

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

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists 189 metacells. Most metacells have a false positive count of 1, with a few having counts of 2 or 3. Metacells 175, 176, and 179 have the highest counts, all at 4.

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

Bar chart showing the frequency (fp) of metacells. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists 180 metacells. Most metacells have a frequency of 1, indicated by a horizontal line. A few metacells have higher frequencies: metacell 121 has fp=2, metacell 161 has fp=2, metacell 171 has fp=1, metacell 172 has fp=4, metacell 173 has fp=3, metacell 174 has fp=2, metacell 175 has fp=1, metacell 176 has fp=1, metacell 177 has fp=1, metacell 178 has fp=1, metacell 179 has fp=1, metacell 180 has fp=1.

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists 180 metacells. Most metacells have 0 false positives, but some have 1, 2, or more. The bars are colored in a repeating pattern of black, purple, and green.

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists 189 metacells. The bars are colored in a gradient from blue to red. Most metacells have a false positive count of 1, with a few having counts of 2 or 3.

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

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

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

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 217. The bars are colored in a repeating pattern of light blue, light green, and light red. Most metacells have a false positive count of 1, with some having 2 or 3. Metacells 178 and 181 have the highest false positive counts, at 5 and 4 respectively.

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

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

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

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

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

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

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

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

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

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

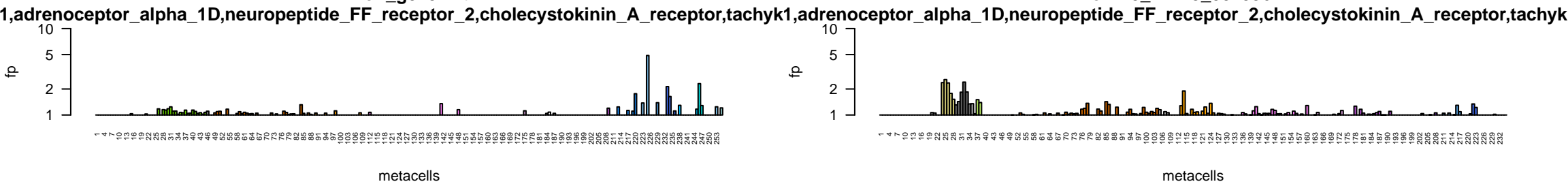
A bar chart showing the number of false positives (fp) for each metacell. The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 0 to 10. The chart shows a distribution of false positives across metacells, with most having 0 or 1 false positive, and a few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

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

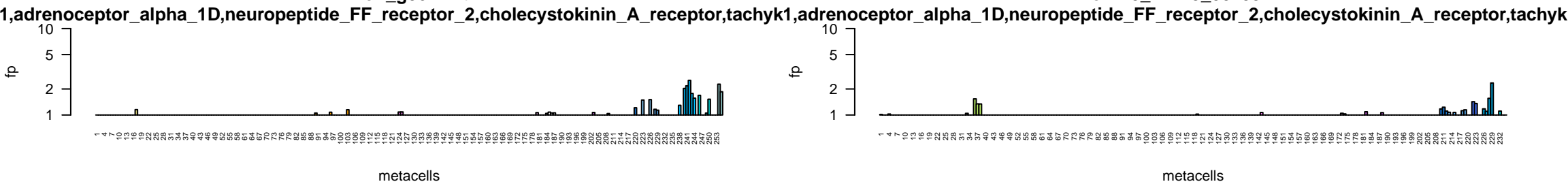
A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis ranges from 1 to 250, and the y-axis ranges from 1 to 10. Most metacells have a frequency of 1, with a small cluster of higher frequencies around metacell 240.

metacells	fp
1	1
7	1
10	1
16	1
18	1
19	1
25	1
26	1
28	1
34	1
37	1
43	1
46	1
55	1
61	1
64	1
70	1
76	1
79	1
85	1
88	1
94	1
98	1
103	1
112	1
115	1
121	1
122	1
127	1
130	1
136	1
145	1
148	1
154	1
157	1
163	1
166	1
172	1
175	1
181	1
184	1
187	1
190	1
196	1
199	1
205	1
208	1
214	1
217	1
223	1
228	1
232	1
235	1
236	1
241	1
242	1
244	1
247	1
250	1
253	1

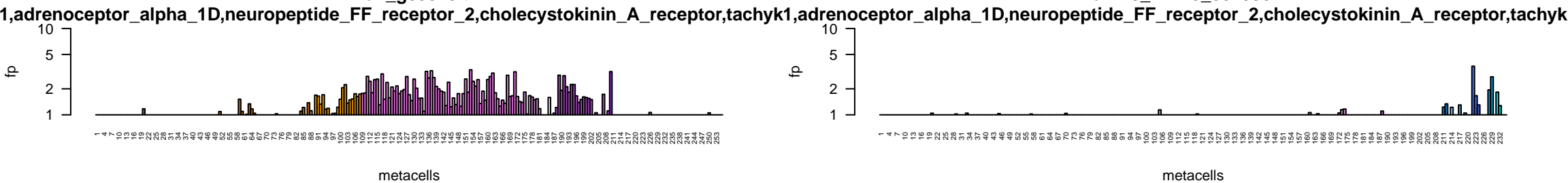
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Hhon_g04574.t1



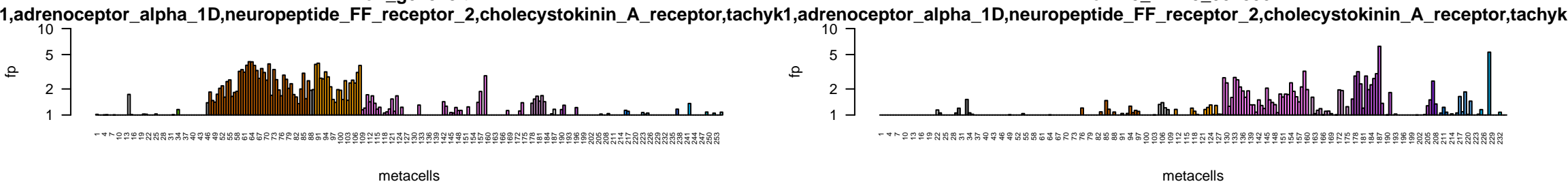
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Hhon_g06114.t1



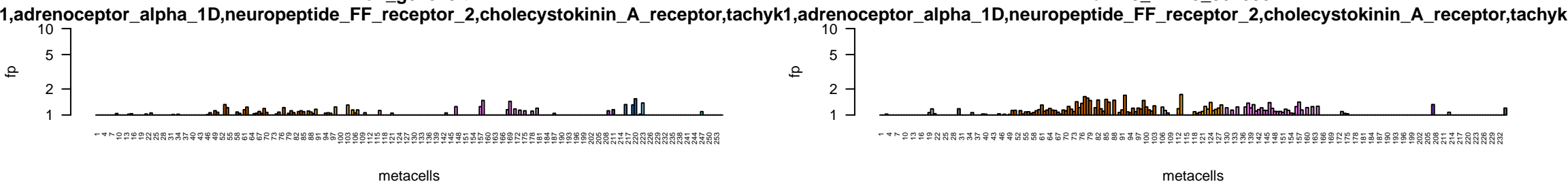
Hhon OG_5725
Hhon_g09573.t1



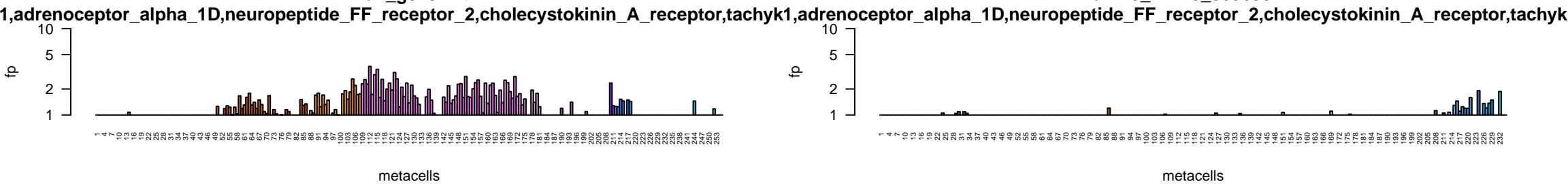
Hhon OG_5725
Hhon_g07528.t1



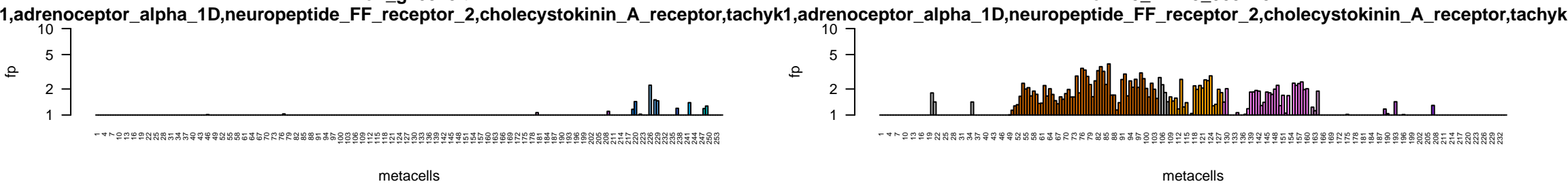
Hhon OG_5725
Hhon_g07529.t1



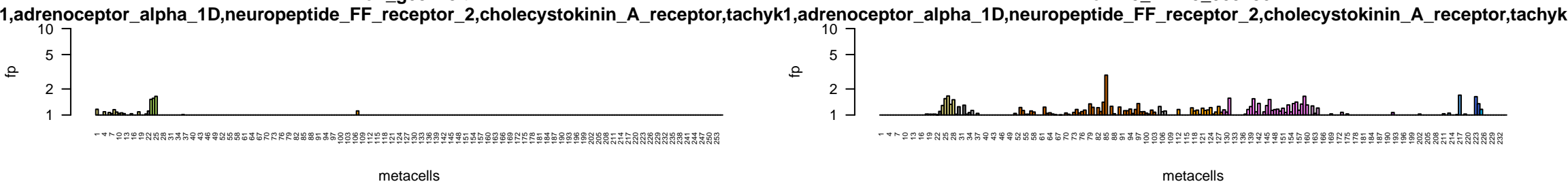
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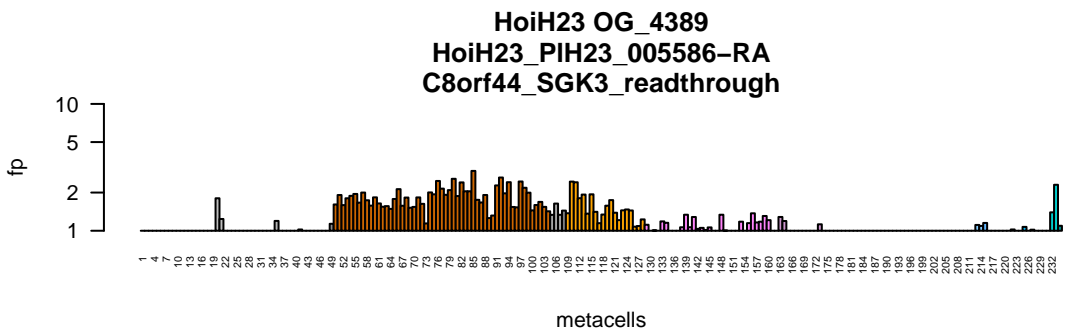
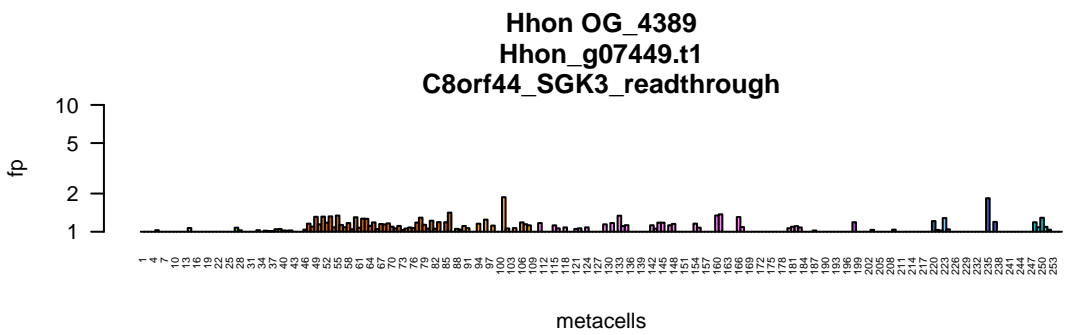
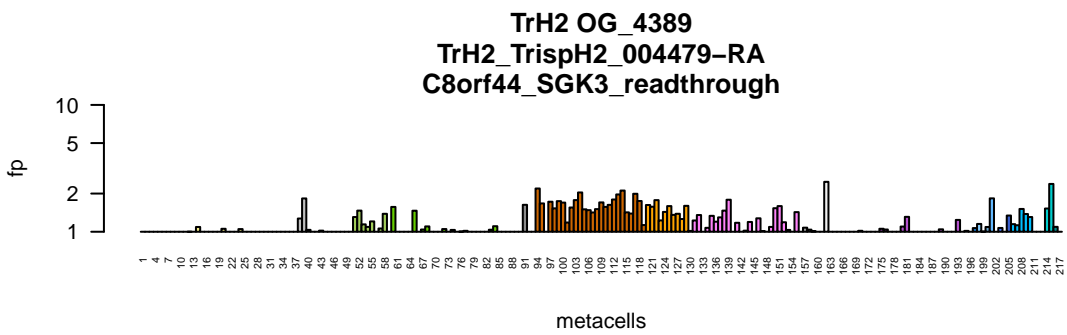
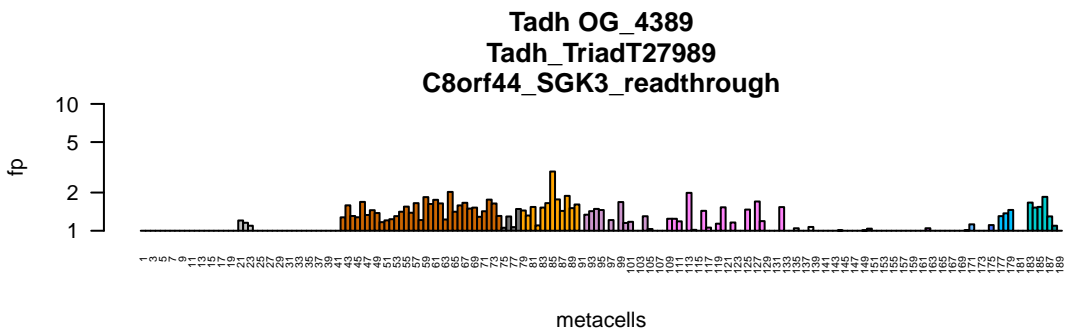


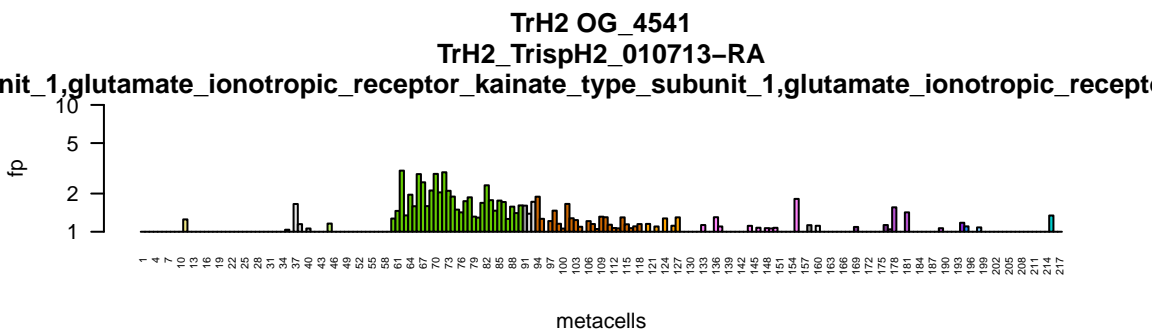
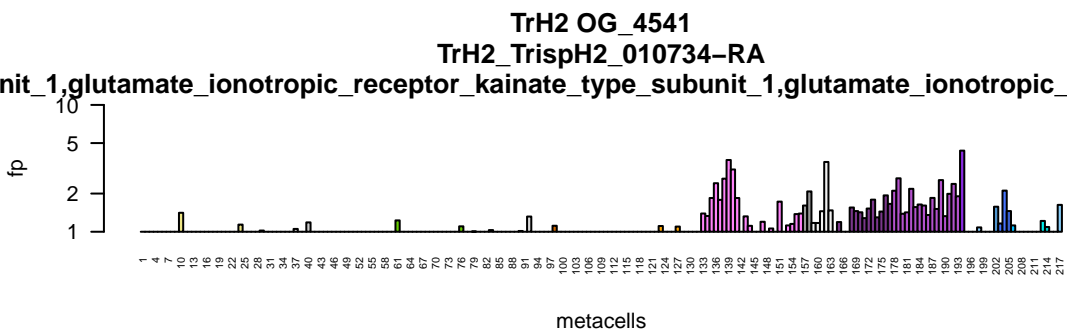
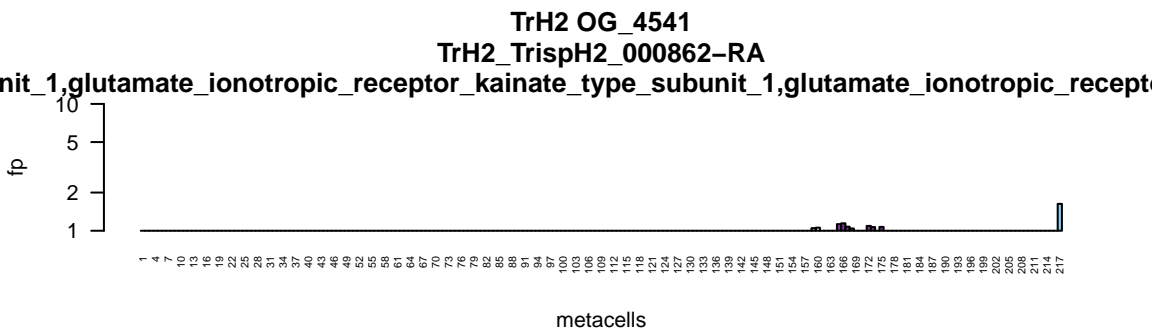
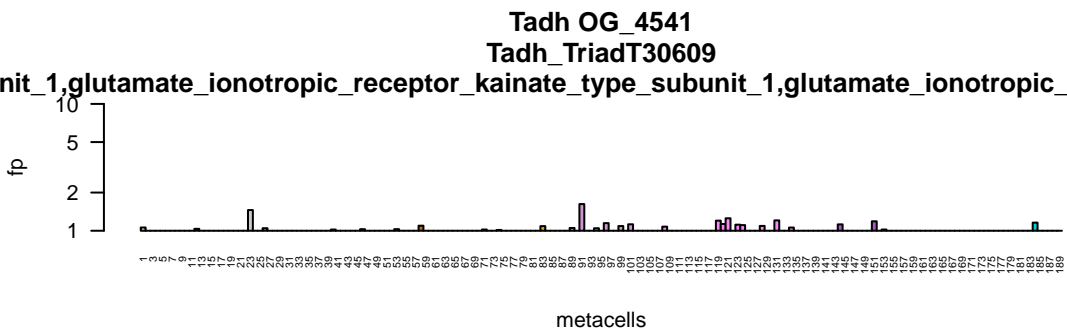
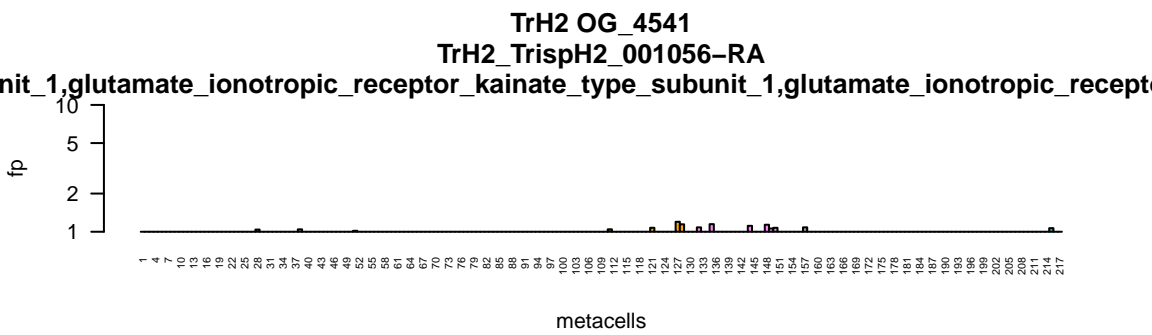
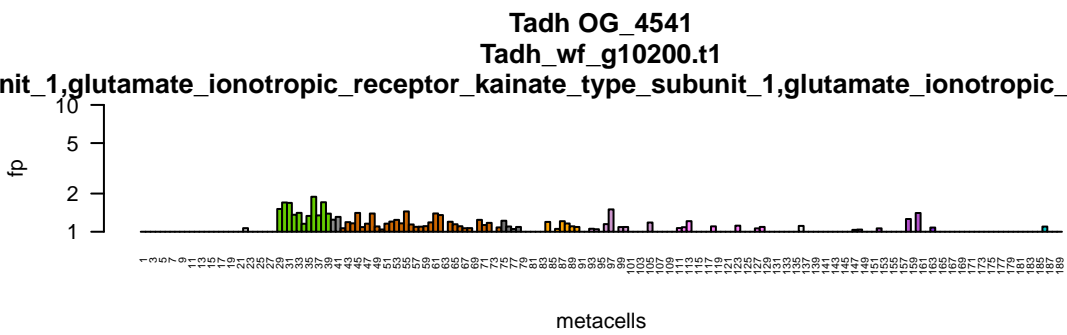
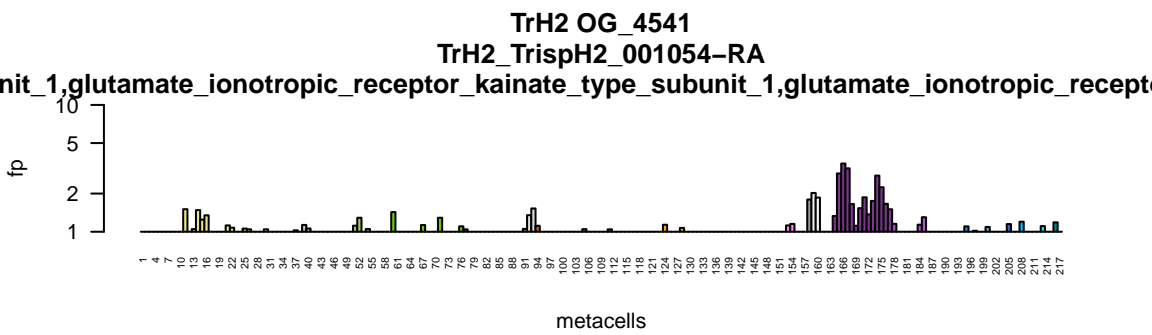
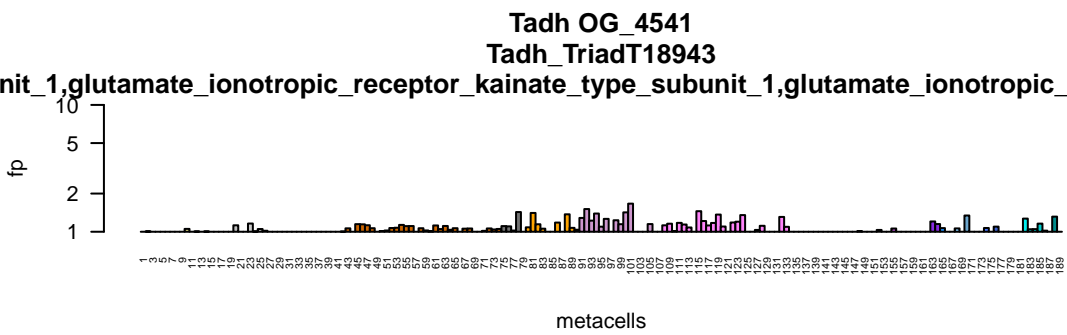
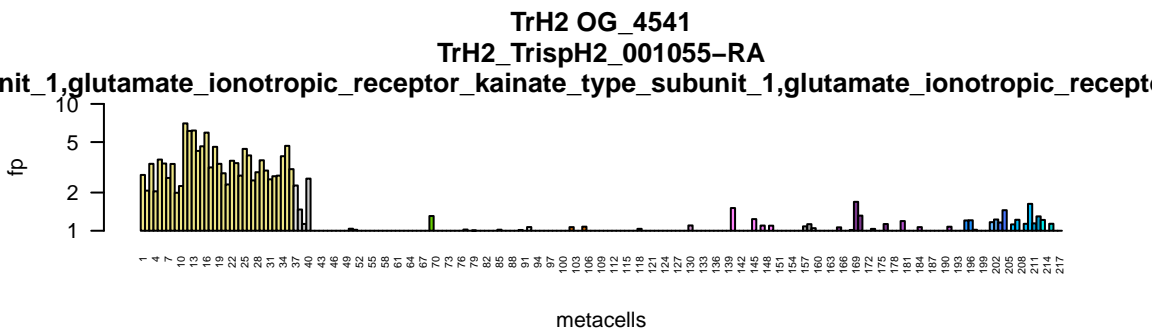
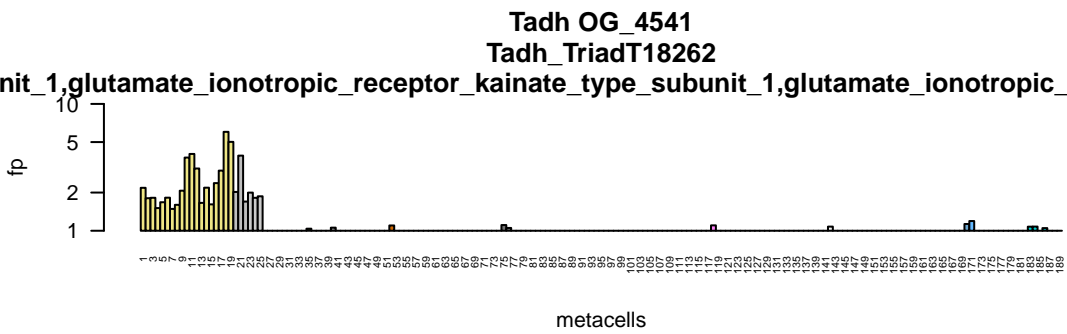
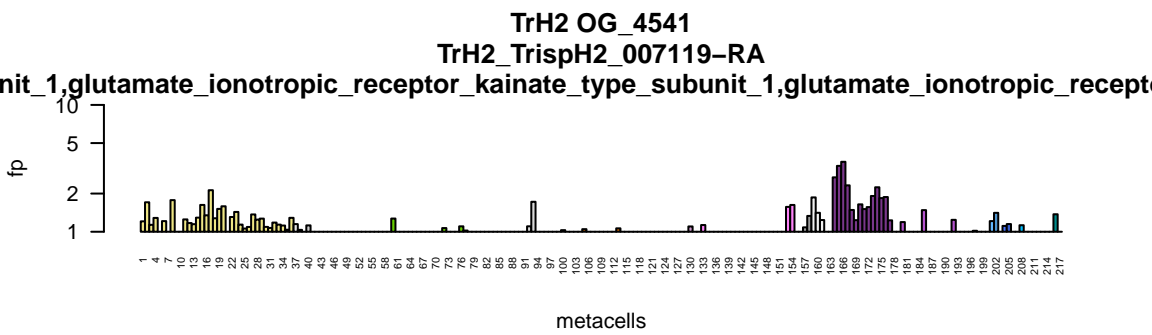
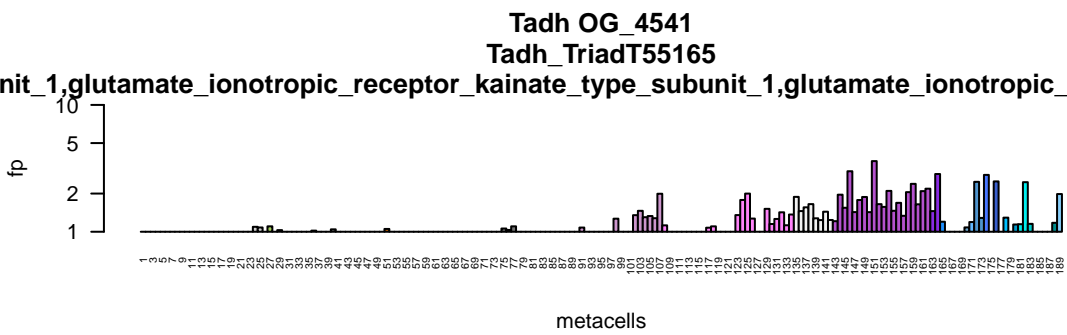
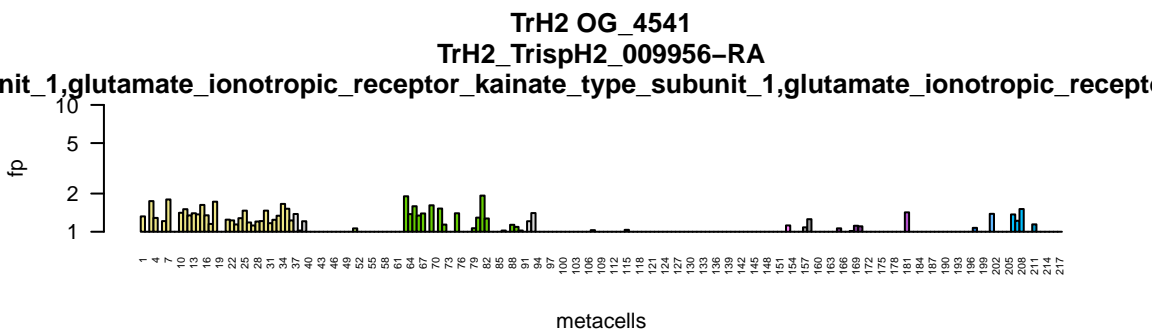
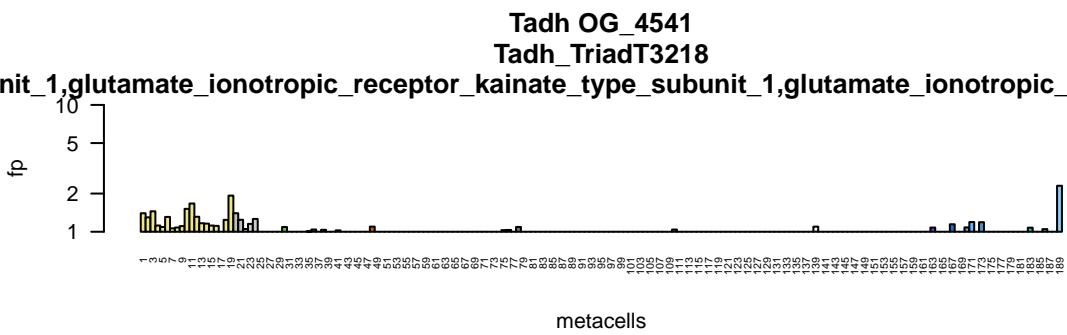
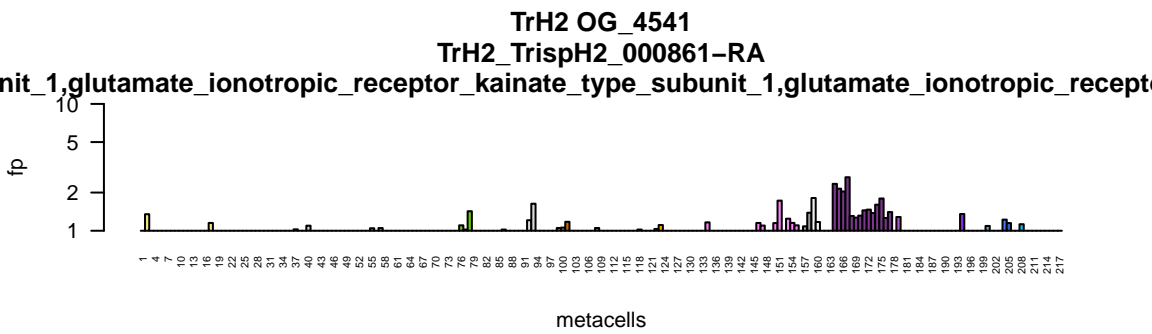
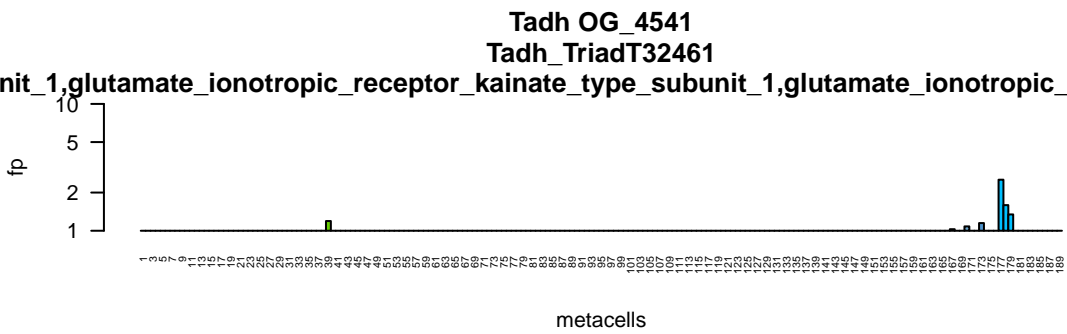
Hhon OG_5725
Hhon_g10923.t1

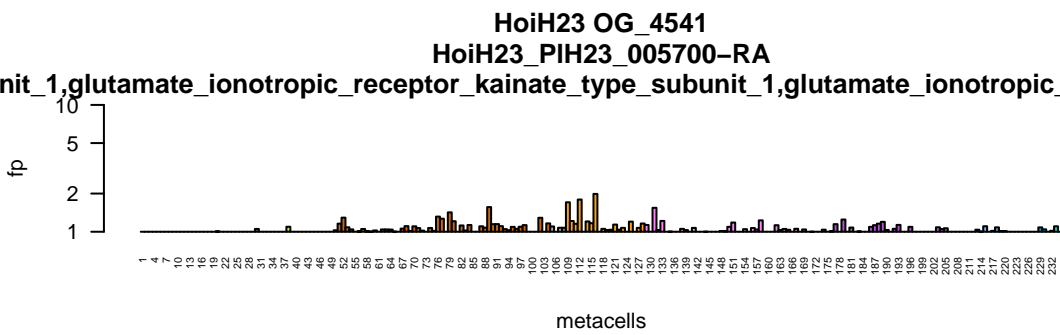
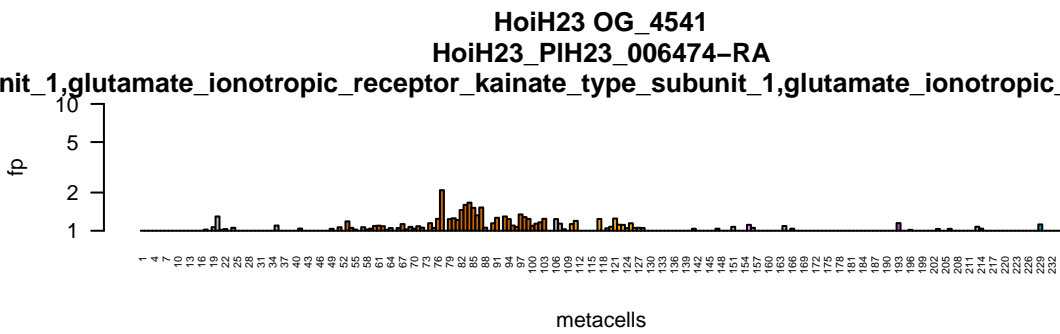
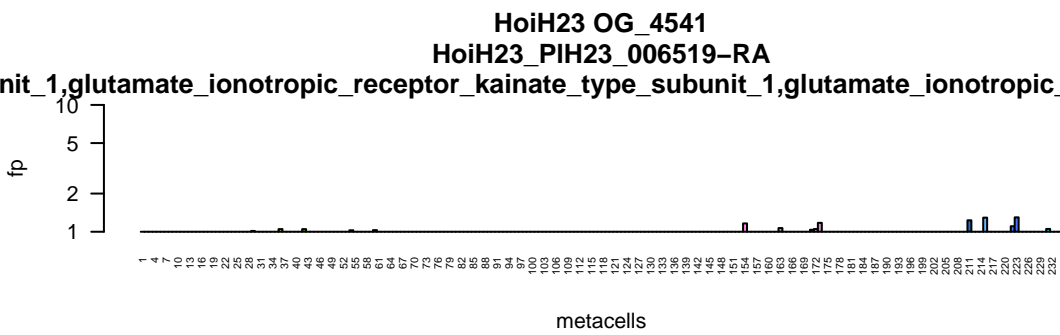
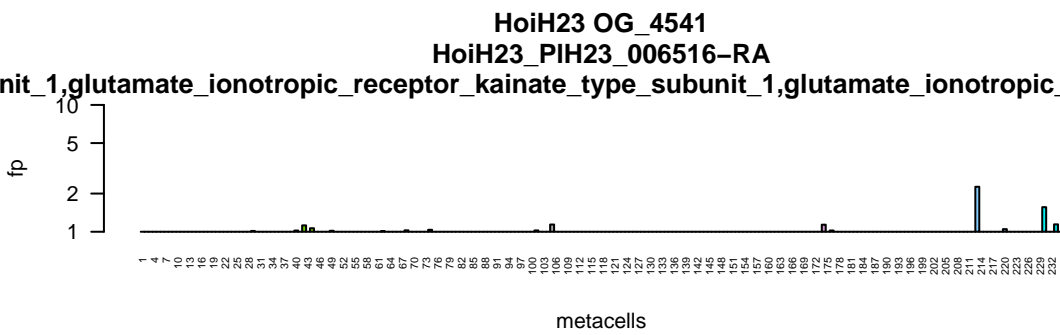
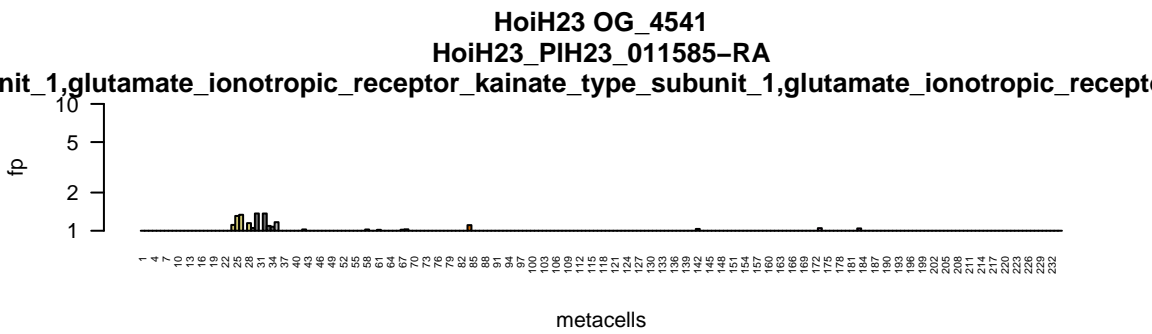
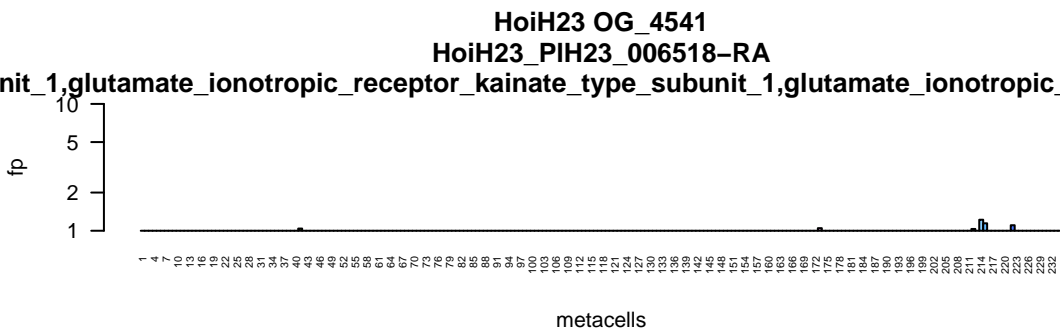
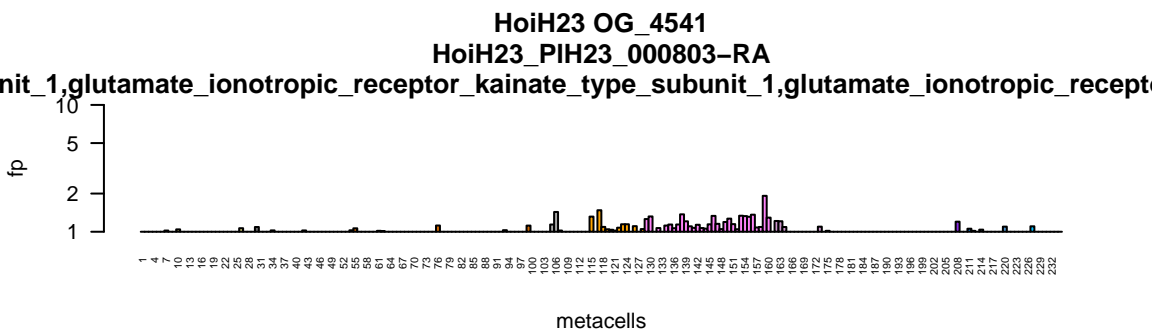
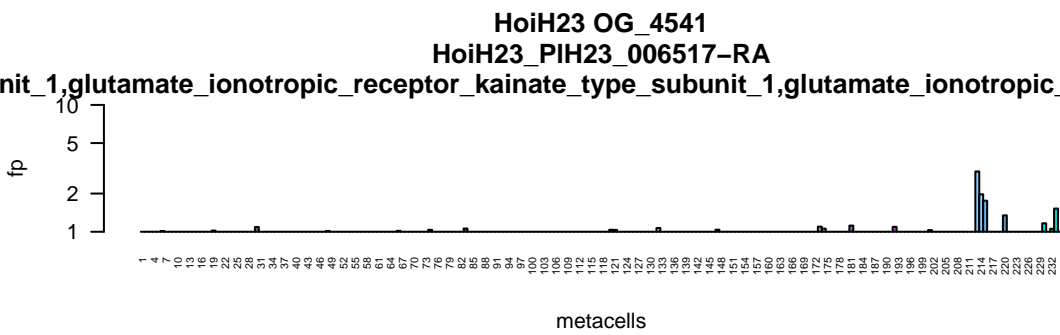
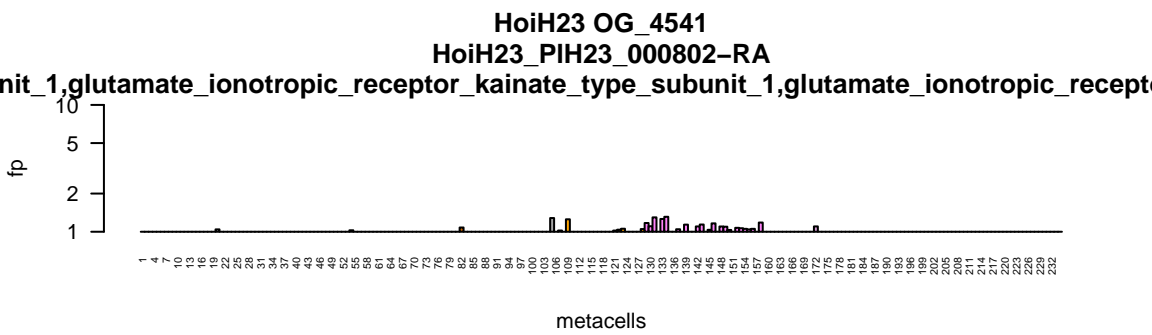
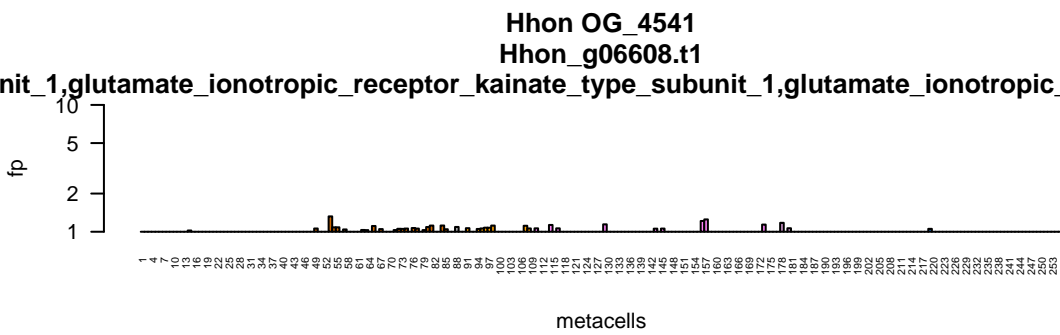
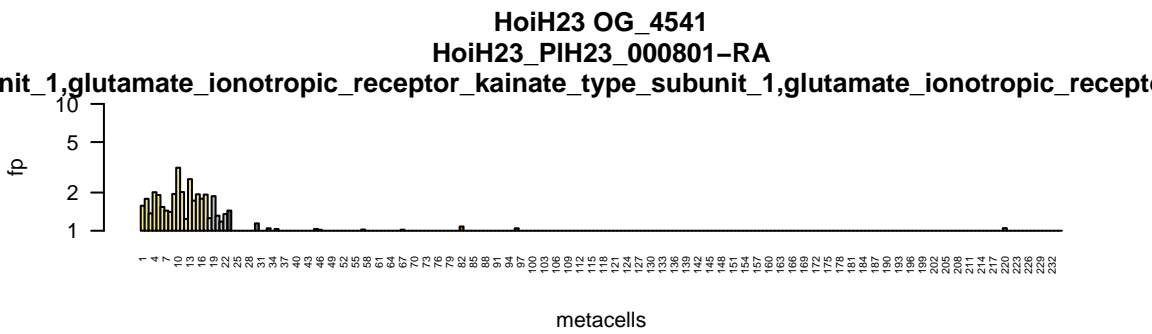
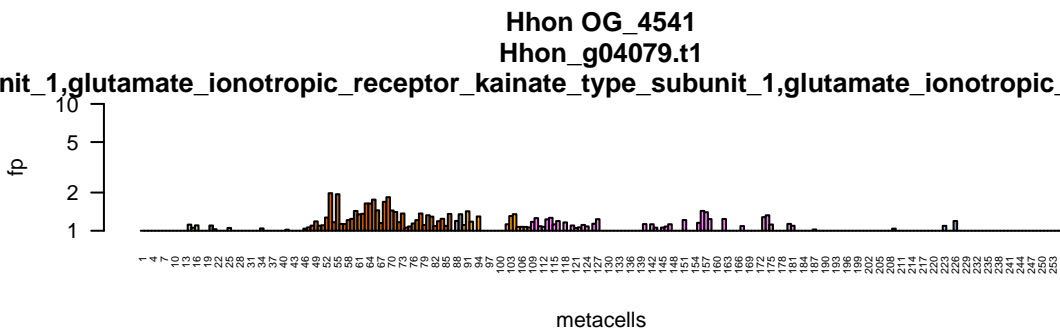


Hhon OG_5725
Hhon_g08418.t1





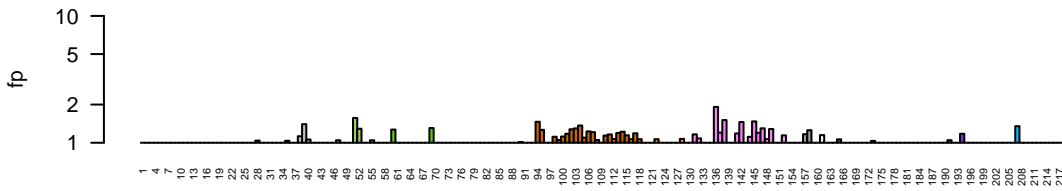




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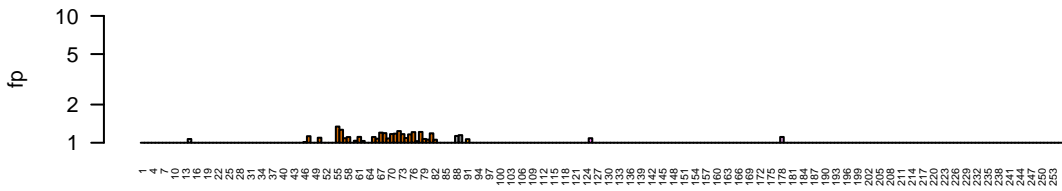


TrH2 OG_7017
TrH2_TrispH2_003191-RA



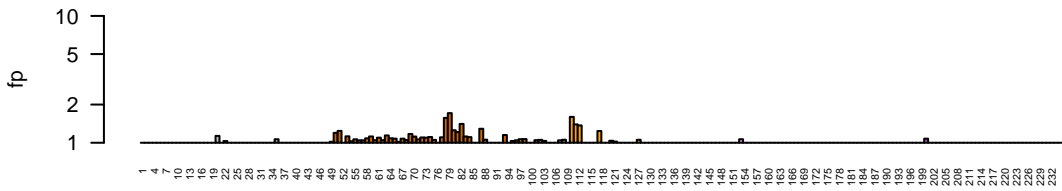
metacells

Hhon OG_7017
Hhon_g03674.t1



metacells

HoiH23 OG_7017
HoiH23_PIH23_002076-RA



metacells

Bar chart showing the frequency of metacells (x-axis) versus frequency (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 214. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows a distribution of frequencies across metacells, with most metacells having a frequency of 1, and a few having higher frequencies up to 10.

Bar chart showing the frequency of metacells (x-axis) versus frequency (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 214. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows a distribution of frequencies across metacells, with most metacells having a frequency of 1, and a few having higher frequencies (up to 2).

TrH2 OG_2962
TrH2_TripSH2_011702-RA
CASP2_and_RIPK1_domain_containing_adaptor_with_death_domain,Fas_associat

fp

metacells

TrH2 OG_2962
TrH2_TripSH2_011686-RA
CASP2_and_RIPK1_domain_containing_adaptor_with_death_domain,Fas_associat

metacell	fp
1	1
4	1
10	1
13	1
16	1
19	1
21	1
25	1
28	1
31	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
59	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	1.5
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1.2
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	1.5

TrH2 OG_2962
TrH2_TripSH2_011703-RA
CASP2_and_RIPK1_domain_containing_adaptor_with_death_domain,Fas_associat

fp

metacells

TrH2 OG_2962
TrH2_TripSH2_011880-RA
CASP2_and_RIPK1_domain_containing_adaptor_with_death_domain,Fas_associat

fp

metacells

TrH2 OG_2962
TrH2_TripSH2_003510-RA
CASP2_and_RIPK1_domain_containing_adaptor_with_death_domain,Fas_associat

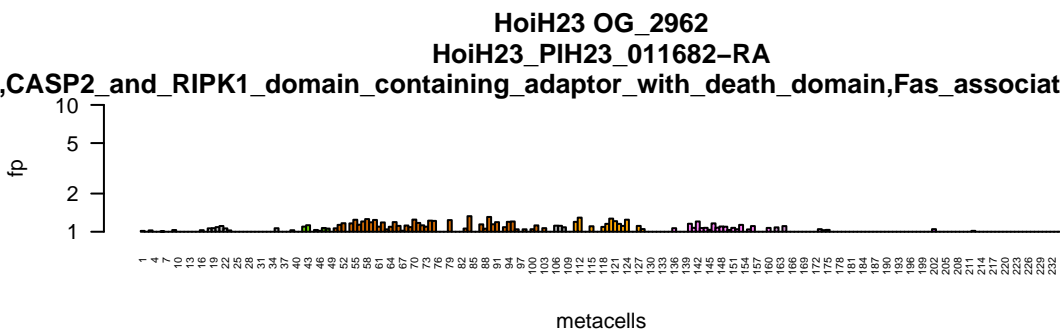
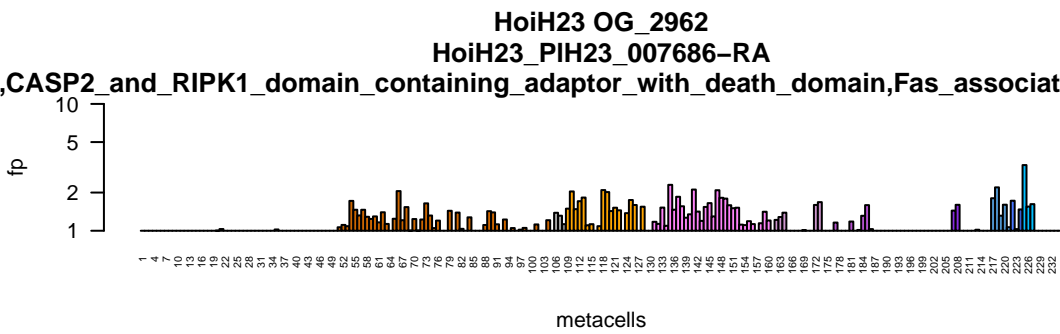
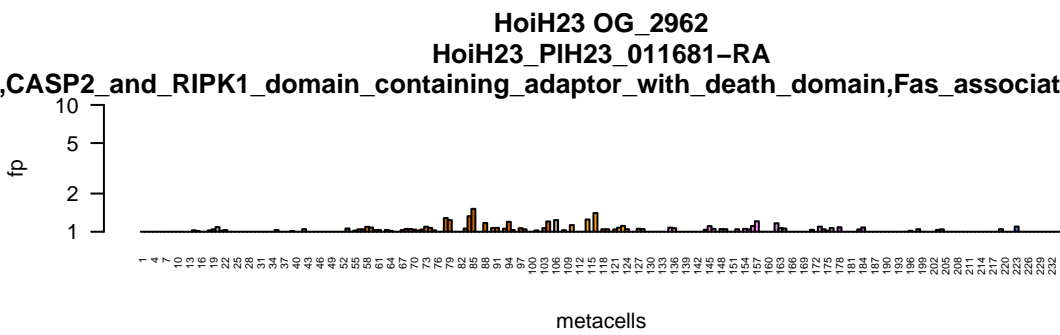
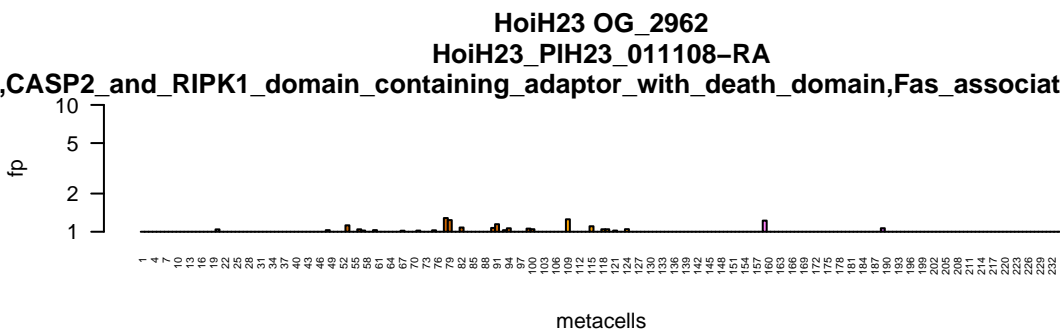
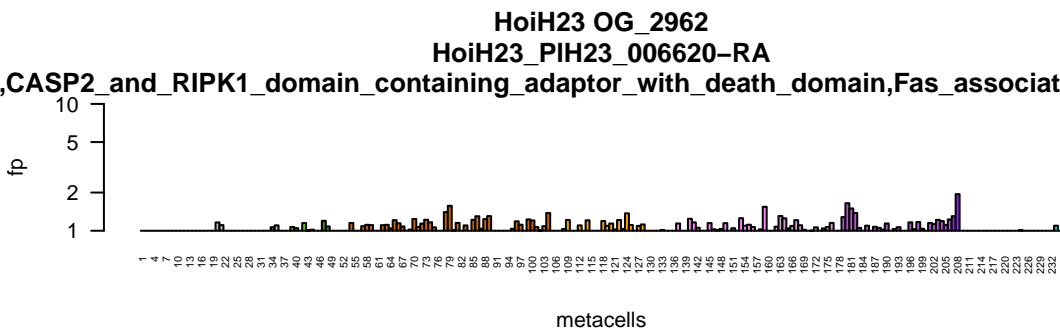
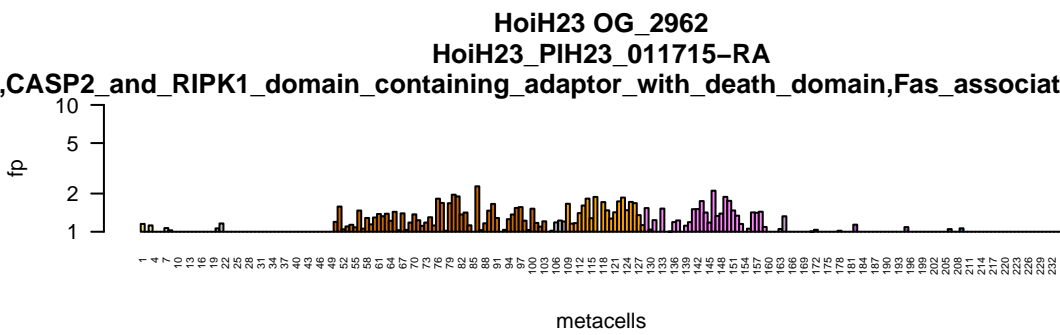
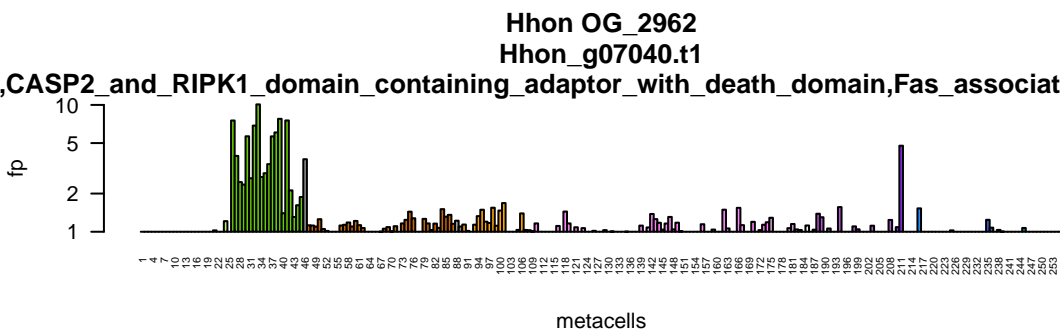
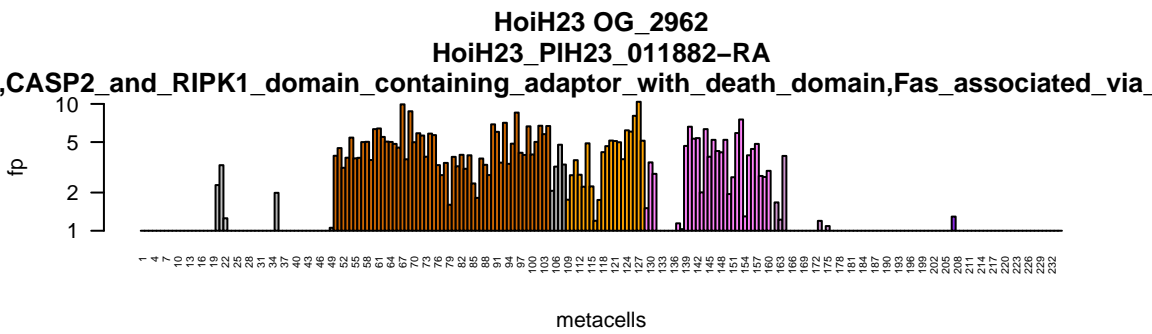
fp

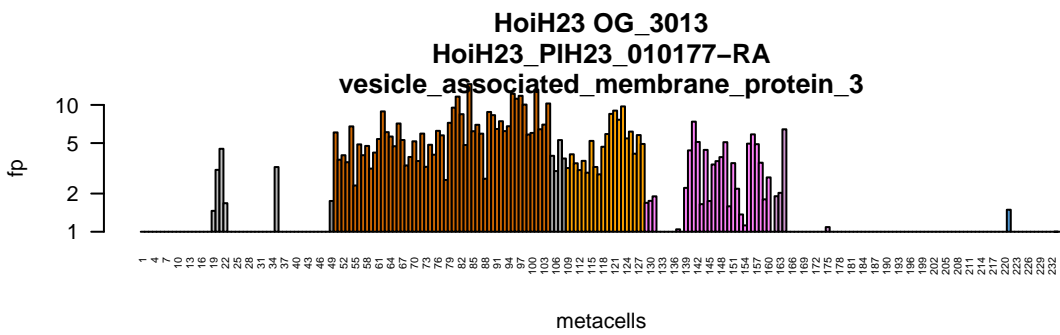
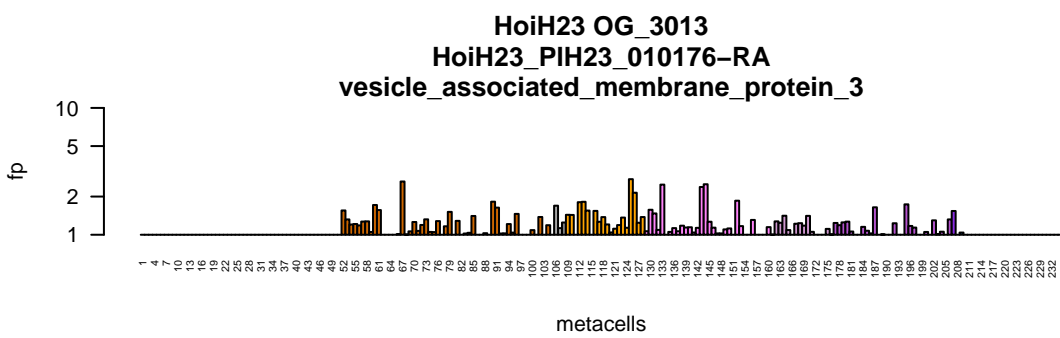
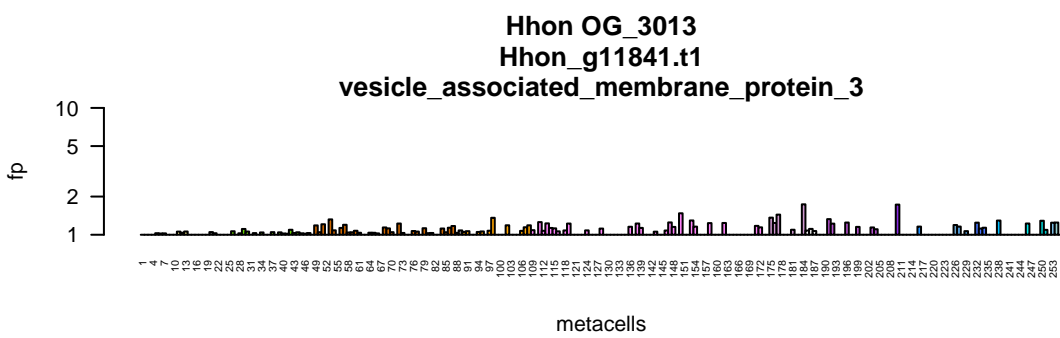
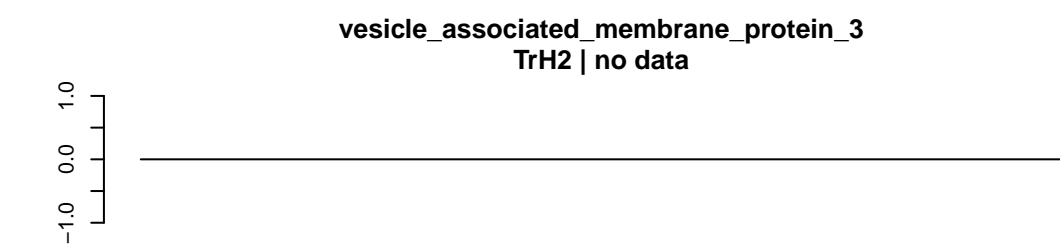
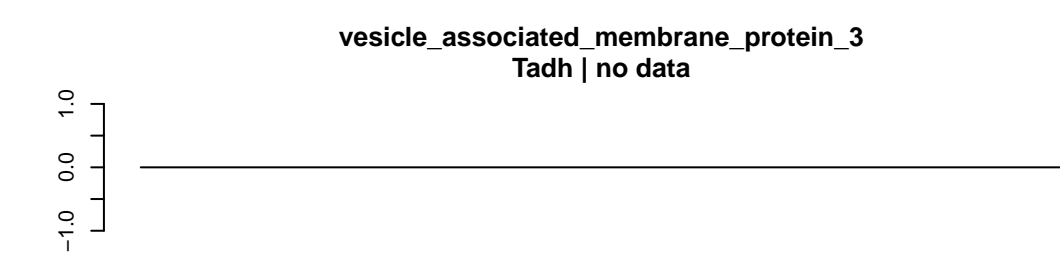
metacells

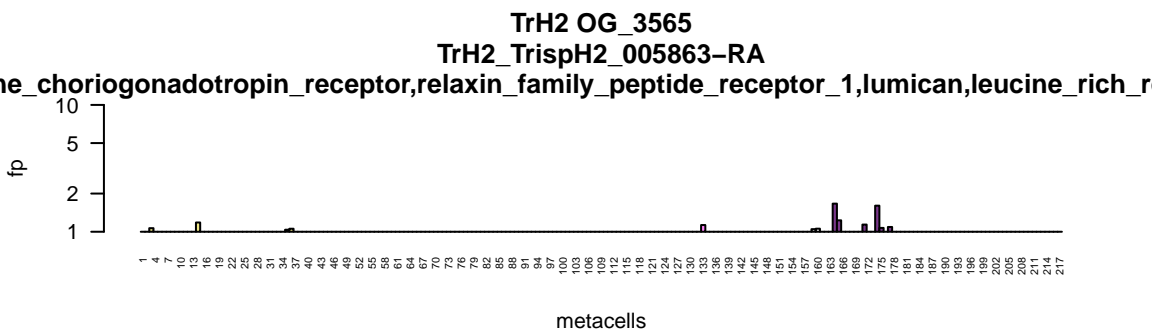
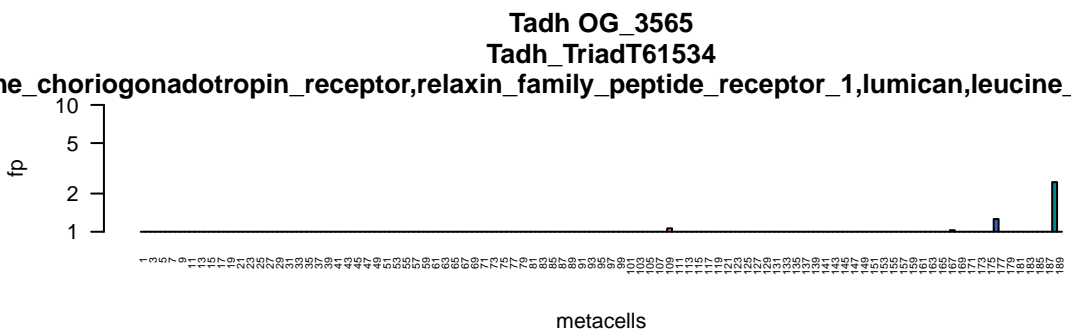
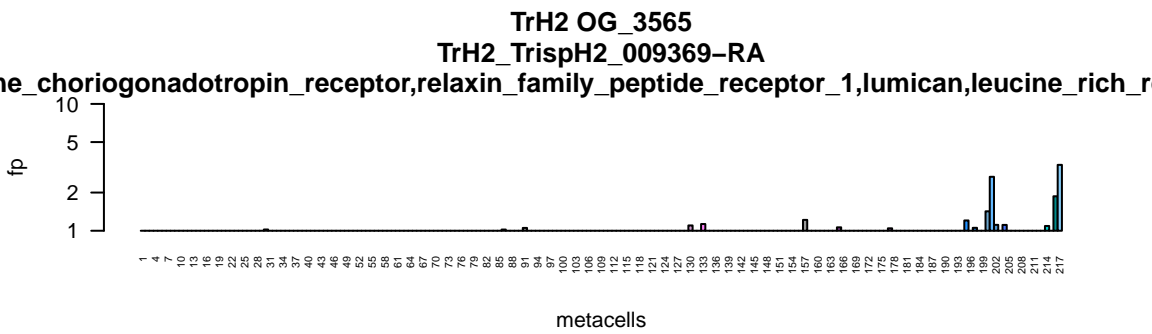
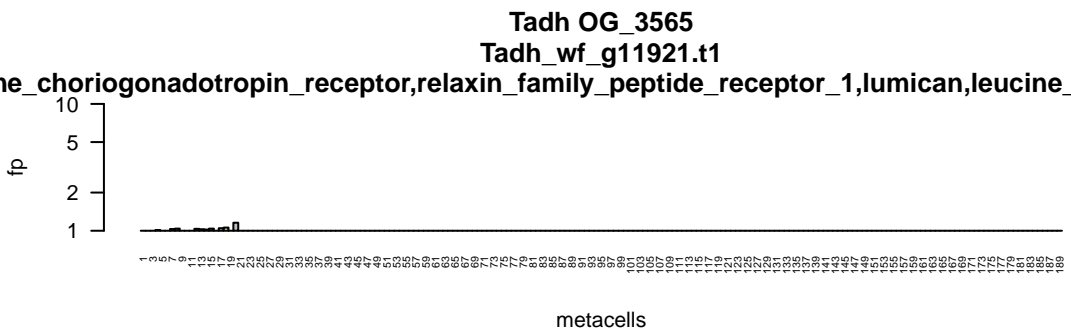
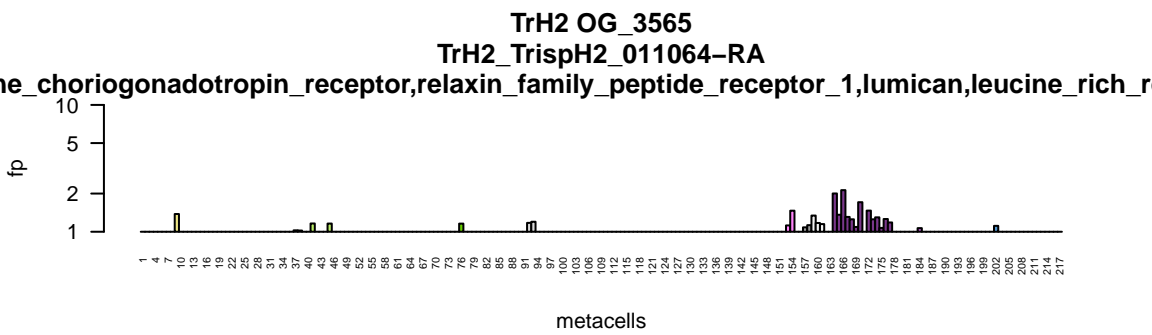
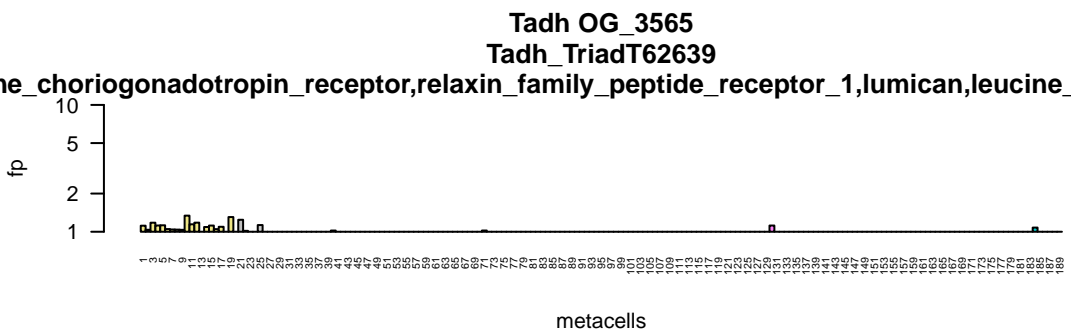
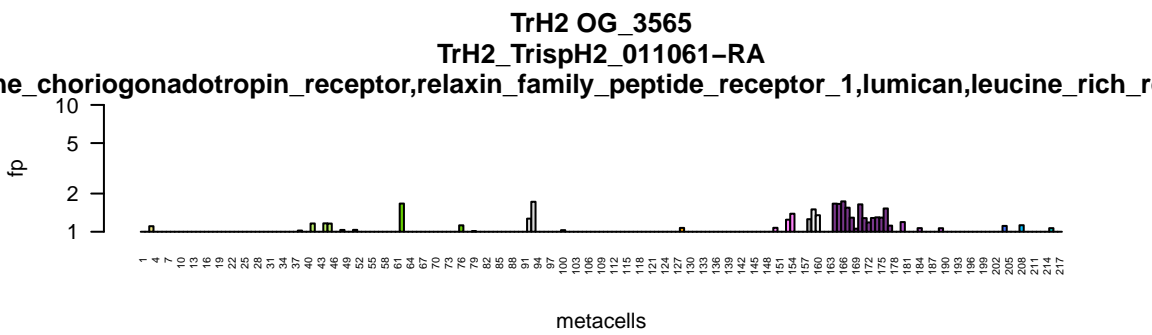
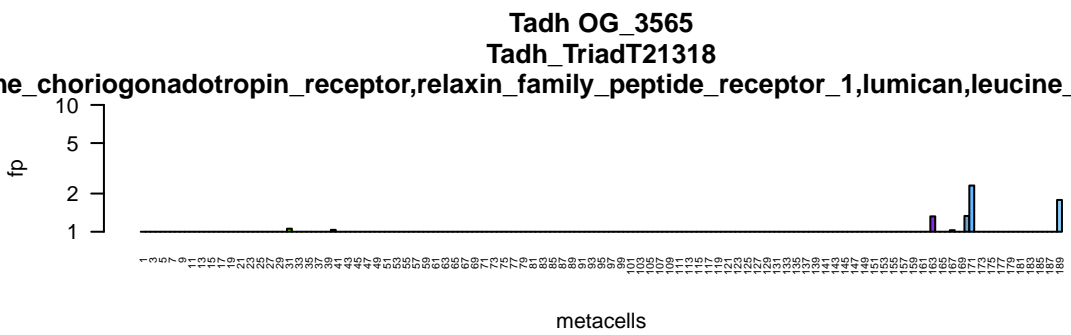
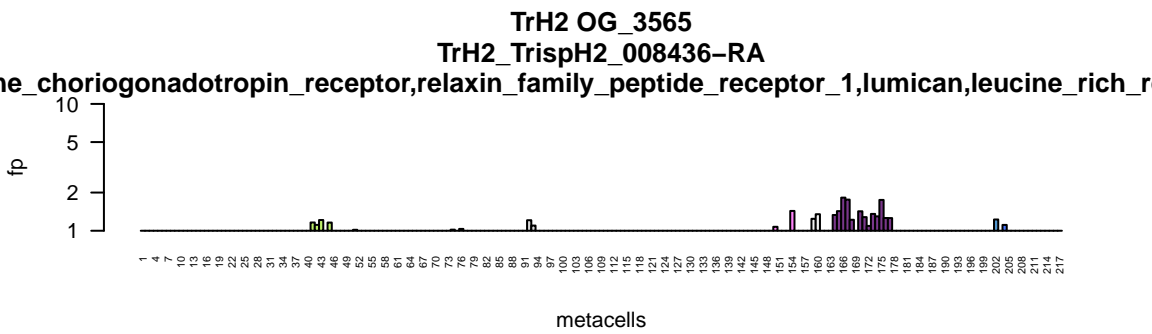
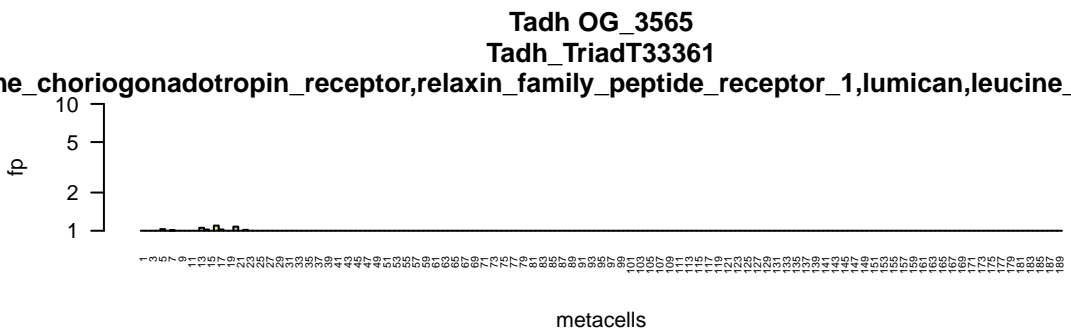
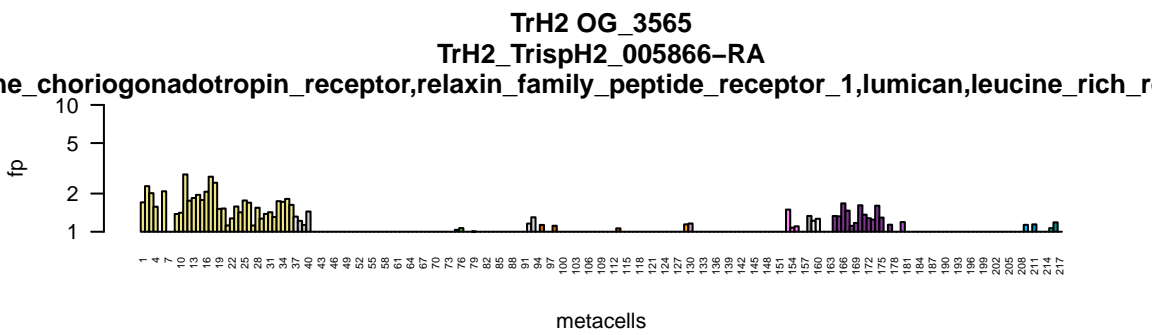
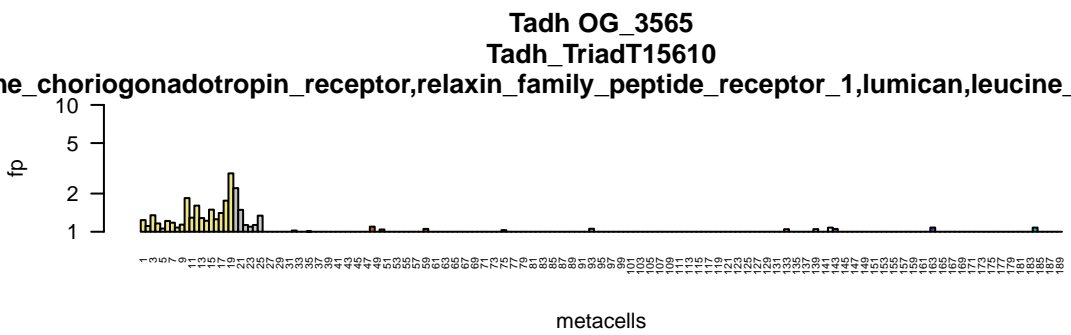
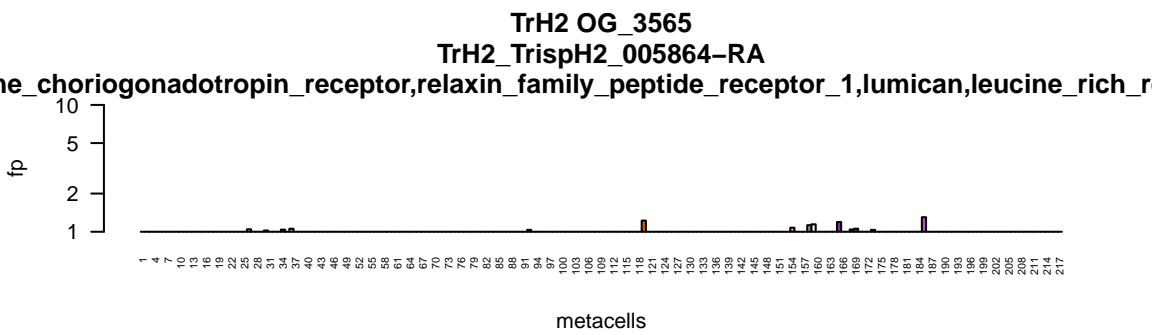
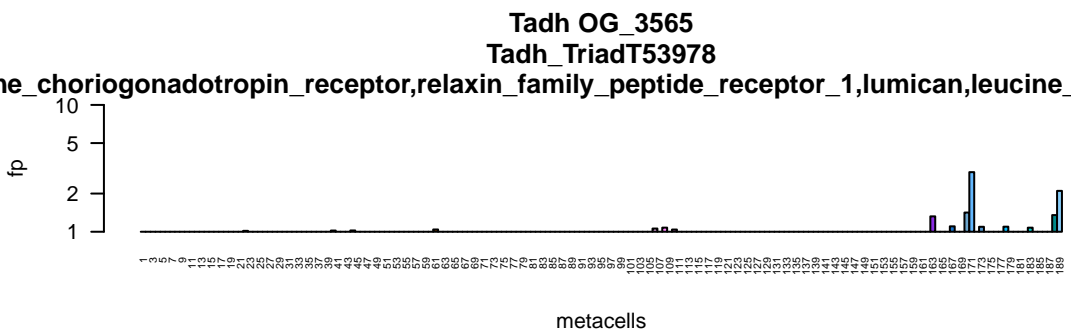
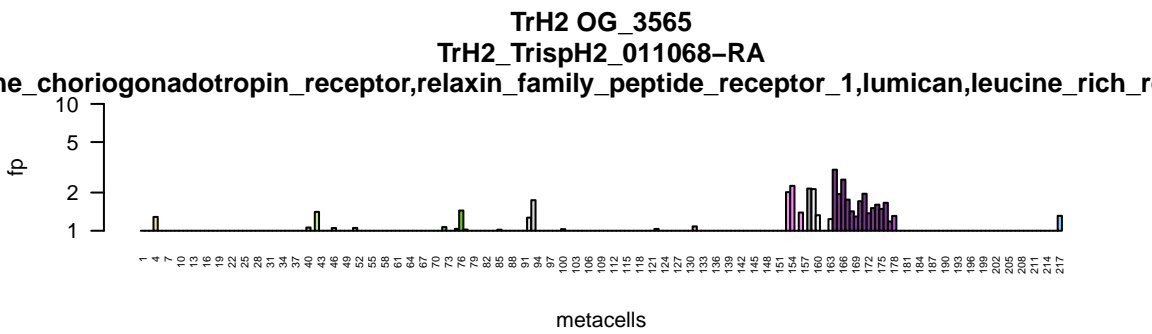
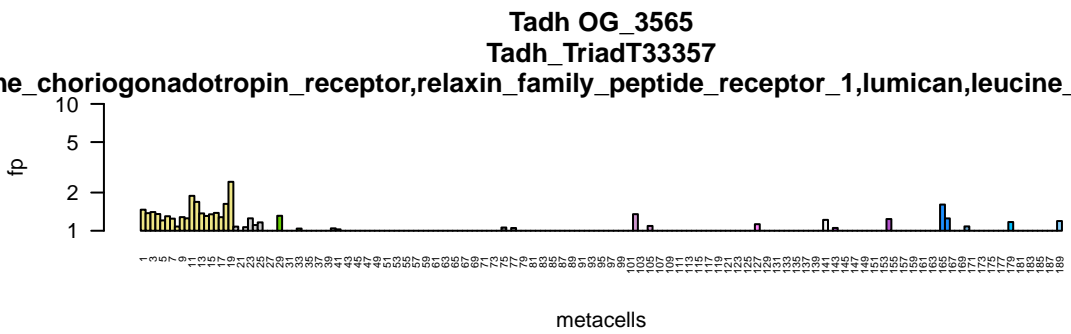
TrH2 OG_2962
TrH2_TripSH2_011821-RA
CASP2_and_RIPK1_domain_containing_adaptor_with_death_domain,Fas_associat

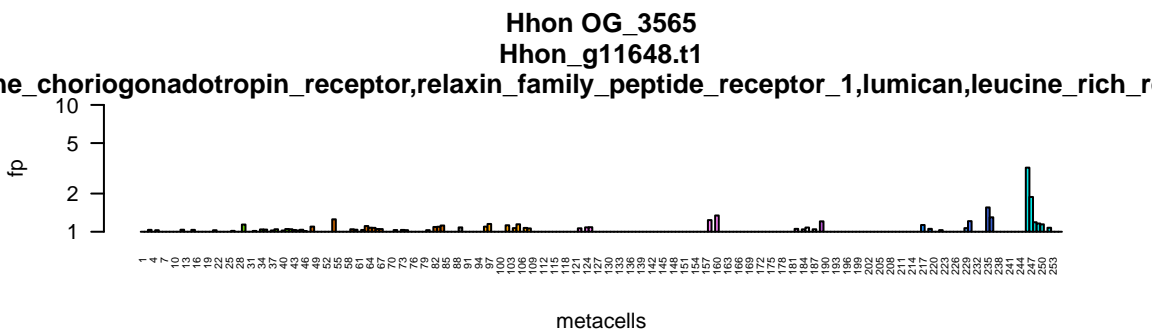
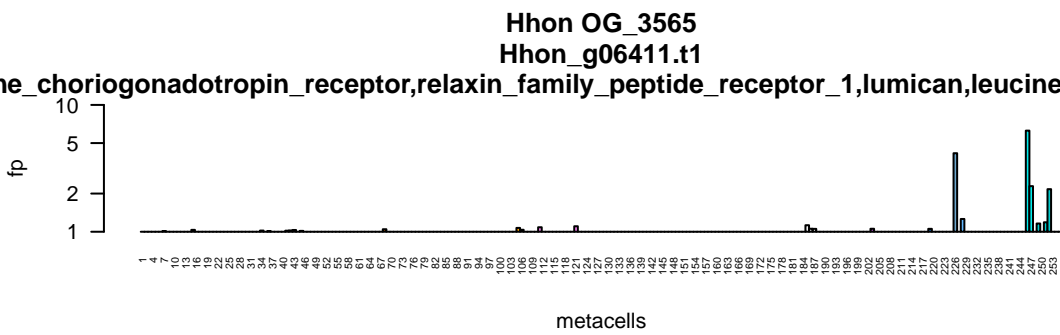
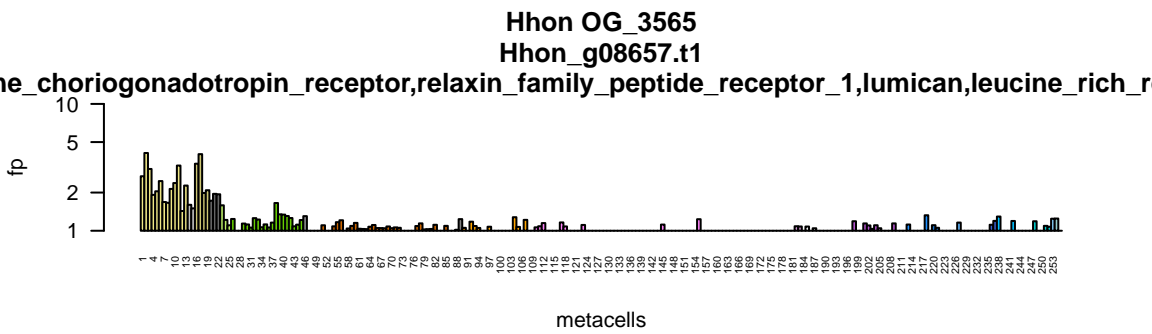
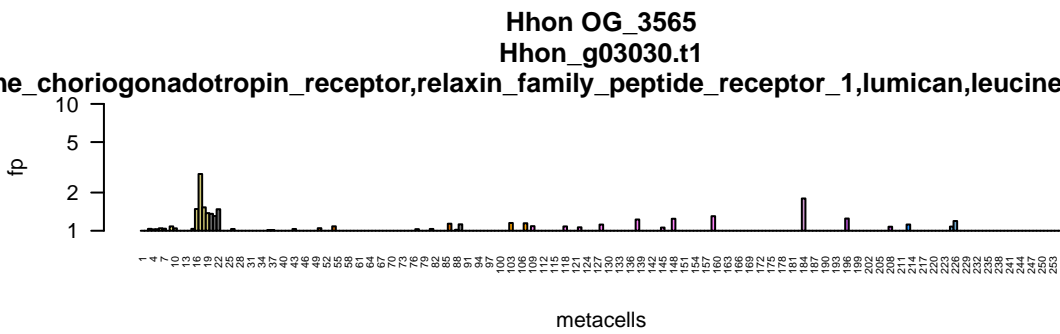
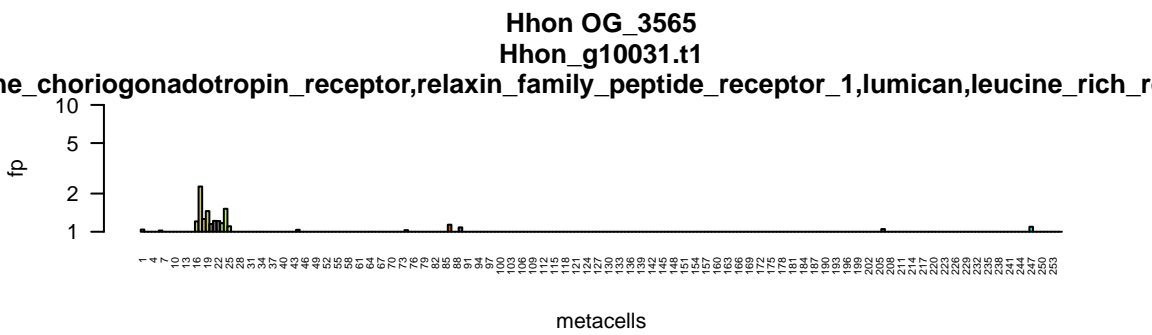
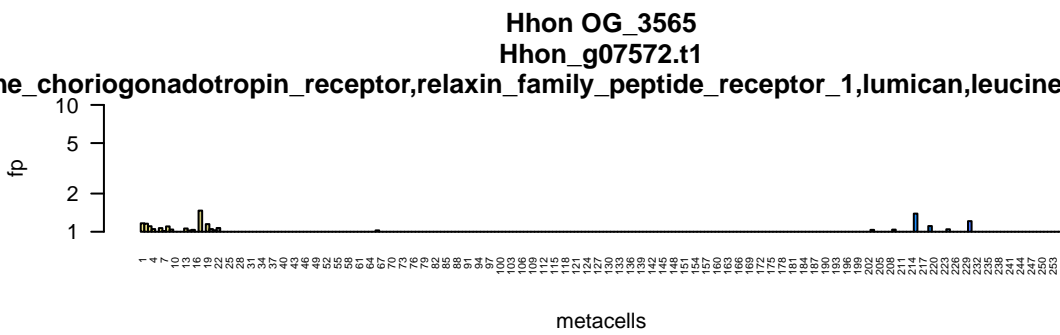
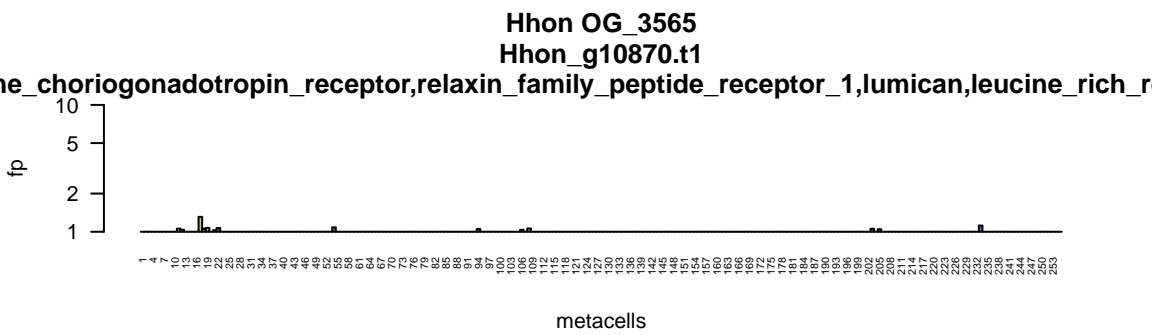
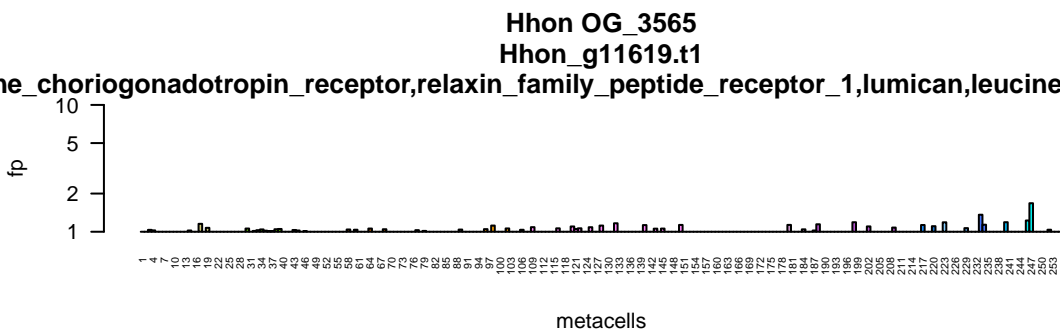
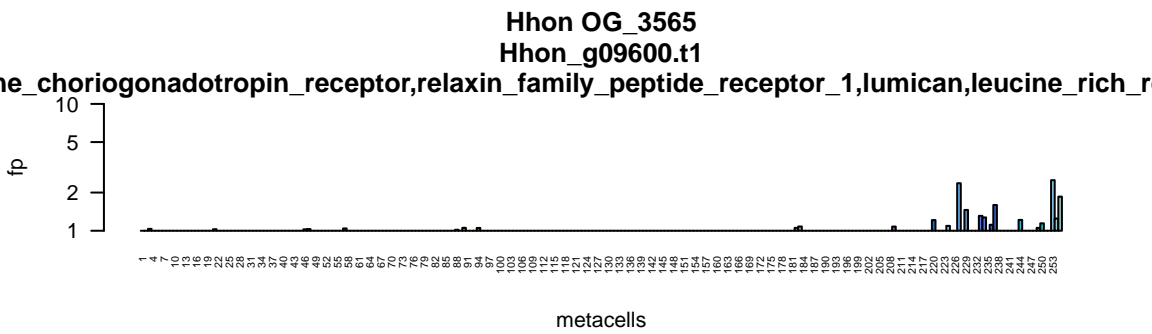
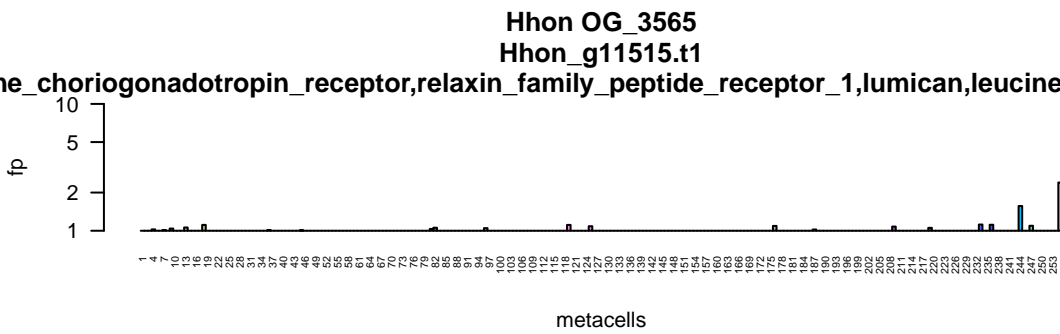
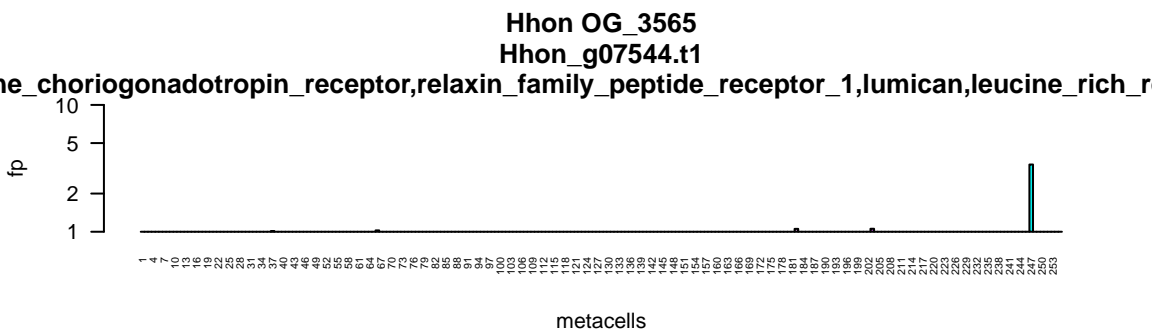
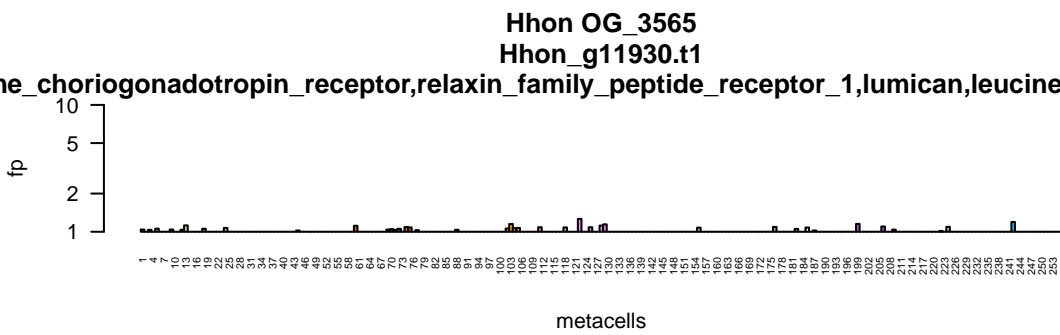
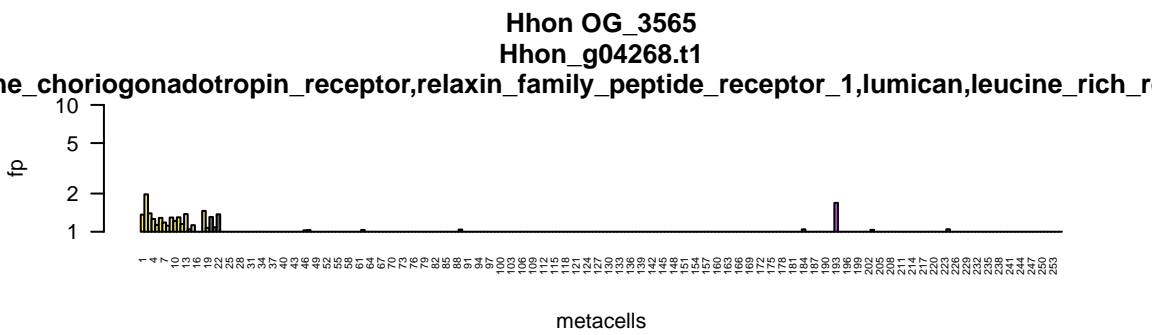
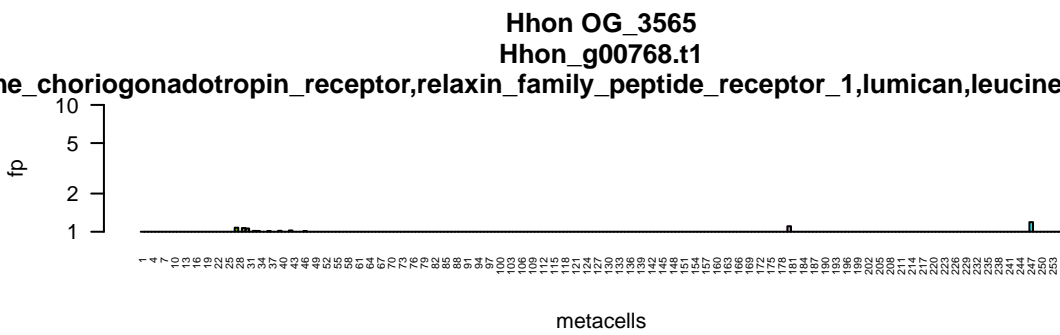
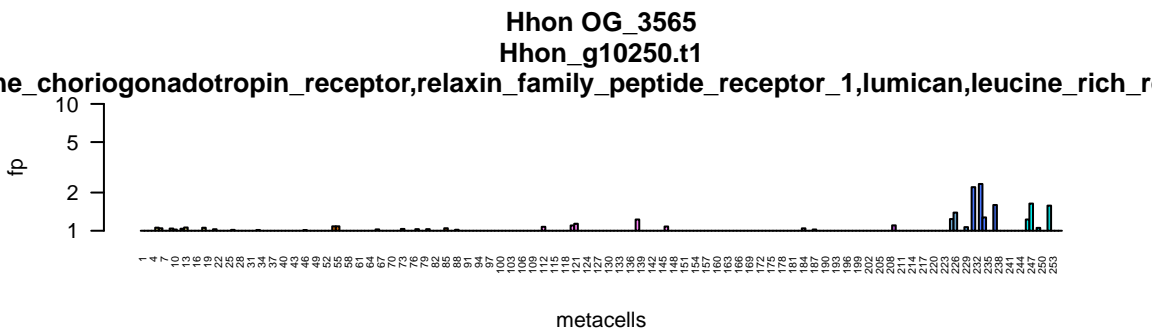
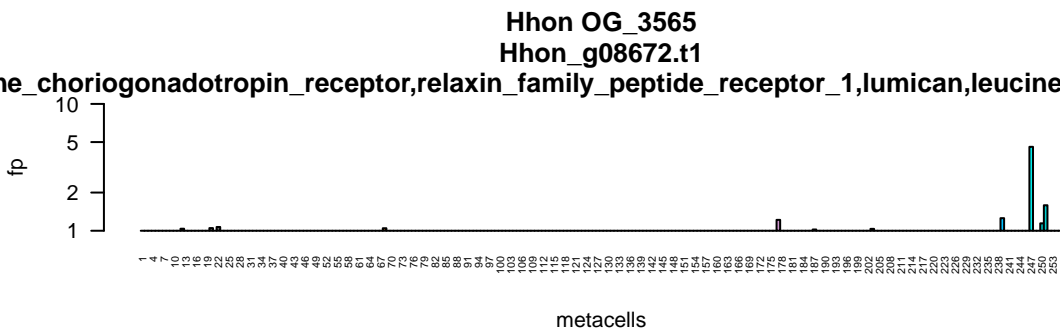
fp

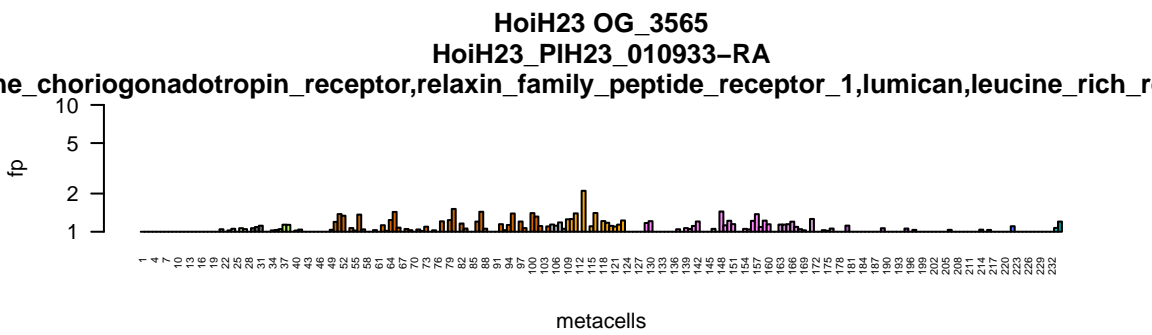
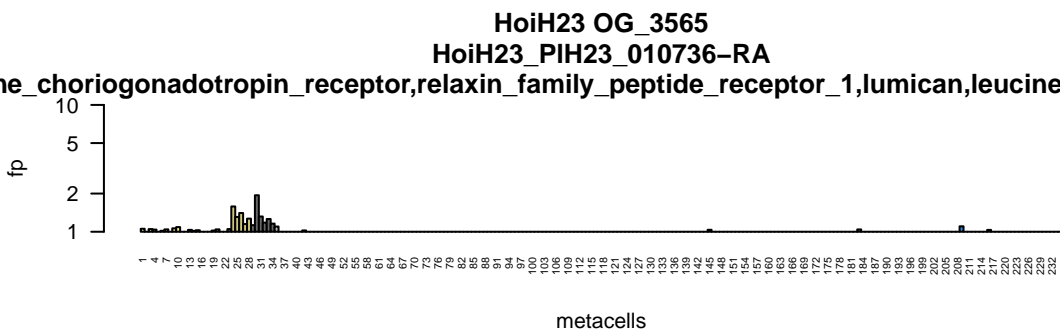
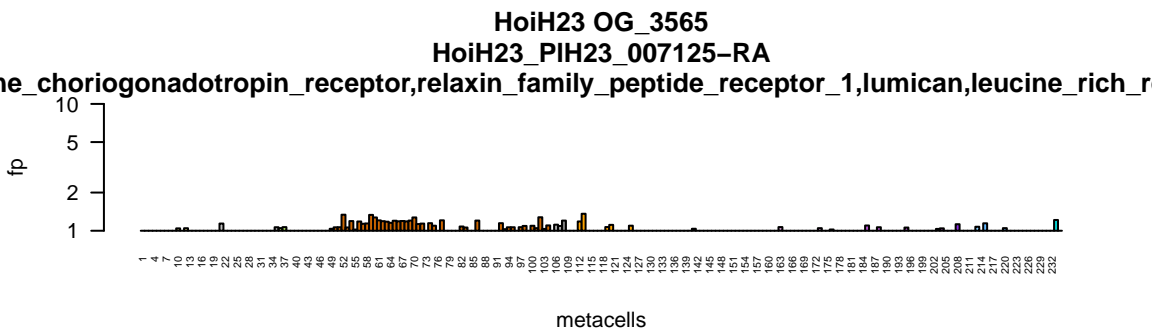
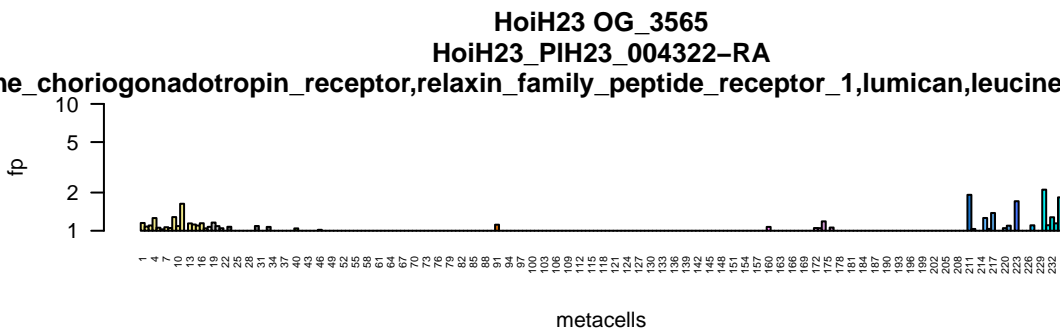
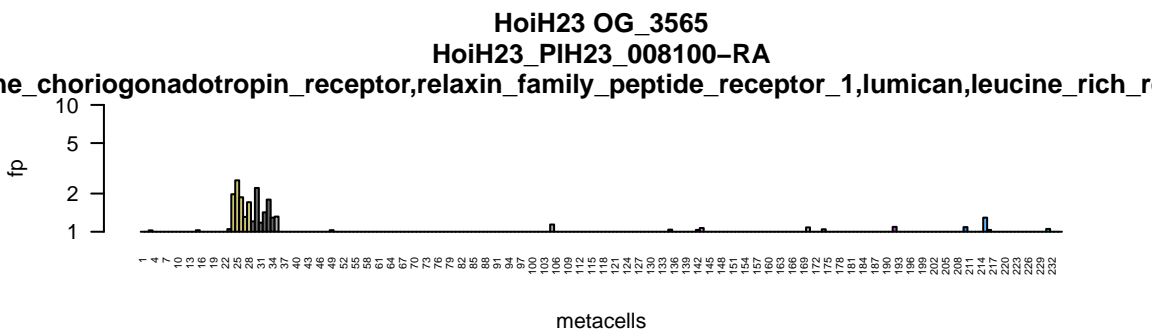
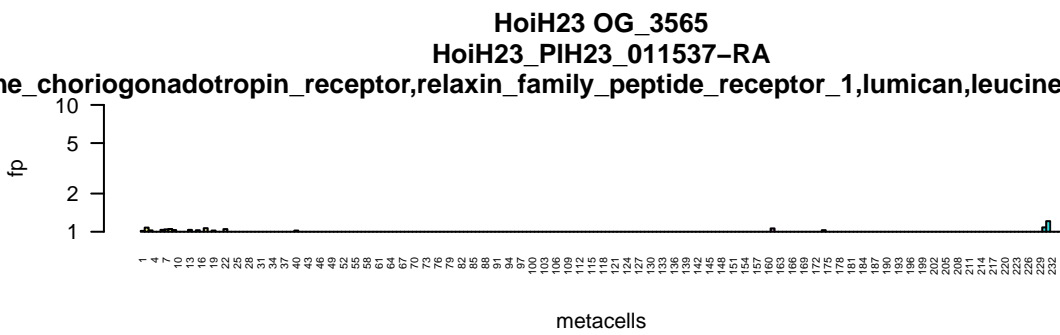
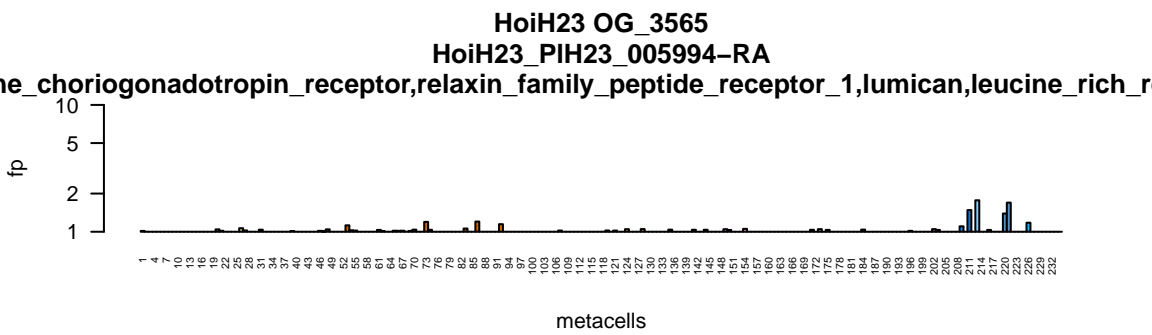
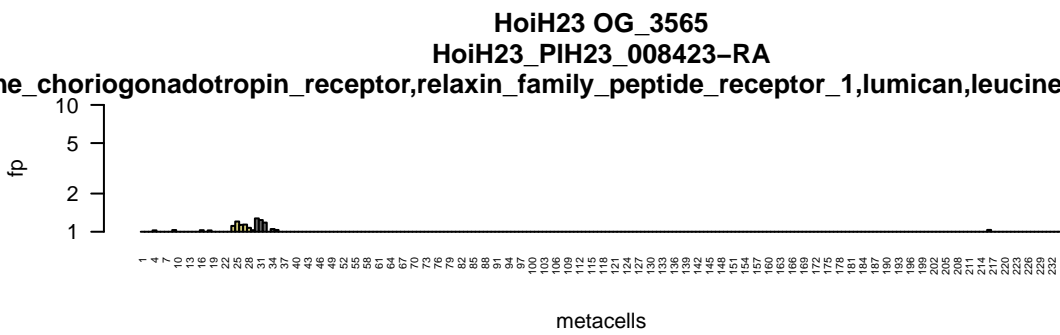
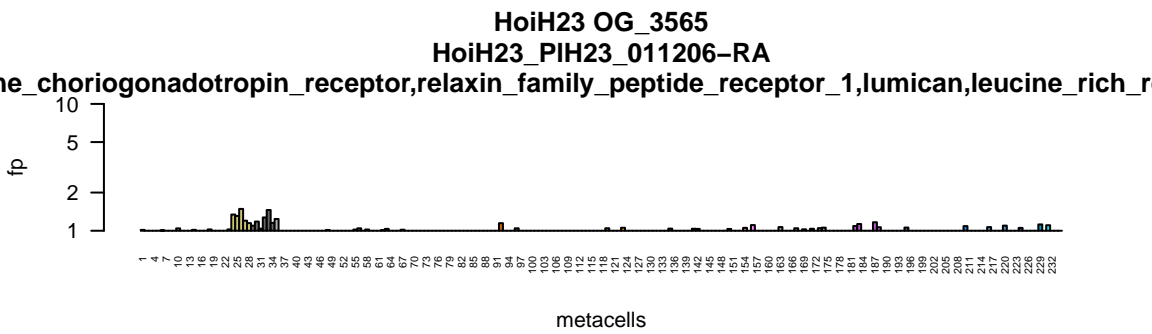
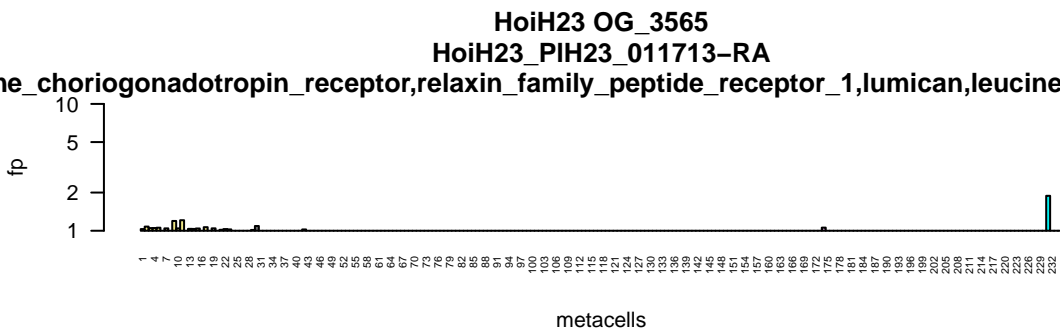
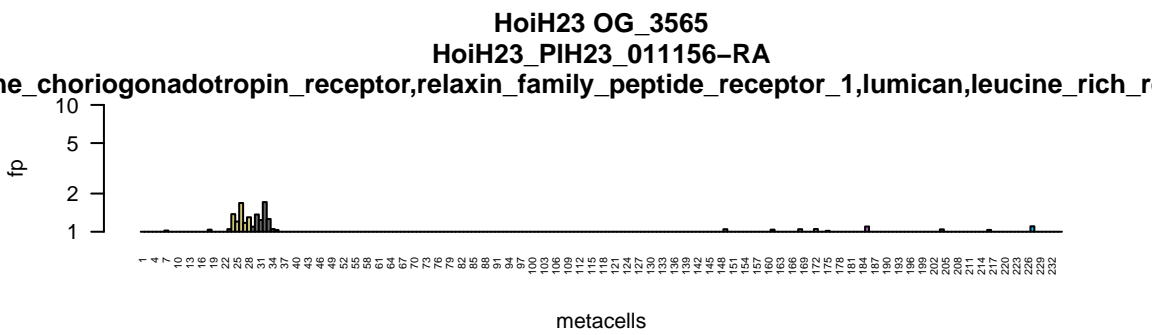
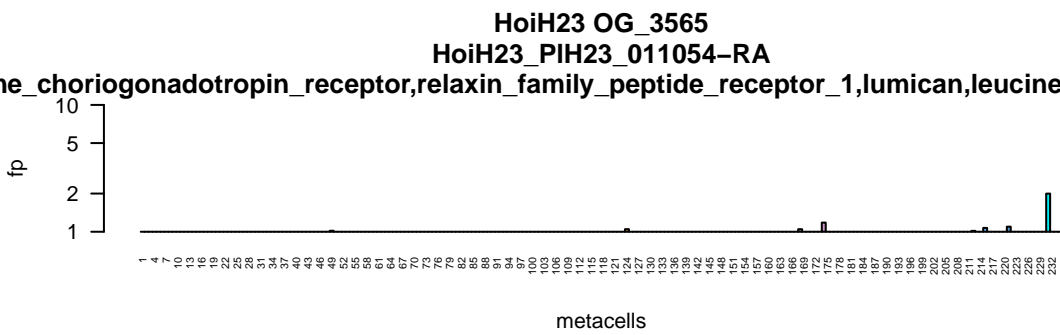
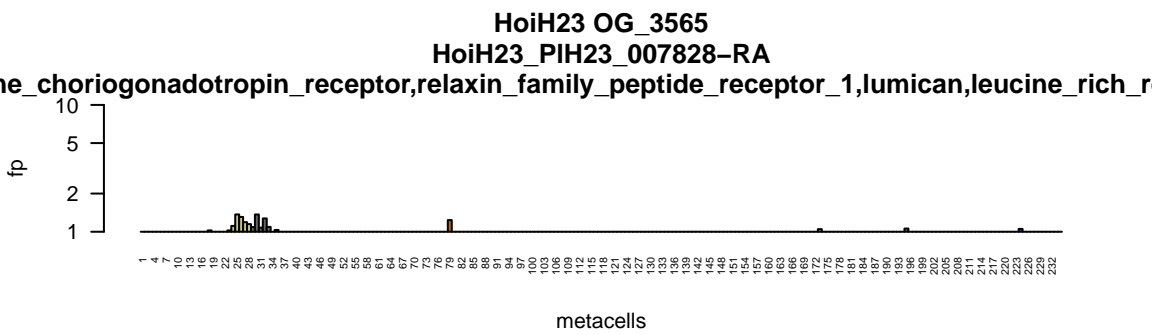
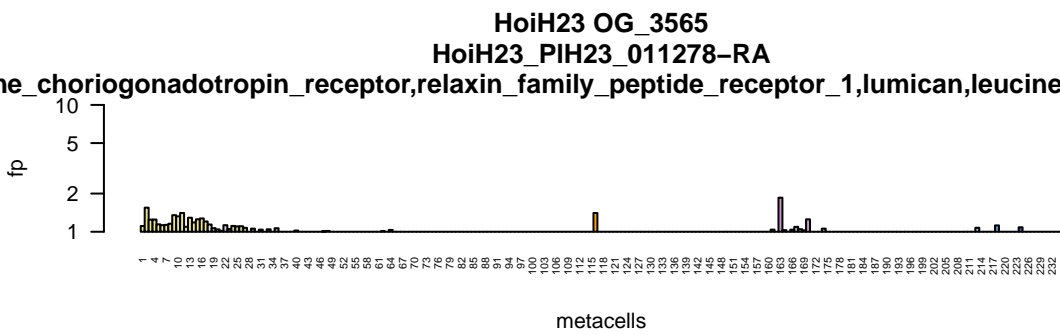
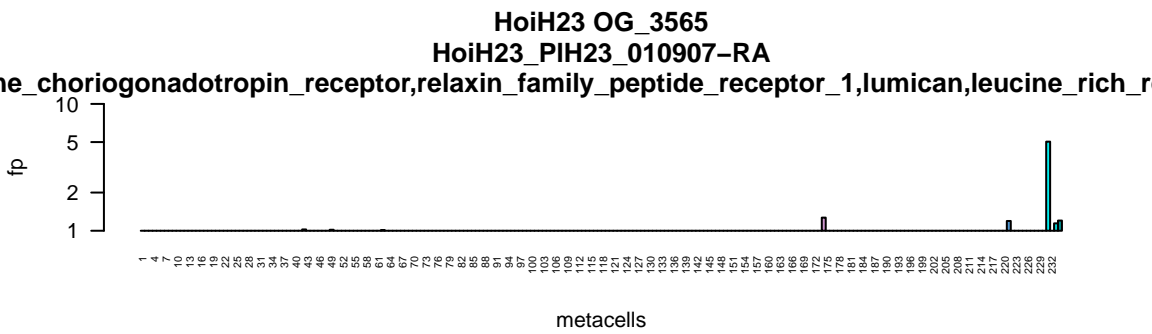
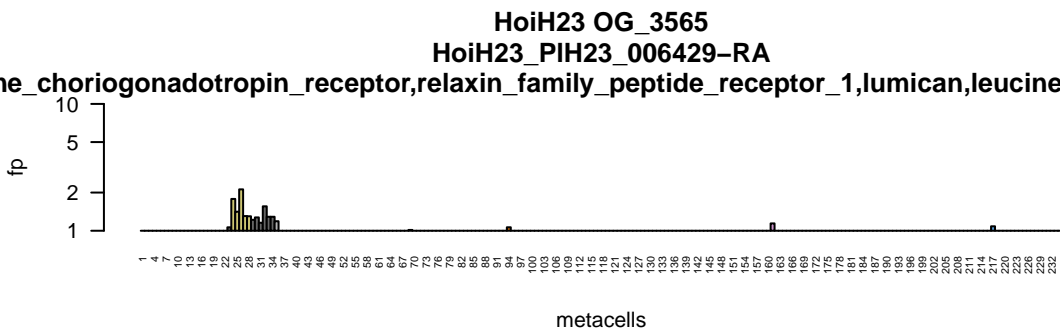
metacells

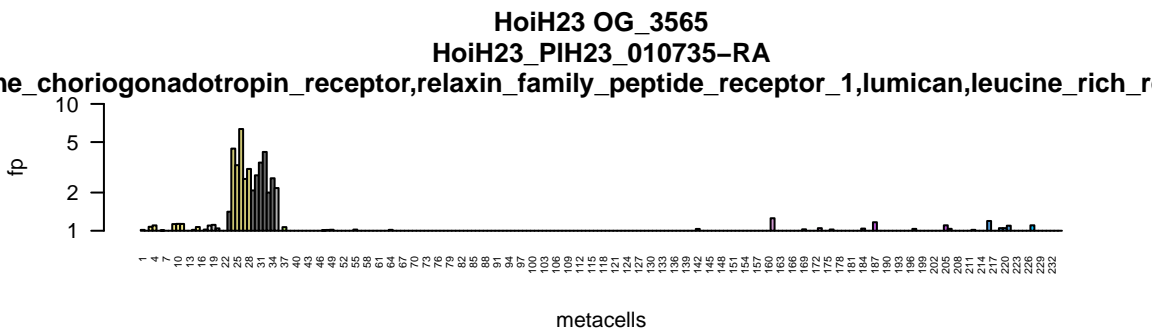
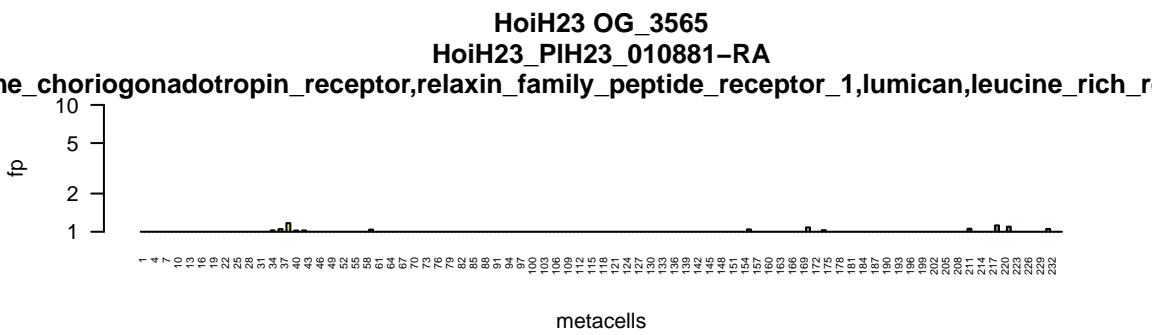
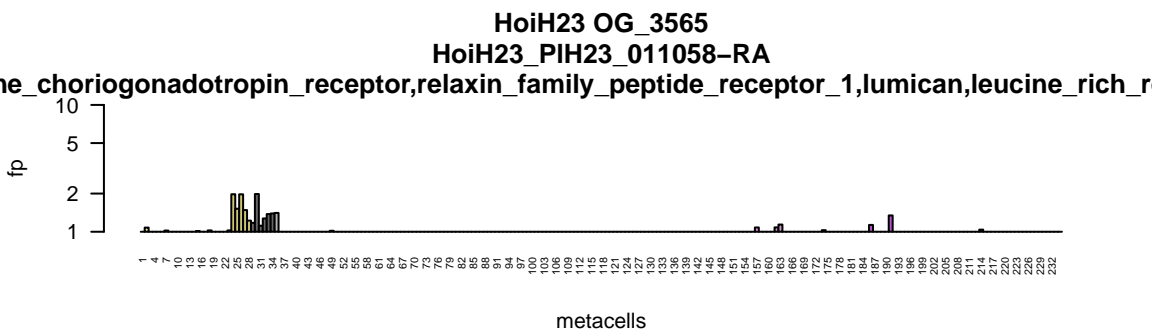
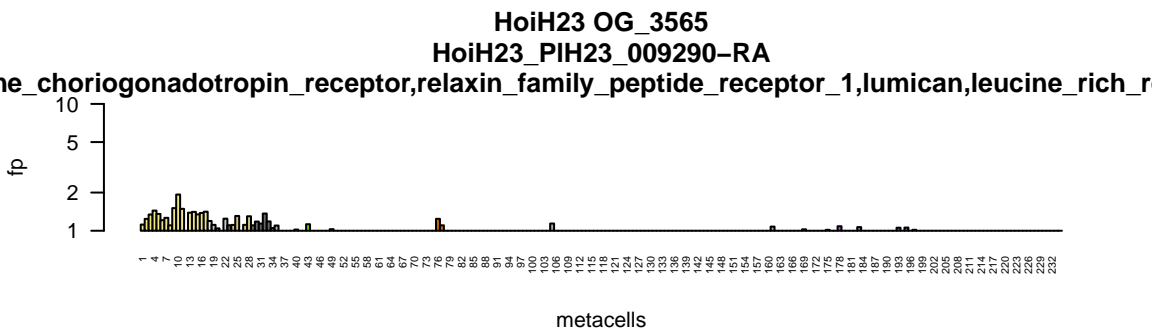


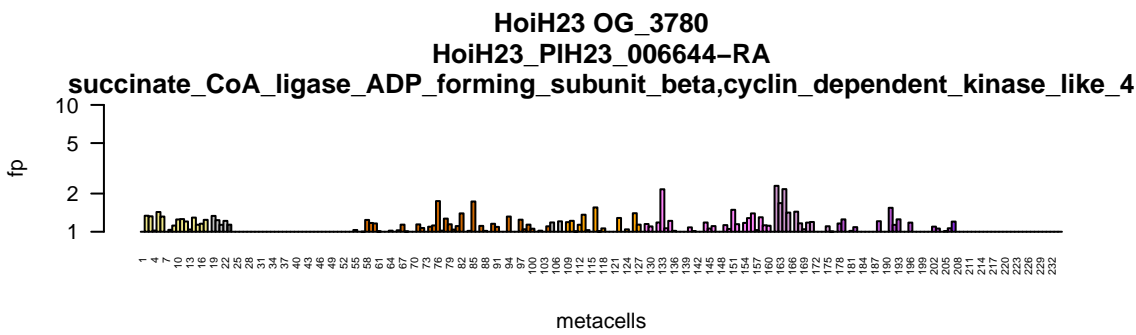
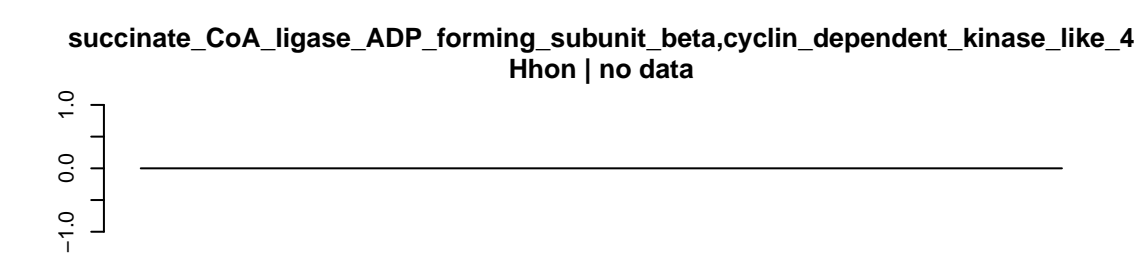
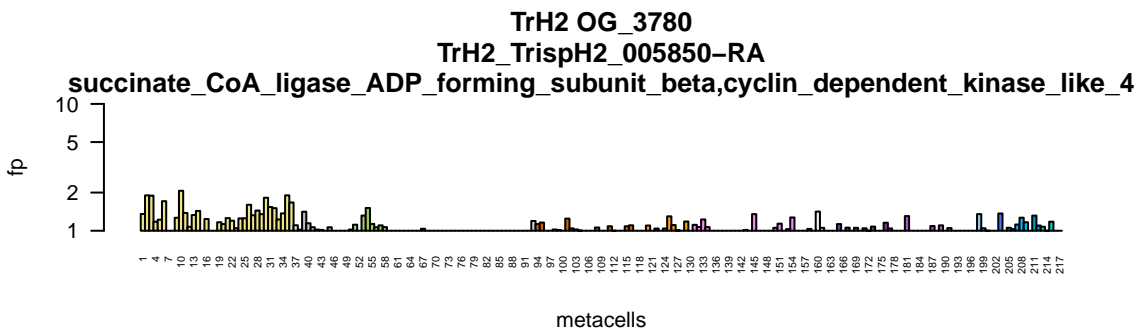
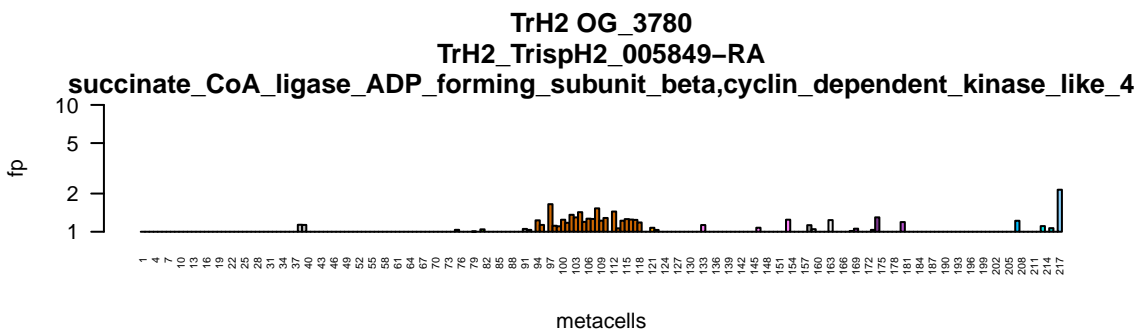
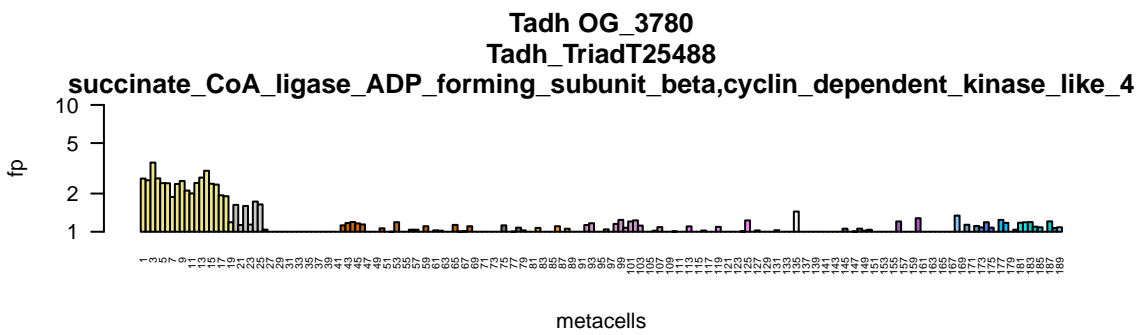
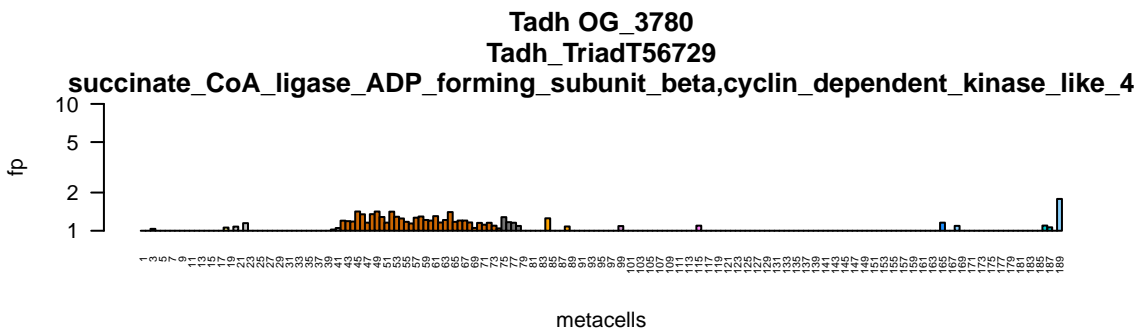








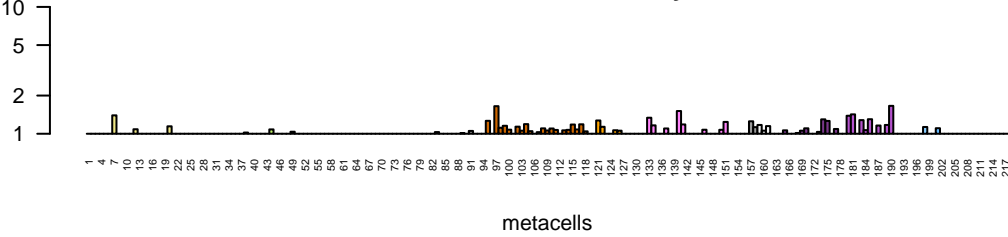




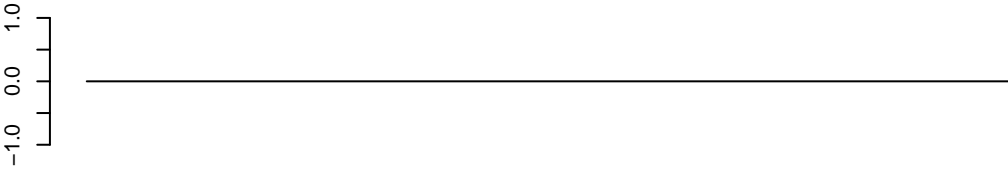
IL2_inducible_T_cell_kinase, TXK_tyrosine_kinase
Tadh | no data



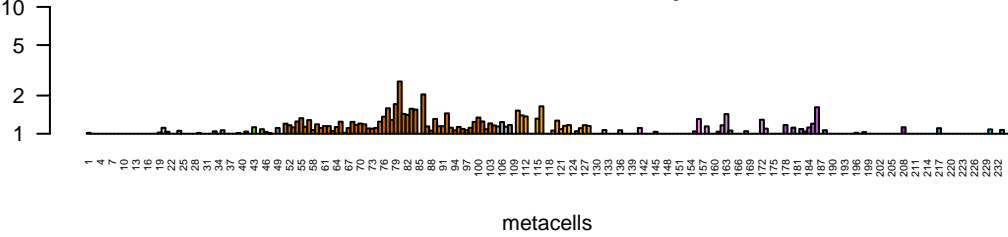
TrH2 OG_3781
TrH2_TrispH2_005851-RA
IL2_inducible_T_cell_kinase, TXK_tyrosine_kinase

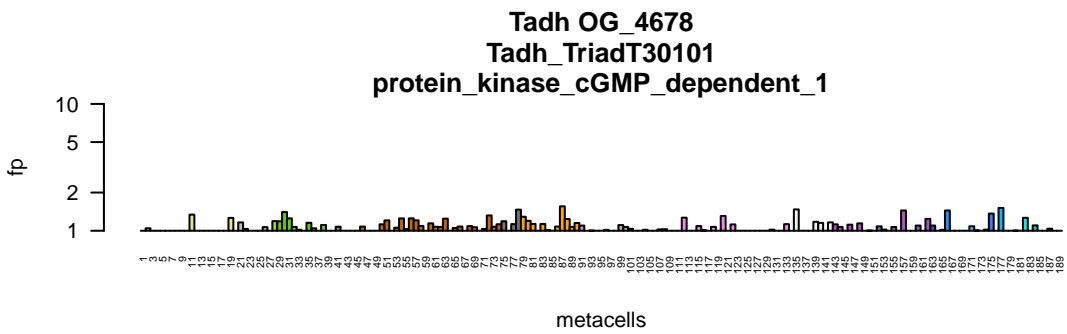


IL2_inducible_T_cell_kinase, TXK_tyrosine_kinase
Hhon | no data



HoiH23 OG_3781
HoiH23_PIH23_006645-RA
IL2_inducible_T_cell_kinase, TXK_tyrosine_kinase





TrH2 OG_4762
TrH2_TrispH2_010632-RA
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine_

fp

metacells

TrH2 OG_4762
TrH2_TrispH2_010635-RA
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

metacells	fp
1	1
7	1
13	1
16	1
19	1
25	1
28	1
31	1
34	1
37	1
40	1.2
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	1.1
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	1.2
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	1.8
202	1
205	1.1
208	1
211	1
214	1.8
217	1

TrH2 OG_4762
TrH2_TrispH2_010271-RA
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

fp

metacells

Hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

metacells

Hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

fp

metacells

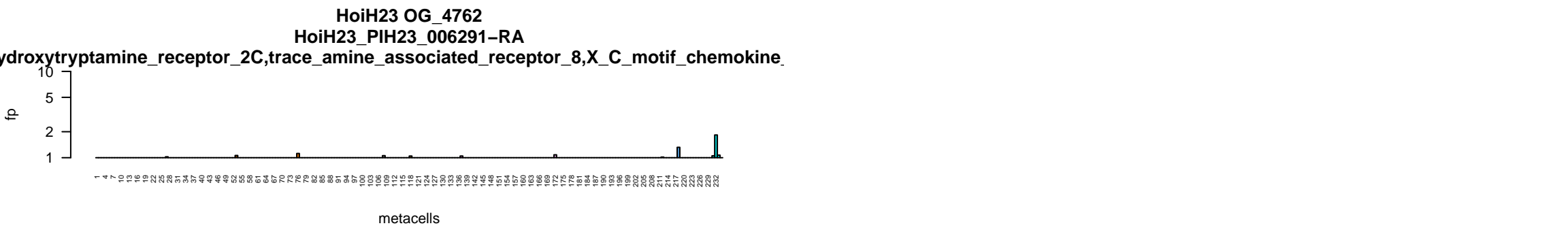
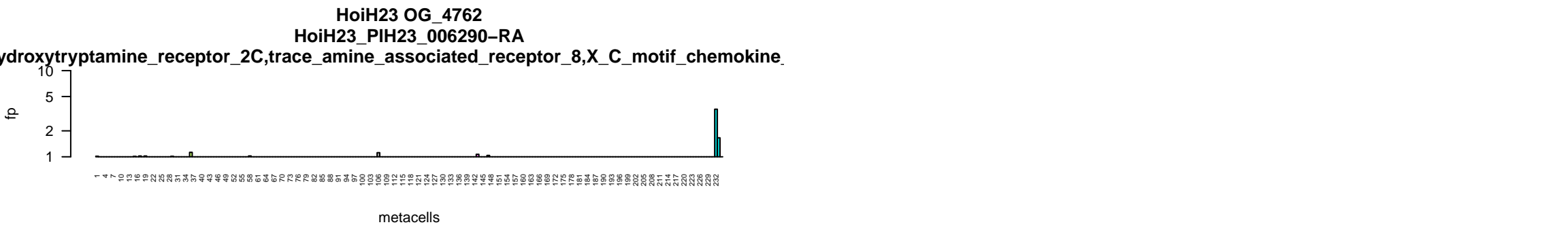
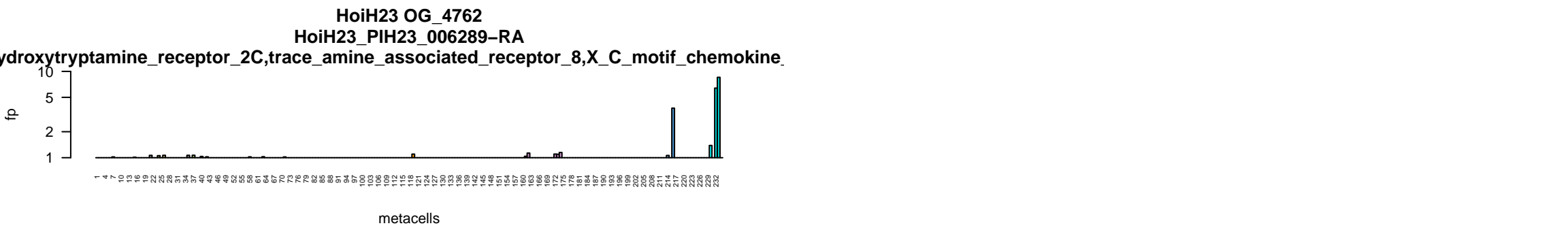
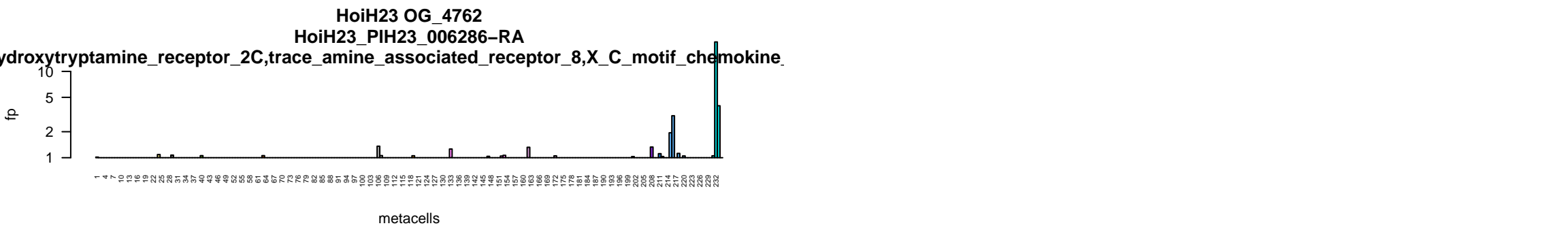
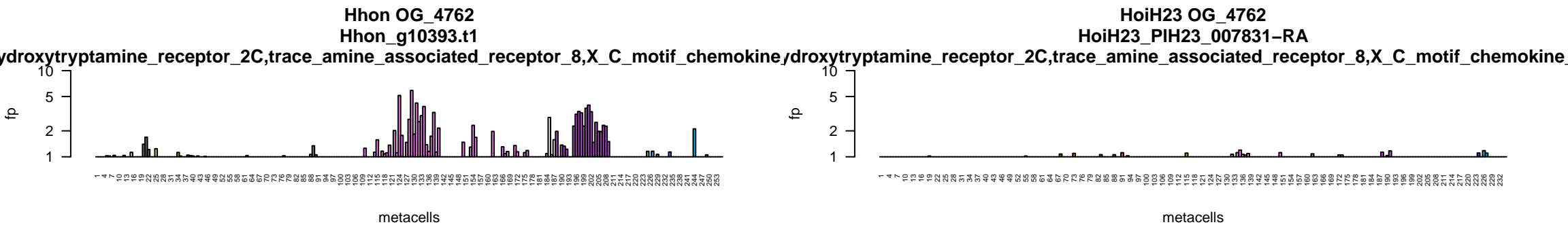
Hhox OG_4762
Hhox_g01703.t1

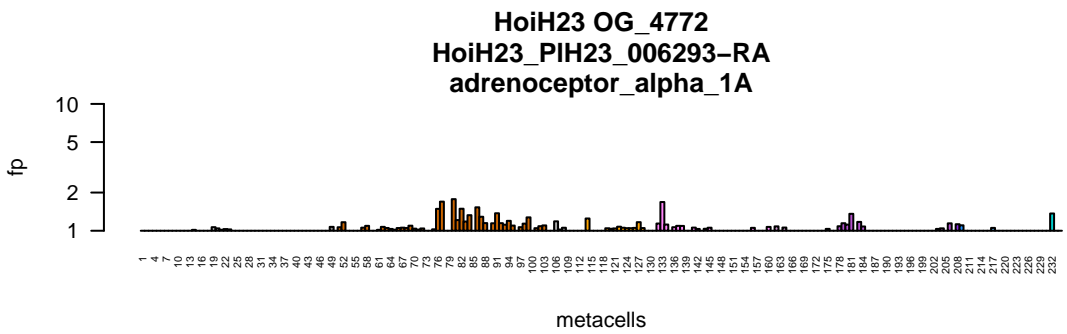
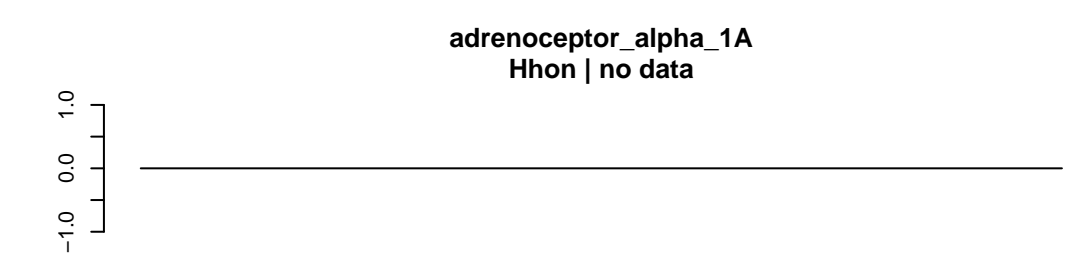
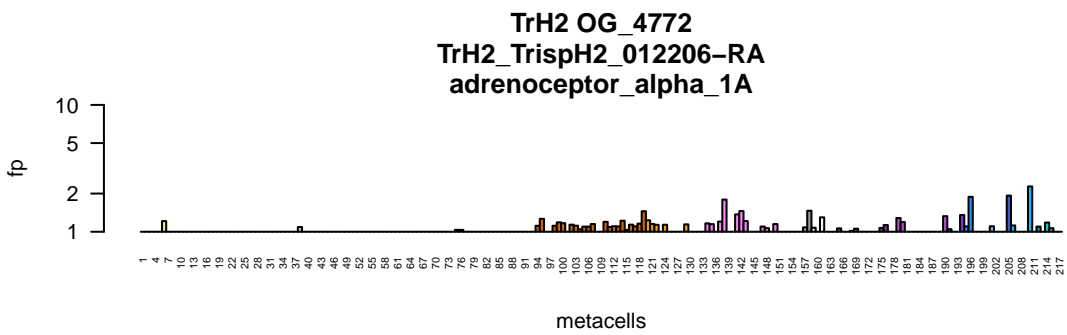
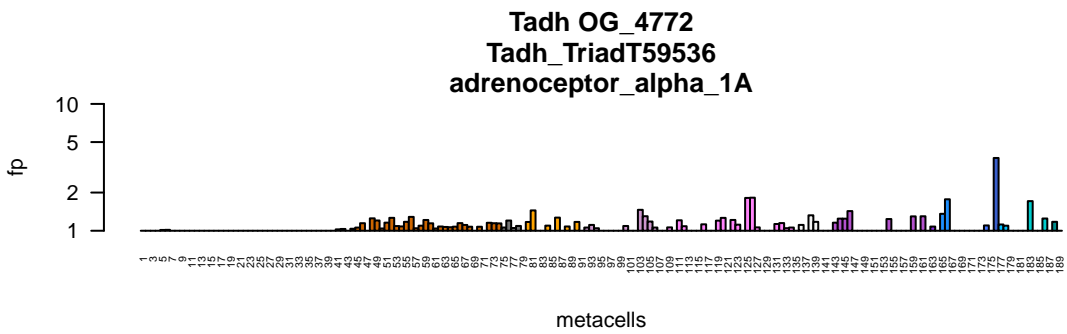
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

fp

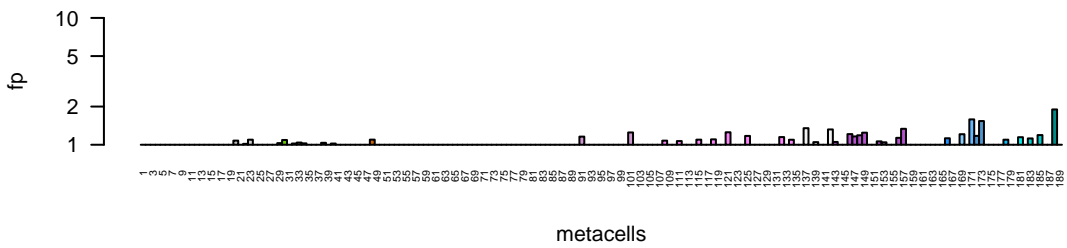
metacells

metacell	fp
1	1
10	1
13	1
19	1
22	1
28	1
31	1
37	1
43	1
49	1
52	1
55	1
61	1
64	1
70	1
73	1
79	1
88	1
91	1
97	1
100	1
109	1
108	1
115	1
119	1
127	1
124	1
133	1
136	1
142	1
146	1
151	1
154	1
160	1
163	1
169	1
175	1
178	1
184	1
187	1
193	1
196	1
202	1
205	1
211	1
214	1
220	2
223	1
229	1
232	1
238	1
241	1
247	1
250	2
253	2

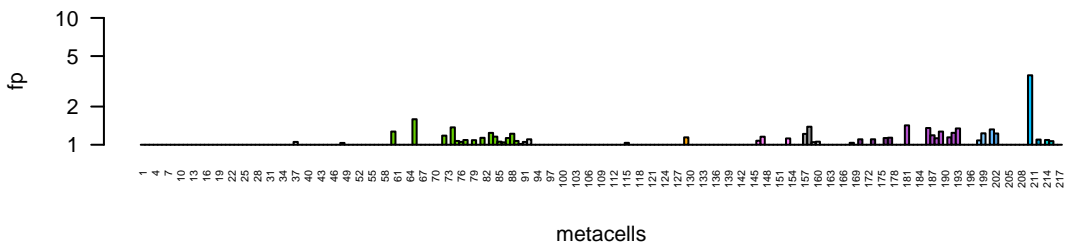




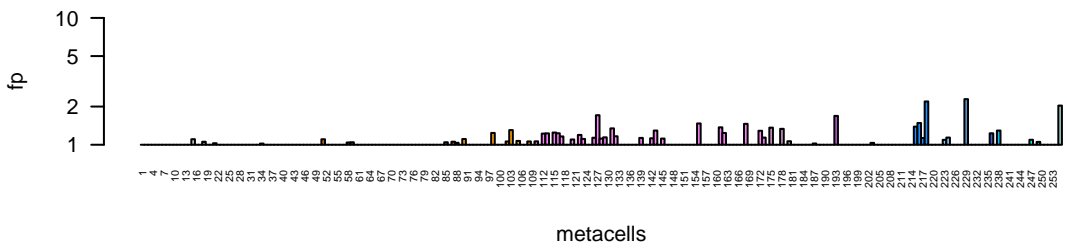
Tadh OG_5125
Tadh_wf_g10501.t1



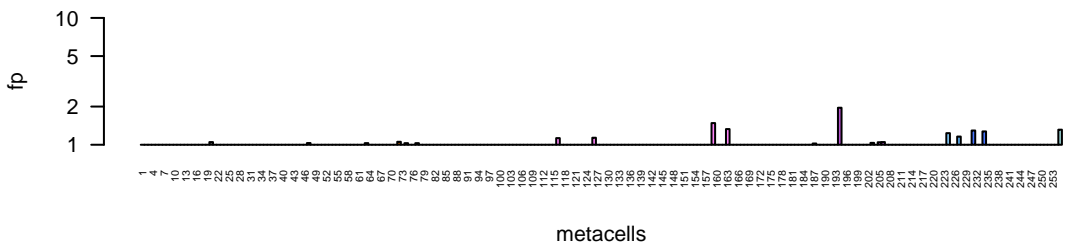
TrH2 OG_5125
TrH2_TrispH2_003544-RA



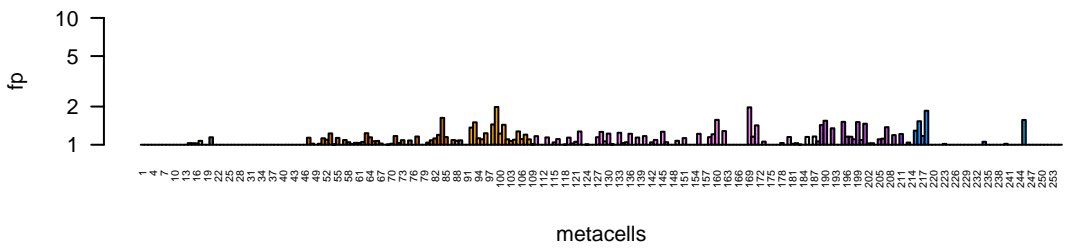
Hhon OG_5125
Hhon_g07867.t1



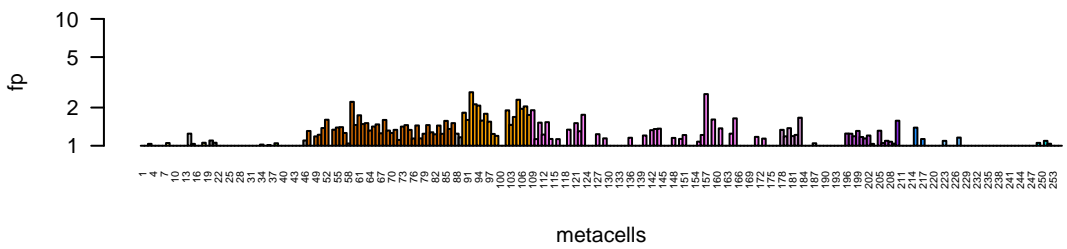
Hhon OG_5125
Hhon_g07868.t1



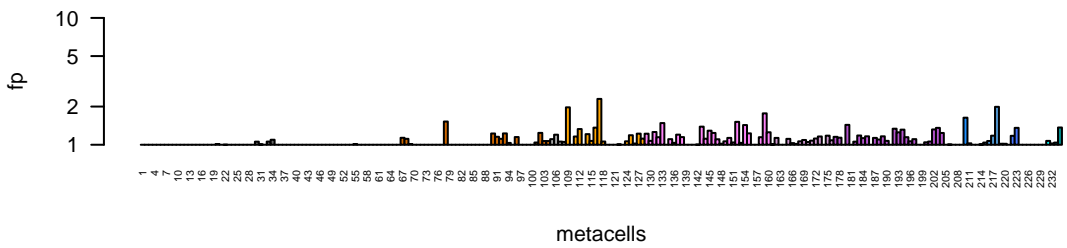
Hhon OG_5125
Hhon_g07869.t1

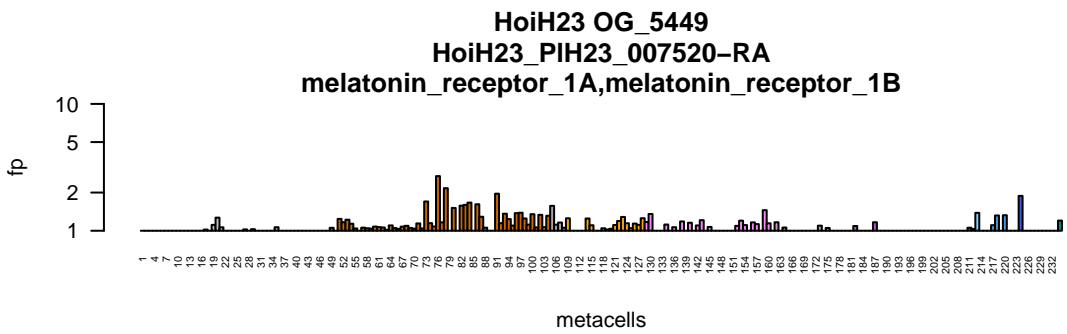
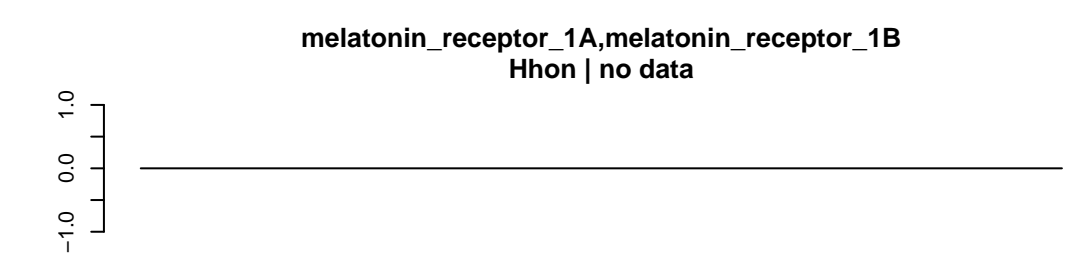
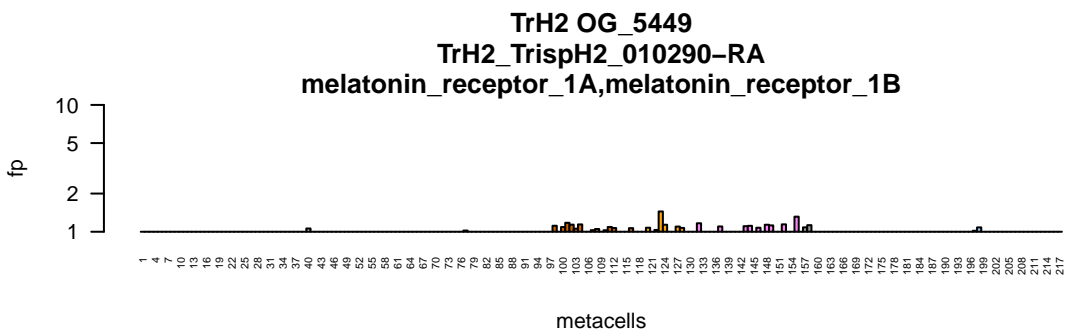
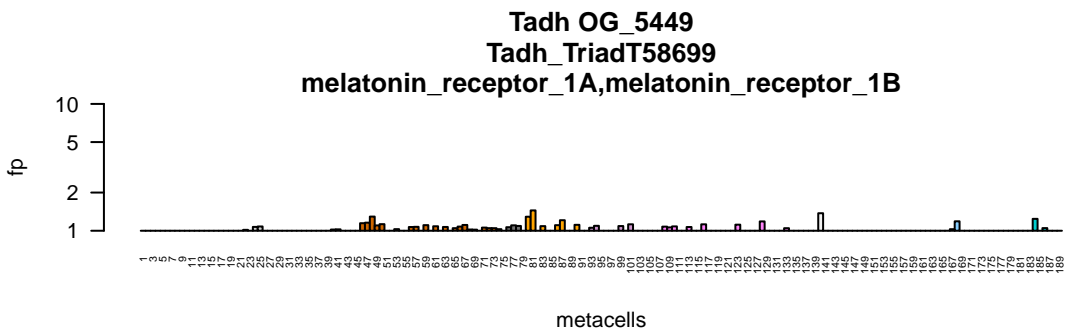


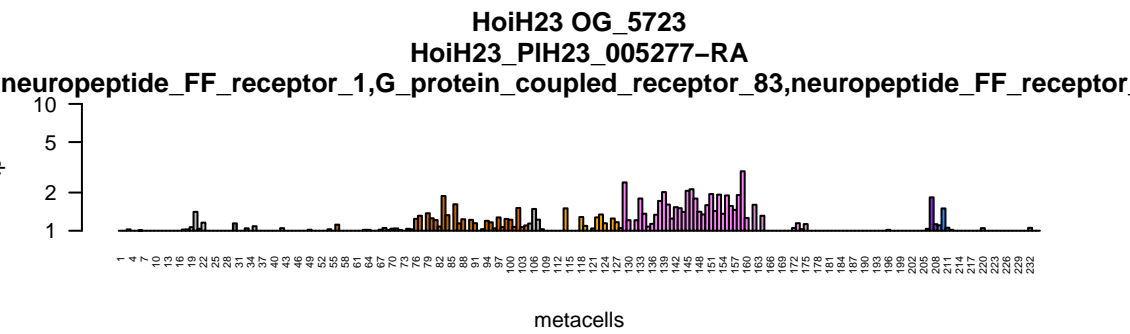
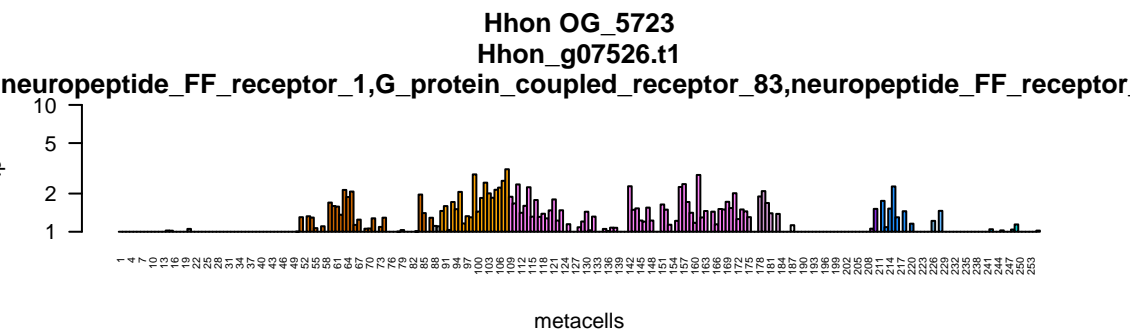
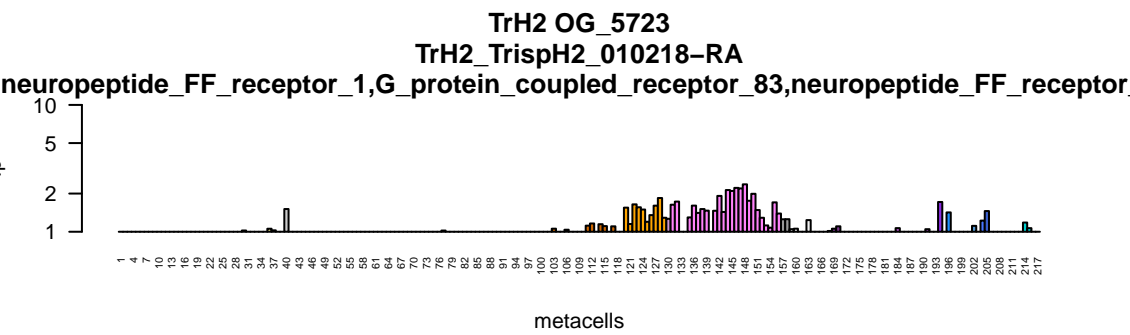
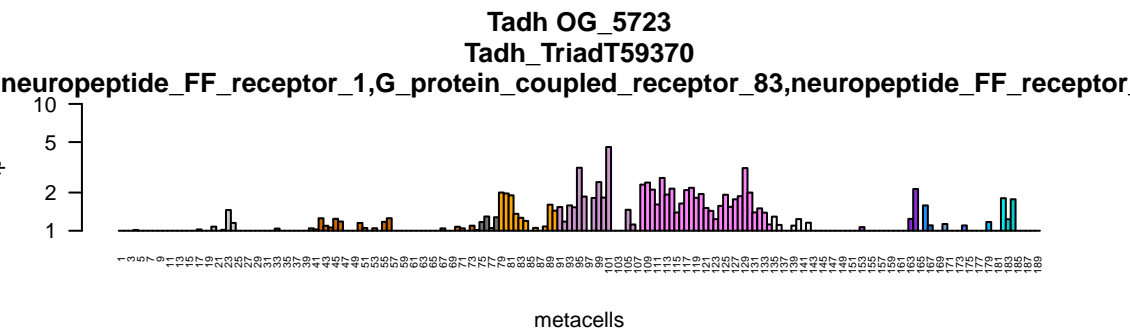
Hhon OG_5125
Hhon_g07870.t1

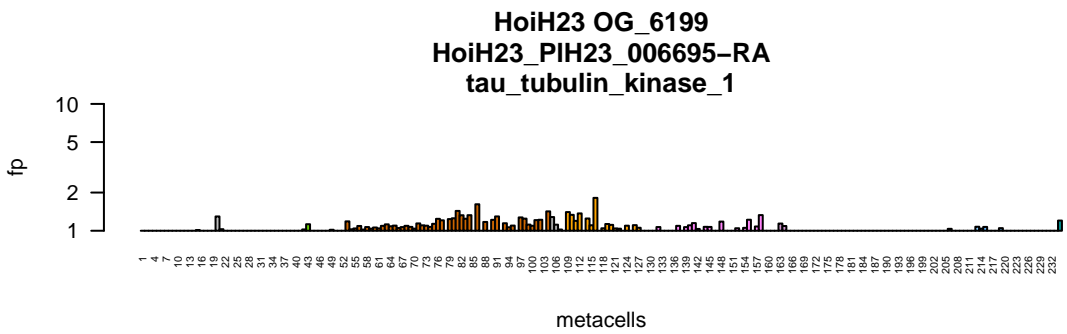
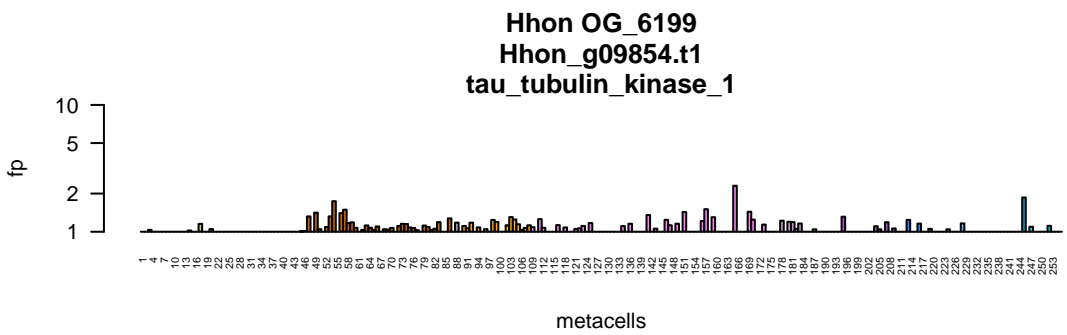
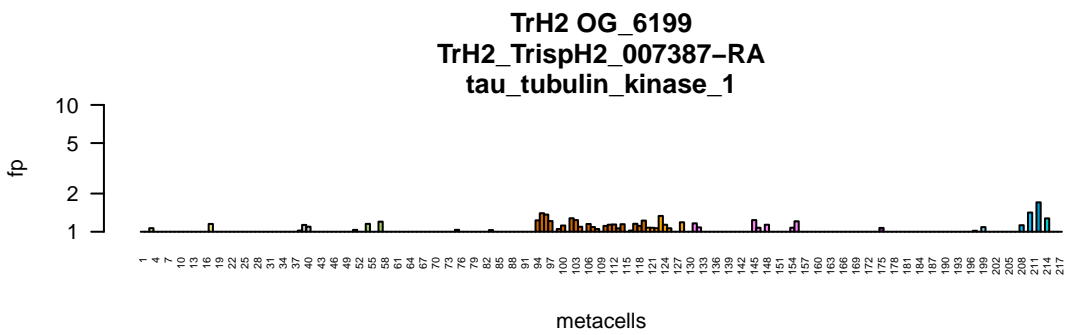
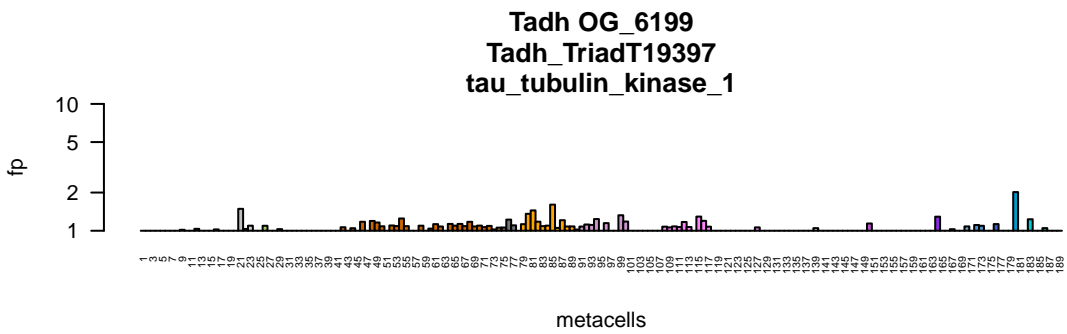


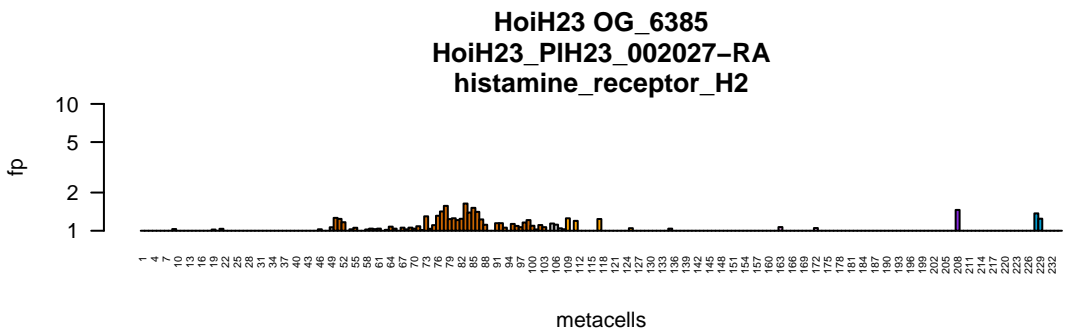
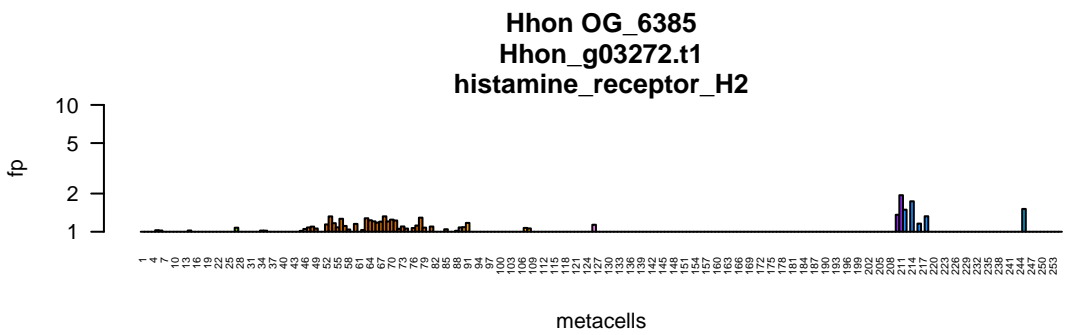
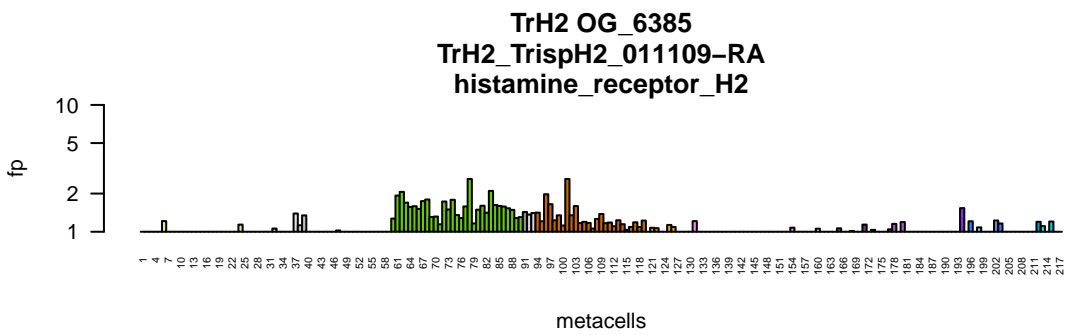
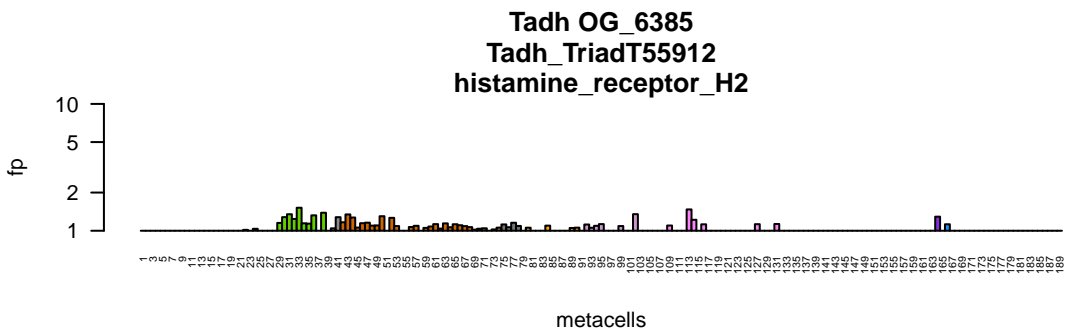
HoiH23 OG_5125
HoiH23_PIH23_007715-RA

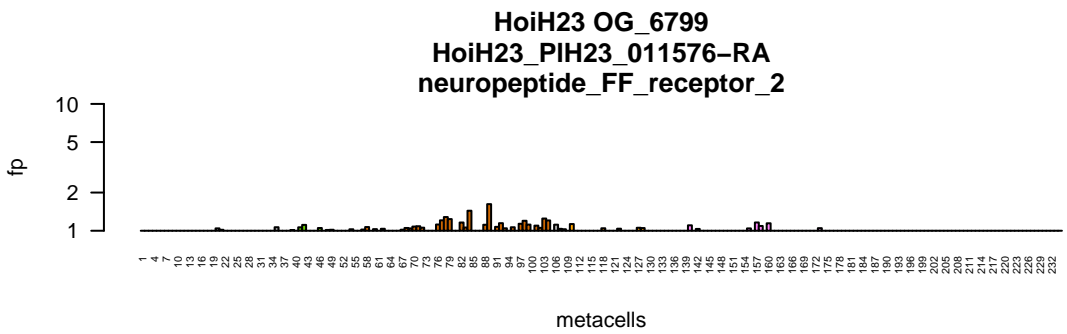
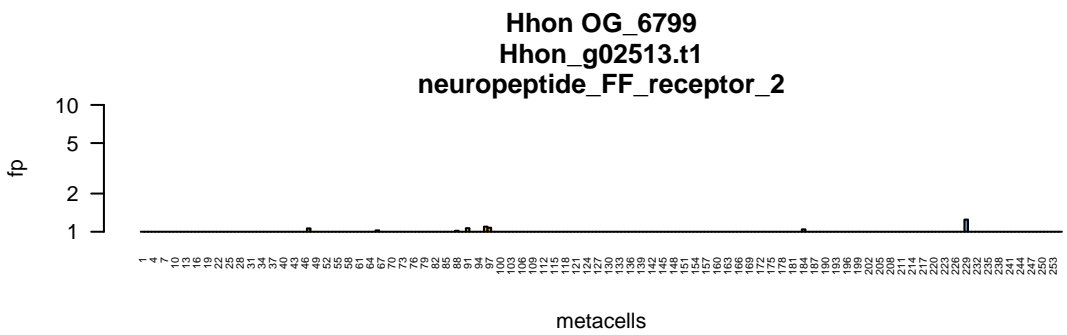
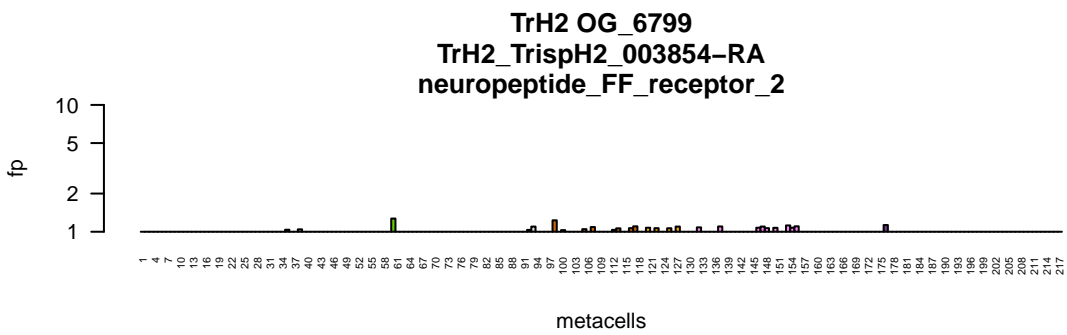
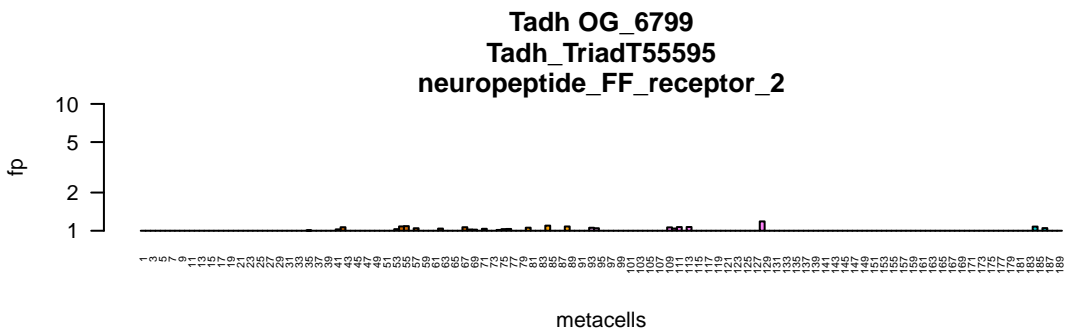








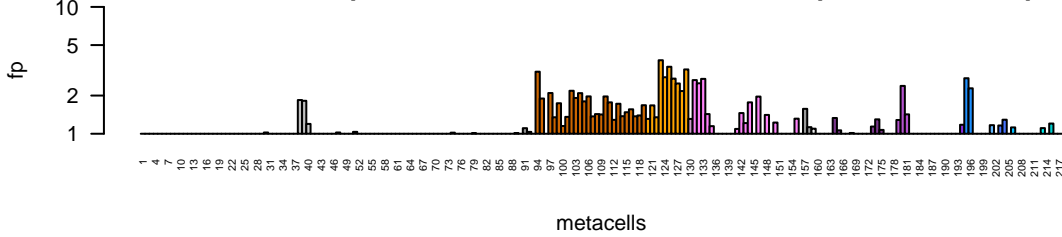




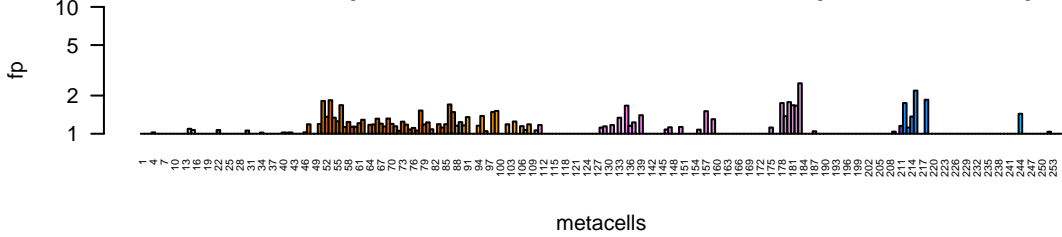
C_C_motif_chemokine_receptor_1,trace_amine_associated_receptor_8,adrenoceptor_alpha2
Tadh | no data



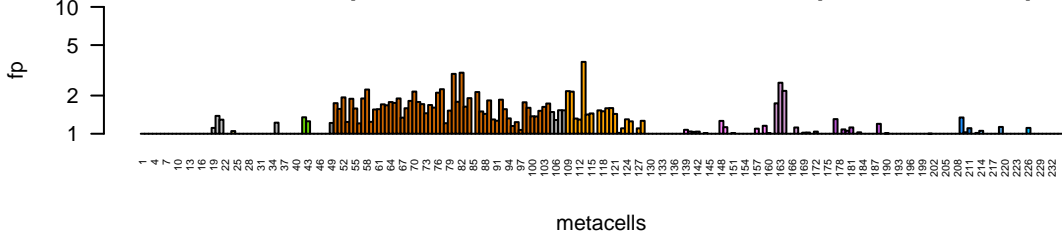
TrH2 OG_7015
TrH2_TrispH2_003193-RA
C_C_motif_chemokine_receptor_1,trace_amine_associated_receptor_8,adrenoceptor_alpha2

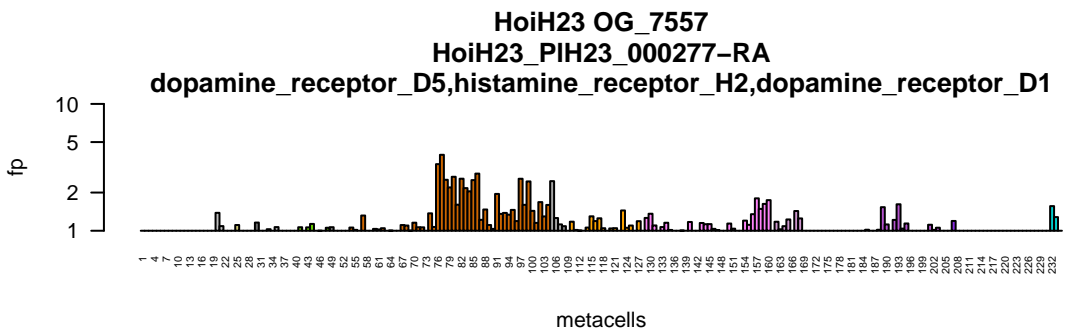
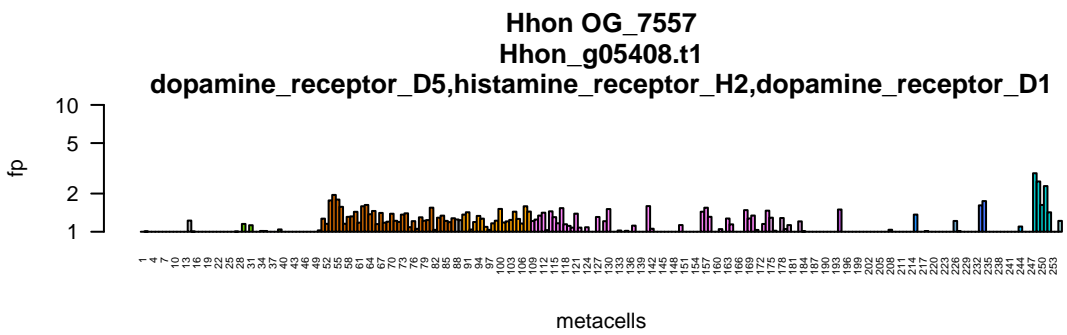
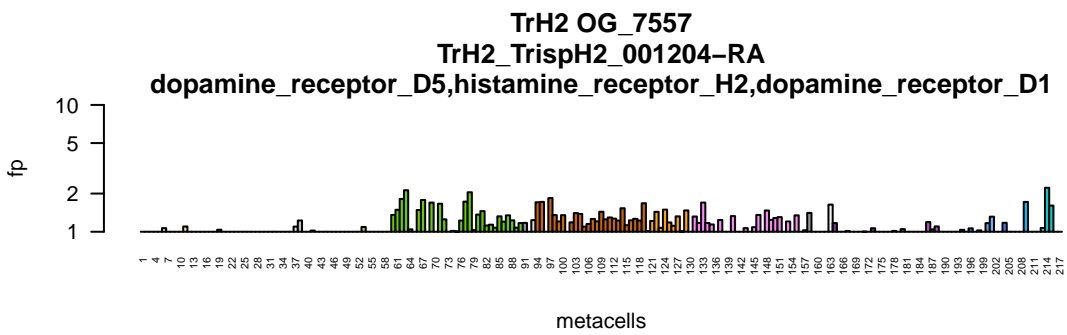
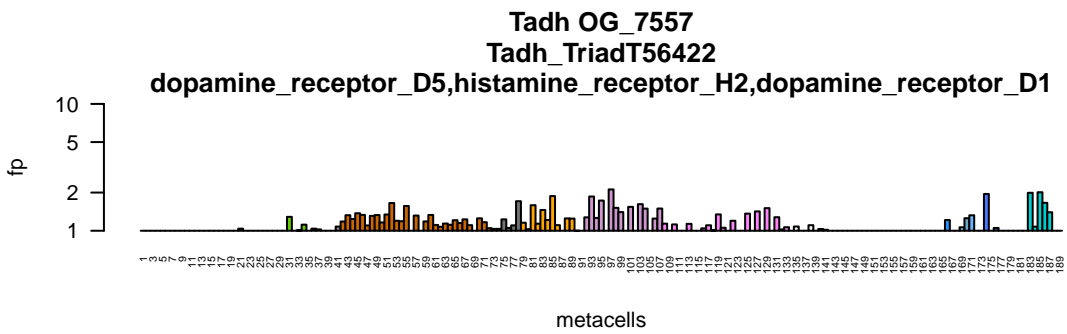


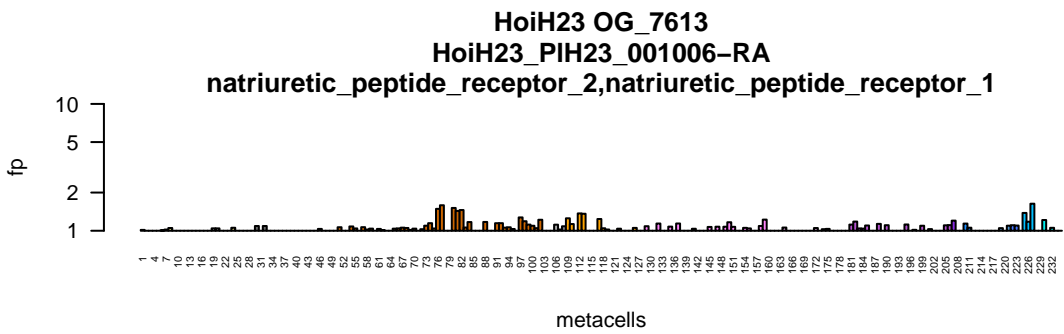
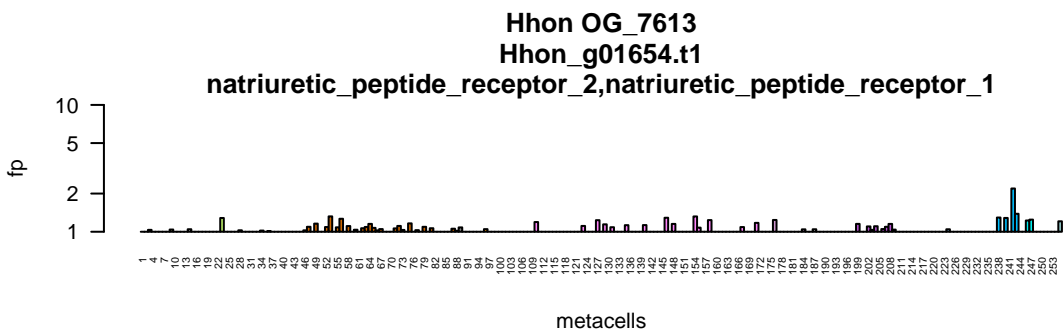
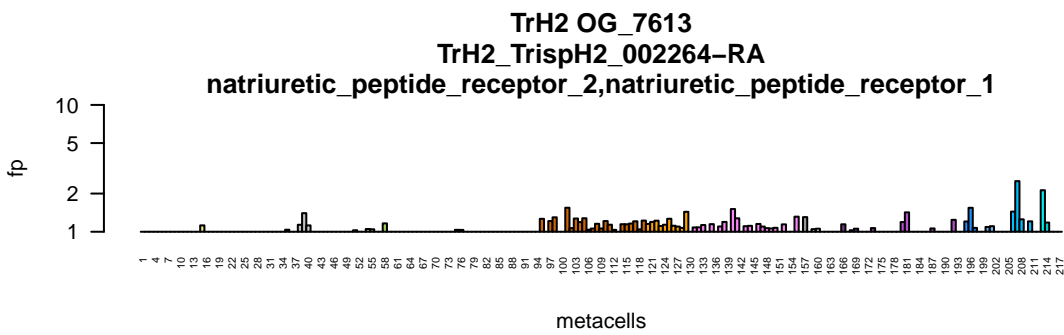
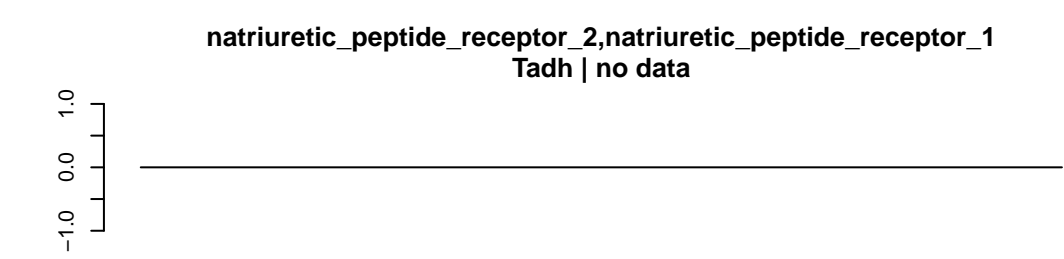
Hhon OG_7015
Hhon_g03672.t1
C_C_motif_chemokine_receptor_1,trace_amine_associated_receptor_8,adrenoceptor_alpha2

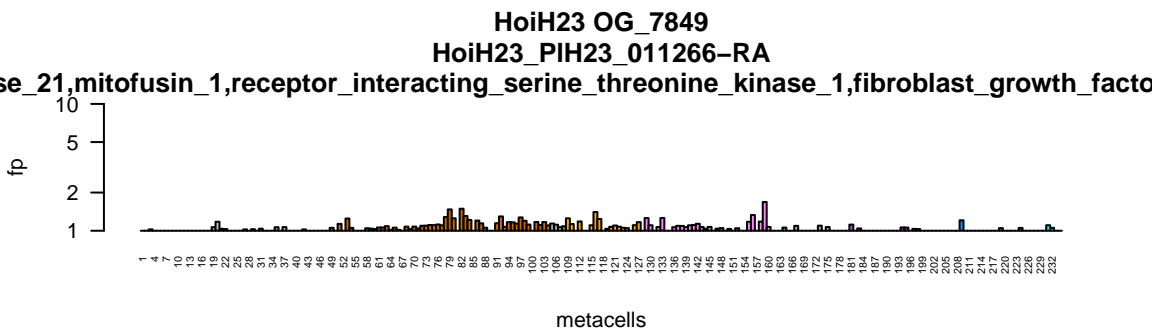
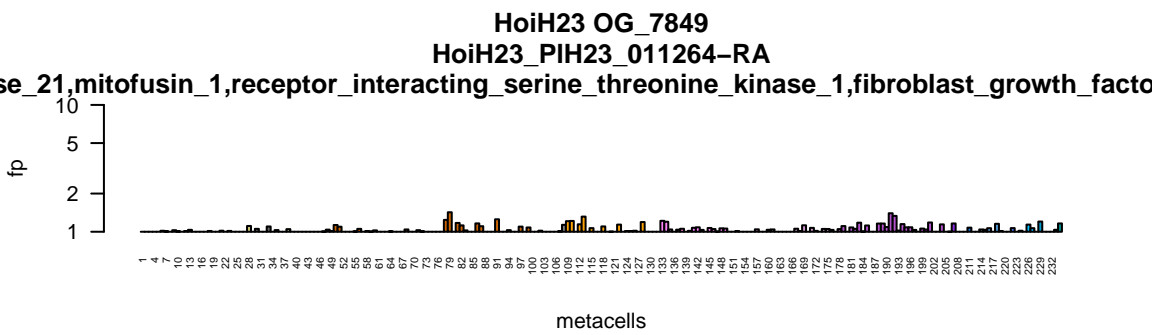
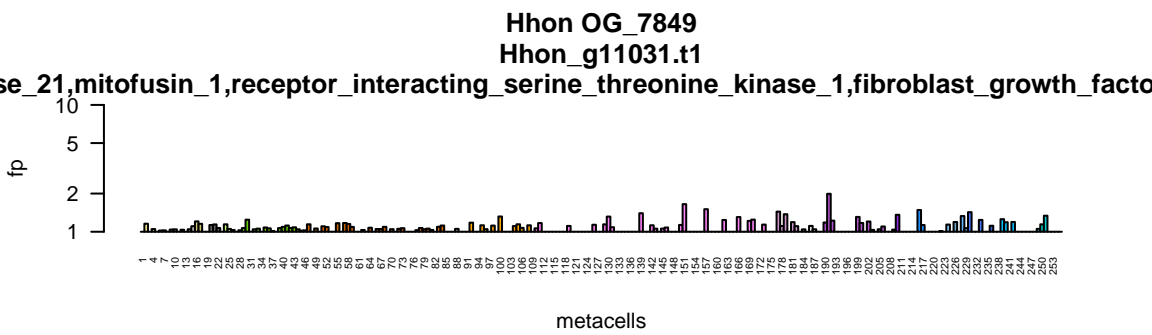
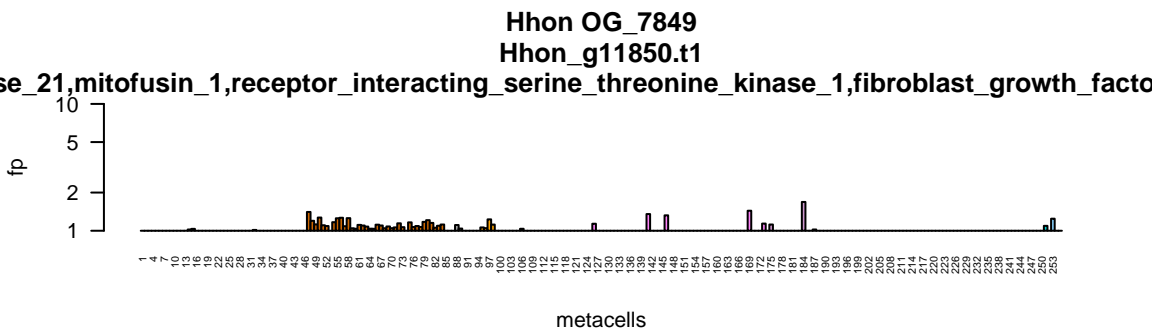
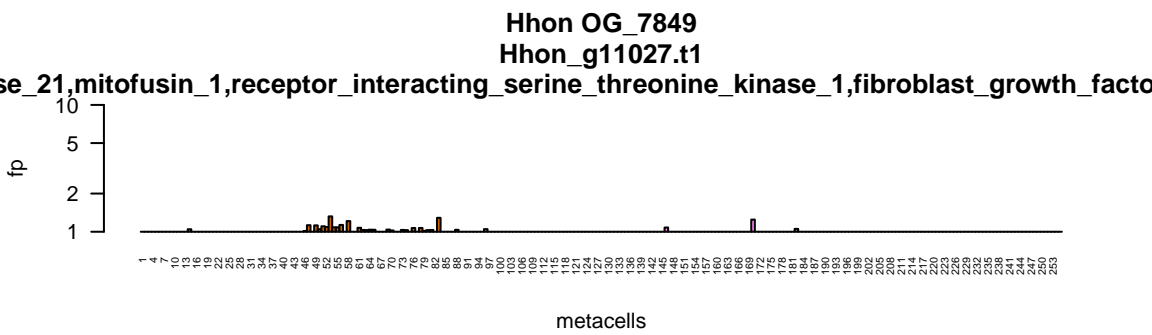
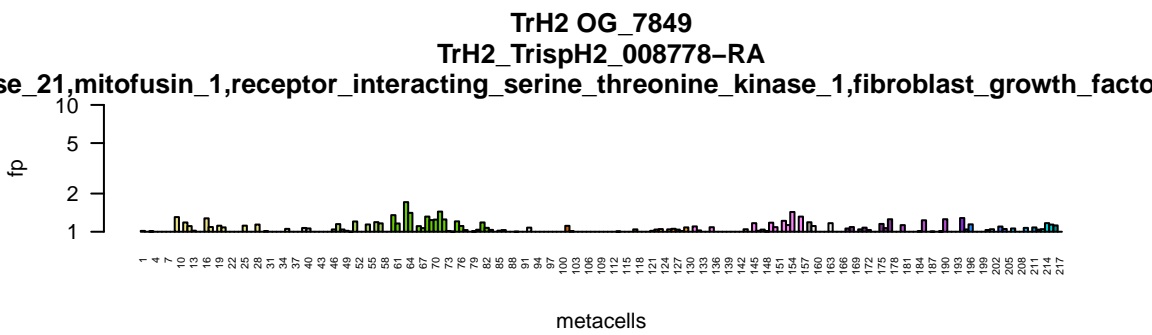
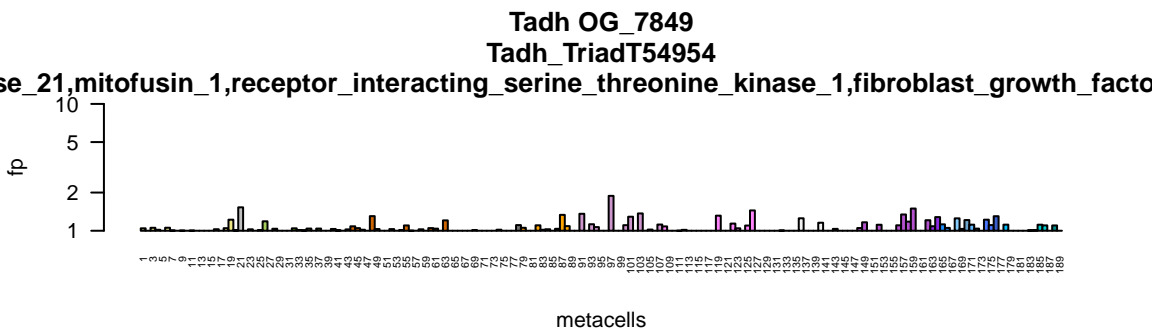


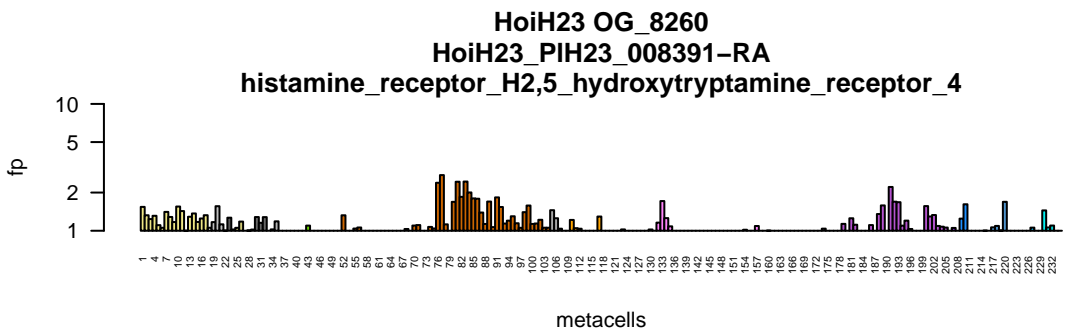
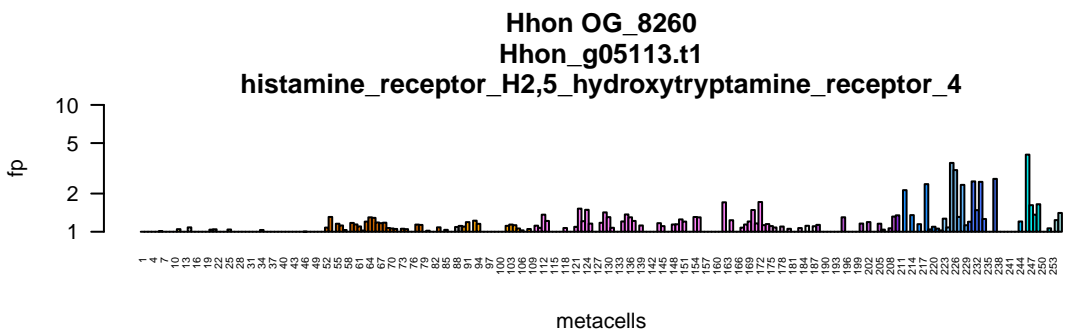
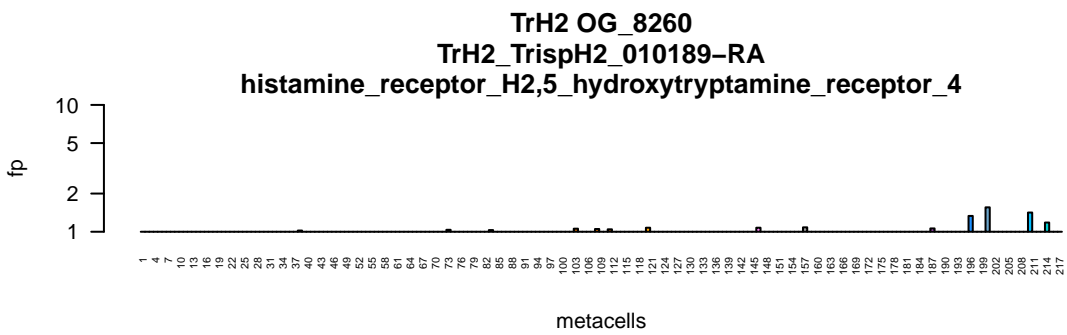
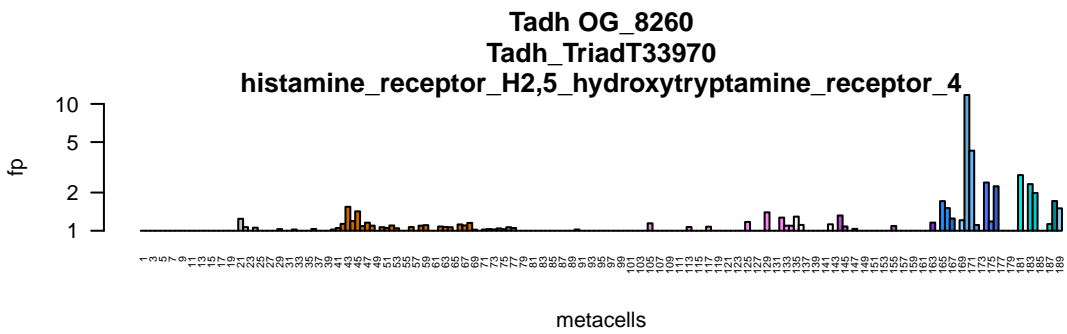
HoiH23 OG_7015
HoiH23_PIH23_002074-RA
C_C_motif_chemokine_receptor_1,trace_amine_associated_receptor_8,adrenoceptor_alpha2

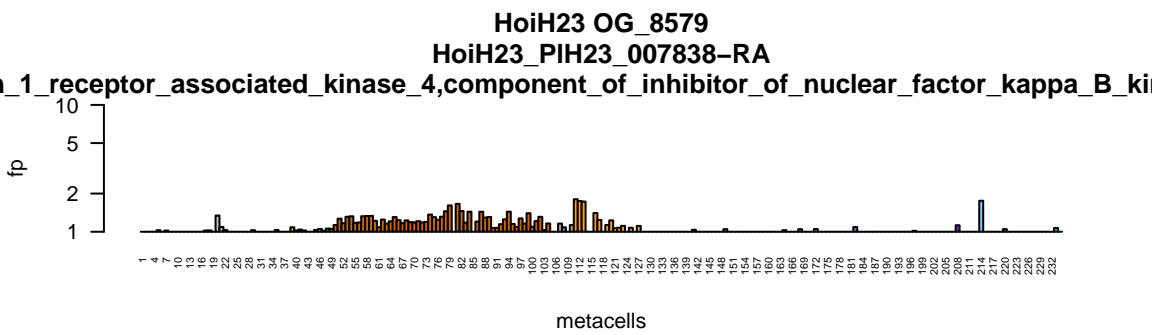
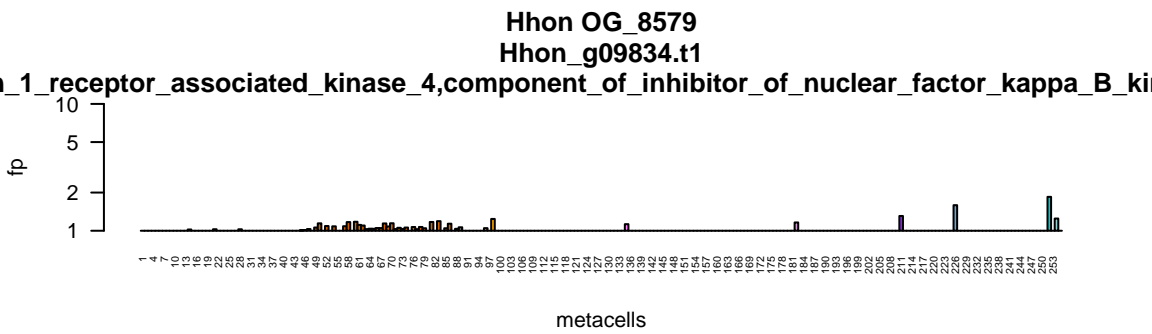
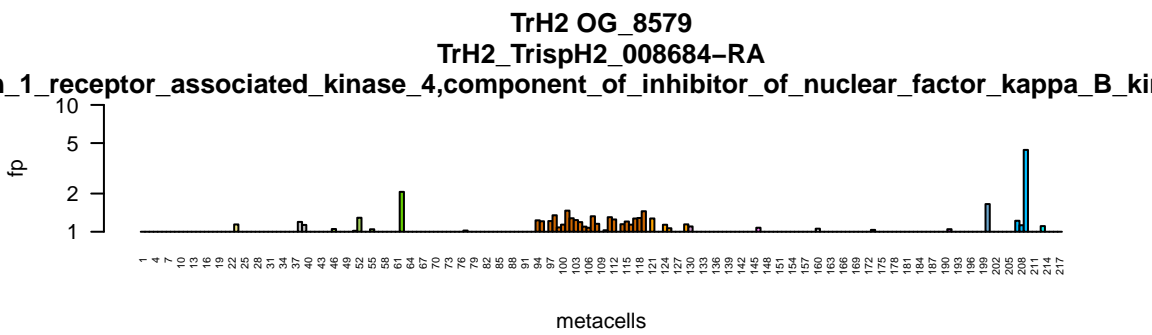
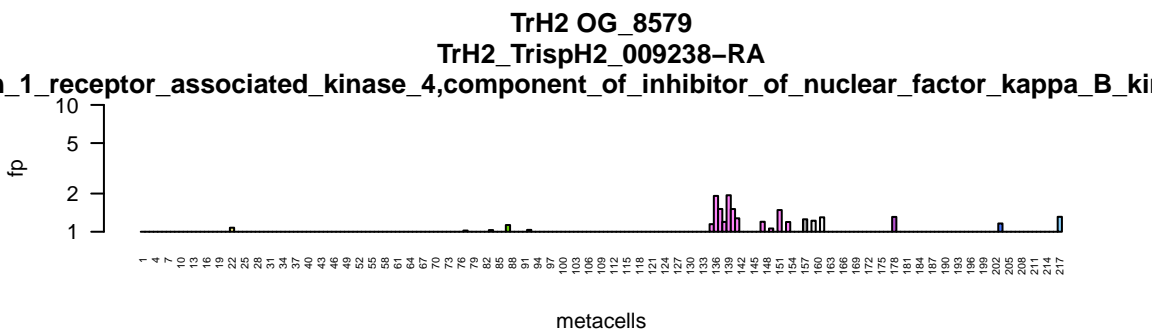
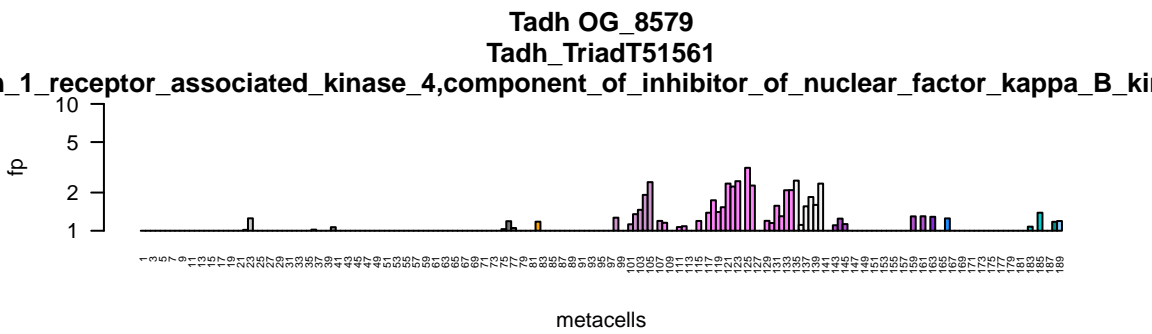








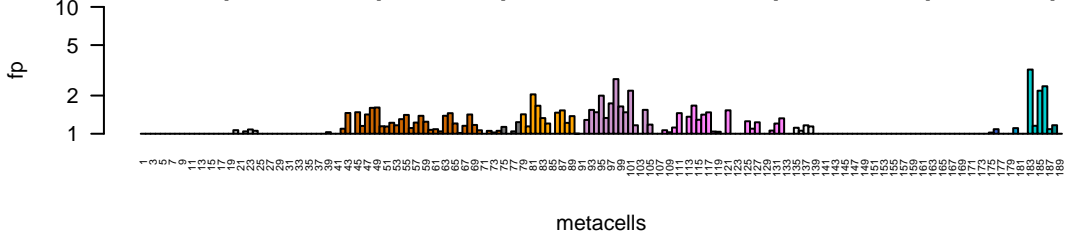




Tadh OG_9705

Tadh_TriadT64227

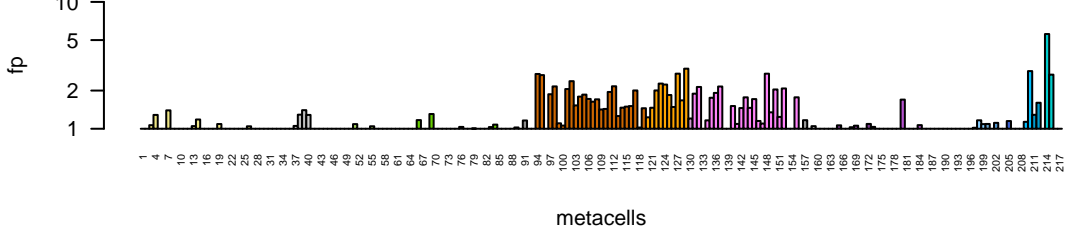
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TrH2 OG_9705

TrH2_TrispH2_010944-RA

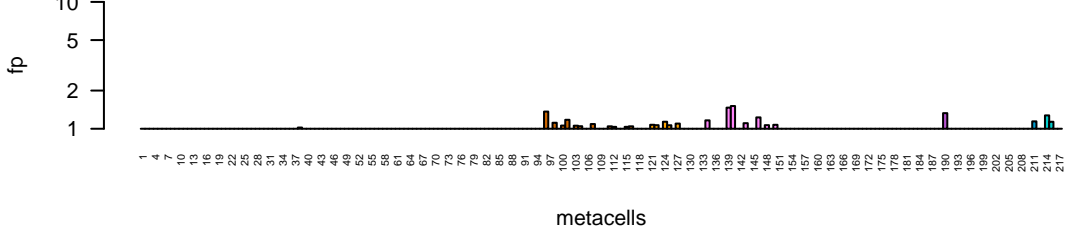
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TrH2 OG_9705

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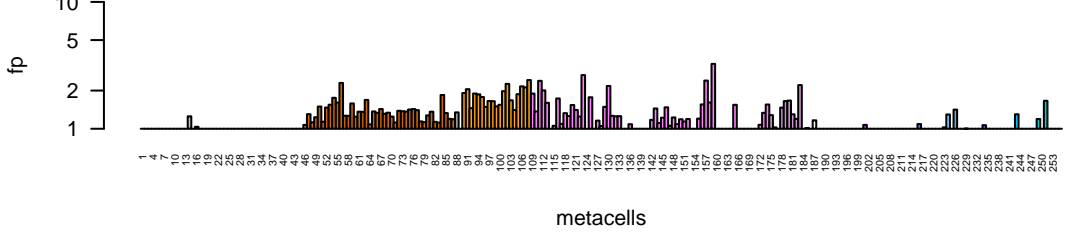
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Hhon OG_9705

Hhon_g09977.t1

adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_coupled_receptor_L1



HoiH23 OG_9705

HoiH23_PIH23_005603-RA

adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_coupled_receptor_L1

