

Bar chart showing the frequency of metacells (x-axis) versus frequency (fp, y-axis). The x-axis lists metacells from 1 to 189. The y-axis ranges from 1 to 10. The chart shows a distribution of frequencies across the metacells, with a peak around metacell 137.

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. The highest counts are for metacells 175, 176, and 179, which 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	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 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. The bars are colored in a repeating pattern of black, purple, and green. Most metacells have a false positive count of 1, with a few having counts of 2 or 3.

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists 189 metacells. 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.

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
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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

A bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows a distribution of metacell frequencies, with most metacells having a frequency of 1, and a few having higher frequencies up to 5.

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

metacell	fp
1	1
4	1
10	1
13	1
16	1
22	1
23	1
25	1
28	1
31	1
32	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
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 4.

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

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

metacells	fp
1	1
4	1
10	1
13	1
16	1.5
22	1
25	1
28	1
31	1
32	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1.5
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1.5
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1
202	2.5
205	1.5
208	6
211	4
214	4
217	1

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

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

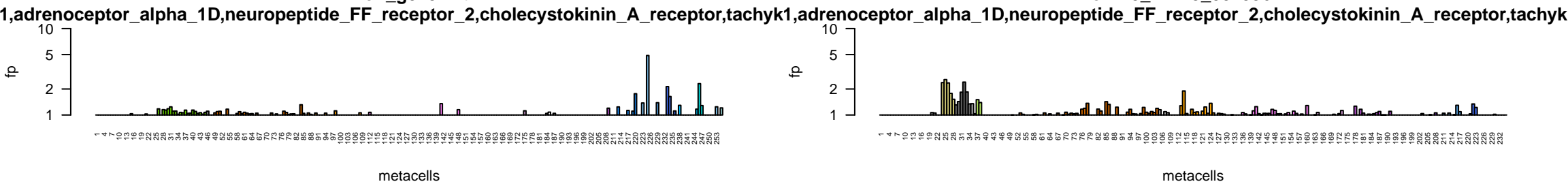
Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists metacells from 1 to 217. Most metacells have a false positive count of 1, with a few having counts of 2 or 3. Metacells 205 and 206 show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

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

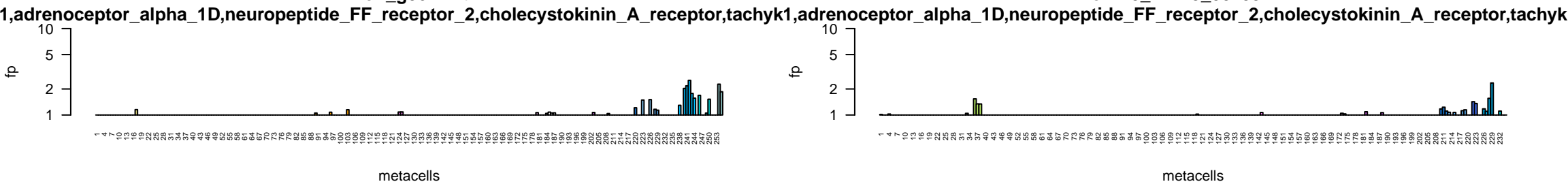
Bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and the y-axis is labeled 'fp'. The x-axis has 253 categories, and the y-axis ranges from 1 to 10. Most categories have a frequency of 1, with a small cluster of higher frequencies around category 241.

metacells	fp
1	1
7	1
10	1
11	1
18	1
19	1
25	1
28	1
34	1
35	1
43	1
46	1
52	1
55	1
61	1
64	1
70	1
76	1
79	1
85	1
89	1
94	1
97	1
103	1
108	1
112	1
115	1
121	1
127	1
130	1
139	1
145	1
148	1
154	1
157	1
166	1
172	1
175	1
181	1
187	1
190	1
193	1
199	1
205	1
208	1
214	1
217	1
223	1
226	1
232	1
235	1
241	2
241	3
241	4
241	5
247	1
250	1
253	1

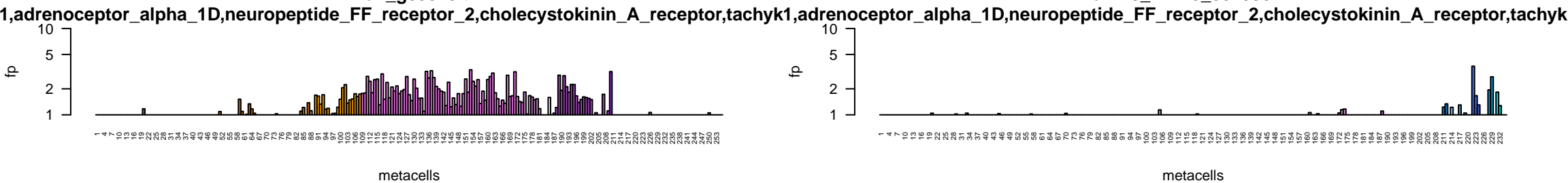
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Hhon\_g04574.t1



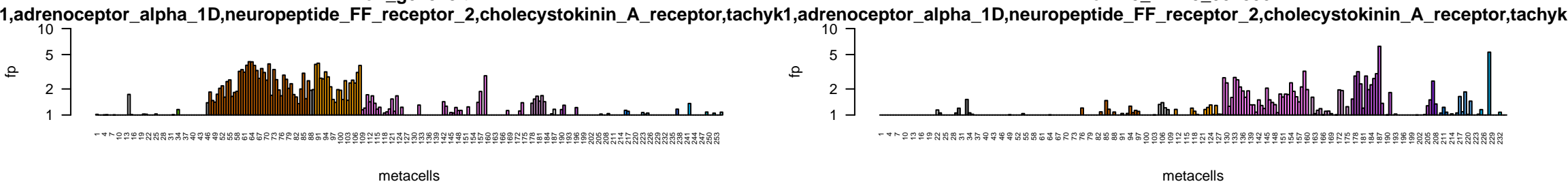
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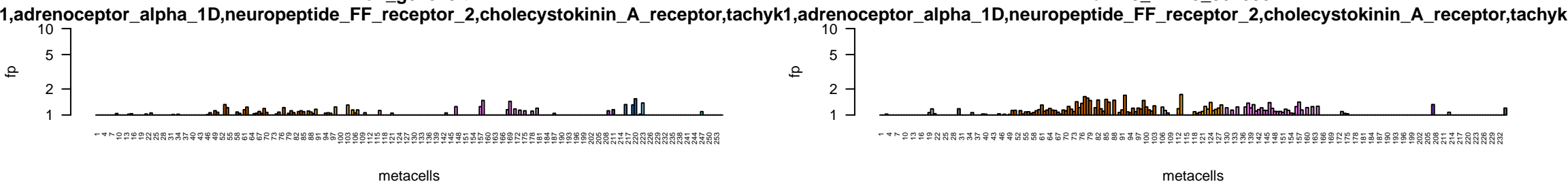
Hhon OG\_5725  
Hhon\_g09573.t1



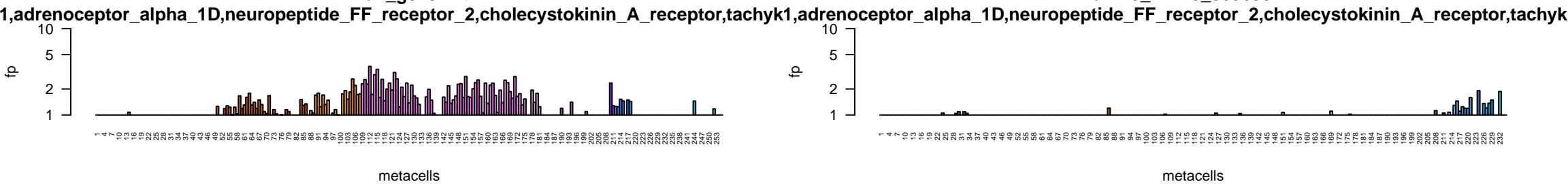
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Hhon\_g07528.t1



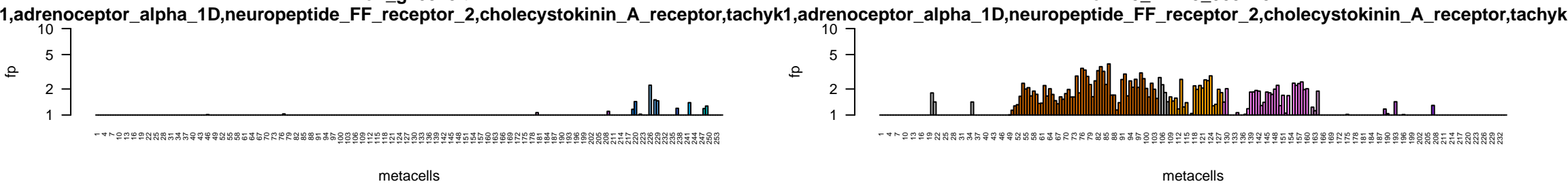
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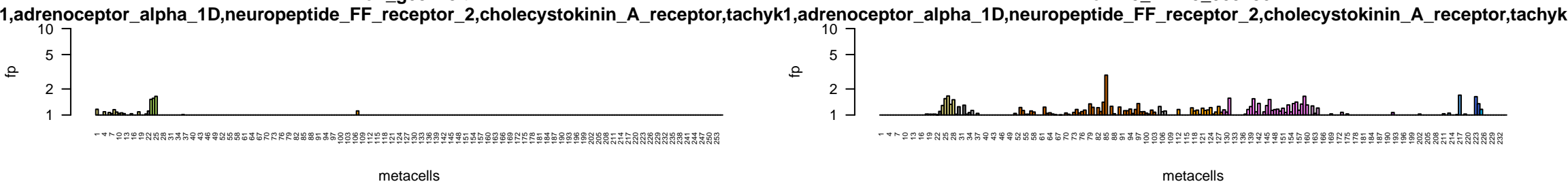
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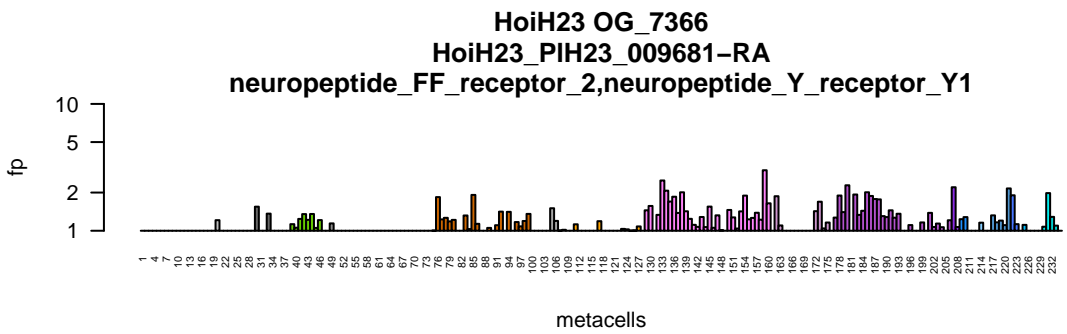
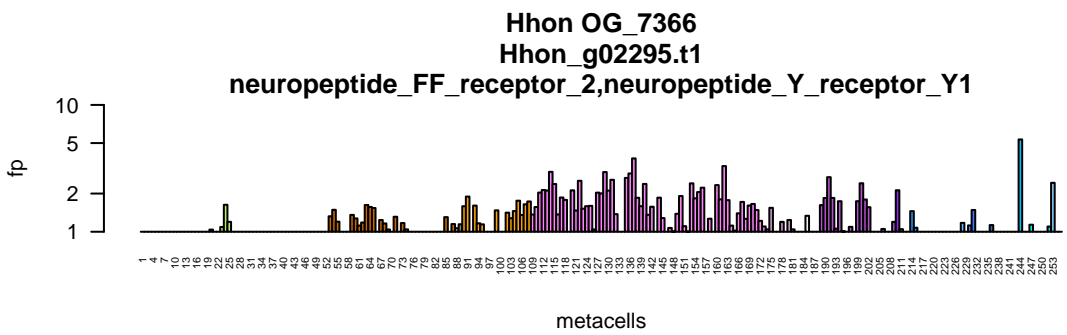
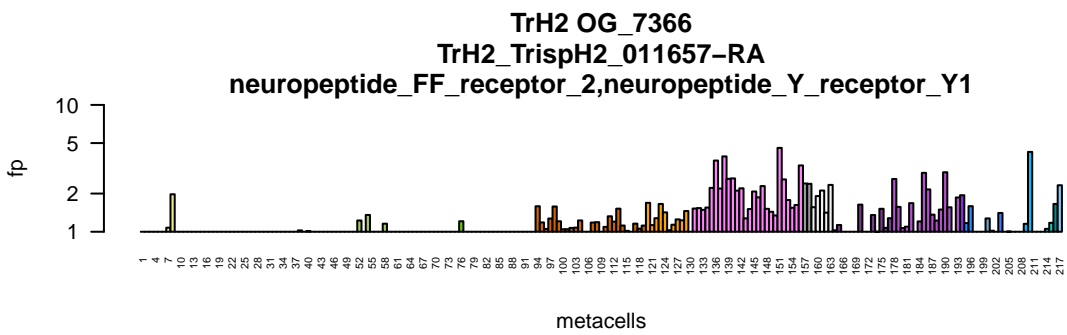
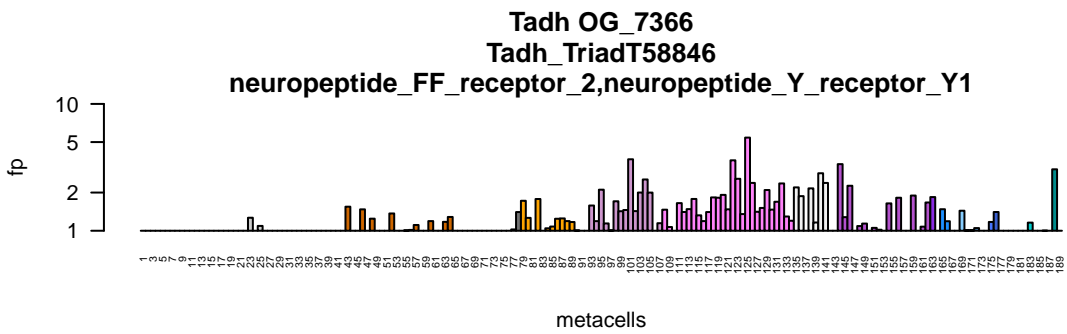


Hhon OG\_5725  
Hhon\_g10923.t1



Hhon OG\_5725  
Hhon\_g08418.t1

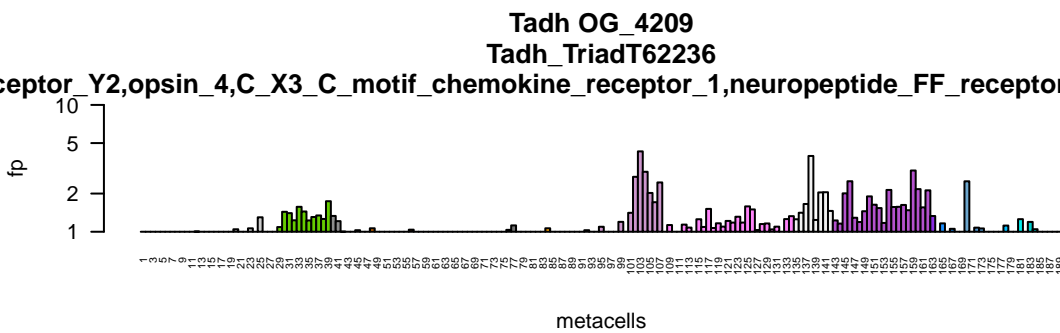




ceptor\_Y2,opsin\_4,C\_X3\_C\_motif\_chemokine\_receptor\_1,neuropeptide\_FF\_receptor

fp

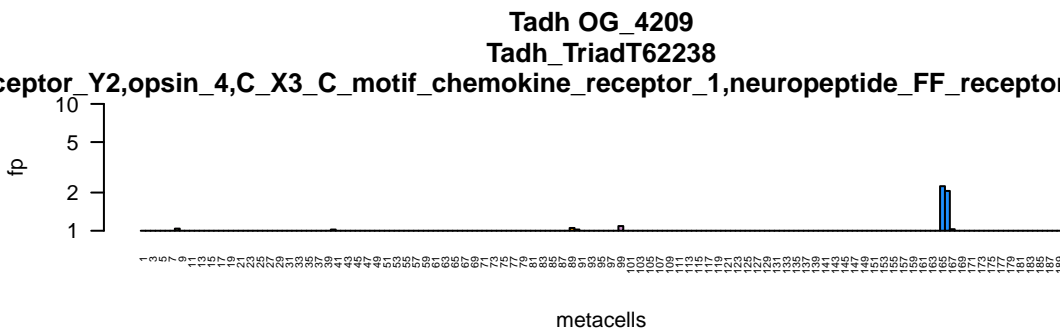
metacells



ceptor\_Y2,opsin\_4,C\_X3\_C\_motif\_chemokine\_receptor\_1,neuropeptide\_FF\_receptor

fp

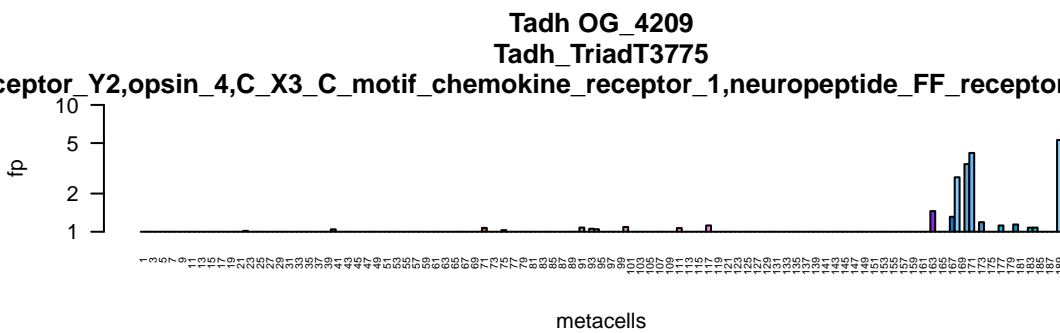
metacells



ceptor\_Y2,opsin\_4,C\_X3\_C\_motif\_chemokine\_receptor\_1,neuropeptide\_FF\_receptor

Motif	fp
1	1
2	1
3	1
4	1
5	1
6	1
7	1
8	1
9	1
10	1
11	1
12	1
13	1
14	1
15	1
16	2
17	1
18	1
19	1
20	1.5
21	1.2
22	1.1
23	1.0
24	0.9
25	0.8
26	0.7
27	0.6
28	0.5
29	0.4
30	0.3
31	0.2
32	0.1
33	0.1
34	0.1
35	0.1
36	0.1
37	0.1
38	0.1
39	0.1
40	0.1

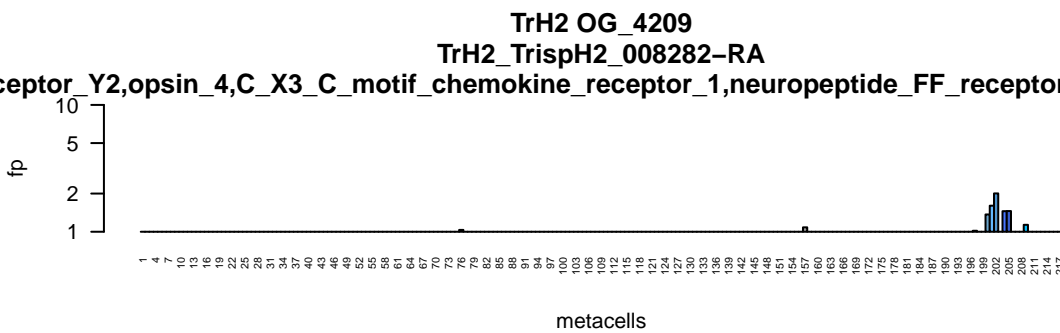
metacells



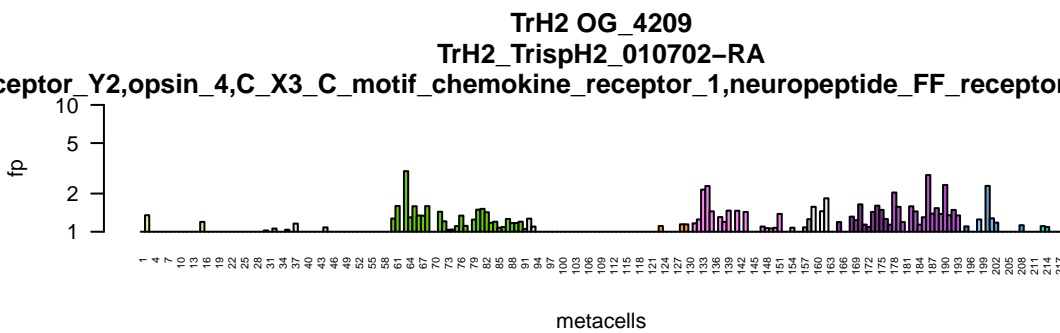
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fp

metacells



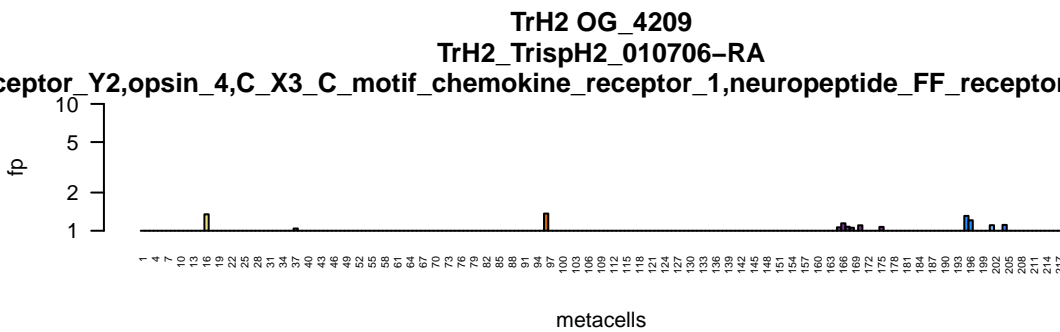
ceptor\_Y2,opsin\_4,C\_X3\_C\_motif\_chemokine\_receptor\_1,neuropeptide\_FF\_receptor



ceptor\_Y2,opsin\_4,C\_X3\_C\_motif\_chemokine\_receptor\_1,neuropeptide\_FF\_receptor

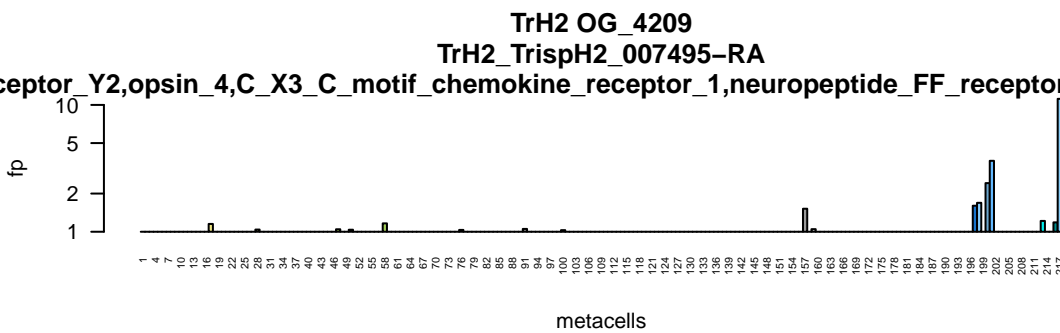
fp

metacells



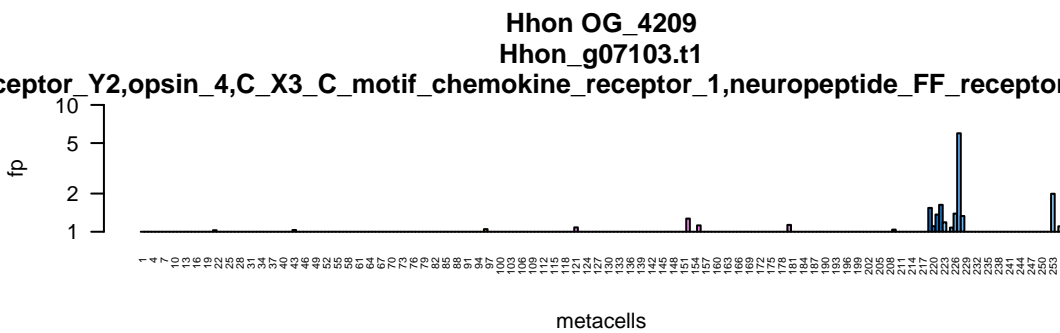
ceptor\_Y2,opsin\_4,C\_X3\_C\_motif\_chemokine\_receptor\_1,neuropeptide\_FF\_receptor

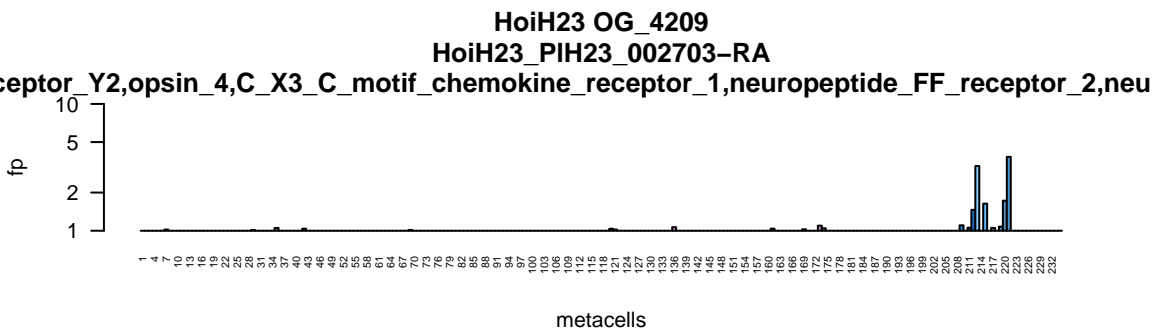
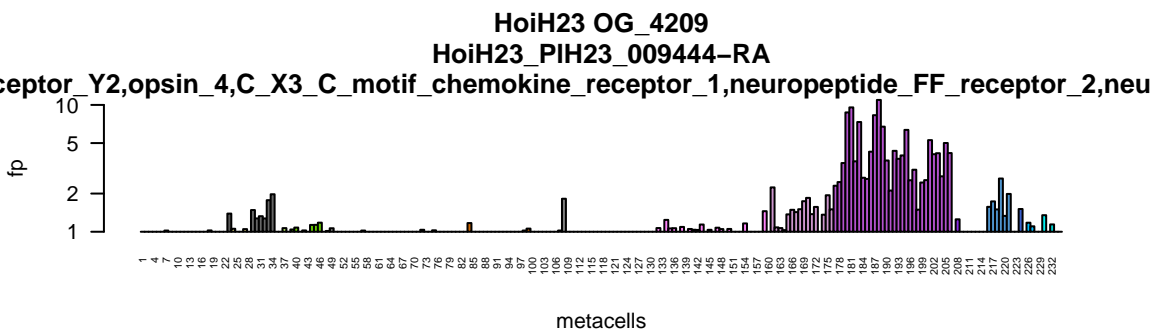
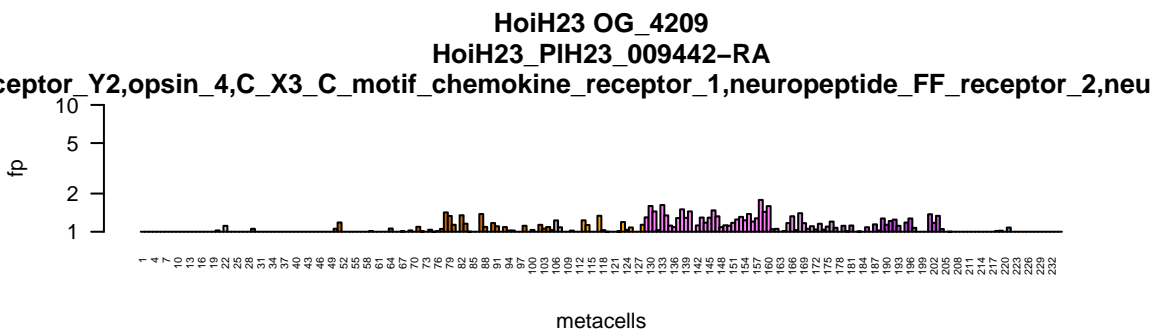
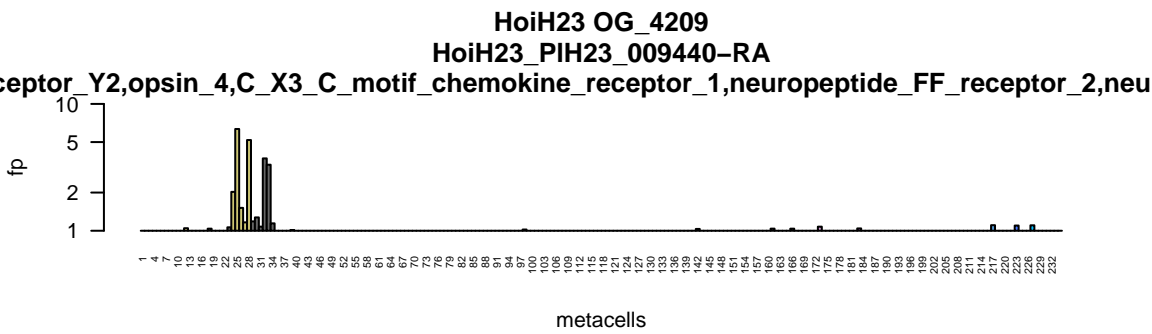
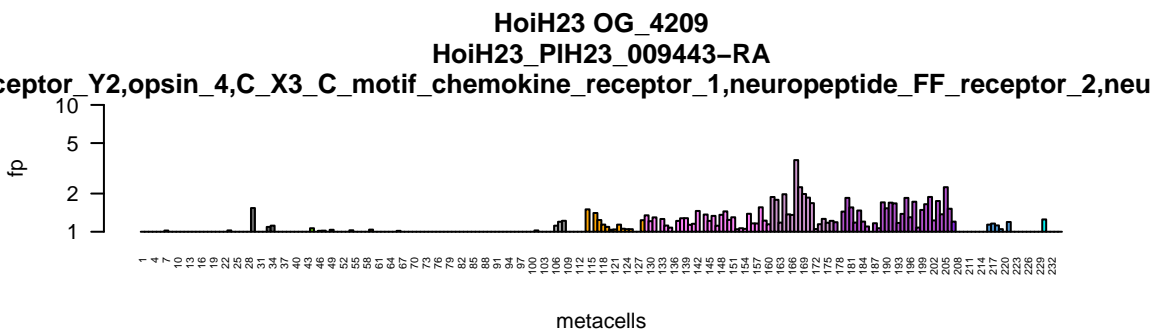
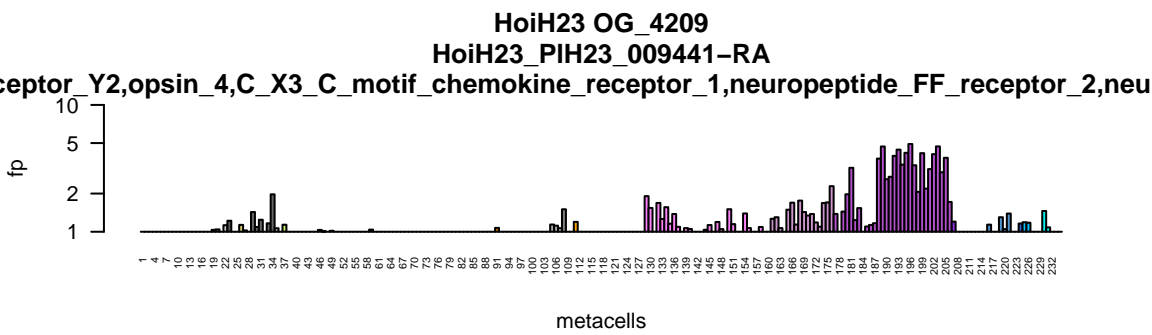
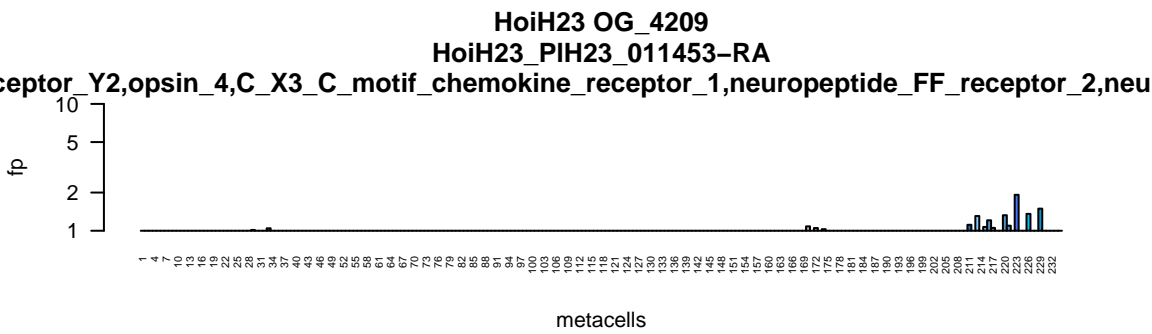
metacells	fp
1	1
4	1
10	1
13	1
16	1
19	1
21	1
25	1
28	1
31	1
31	1
37	1
40	1
43	1
46	1
52	1
55	1
59	1
61	1
64	1
67	1
70	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
103	1
106	1
109	1
111	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
169	1
172	1
175	1
181	1
184	1
187	1
193	1
196	1
199	1.2
202	1.5
208	1
211	1
214	1.8

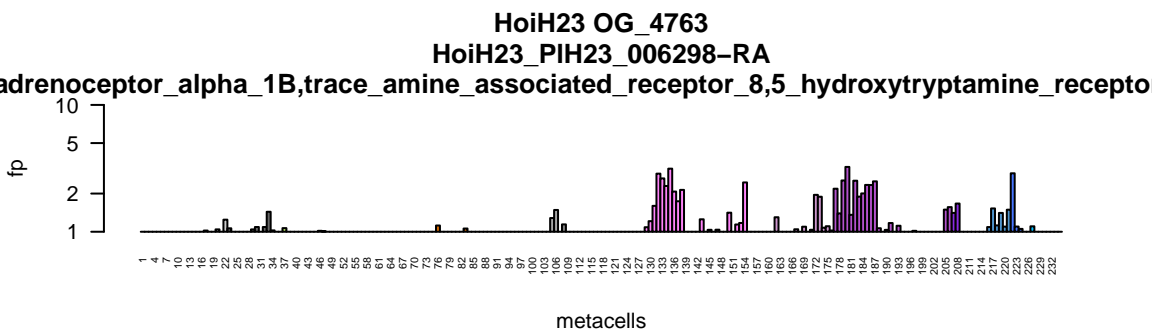
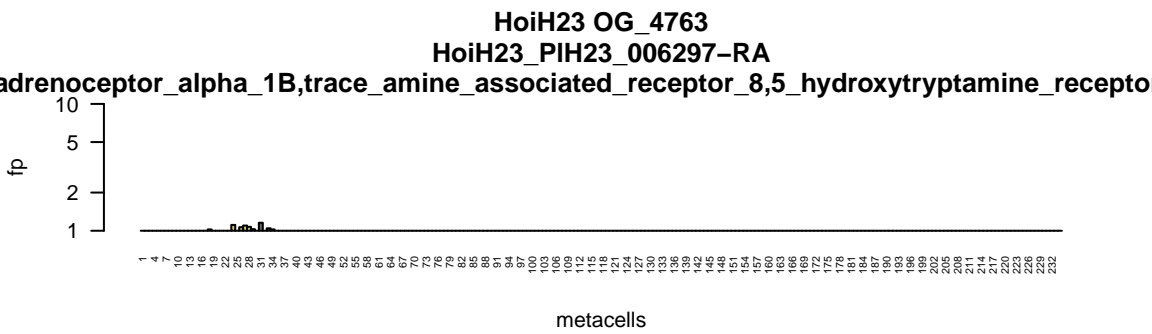
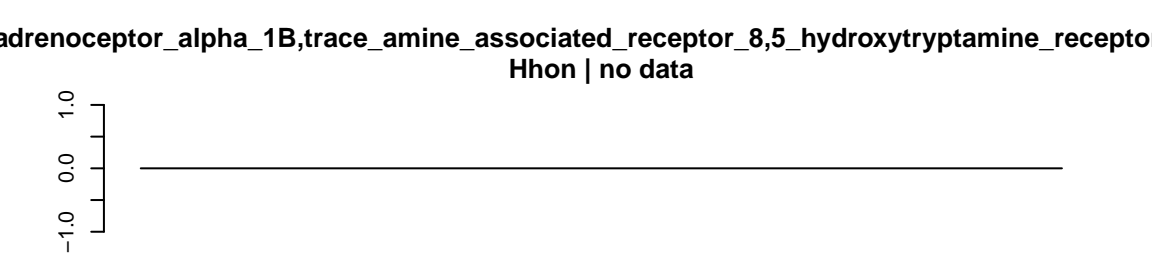
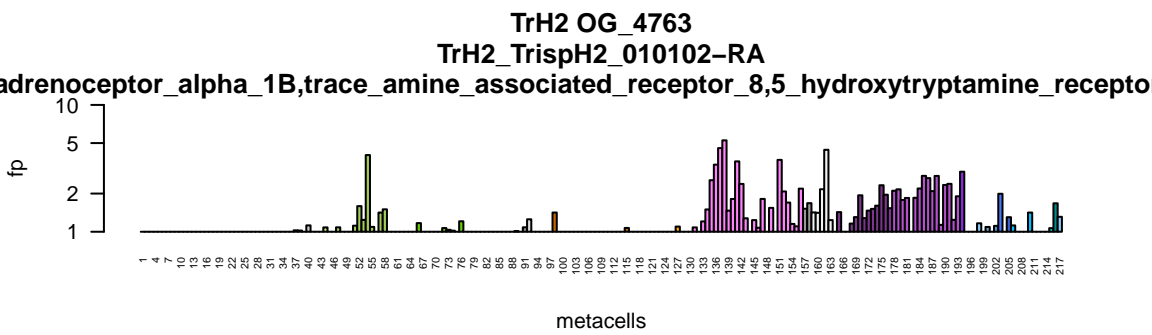
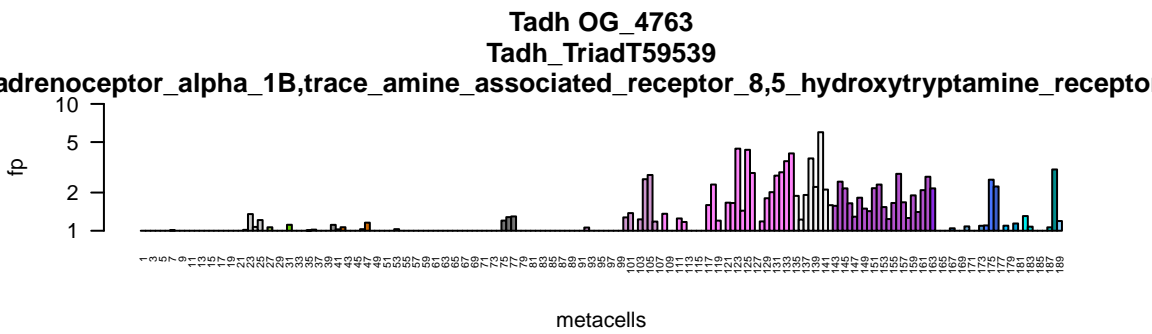


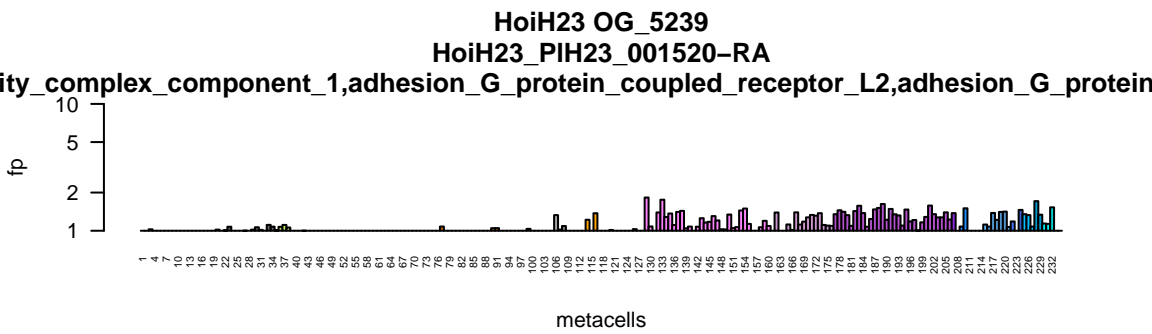
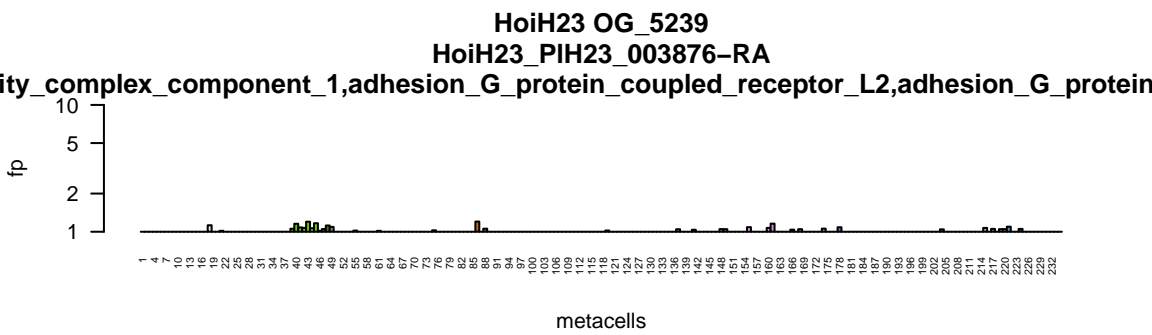
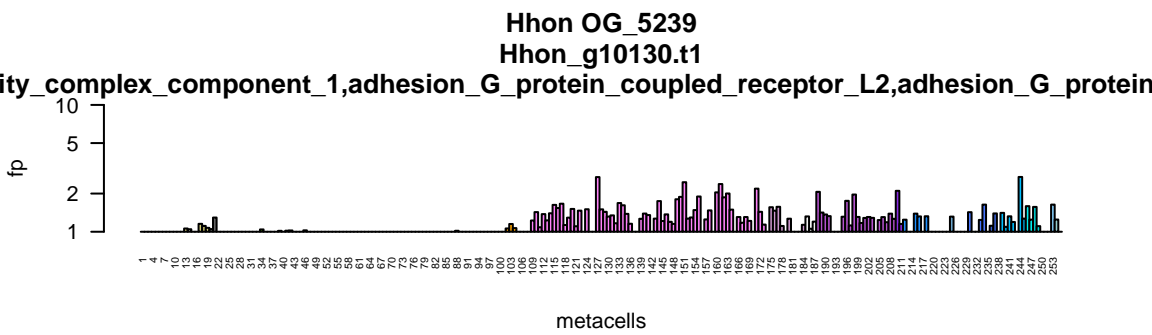
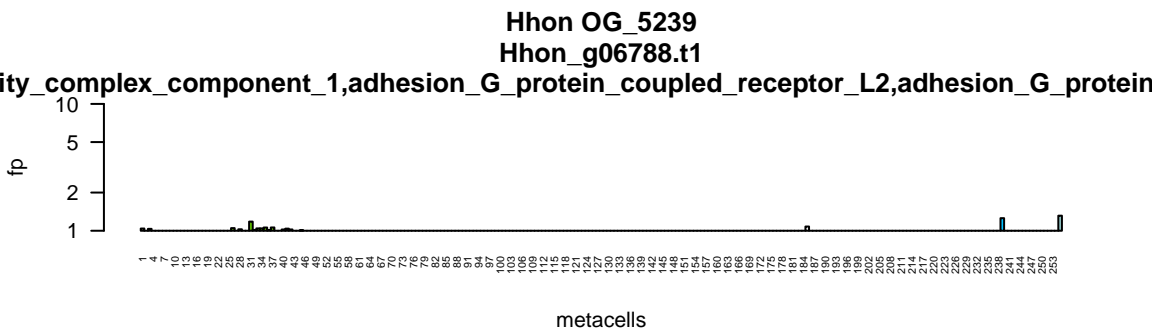
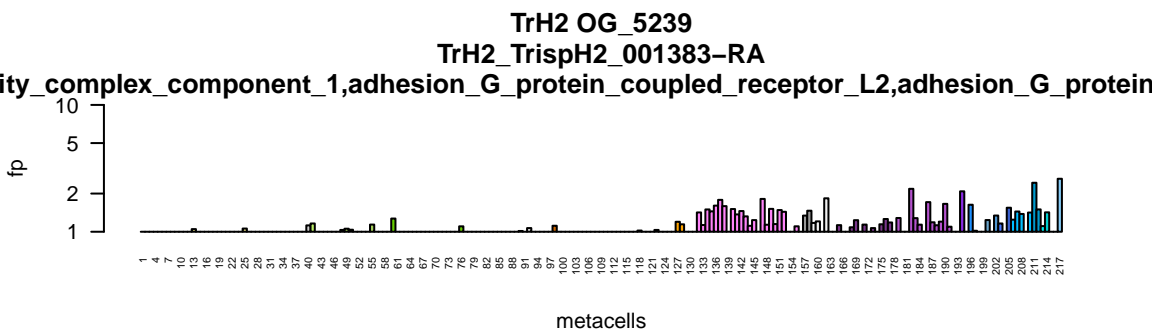
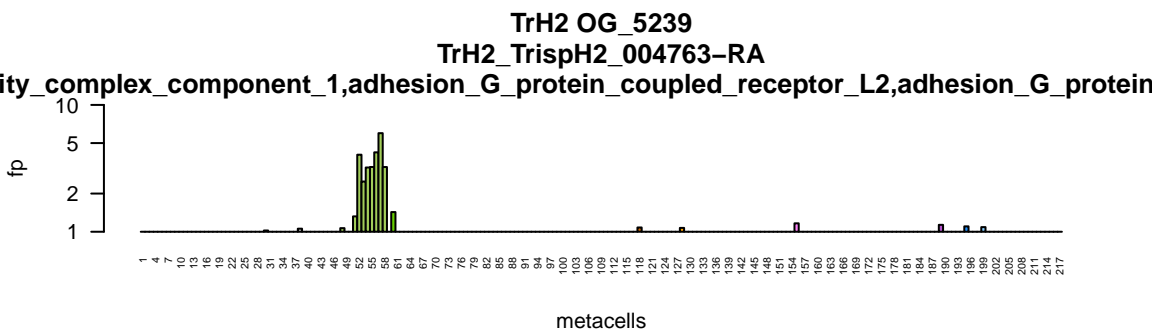
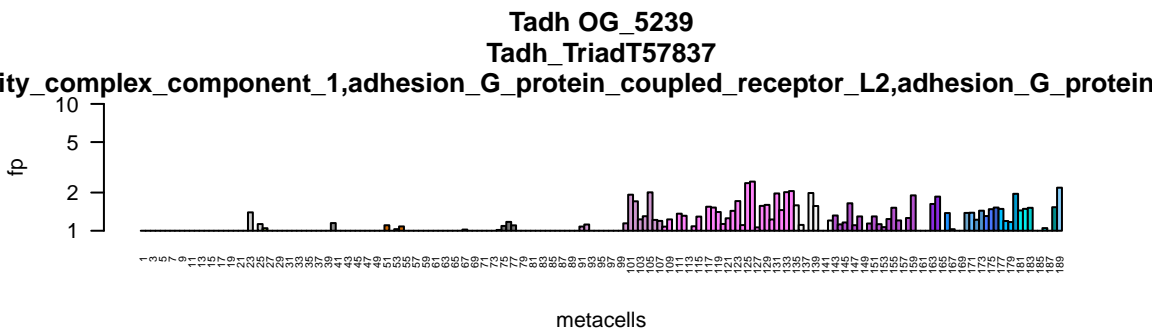
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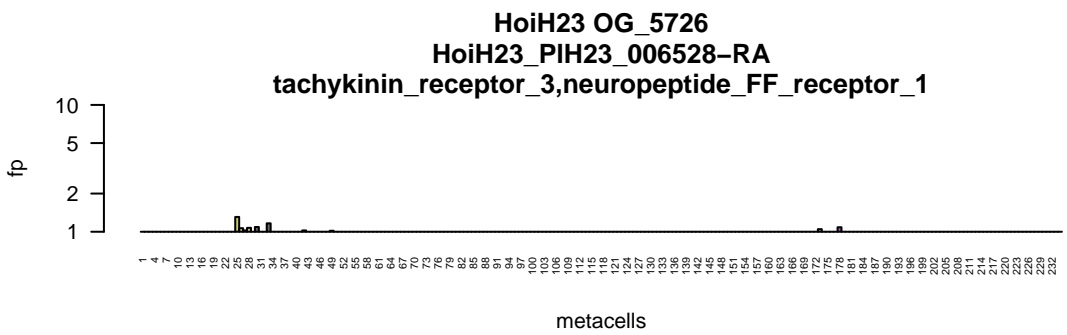
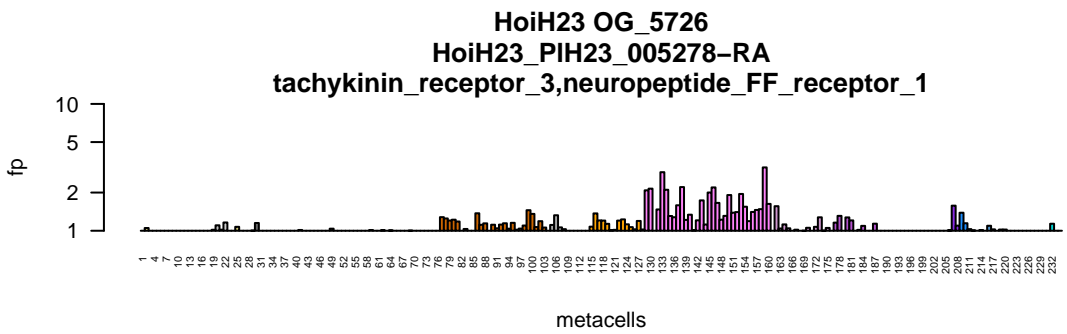
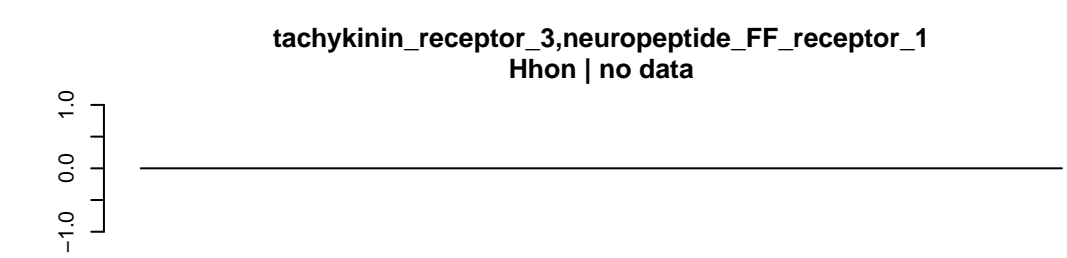
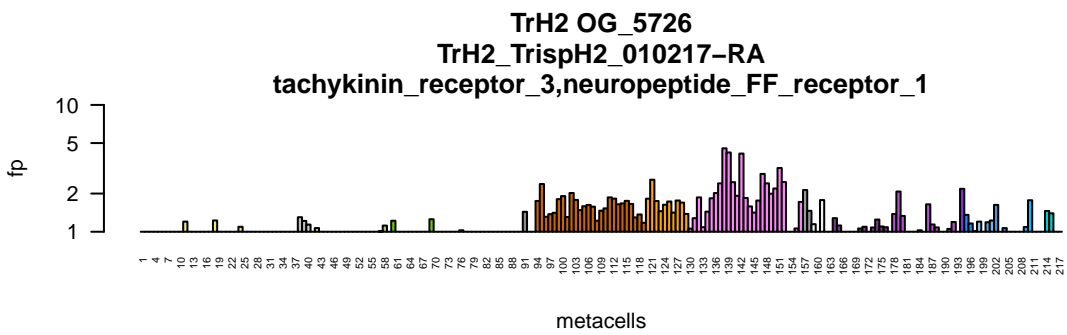
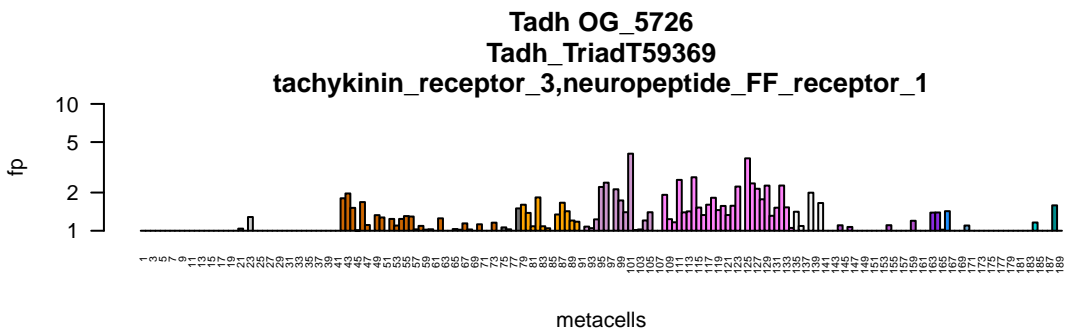
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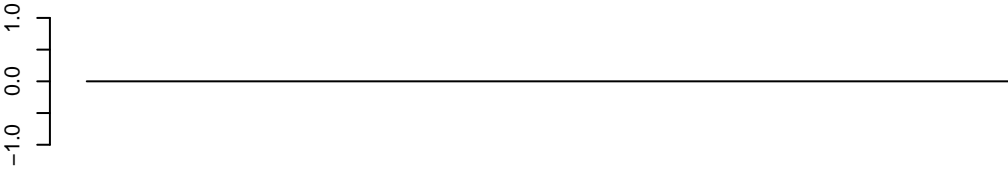




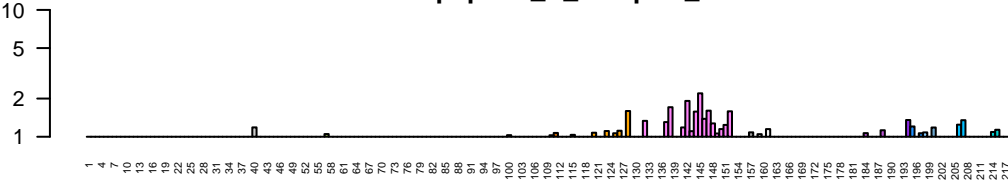




neuropeptide\_Y\_receptor\_Y2  
Tadh | no data

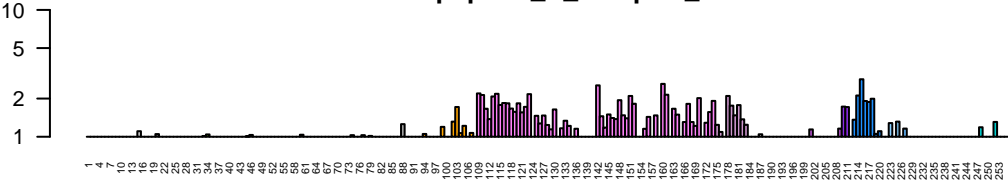


TrH2 OG\_7764  
TrH2\_TrispH2\_010449-RA  
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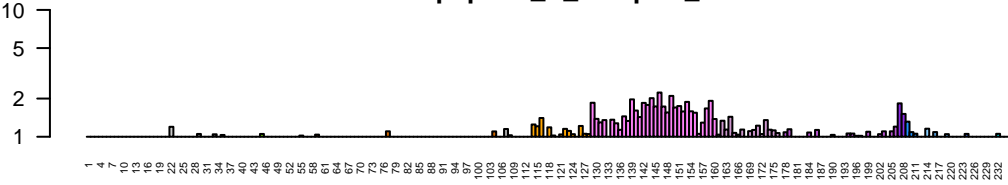
metacells

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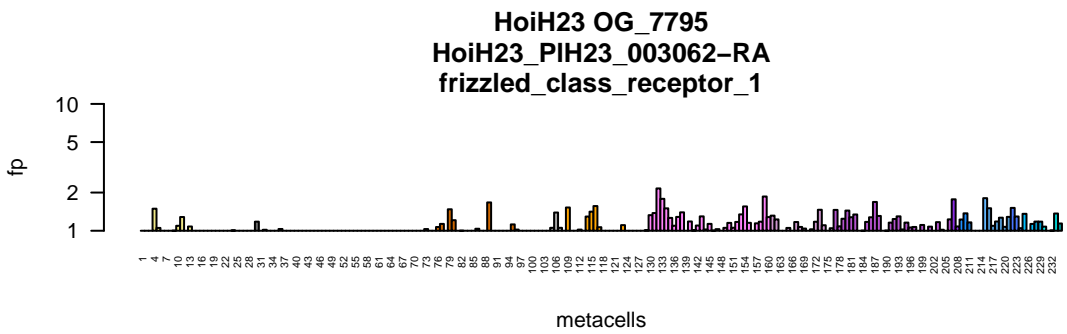
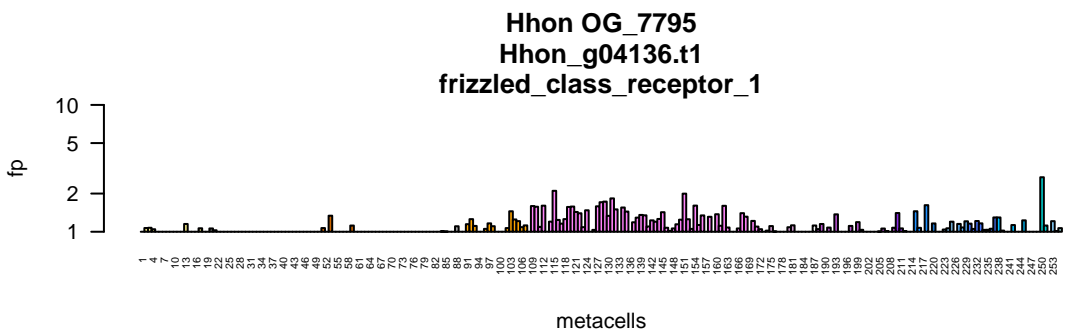
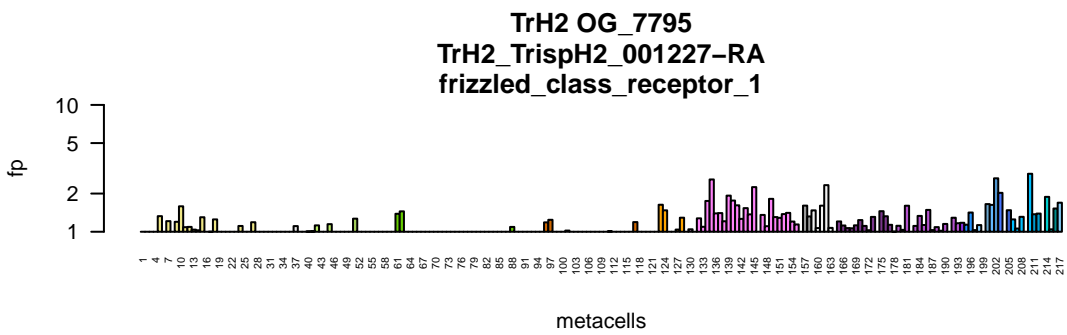
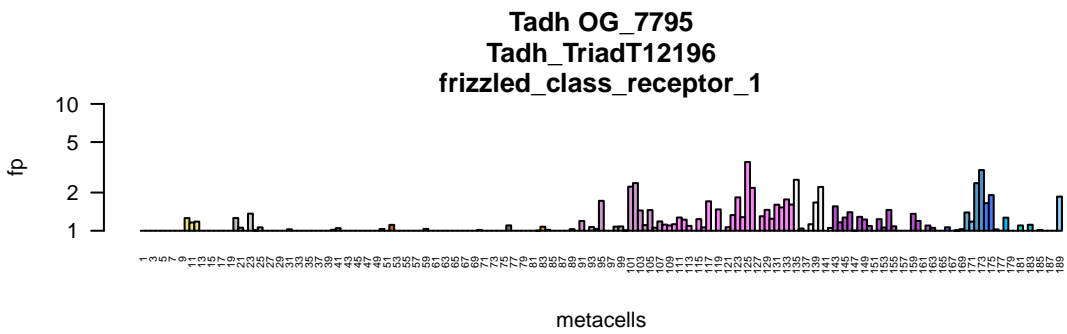


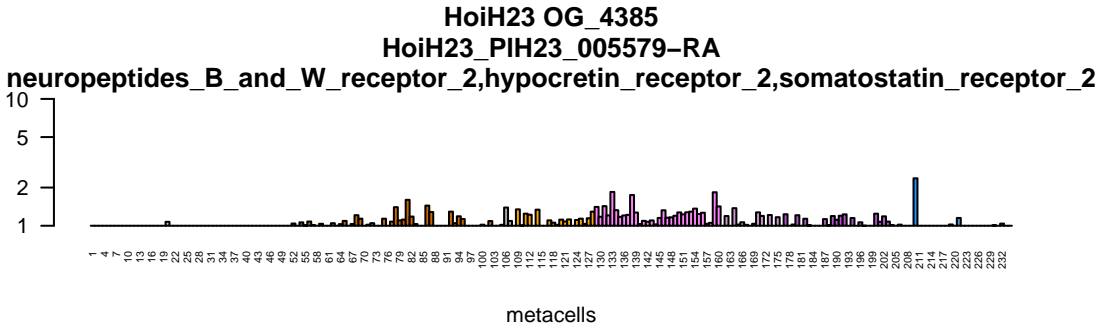
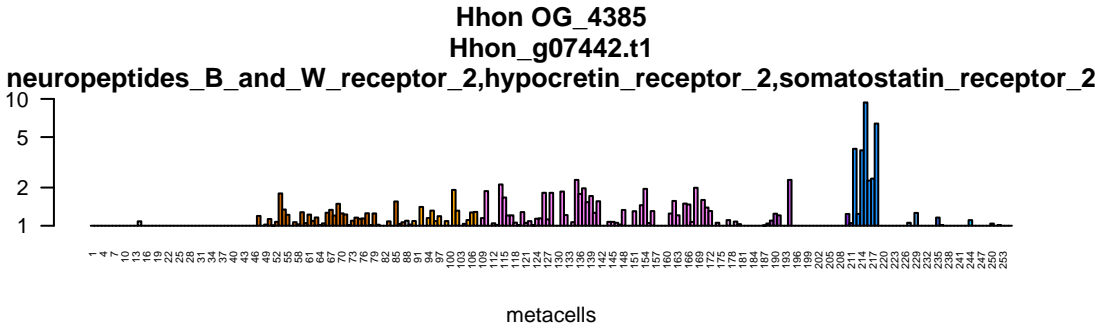
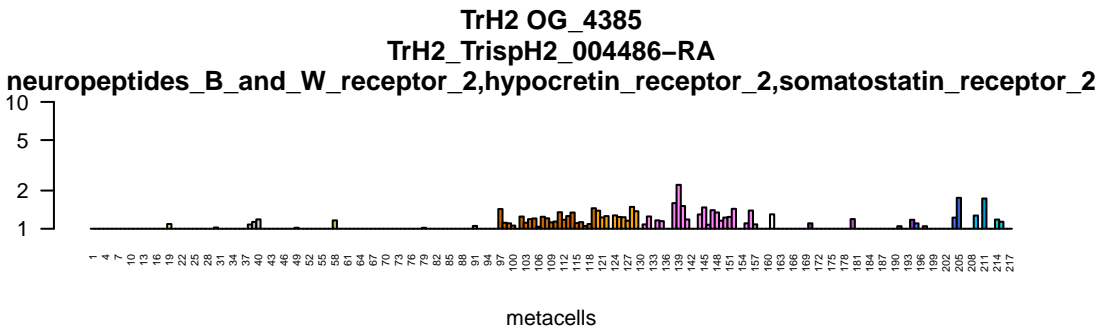
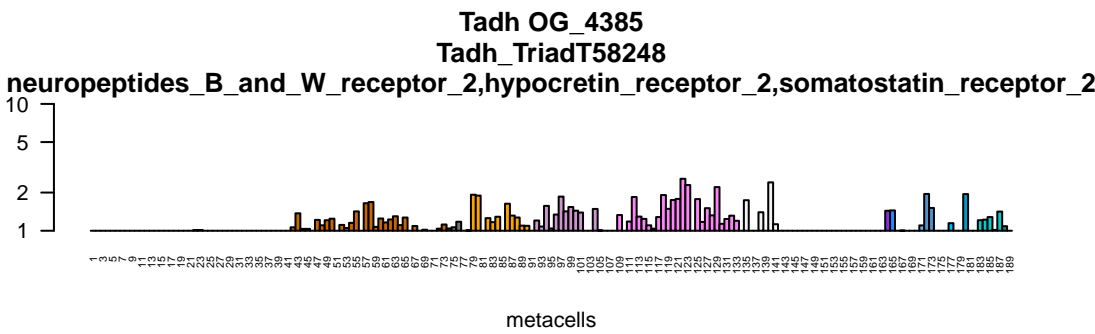
metacells

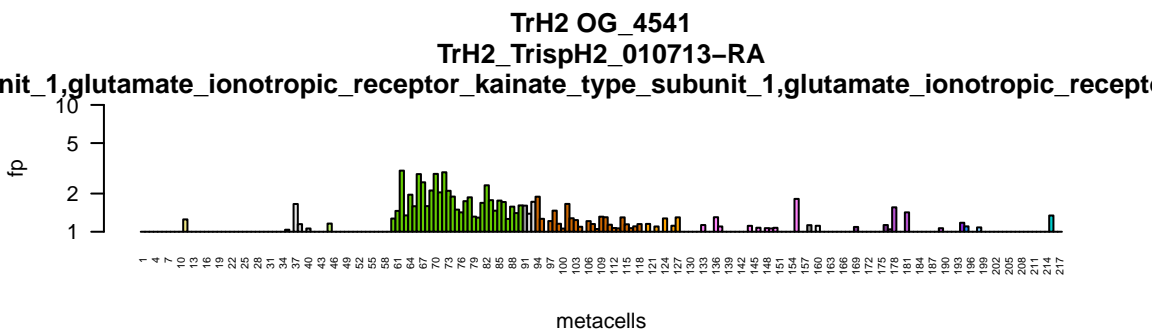
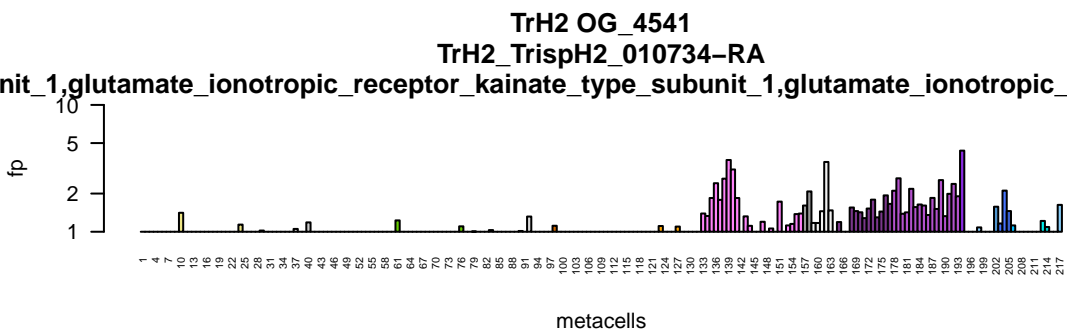
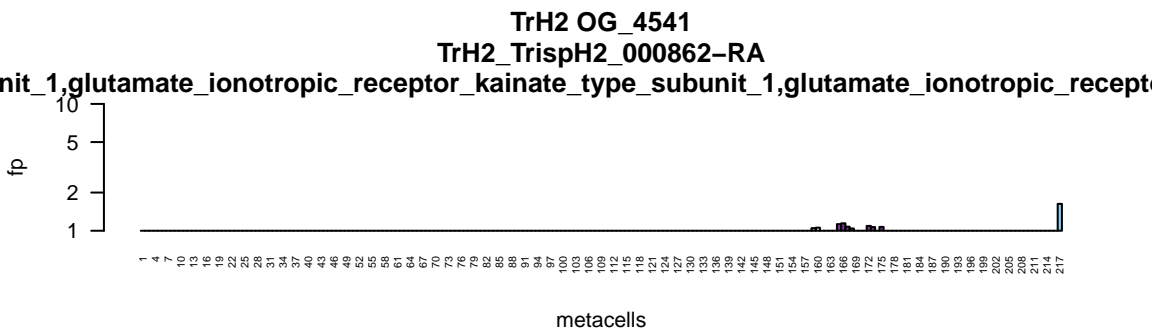
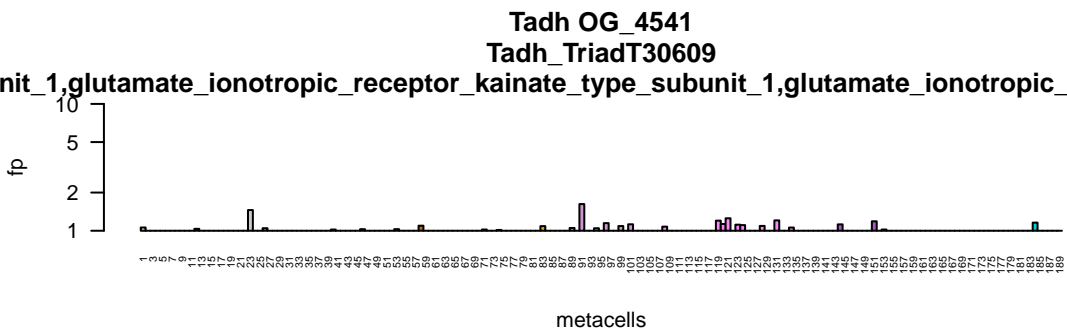
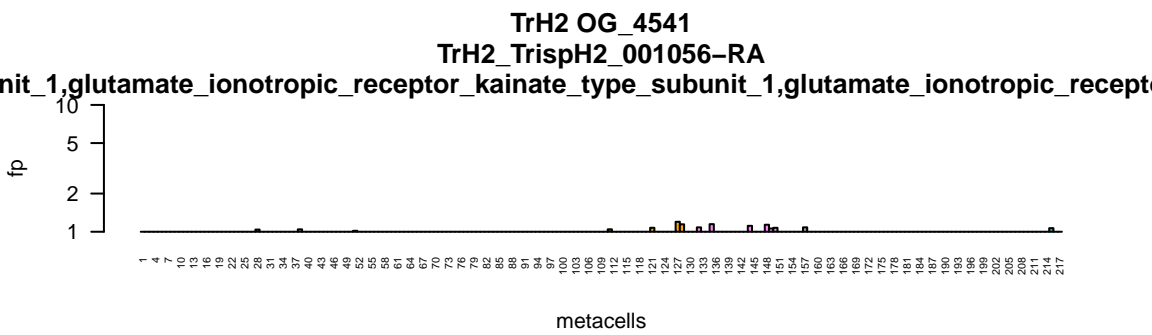
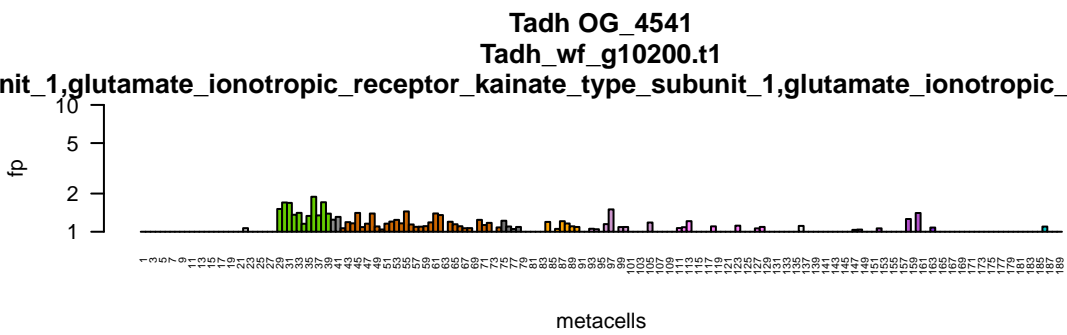
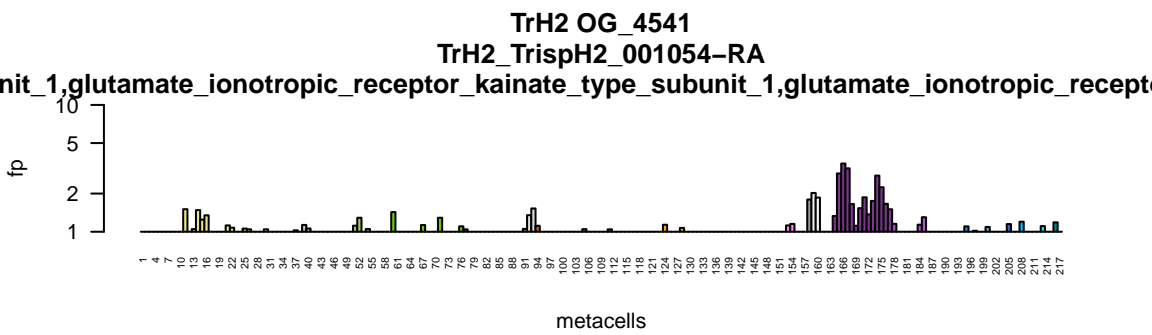
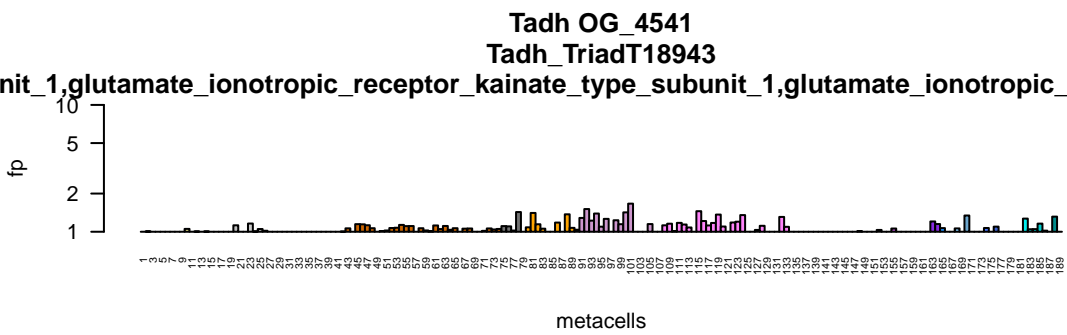
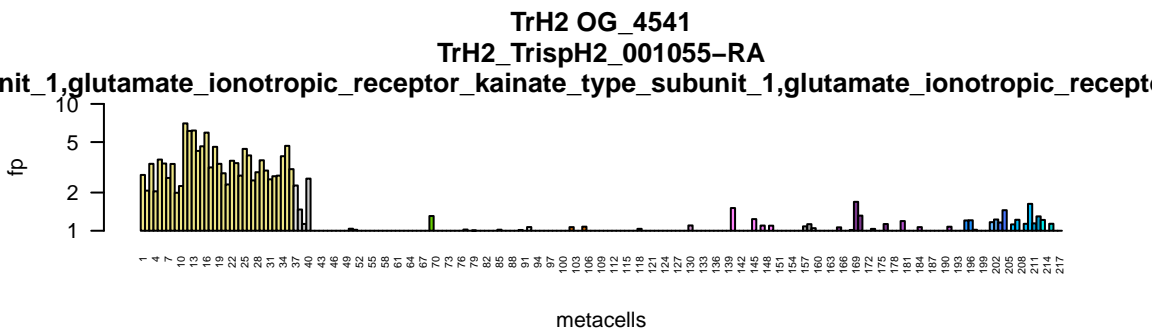
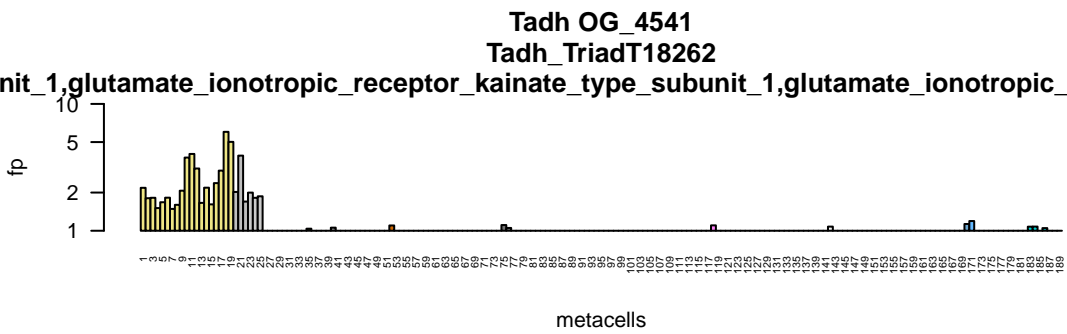
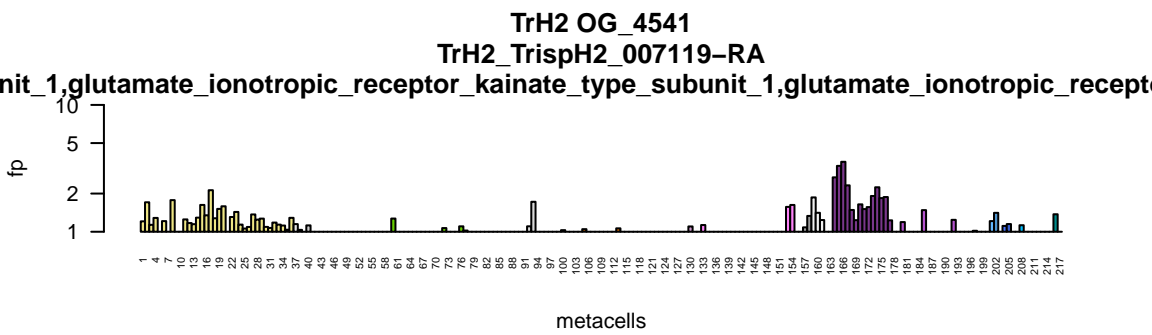
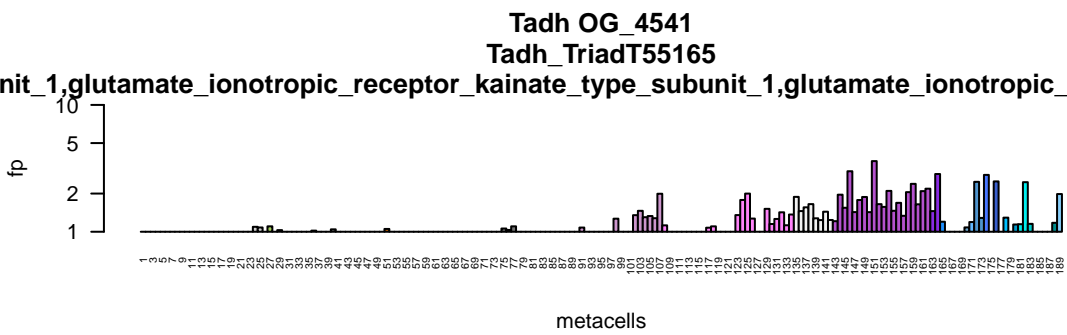
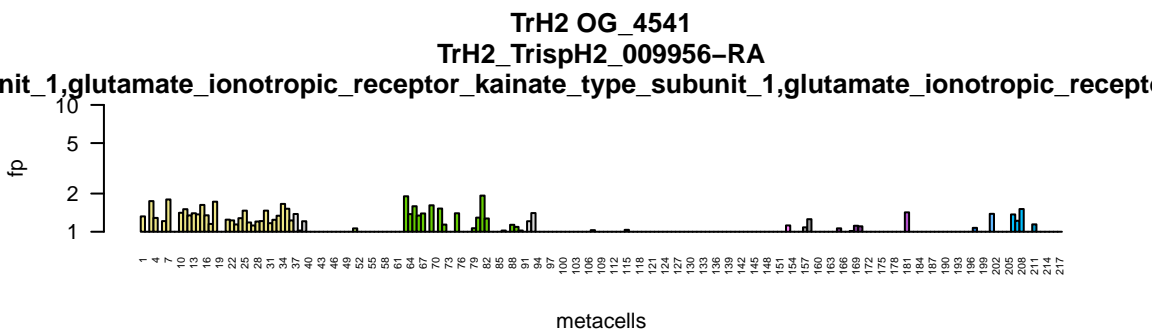
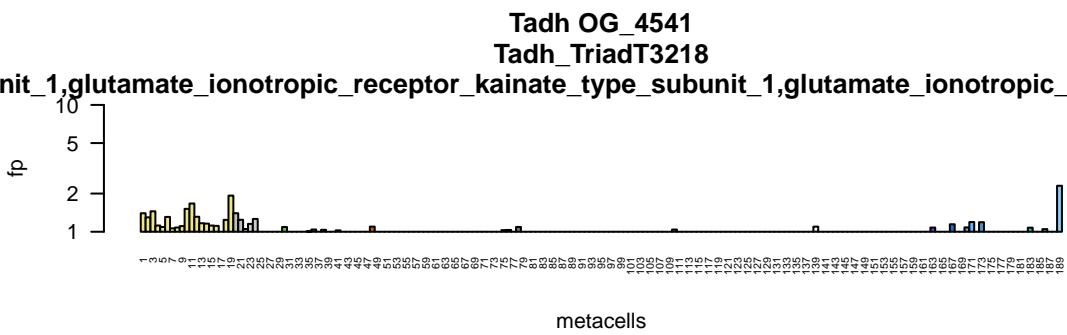
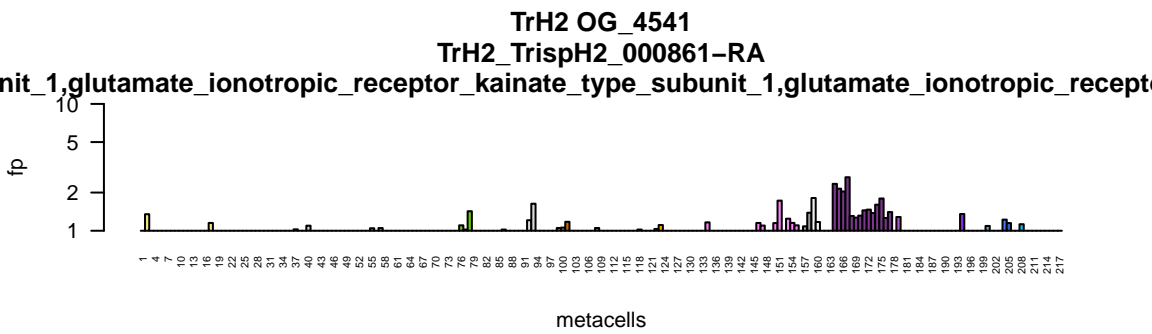
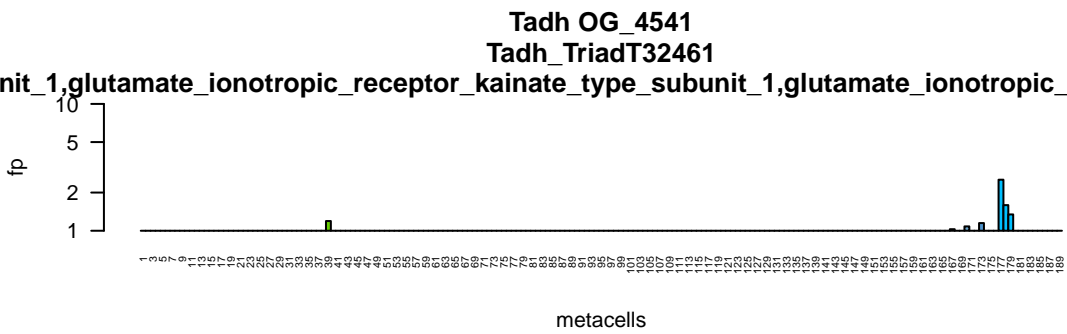
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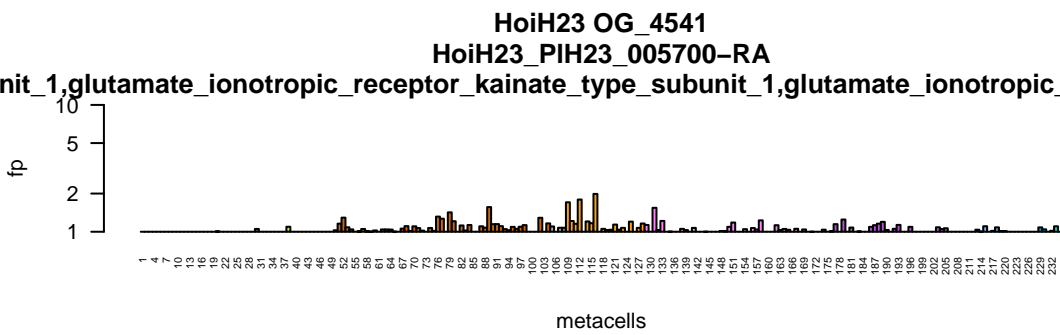
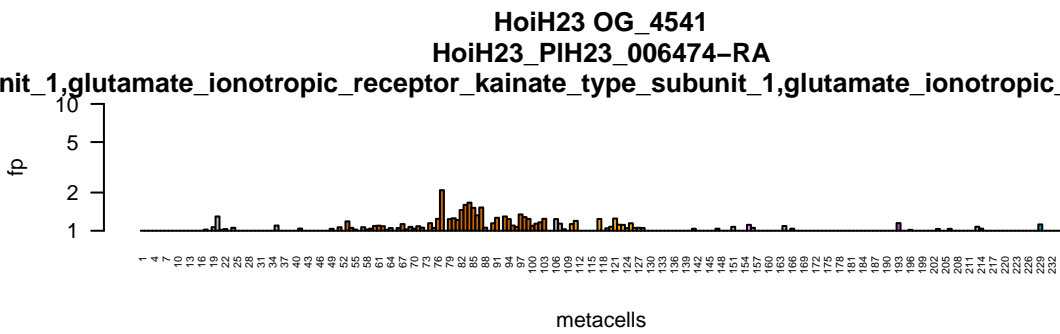
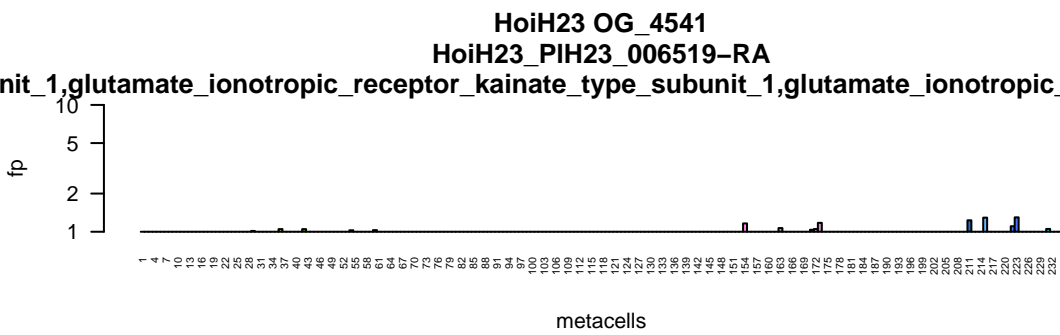
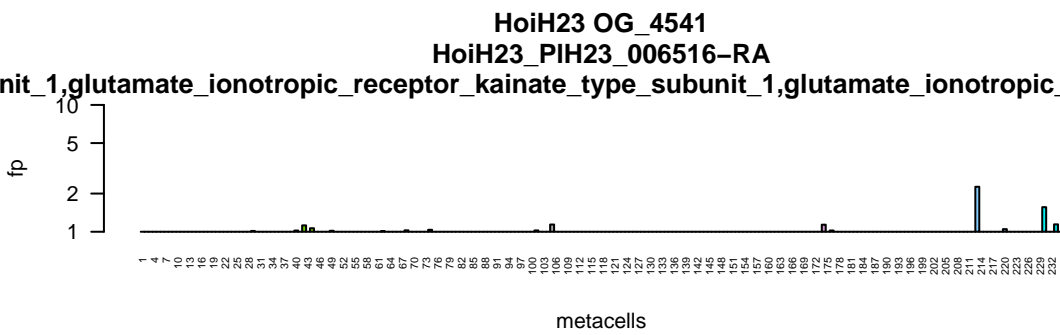
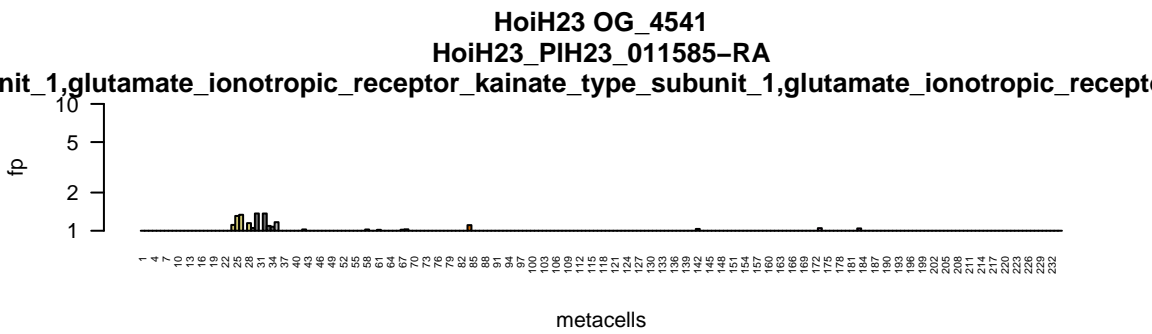
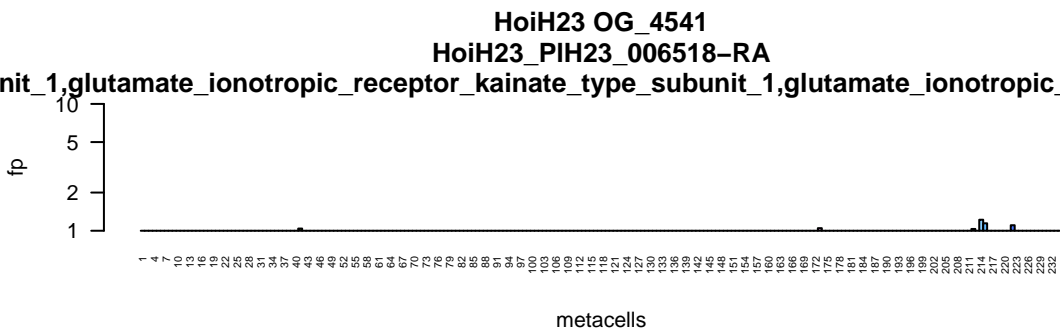
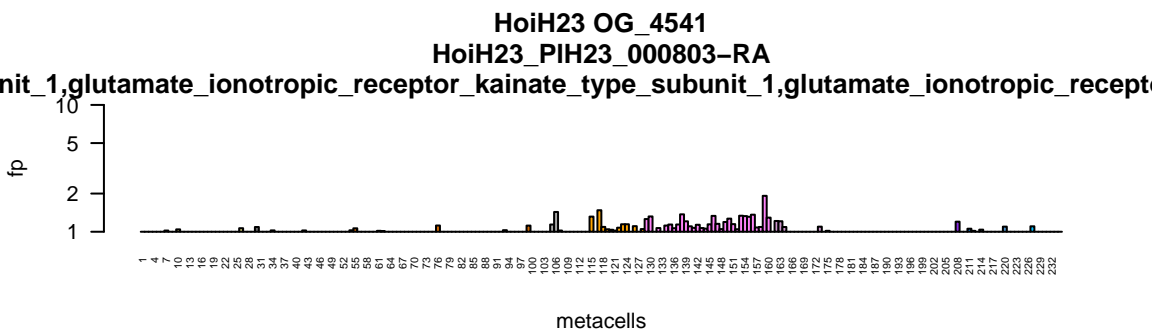
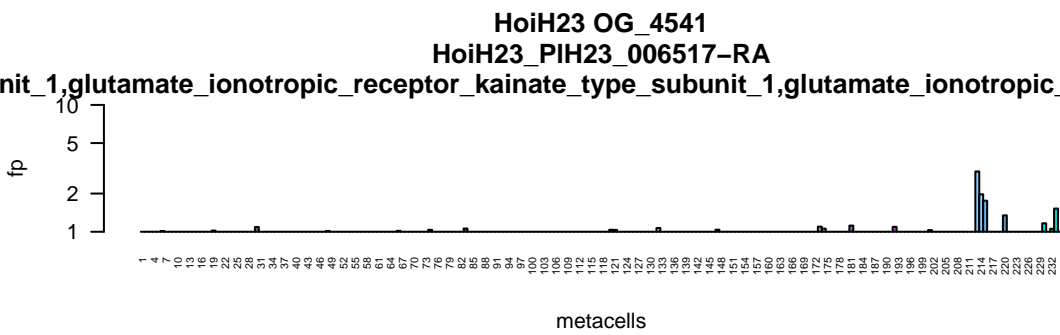
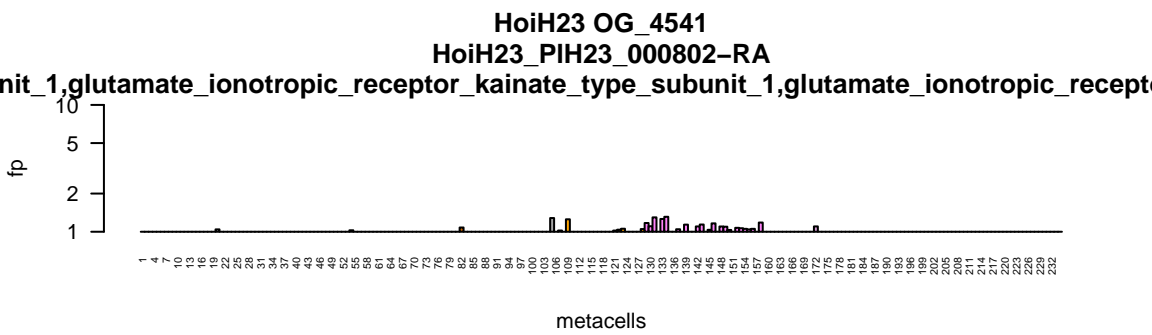
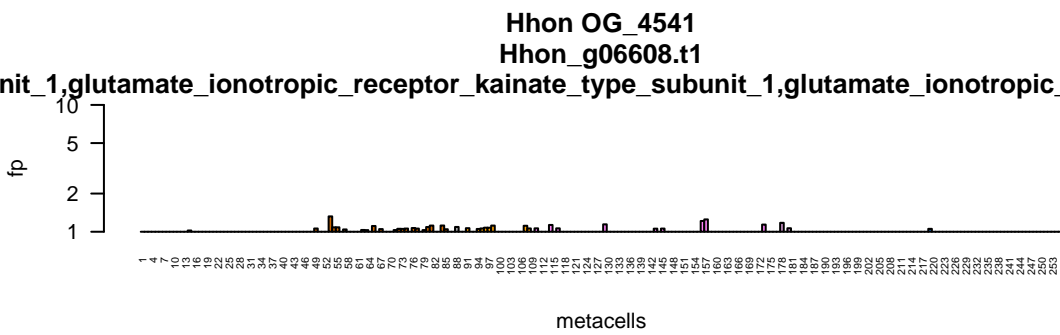
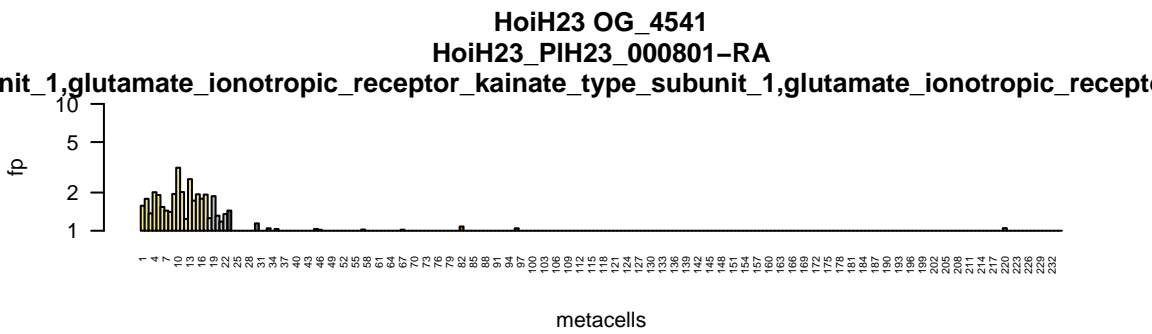
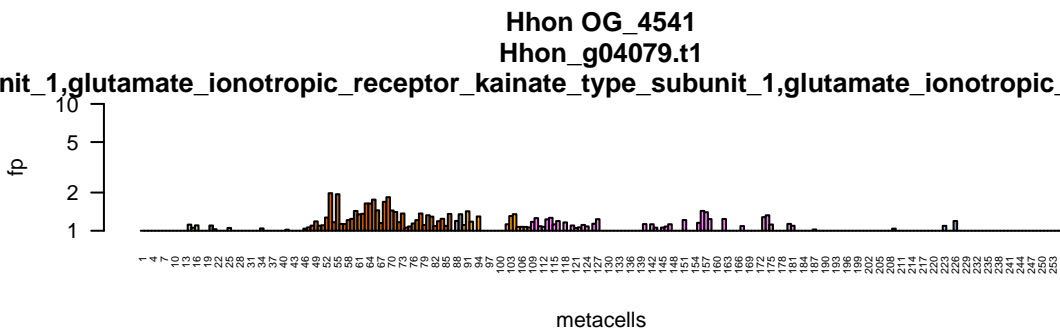


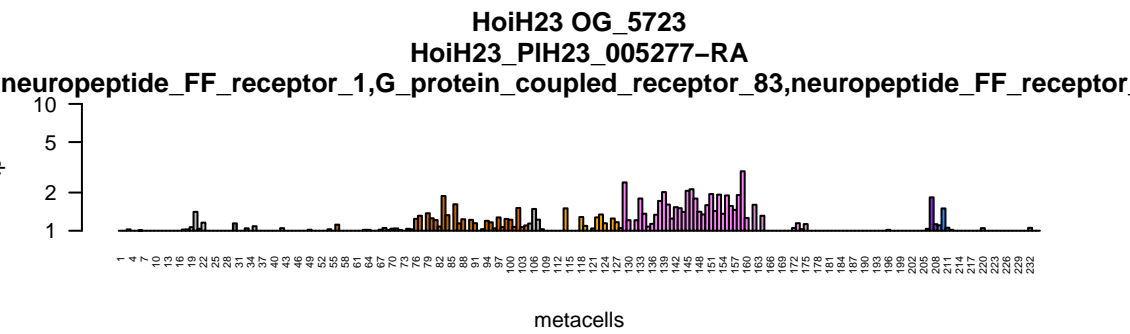
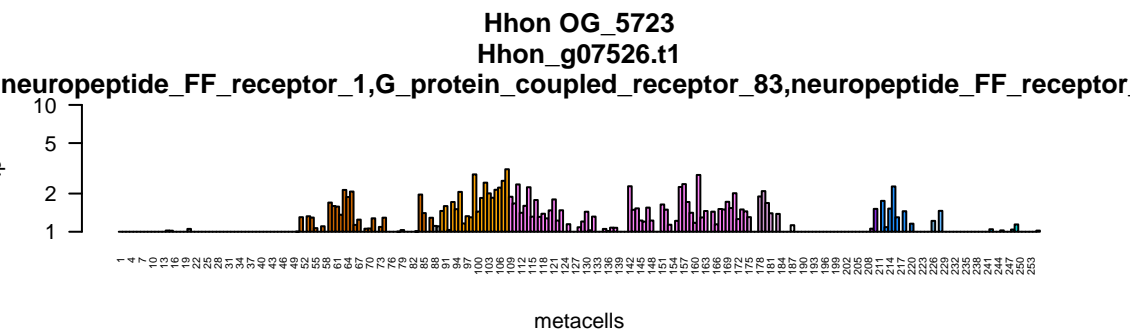
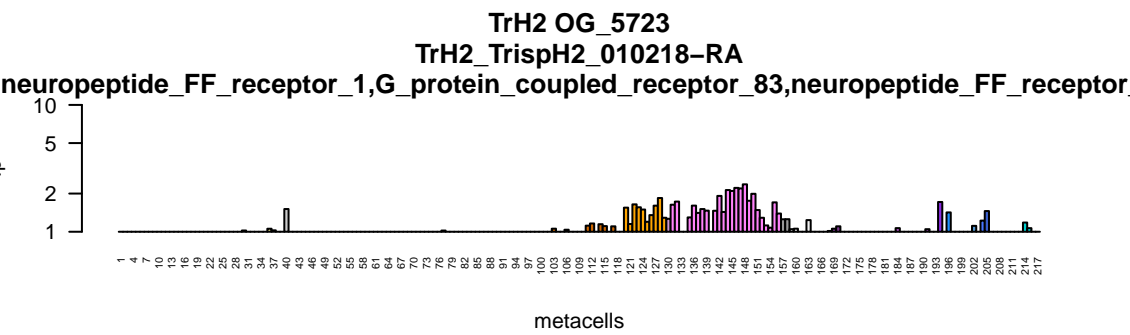
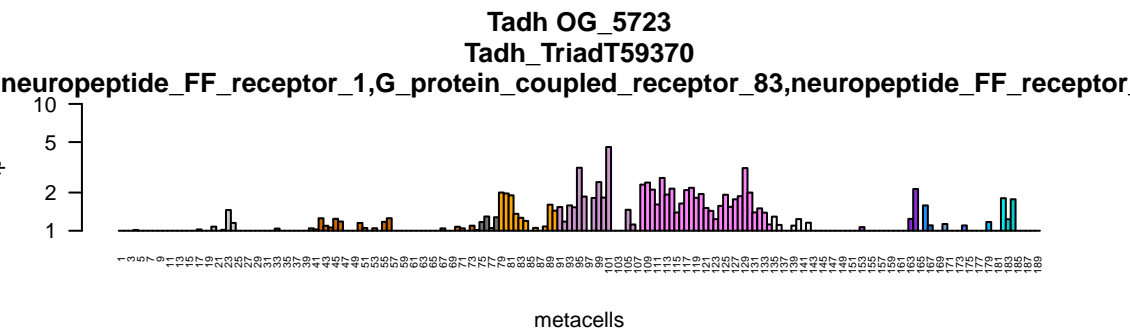
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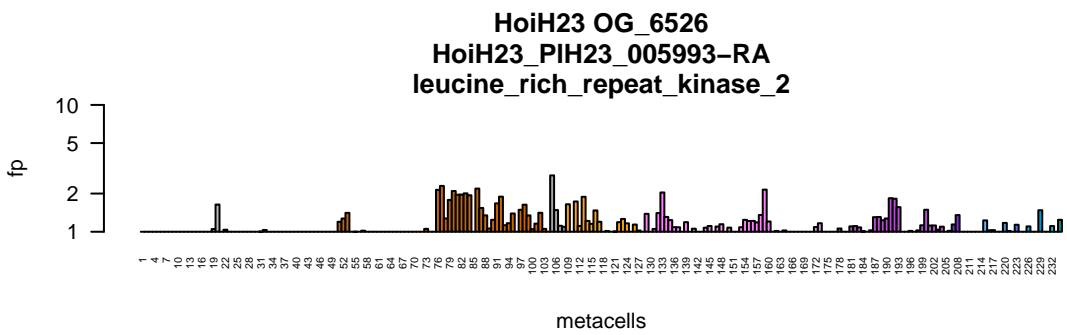
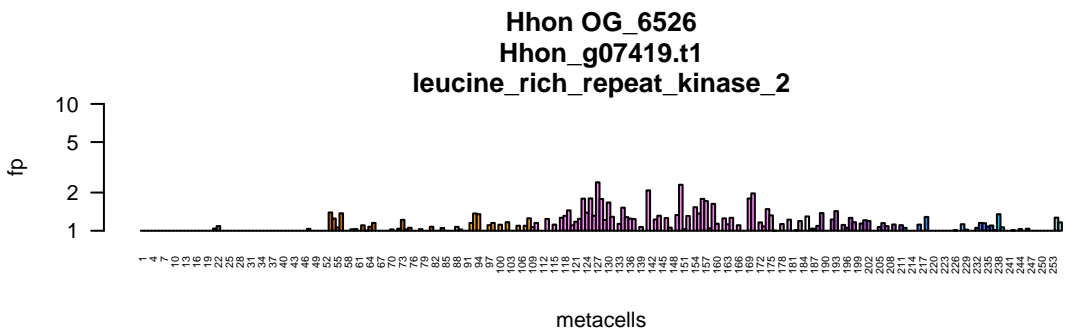
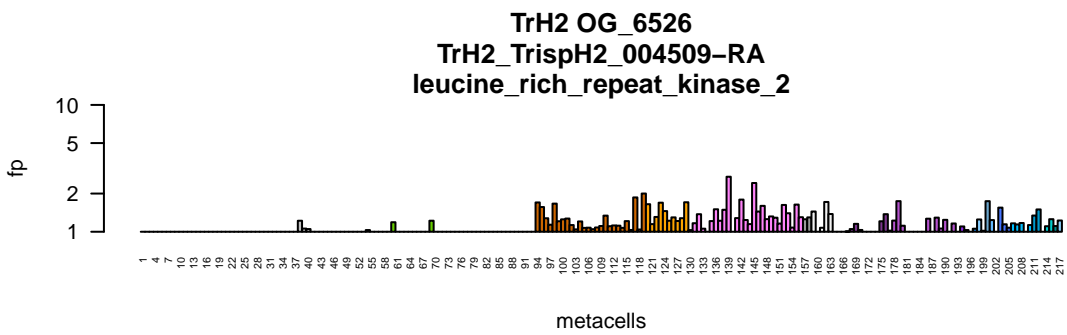
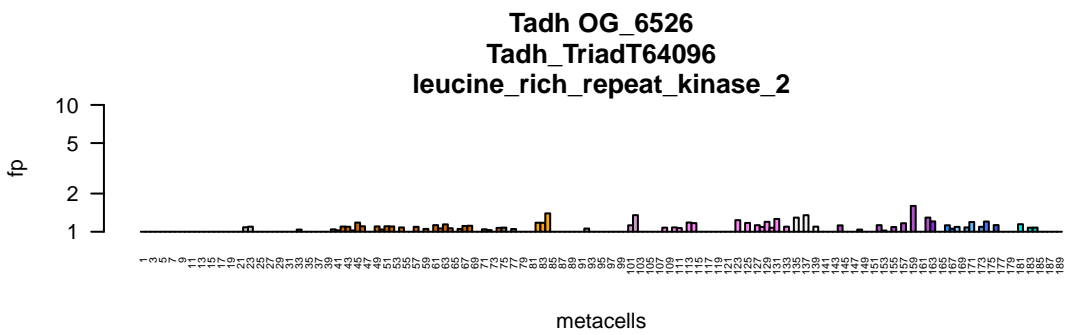
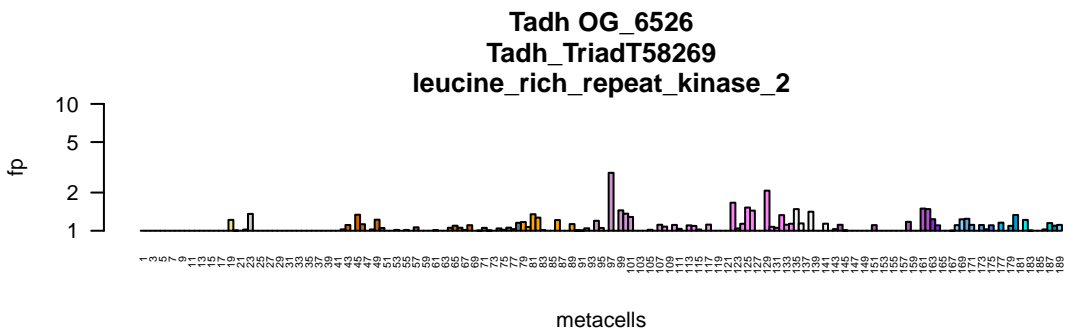


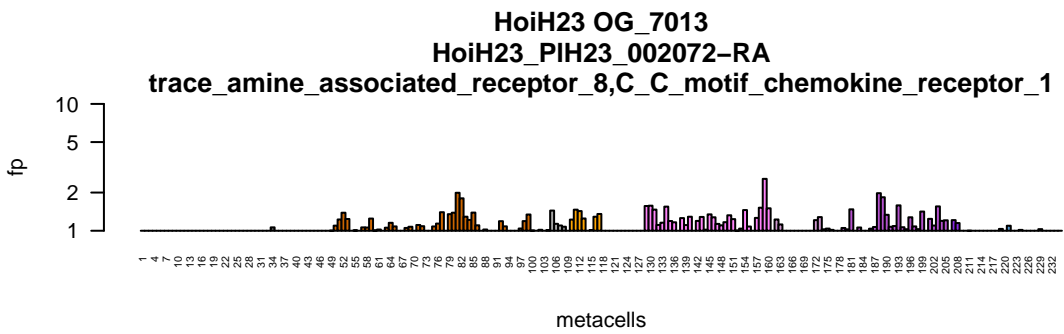
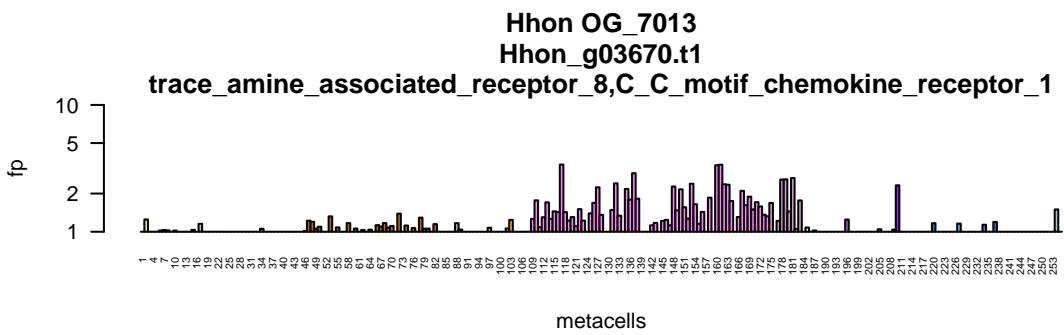
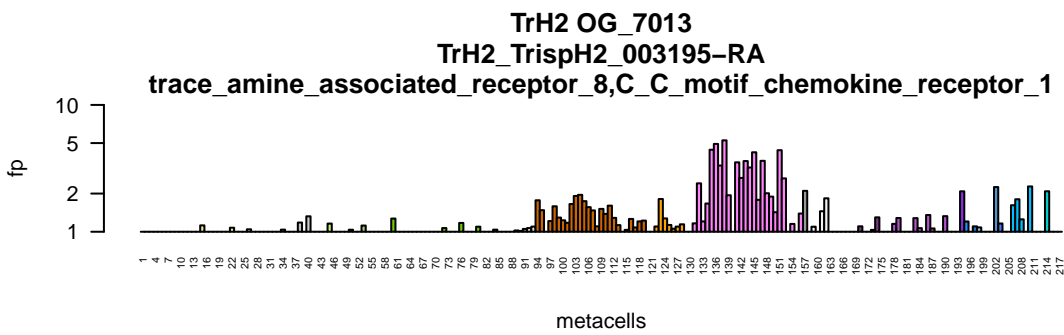
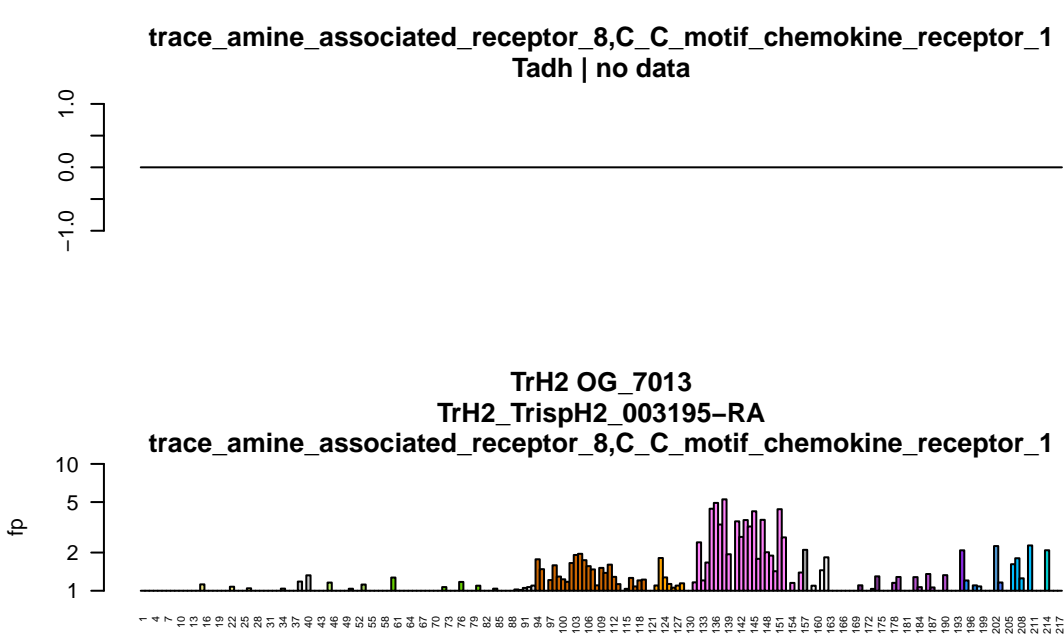




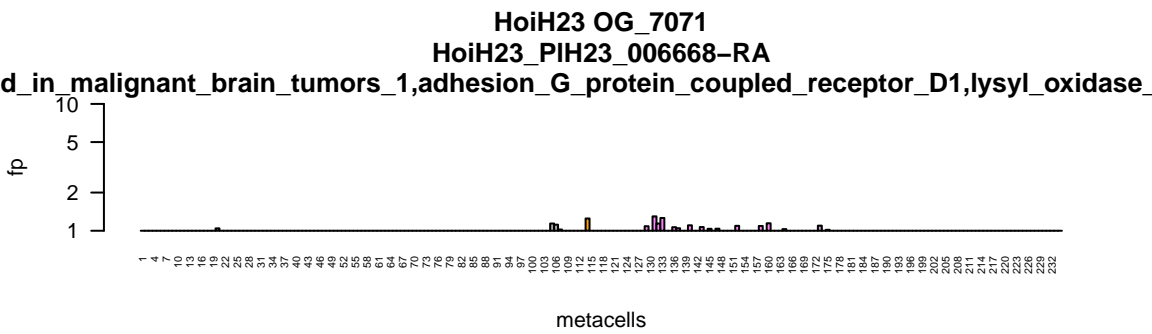
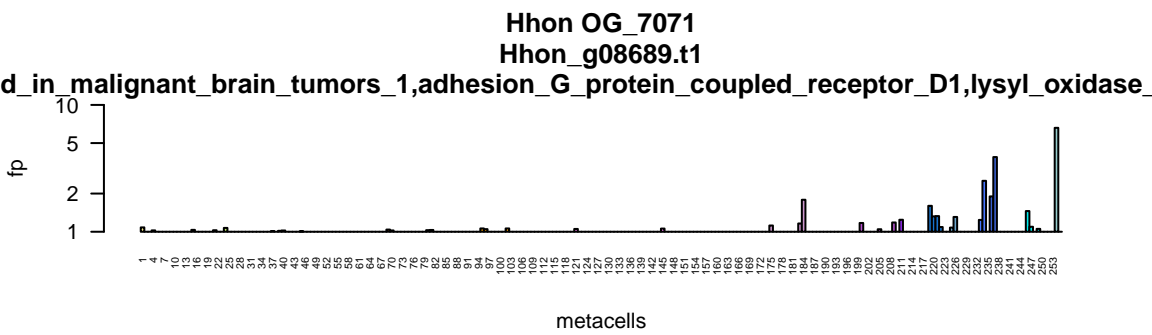
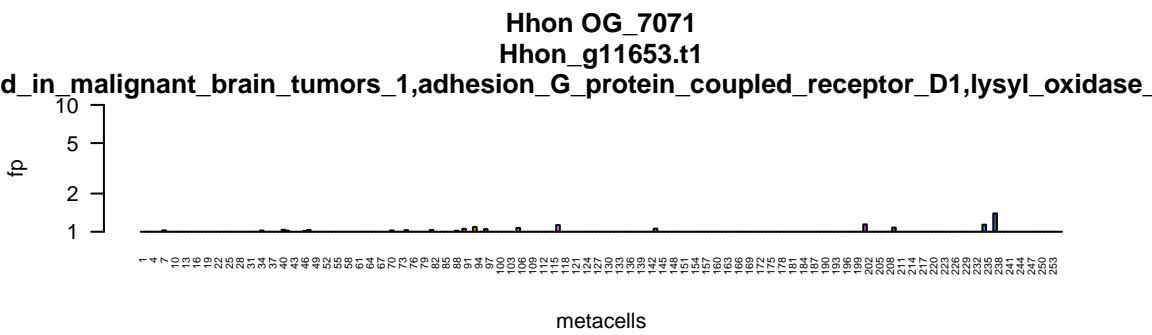
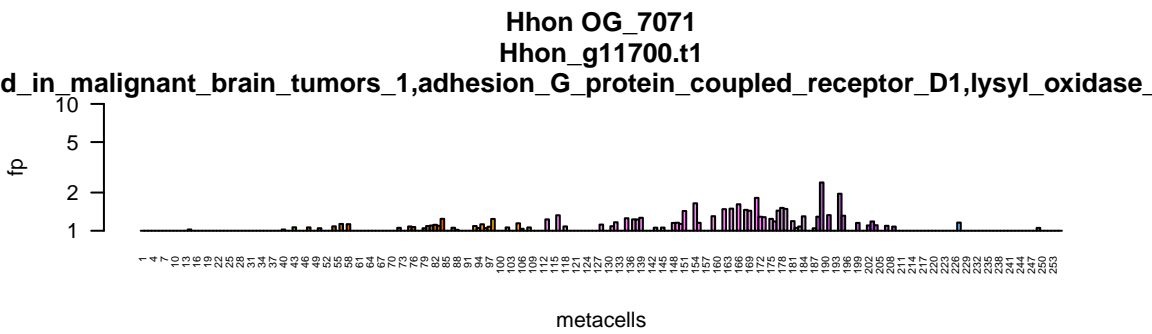
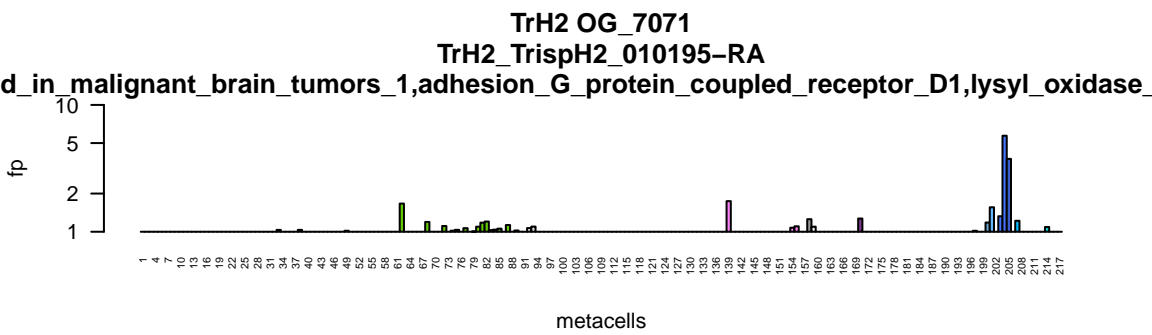
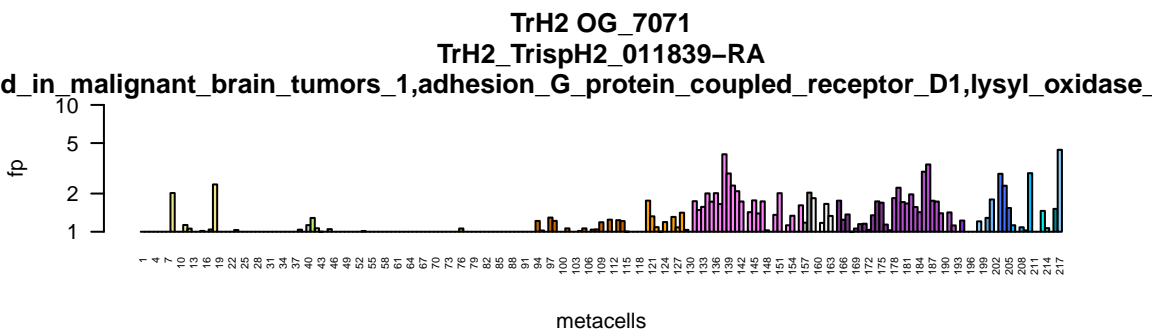
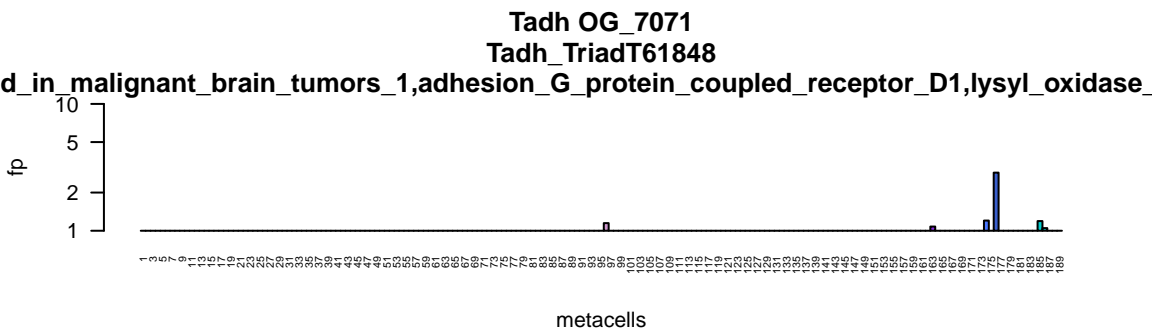


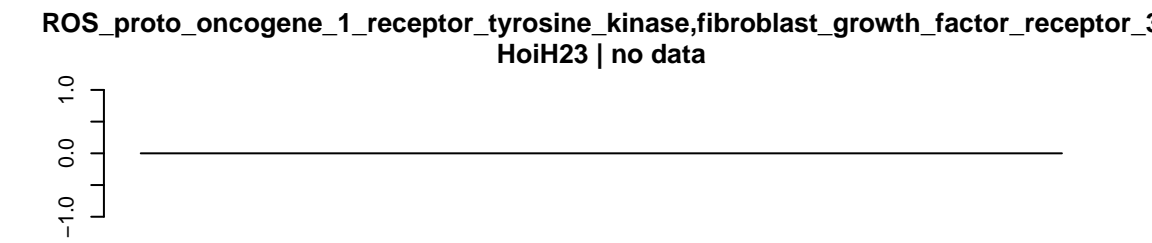
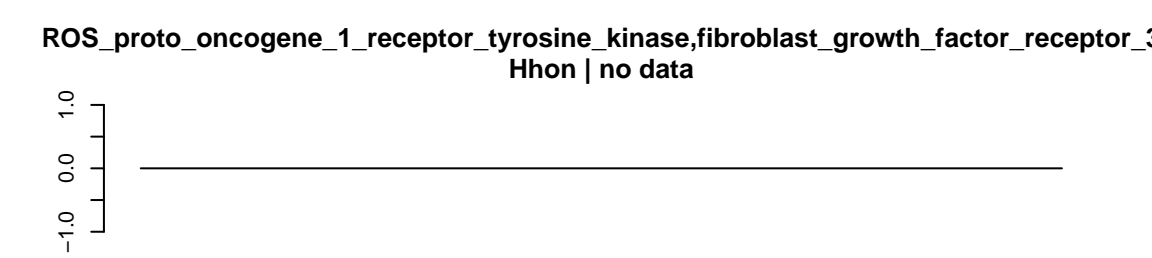
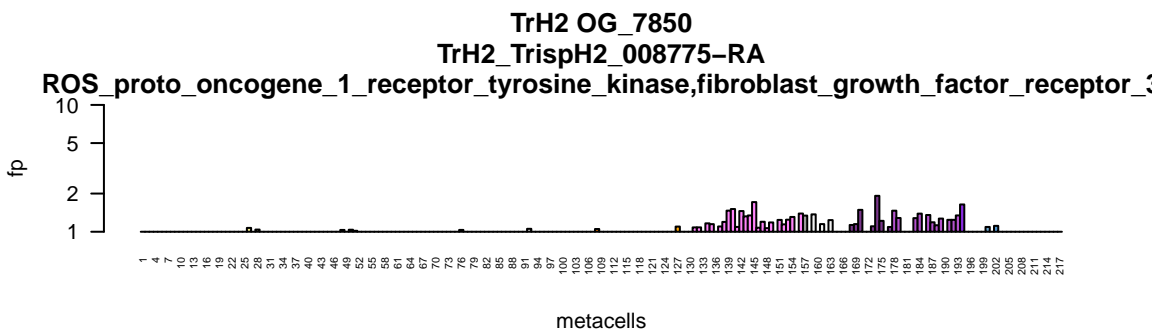
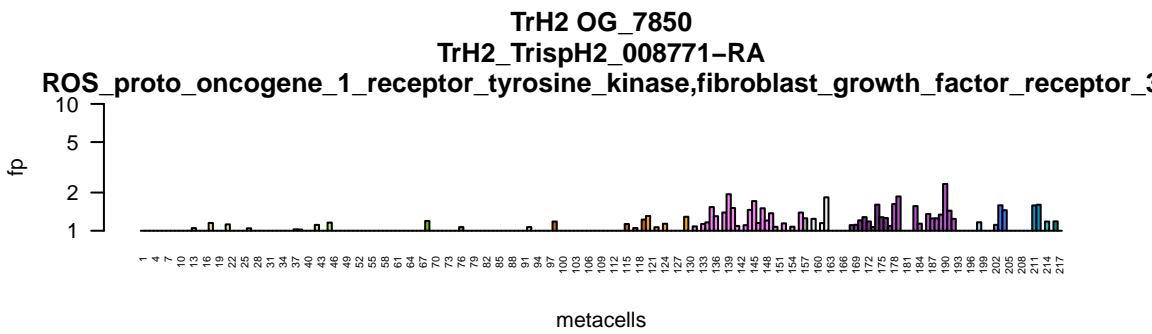
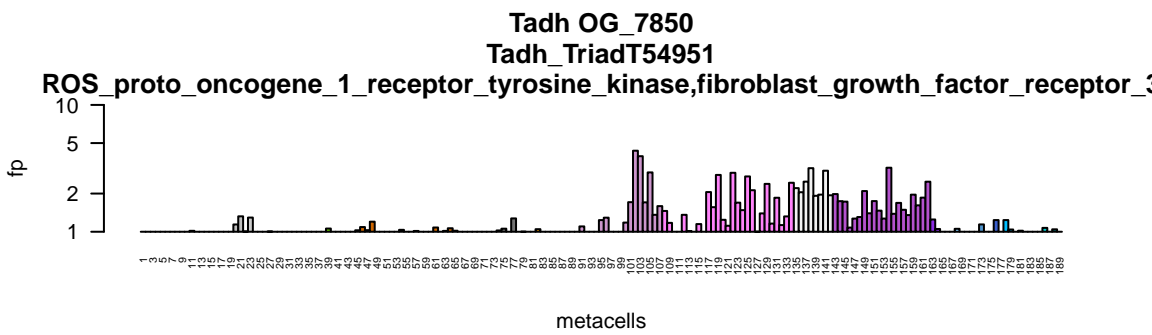
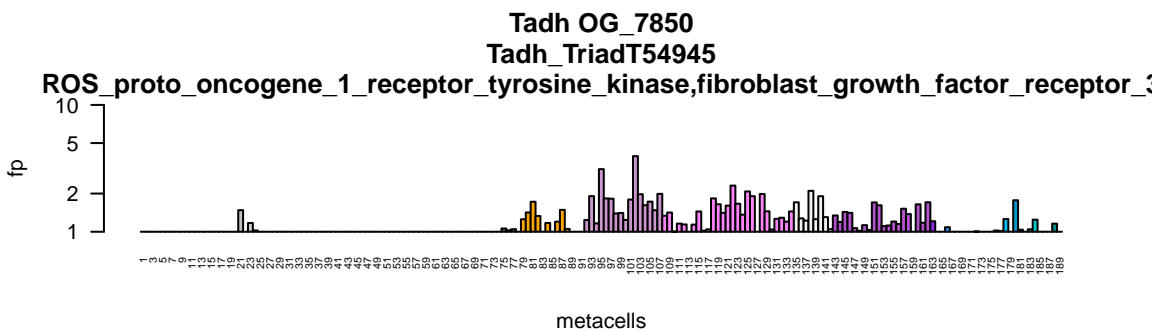
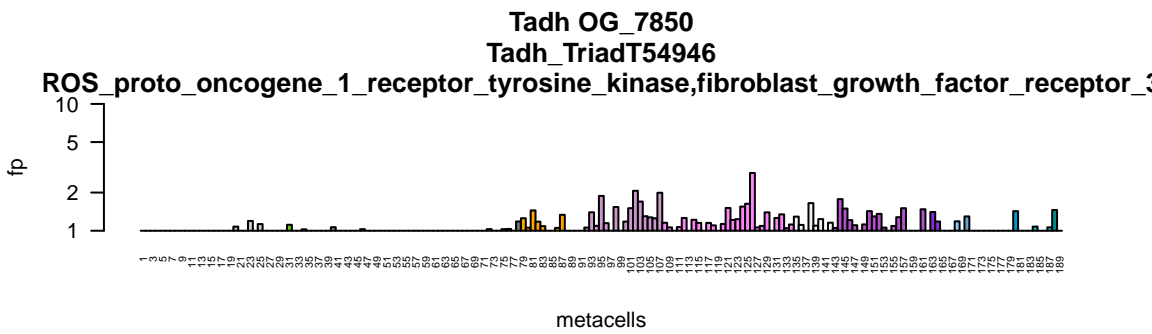


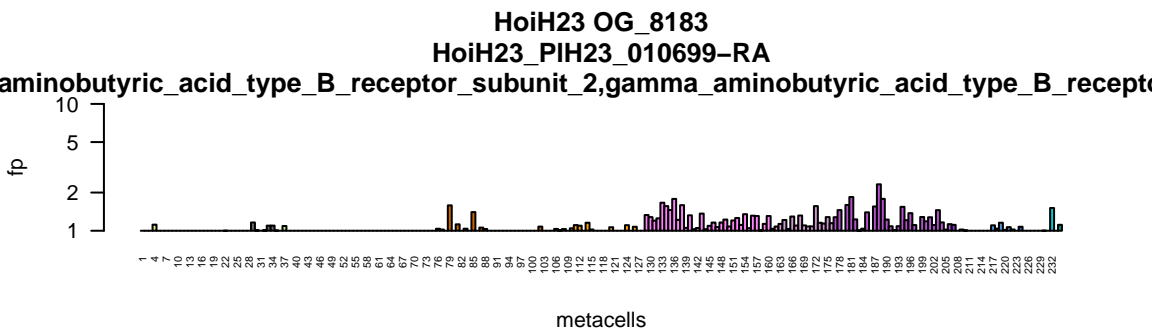
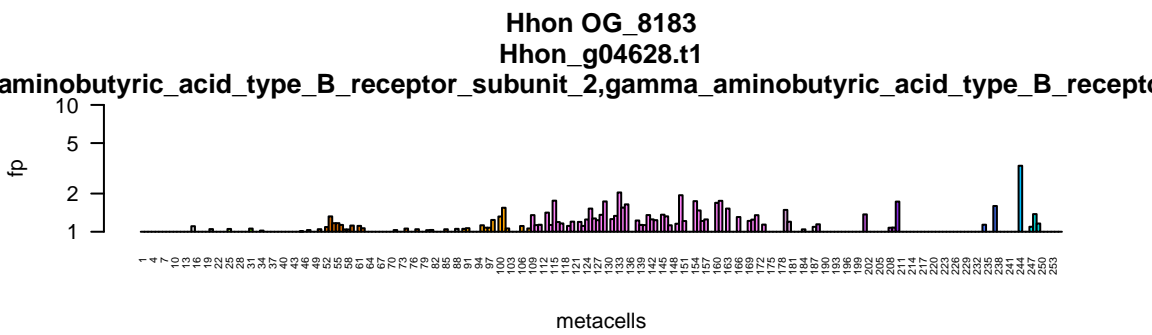
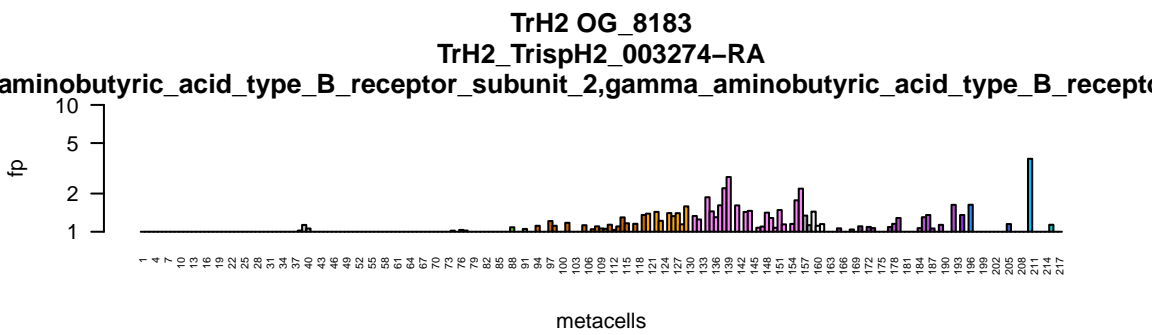
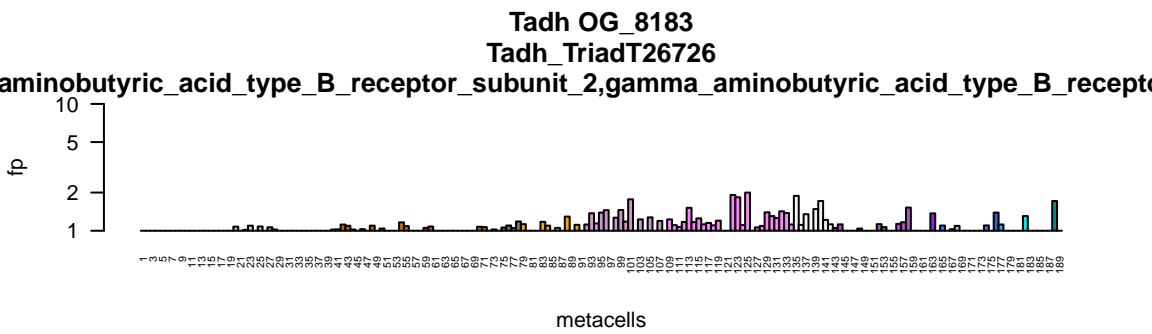


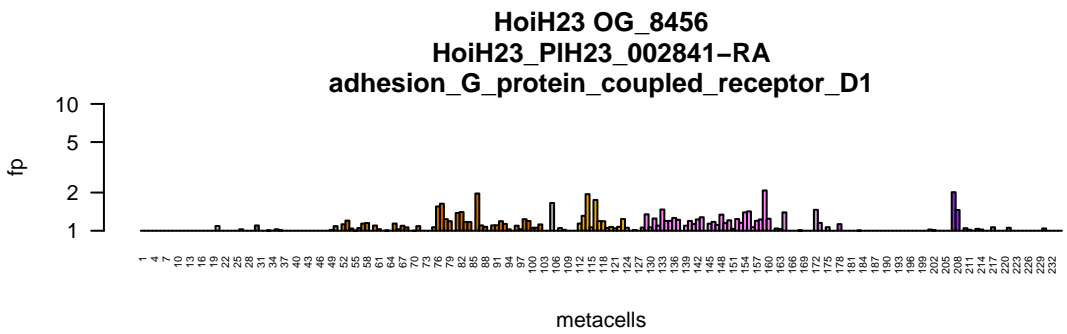
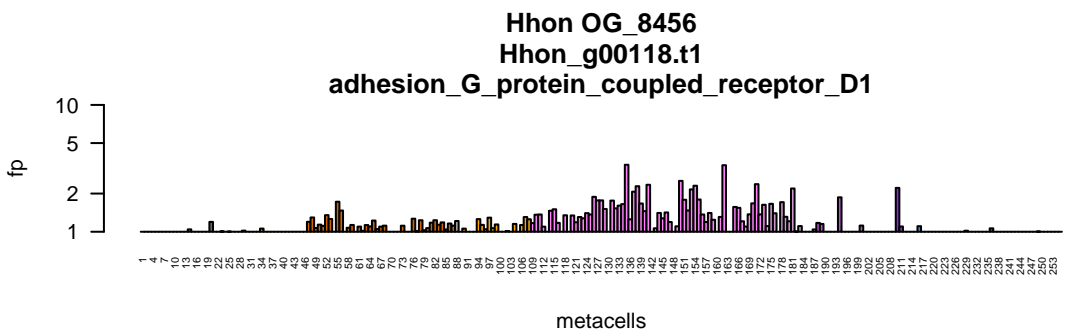
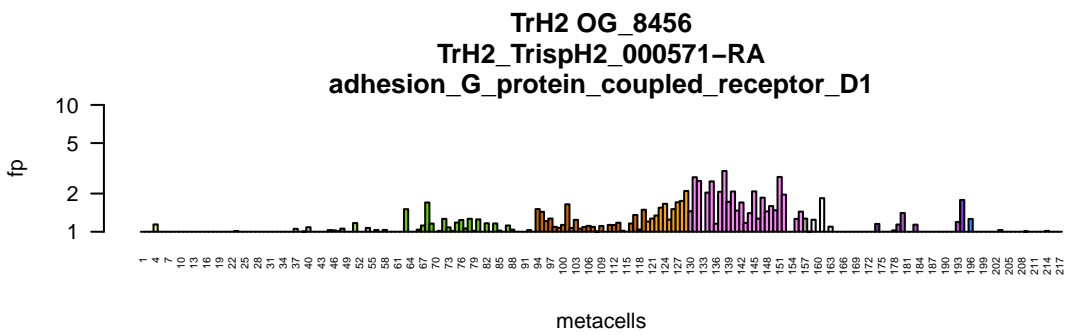
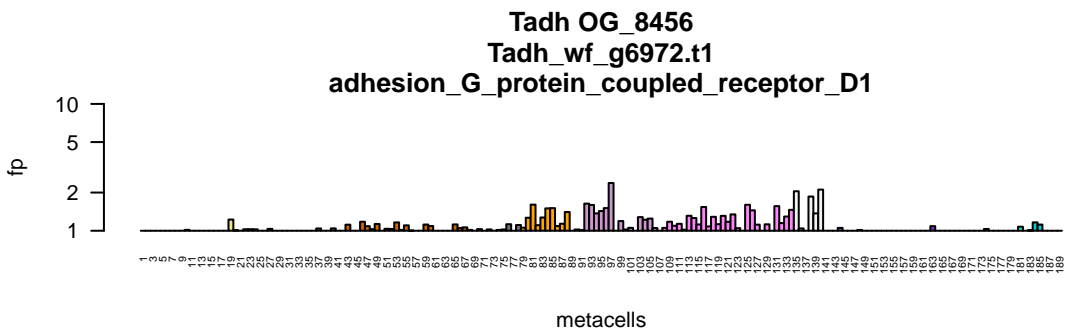


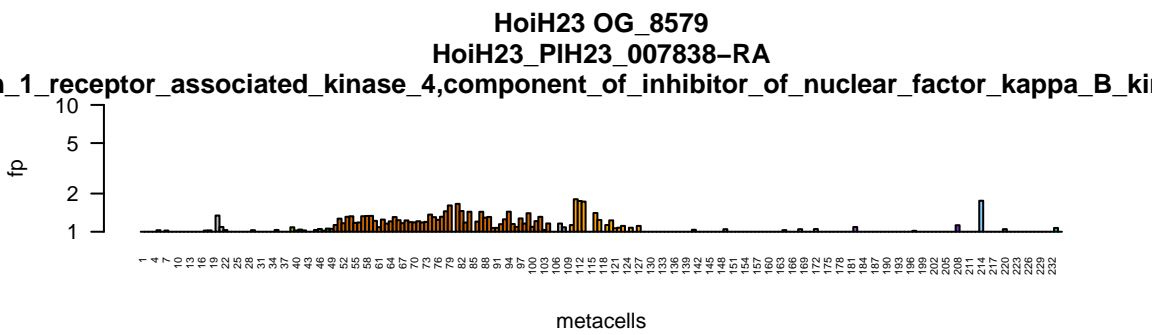
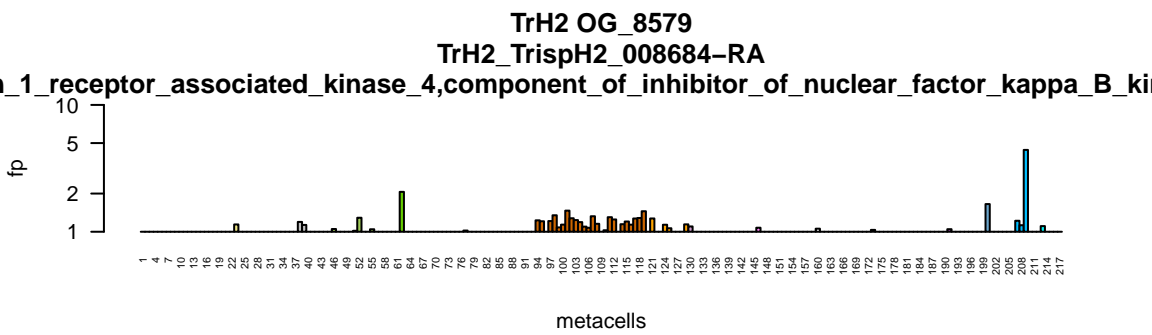
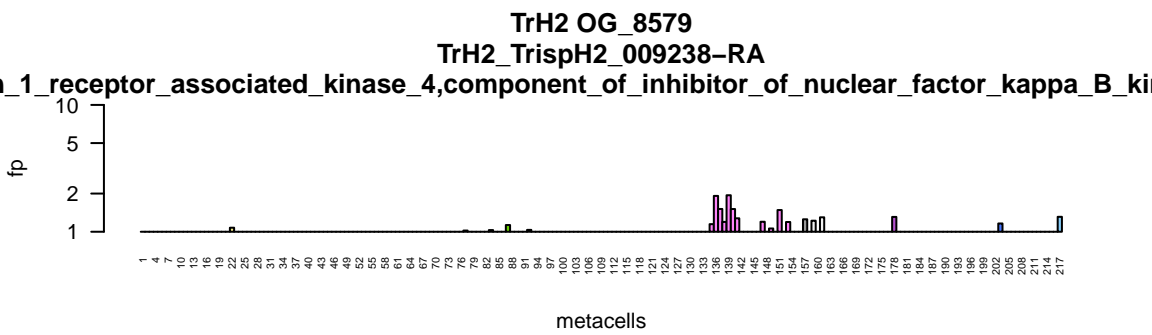
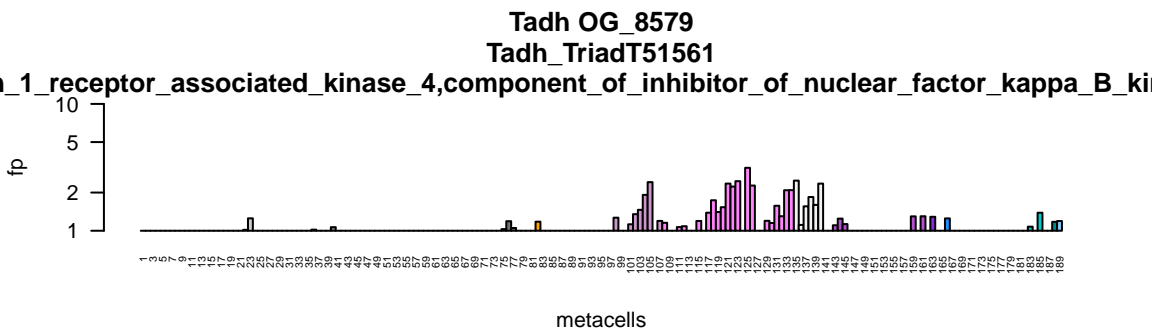


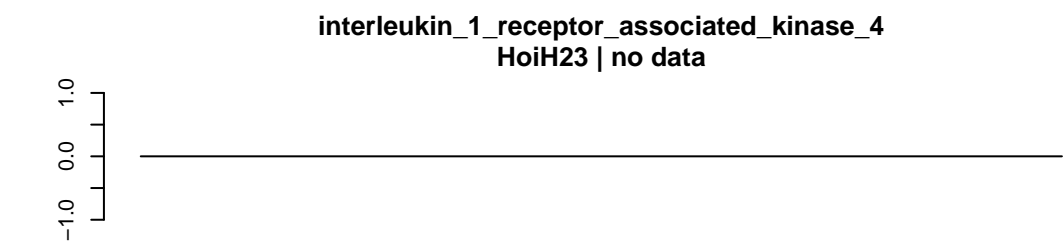
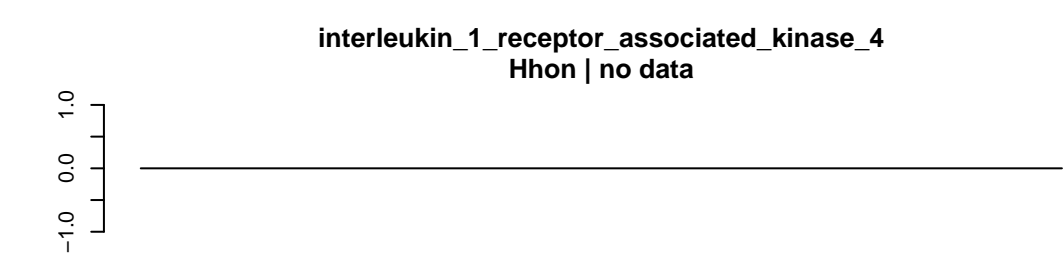
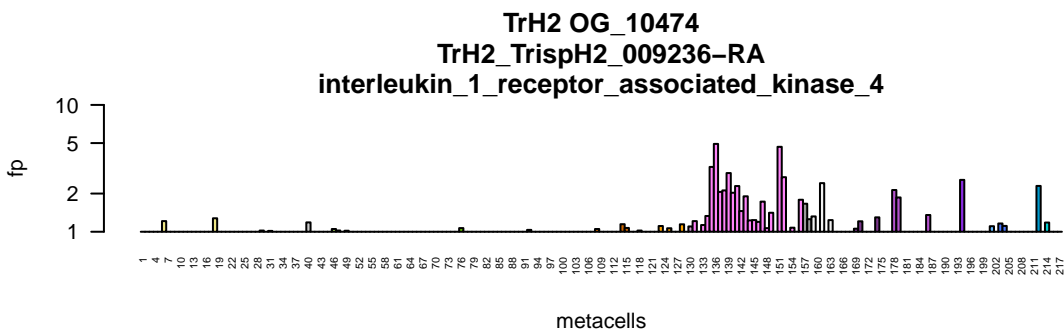
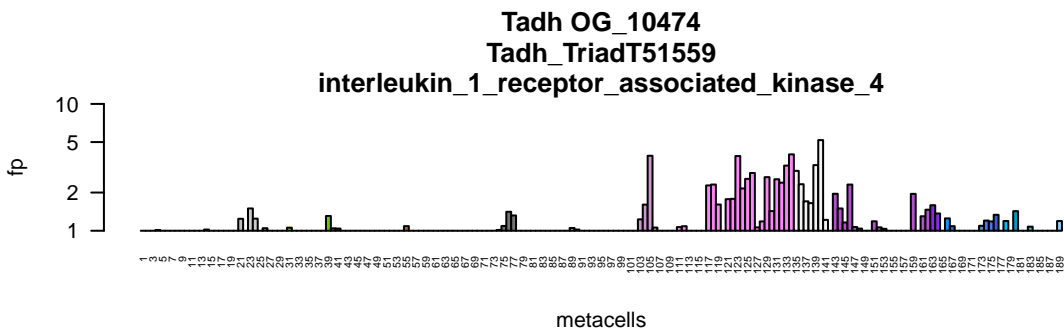












TrH2 OG\_2962  
TrH2\_TripH2\_008615-RA  
CASP2\_and\_RIPK1\_domain\_containing\_adaptor\_with\_death\_domain,Fas\_associated\_via

fp

metacells

TrH2 OG\_2962  
TrH2\_TrispH2\_011883-RA  
CASP2\_and\_RIPK1\_domain\_containing\_adaptor\_with\_death\_domain,Fas\_associated\_via

fp

metacells

TrH2 OG\_2962  
TrH2\_TripH2\_011702-RA  
CASP2\_and\_RIPK1\_domain\_containing\_adaptor\_with\_death\_domain,Fas\_associated\_via

fp

metacells

TrH2 OG\_2962  
TrH2\_TripH2\_011686-RA  
CASP2\_and\_RIPK1\_domain\_containing\_adaptor\_with\_death\_domain,Fas\_associated\_via

fp

metacells

TrH2 OG\_2962  
TrH2\_TripH2\_011703-RA  
CASP2\_and\_RIPK1\_domain\_containing\_adaptor\_with\_death\_domain,Fas\_associated\_via

metacells	fp
94	10
184	1

TrH2 OG\_2962  
TrH2\_TrispH2\_003510-RA  
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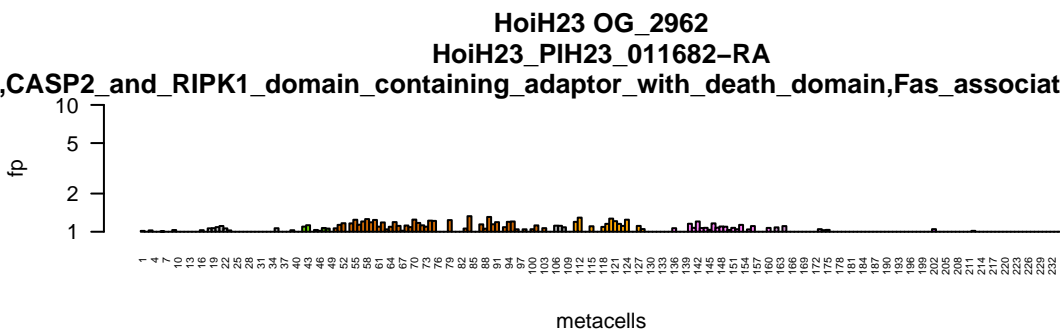
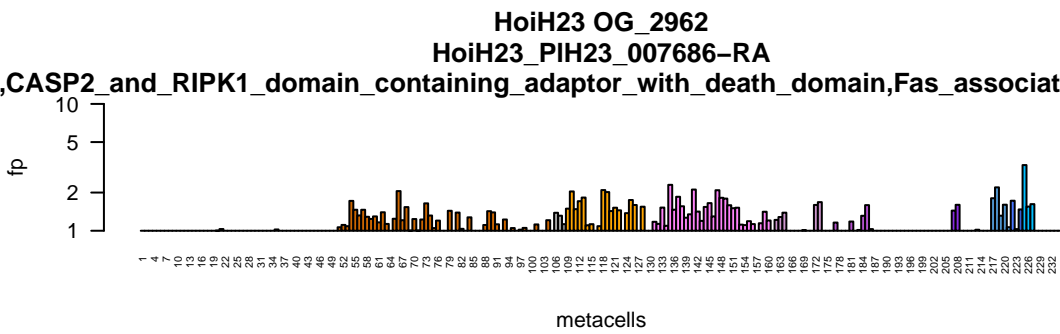
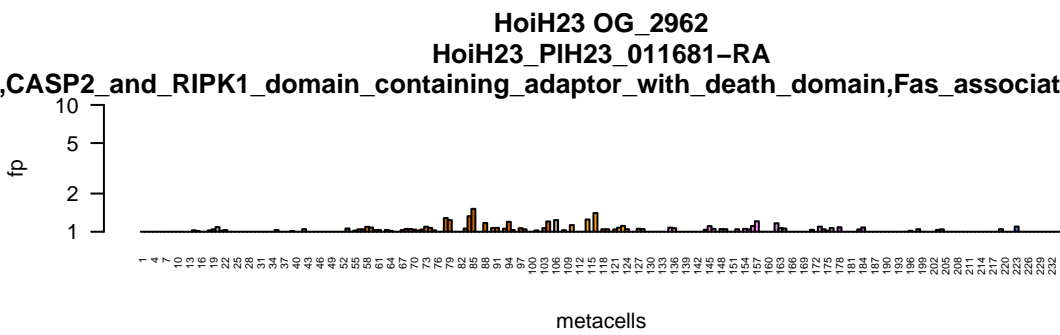
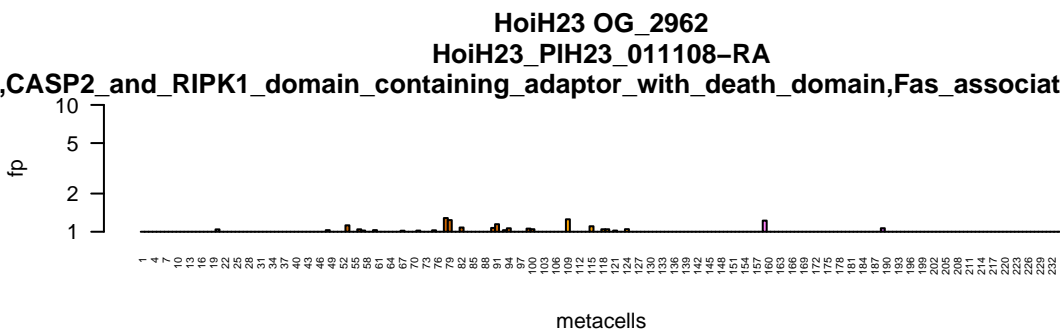
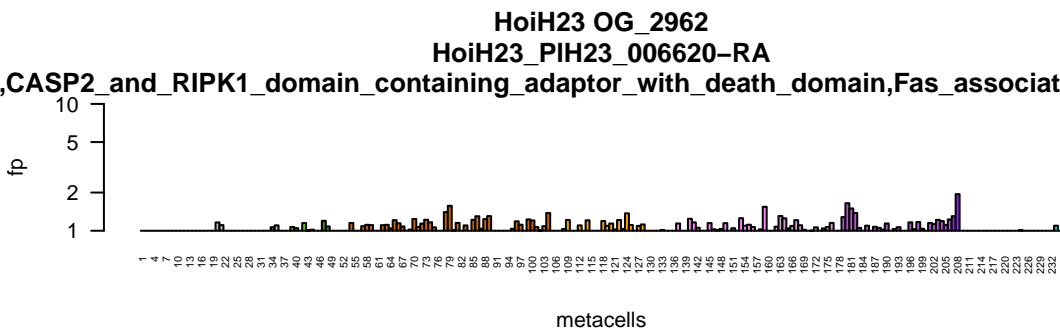
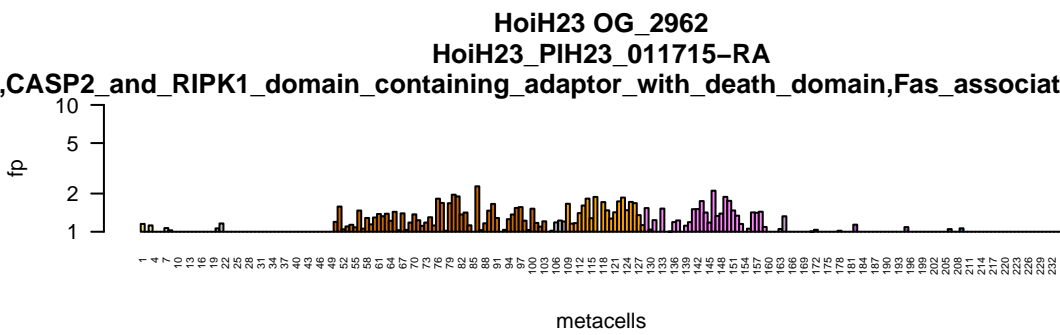
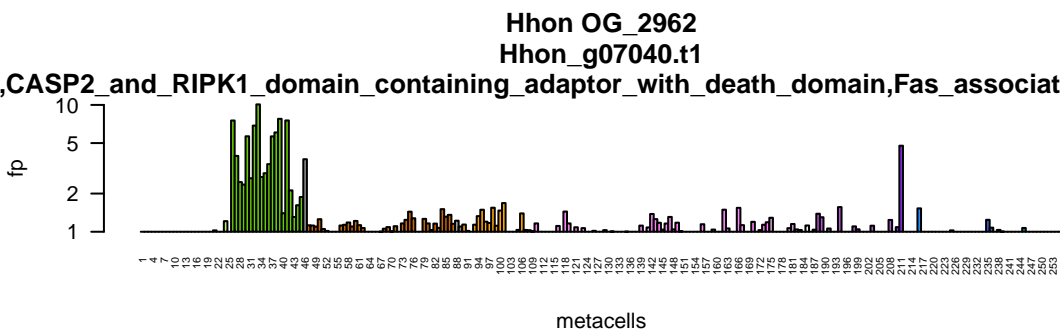
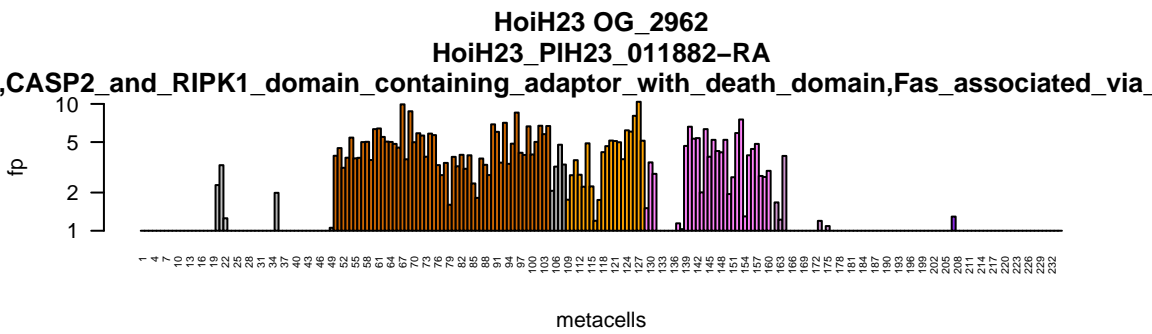
fp

metacells

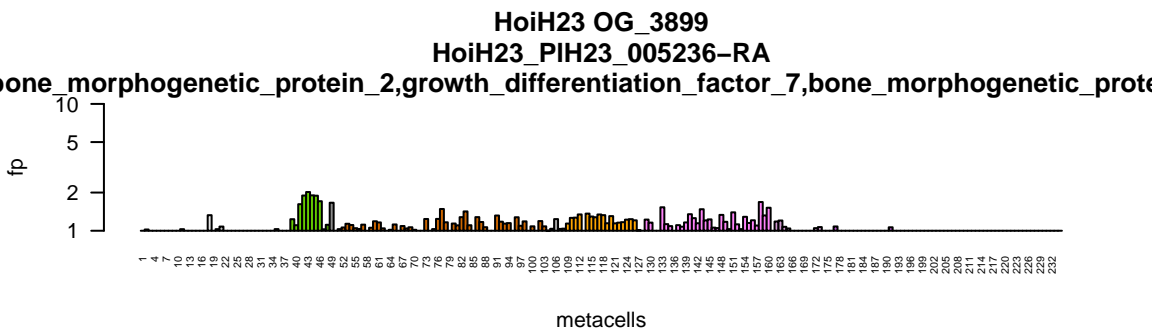
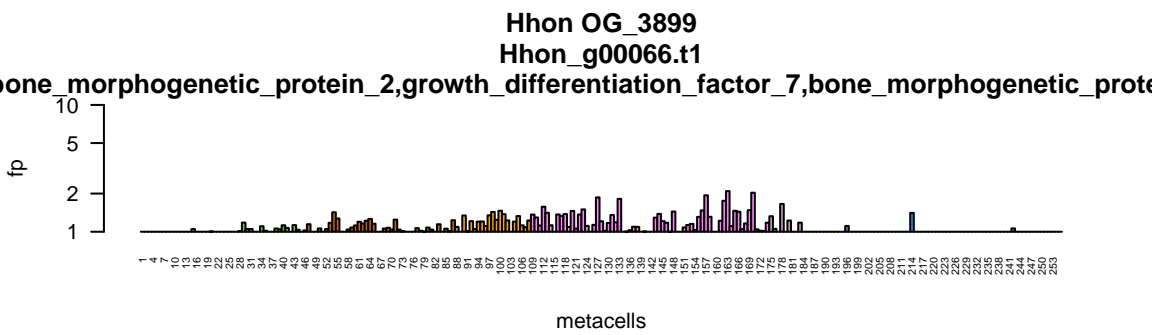
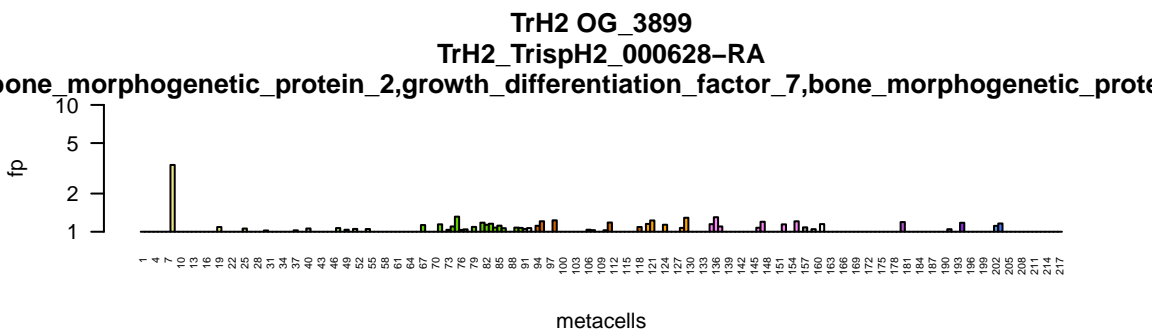
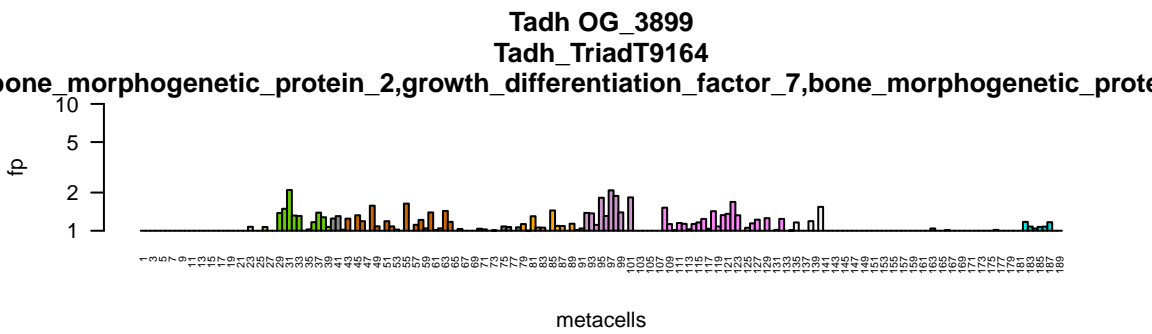
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TrH2\_TrispH2\_011821-RA  
CASP2\_and\_RIPK1\_domain\_containing\_adaptor\_with\_death\_domain,Fas\_associated\_via

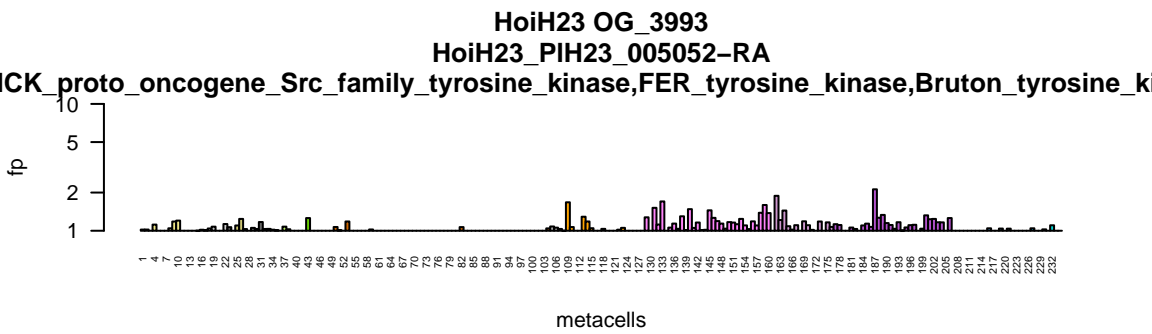
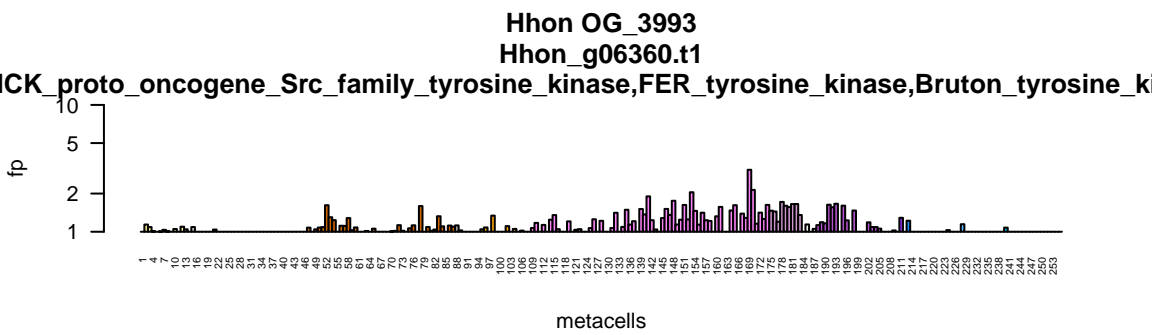
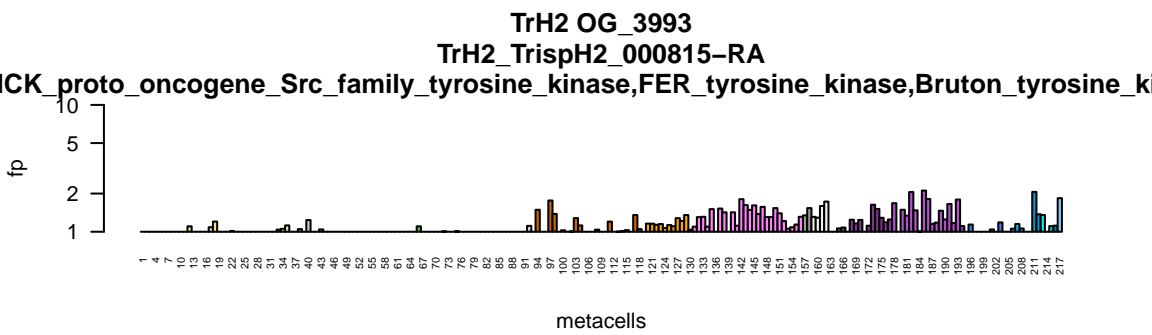
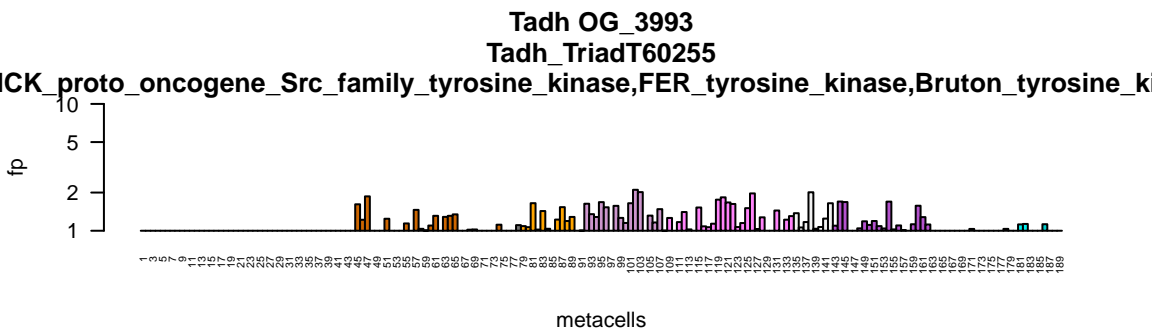
fp

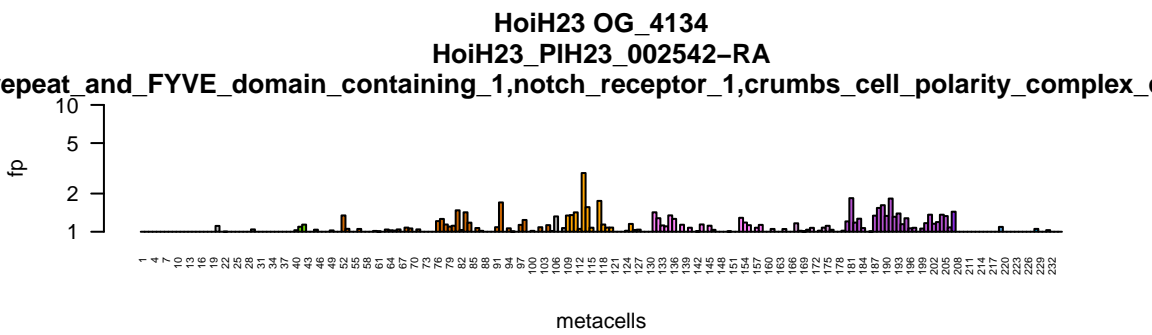
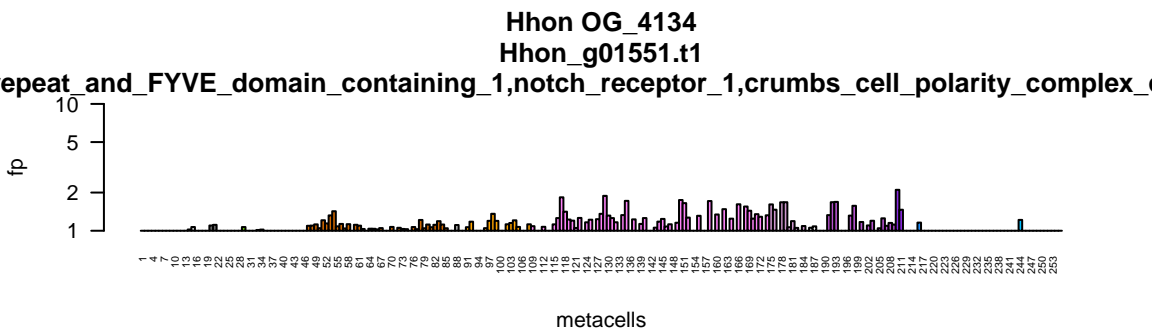
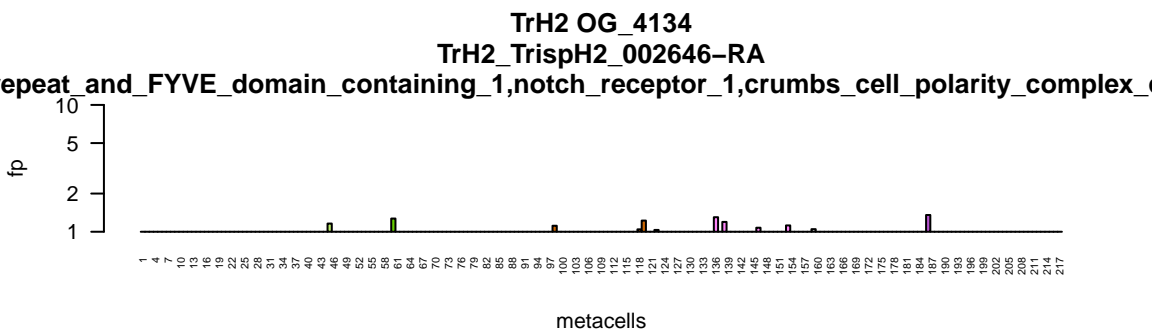
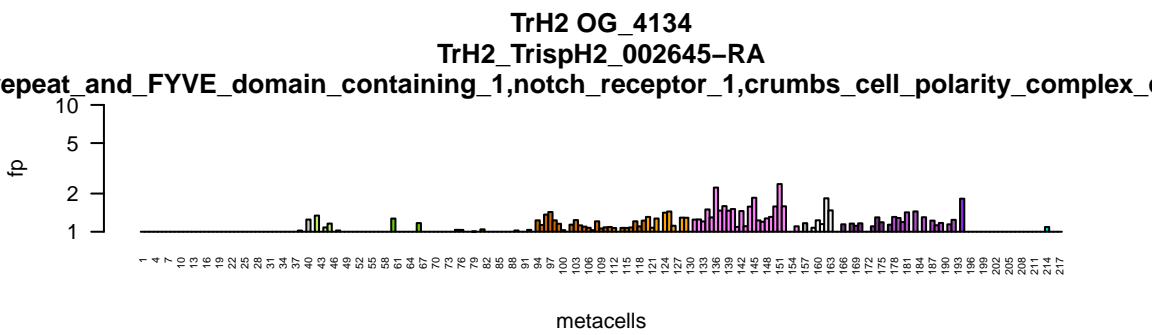
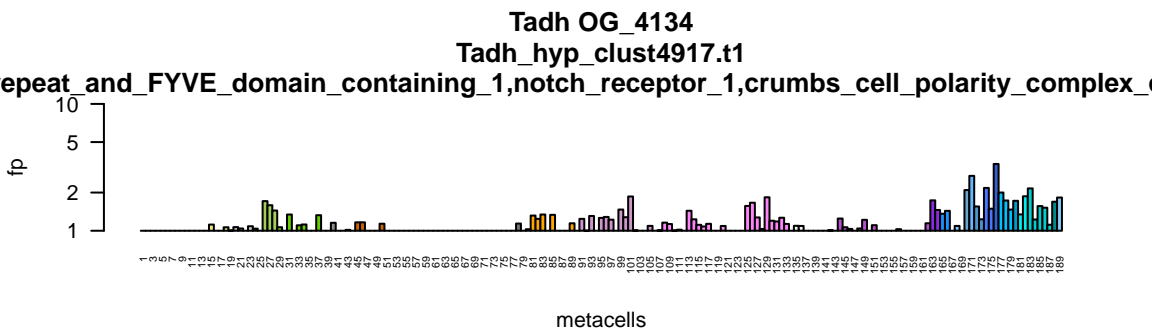
metacells

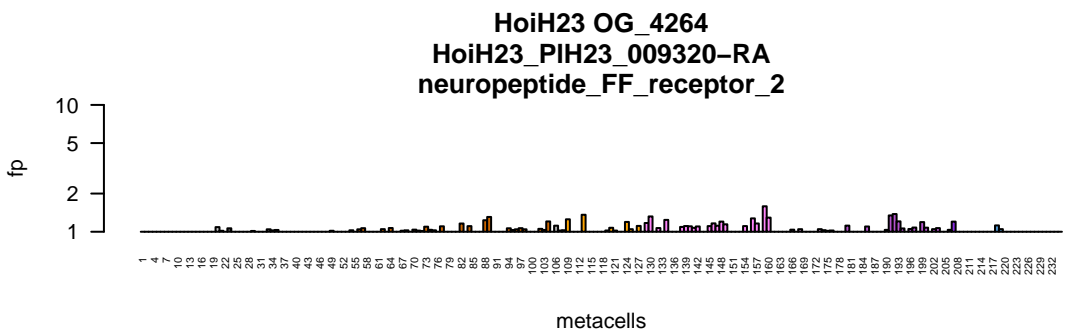
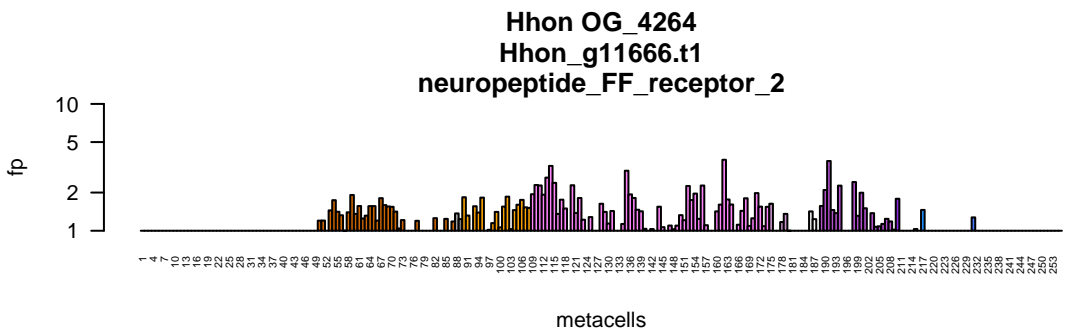
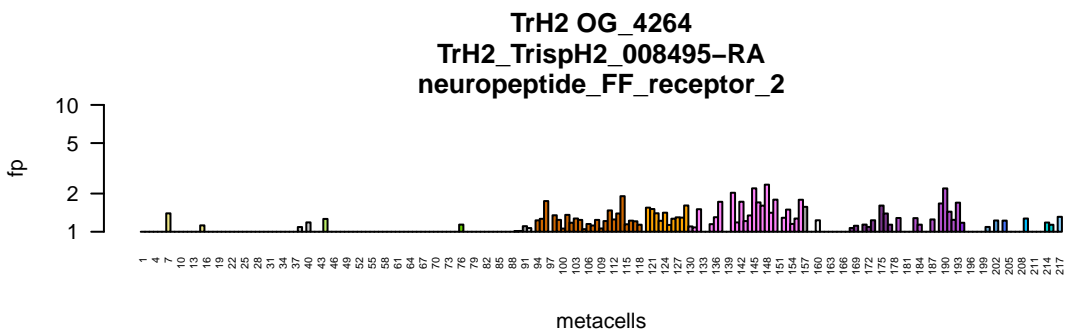
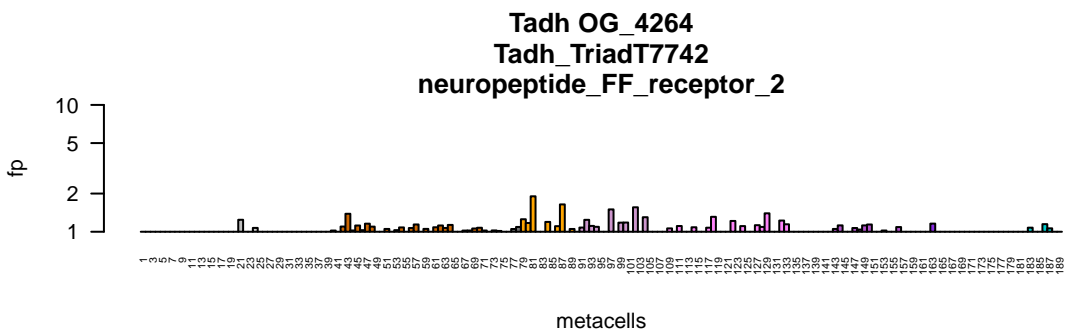
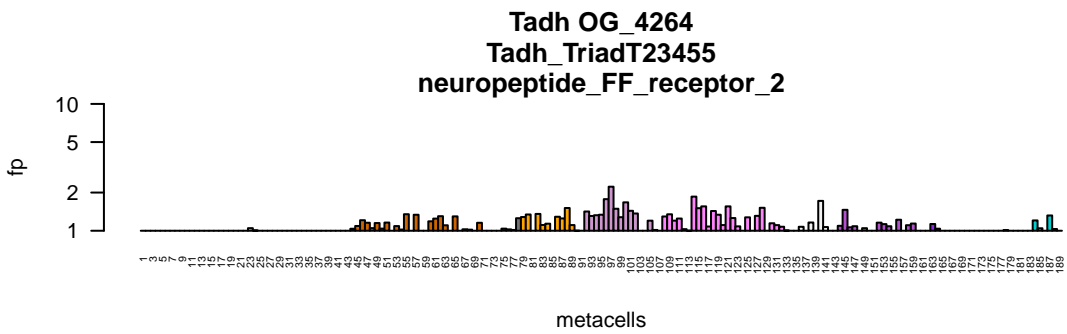


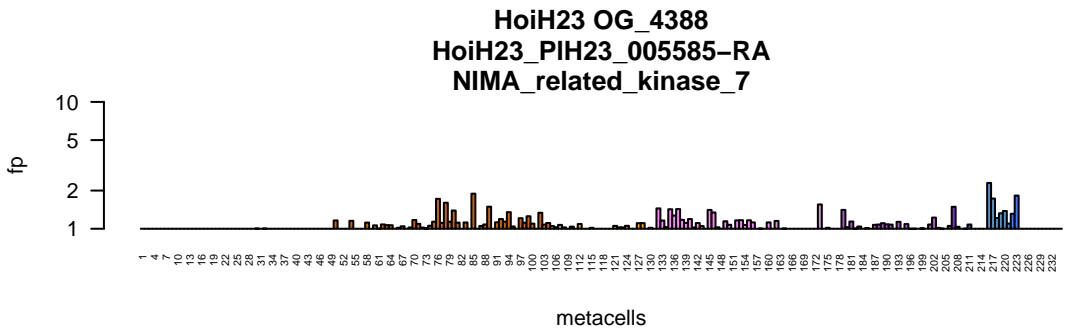
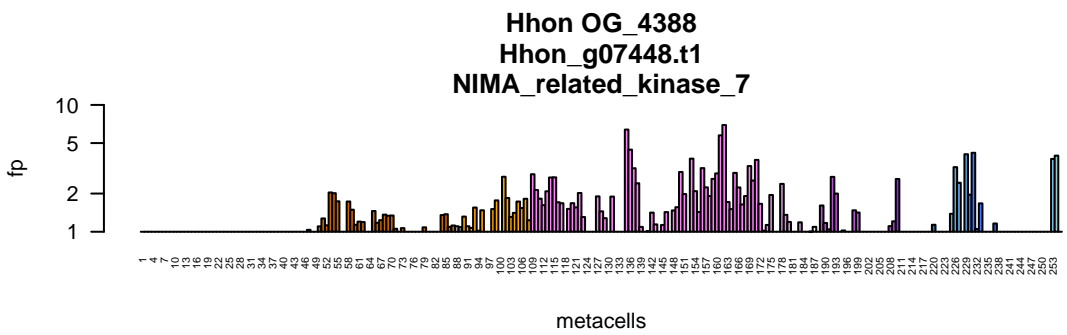
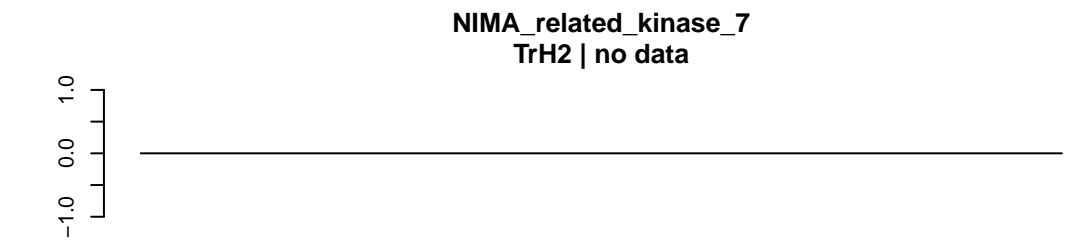
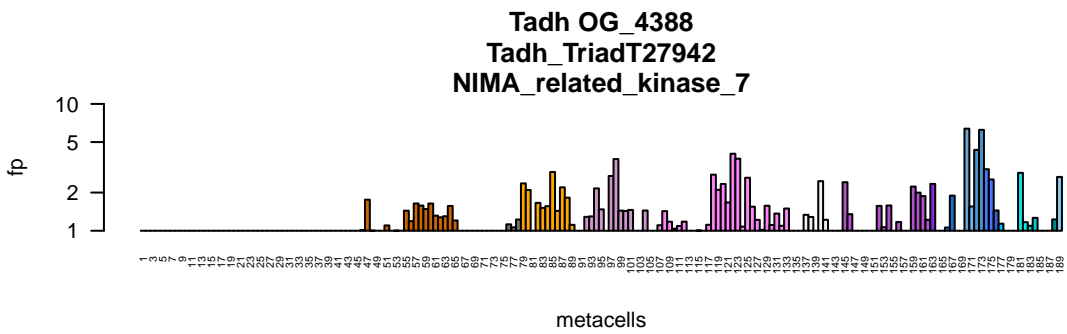


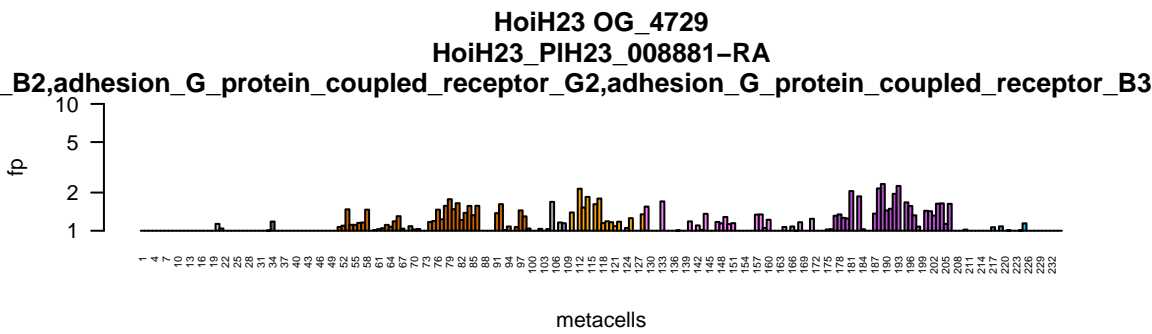
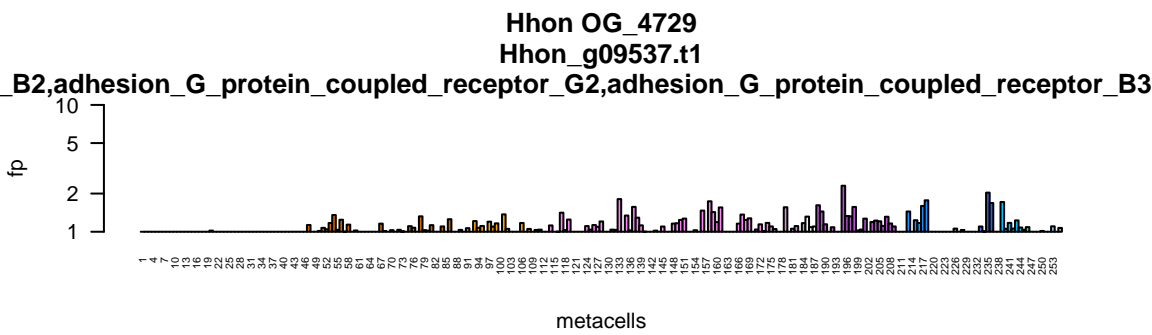
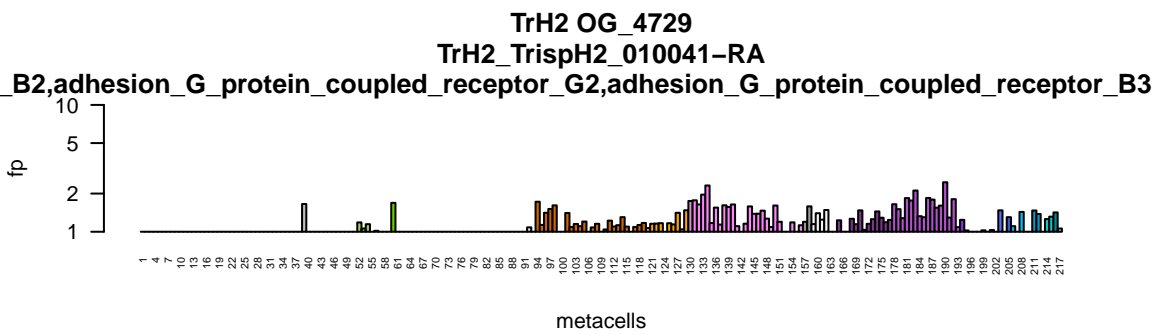
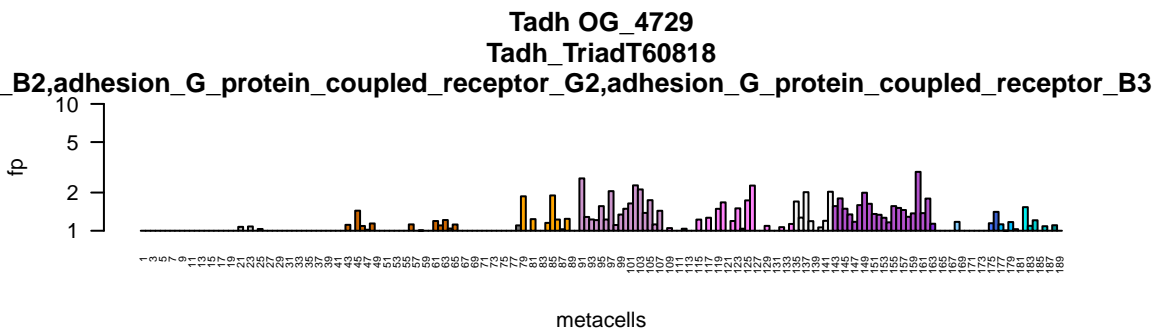


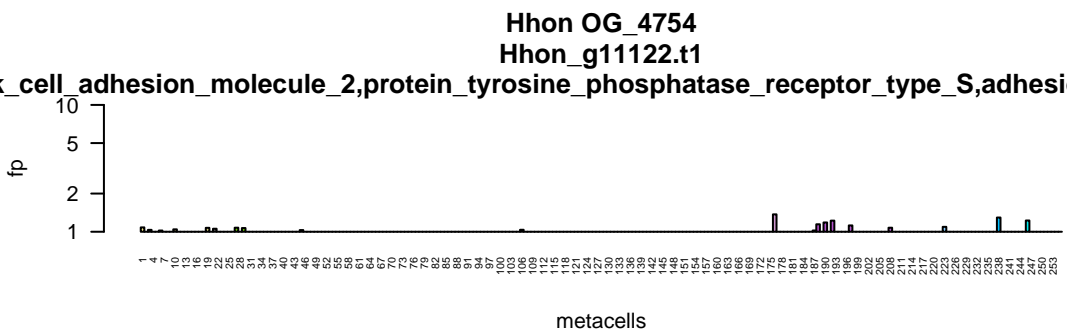
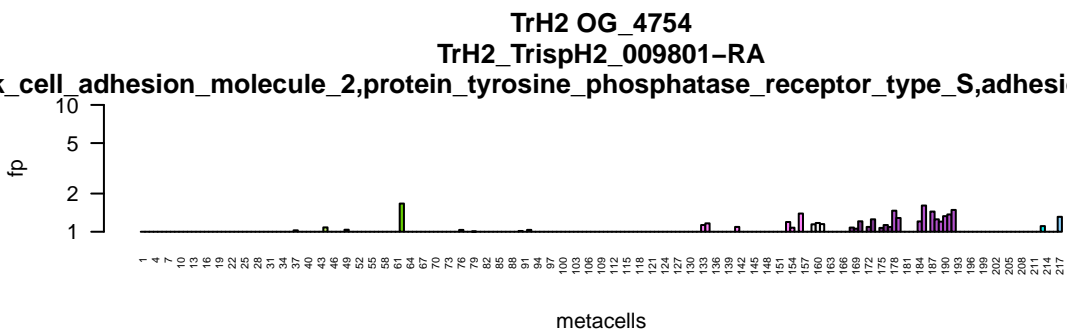
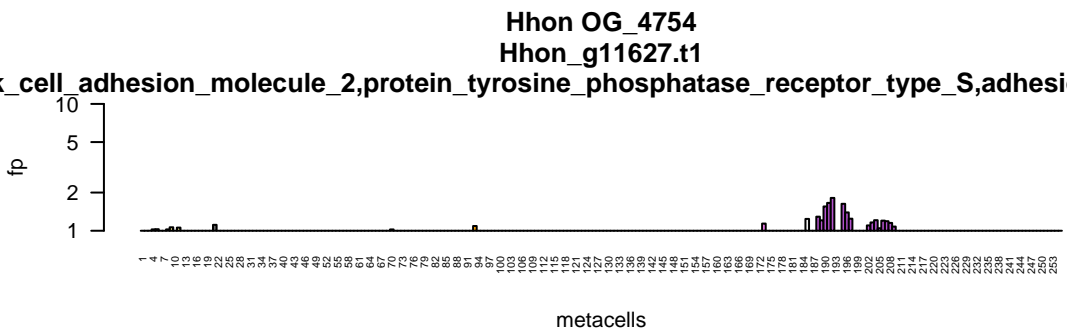
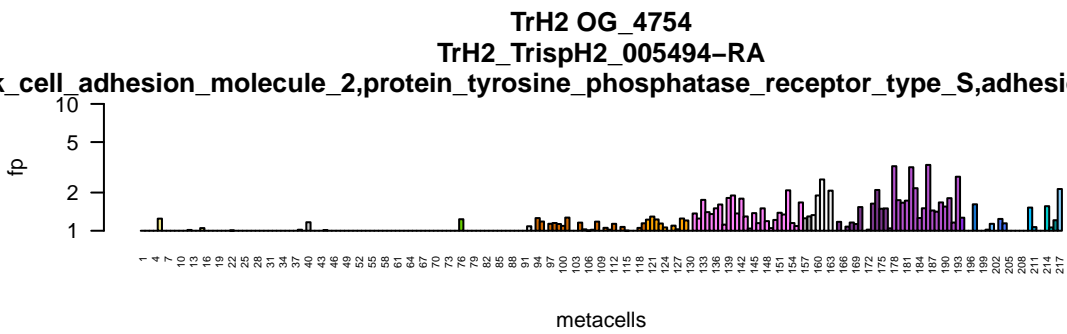
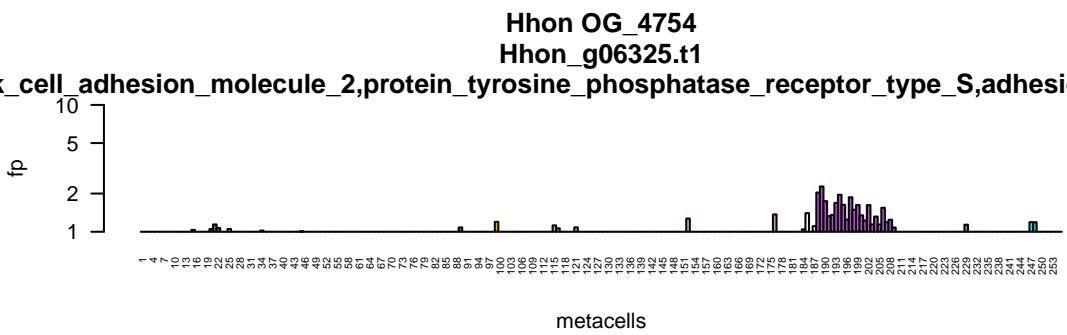
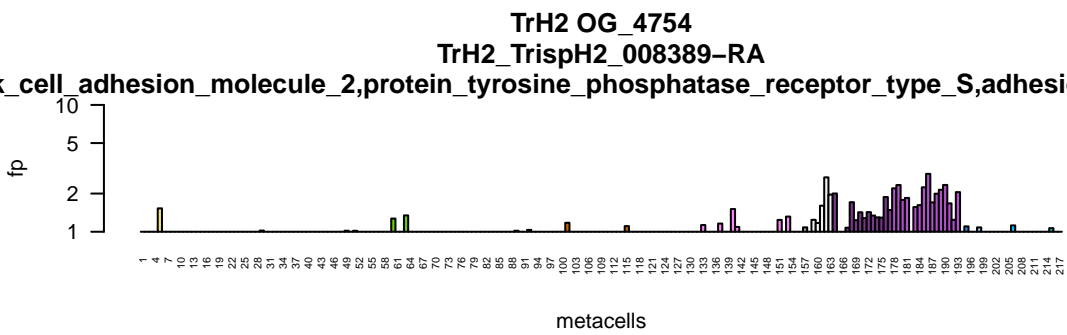
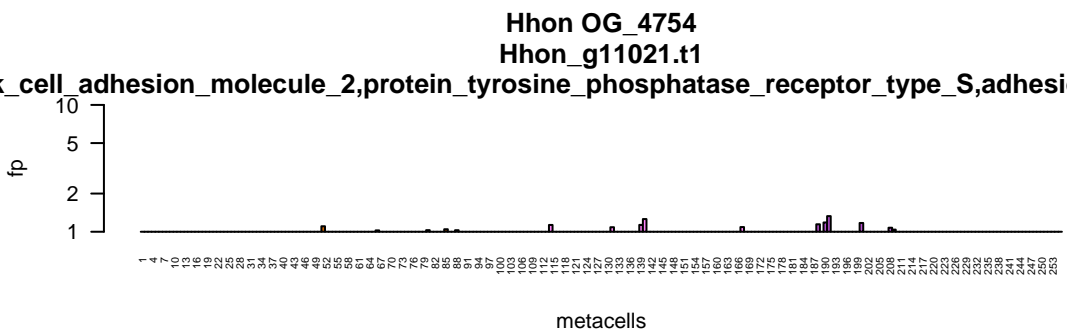
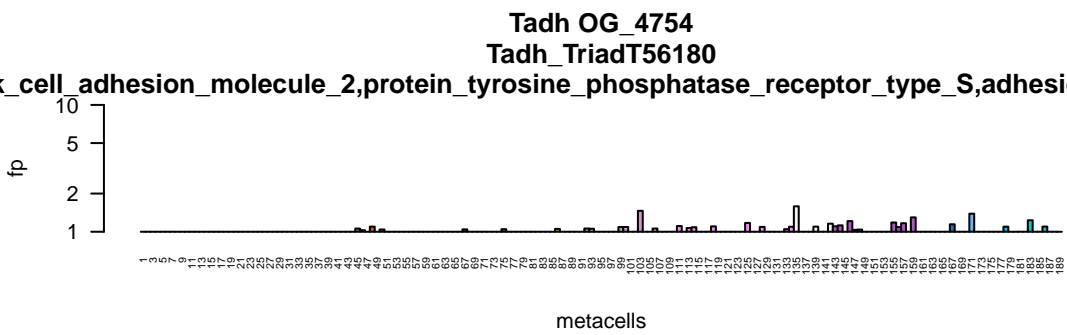
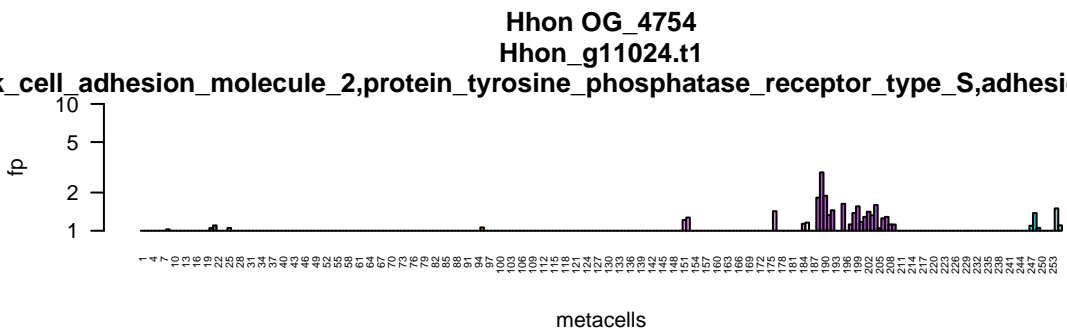
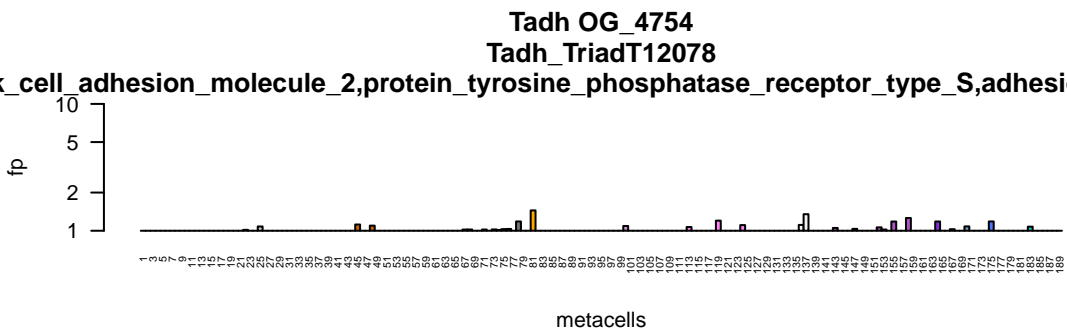
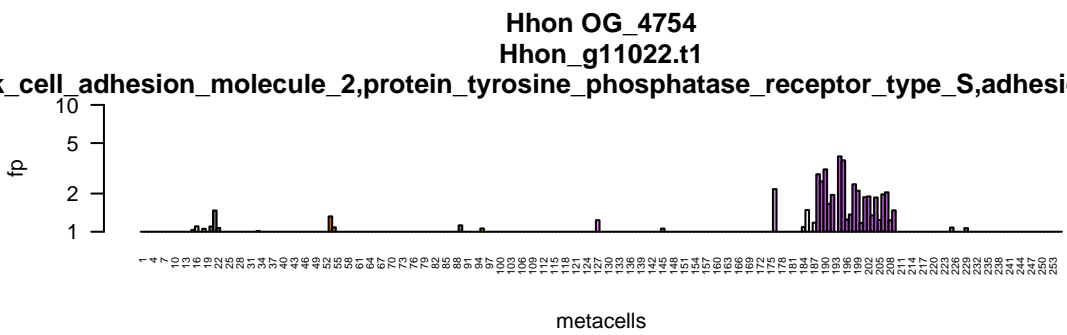
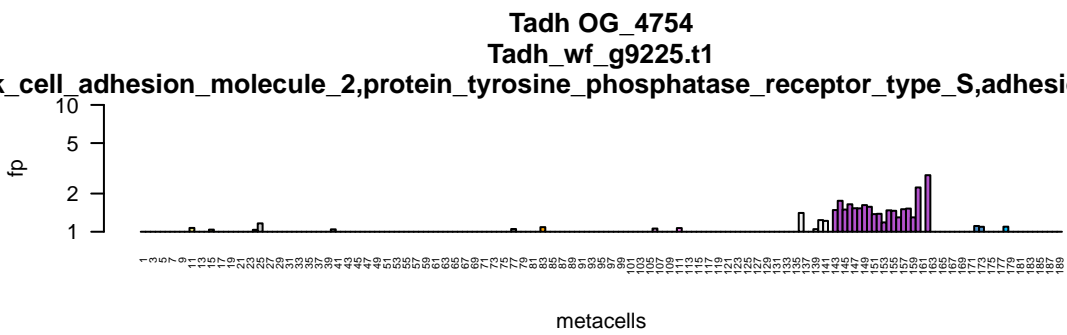
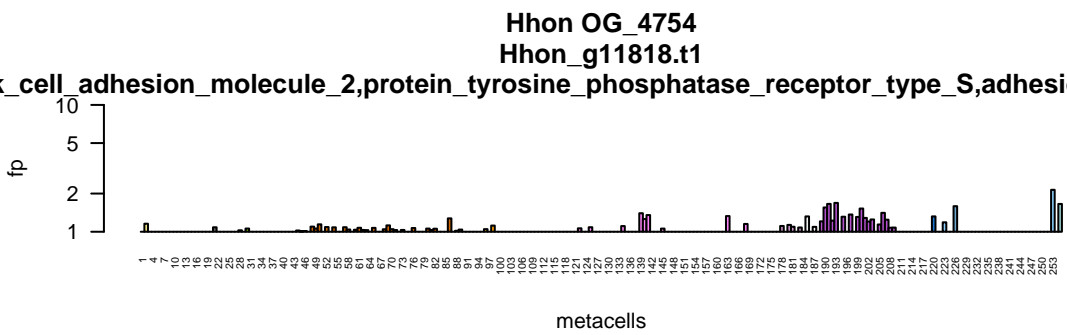
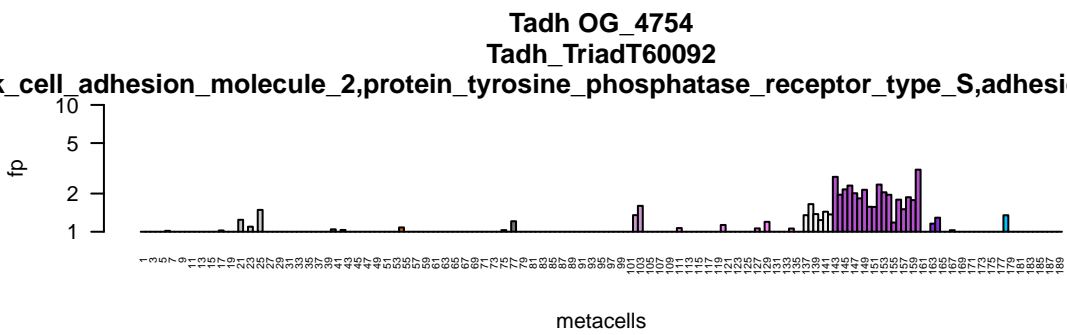
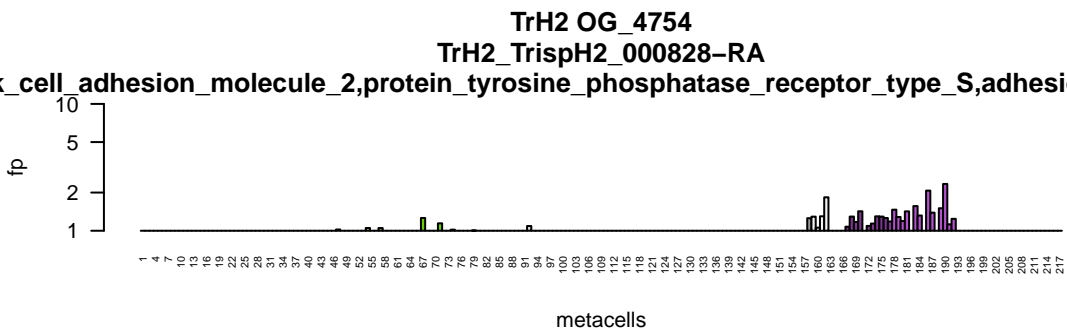
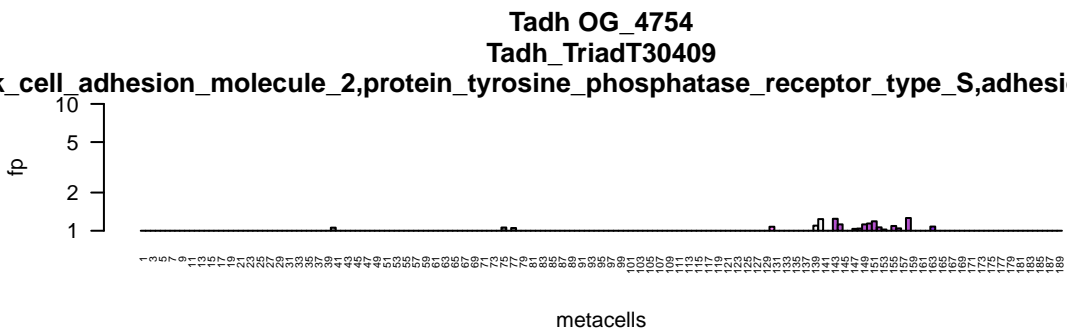


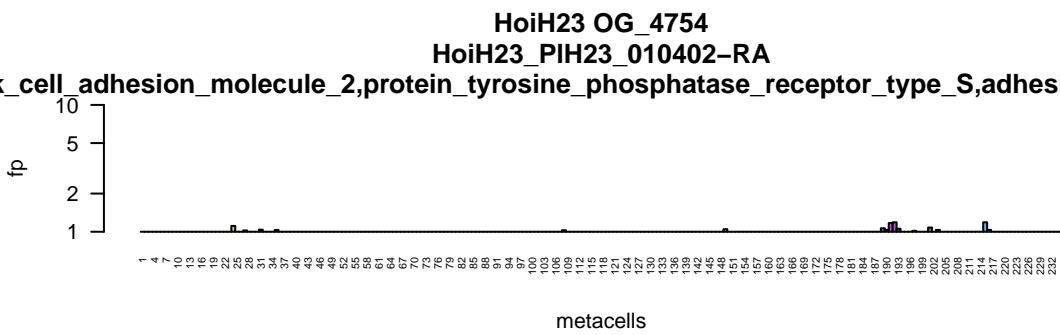
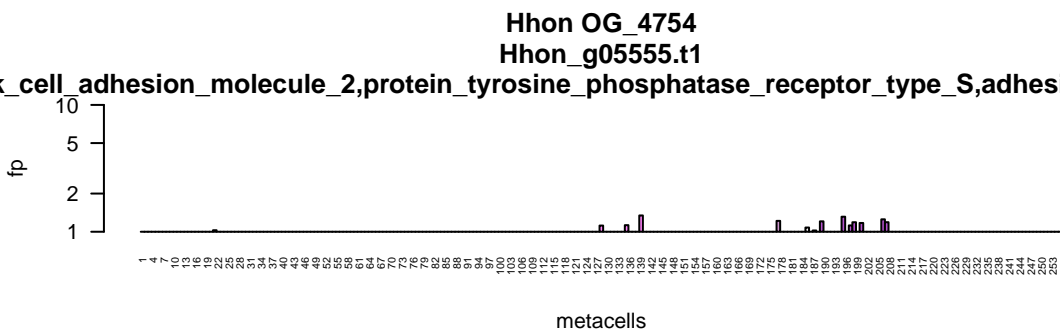
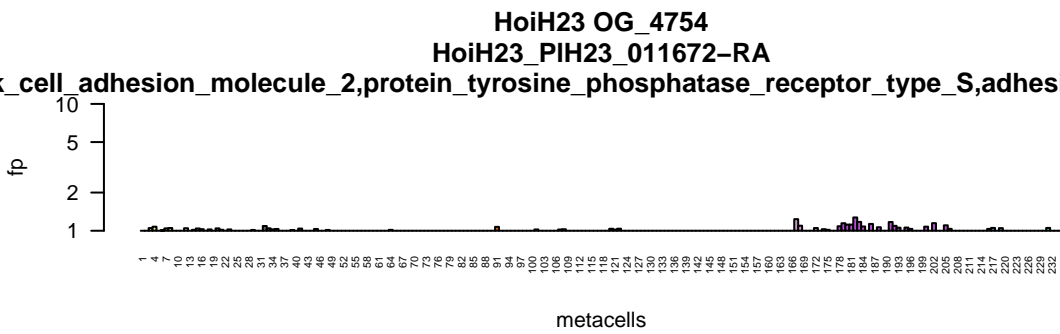
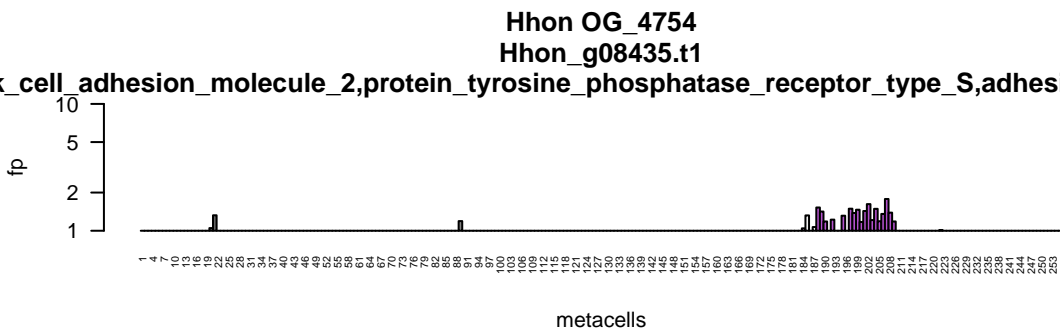
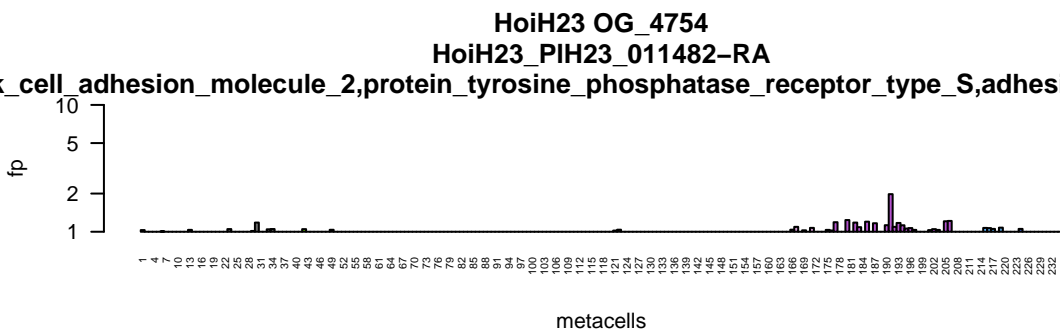
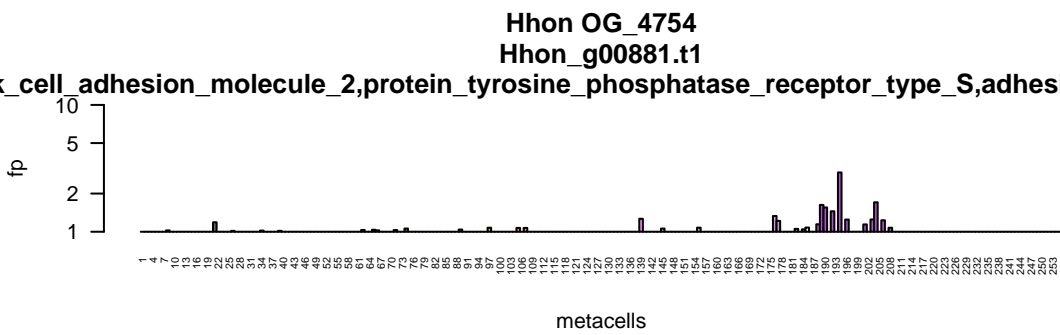
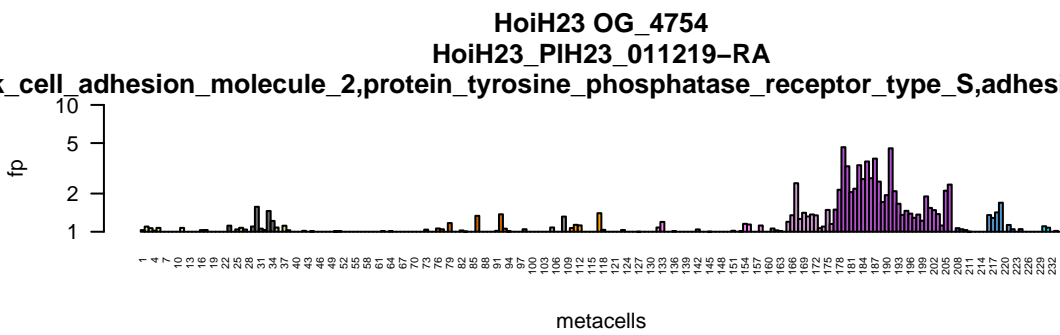
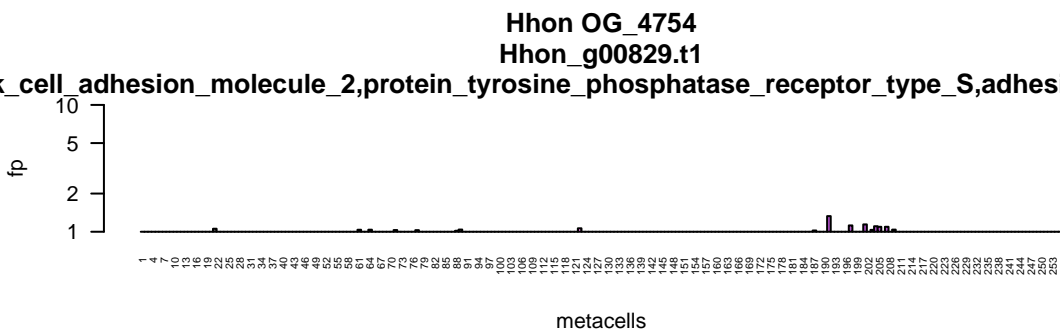
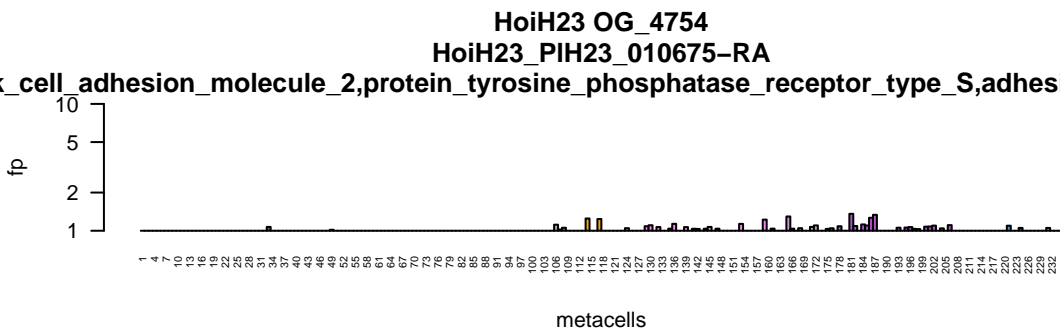
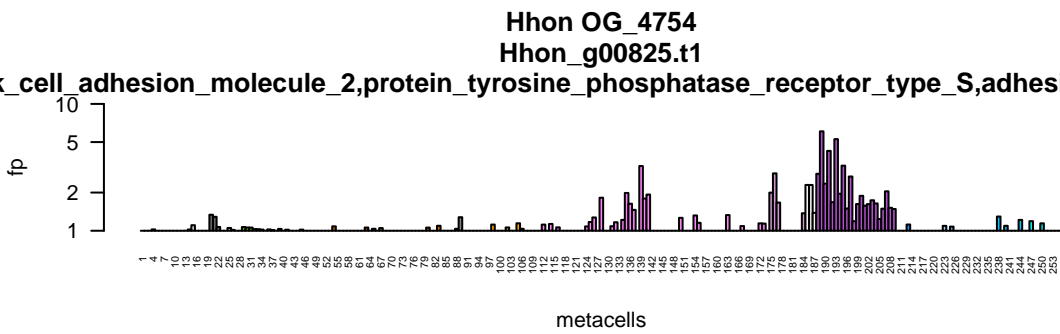
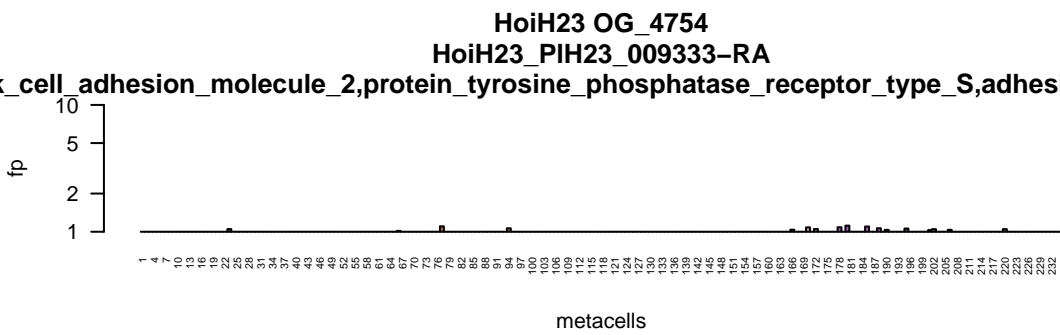
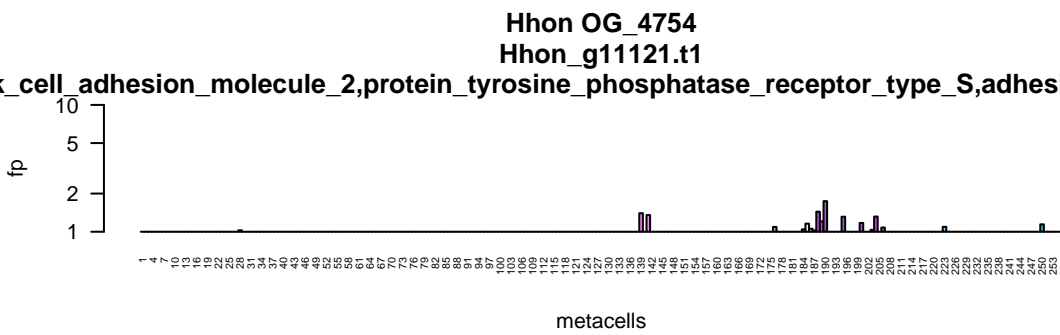
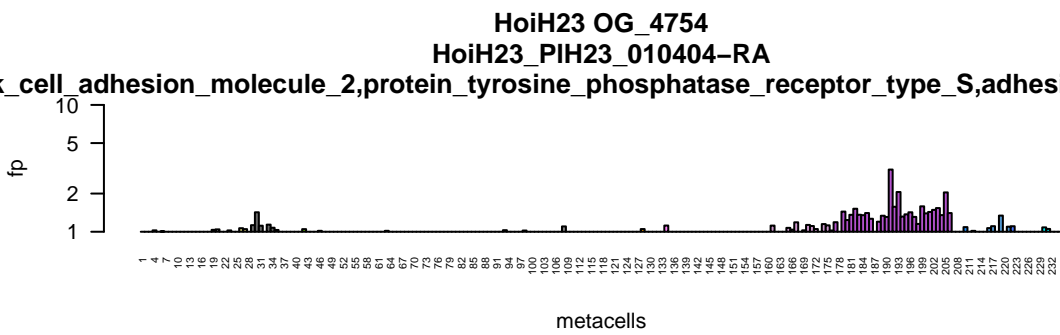
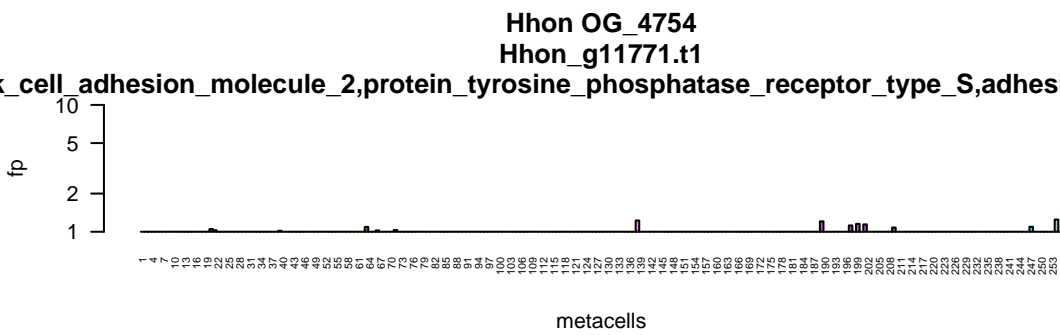
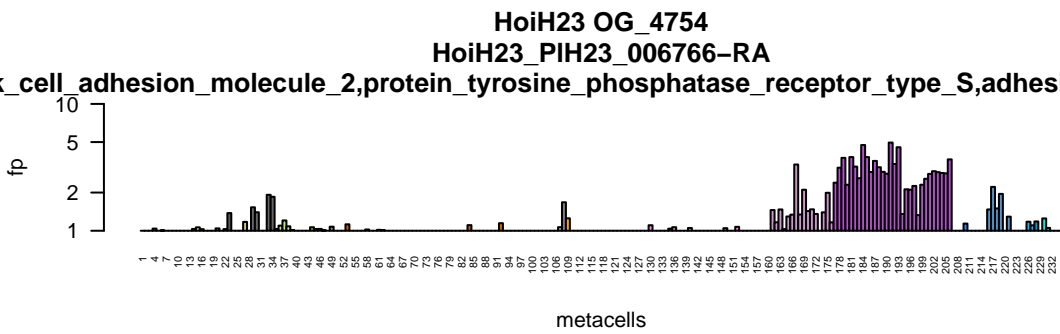
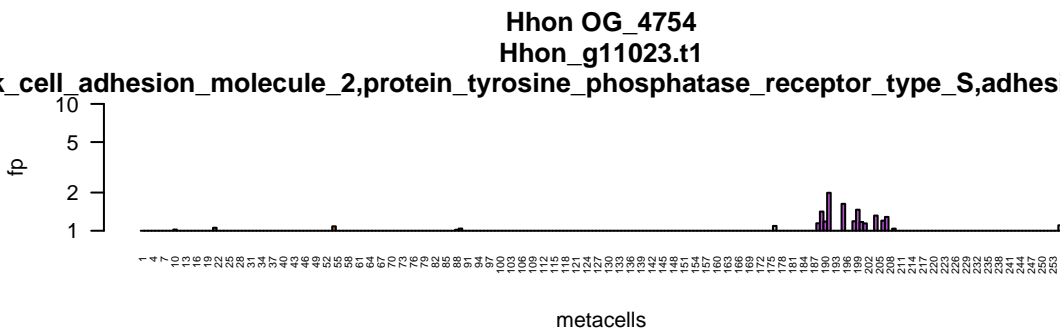












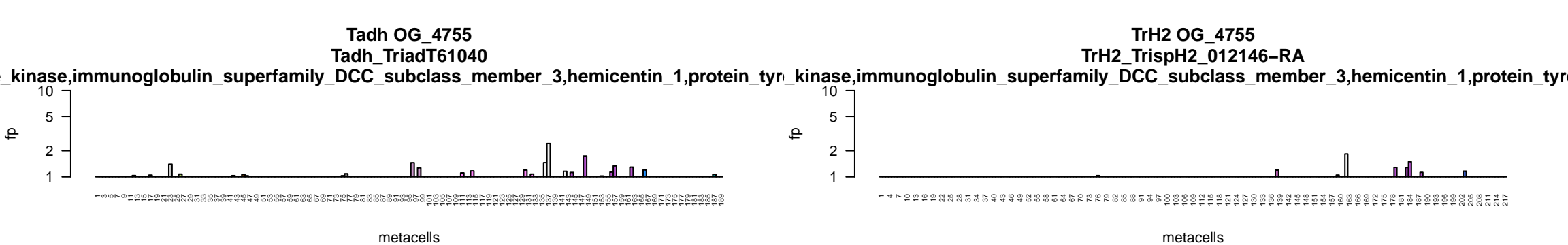


fp

metacells

fp

metacells



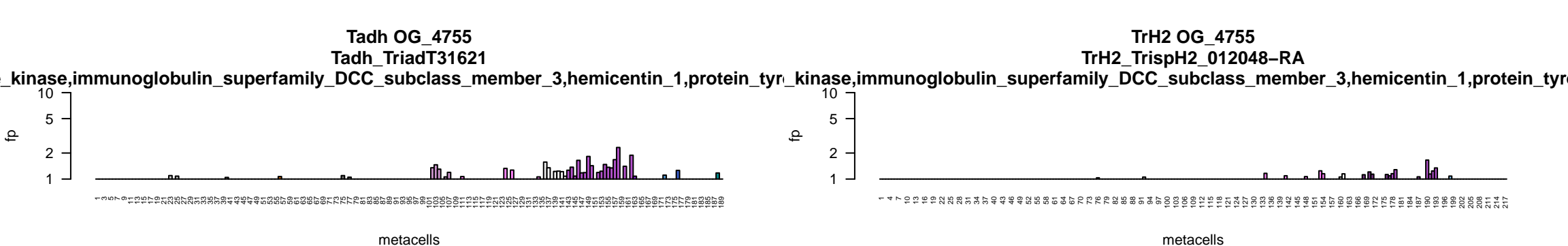
kinase,immunoglobulin\_superfamily\_DCC\_subclass\_member\_3,hemicentin\_1,protein\_tyr\_kinase,immunoglobulin\_superfamily\_DCC\_subclass\_member\_3,hemicentin\_1,protein\_tyr\_kinase,immunoglobulin\_superfamily\_DCC\_subclass\_member\_3,hemicentin\_1,protein\_tyr\_kinase

fp

metacells

fp

metacells



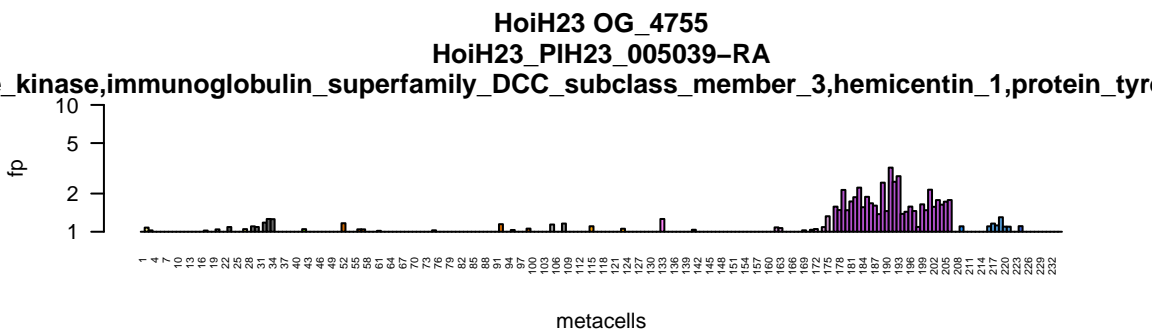
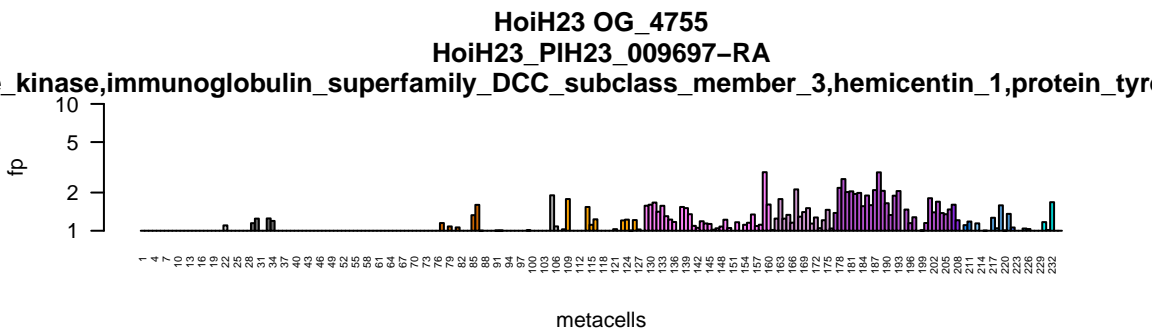
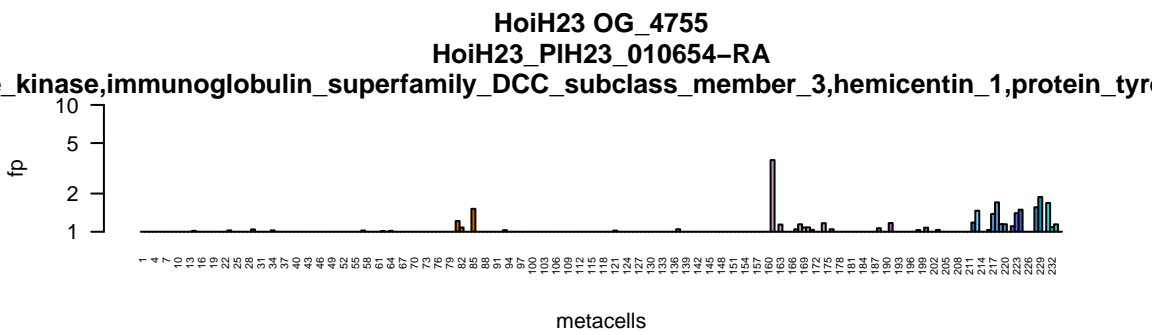
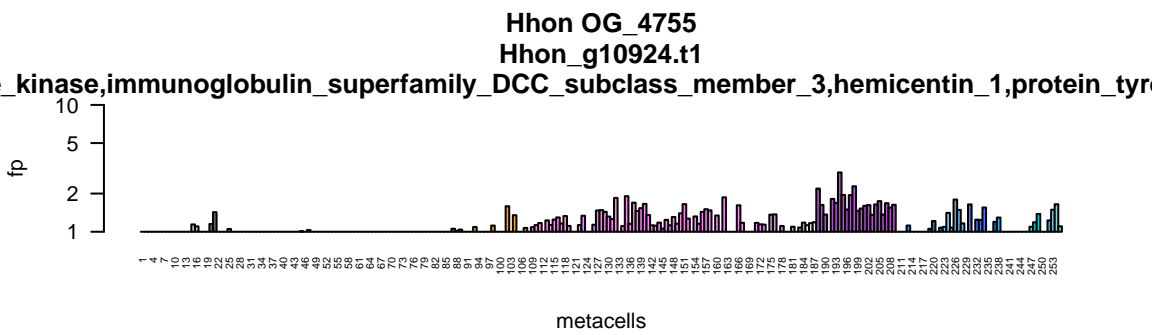
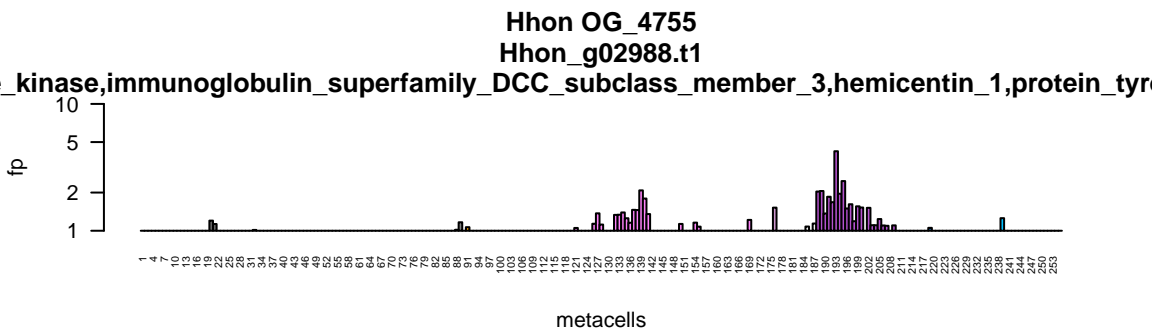
kinase,immunoglobulin\_superfamily\_DCC\_subclass\_member\_3,hemicentin\_1,protein\_tyr\_kinase,immunoglobulin\_superfamily\_DCC\_subclass\_member\_3,hemicentin\_1,protein\_tyr\_kinase

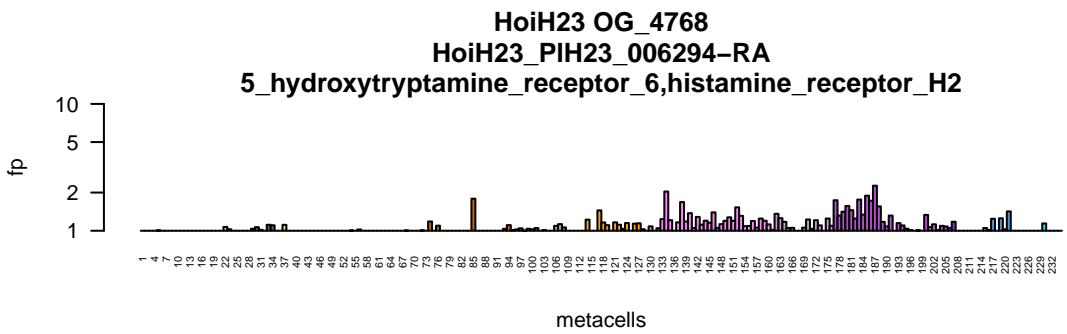
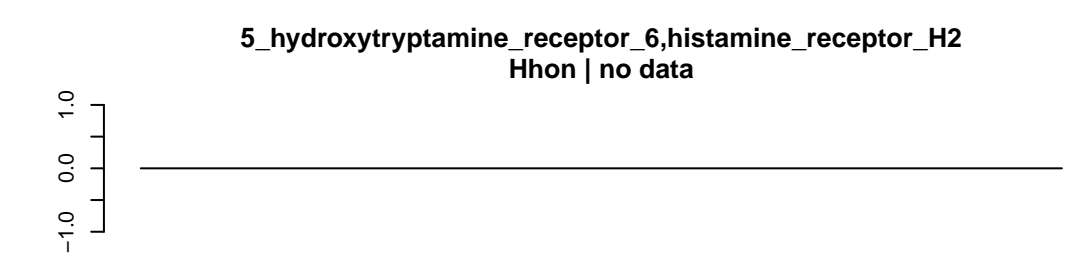
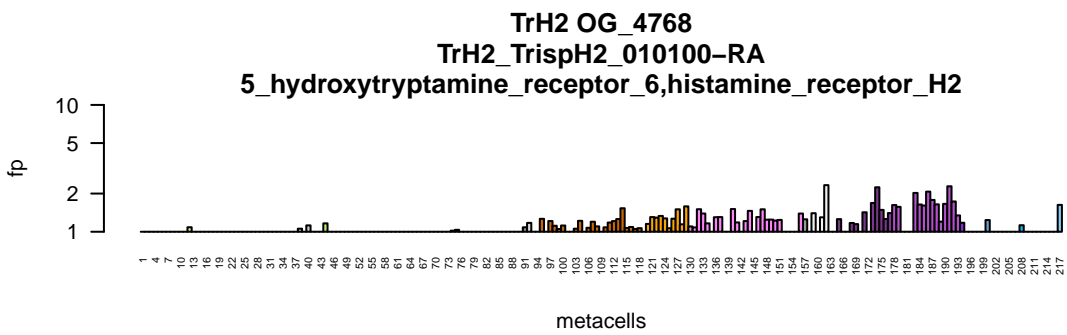
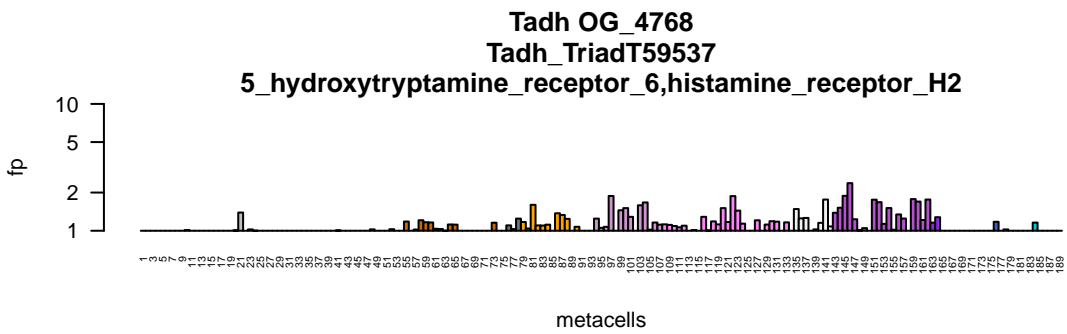
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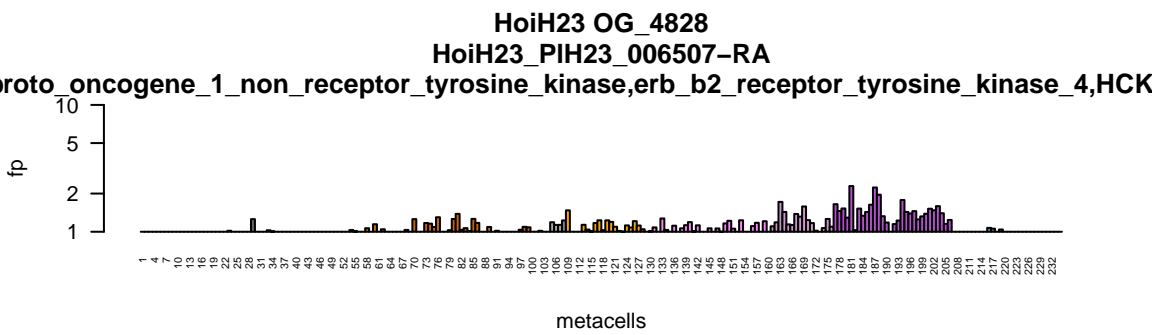
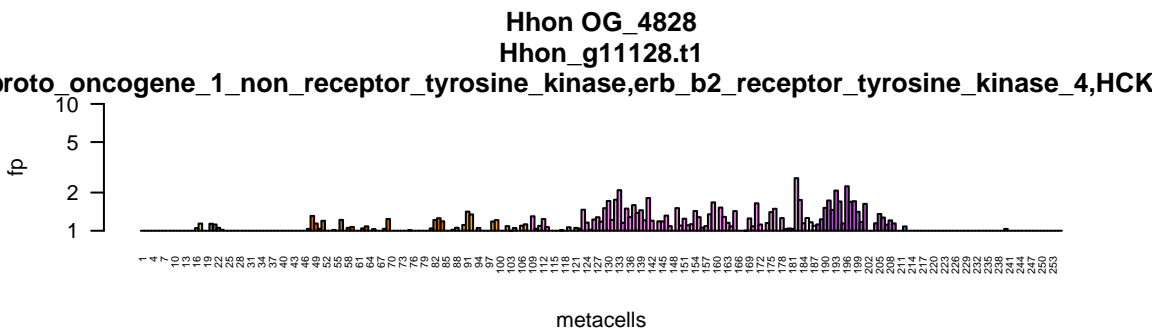
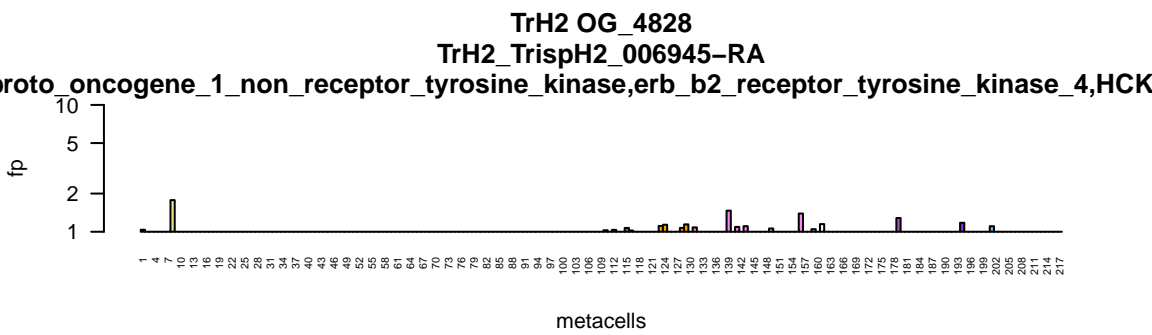
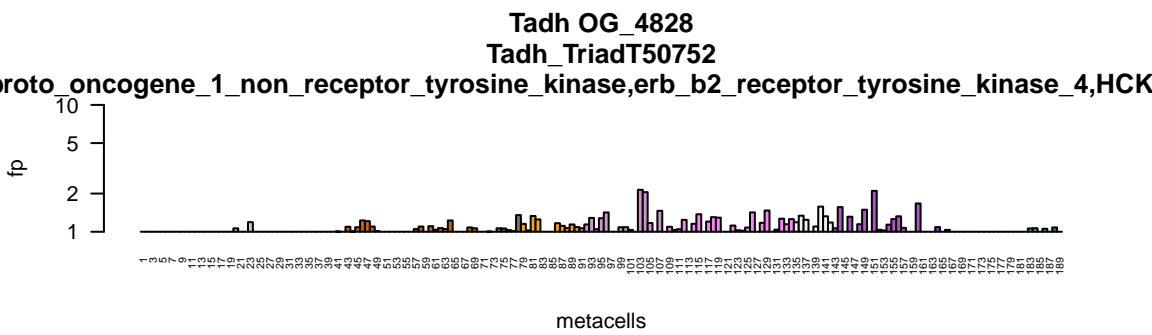
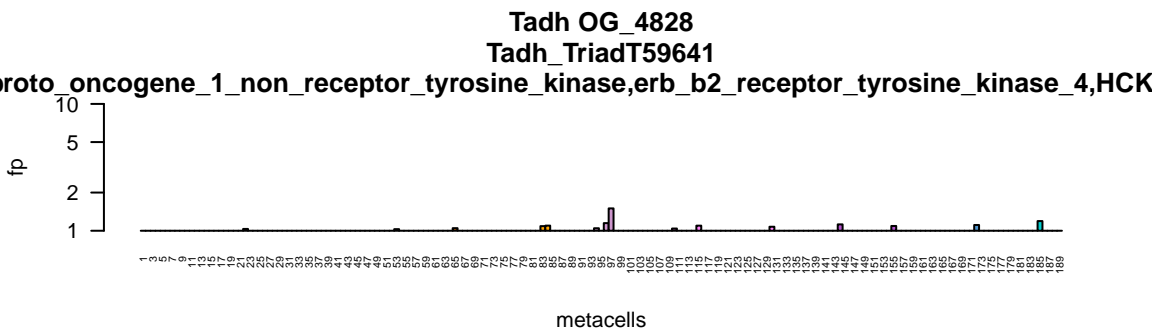
metacells

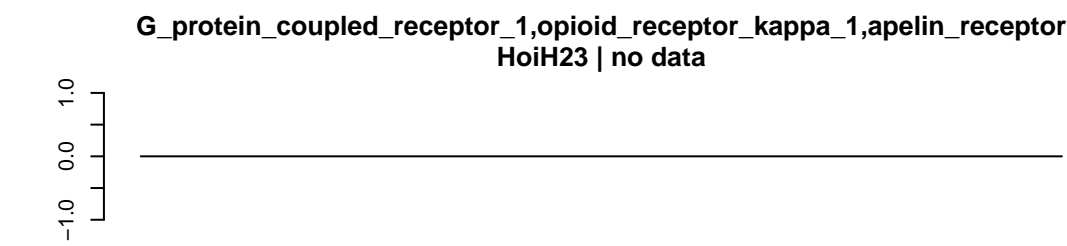
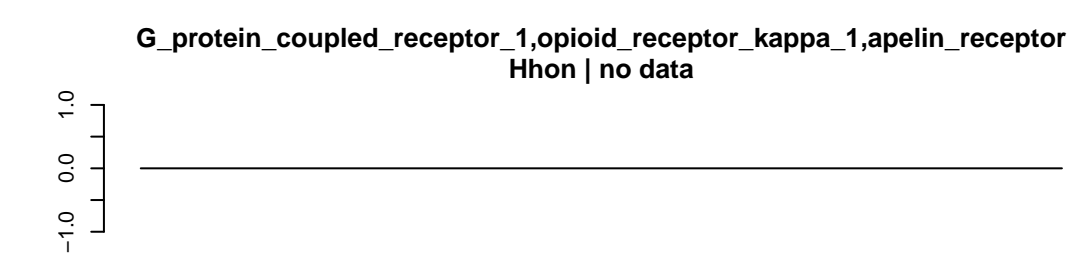
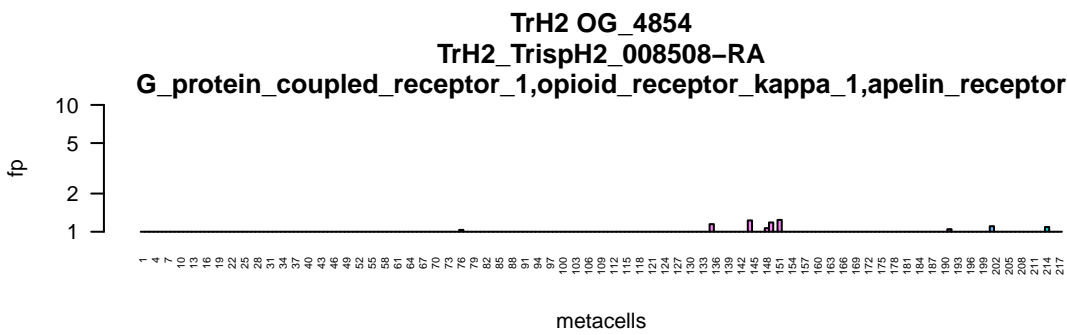
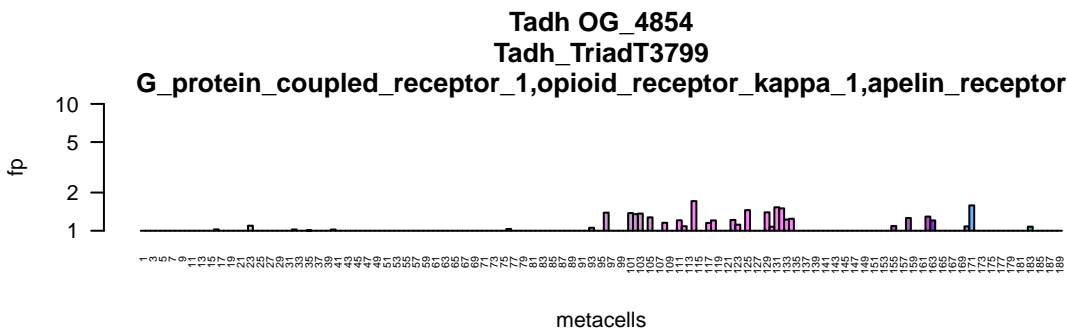
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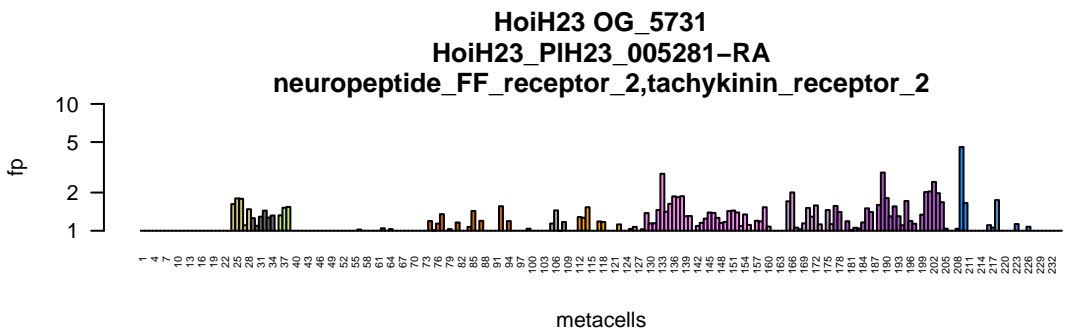
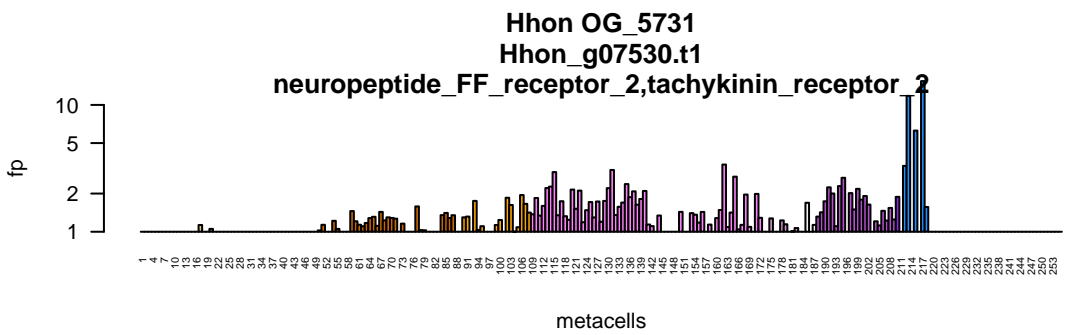
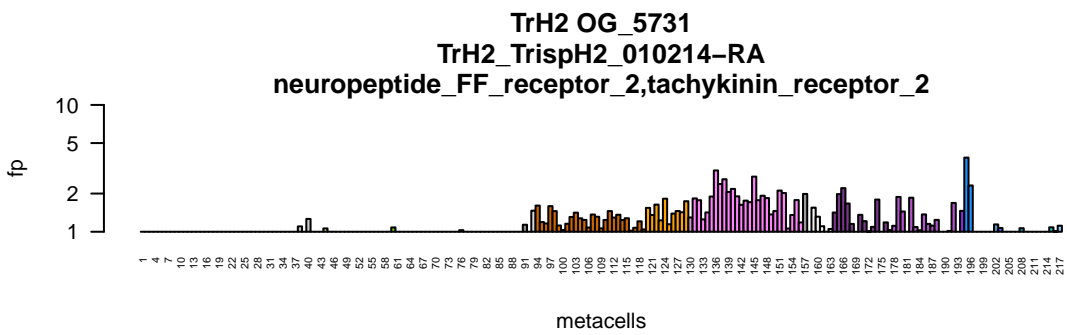
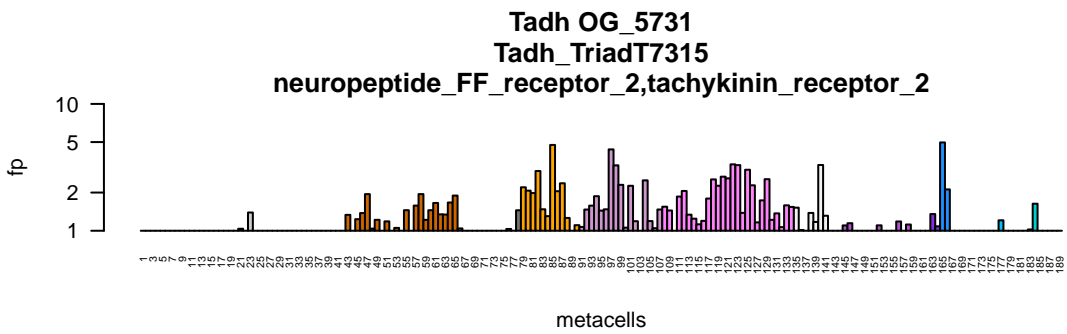
metacells



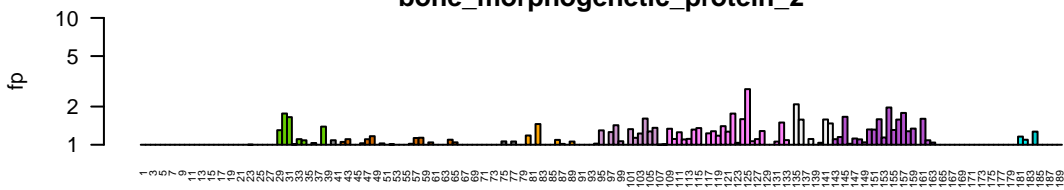






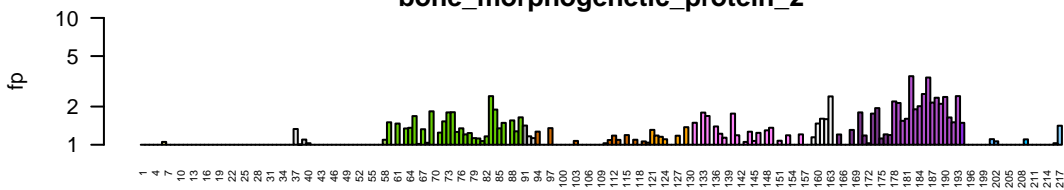


**Tadh OG\_5897**  
**Tadh\_TriadT57057**  
**bone\_morphogenetic\_protein\_2**



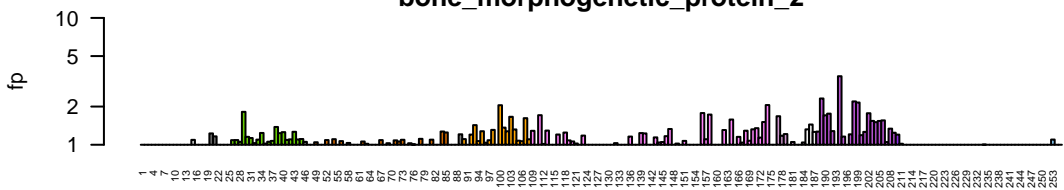
metacells

**TrH2 OG\_5897**  
**TrH2\_TrispH2\_002393-RA**  
**bone\_morphogenetic\_protein\_2**



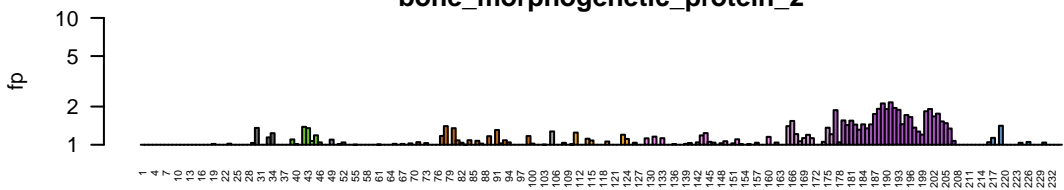
metacells

**Hhon OG\_5897**  
**Hhon\_g07488.t1**  
**bone\_morphogenetic\_protein\_2**

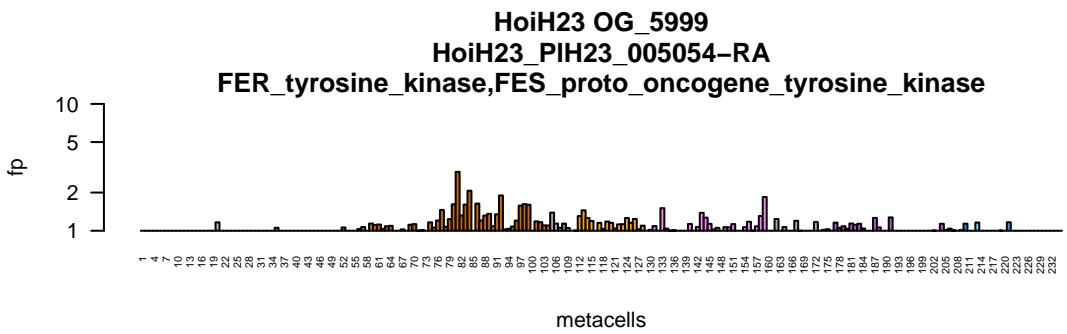
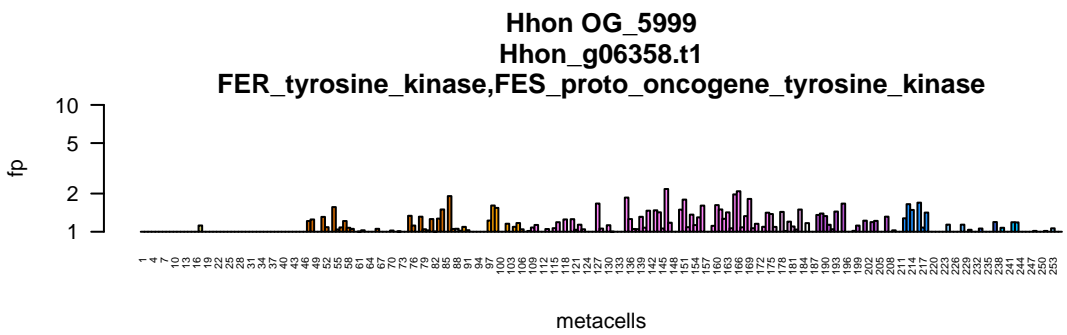
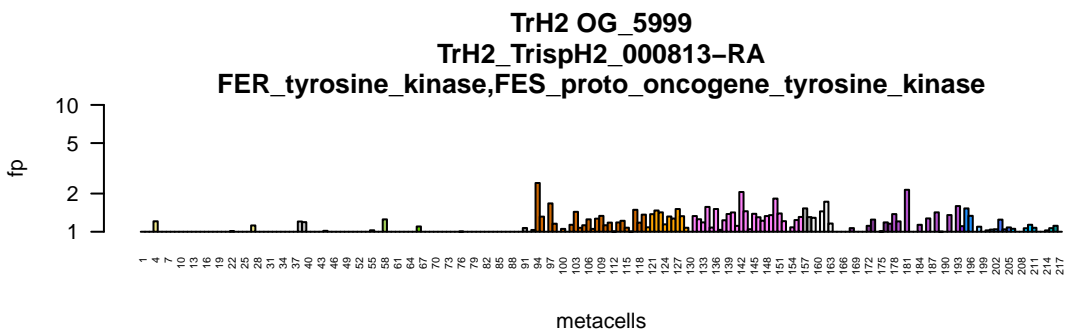
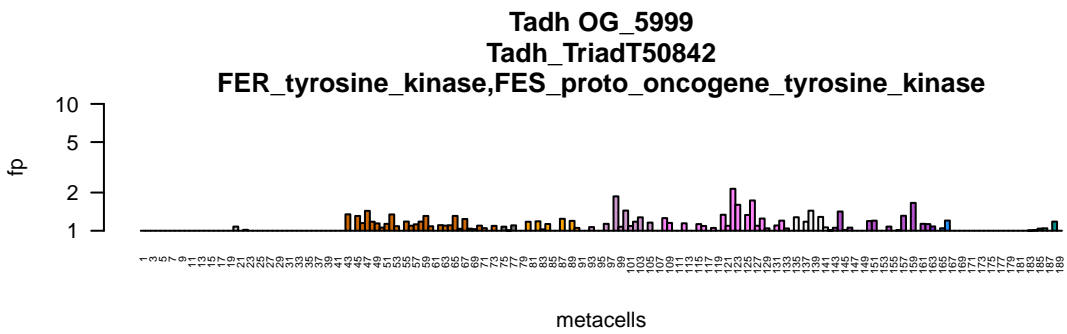


metacells

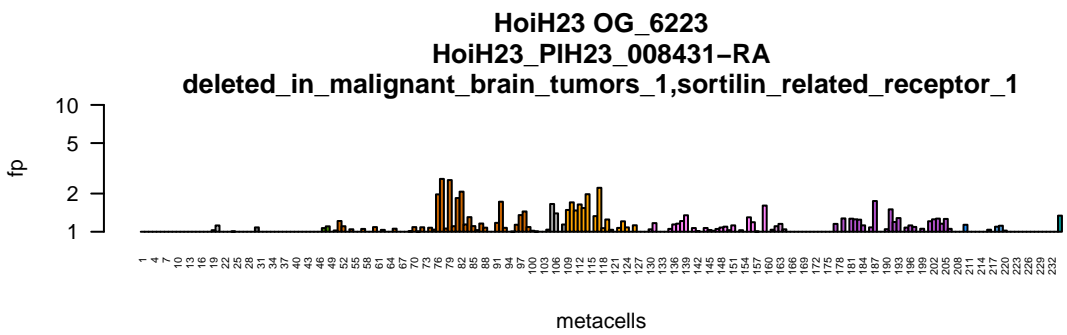
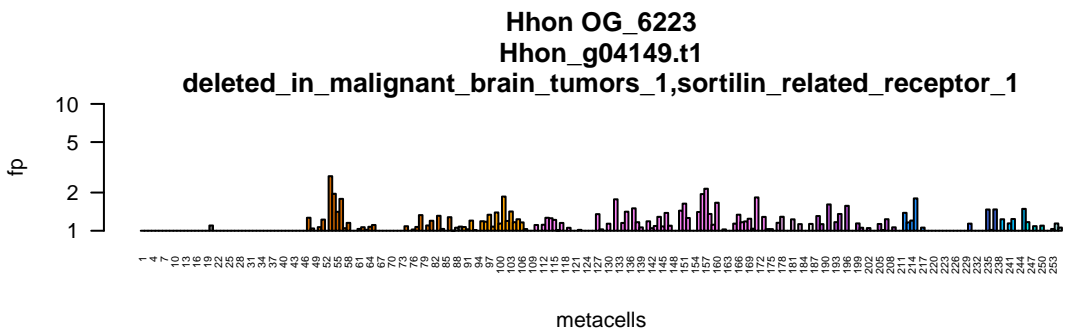
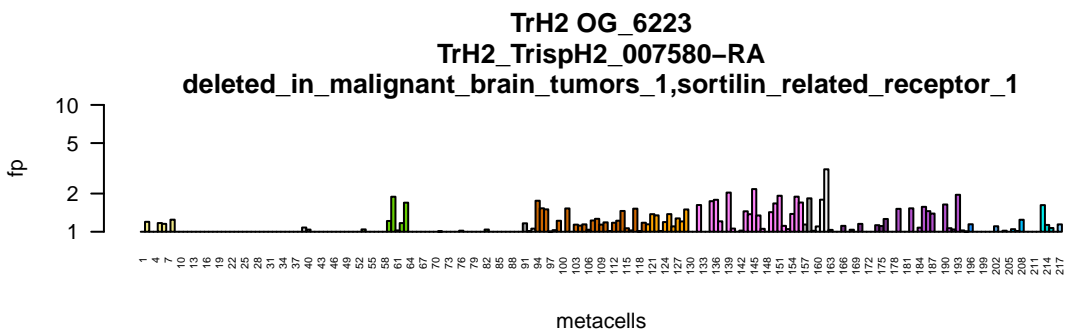
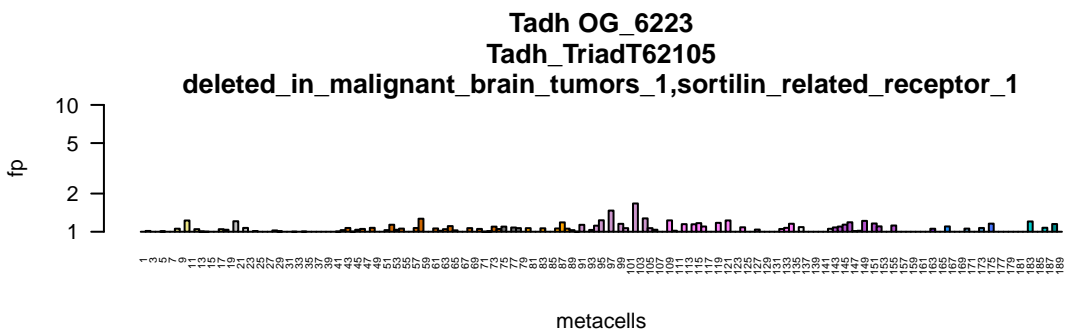
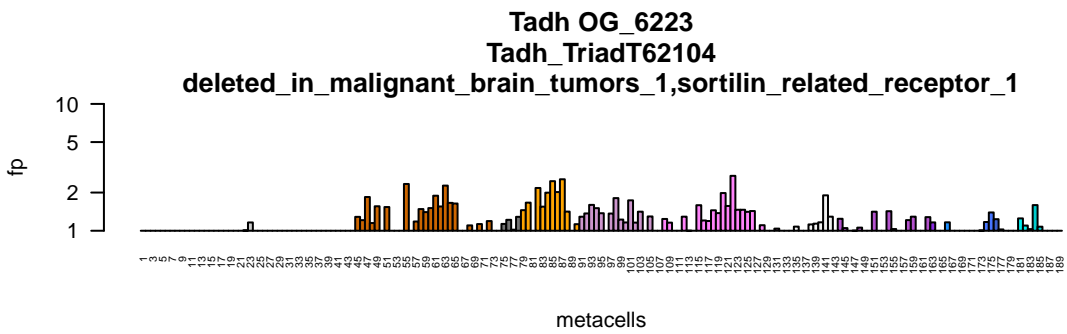
**HoiH23 OG\_5897**  
**HoiH23\_PIH23\_002971-RA**  
**bone\_morphogenetic\_protein\_2**

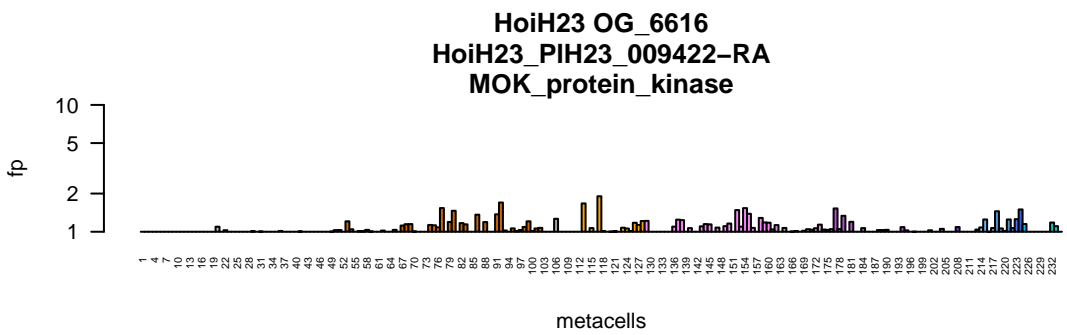
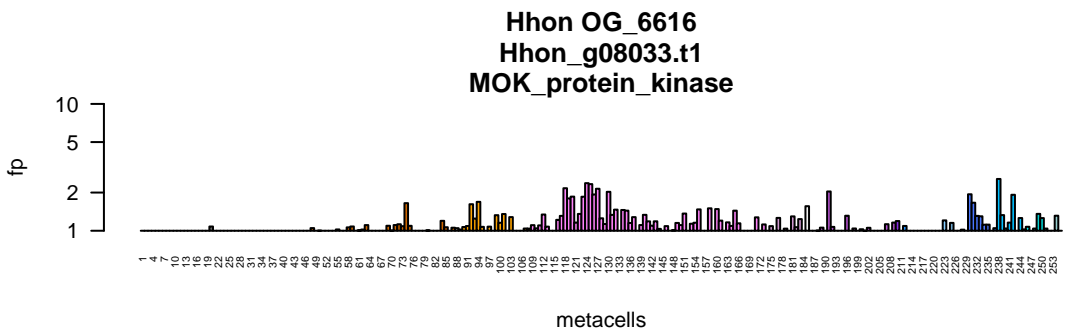
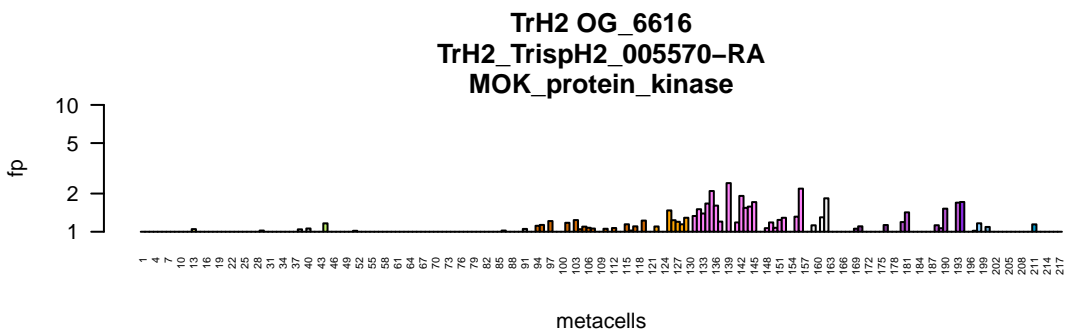
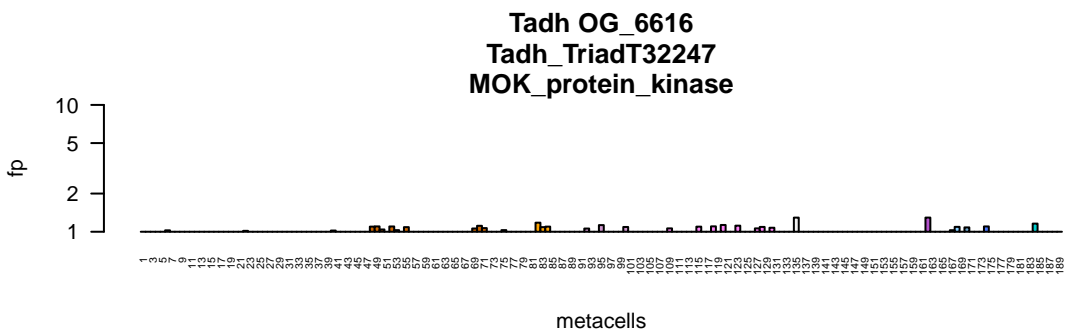
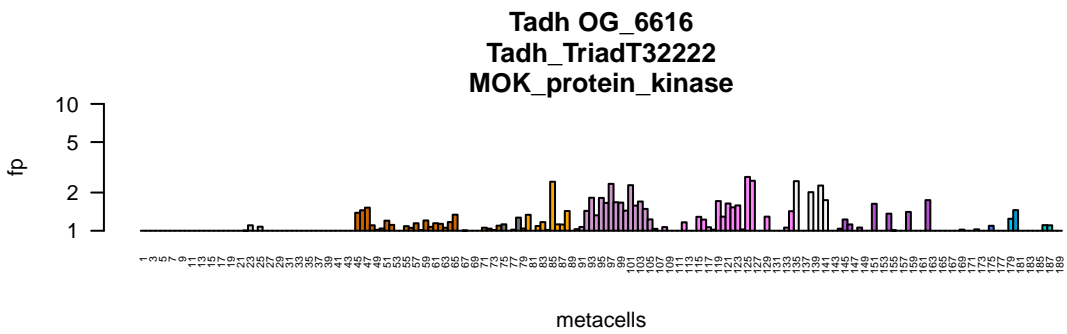


metacells

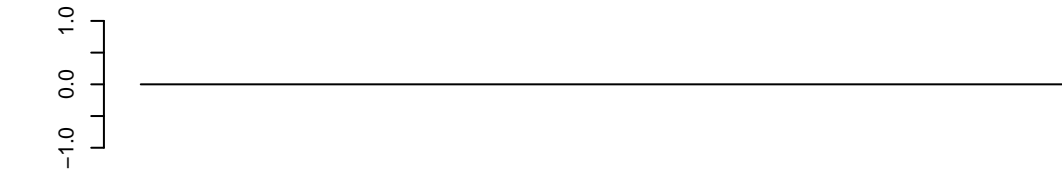




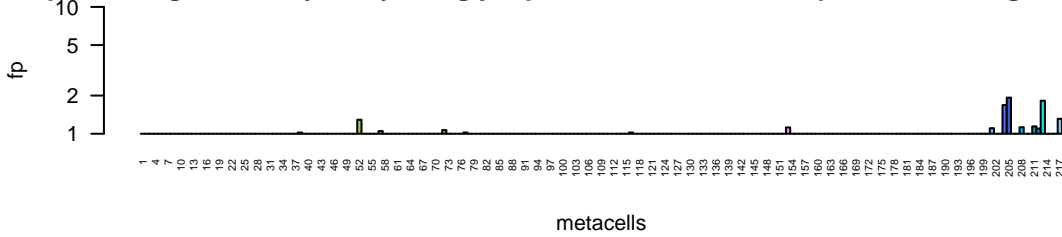




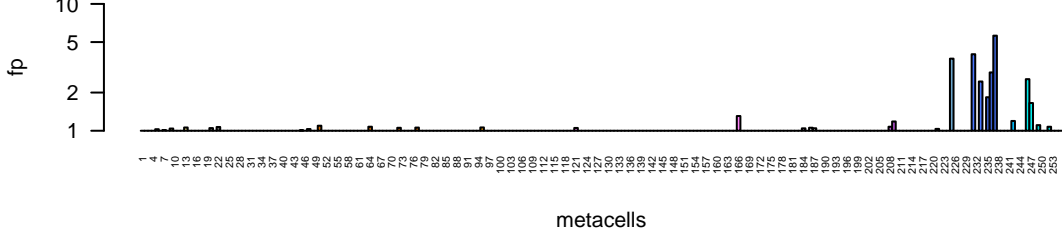
receptor\_1,oligodendrocyte\_myelin\_glycoprotein,leucine\_rich\_repeat\_containing\_40,relaxir  
Tadh | no data



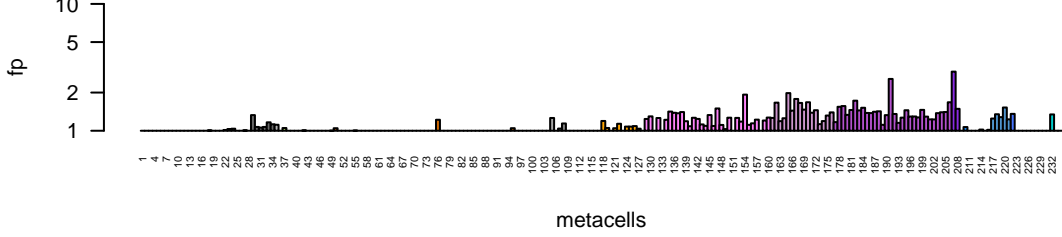
TrH2 OG\_6800  
TrH2\_TrispH2\_006274-RA  
receptor\_1,oligodendrocyte\_myelin\_glycoprotein,leucine\_rich\_repeat\_containing\_40,relaxir



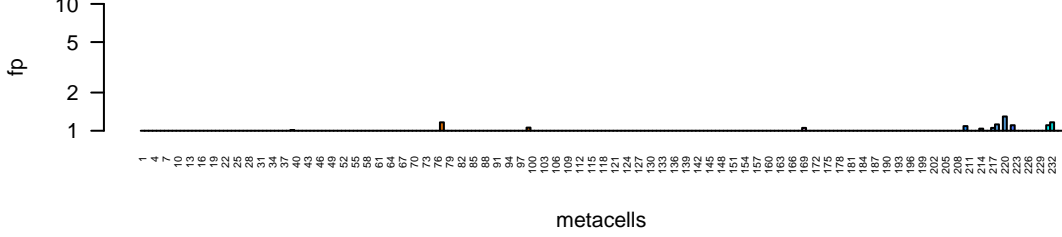
Hhon OG\_6800  
Hhon\_g04761.t1  
receptor\_1,oligodendrocyte\_myelin\_glycoprotein,leucine\_rich\_repeat\_containing\_40,relaxir

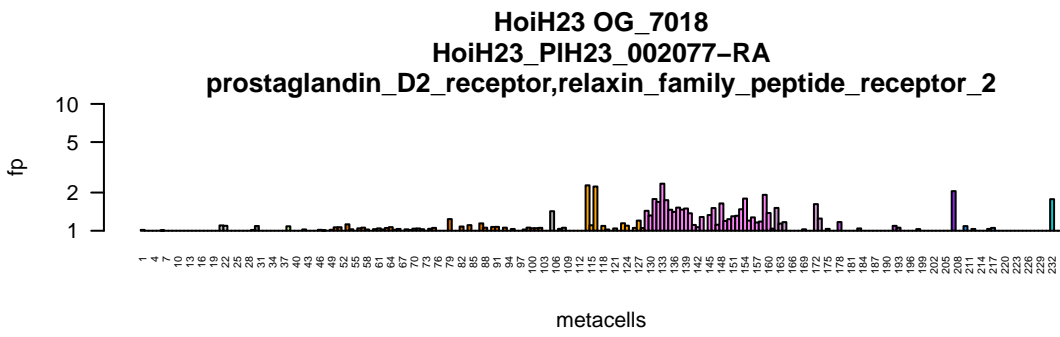
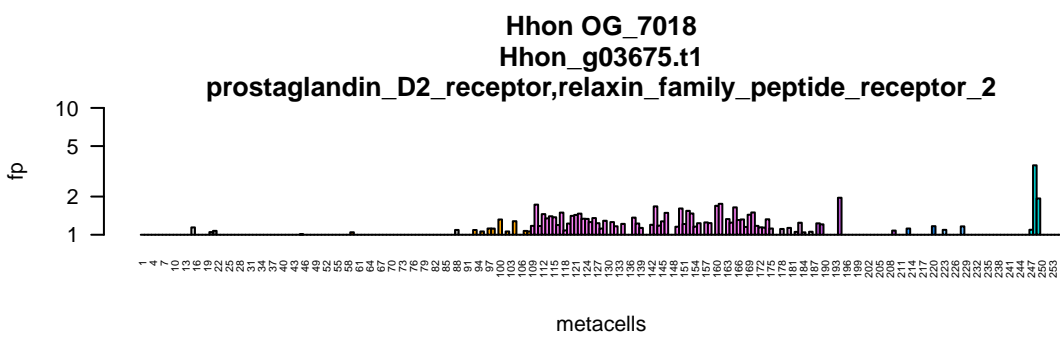
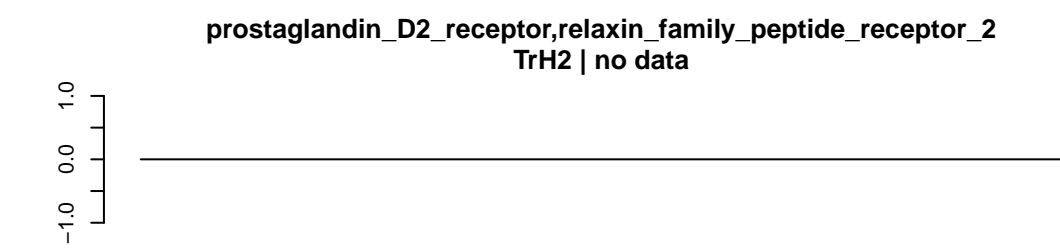
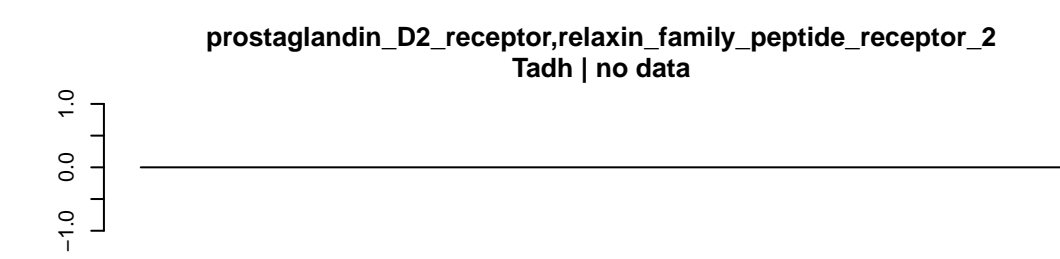


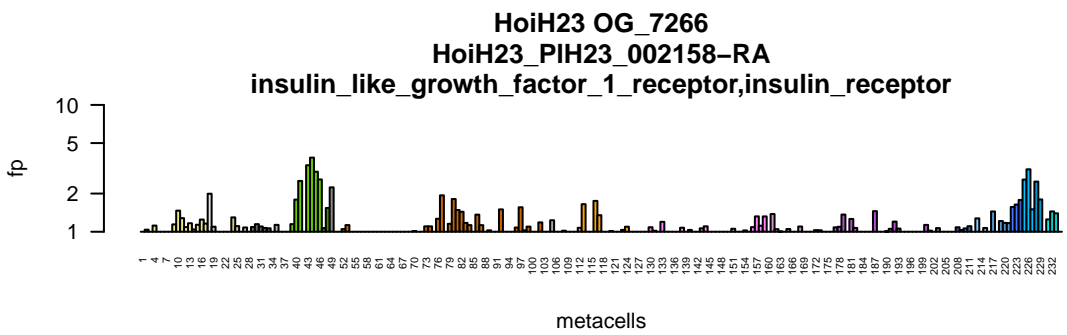
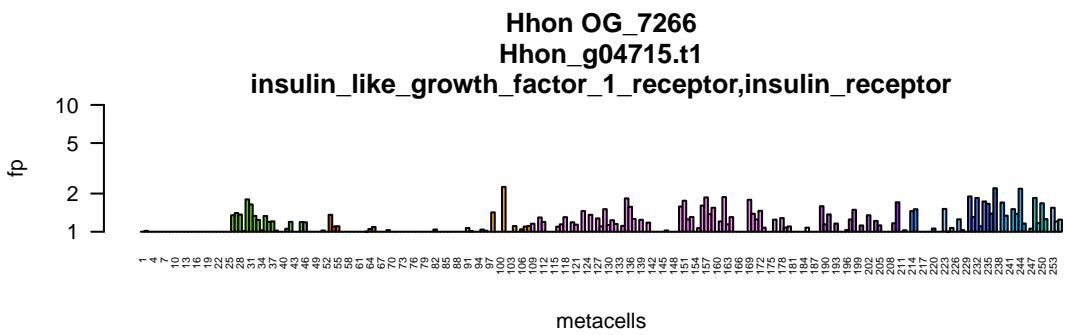
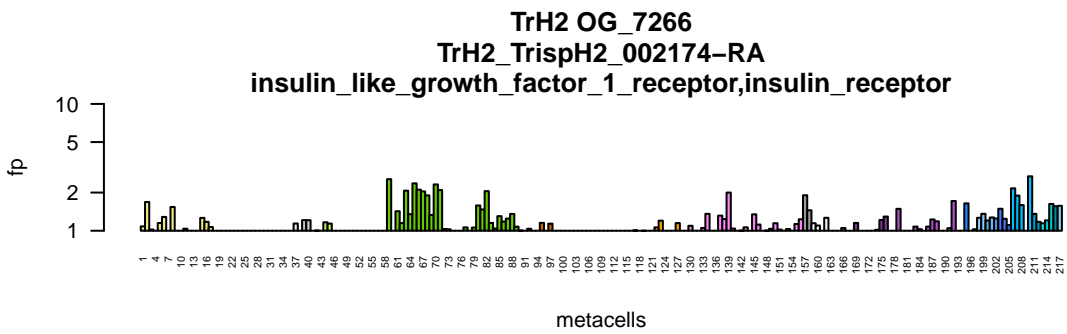
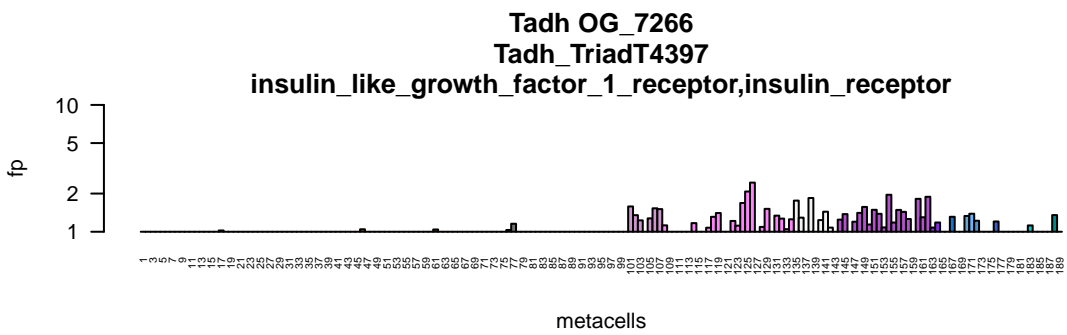
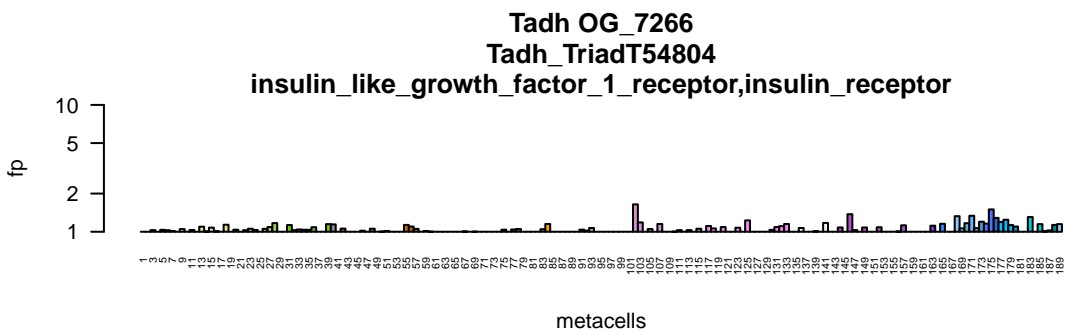
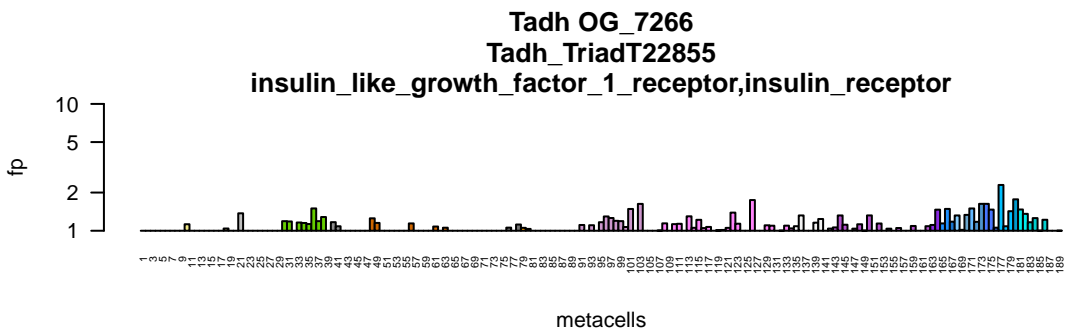
HoiH23 OG\_6800  
HoiH23\_PIH23\_001555-RA  
receptor\_1,oligodendrocyte\_myelin\_glycoprotein,leucine\_rich\_repeat\_containing\_40,relaxir



HoiH23 OG\_6800  
HoiH23\_PIH23\_001554-RA  
receptor\_1,oligodendrocyte\_myelin\_glycoprotein,leucine\_rich\_repeat\_containing\_40,relaxir



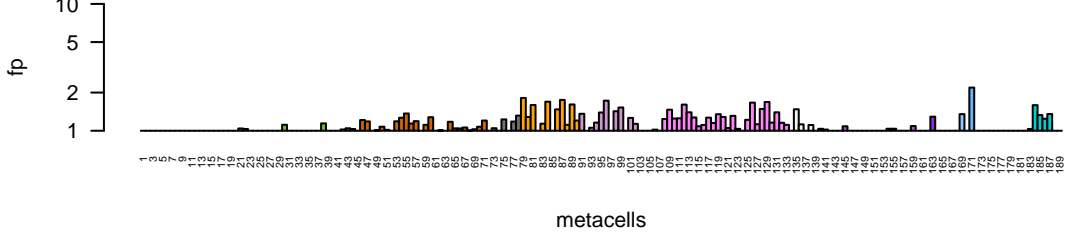




Tadh OG\_7344

Tadh\_TriadT57447

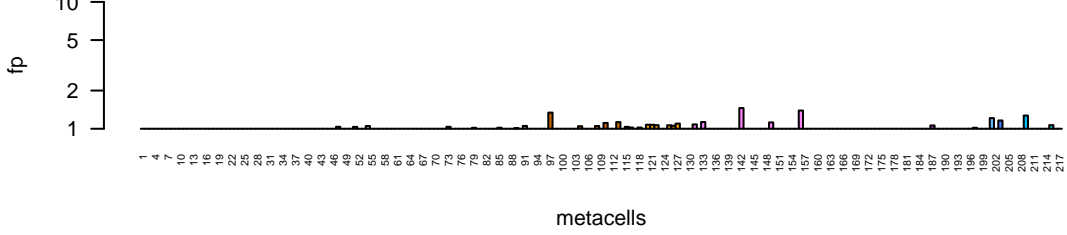
5\_hydroxytryptamine\_receptor\_1E,adrenoceptor\_alpha\_1A,angiotensin\_II\_receptor\_type



TrH2 OG\_7344

TrH2\_TrispH2\_009142-RA

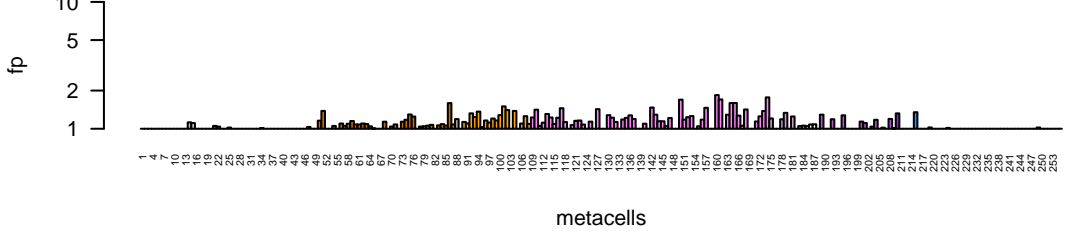
5\_hydroxytryptamine\_receptor\_1E,adrenoceptor\_alpha\_1A,angiotensin\_II\_receptor\_type



Hhon OG\_7344

Hhon\_g01918.t1

5\_hydroxytryptamine\_receptor\_1E,adrenoceptor\_alpha\_1A,angiotensin\_II\_receptor\_type



HoiH23 OG\_7344

HoiH23\_PIH23\_001482-RA

5\_hydroxytryptamine\_receptor\_1E,adrenoceptor\_alpha\_1A,angiotensin\_II\_receptor\_type

