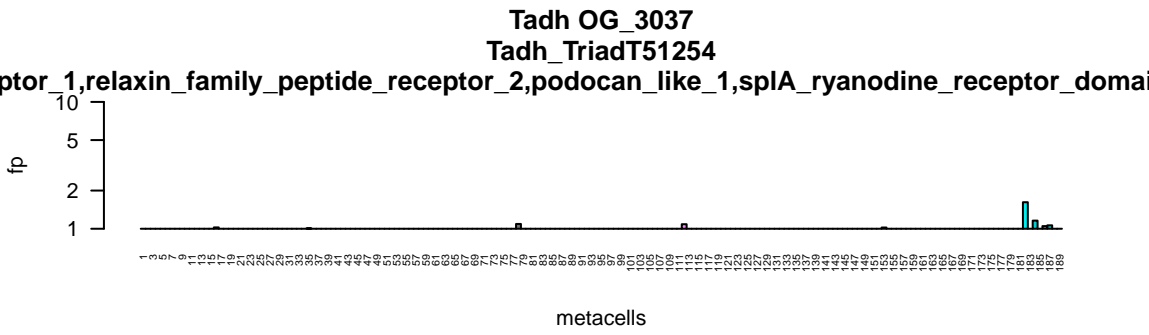
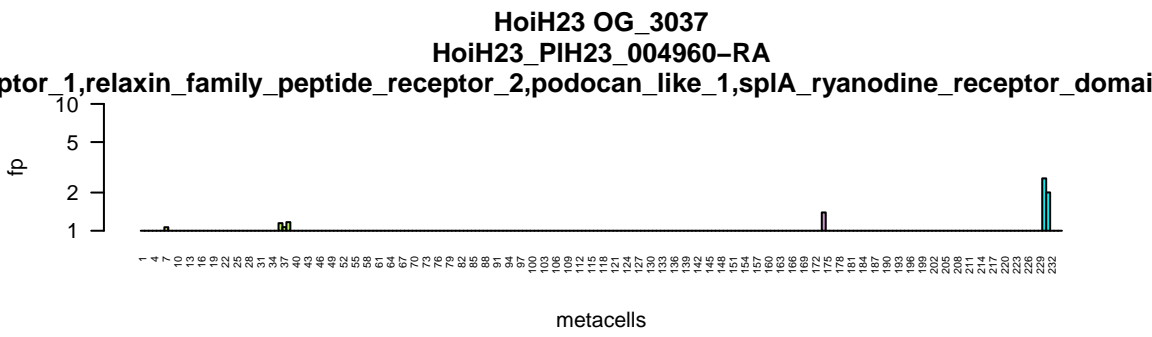
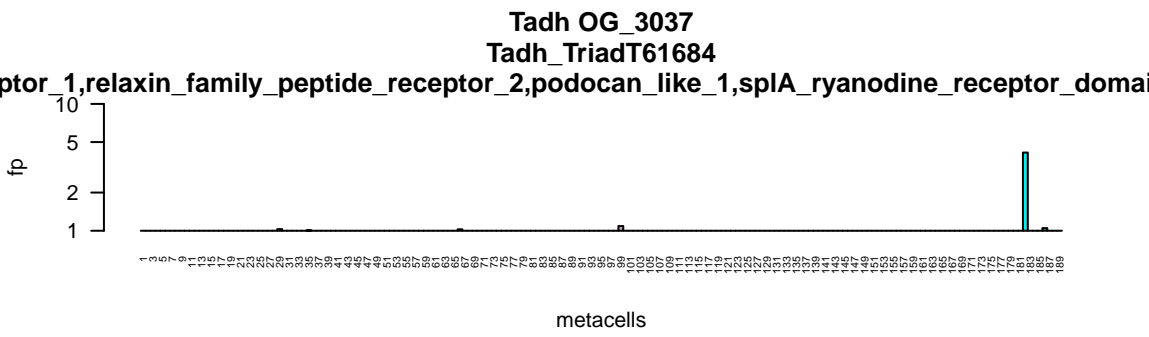
metacells

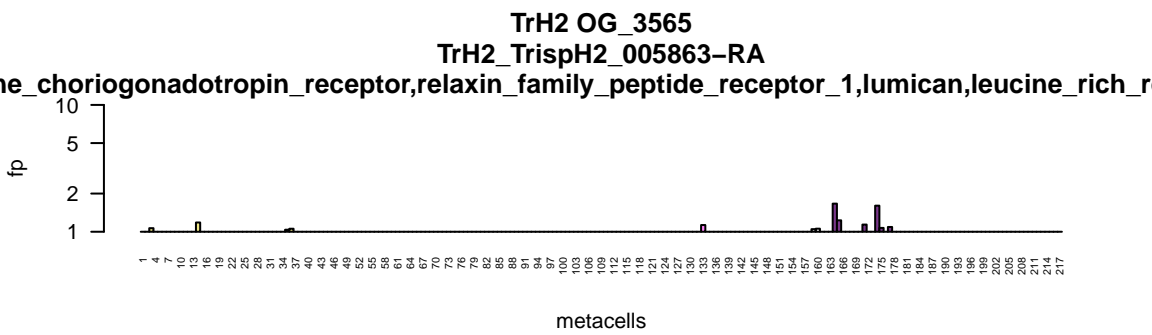
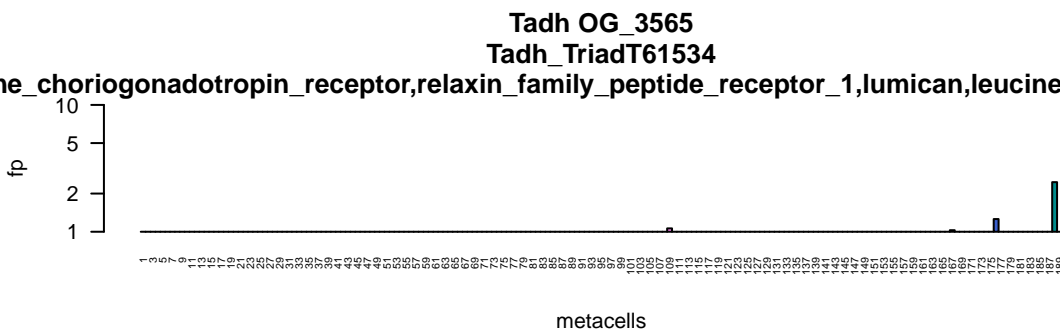
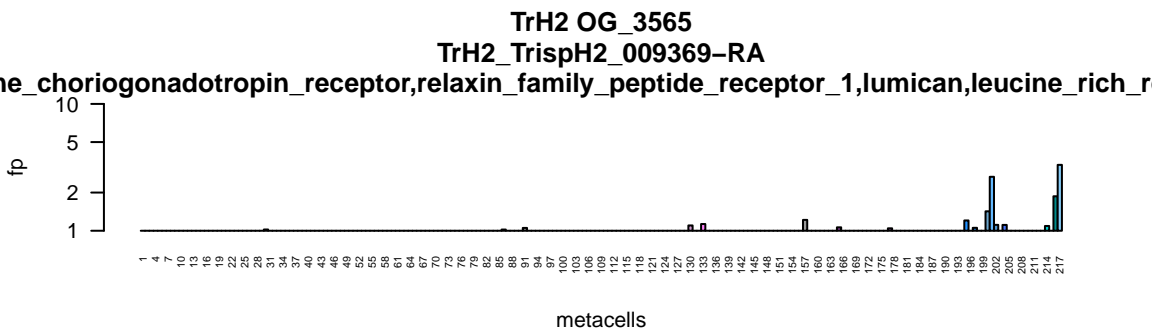
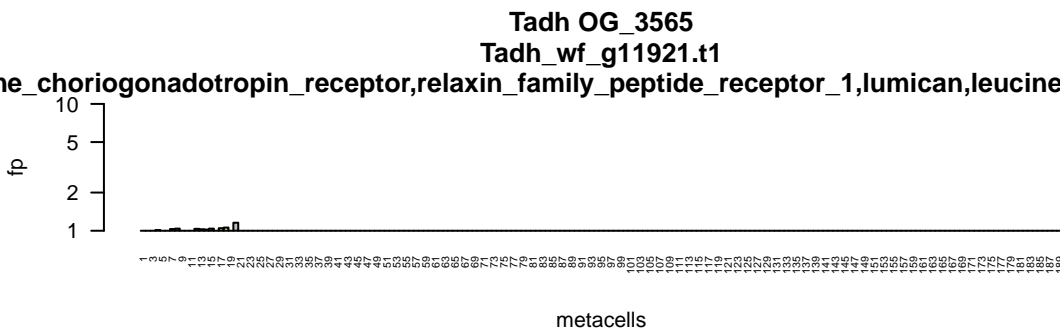
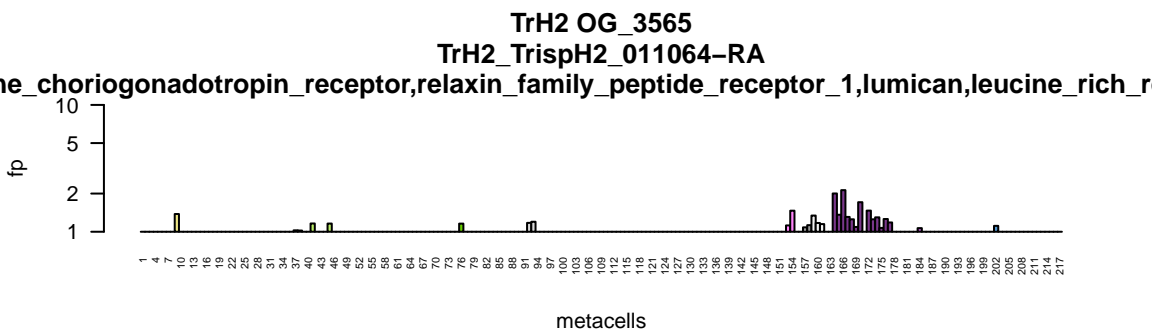
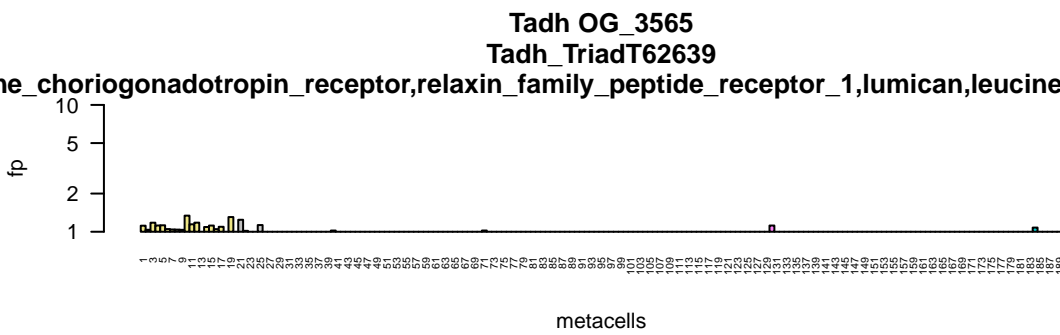
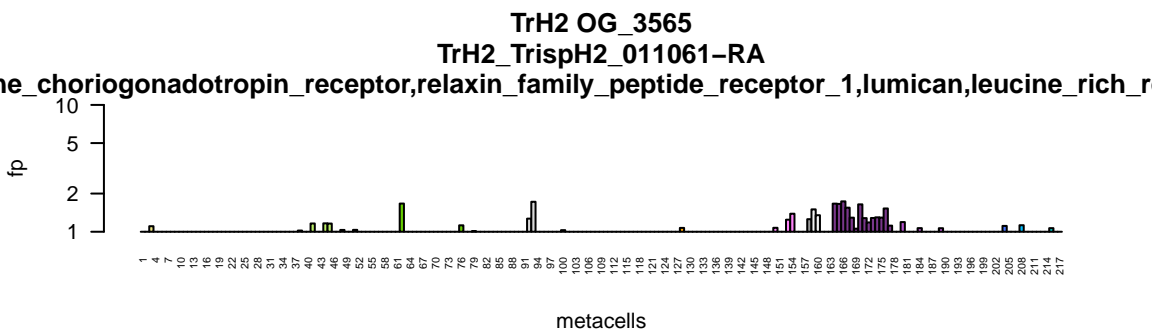
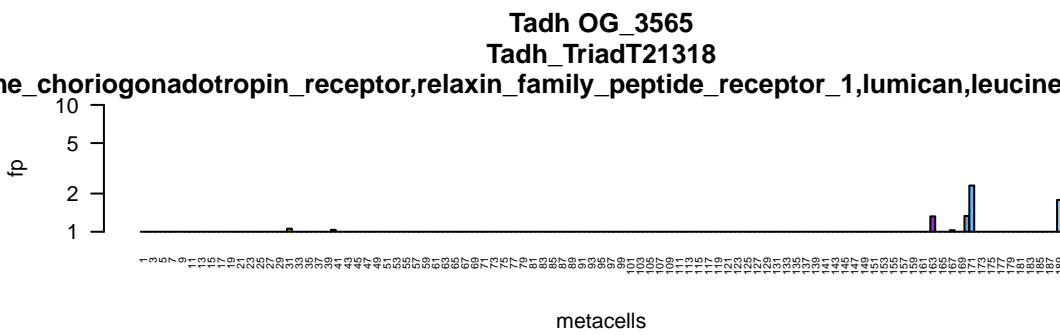
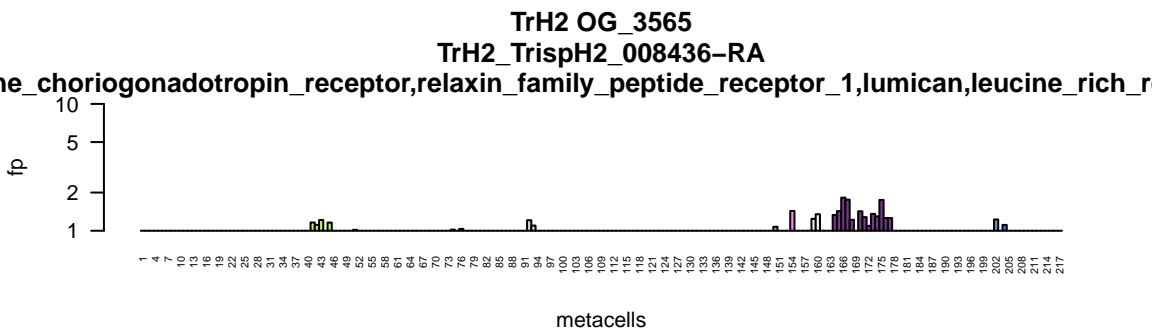
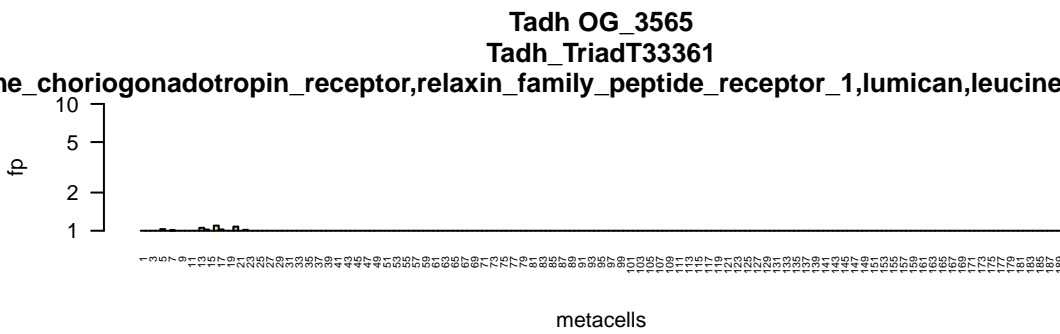
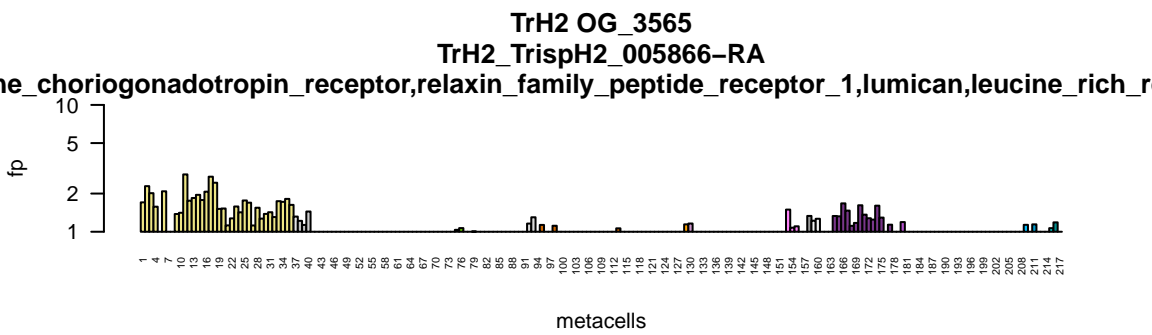
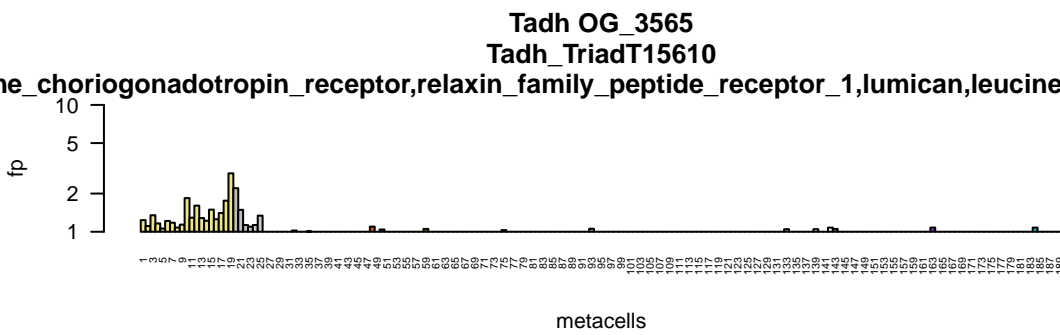
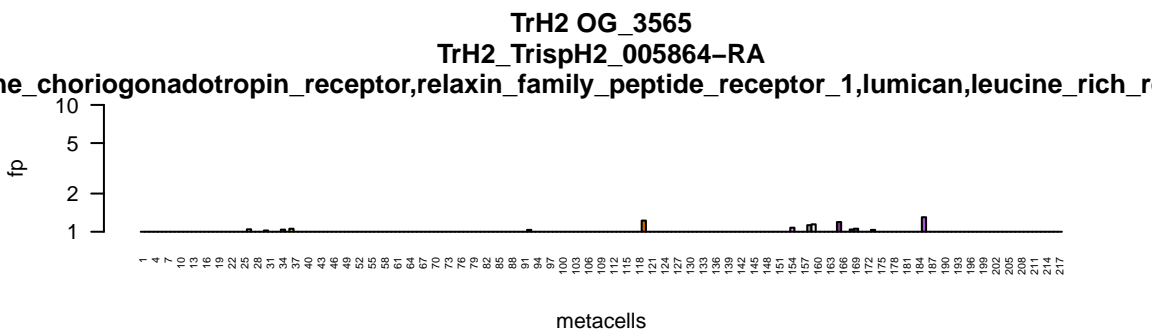
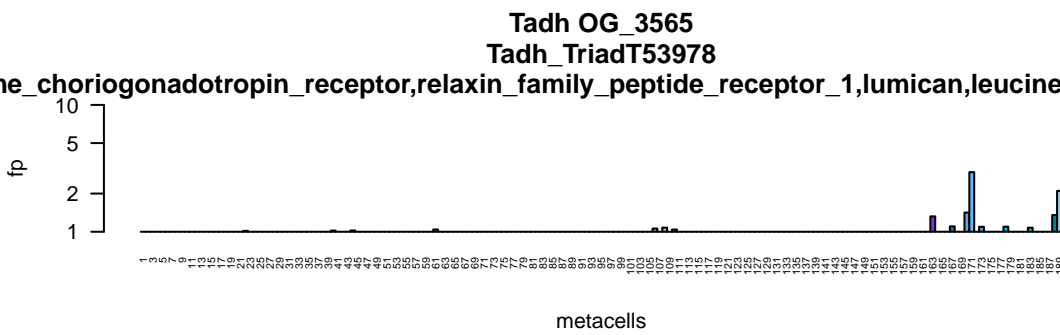
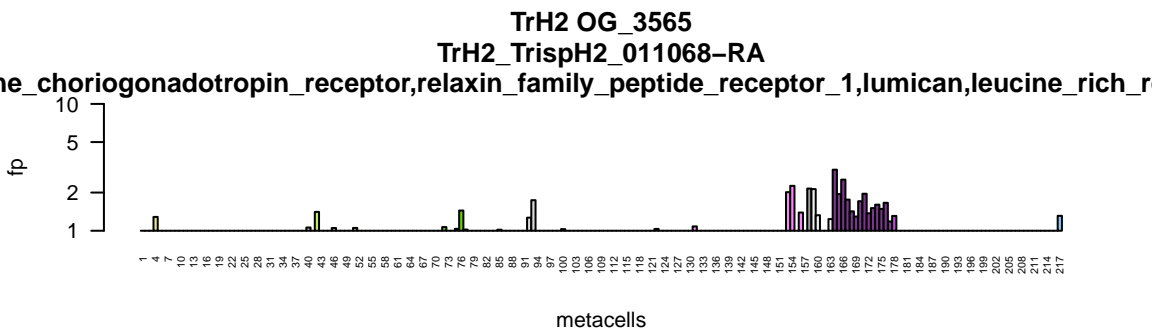
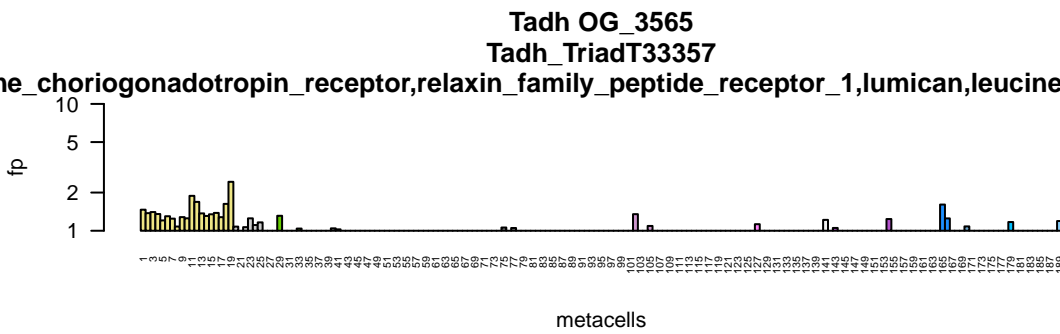
HoiH23 OG_6120
HoiH23_PIH23_008464-RA

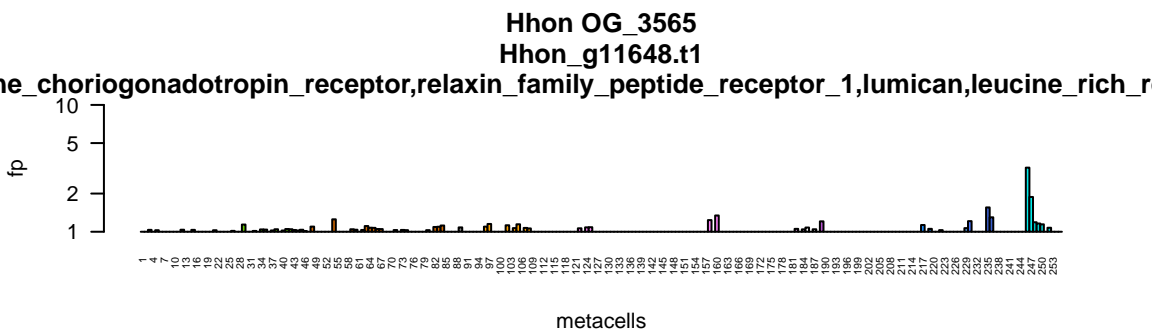
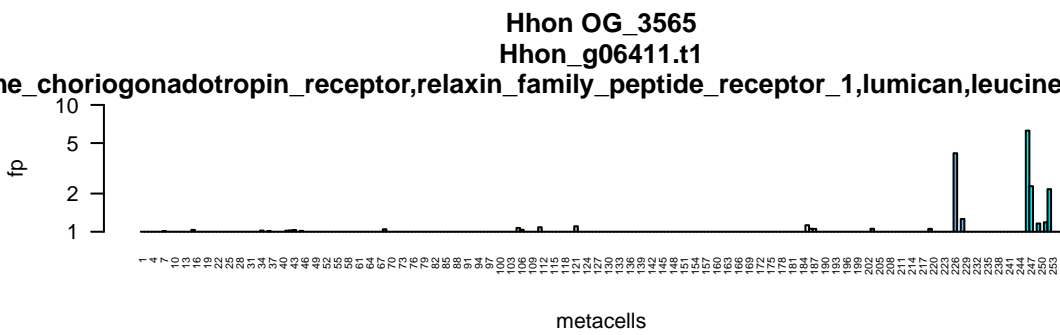
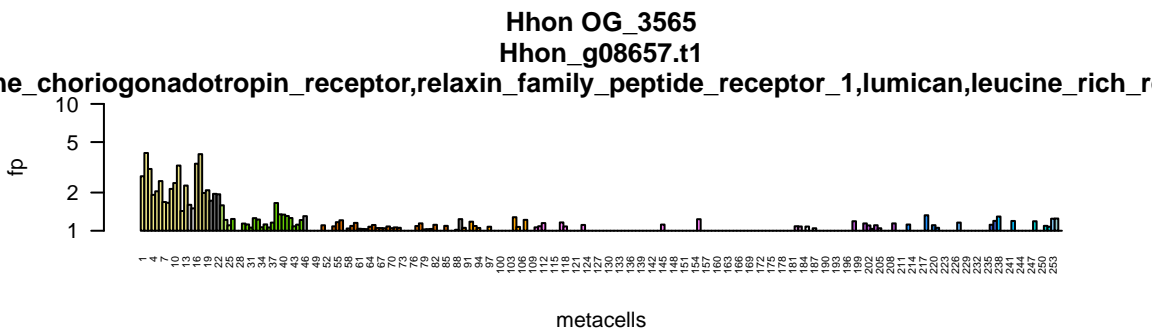
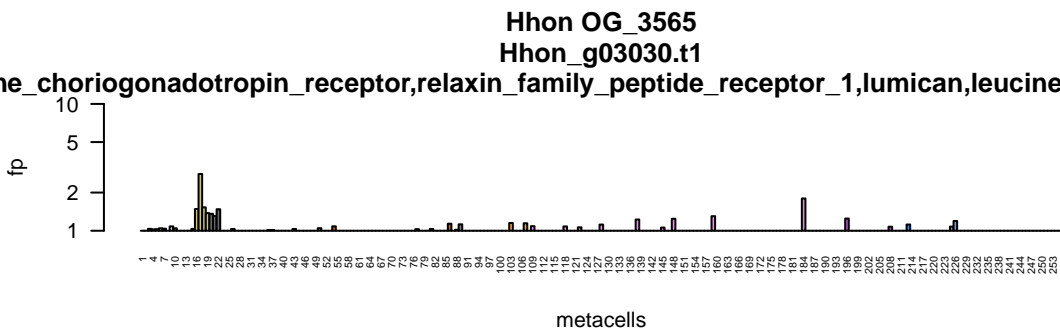
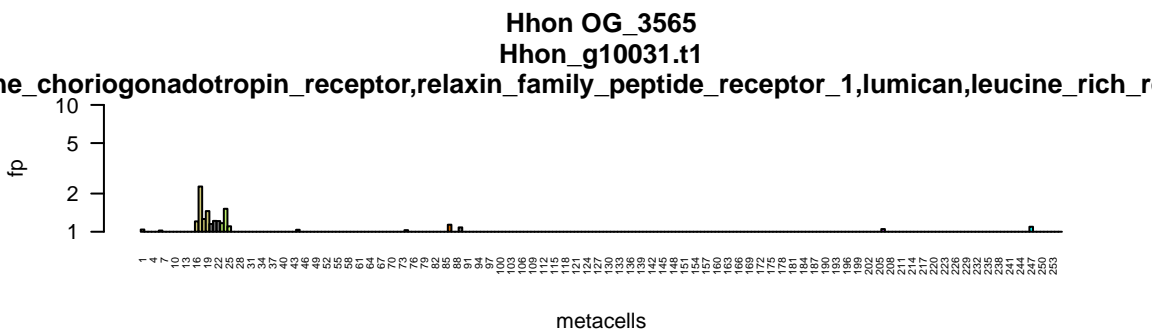
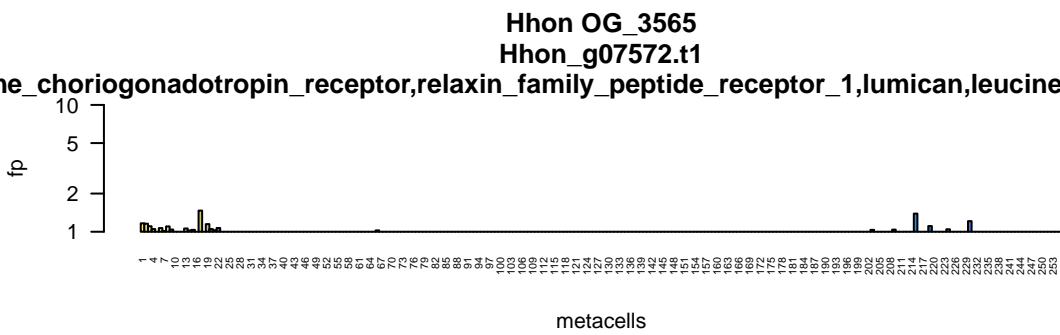
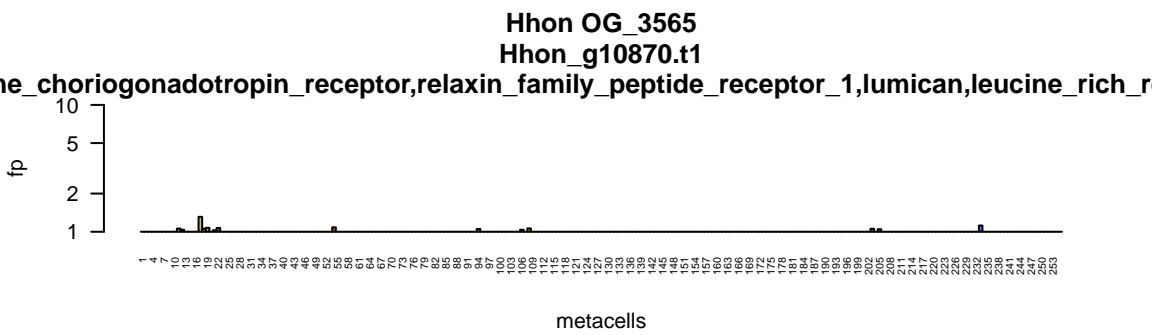
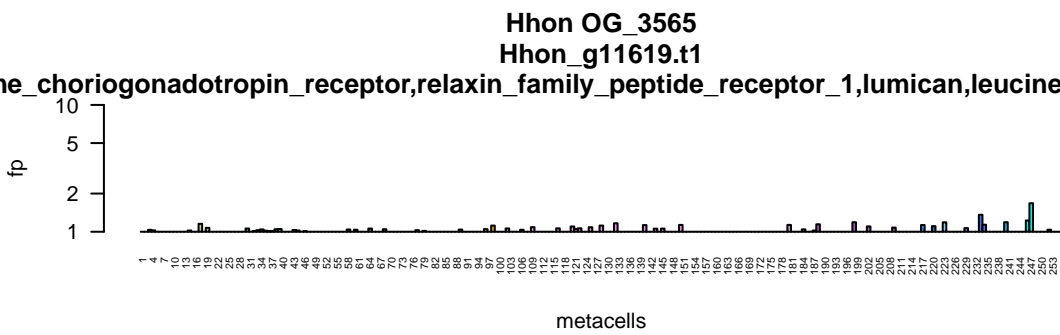
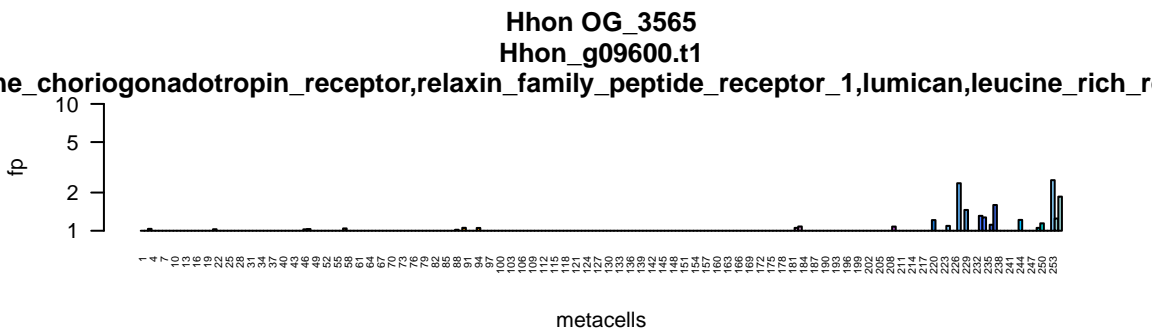
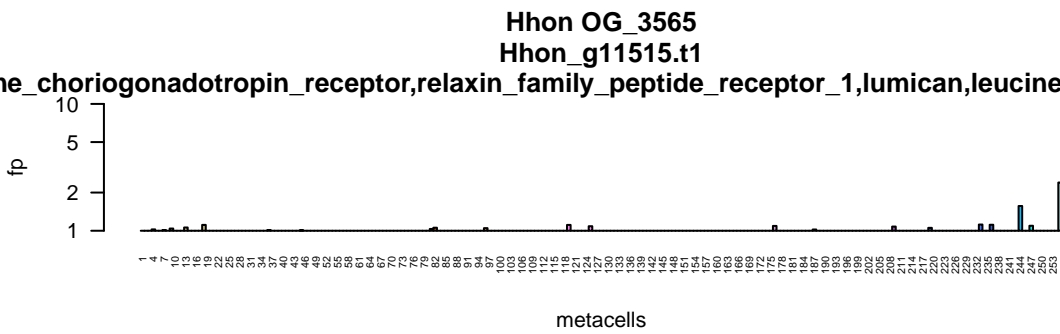
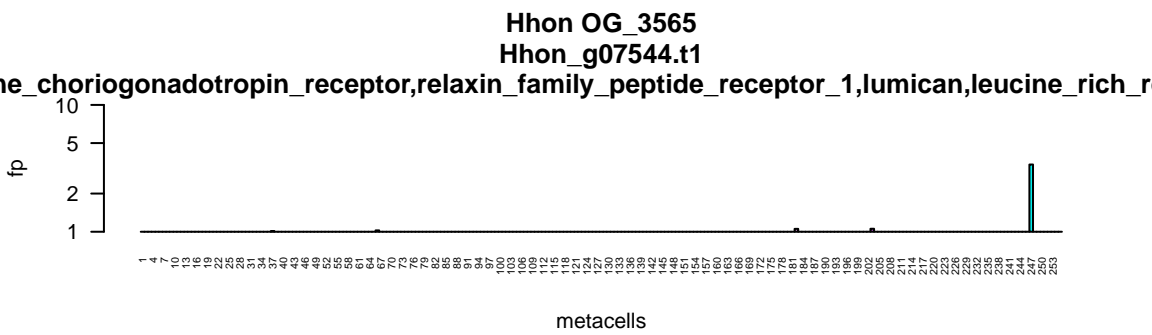
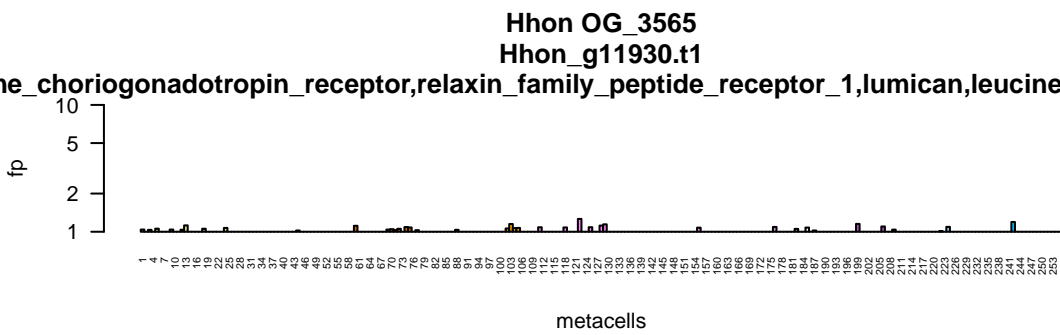
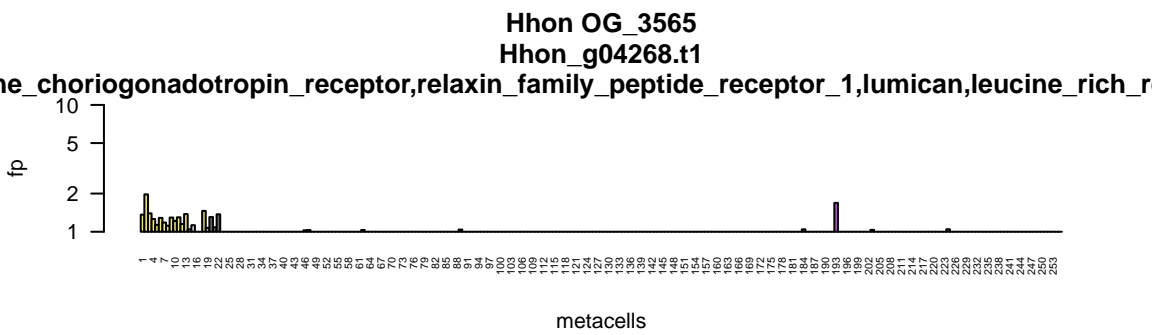
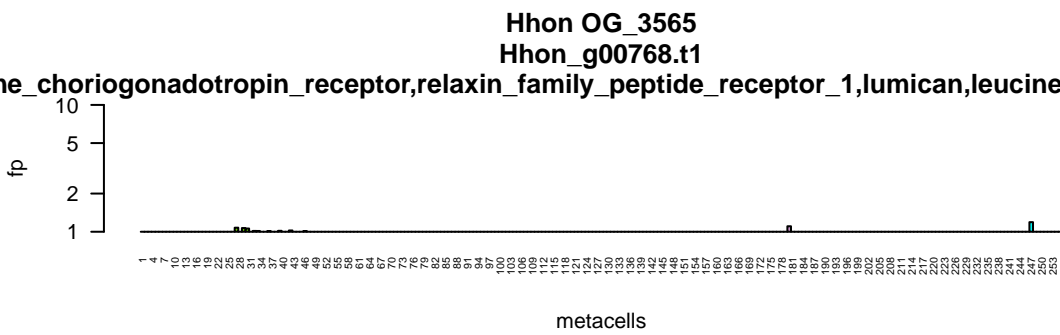
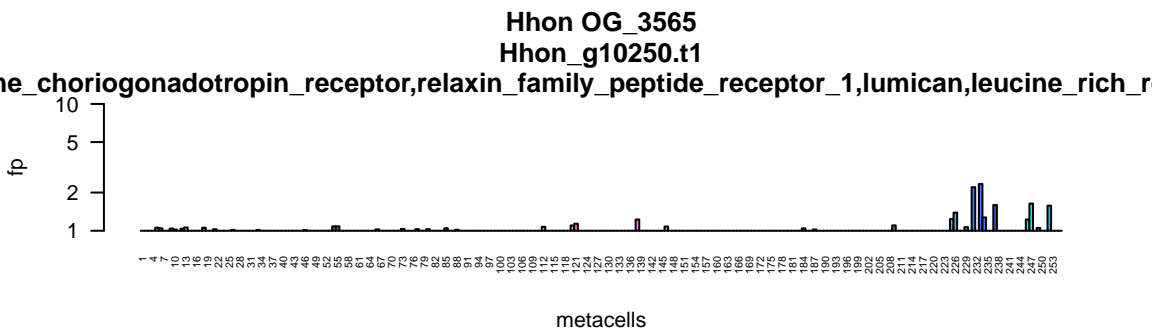
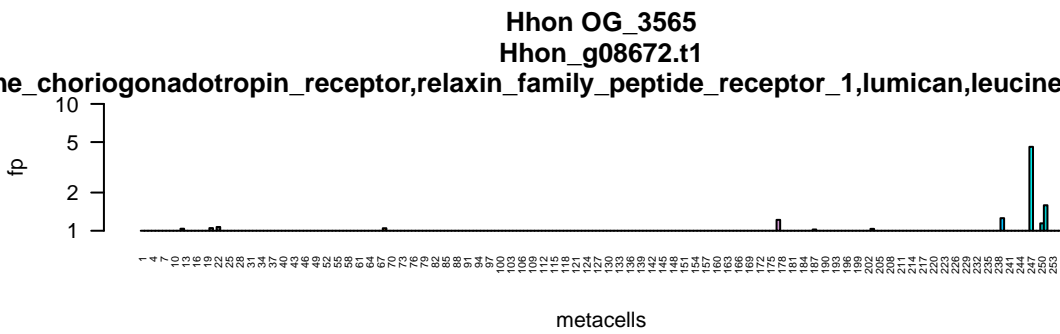


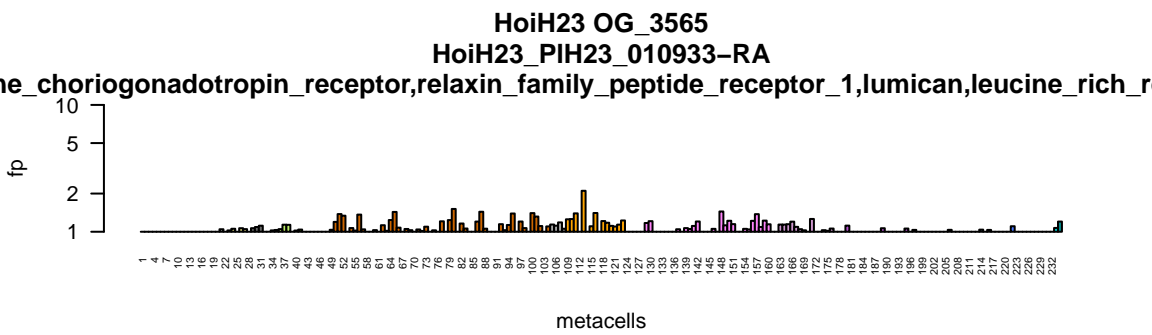
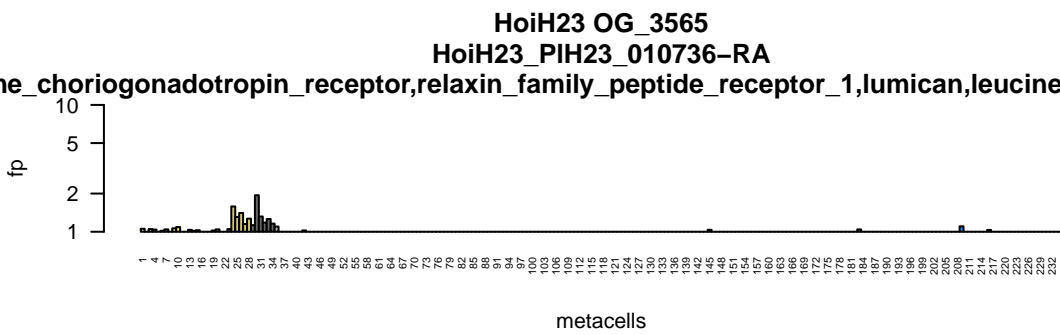
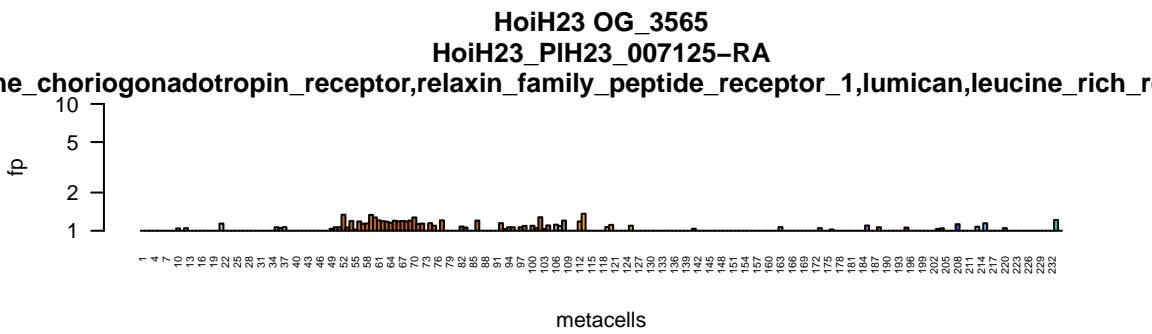
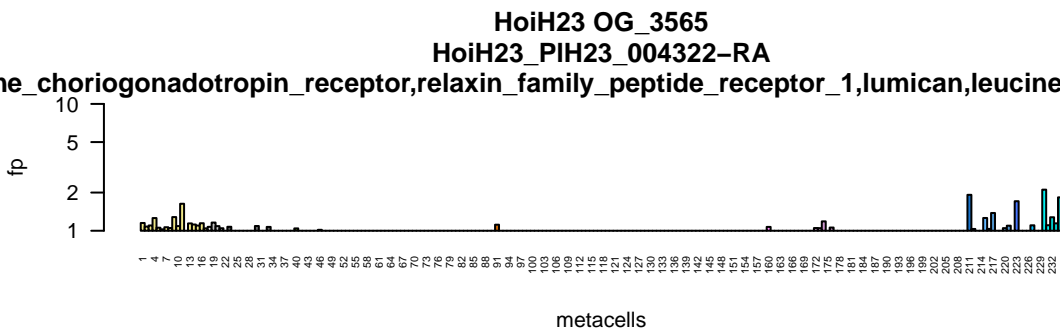
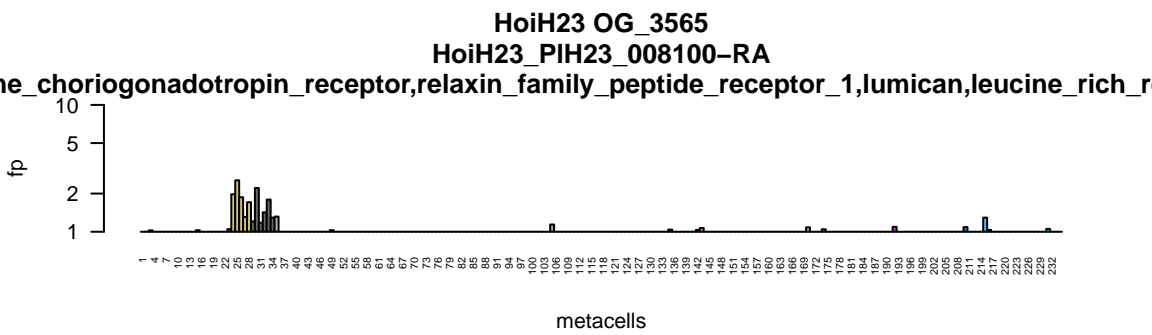
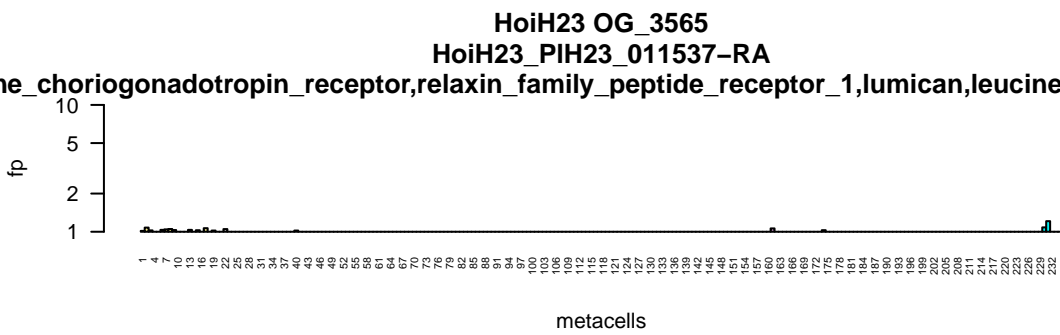
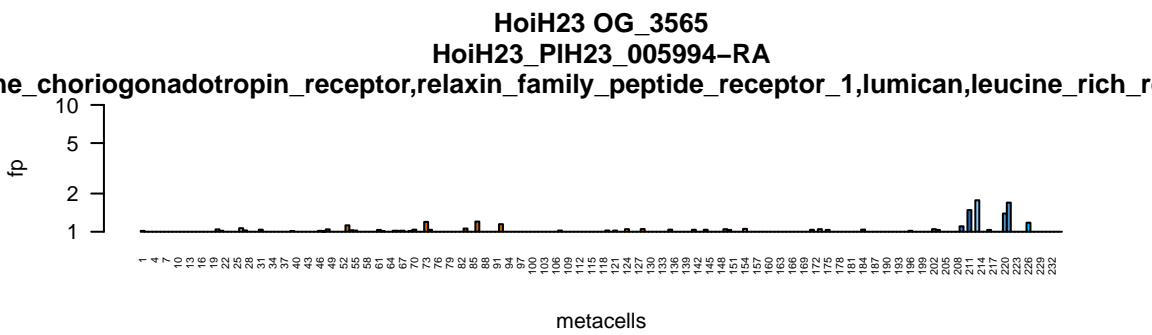
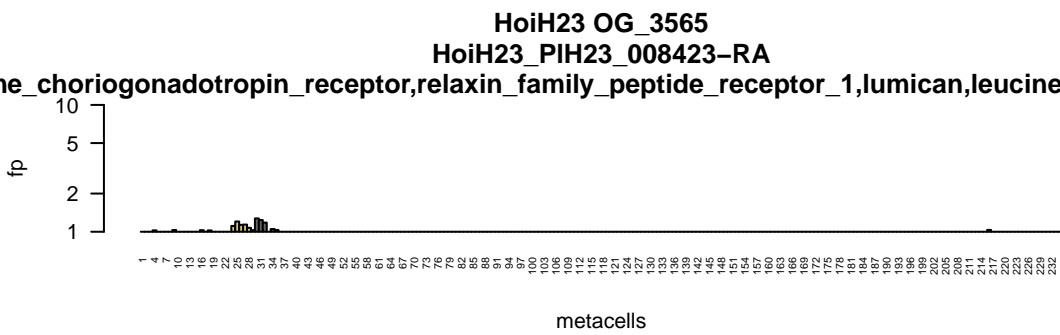
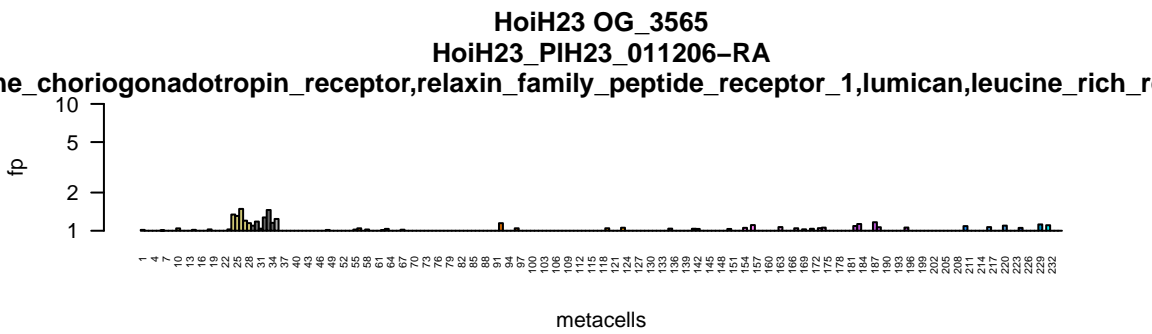
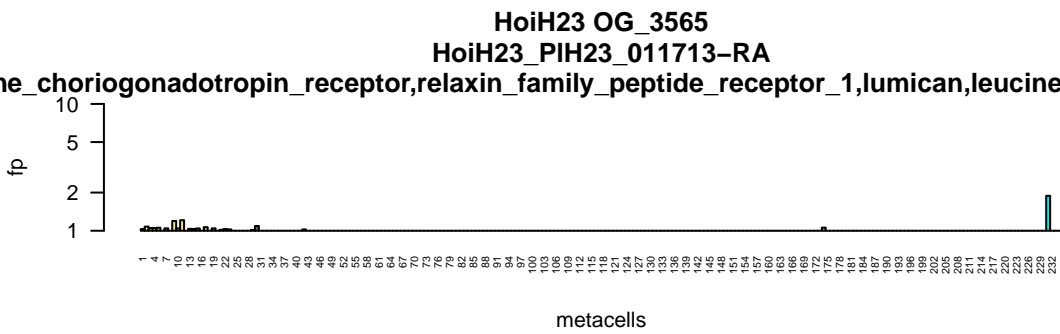
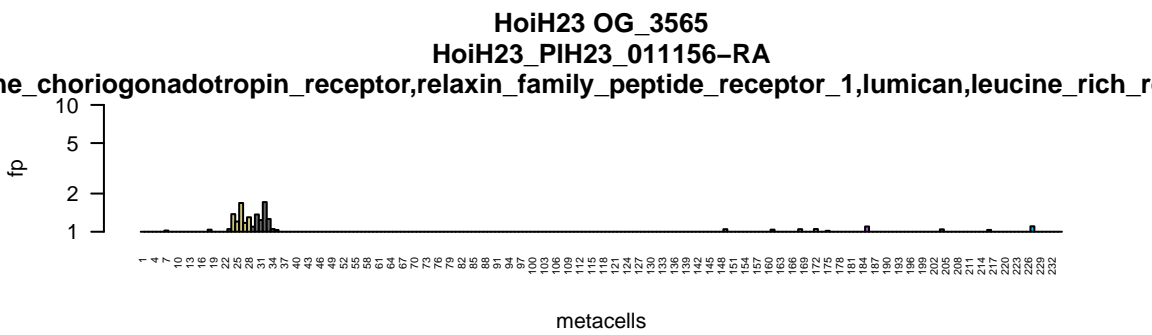
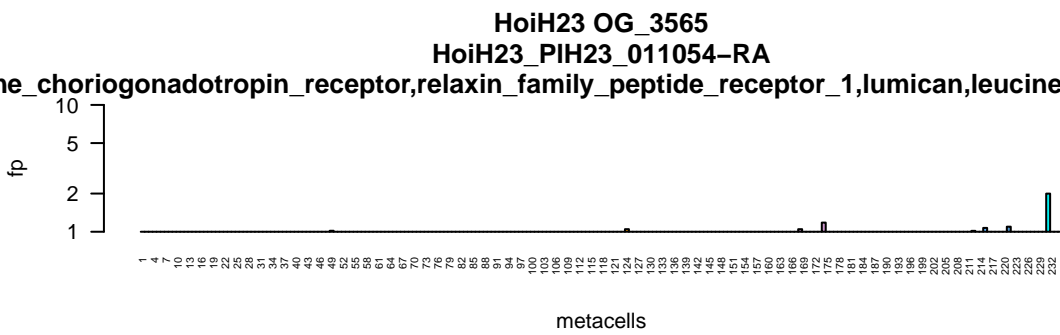
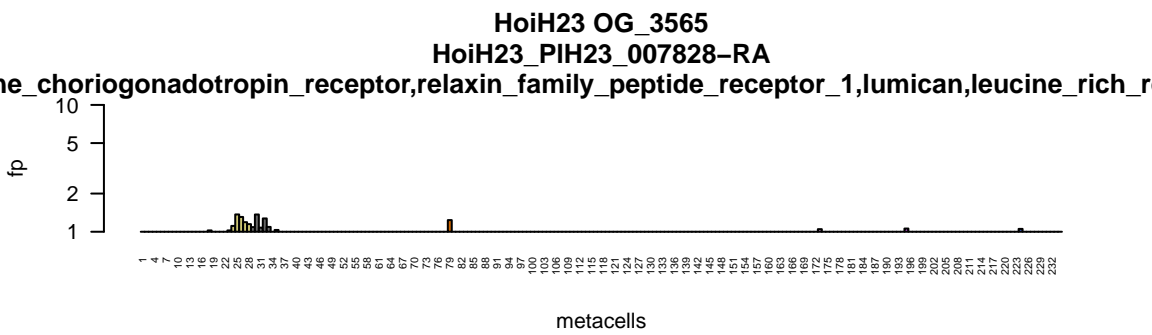
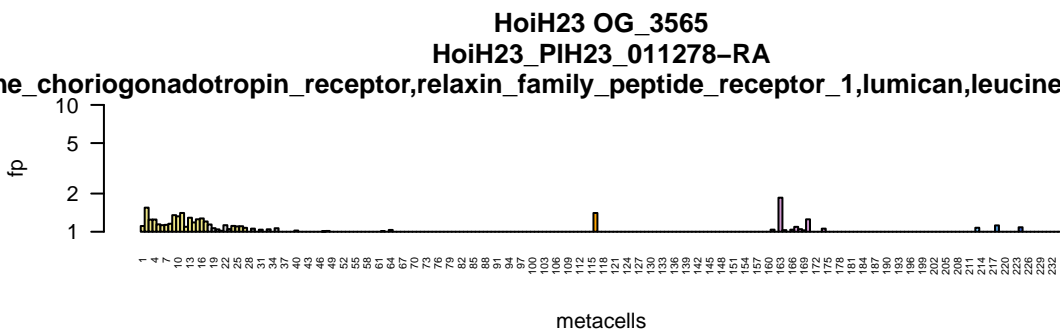
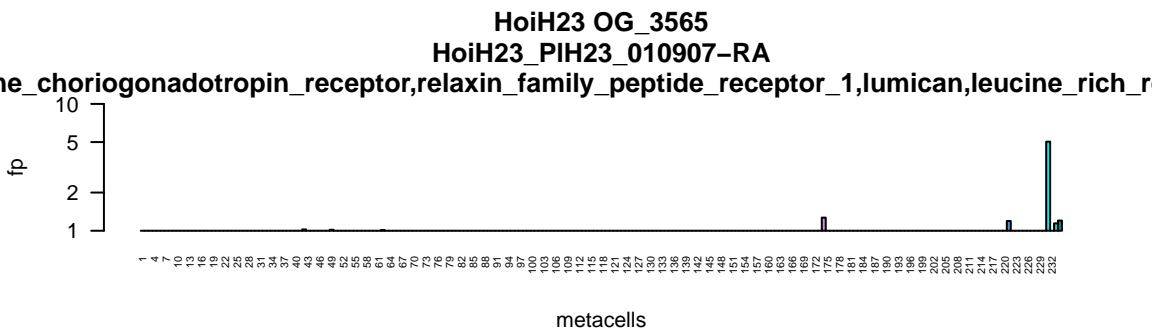
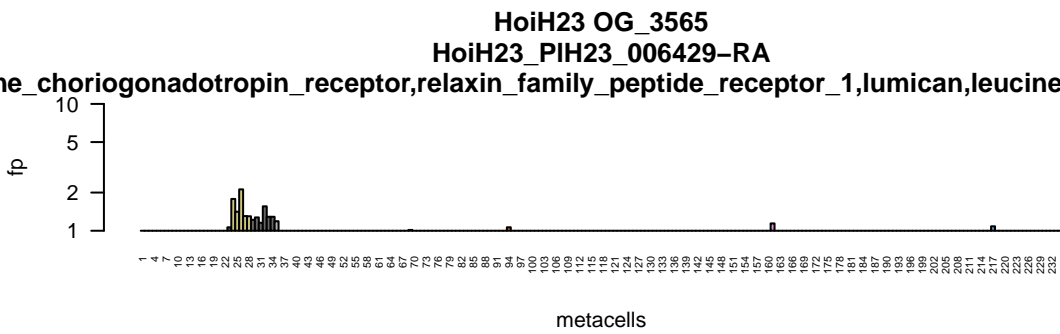
metacells

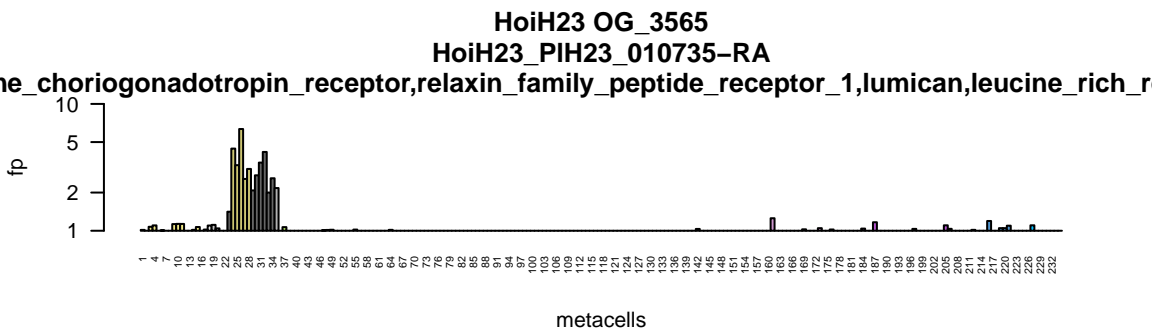
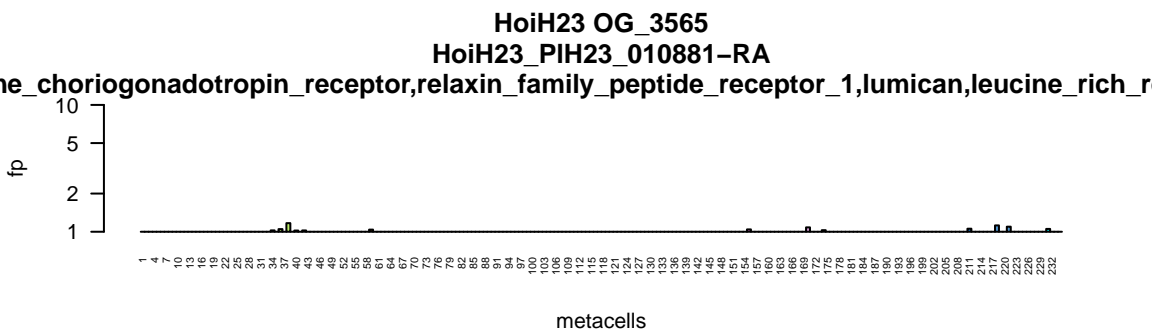
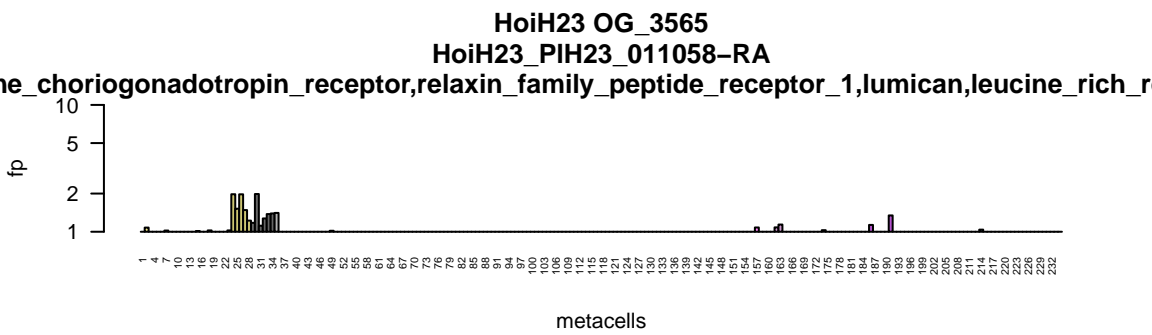
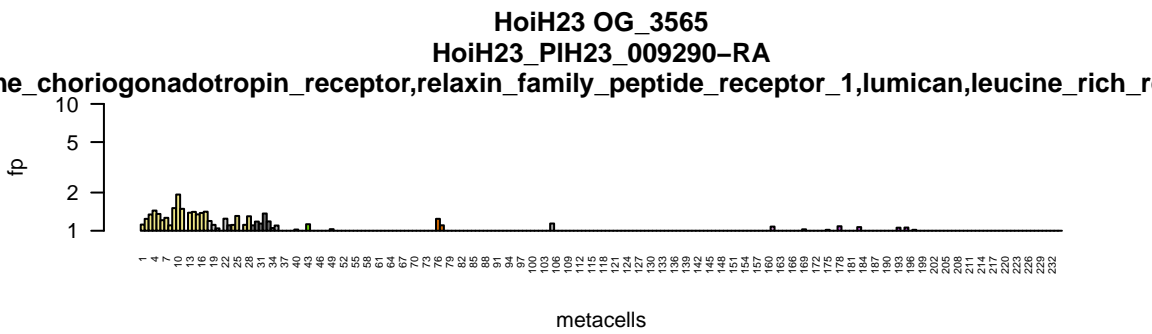


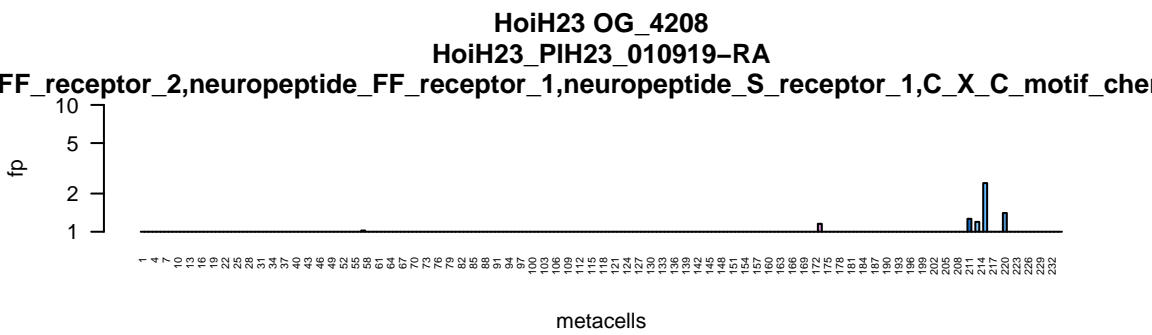
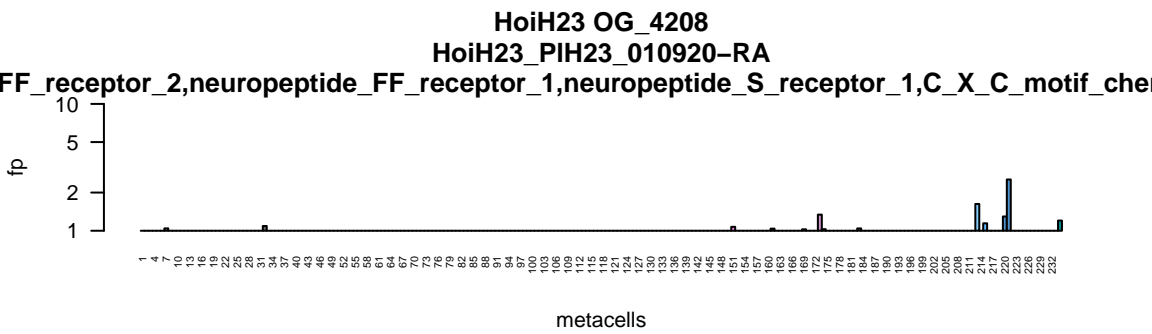
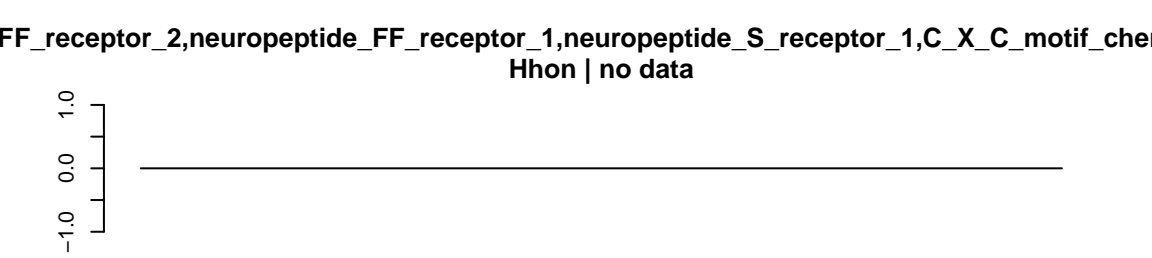
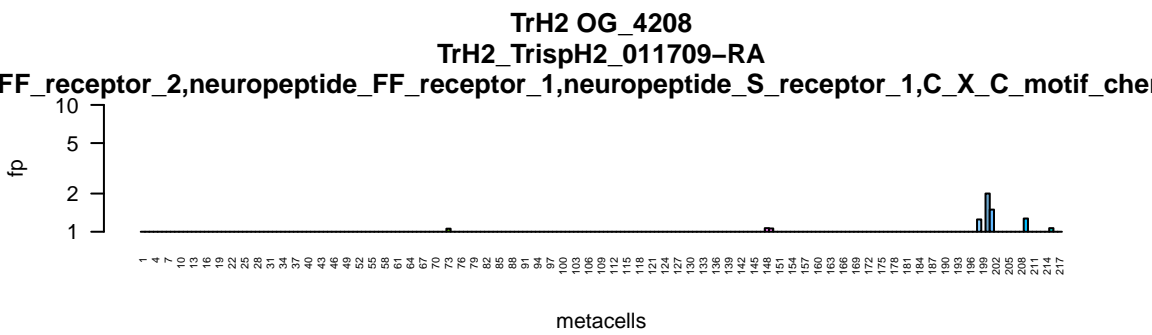
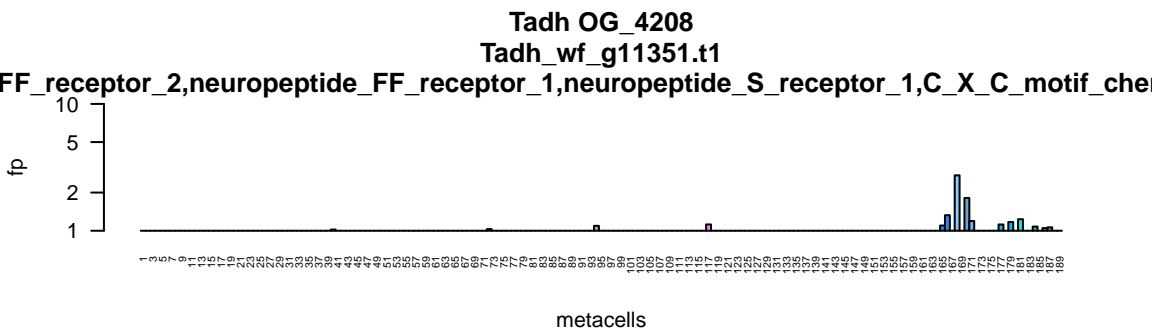








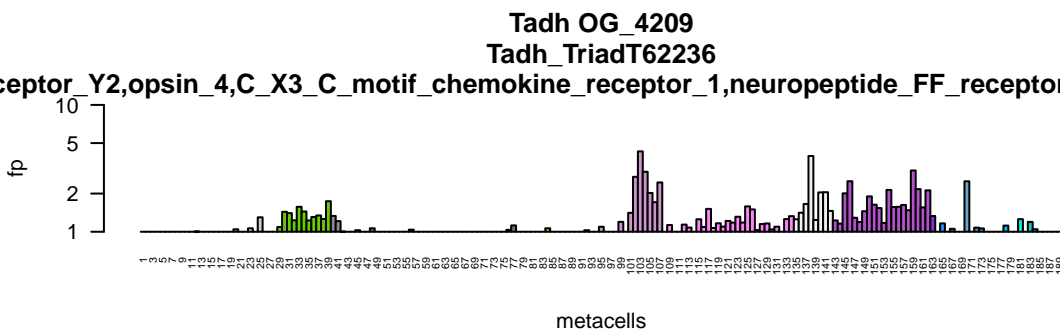




ceptor_Y2,opsin_4,C_X3_C_motif_chemokine_receptor_1,neuropeptide_FF_receptor

fp

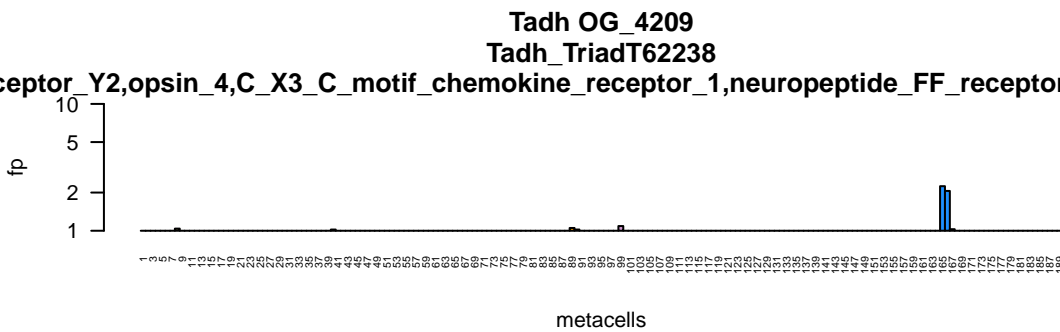
metacells



ceptor_Y2,opsin_4,C_X3_C_motif_chemokine_receptor_1,neuropeptide_FF_receptor

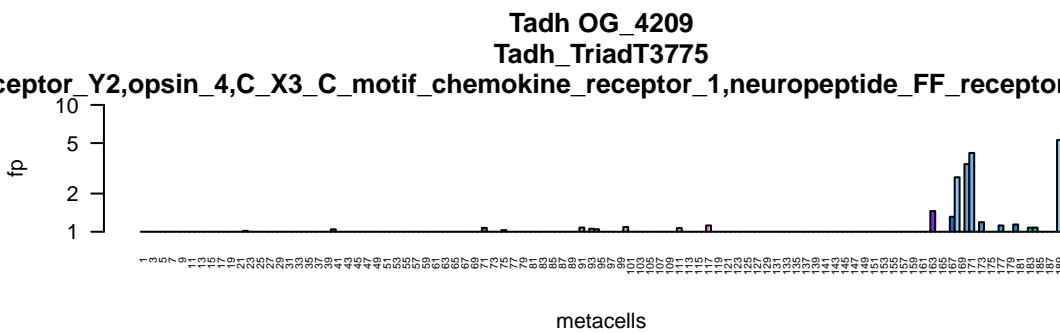
fp

metacells



ceptor_Y2,opsin_4,C_X3_C_motif_chemokine_receptor_1,neuropeptide_FF_receptor

metacells

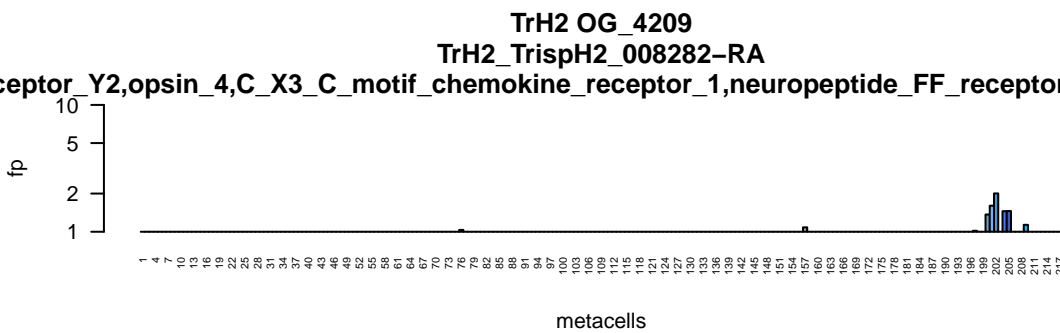


ceptor_Y2,opsin_4,C_X3_C_motif_chemokine_receptor_1,neuropeptide_FF_receptor

fp

metacells

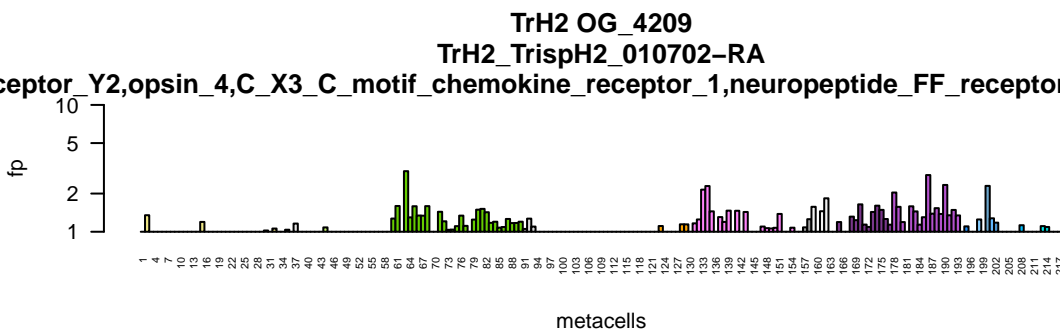
Motif	fp
1	1
2	1
3	1
4	1
5	1
6	1
7	1
8	1
9	1
10	1
11	1
12	1
13	1
14	1
15	1
16	1
17	1
18	1
19	1
20	1
21	1
22	1
23	1
24	1
25	1
26	1
27	1
28	1
29	1
30	1
31	1
32	1
33	1
34	1
35	1
36	1
37	1
38	1
39	1
40	1
41	1
42	1
43	1
44	1
45	1
46	1
47	1
48	1
49	1
50	1
51	1
52	1
53	1
54	1
55	1
56	1
57	1
58	1
59	1
60	1
61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



ceptor_Y2,opsin_4,C_X3_C_motif_chemokine_receptor_1,neuropeptide_FF_receptor

fp

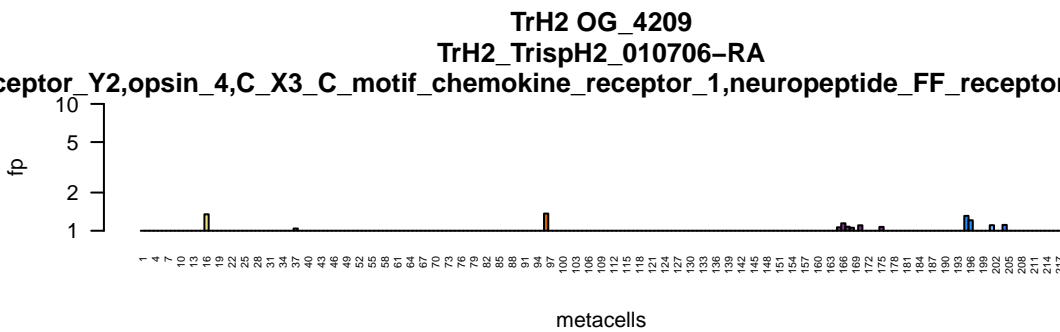
metacells



ceptor_Y2,opsin_4,C_X3_C_motif_chemokine_receptor_1,neuropeptide_FF_receptor

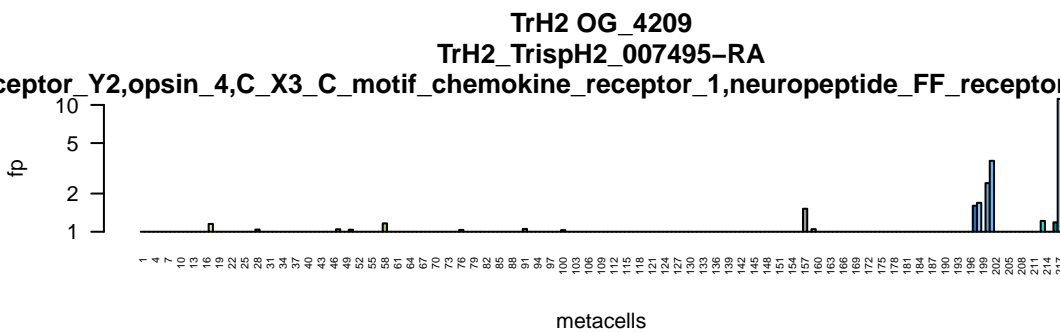
fp

metacells



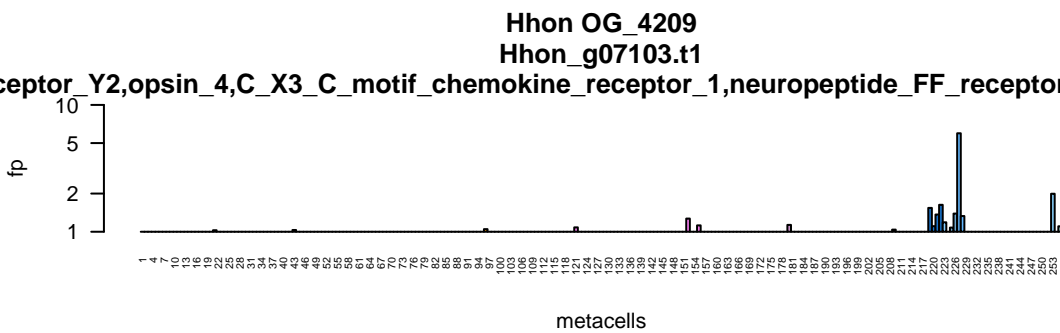
ceptor_Y2,opsin_4,C_X3_C_motif_chemokine_receptor_1,neuropeptide_FF_receptor

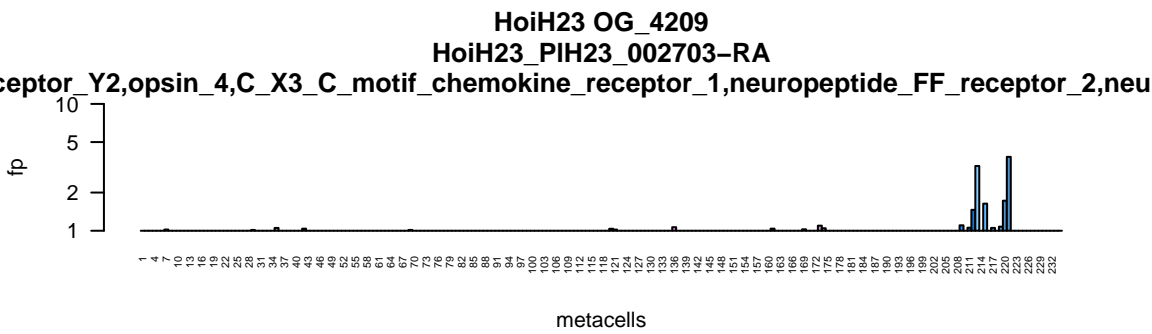
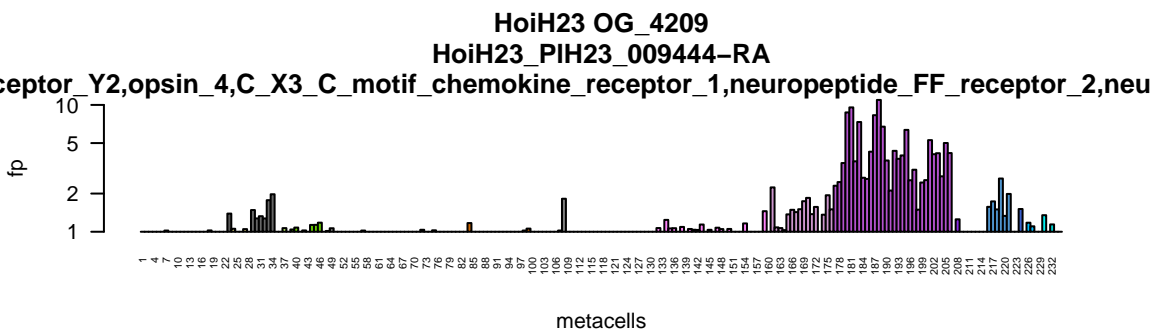
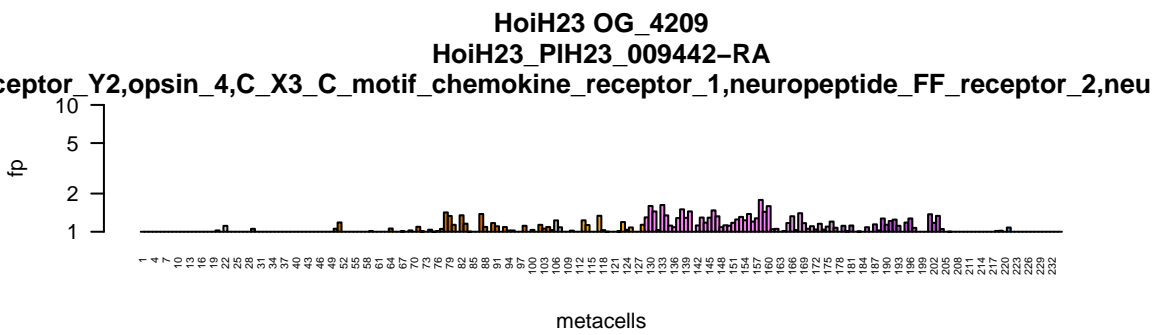
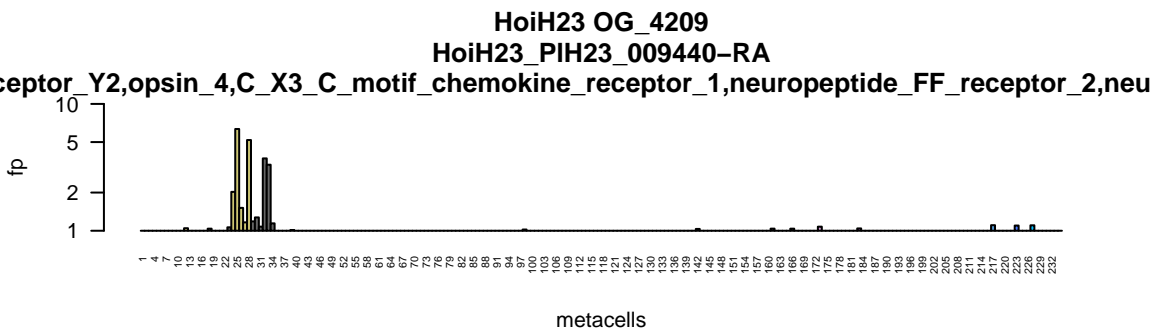
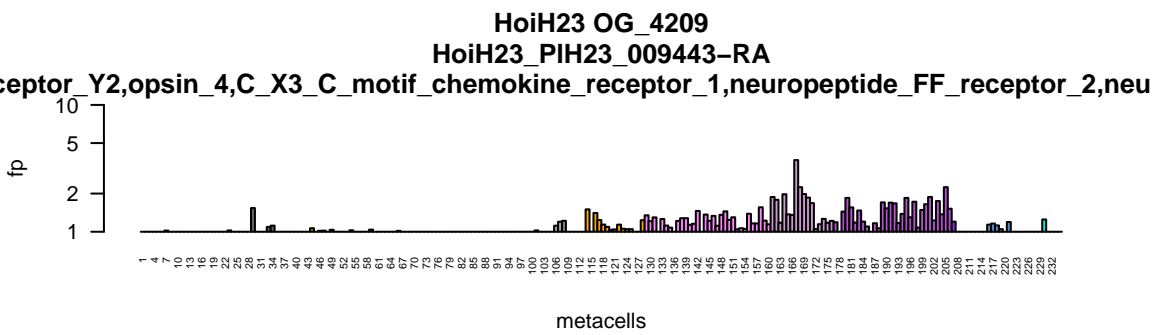
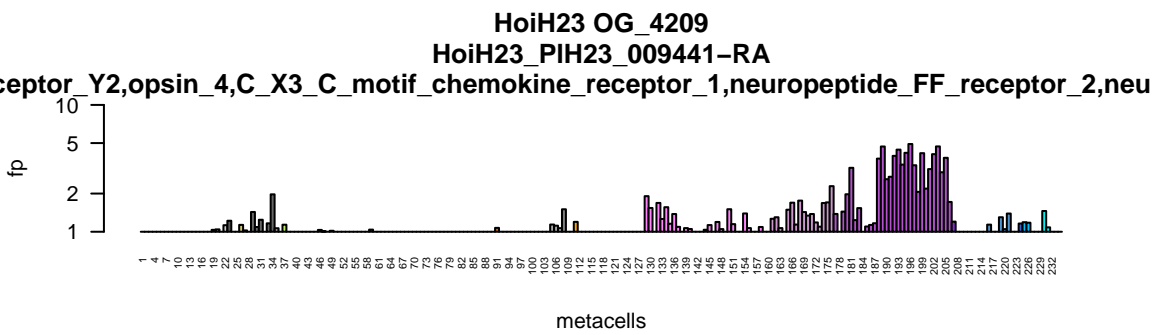
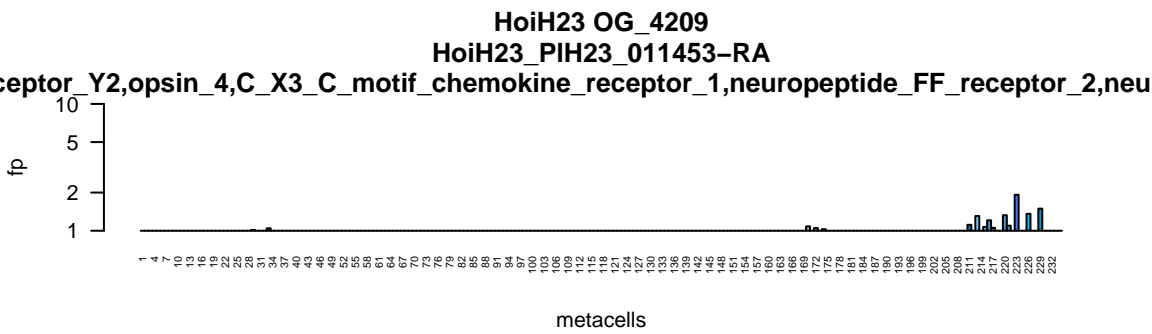
metacells	fp
1	1
4	1
10	1
13	1
16	1
19	1
21	1
25	1
28	1
31	1
31	1
37	1
40	1
43	1
46	1
46	1
52	1
55	1
59	1
61	1
64	1
67	1
70	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
103	1
106	1
109	1
111	1
115	1
118	1
121	1
124	1
124	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
169	1
172	1
175	1
181	1
184	1
187	1
193	1
196	1
199	1.2
202	1.5
208	1
211	1
214	2.5



ceptor_Y2,opsin_4,C_X3_C_motif_chemokine_receptor_1,neuropeptide_FF_receptor

metacells	fp
1	1
7	1
10	1
13	1
19	1
25	1
26	1
28	1
34	1
37	1
43	1
46	1
55	1
61	1
64	1
70	1
73	1
79	1
82	1
88	1
91	1
97	1
103	1
106	1
112	1
115	1
121	1
124	1
130	1
133	1
139	1
148	1
151	1
157	1
160	1
166	1
169	1
175	1
179	1
184	1
187	1
199	1
202	1
208	1
211	1
217	1
220	1
223	1
226	1
229	1
235	1
238	1
244	1
247	1
253	1





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

metacell	fp
1	0
2	0
3	0
4	0
5	0
6	0
7	0
8	0
9	0
10	0
11	0
12	0
13	0
14	0
15	0
16	0
17	0
18	0
19	0
20	0
21	0
22	0
23	0
24	0
25	0
26	0
27	0
28	0
29	0
30	0
31	0
32	0
33	0
34	0
35	0
36	0
37	0
38	0
39	0
40	0
41	0
42	0
43	0
44	0
45	0
46	0
47	0
48	0
49	0
50	0
51	0
52	0
53	0
54	0
55	0
56	0
57	0
58	0
59	0
60	0
61	0
62	0
63	0
64	0
65	0
66	0
67	0
68	0
69	0
70	0
71	0
72	0
73	0
74	0
75	0
76	0
77	0
78	0
79	0
80	0
81	0
82	0
83	0
84	0
85	0
86	0
87	0
88	0
89	0
90	0
91	0
92	0
93	0
94	0
95	0
96	0
97	0
98	0
99	0
100	0
101	0
102	0
103	0
104	0
105	0
106	0
107	0
108	0
109	0
110	0
111	0
112	0
113	0
114	0
115	0
116	0
117	0
118	0
119	0
120	0
121	0
122	0
123	0
124	0
125	0
126	0
127	0
128	0
129	0
130	0
131	0
132	0
133	0
134	0
135	0
136	0
137	0
138	0
139	0
140	0
141	0
142	0
143	0
144	0
145	0
146	0
147	0
148	0
149	0
150	0
151	0
152	0
153	0
154	0
155	0
156	0
157	0
158	0
159	0
160	0
161	0
162	0
163	0
164	0
165	0
166	0
167	0
168	0
169	0
170	0
171	0
172	0
173	1
174	2
175	2
176	3
177	4
178	3
179	2
180	1
181	1
182	1
183	1
184	1
185	1
186	1
187	1
188	1
189	1
190	1
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 187 metacells. Most metacells have a false positive count of 1, with a few having counts of 2 or 4.

metacell	fp
1	1
2	1
3	1
4	1
5	1
6	1
7	1
8	1
9	1
10	1
11	1
12	1
13	1
14	1
15	1
16	1
17	1
18	1
19	1
20	1
21	1
22	1
23	1
24	1
25	1
26	1
27	1
28	1
29	1
30	1
31	1
32	1
33	1
34	1
35	1
36	1
37	1
38	1
39	1
40	1
41	1
42	1
43	1
44	1
45	1
46	1
47	1
48	1
49	1
50	1
51	1
52	1
53	1
54	1
55	1
56	1
57	1
58	1
59	1
60	1
61	1
62	1
63	1
64	1
65	1
66	1
67	1
68	1
69	1
70	1
71	1
72	1
73	1
74	1
75	1
76	1
77	1
78	1
79	1
80	1
81	1
82	1
83	1
84	1
85	1
86	1
87	1
88	1
89	1
90	1
91	1
92	1
93	1
94	1
95	1
96	1
97	1
98	1
99	1
100	1
101	1
102	1
103	1
104	1
105	1
106	1
107	1
108	1
109	1
110	1
111	1
112	1
113	1
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119	1
120	1
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122	1
123	1
124	1
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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
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157	1
158	1
159	1
160	1
161	1
162	1
163	1
164	1
165	1
166	1
167	1
168	1
169	1
170	1
171	1
172	1
173	1
174	1
175	1
176	1
177	1
178	1
179	1
180	1
181	1
182	1
183	1
184	1
185	1
186	1
187	1

A bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and the y-axis is labeled 'fp'. The y-axis has a non-linear scale with values 1, 2, 5, and 10. The x-axis lists 187 metacells, numbered 1 to 187. The bars represent the frequency of each metacell. Most metacells have a frequency of 1, with a few having higher frequencies: metacell 163 has a frequency of 1, 167 has 2, 171 has 3, 172 has 4, 173 has 5, 174 has 6, 175 has 7, 176 has 8, 177 has 9, 178 has 10, 181 has 1, 185 has 2, and 187 has 1.

metacell	fp
1	1
2	1
3	1
4	1
5	1
6	1
7	1
8	1
9	1
10	1
11	1
12	1
13	1
14	1
15	1
16	1
17	1
18	1
19	1
20	1
21	1
22	1
23	1
24	1
25	1
26	1
27	1
28	1
29	1
30	1
31	1
32	1
33	1
34	1
35	1
36	1
37	1
38	1
39	1
40	1
41	1
42	1
43	1
44	1
45	1
46	1
47	1
48	1
49	1
50	1
51	1
52	1
53	1
54	1
55	1
56	1
57	1
58	1
59	1
60	1
61	1
62	1
63	1
64	1
65	1
66	1
67	1
68	1
69	1
70	1
71	1
72	1
73	1
74	1
75	1
76	1
77	1
78	1
79	1
80	1
81	1
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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
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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	2
168	1
169	1
170	1
171	3
172	4
173	5
174	6
175	7
176	8
177	9
178	10
179	1
180	1
181	1
182	1
183	1
184	1
185	2
186	1
187	1

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists 48 metacells. Most metacells have a false positive count of 1, with a few having counts of 2 or 3.

metacells	fp
1	1
4	1
10	1
13	1
16	1
19	1
22	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	2
199	1
202	1
205	2
208	3
211	2
214	1

Bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and the y-axis is labeled 'fp'. The chart shows a very high frequency for metacell 1 (fp=10) and a few other metacells with low frequencies (fp=1).

metacells	fp
1	10
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

metacell	fp
1	1
4	0
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	0
34	0
37	0
40	0
43	0
46	0
49	0
52	0
55	0
58	0
61	0
64	0
67	0
70	0
73	0
76	1
79	1
82	0
85	0
88	0
91	1
94	0
97	0
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	0
130	0
133	0
136	0
139	0
142	0
145	0
148	1
151	0
154	1
157	1
160	1
163	0
166	0
169	1
172	1
175	0
178	0
181	0
184	0
187	0
190	0
193	0
196	0
199	2
202	4
205	2
208	5
211	1
214	1

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

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

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

metacell	fp
1	0
4	1
10	0
13	0
19	0
22	0
31	0
34	1
40	0
43	0
46	0
52	0
55	0
58	0
64	0
67	0
73	0
76	1
82	0
85	0
88	0
94	1
97	0
103	0
106	1
109	0
115	0
118	0
124	0
127	1
130	0
136	1
139	0
145	1
151	0
157	0
160	0
166	0
169	0
172	1
178	0
181	0
187	0
190	0
193	0
199	0
202	0
208	0
211	1
214	0
217	3
220	2
221	1
222	6
223	10
229	8
232	1

metacell	fp
1	0
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	1
34	1
37	0
40	0
43	0
46	0
49	0
52	1
55	0
58	0
61	0
64	0
67	0
70	0
73	0
76	0
79	0
82	0
85	0
88	0
91	0
94	0
97	0
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	0
130	0
133	0
136	0
139	0
142	0
145	0
148	0
151	1
154	0
157	0
160	0
163	0
166	1
169	1
172	0
175	0
178	0
181	0
184	0
187	0
190	0
193	0
196	0
199	0
202	0
205	0
208	0
211	0
214	2
215	1
218	1
220	3
221	4
222	3
223	2
224	3
225	1
226	1
227	1
228	1
229	1
230	0
231	0
232	0

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. The bars are colored in a repeating pattern of blue, green, and red.

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

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

Bar chart showing the frequency of metacells across different frequency bins. The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows a distribution with a peak at metacell 202 (frequency 2) and a smaller peak at metacell 199 (frequency 1).

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	2
205	1
208	1
211	1
214	1
217	1

TrH2 OG_4305
TrH2_TrispH2_000683-RA

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

fp

metacells

TrH2 OG_4305
TrH2_TrispH2_011071-RA

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

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	2
205	1
208	1
211	1
214	1
217	2

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

Hhon OG_4305

Hhon_g11834.t1

fp

metacells

Hhon OG_4305
Hhon_g09074.t1

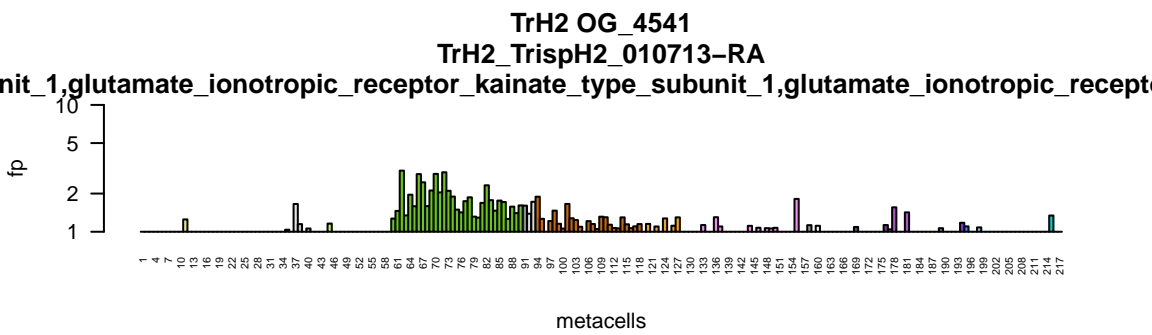
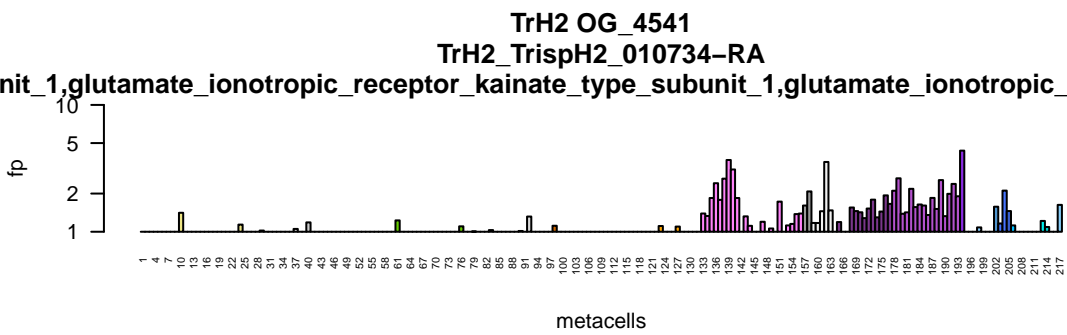
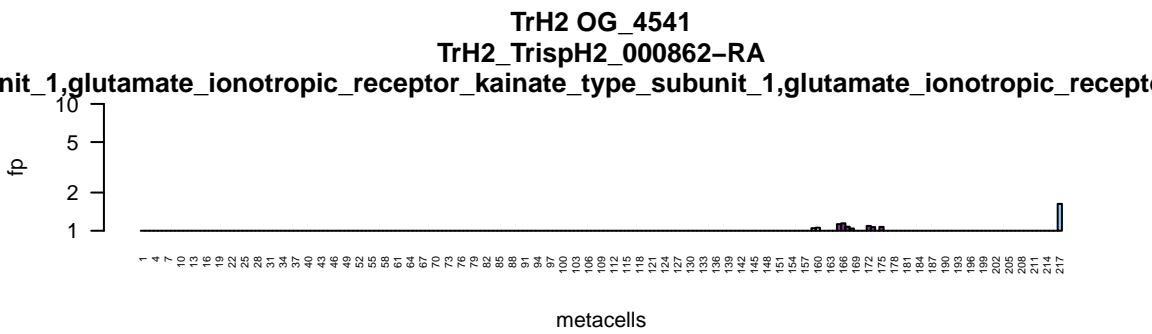
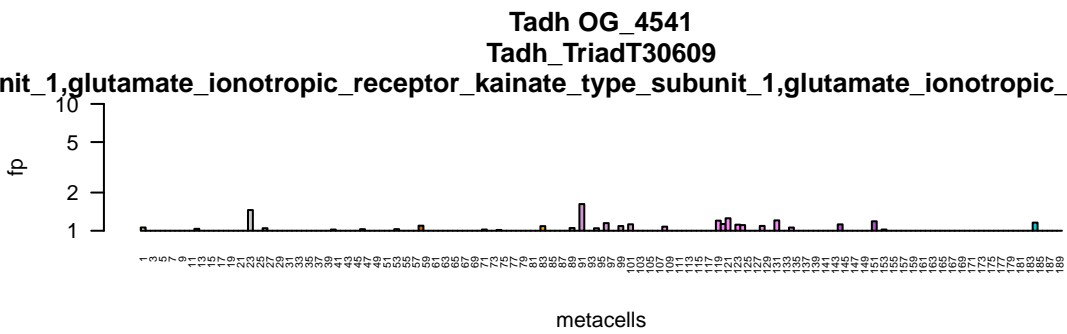
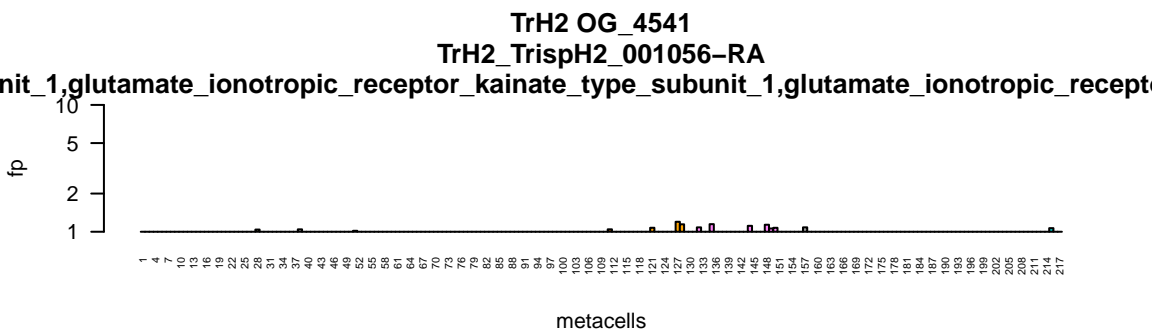
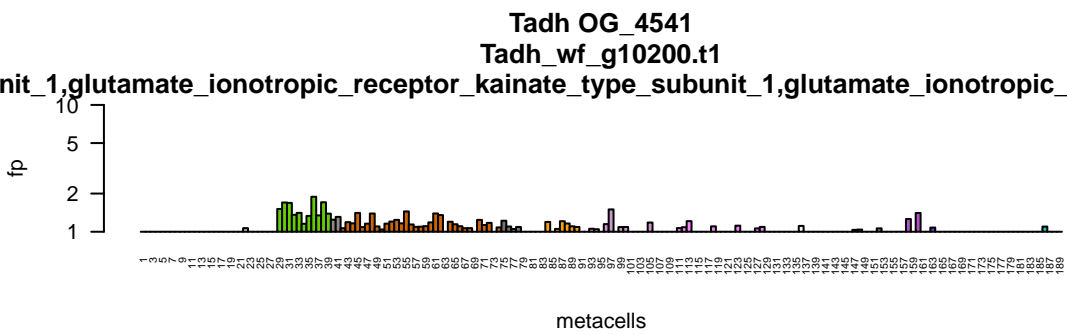
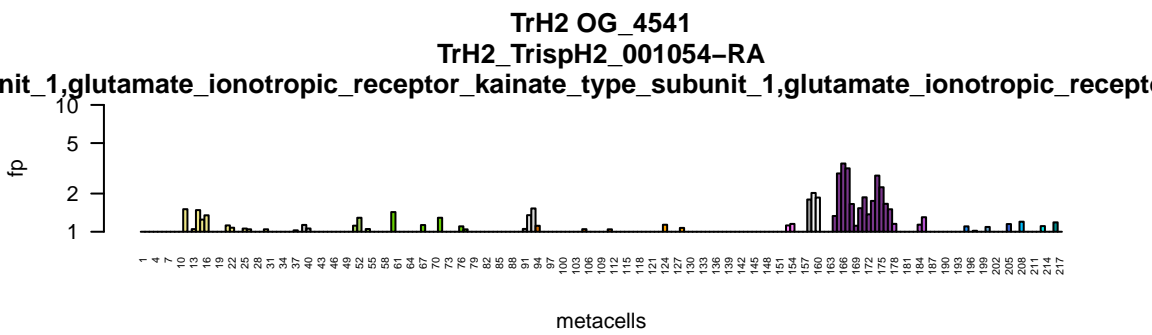
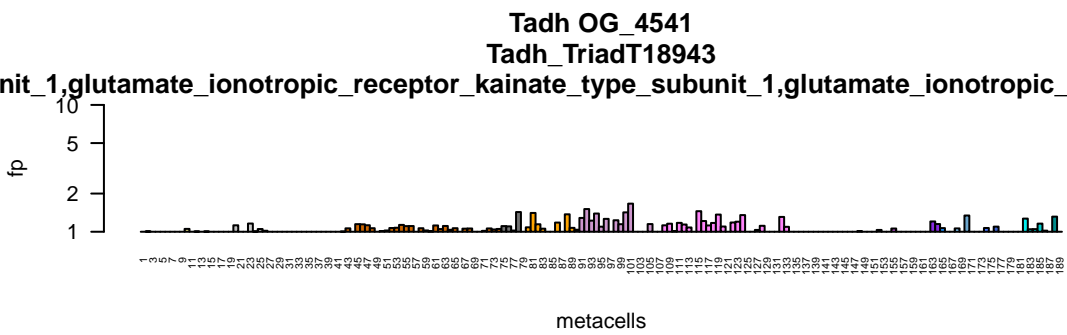
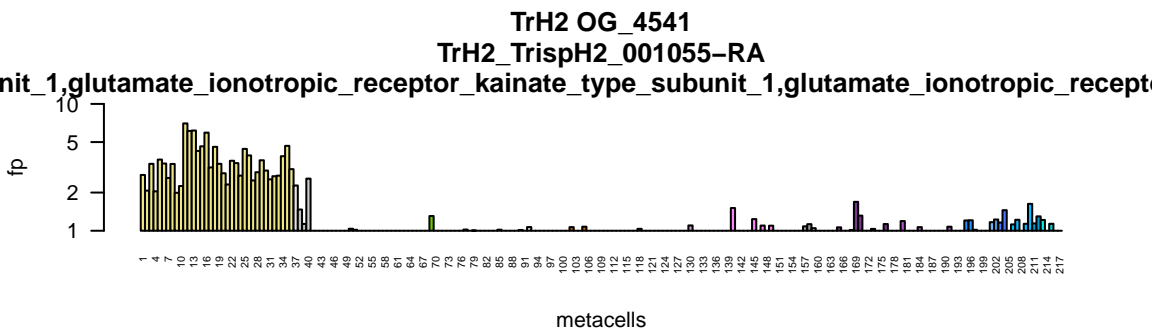
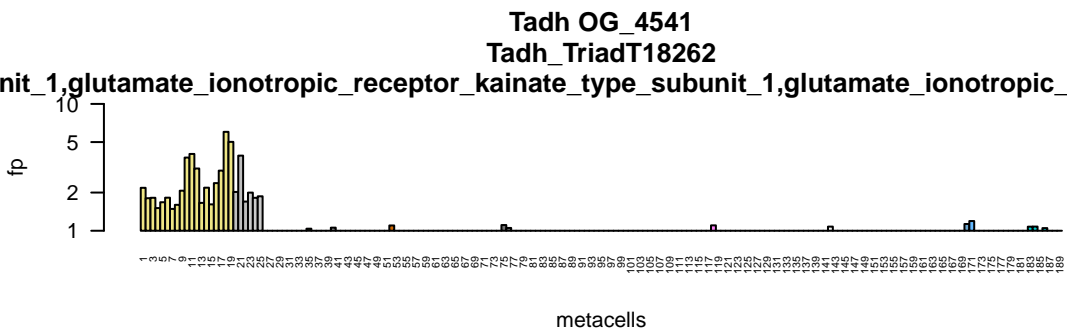
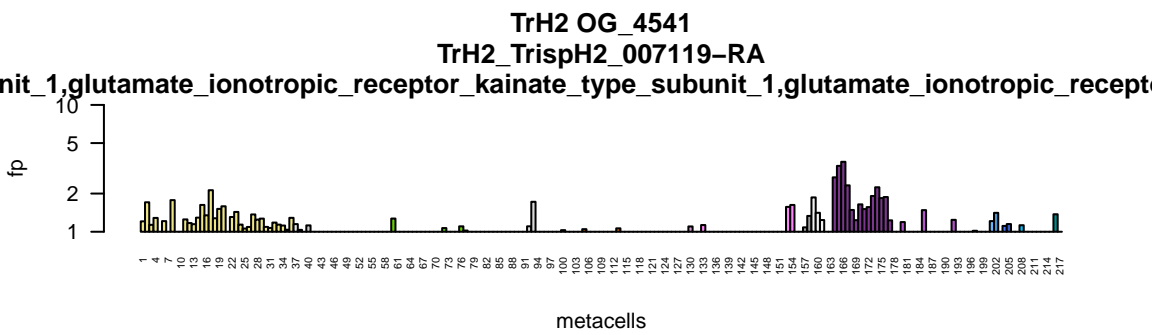
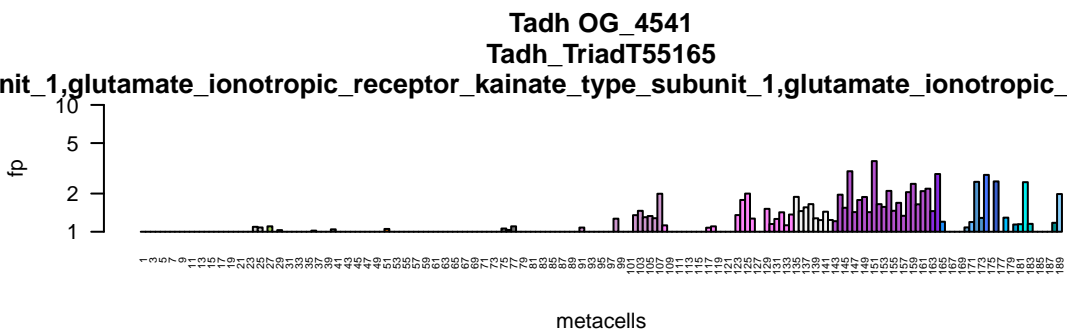
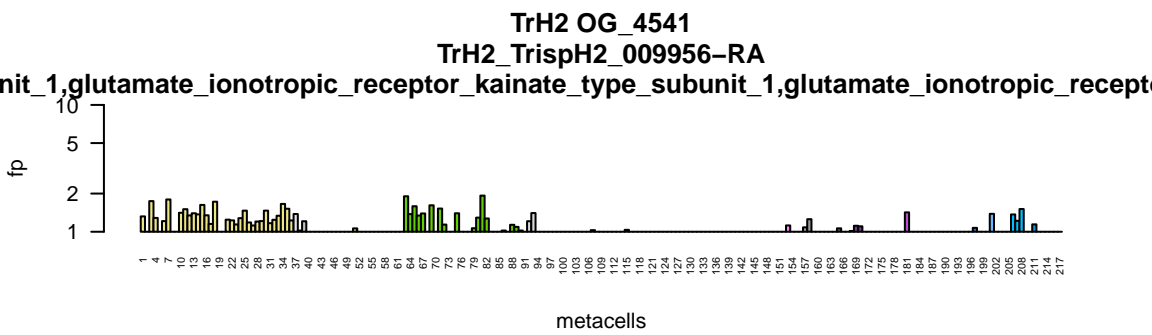
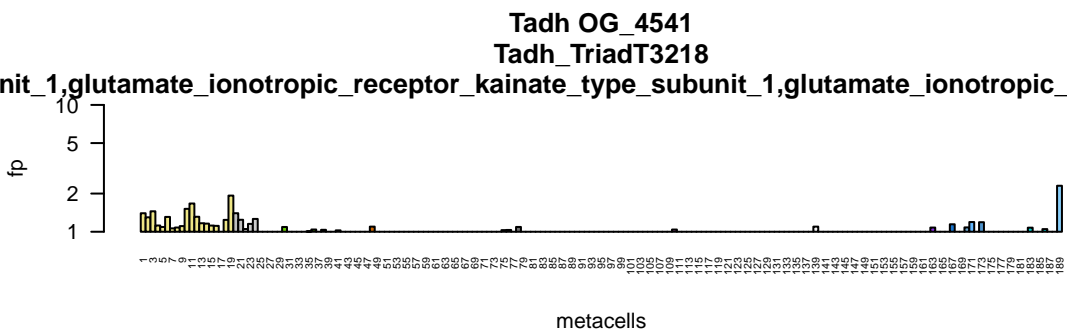
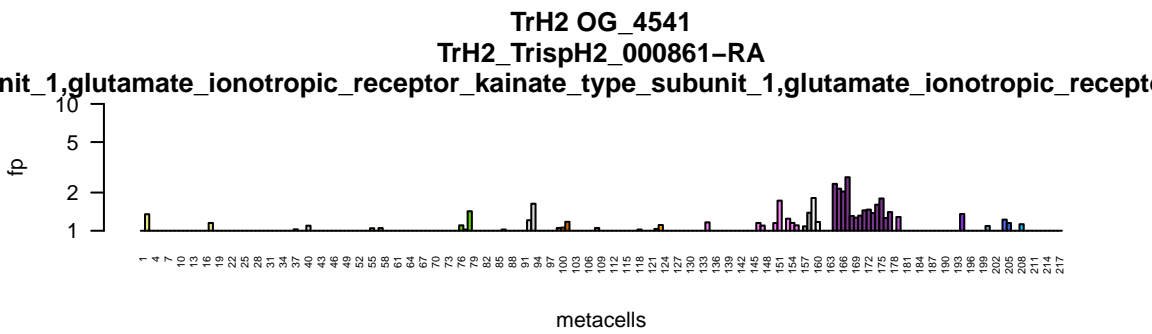
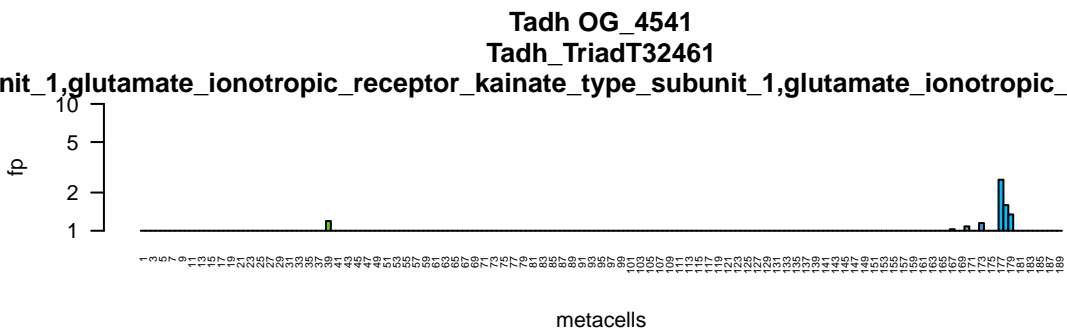
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

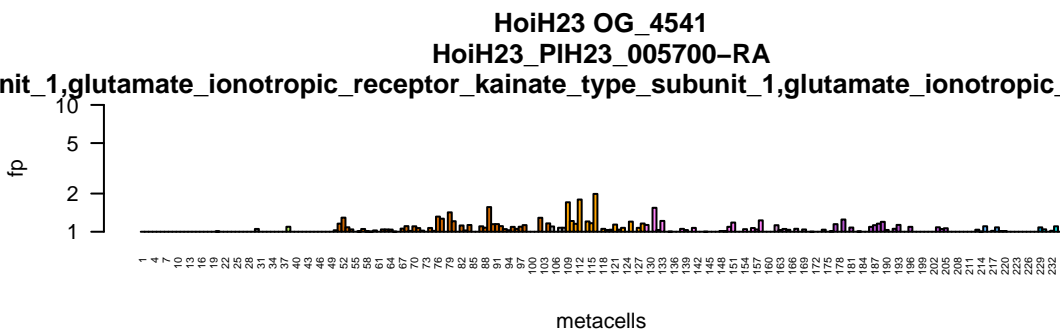
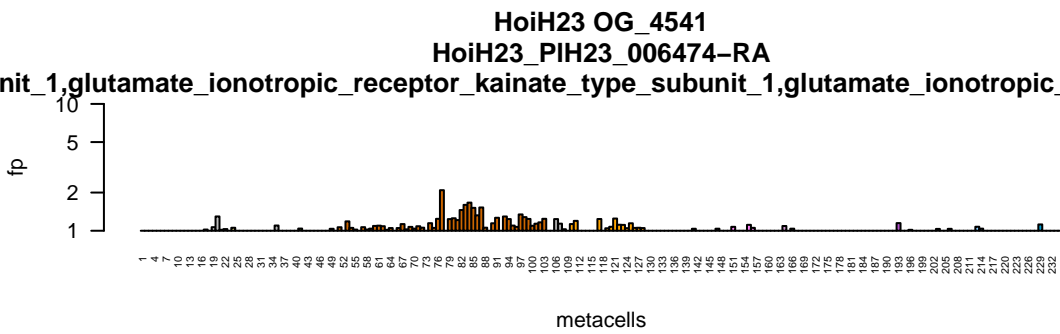
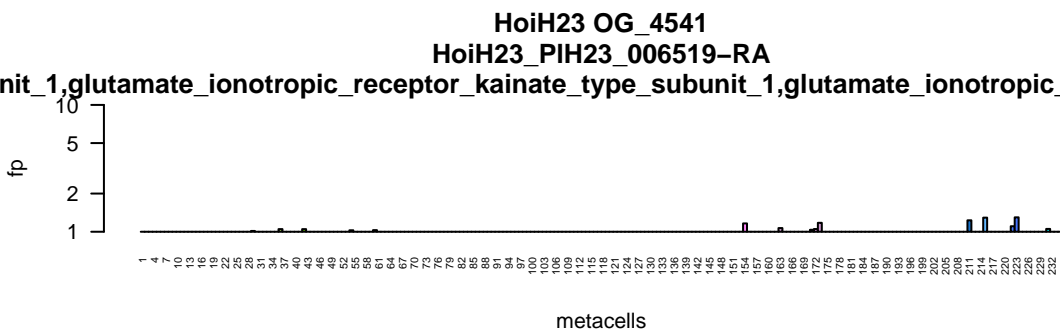
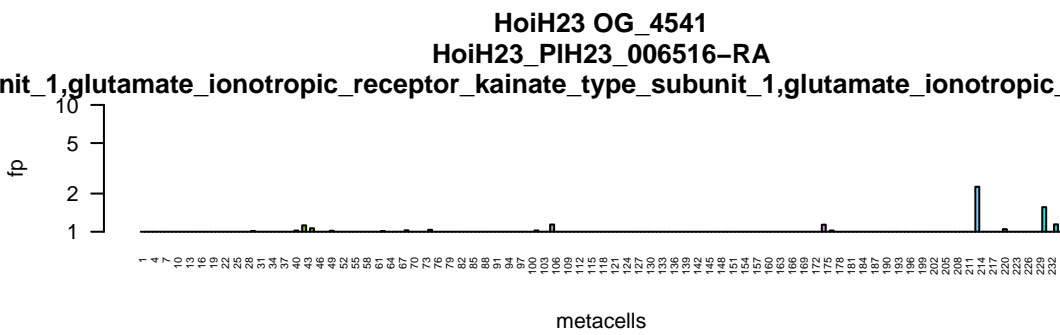
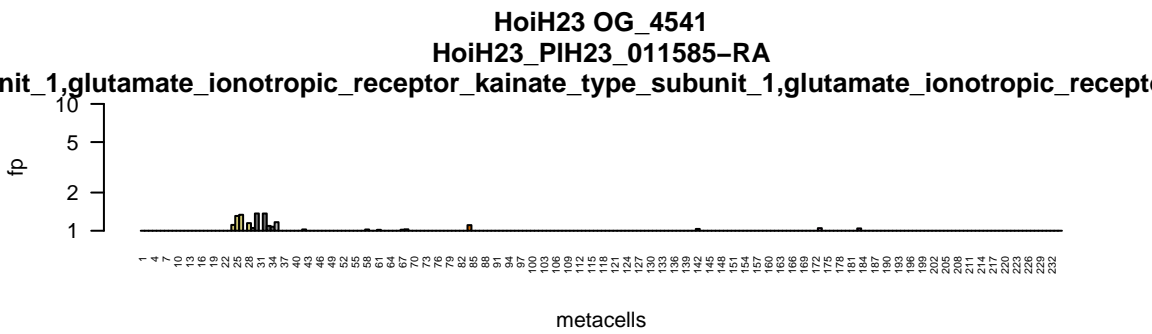
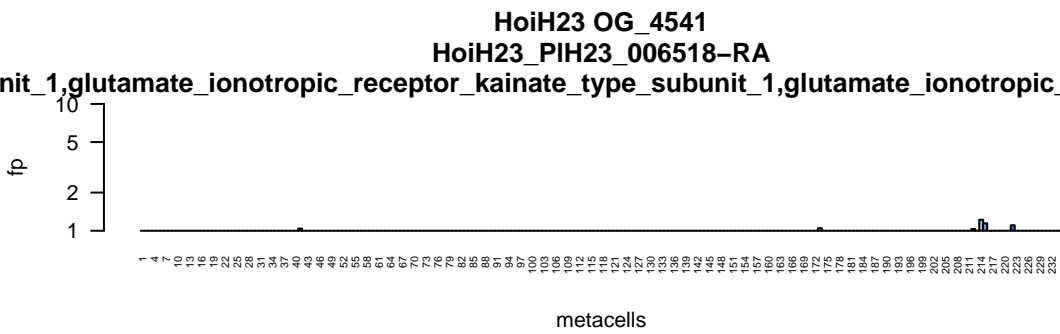
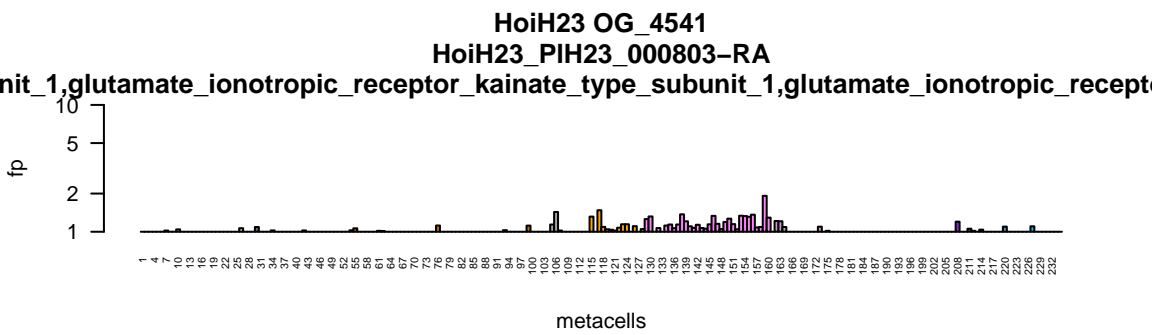
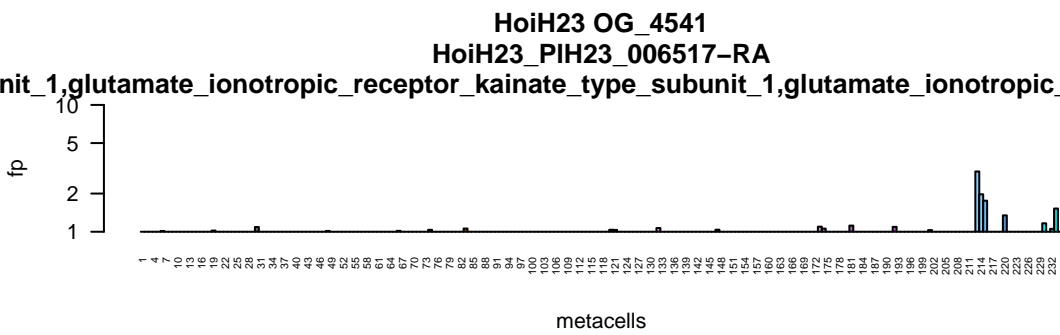
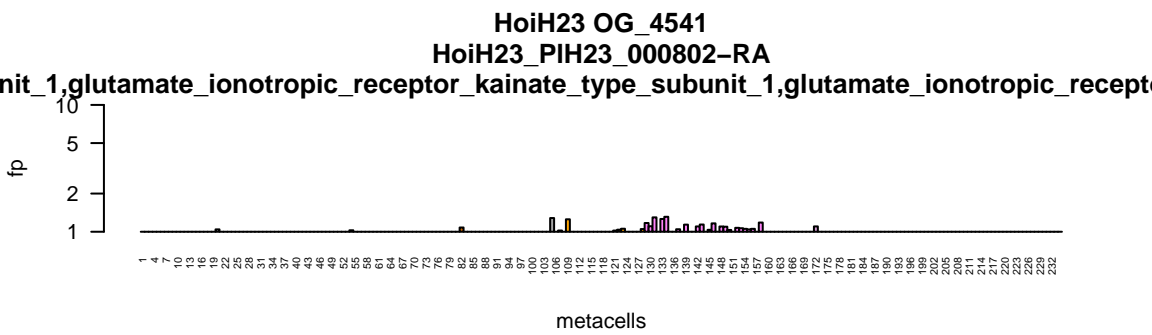
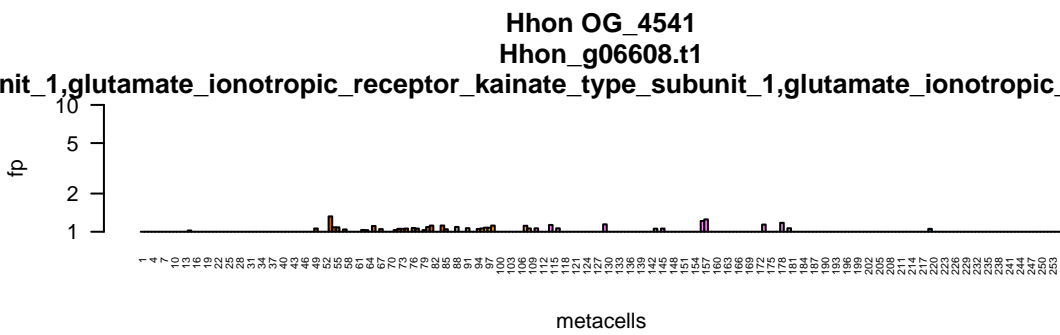
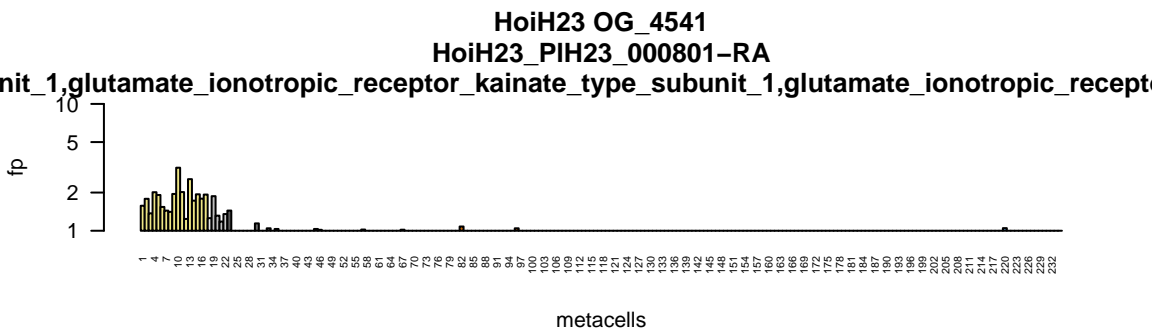
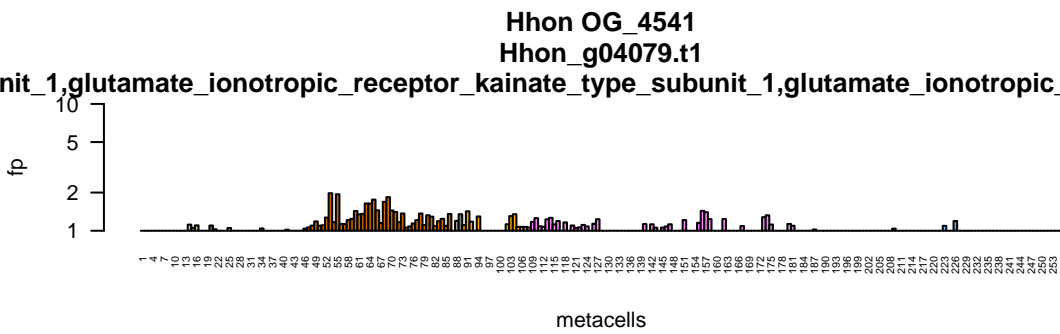
fp

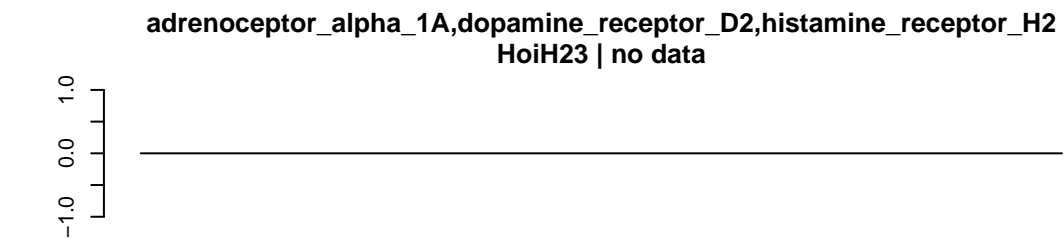
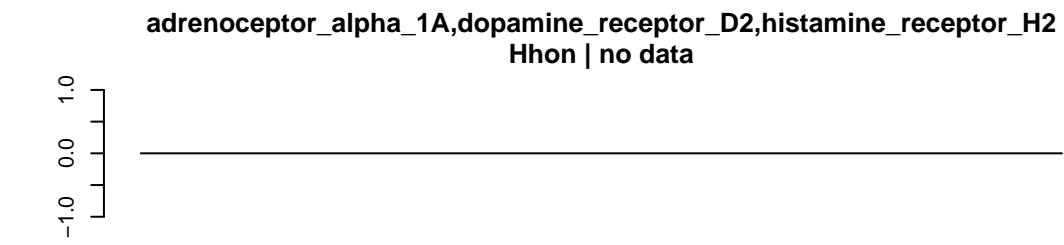
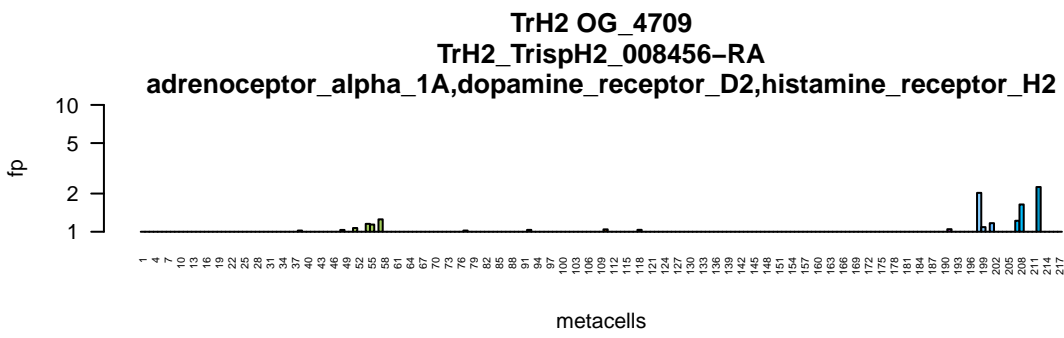
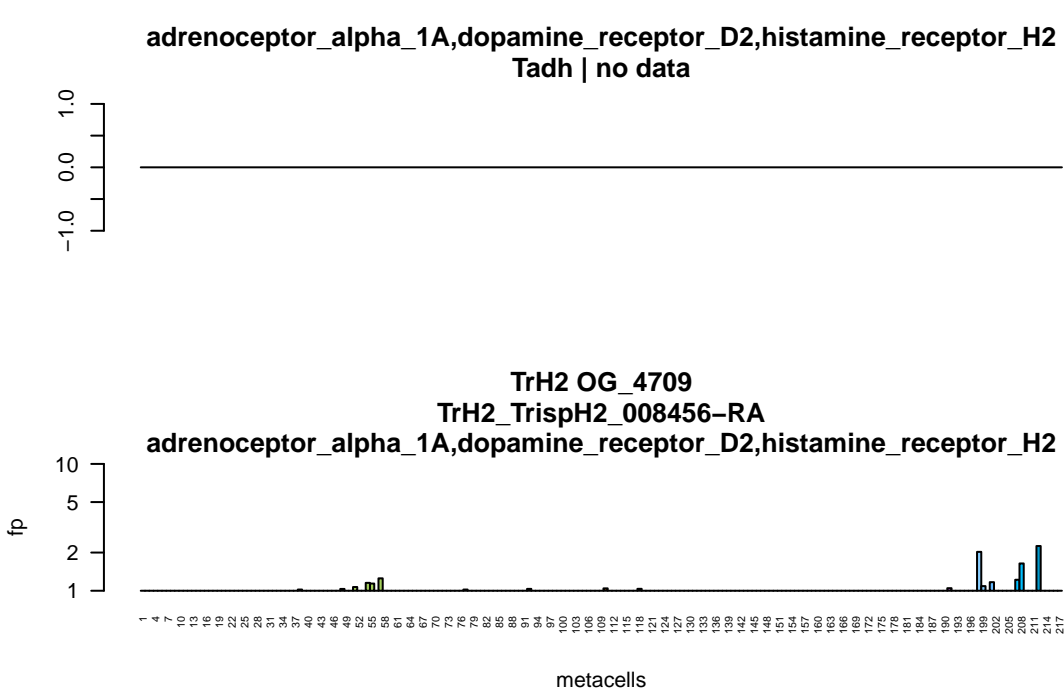
metacells

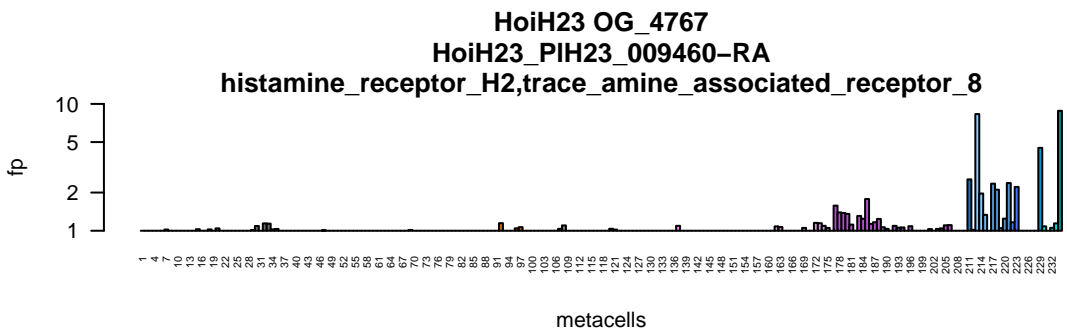
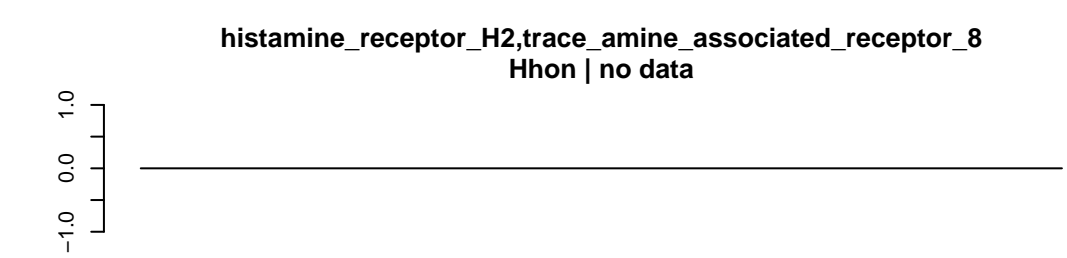
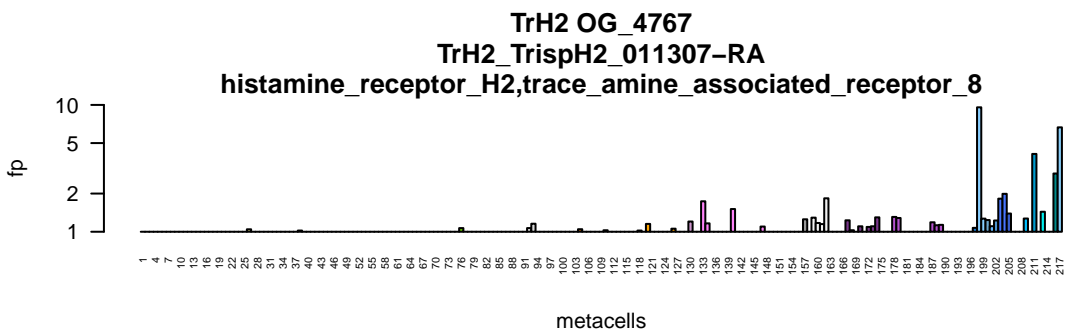
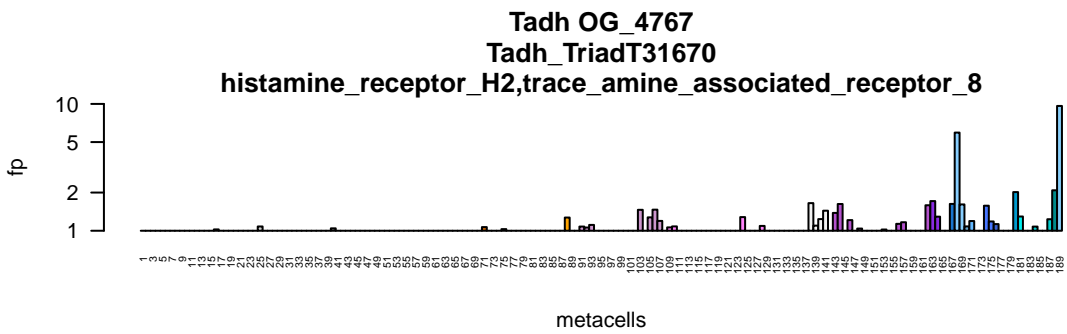
Bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and lists indices from 1 to 232. The y-axis is labeled 'fp' and has a logarithmic scale with values 1, 2, 5, and 10. Most metacells have a frequency of 1. Metacells 214, 223, and 232 show higher frequencies, with 214 being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

A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 232. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a notable peak at metacell 223 having a frequency of 4.





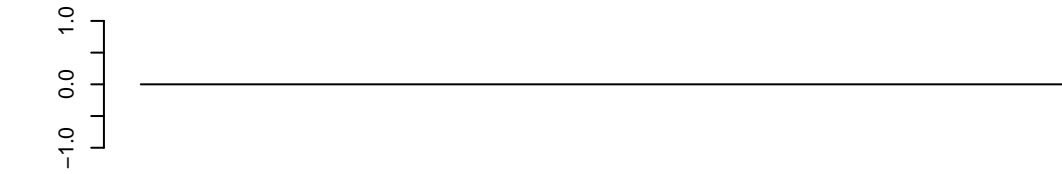




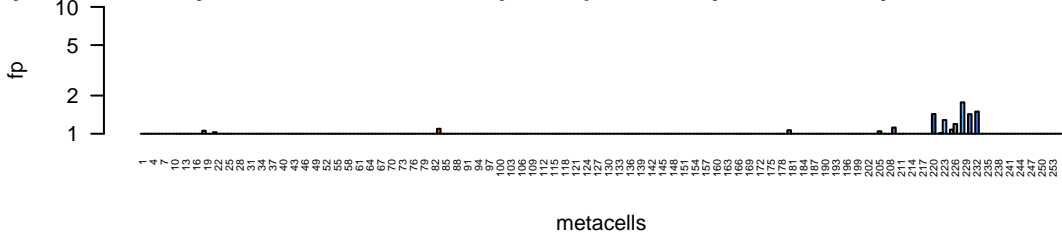
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Tadh | no data



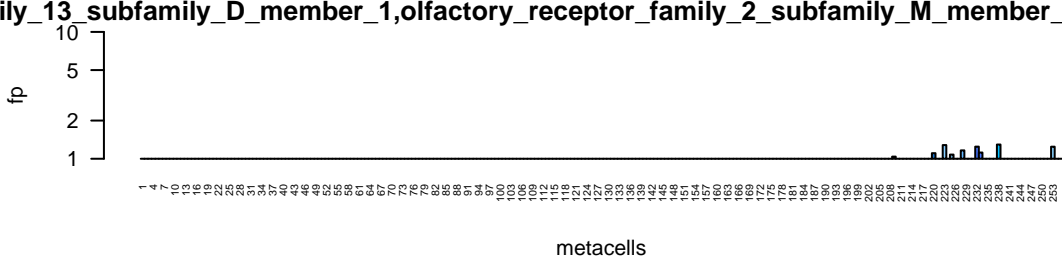
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi
TrH2 | no data



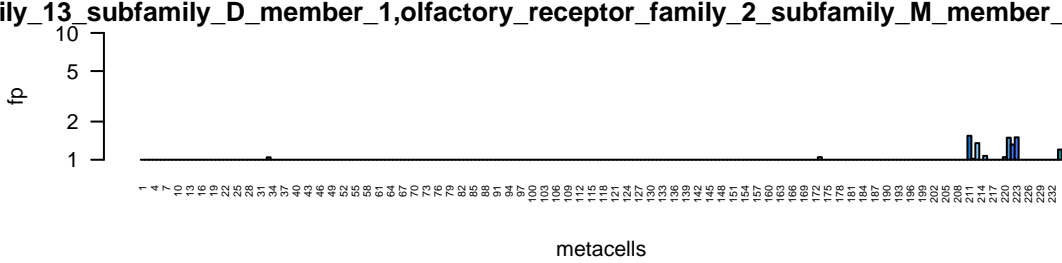
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Hhon_g08016.t1
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi



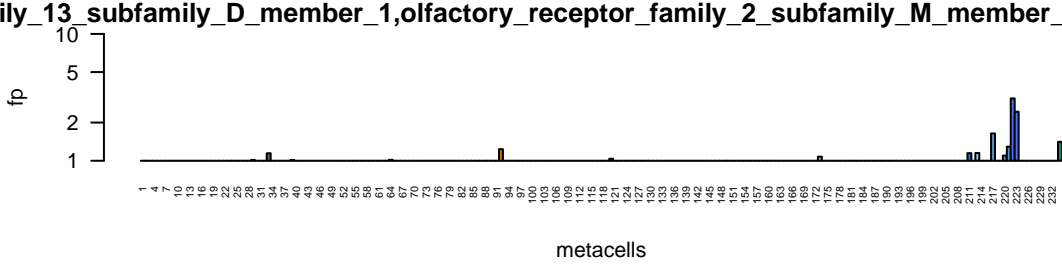
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ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi

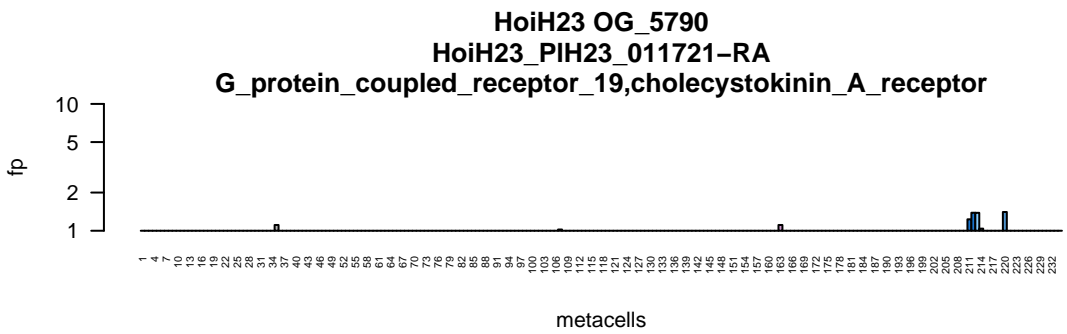
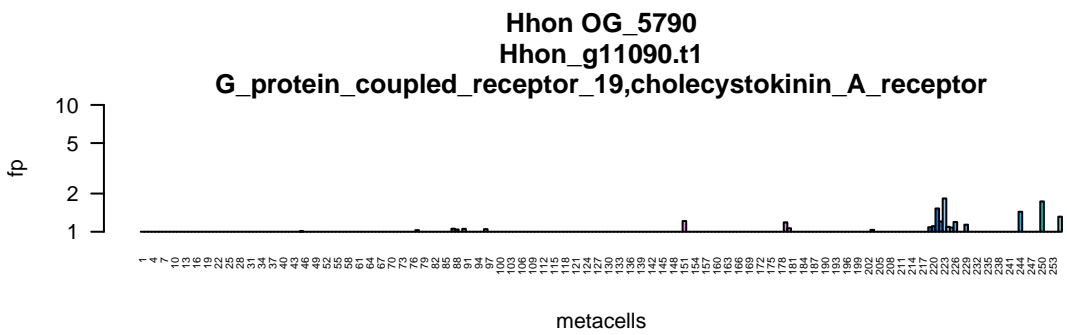
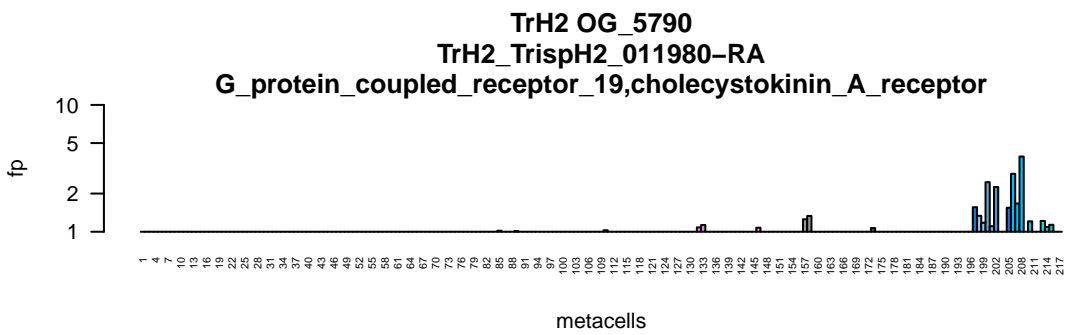
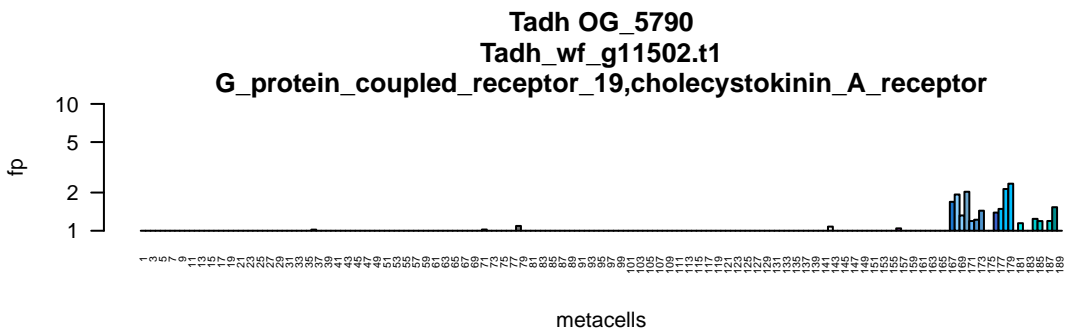


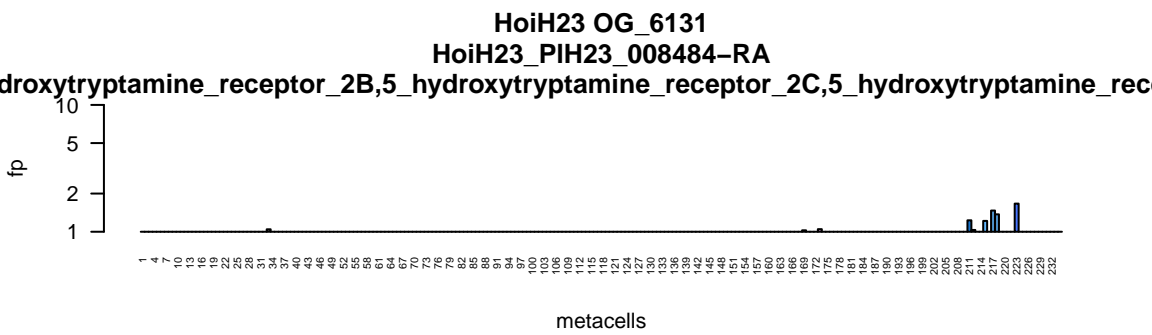
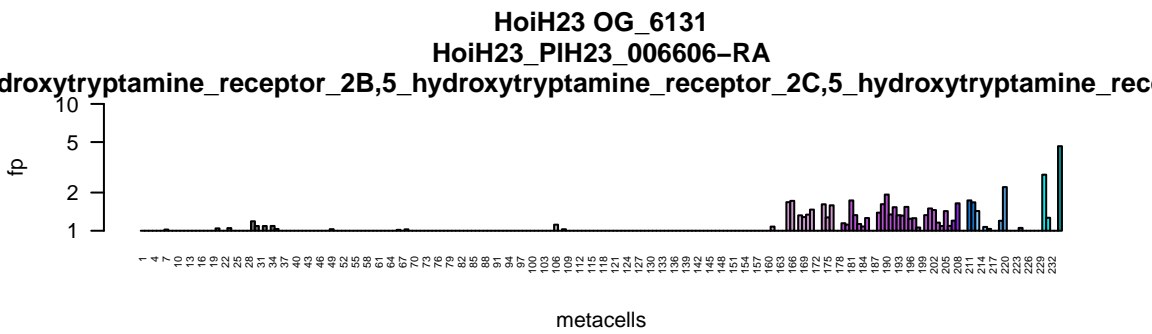
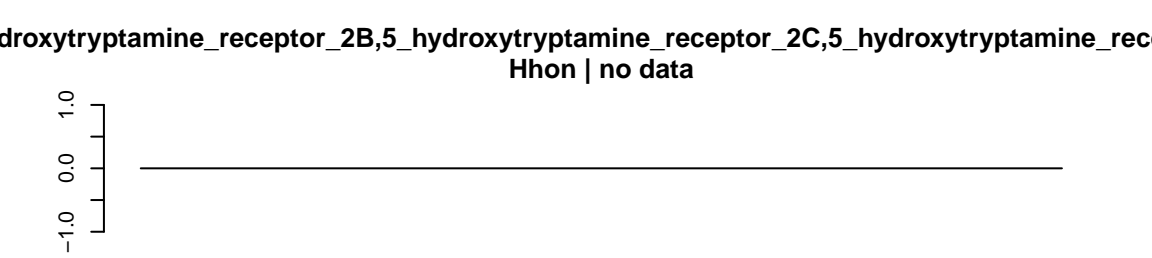
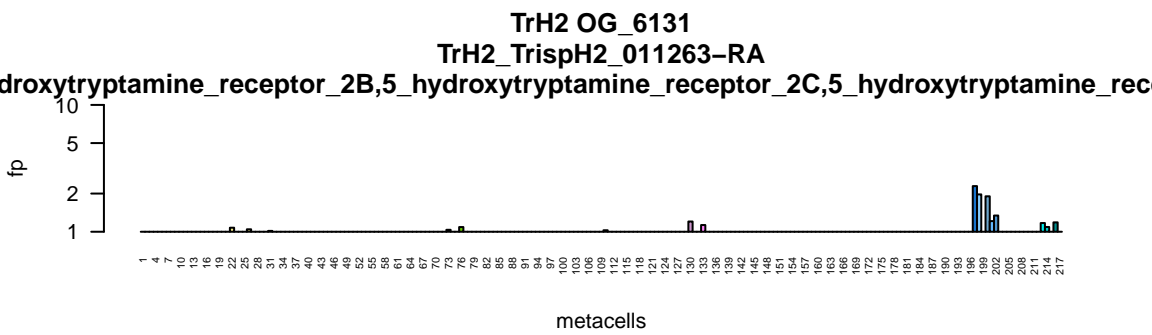
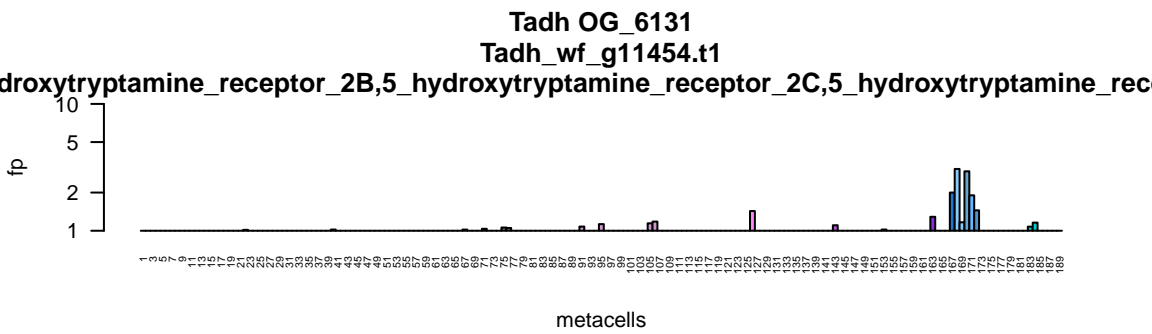
HoiH23 OG_4788
HoiH23_PIH23_005444-RA
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi

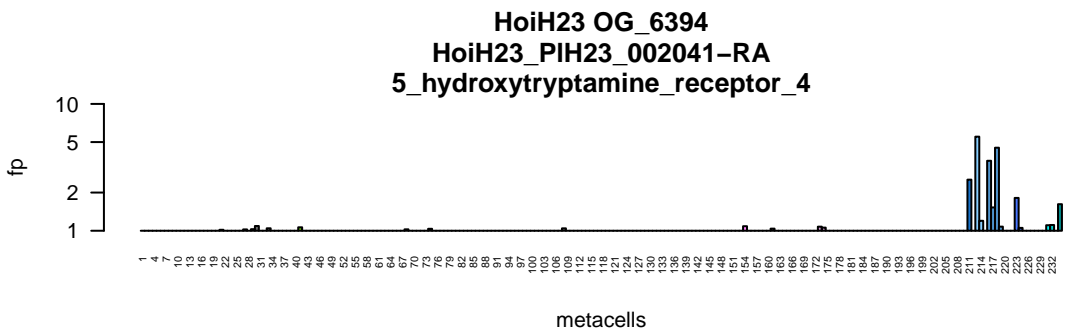
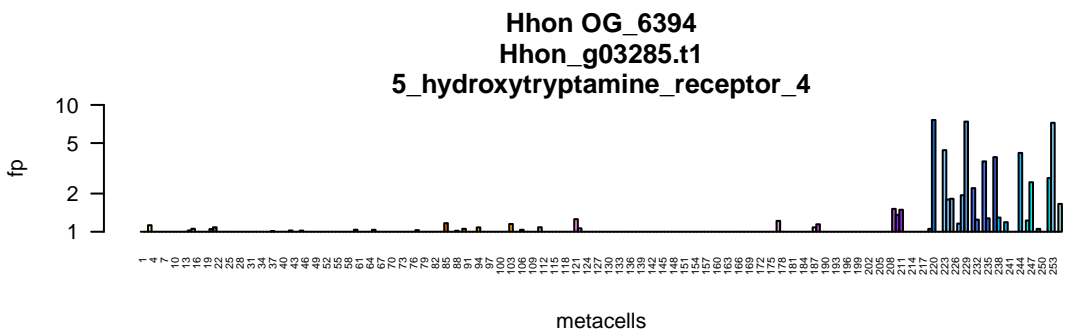
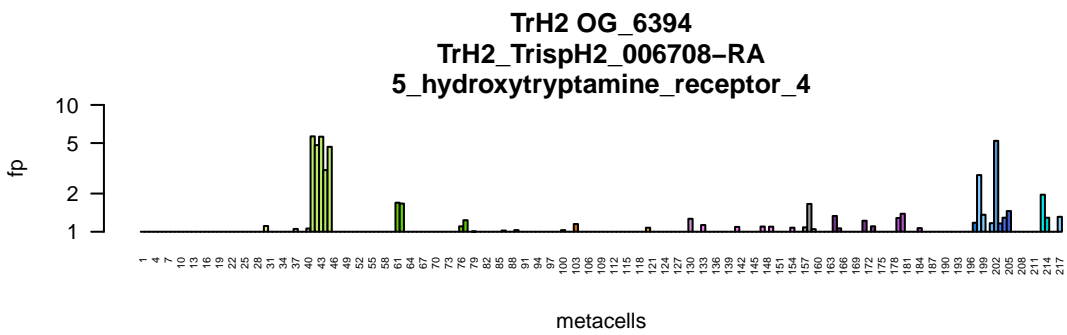
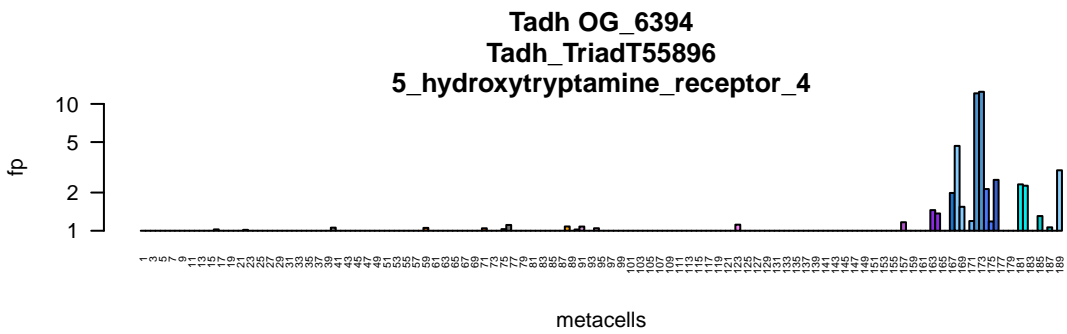


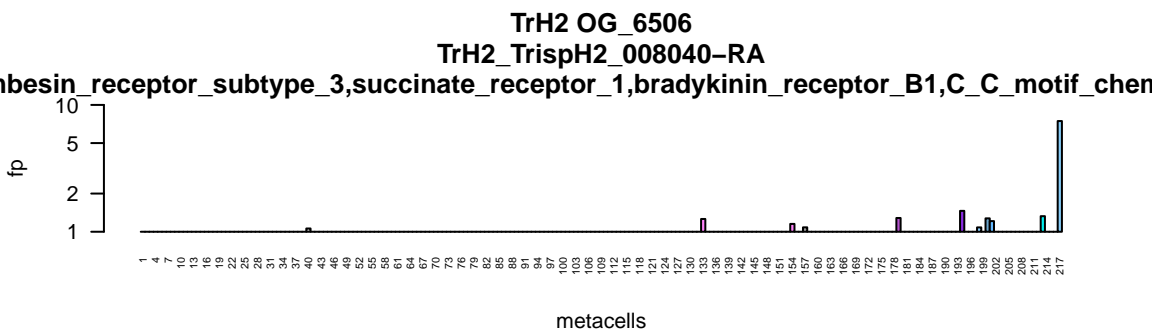
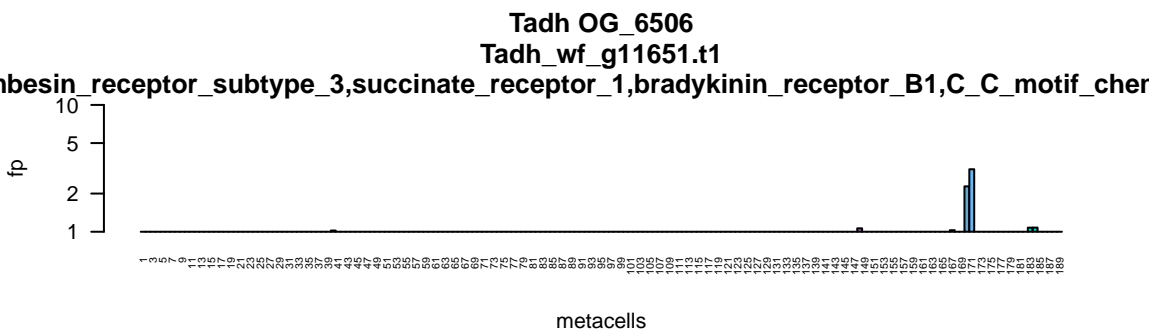
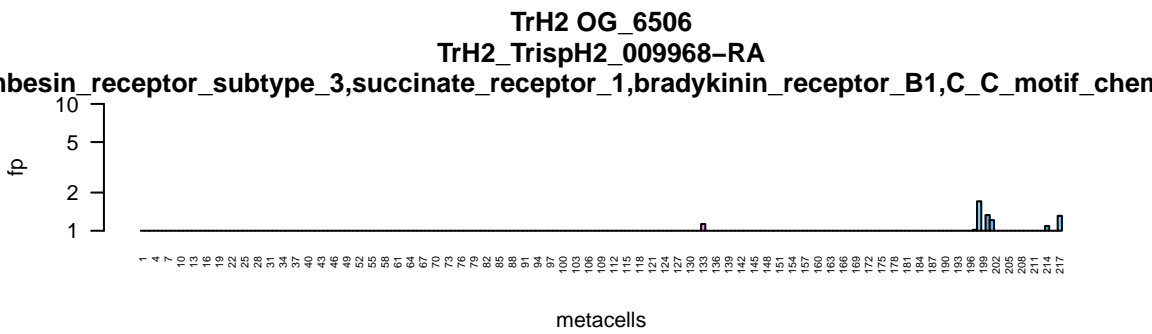
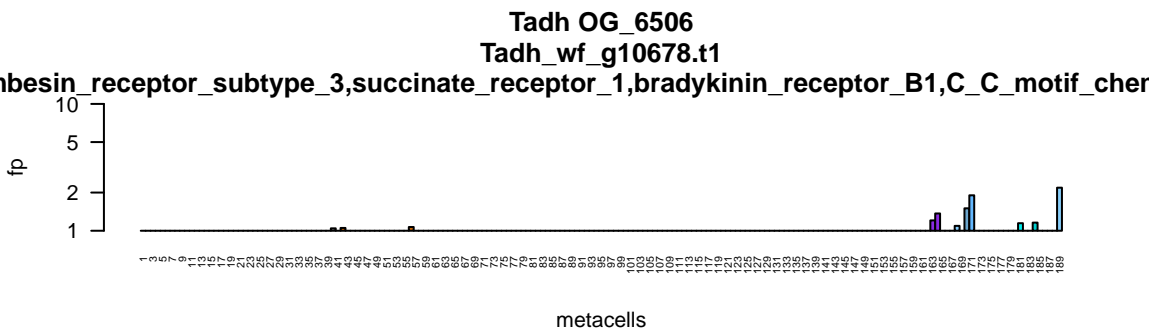
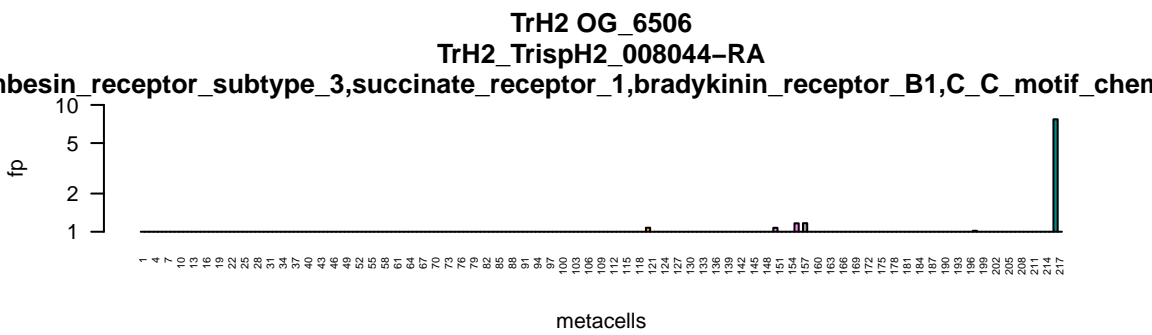
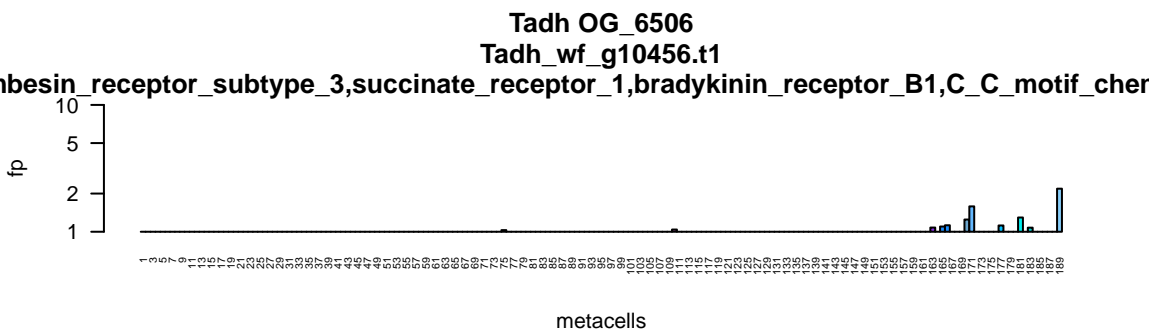
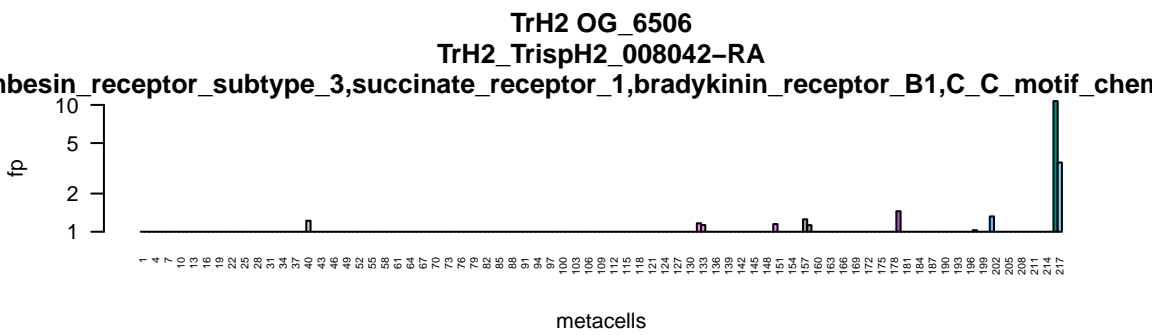
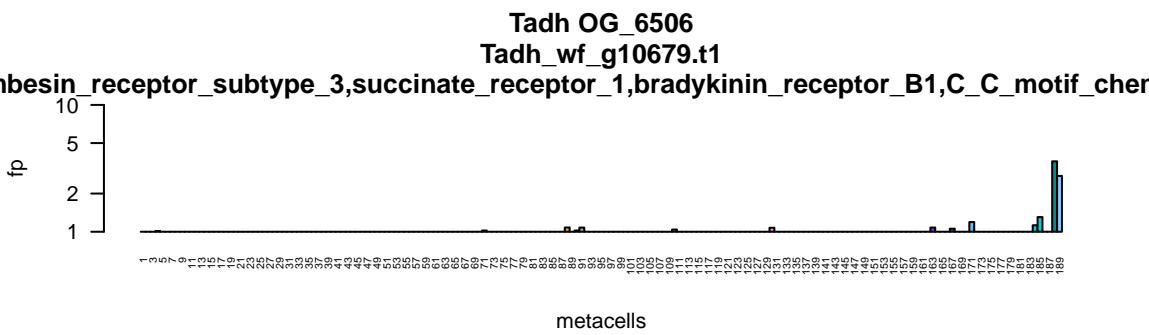
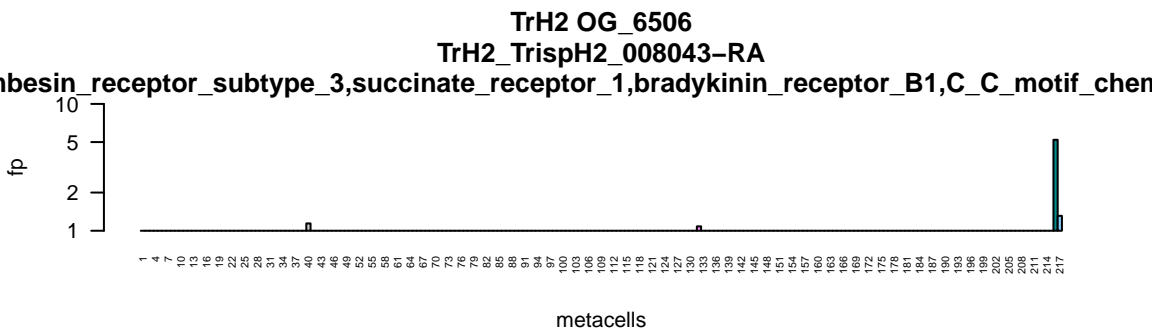
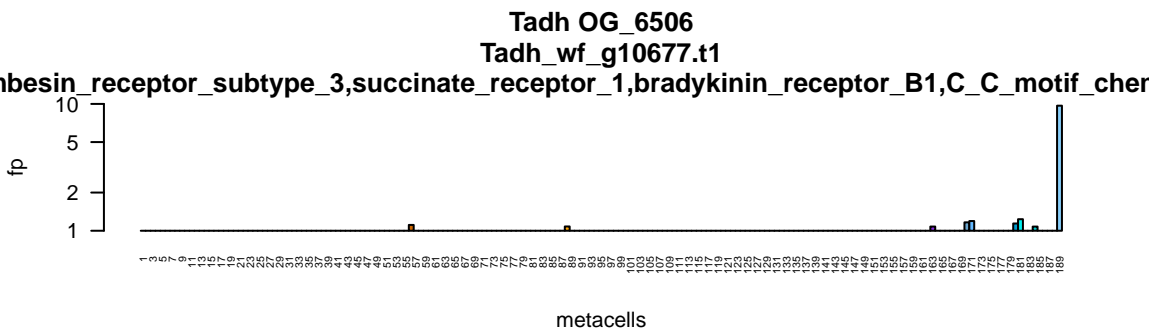
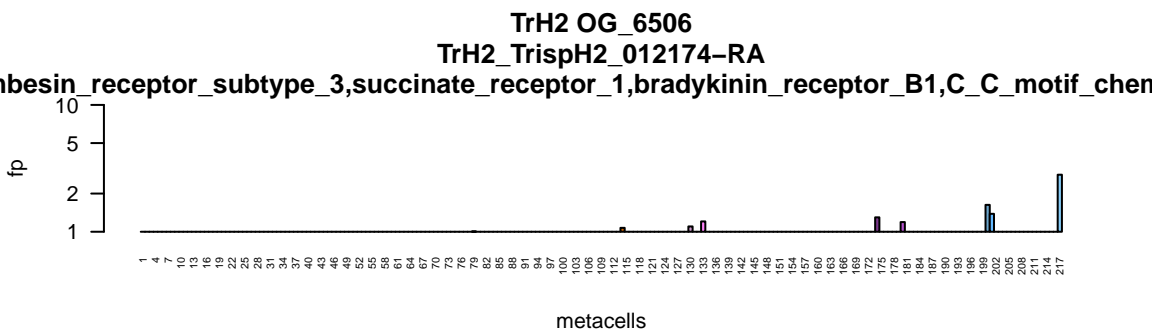
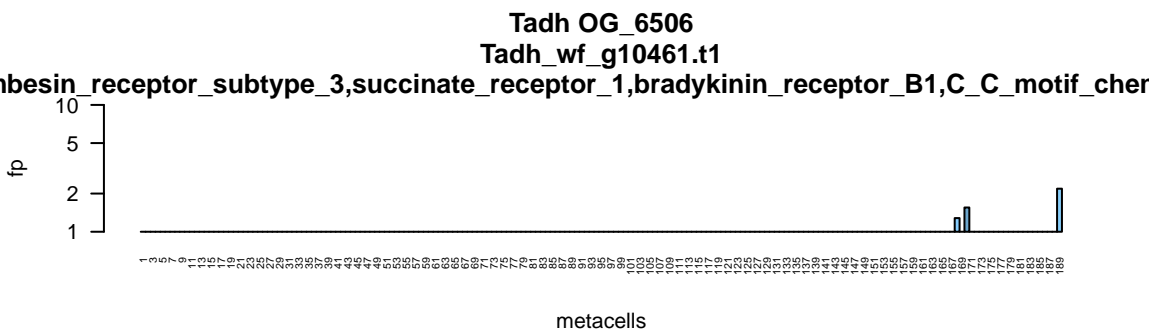
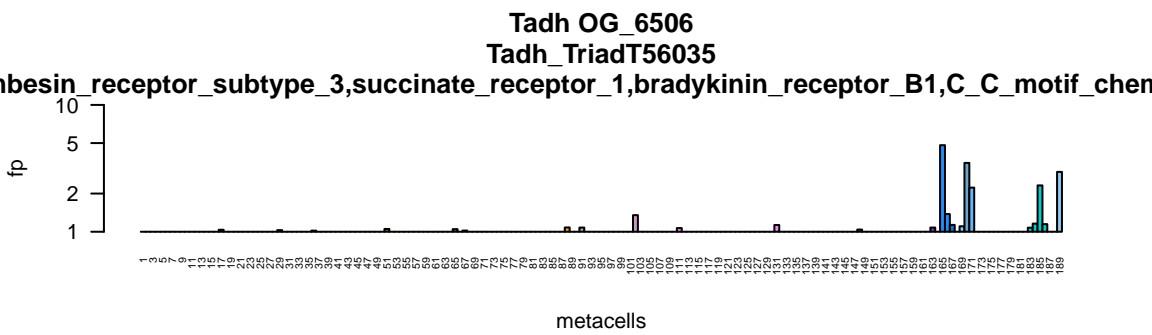
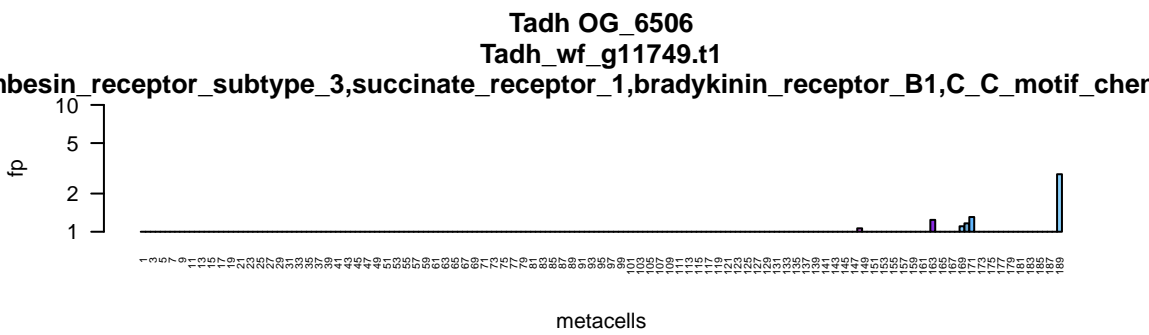
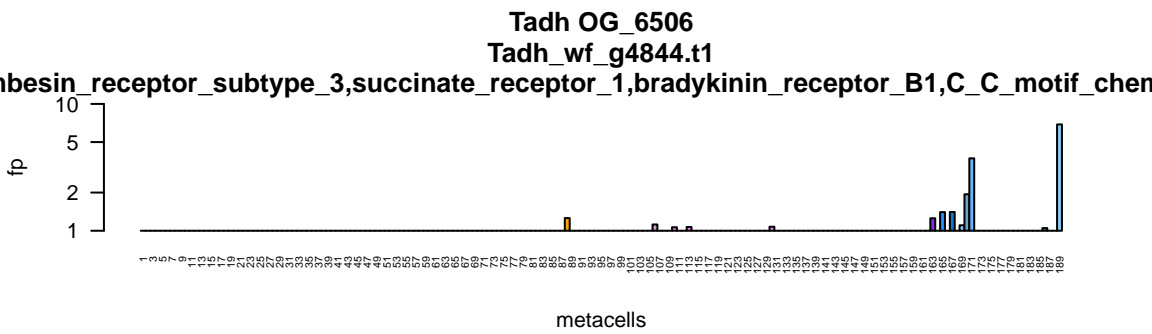
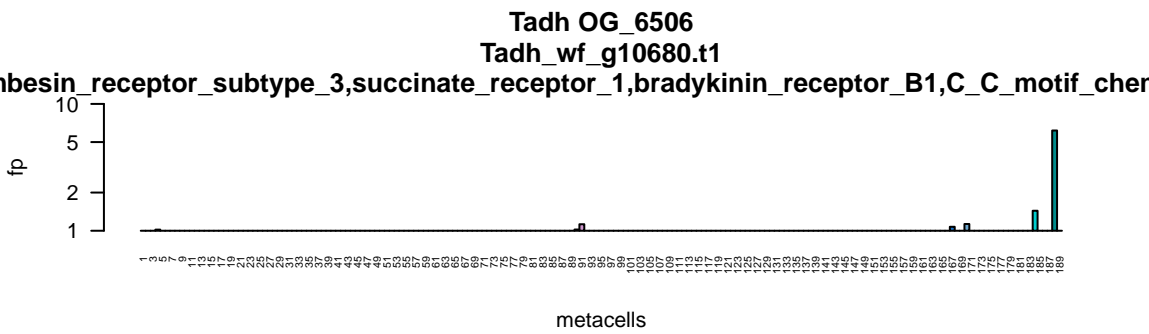
HoiH23 OG_4788
HoiH23_PIH23_005445-RA
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi



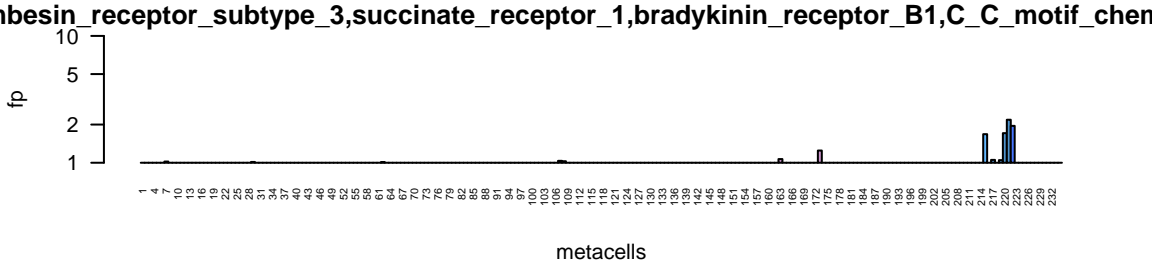




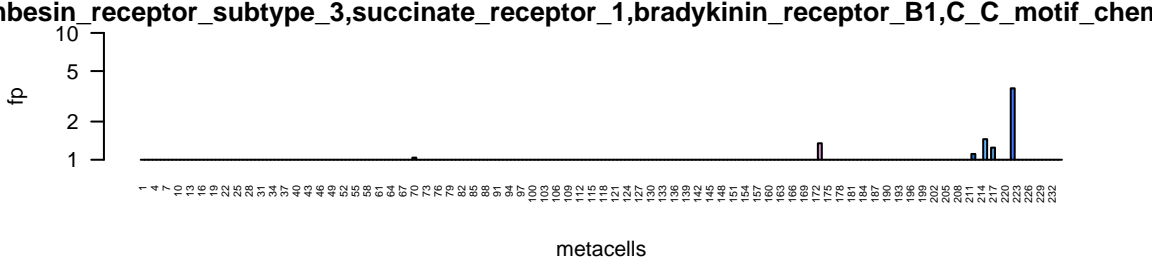


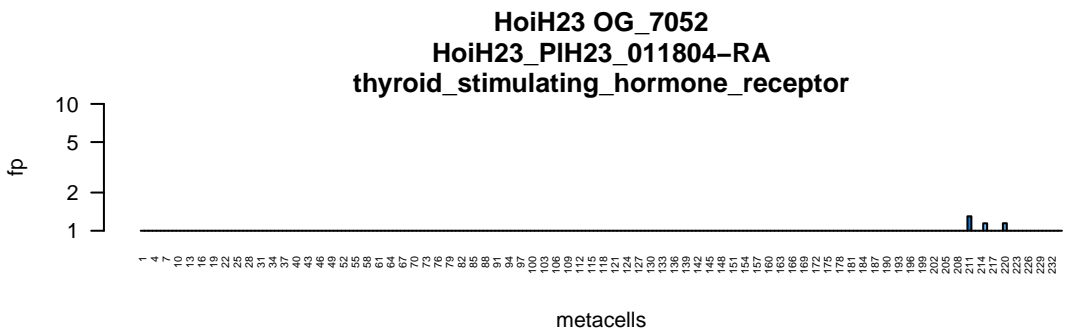
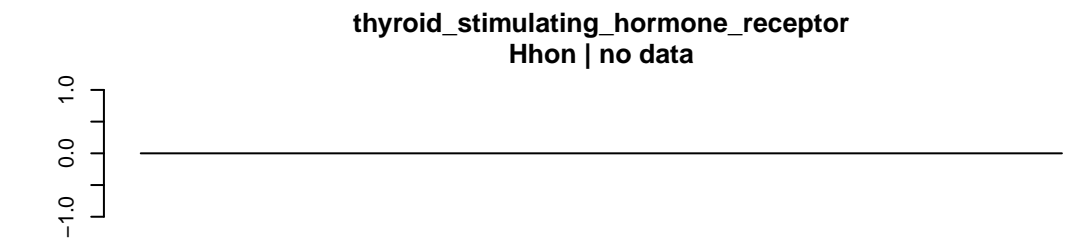
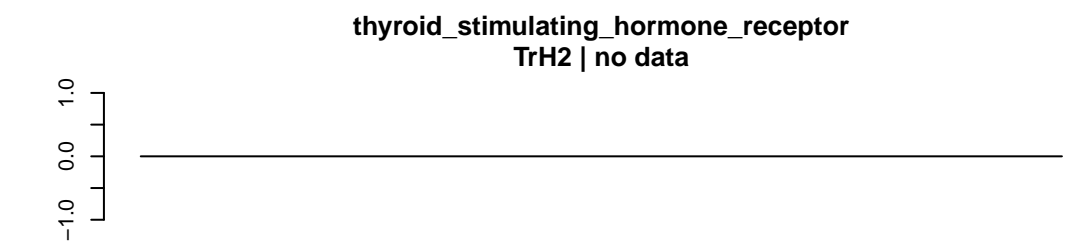
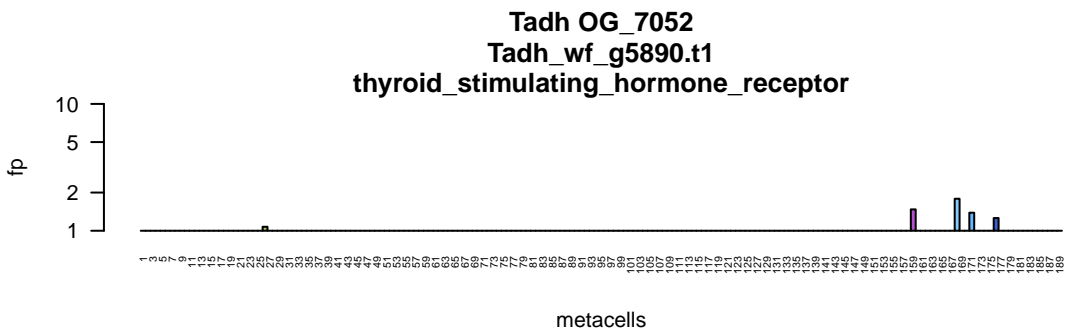


HoiH23 OG_6506
HoiH23_PIH23_008211-RA



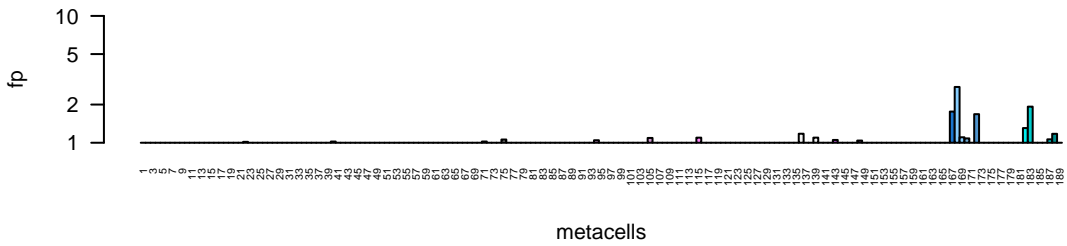
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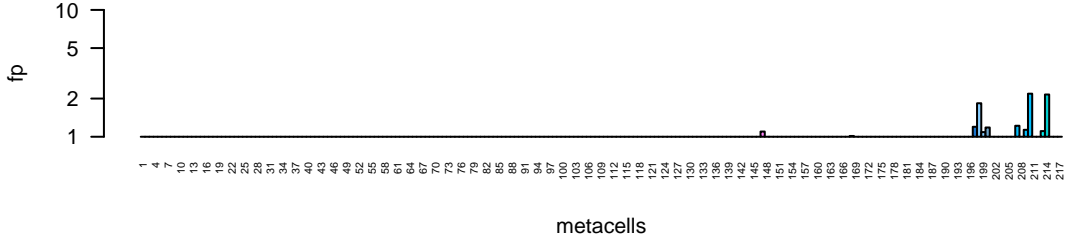
Tadh OG_9033

Tadh_TriadT57072



TrH2 OG_9033

TrH2_TrispH2_002408-RA



Hhon | no data



HoiH23 | no data



