

Bar chart showing the frequency of metacells (x-axis) versus the frequency of pairs (fp, y-axis). The x-axis lists metacells from 1 to 189. The y-axis ranges from 0 to 10. Most metacells have a frequency of 1, with a few having a frequency of 2. The bars are colored in a repeating pattern of orange, green, and blue.

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 232. Most metacells have 0 false positives, but a few have 1 or 2. Metacells 208, 211, 214, 217, 220, 223, 228, and 232 have 1 false positive. Metacells 220, 228, and 232 have 2 false positives.

and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

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	1.5
211	1
214	1.2
217	1

and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

fp

metacells

and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

fp

metacells

and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

metacell	fp
1	1
4	1
10	1.5
13	1
16	1
19	1
22	1.2
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	1.1
220	1
223	1
226	1
229	1
232	2.5

and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

fp

metacells

and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

metacells	fp
1	1
4	1
10	1
13	1
16	1
19	1
22	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
202	1
205	1
208	1
211	1
214	1.5
217	1.8
220	1.2
223	3.5
226	2.5
229	2.2
232	1.2

and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

metacell	fp
1	1
7	1
10	1
16	1
19	1
25	1
34	1
37	1
43	1
46	1
52	1
55	1
61	1
67	1
70	1
76	1
79	1
85	1
88	1
94	1
97	1
103	1
106	1
112	1
115	1
121	2
127	1
130	1
138	1
145	1
148	1
154	1
167	1
168	1
169	1
172	1
175	1
181	1
187	1
190	1
198	1
199	1
205	1.5
211	1
214	1
217	1
223	1
226	1
232	1
235	1
236	1
241	1
244	1
247	4
250	4
253	1

and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

fp

metacells

and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

fp

metacells

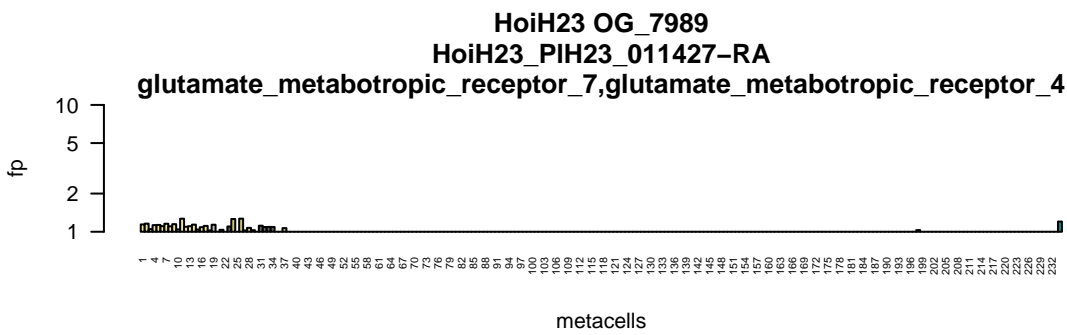
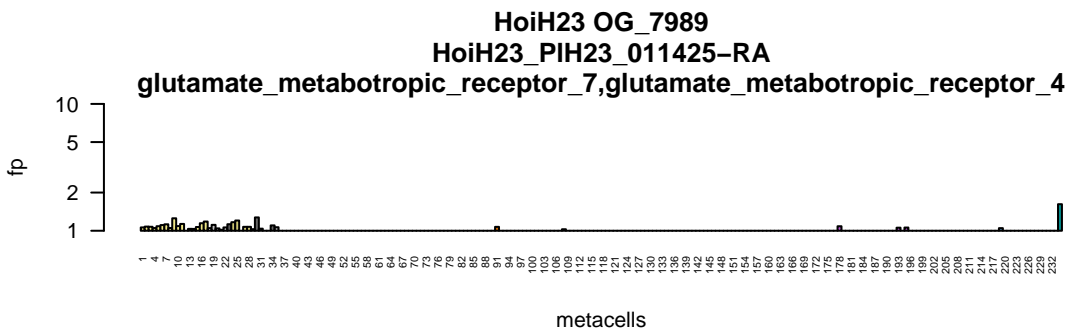
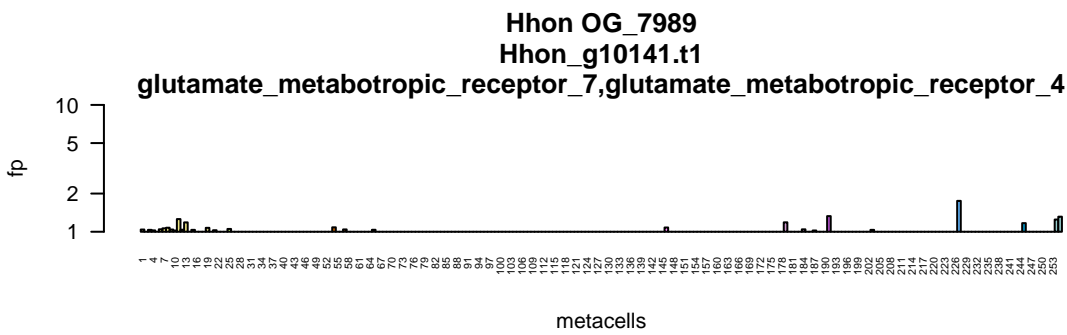
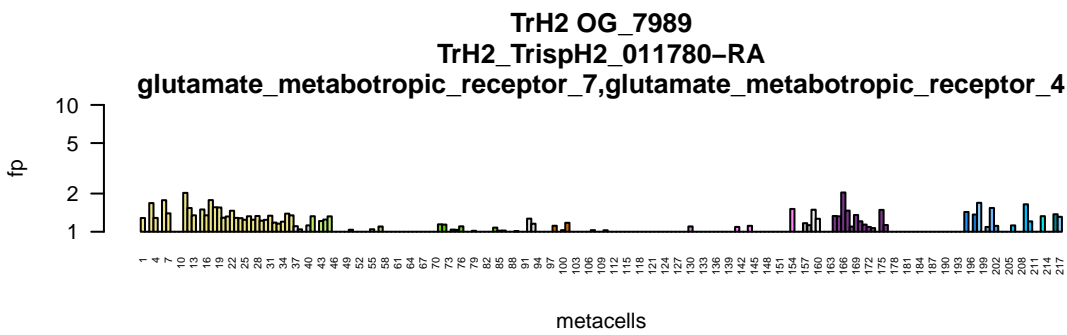
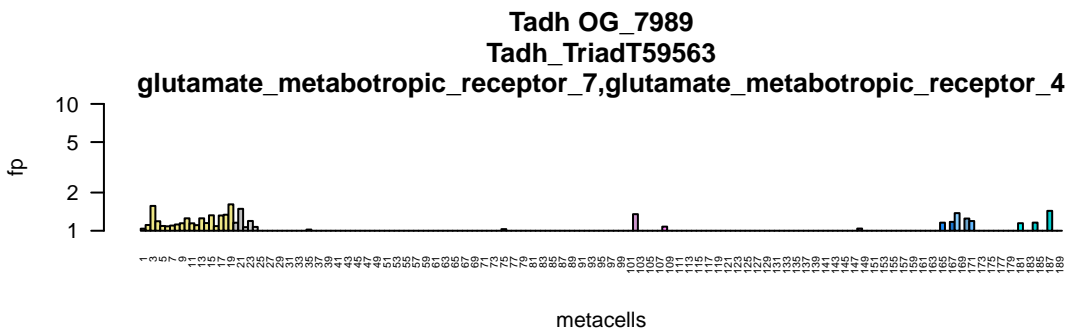
and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

fp

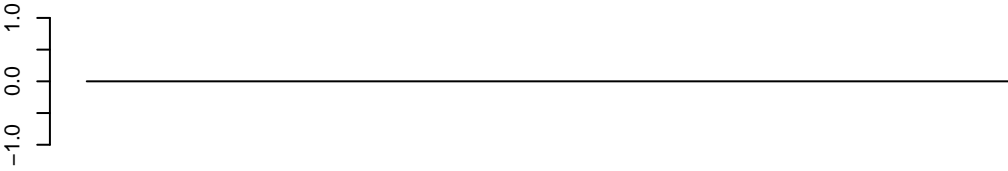
metacells

and PNKP_like_factor,G_protein_coupled_receptor_183,somatostatin_receptor_5,

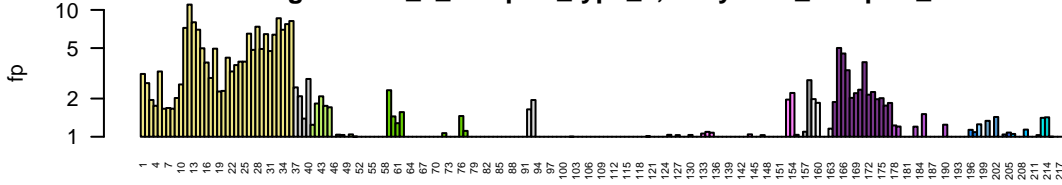
metacells	fp
1	1
4	1
10	1
13	1
16	1
19	1
22	1
25	1
28	1
31	1
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37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	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	2
202	1
205	1
208	1
211	1
214	1
217	1
220	1
223	1
226	1
229	1
232	2.5



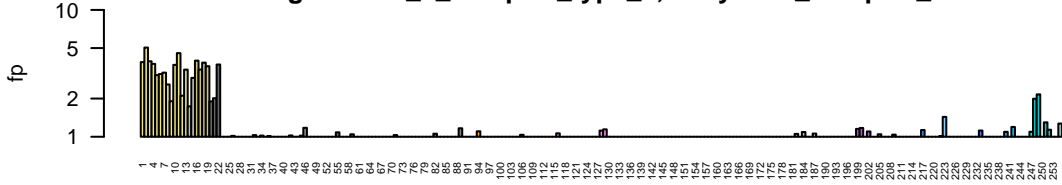
angiotensin_II_receptor_type_2,tachykinin_receptor_1
Tadh | no data



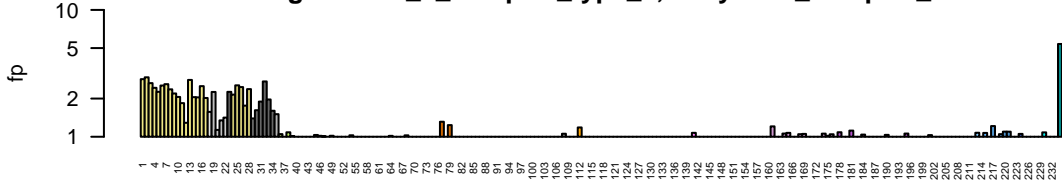
TrH2 OG_9264
TrH2_TrispH2_002603-RA
angiotensin_II_receptor_type_2,tachykinin_receptor_1

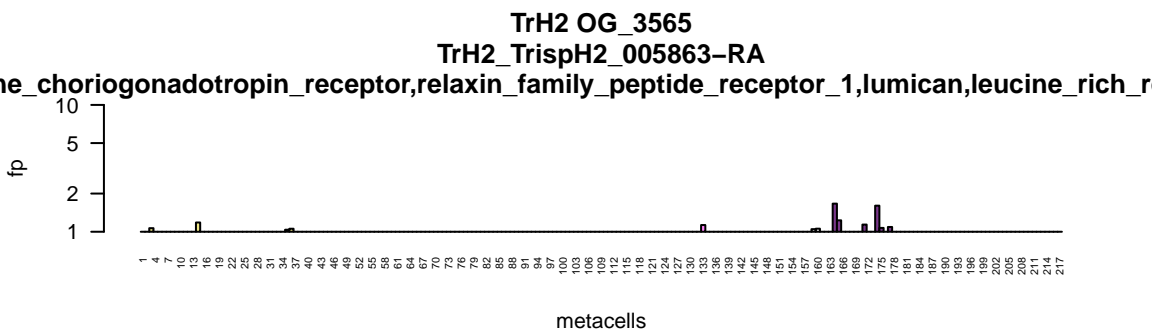
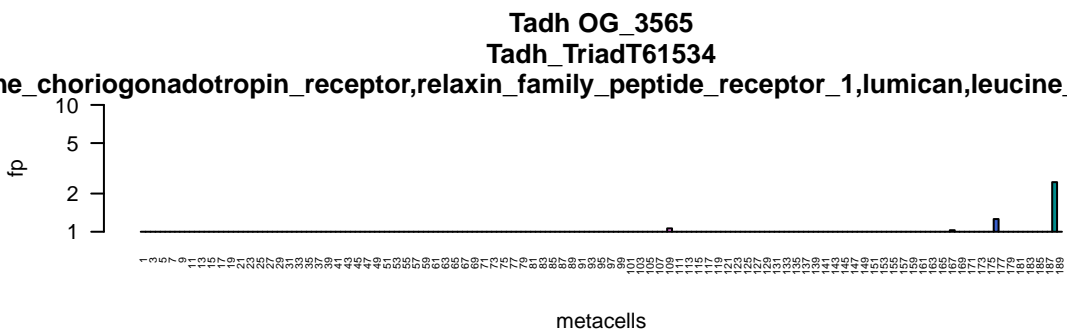
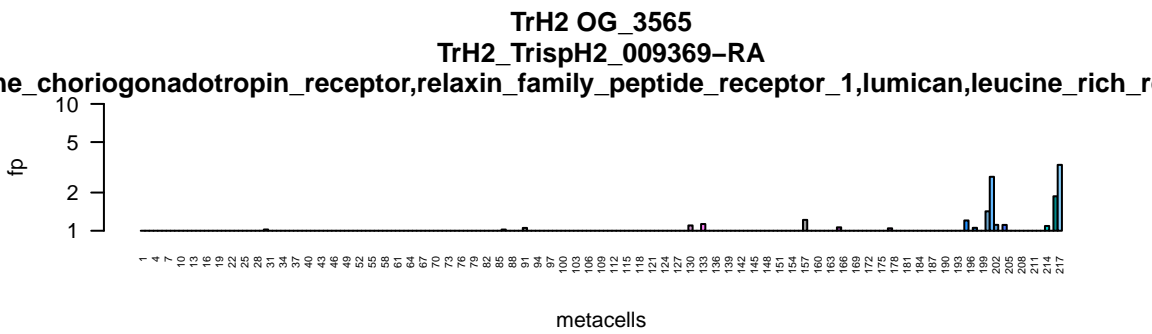
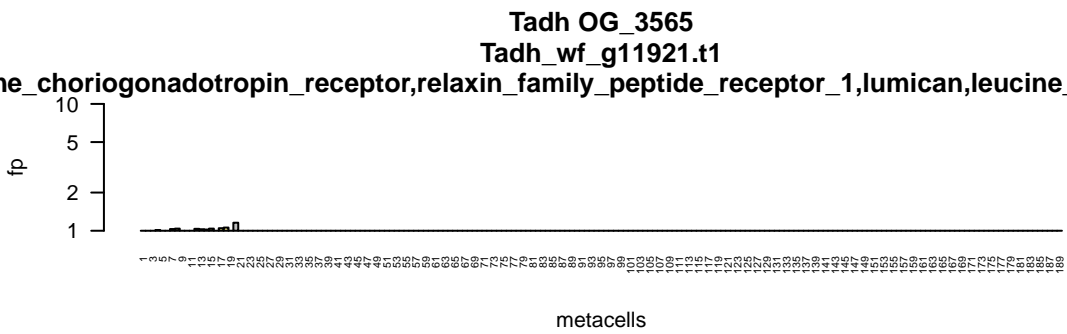
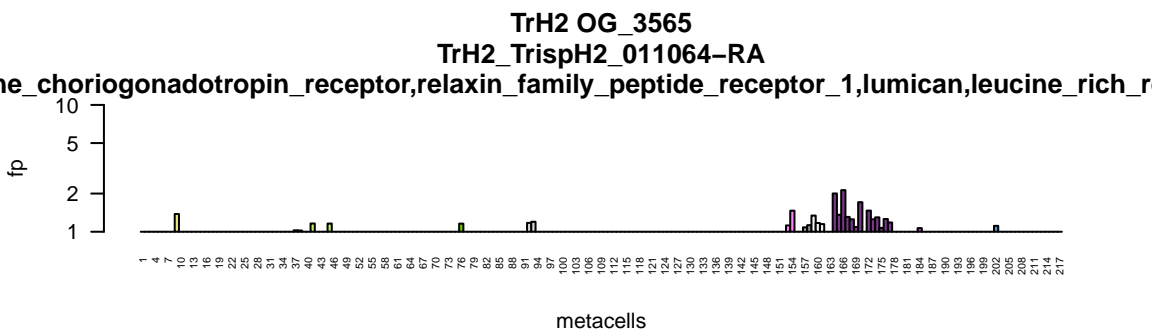
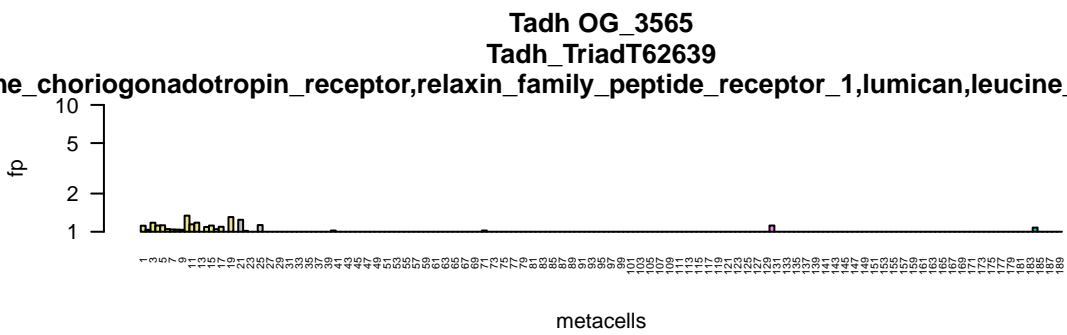
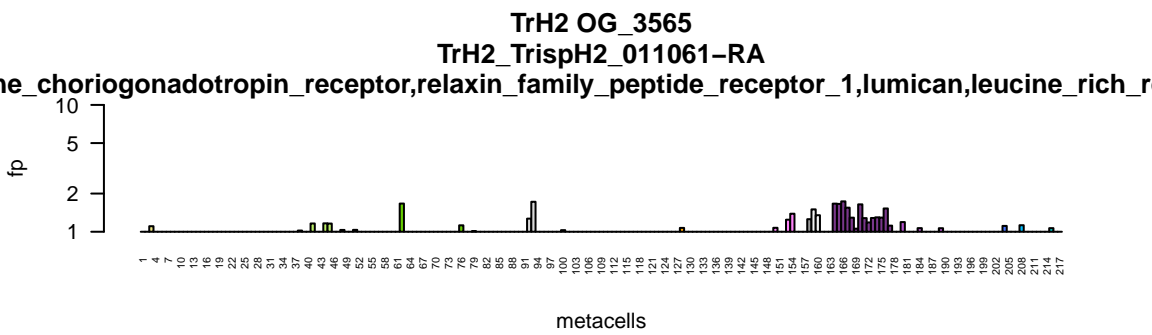
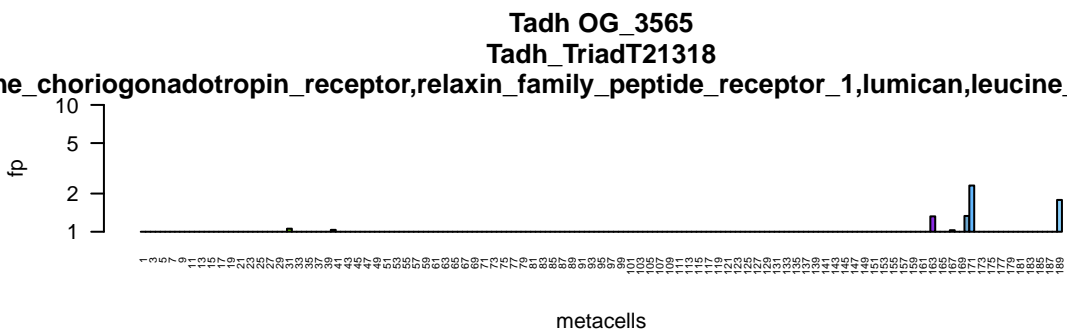
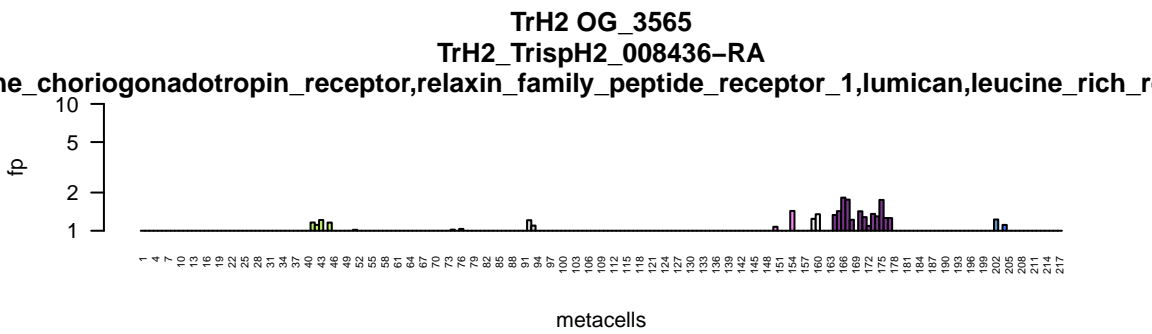
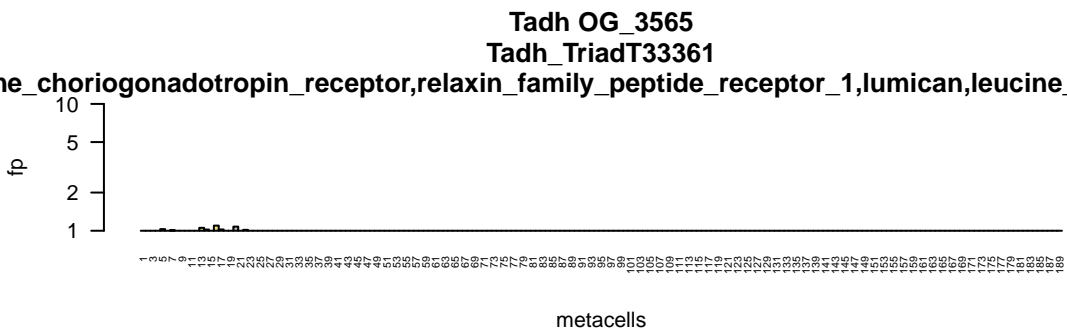
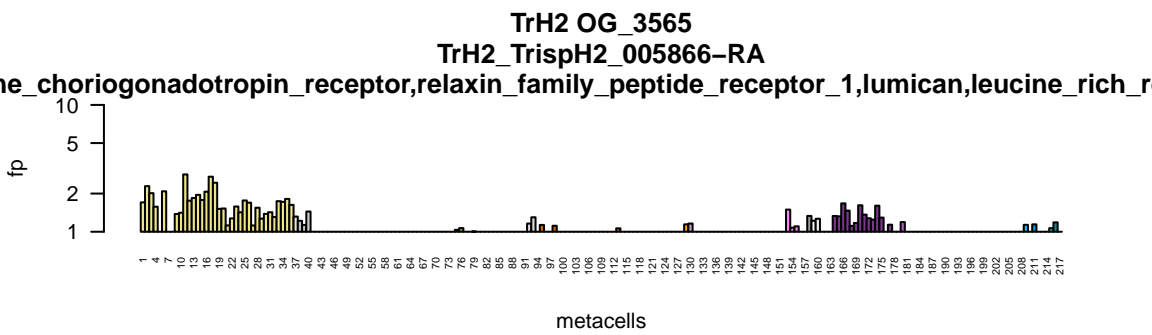
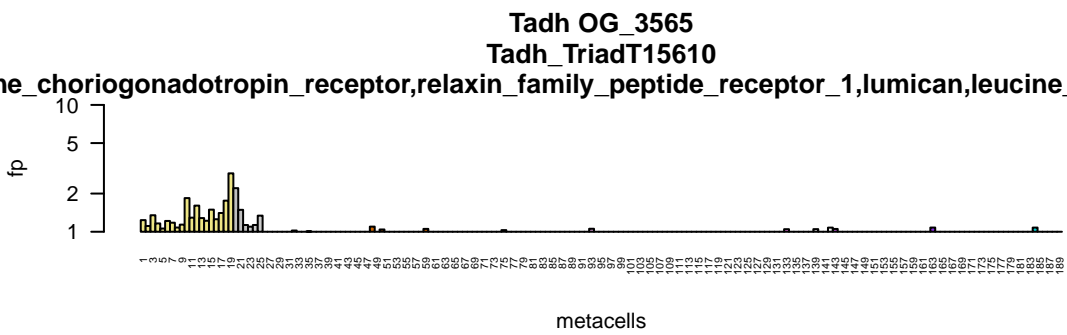
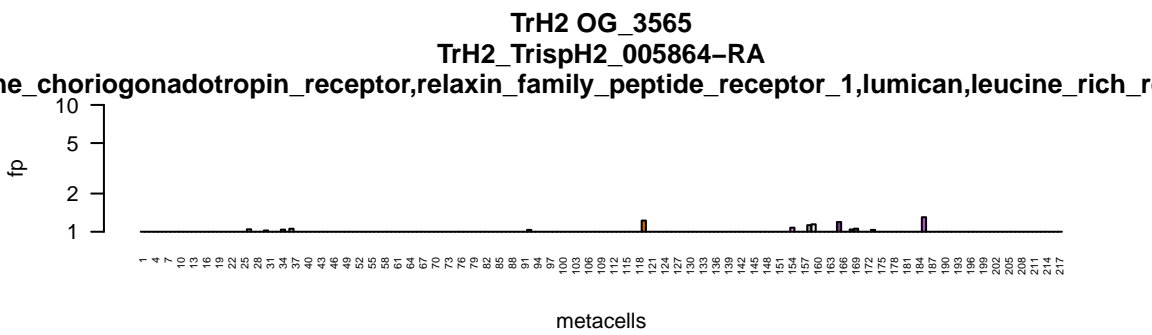
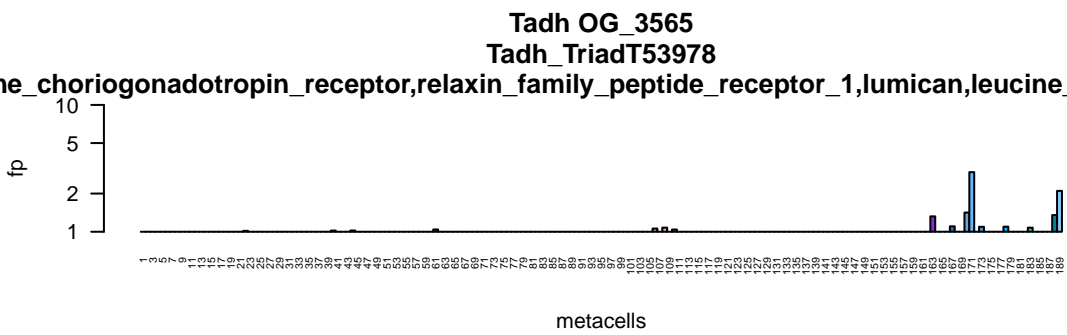
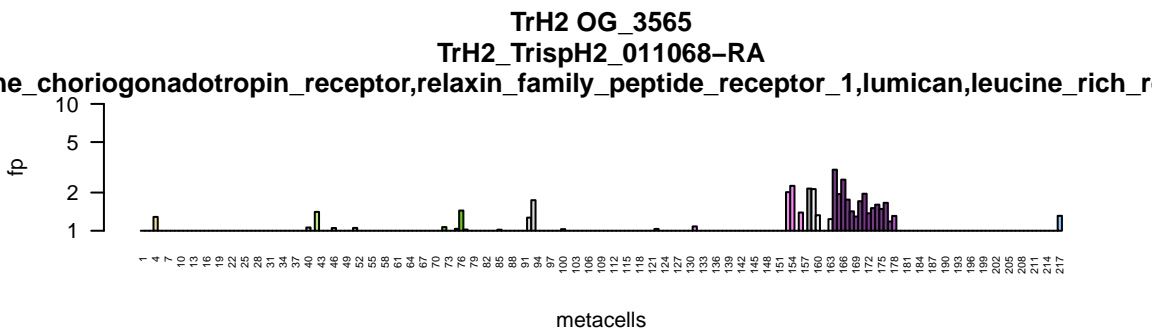
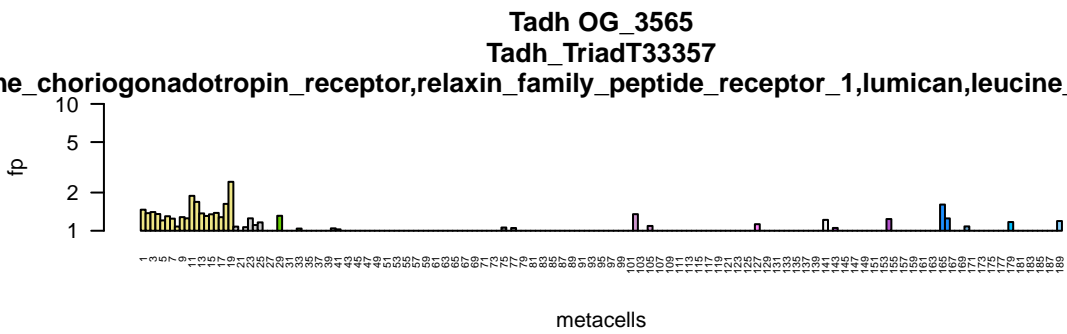


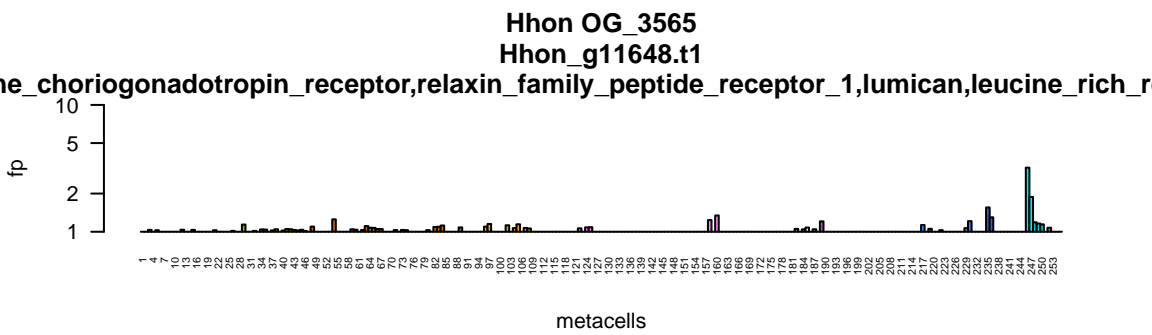
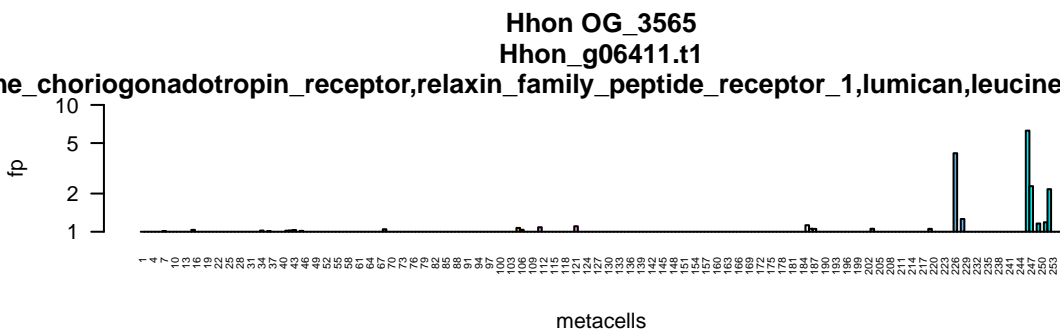
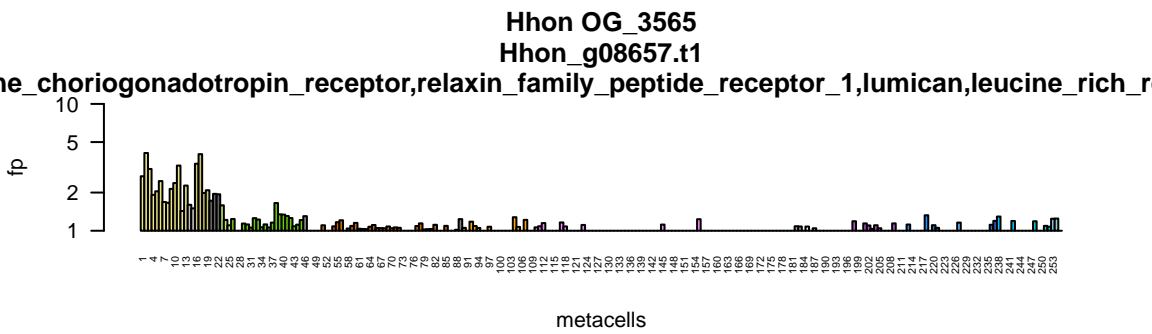
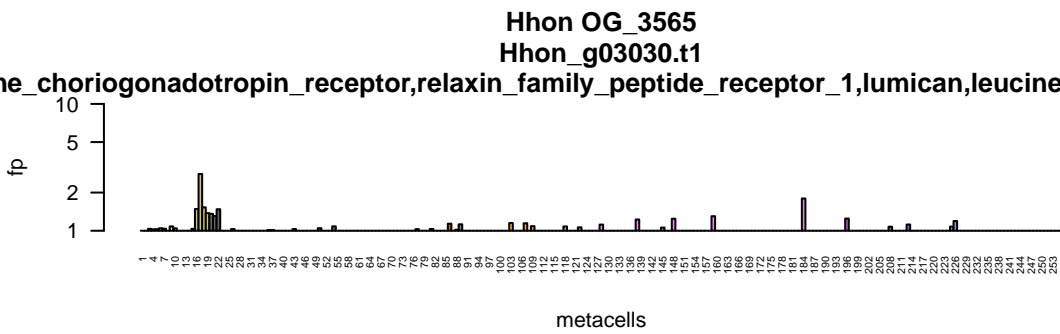
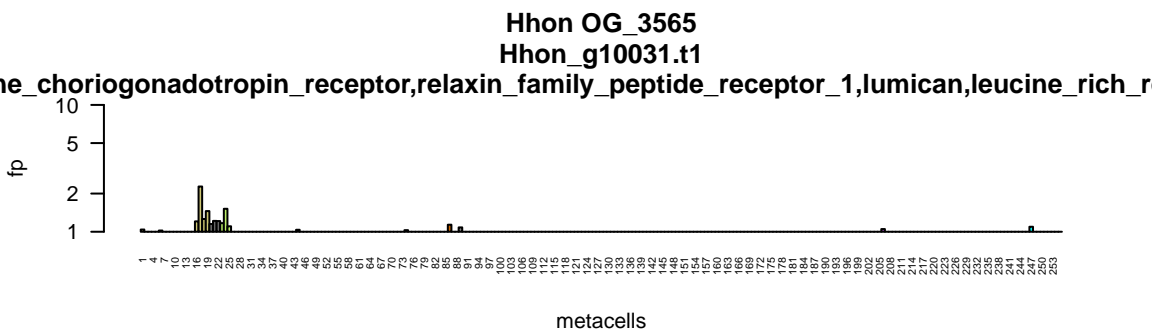
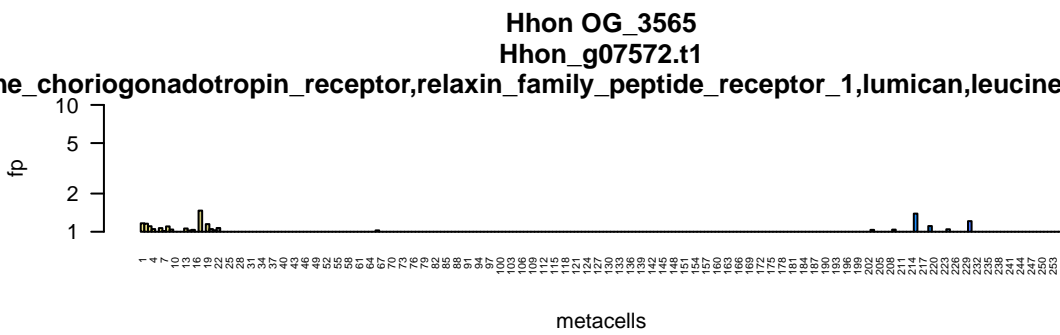
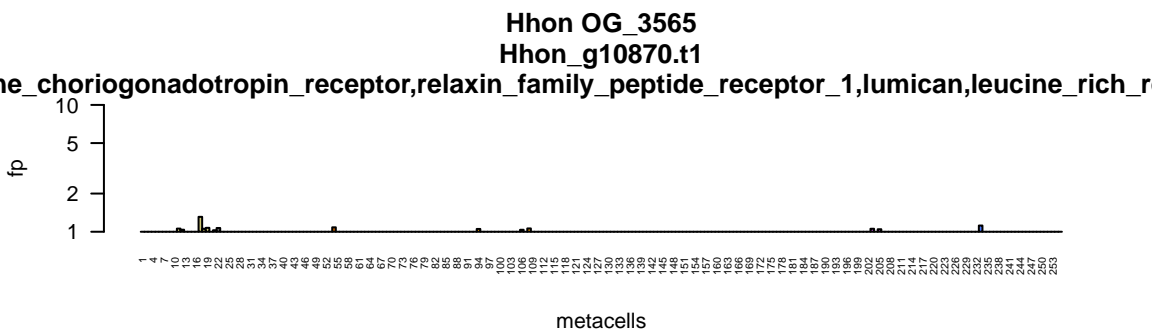
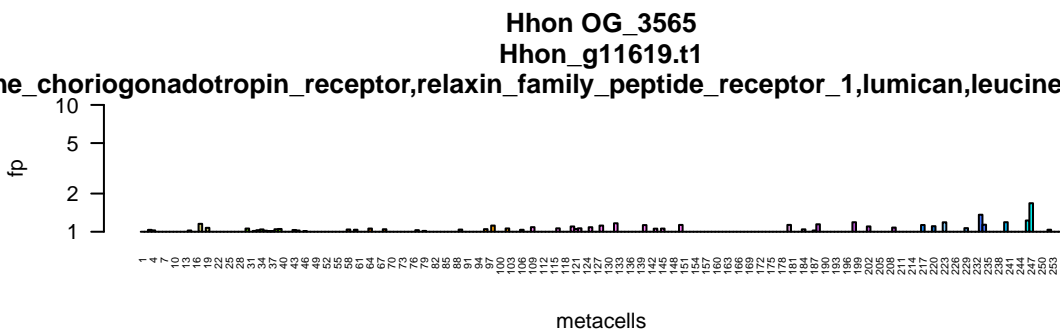
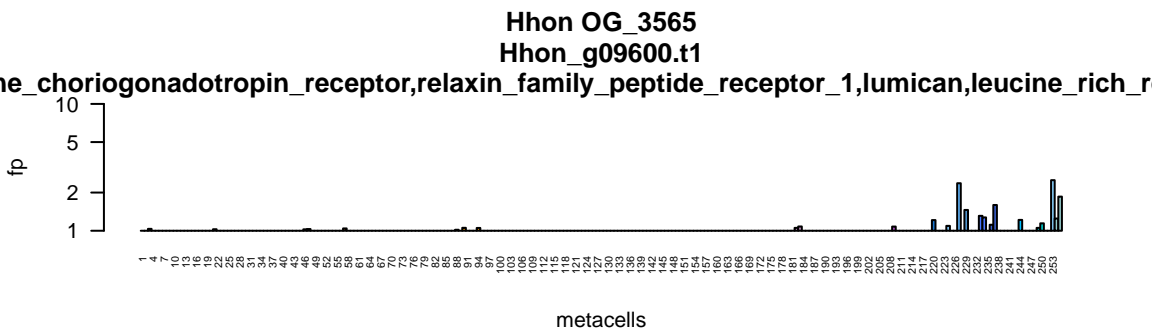
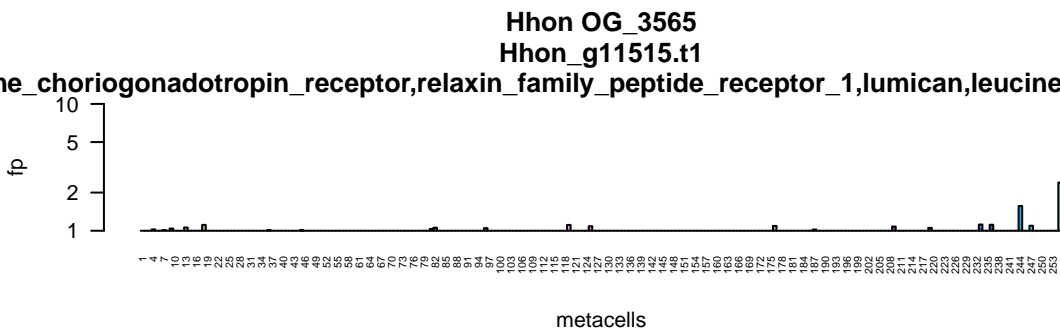
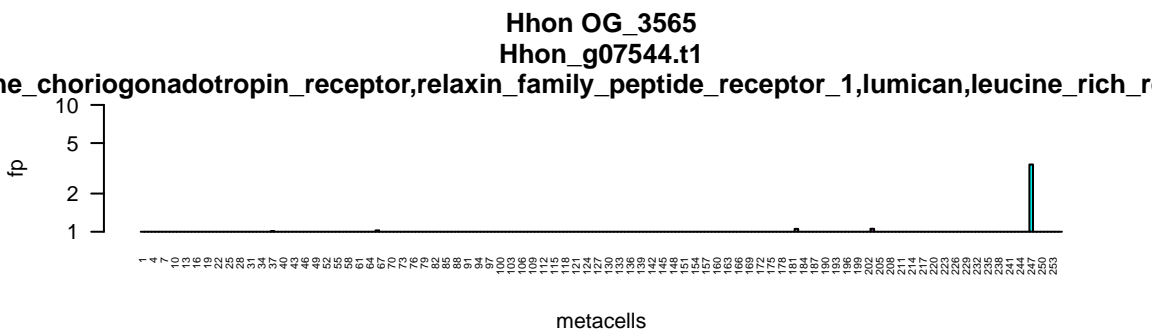
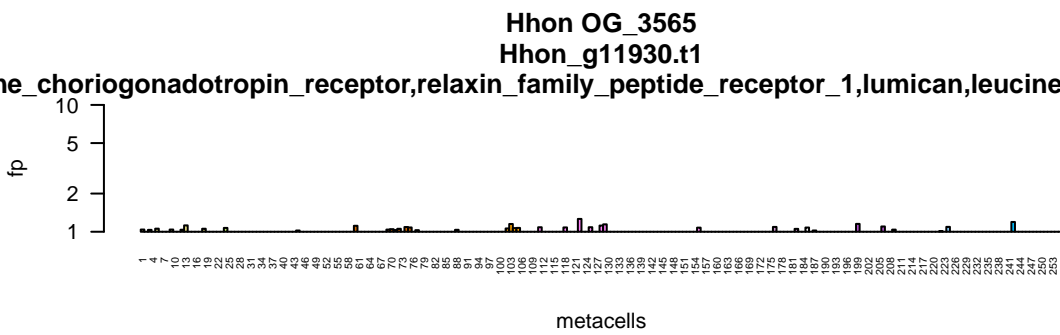
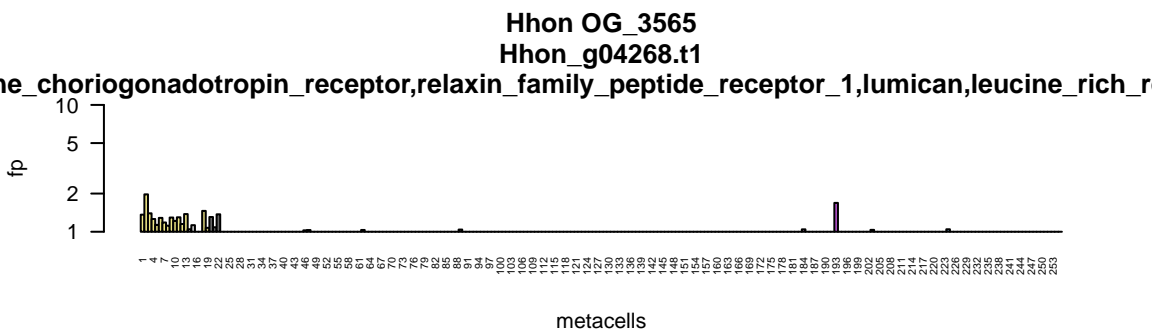
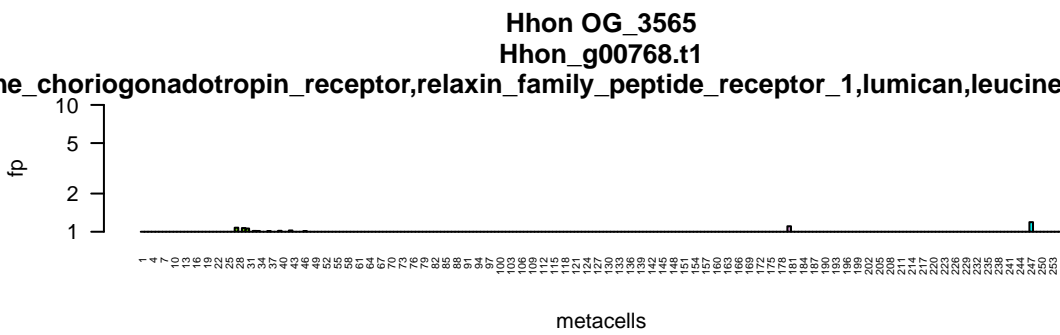
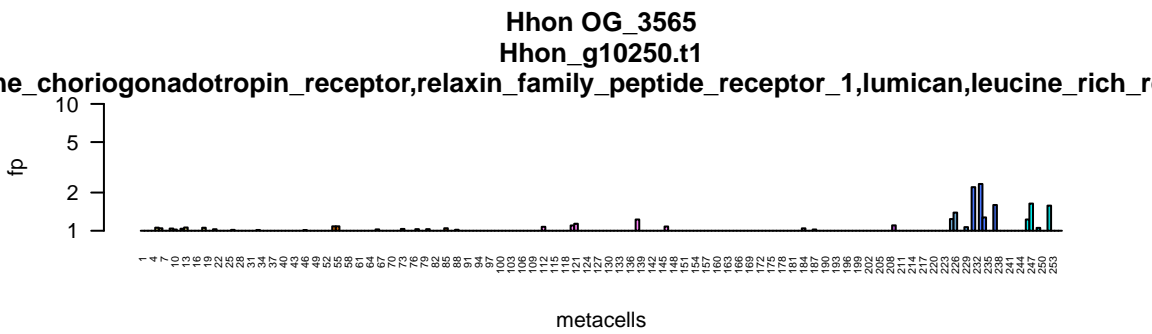
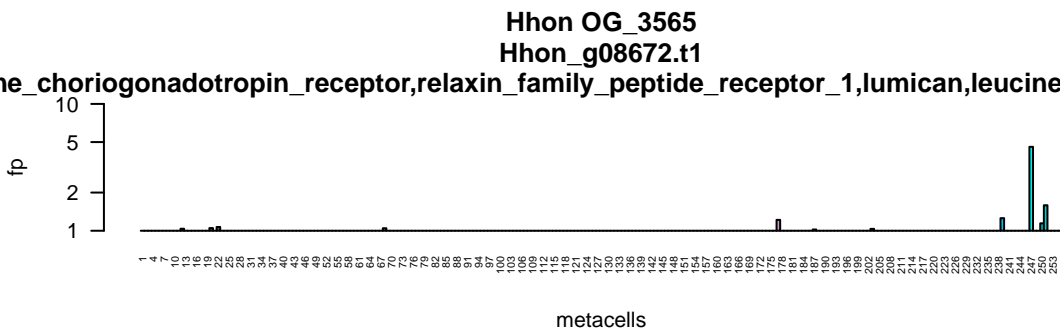
Hhon OG_9264
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angiotensin_II_receptor_type_2,tachykinin_receptor_1

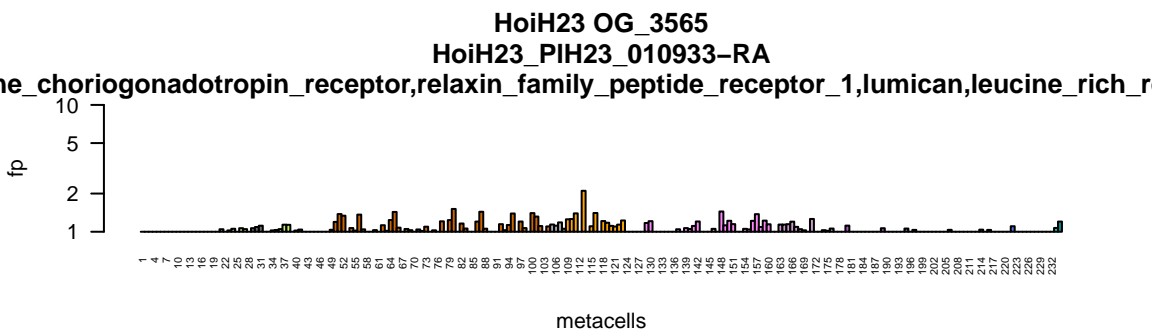
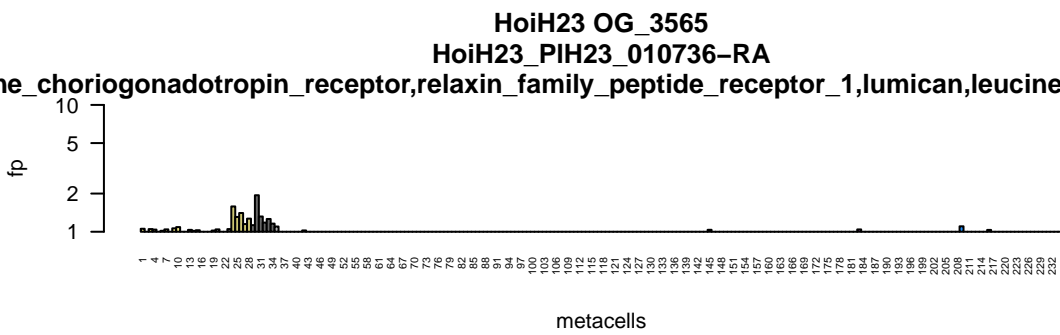
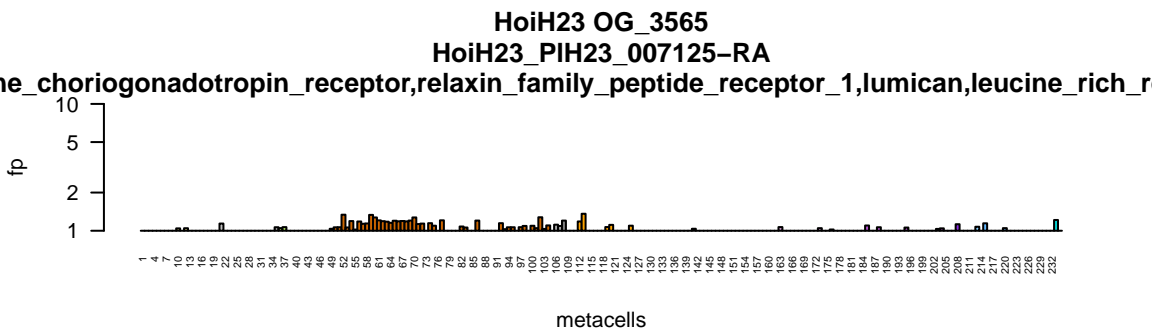
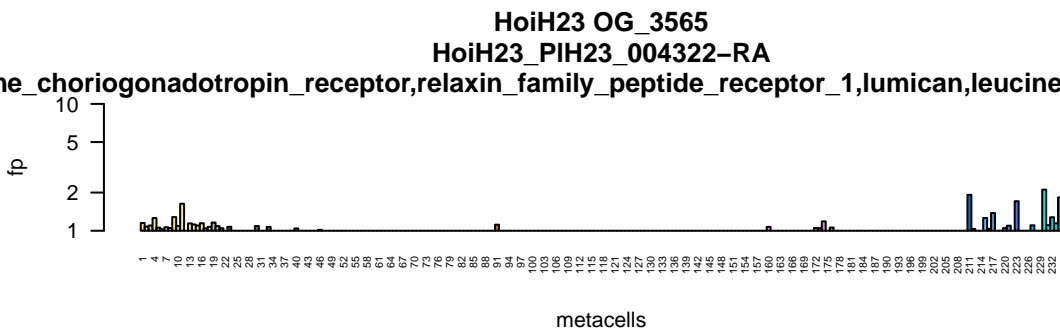
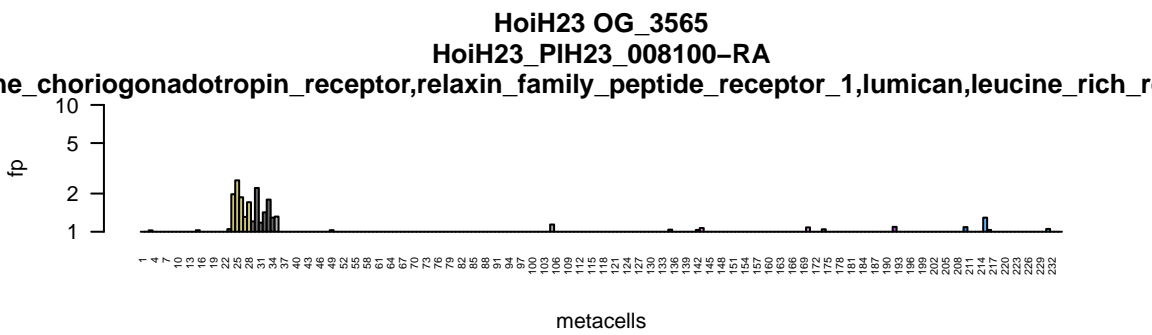
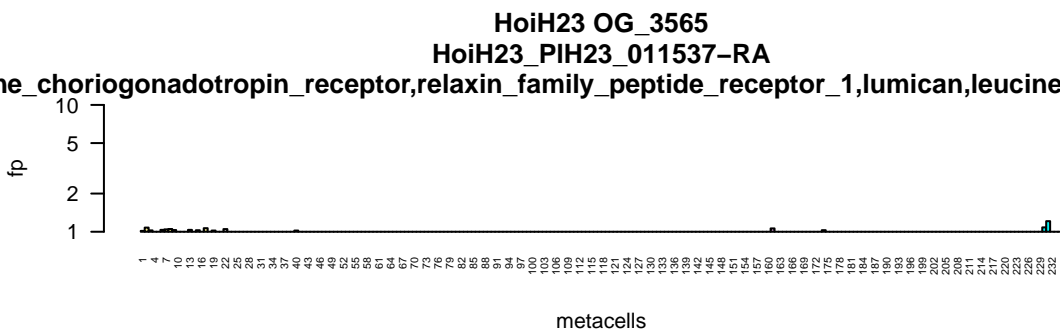
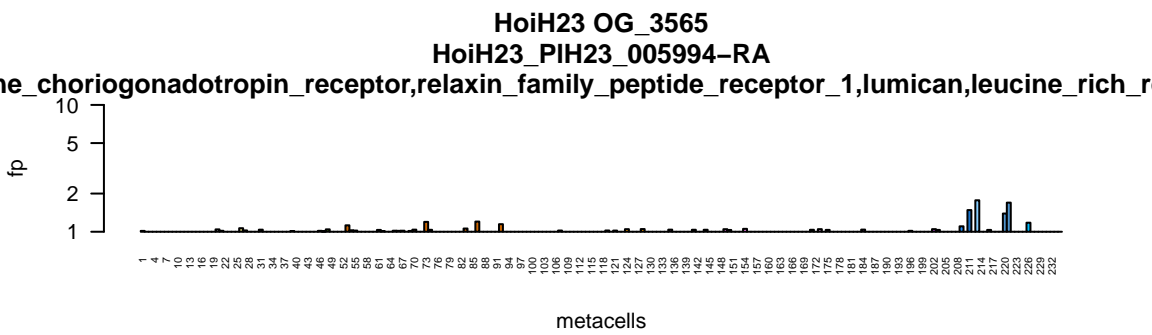
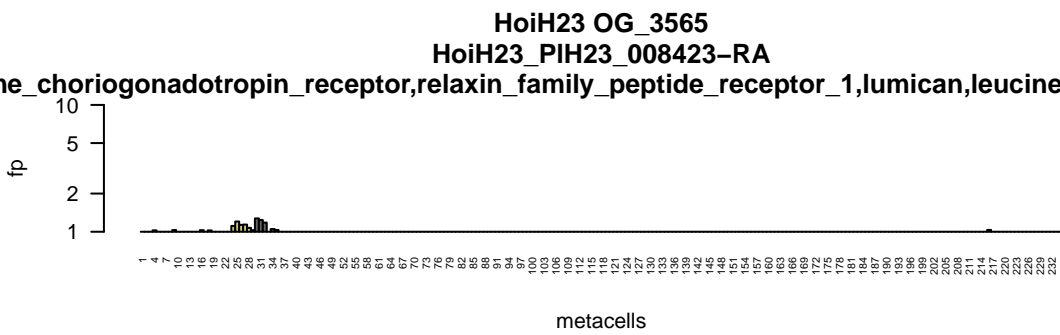
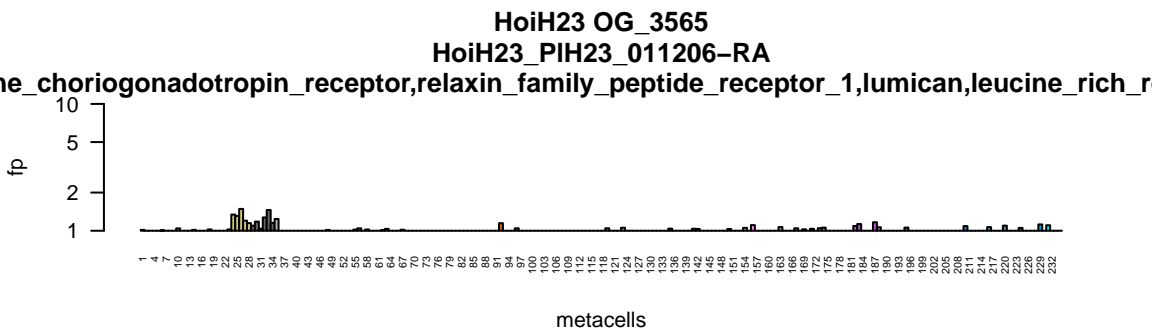
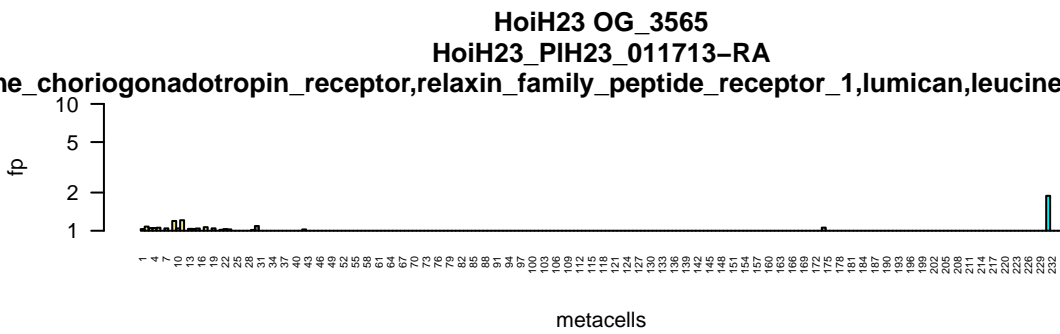
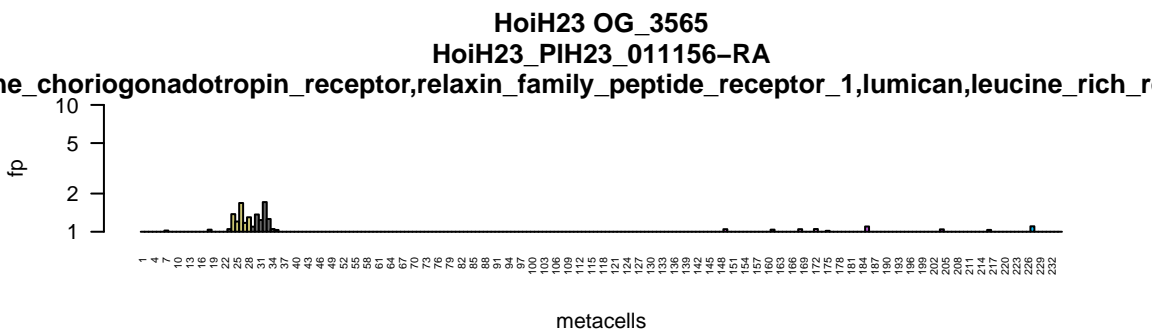
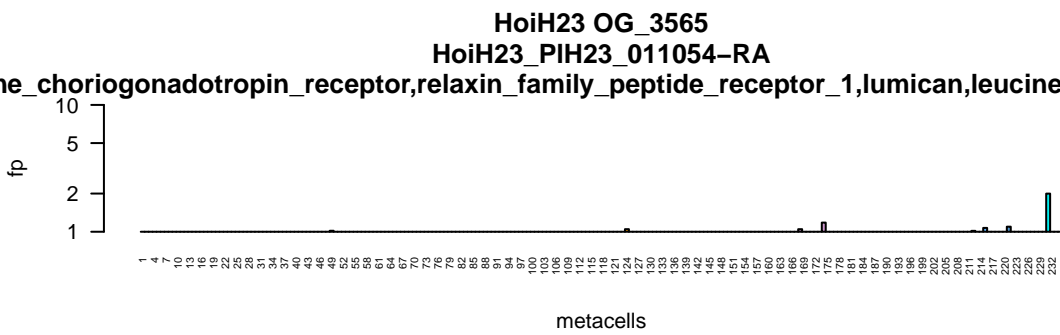
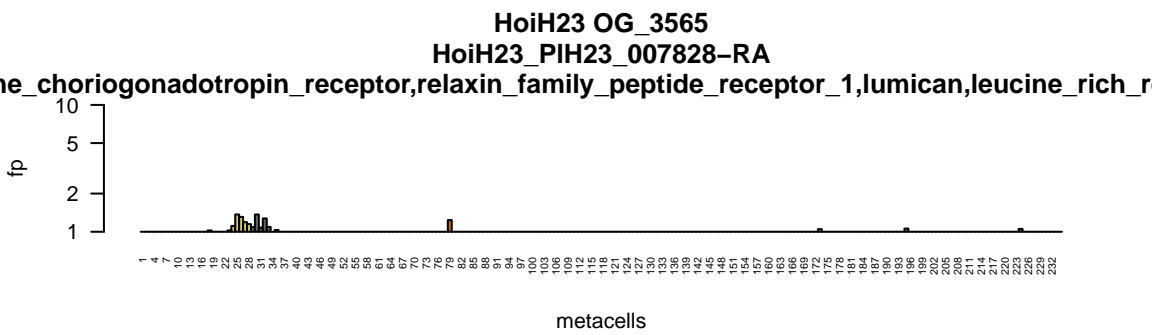
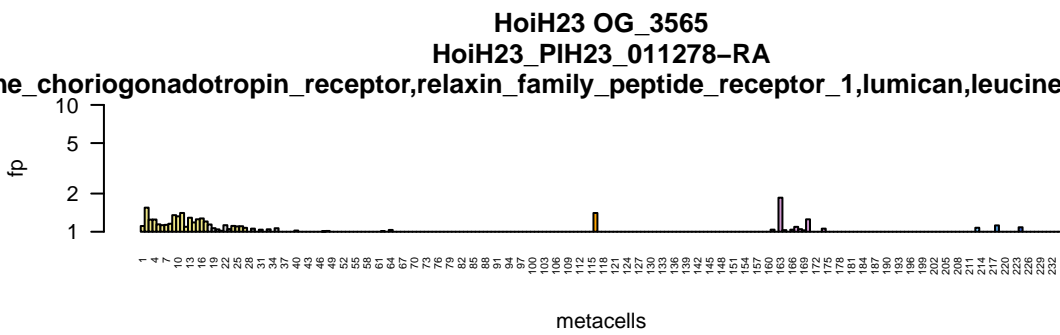
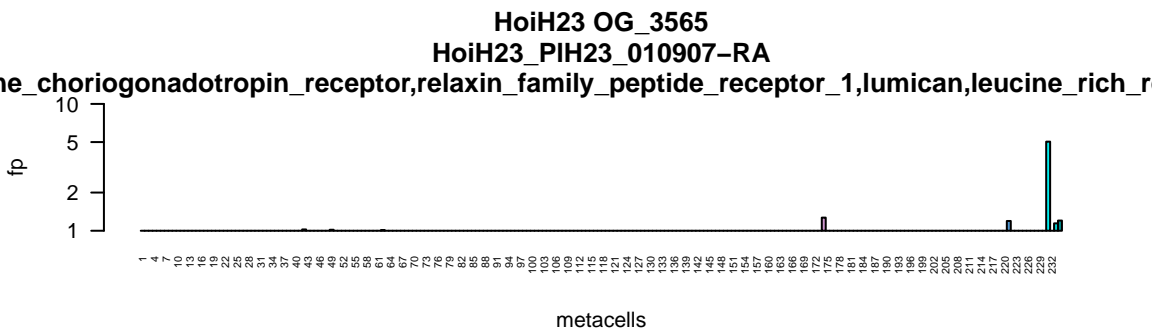
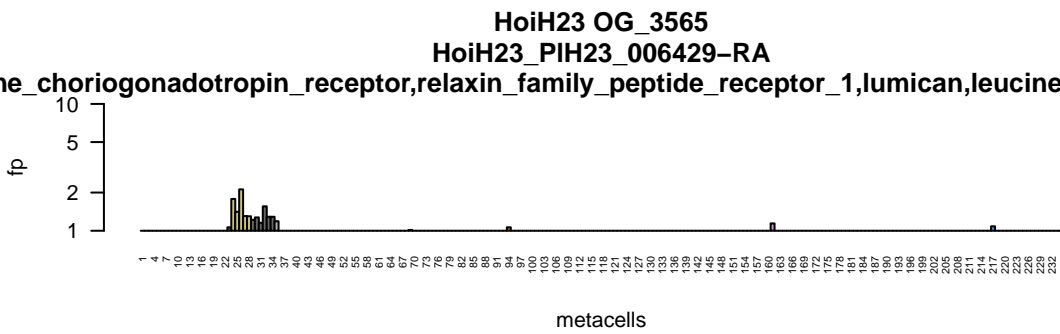


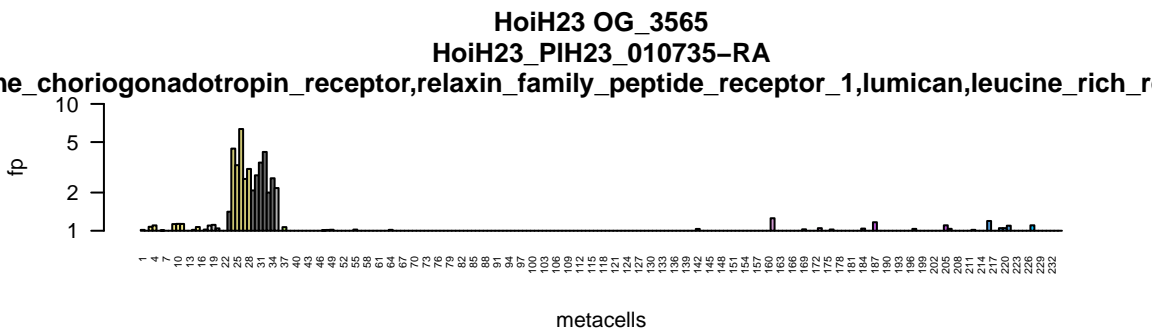
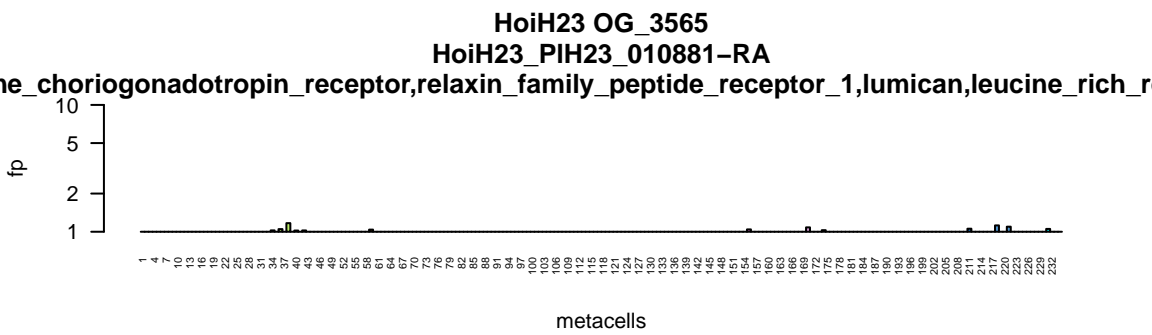
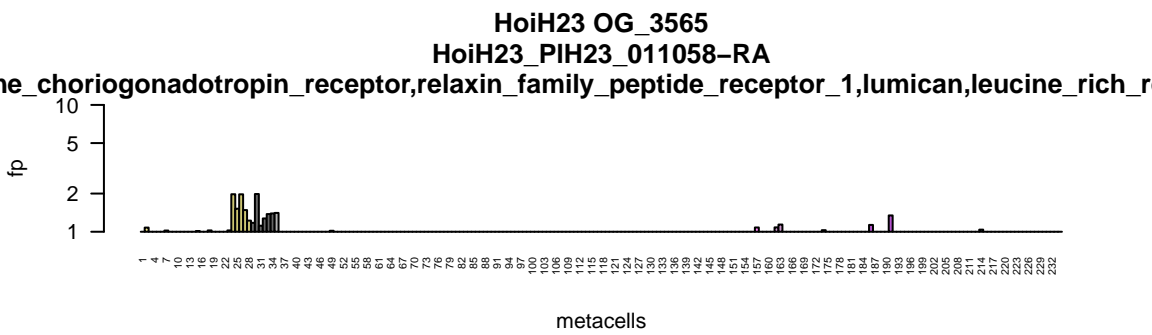
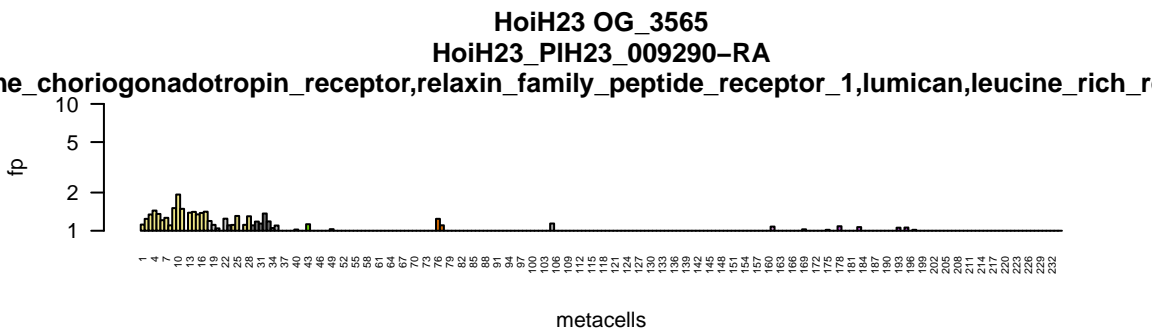
HoiH23 OG_9264
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angiotensin_II_receptor_type_2,tachykinin_receptor_1

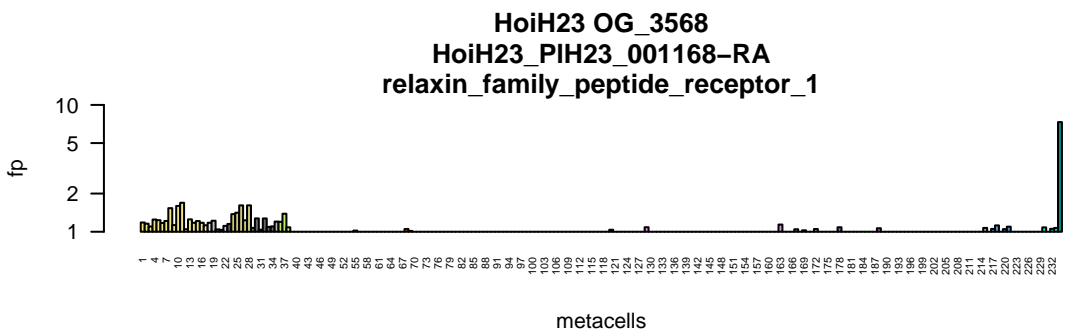
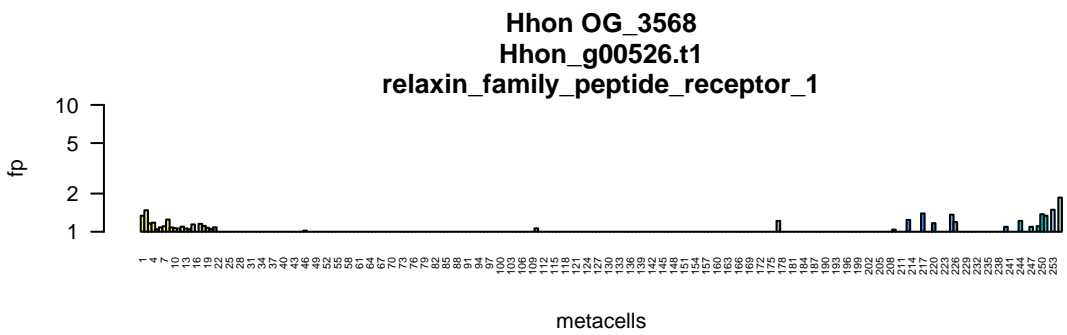
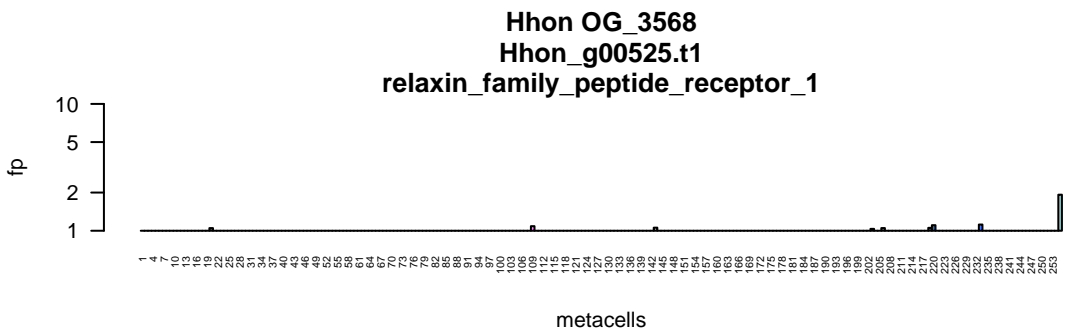
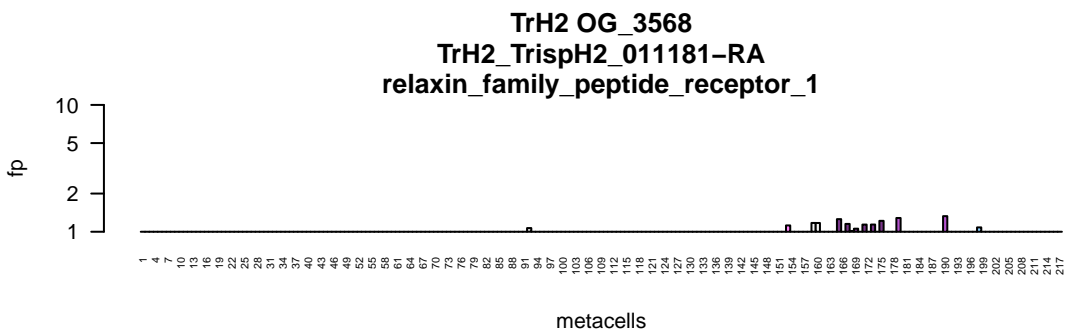
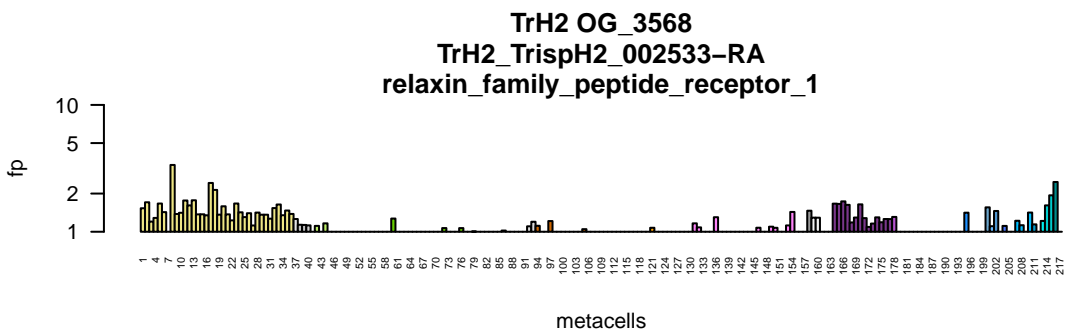
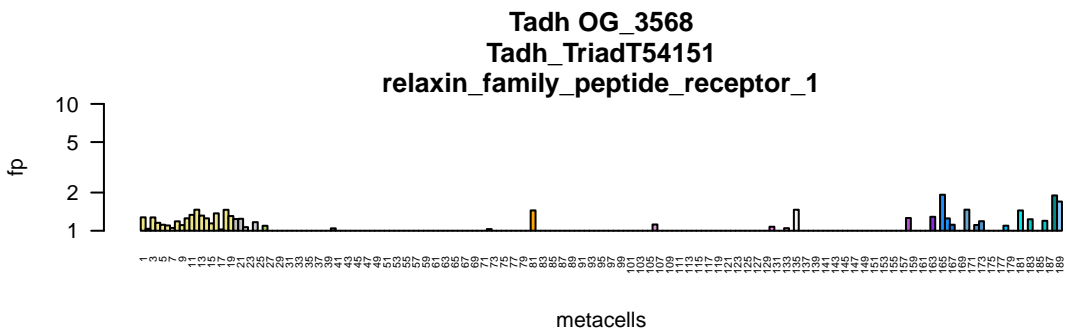


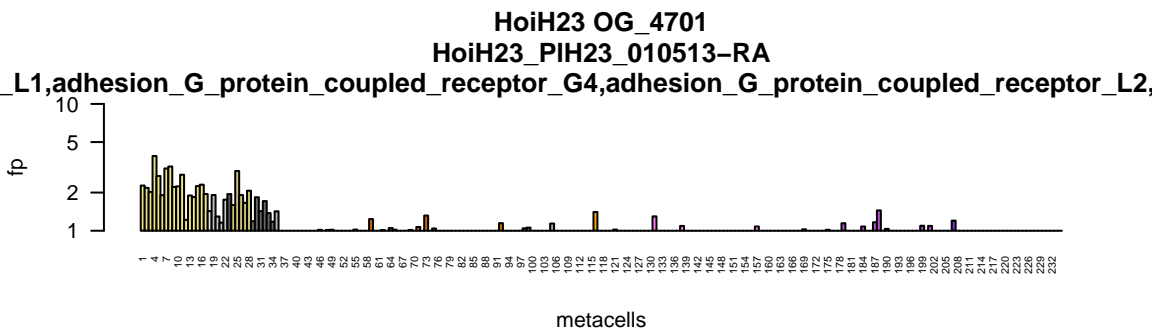
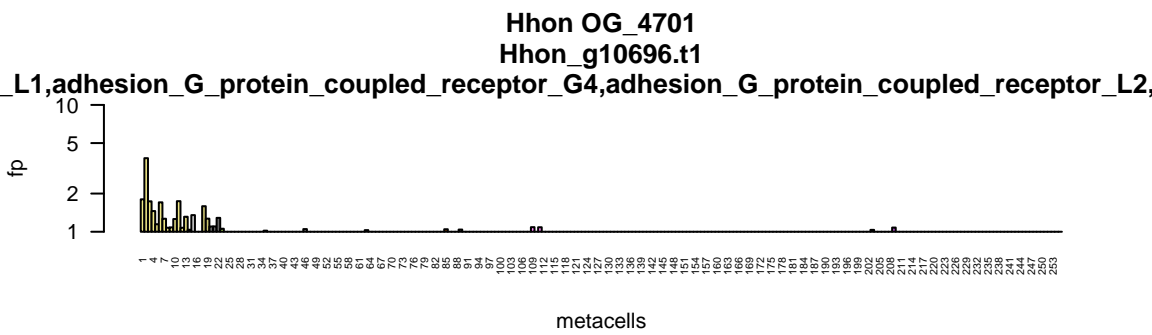
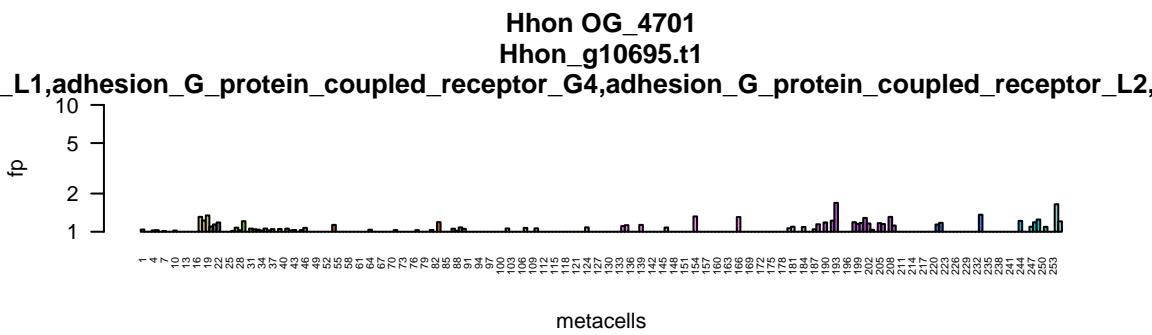
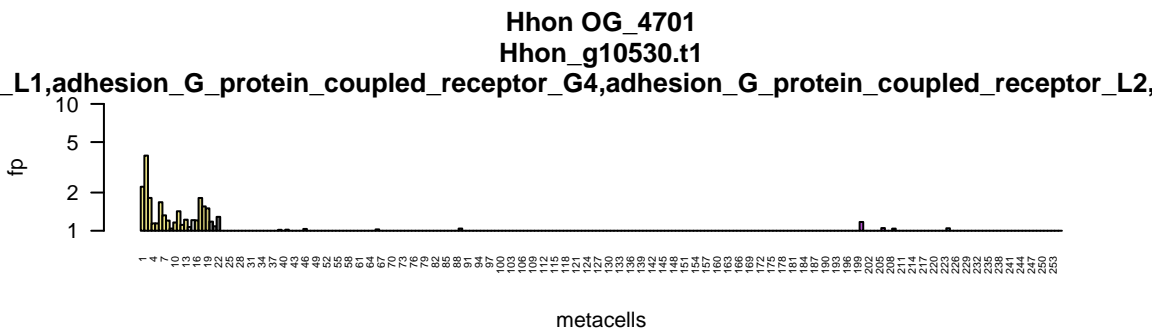
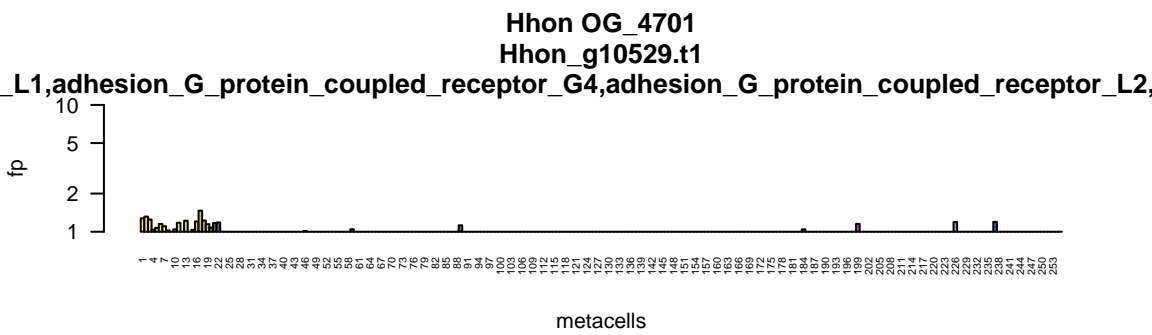
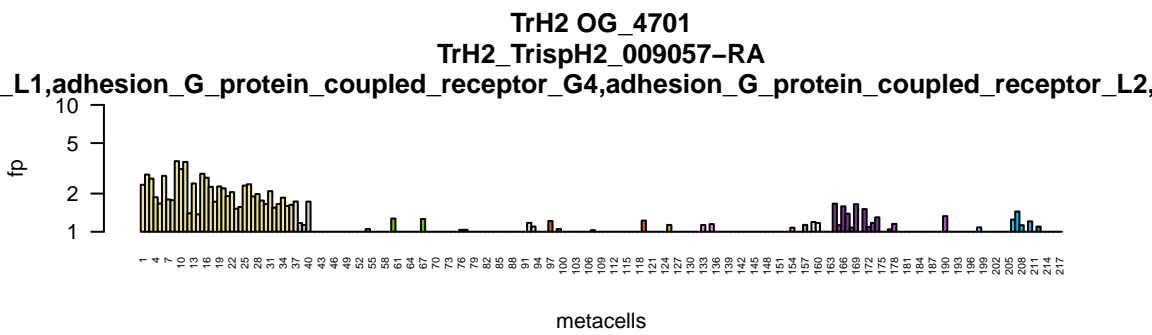
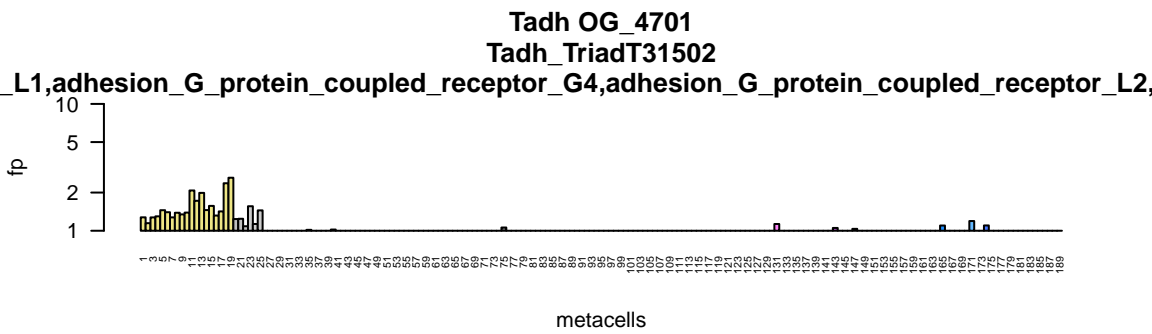


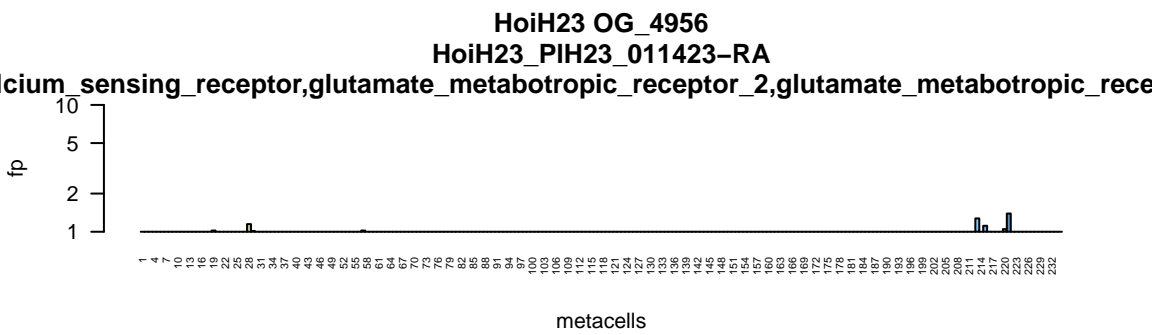
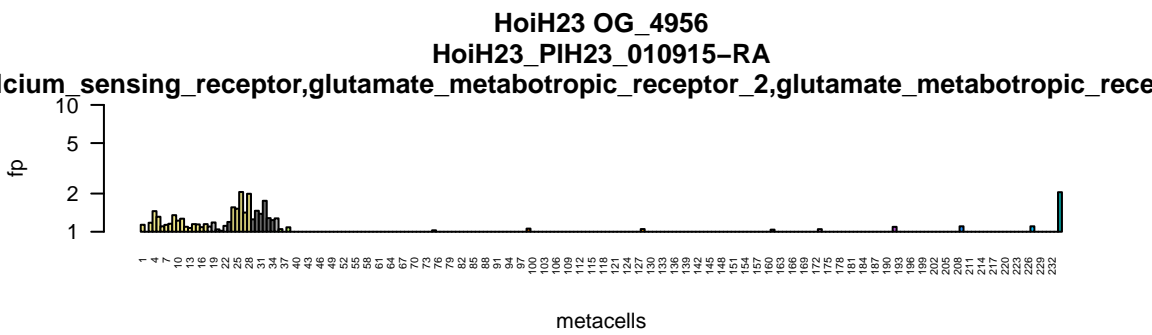
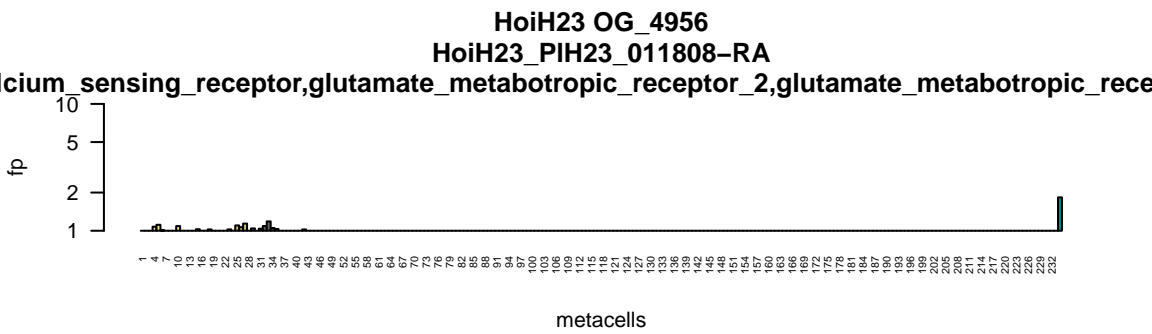
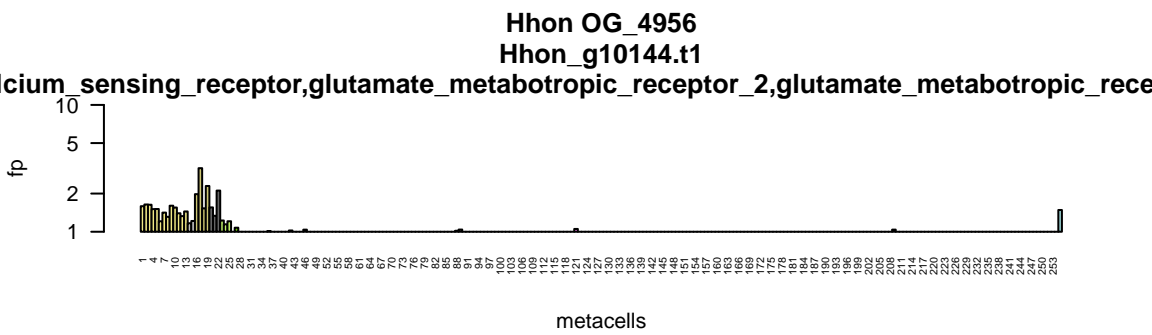
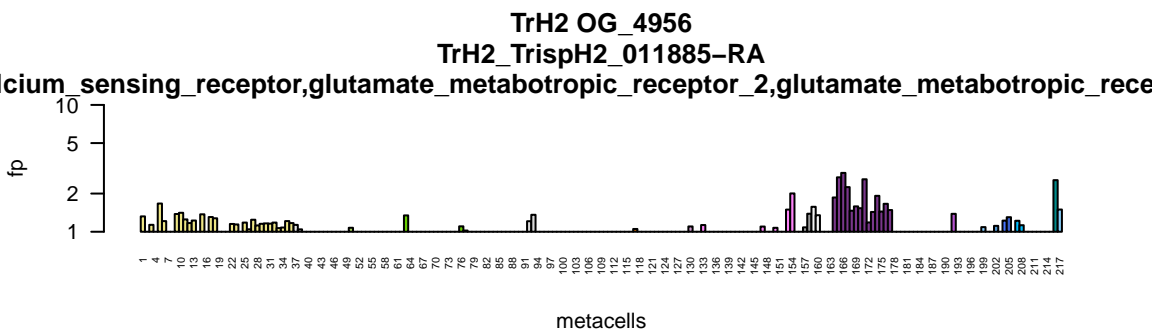
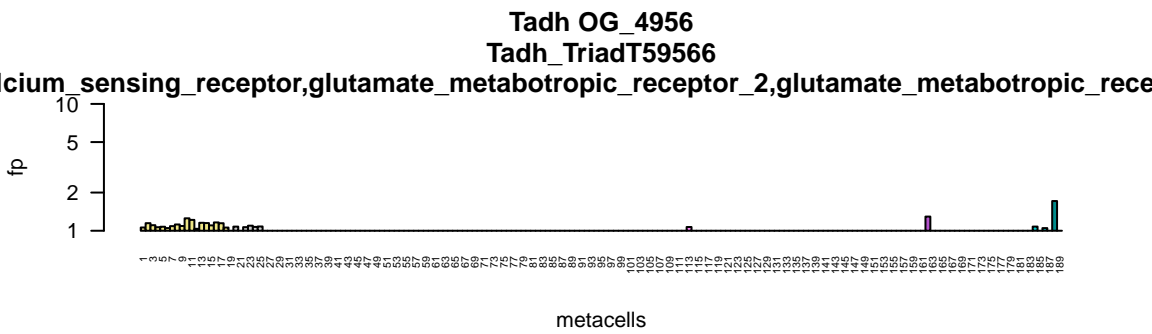


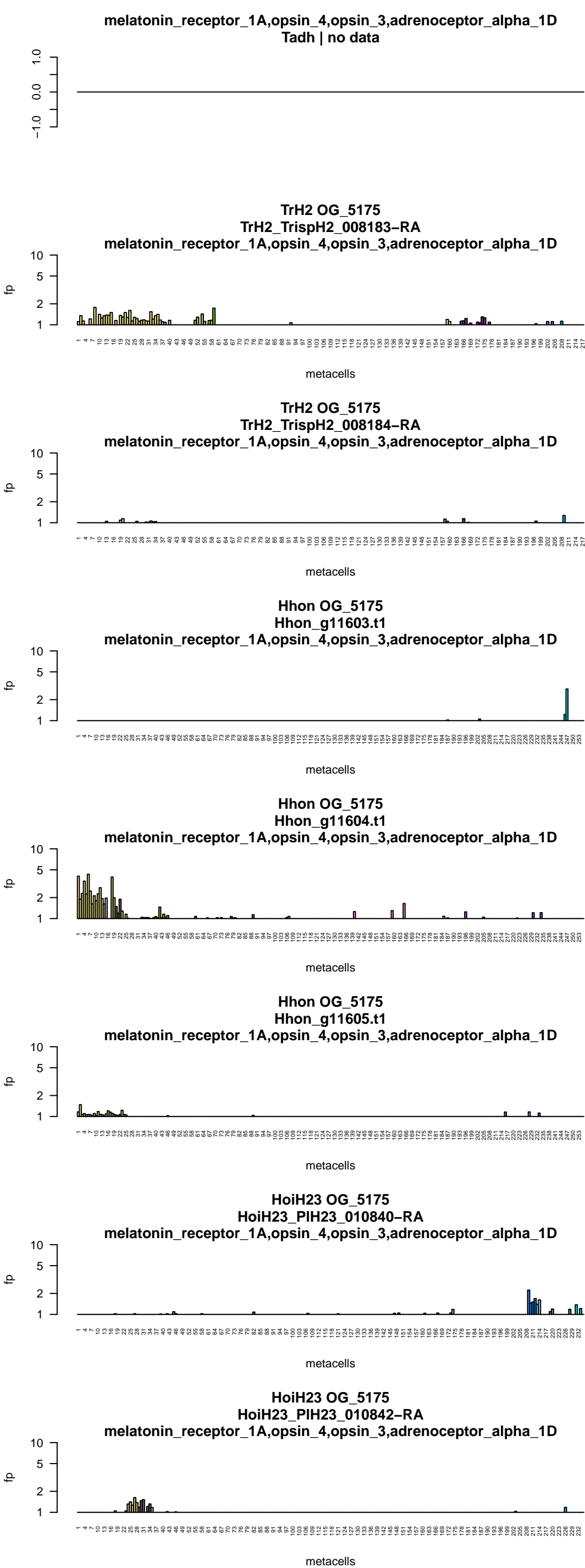


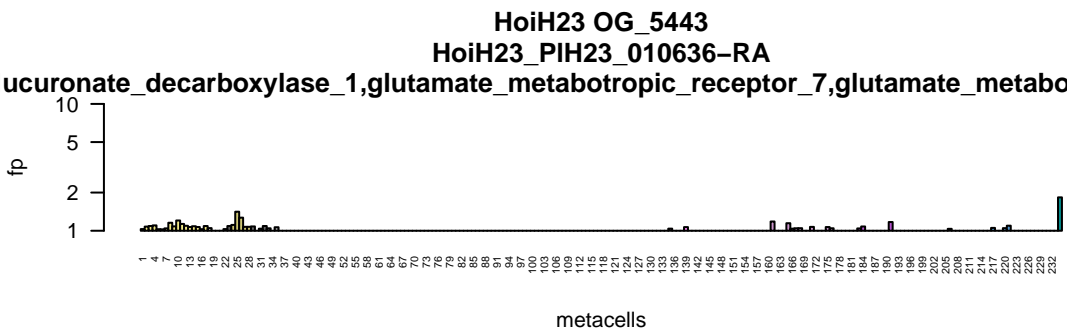
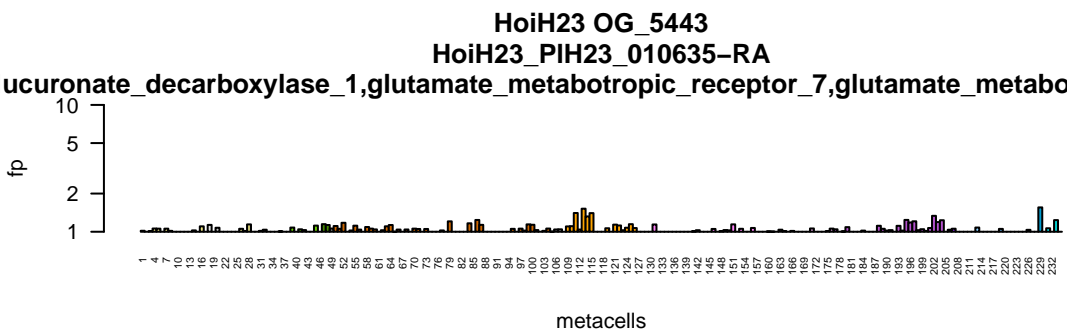
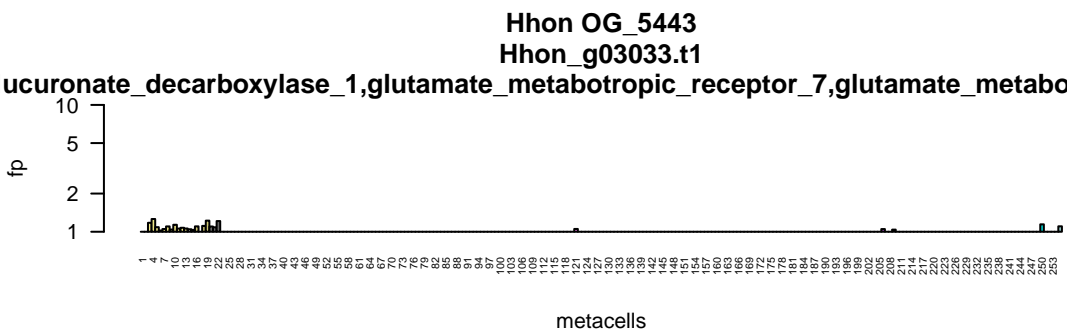
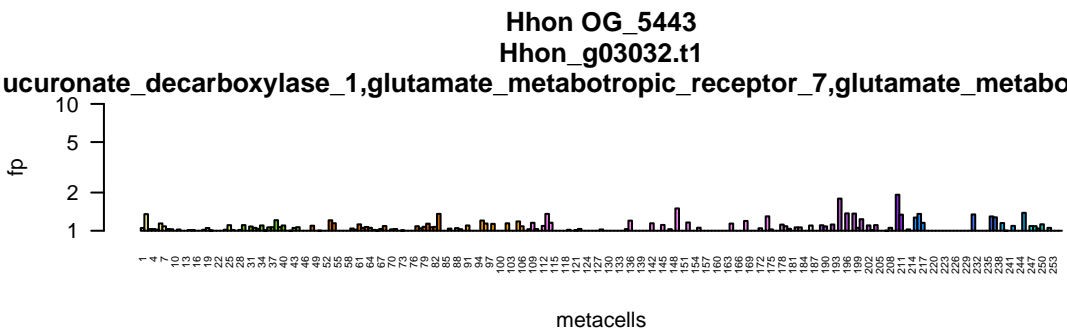
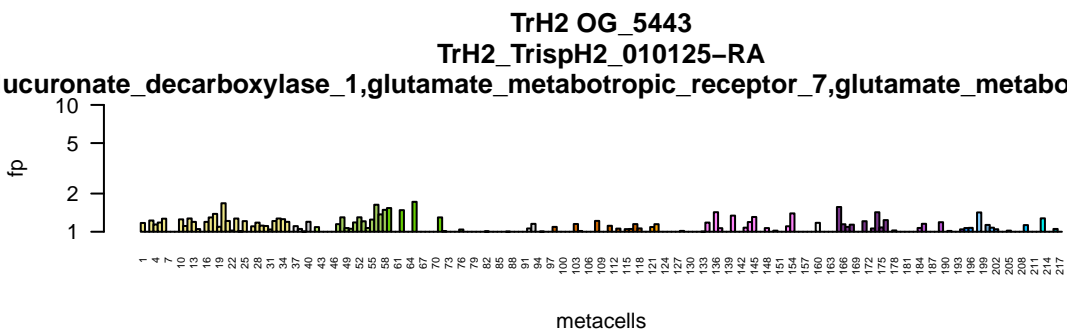
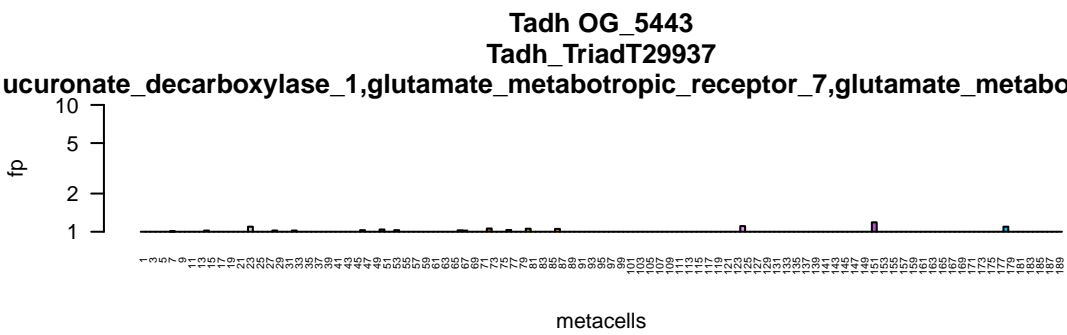
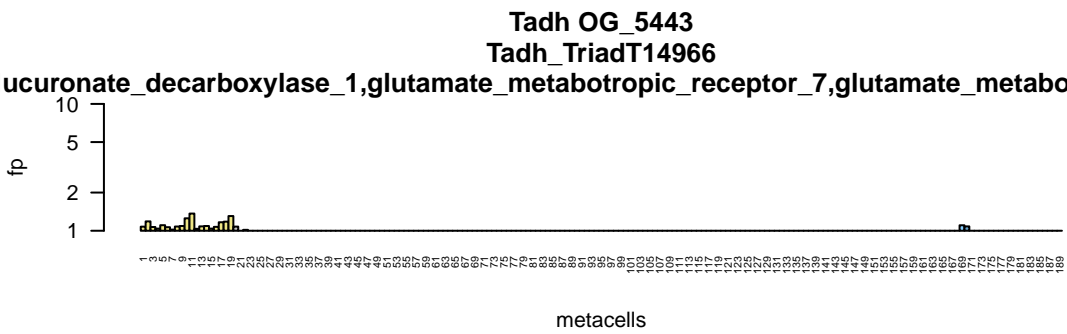


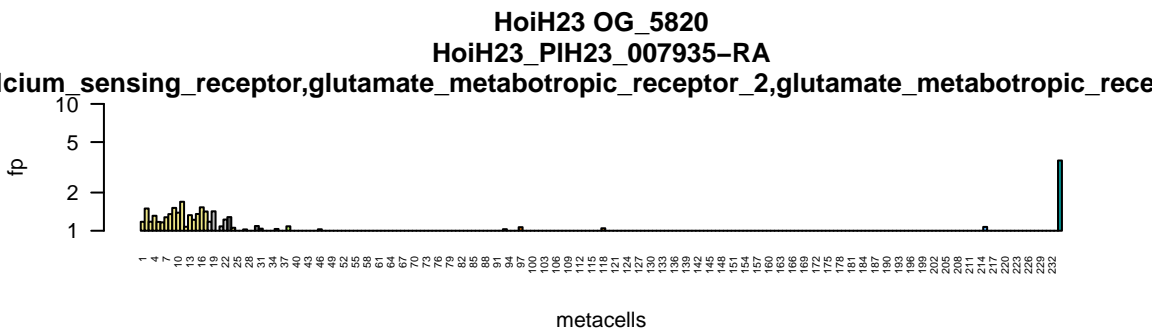
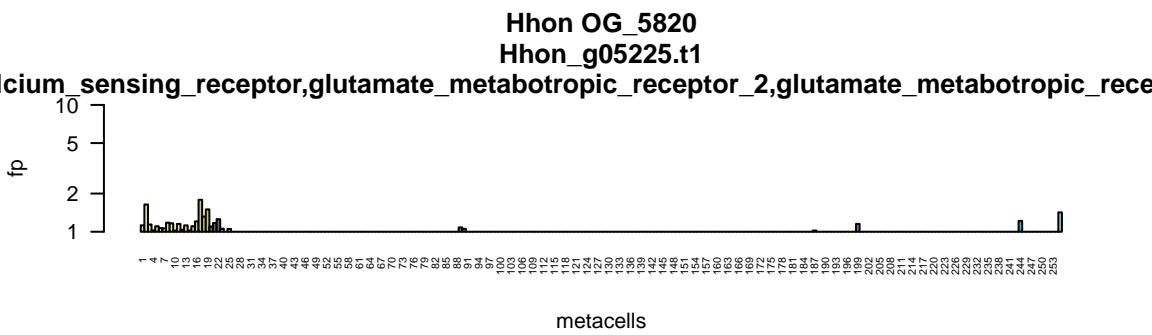
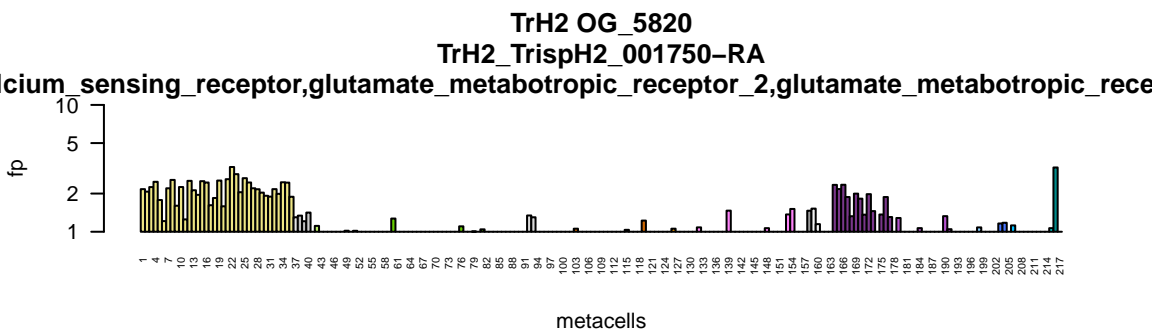
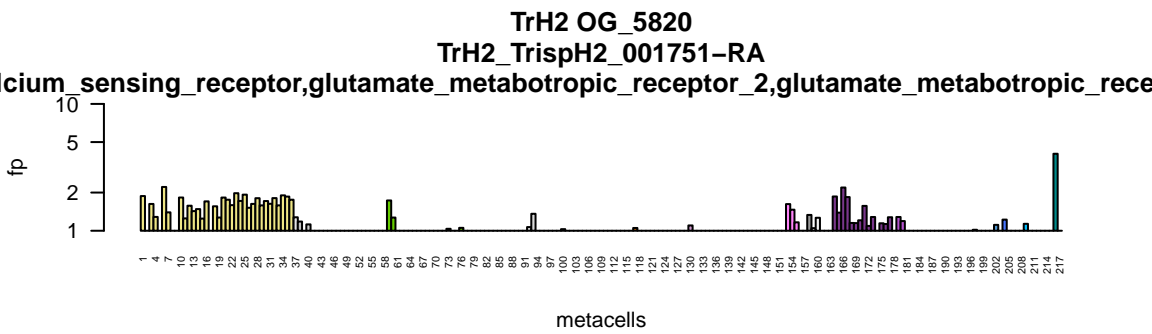
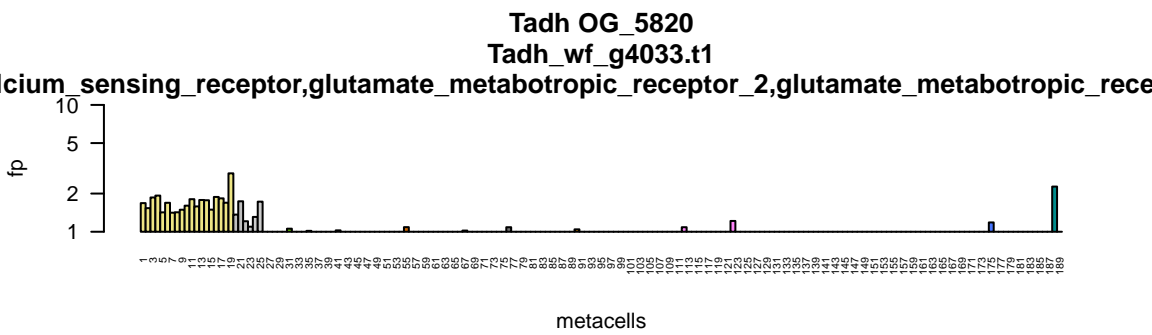
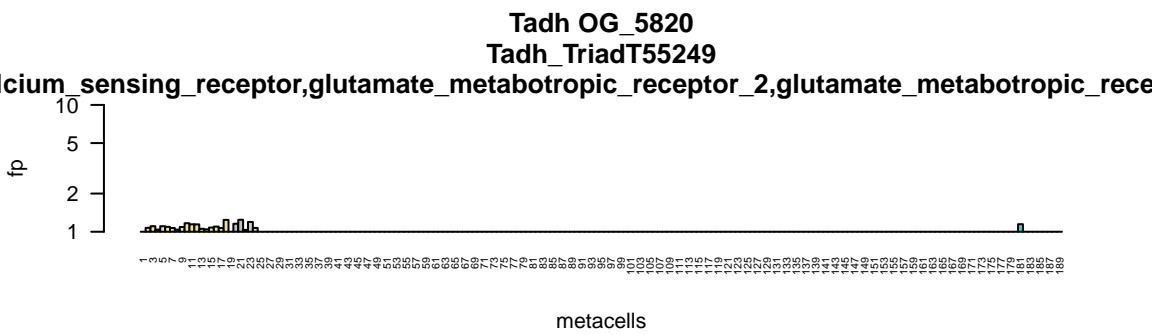
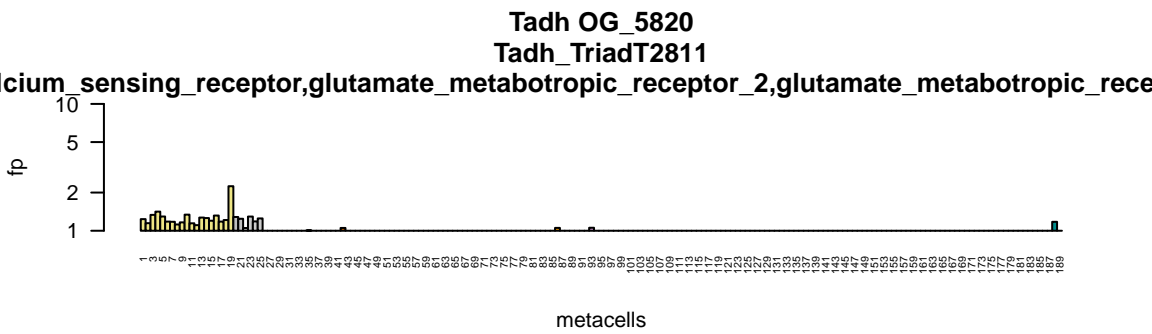












Tadh_TriadT62222

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 0 false positives, but metacell 169 has 10 false positives, and metacell 168 has 1 false positive. Other metacells have 1 or 2 false positives.

metacell	fp
1	0
3	0
7	0
11	0
13	0
17	0
21	1
23	1
27	0
31	0
35	0
37	0
41	0
45	0
47	0
51	1
55	0
57	0
61	0
65	0
69	0
71	1
75	0
77	0
81	0
85	0
89	1
91	1
95	0
99	1
101	0
105	0
109	0
113	0
115	0
119	0
123	0
125	0
129	0
133	0
135	0
139	0
143	0
145	0
149	0
153	0
155	1
159	0
163	0
165	1
167	1
169	10
173	0
177	0
179	0
183	1
187	1
189	2

TrH2_TrispH2_008367-RA

metacell	fp
1	0
4	0
7	0
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	0
34	0
37	0
40	0
43	0
46	0
49	0
52	0
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	0
154	0
157	0
160	1.5
163	1
166	0
169	0
172	0
175	0
179	0
182	0
184	0
187	0
190	0
193	0
196	0
199	2
202	10
205	0
208	0
211	0
213	1
214	0.5
217	0

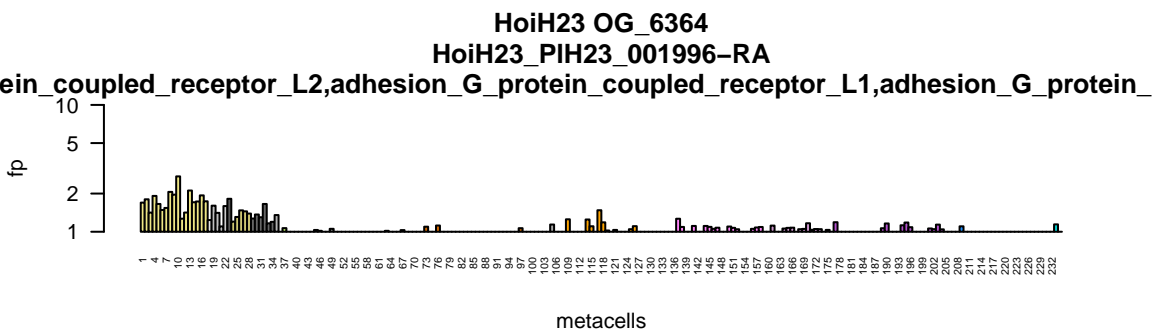
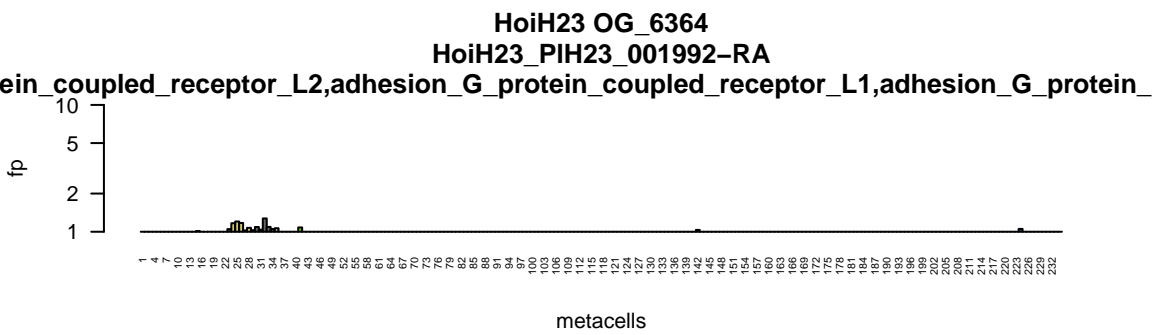
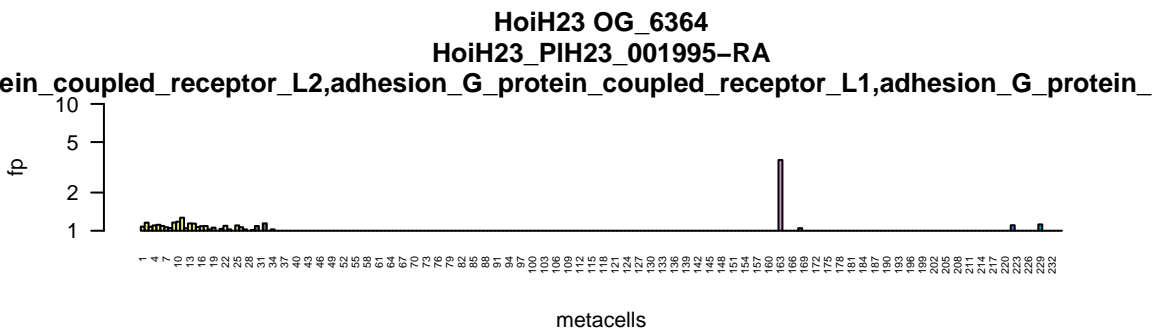
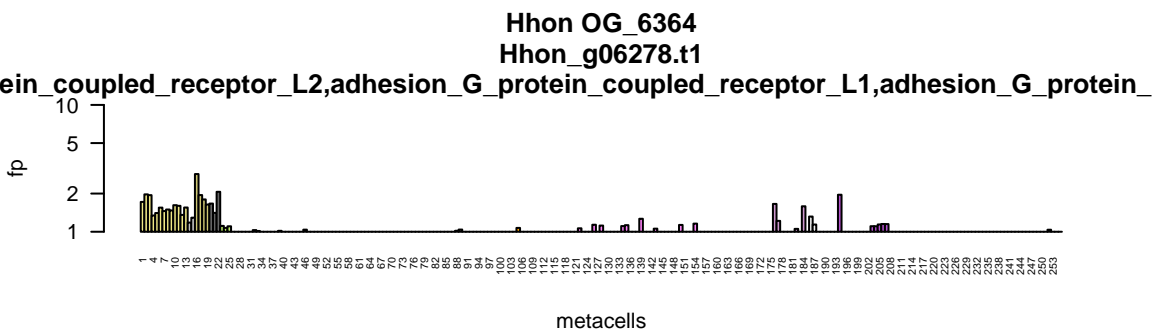
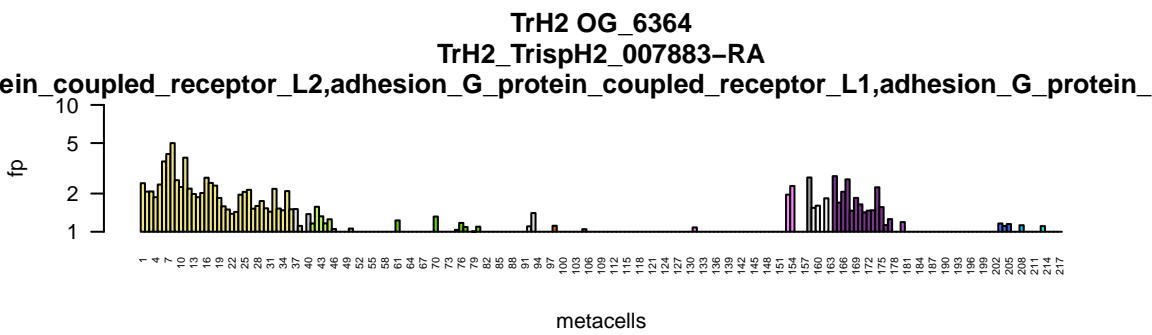
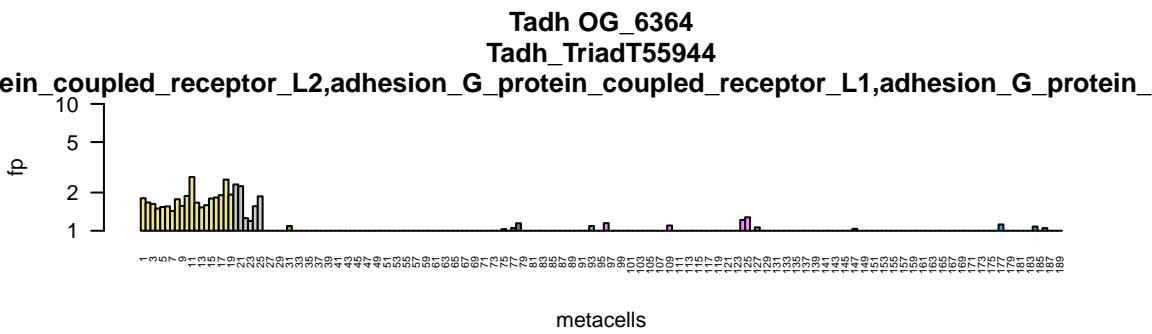
Bar chart showing the frequency of metacells (x-axis) versus the number of features (y-axis, fp). The x-axis ranges from 1 to 253, and the y-axis ranges from 1 to 10. The chart shows a distribution of feature counts across metacells, with most metacells having 1 or 2 features. A notable peak is observed at metacell 223, which has 5 features.

A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 232. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a few outliers reaching up to 2.

Bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis ranges from 1 to 232, and the y-axis ranges from 1 to 10. The distribution is highly skewed, with most metacells having 1 or 2 features. The highest frequency is 2 features, occurring for metacells 25, 26, and 27.

Bar chart showing the frequency of metacells (x-axis) for the number of features (fp, y-axis). The x-axis ranges from 1 to 232, and the y-axis ranges from 1 to 10. The distribution is highly skewed, with most metacells having 1 or 2 features. A notable peak occurs at metacell 218, which has 4 features.

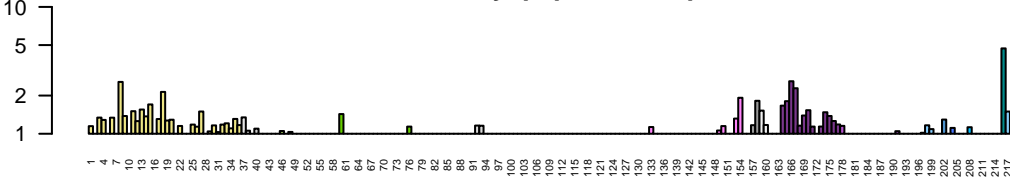
Bar chart showing the number of reads (fp) for each metacell. The y-axis is logarithmic, ranging from 1 to 10. The x-axis lists metacells from 1 to 232. Most metacells have 1 read, but metacell 214 has a significantly higher value, exceeding 10 reads.



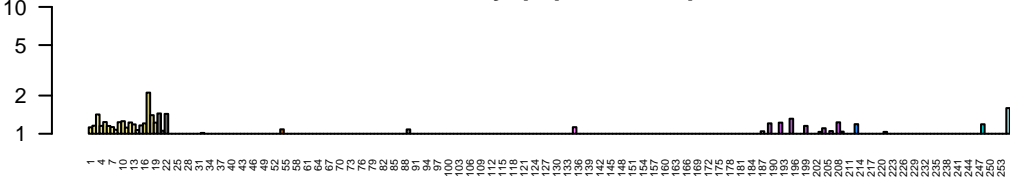
relaxin_family_peptide_receptor_1
Tadh | no data



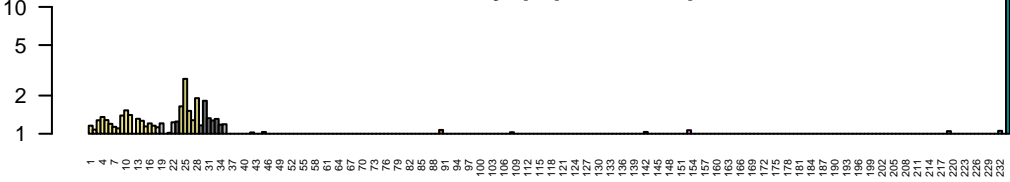
TrH2 OG_8036
TrH2_TrispH2_007969-RA
relaxin_family_peptide_receptor_1

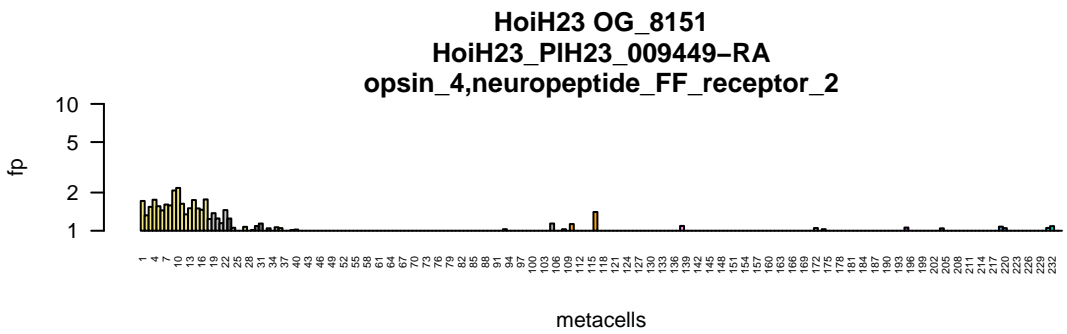
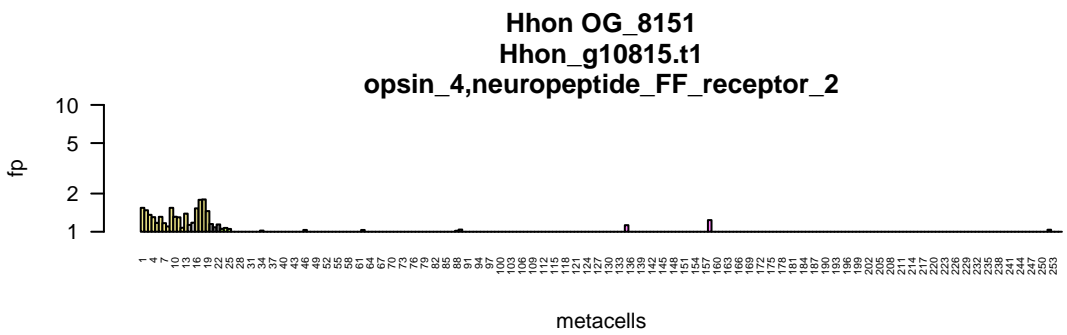
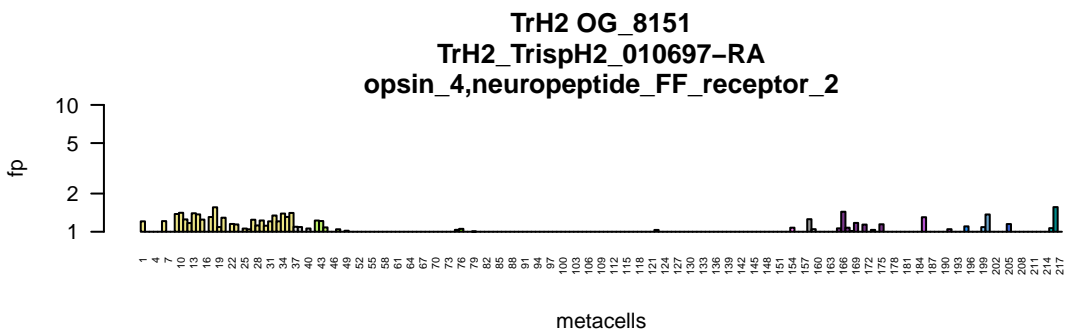
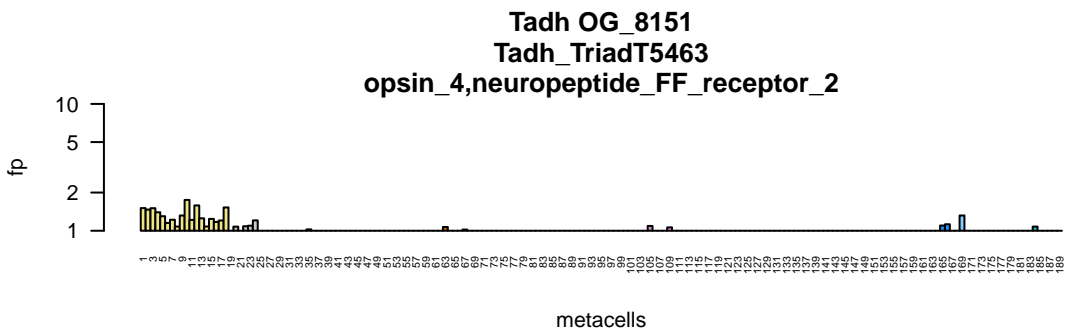


Hhon OG_8036
Hhon_g05641.t1
relaxin_family_peptide_receptor_1



HoiH23 OG_8036
HoiH23_PIH23_005185-RA
relaxin_family_peptide_receptor_1

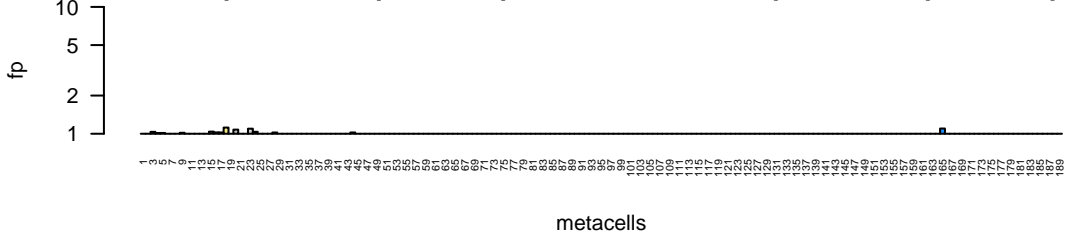




Tadh OG_8767

Tadh_hyp_clust6751.t1

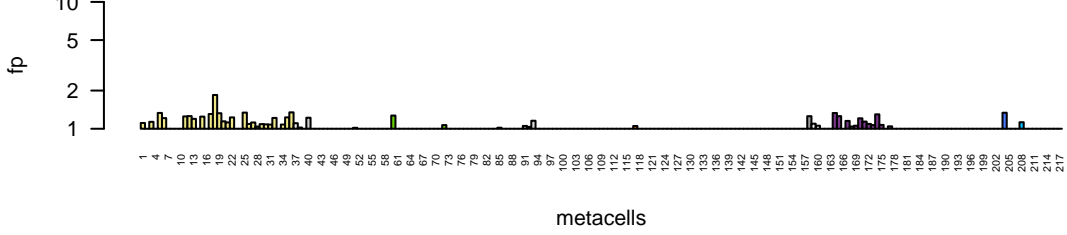
adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_coupled_receptor_L3



TrH2 OG_8767

TrH2_TrispH2_011408-RA

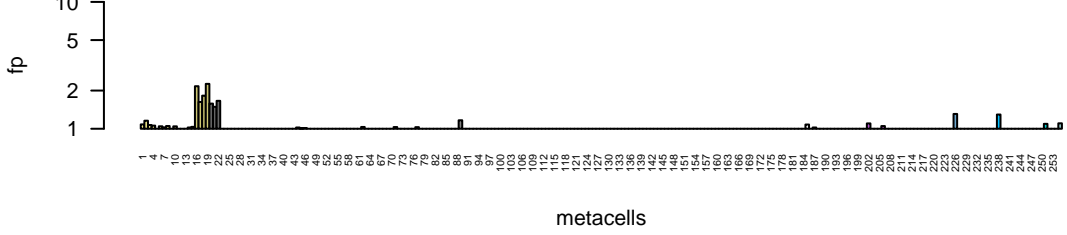
adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_coupled_receptor_L3



Hhon OG_8767

Hhon_g09539.t1

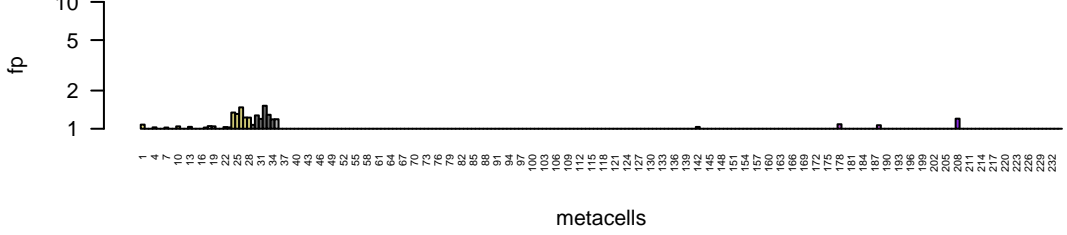
adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_coupled_receptor_L3



HoiH23 OG_8767

HoiH23_PIH23_007683-RA

adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_coupled_receptor_L3



HoiH23 OG_8767

HoiH23_PIH23_007684-RA

adhesion_G_protein_coupled_receptor_L2,adhesion_G_protein_coupled_receptor_L3

