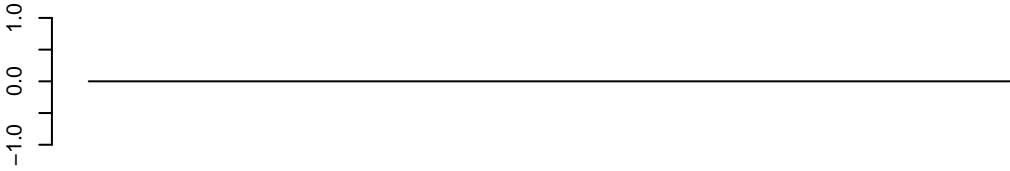
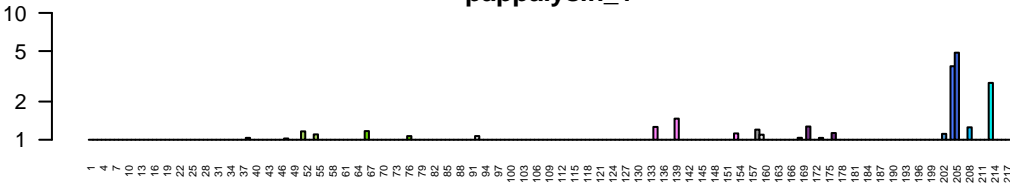


pappalysin\_1  
Tadh | no data

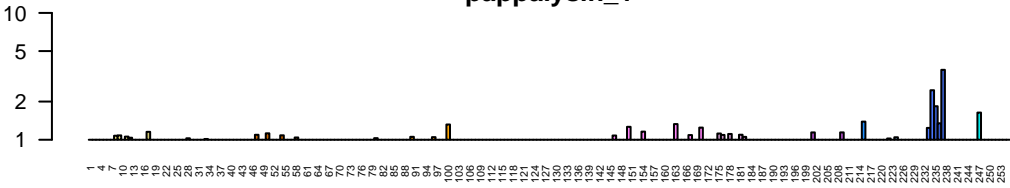


TrH2 OG\_437  
TrH2\_TrispH2\_002674-RA  
pappalysin\_1



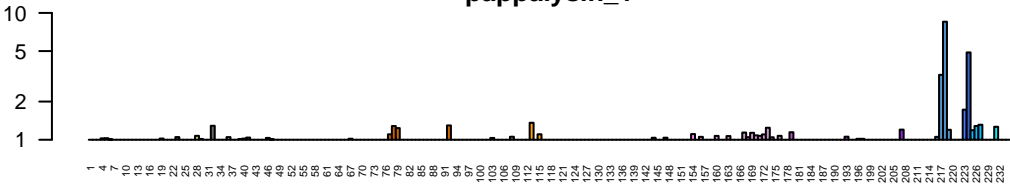
metacells

Hhon OG\_437  
Hhon\_g01583.t1  
pappalysin\_1

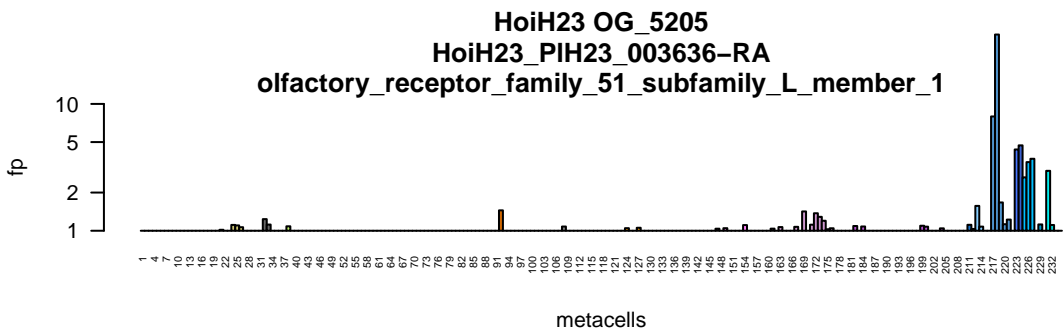
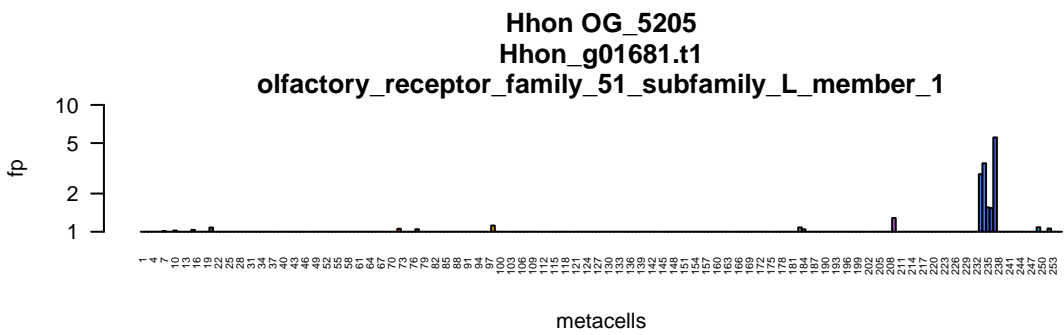
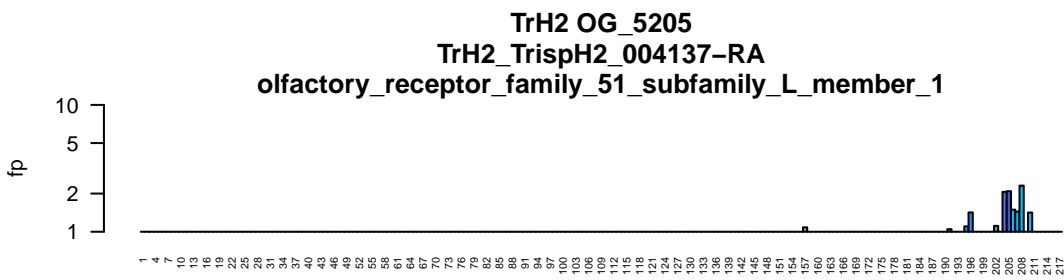
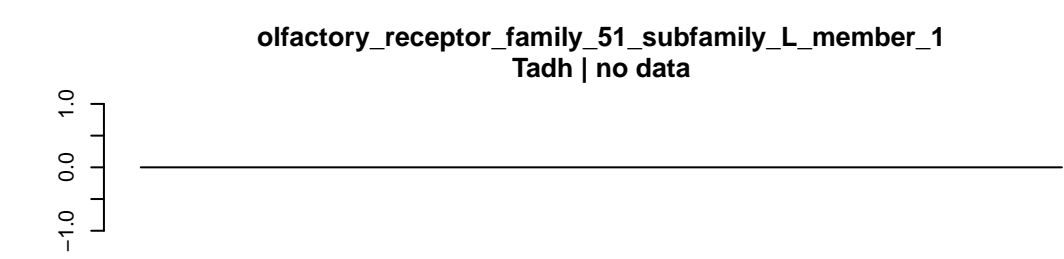


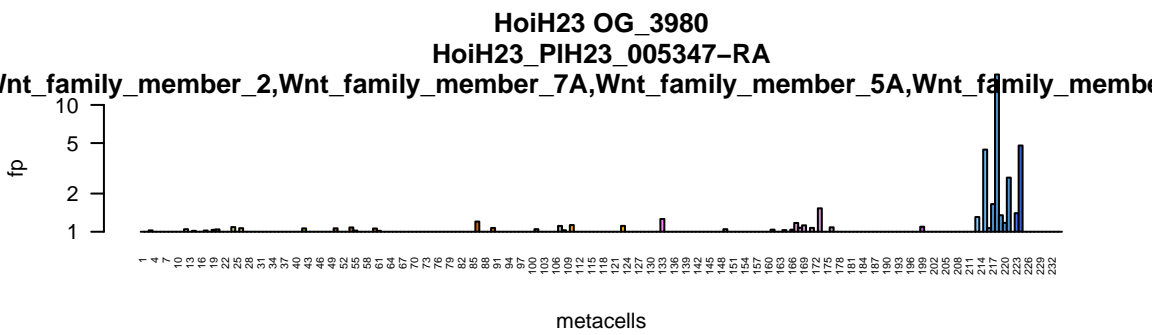
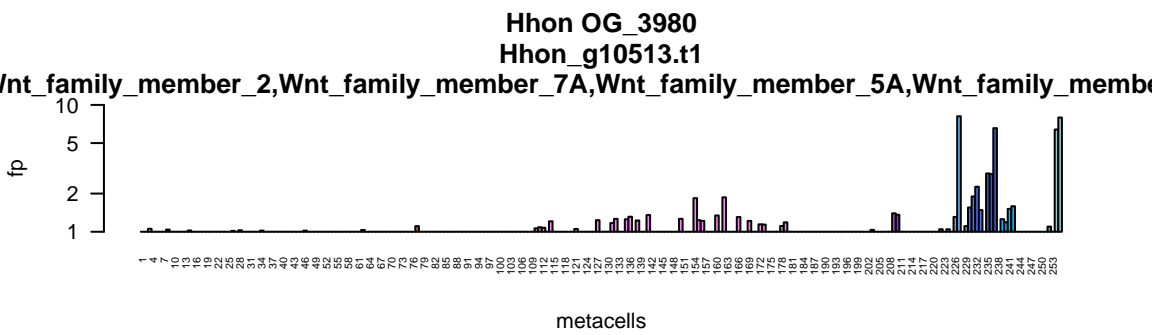
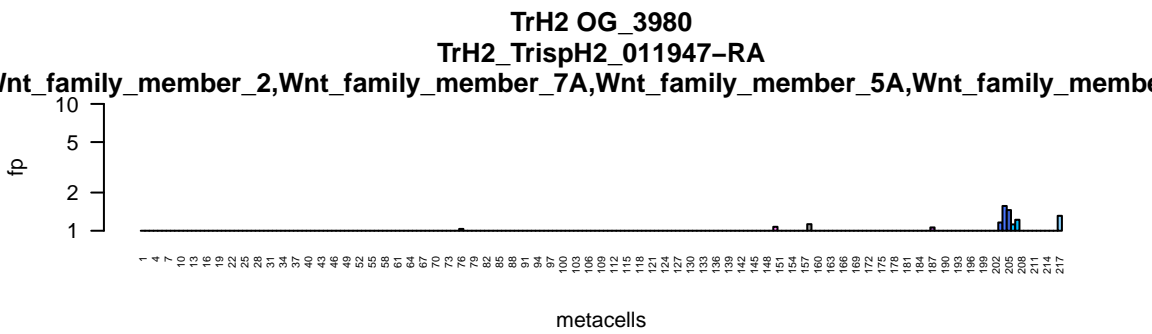
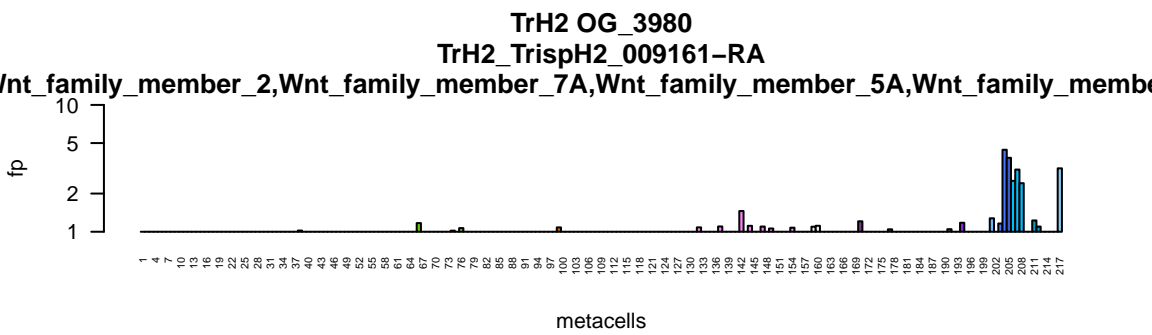
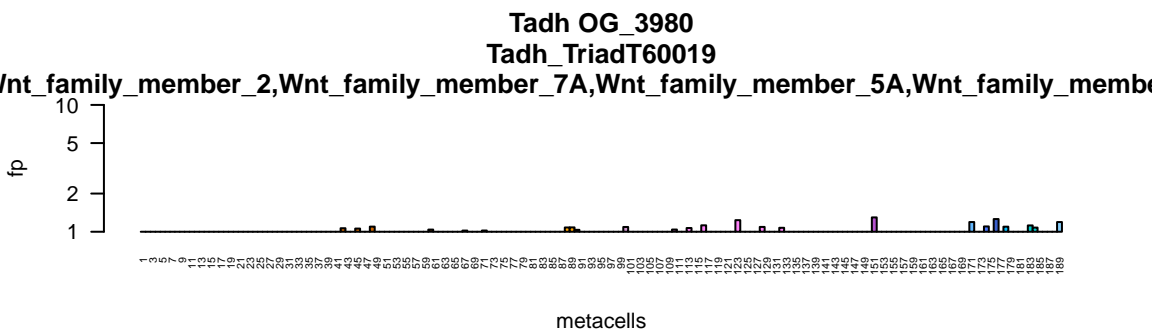
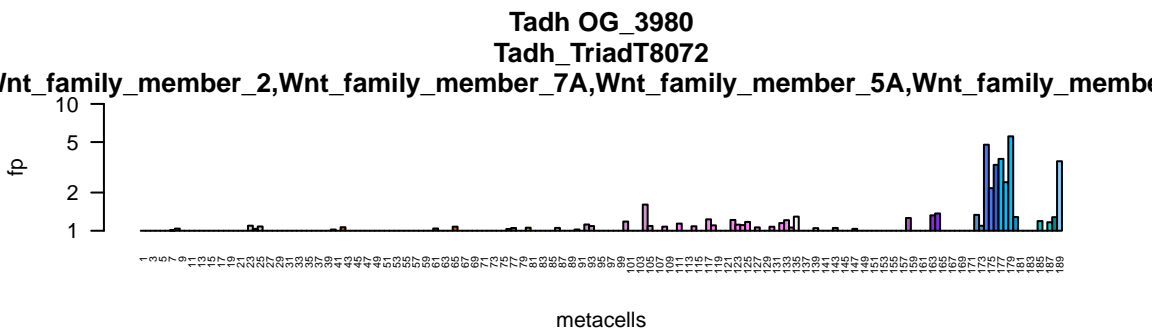
metacells

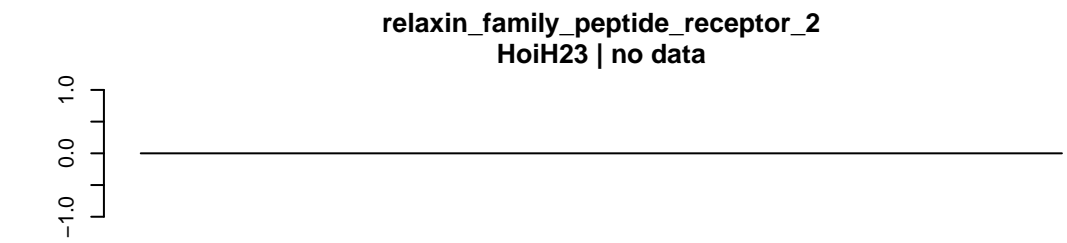
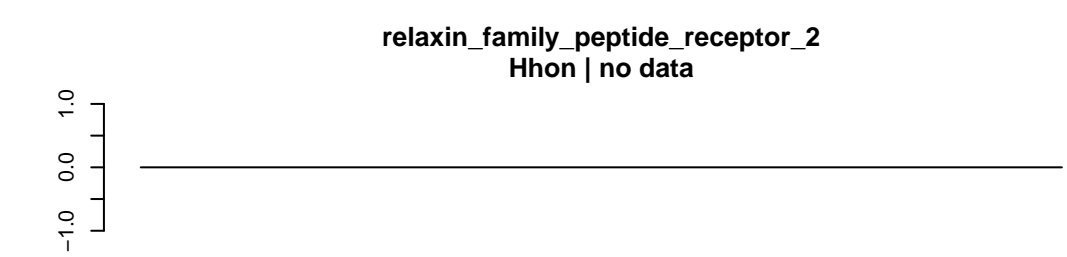
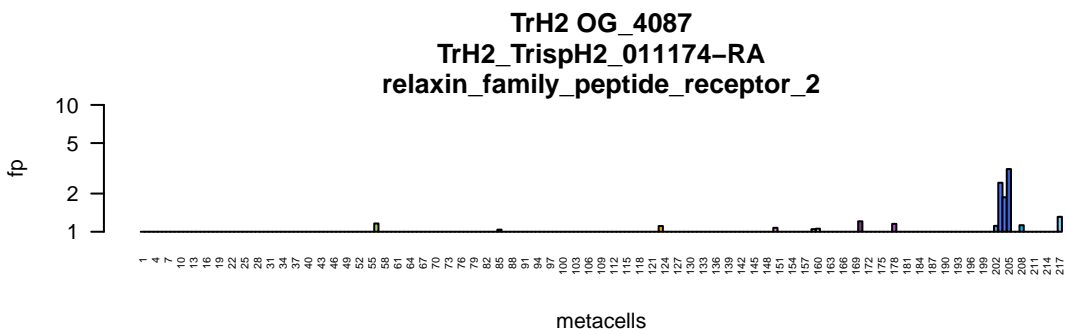
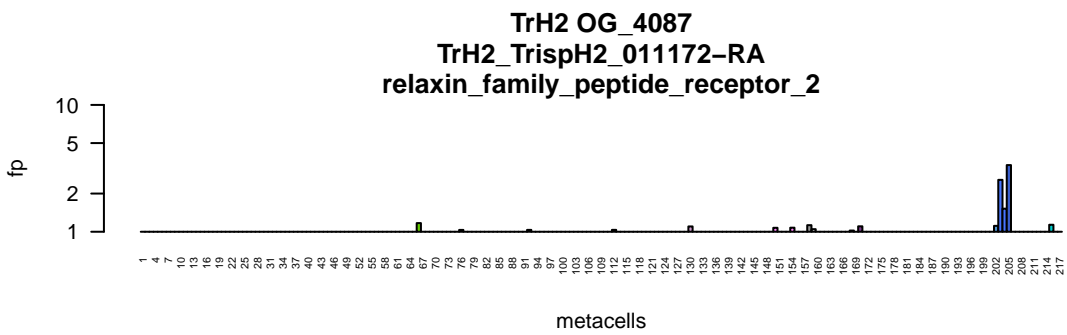
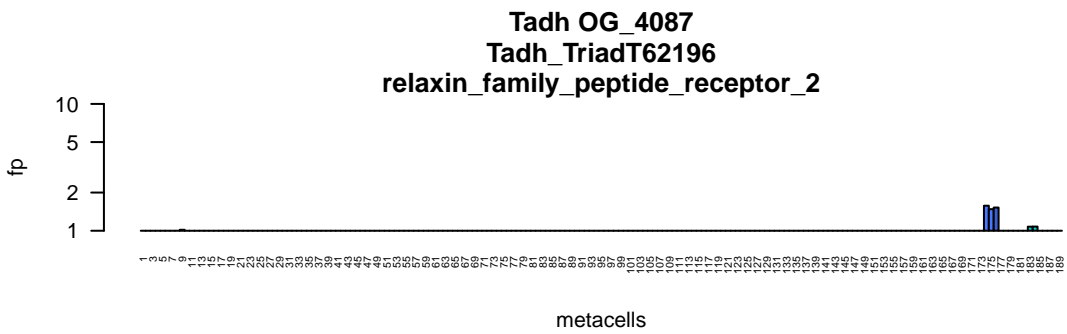
HoiH23 OG\_437  
HoiH23\_PIH23\_002511-RA  
pappalysin\_1

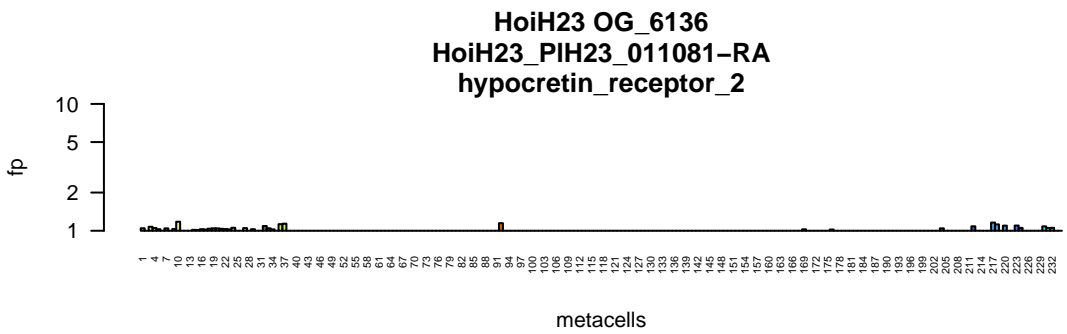
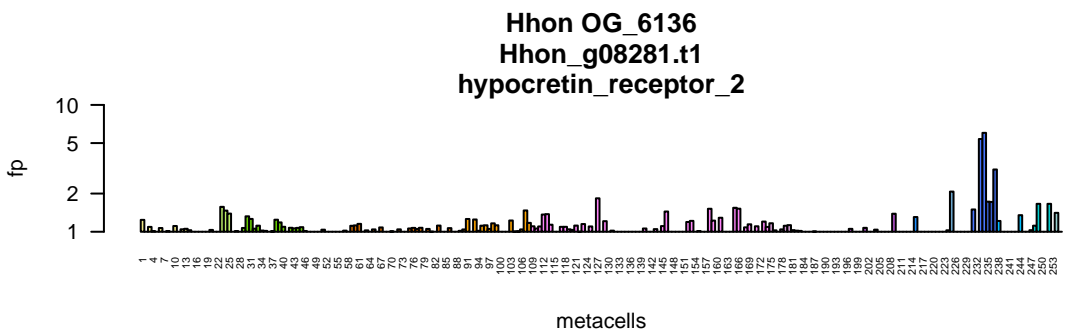
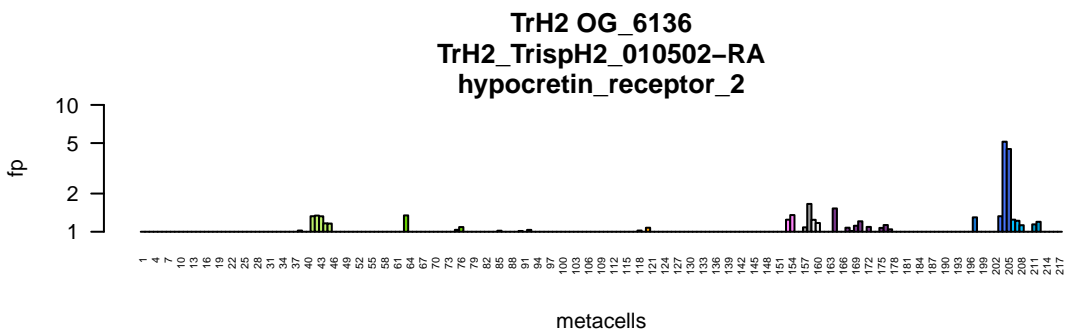
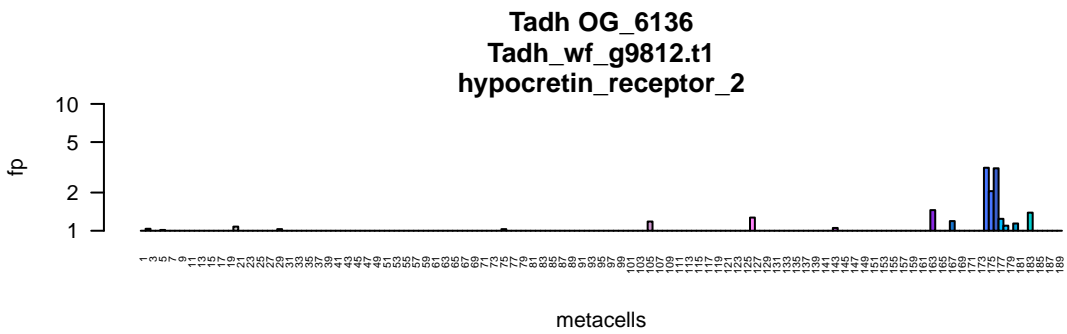


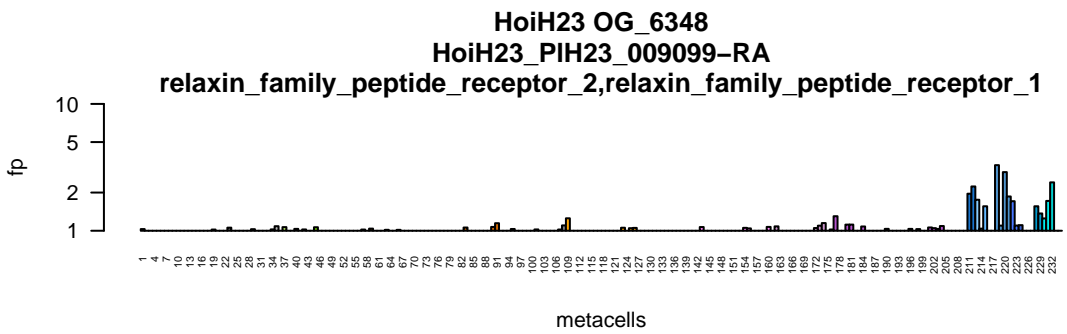
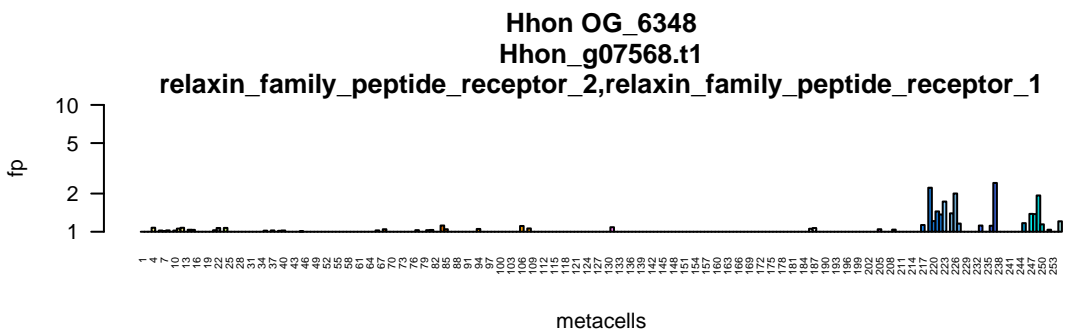
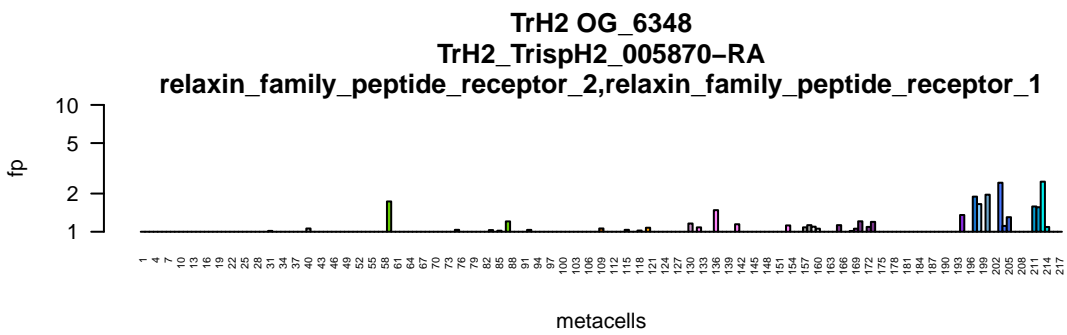
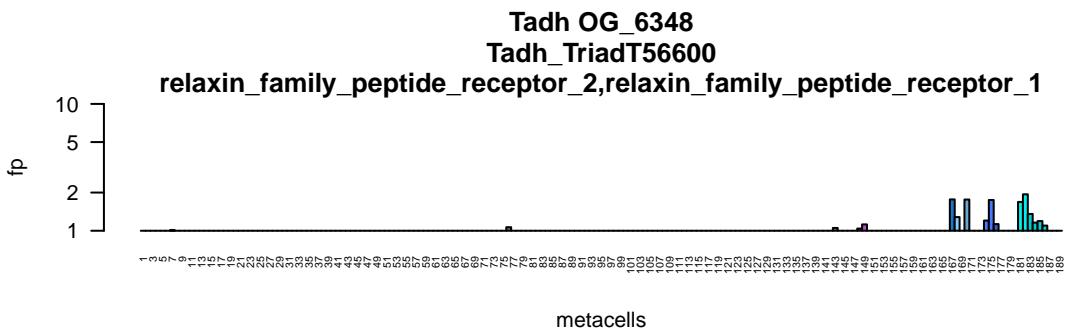
metacells



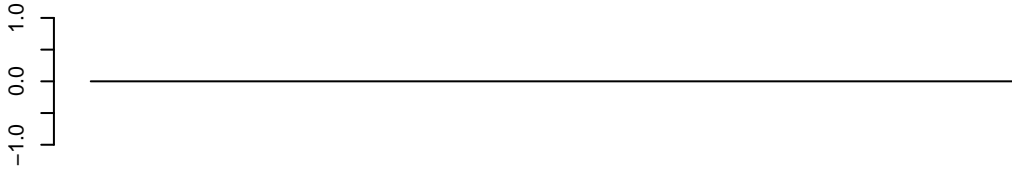




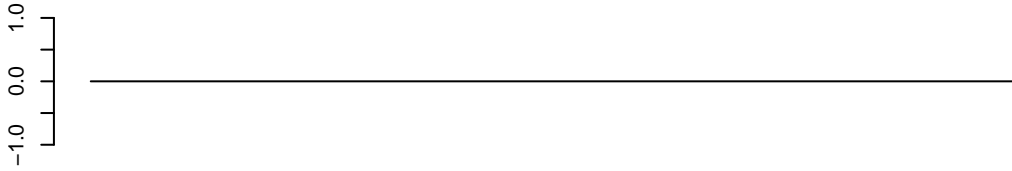




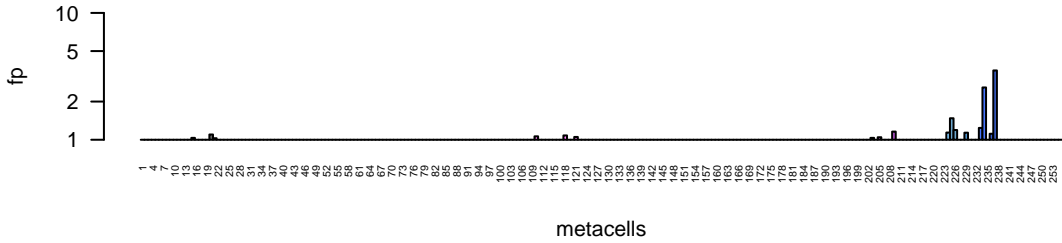
Tadh | no data



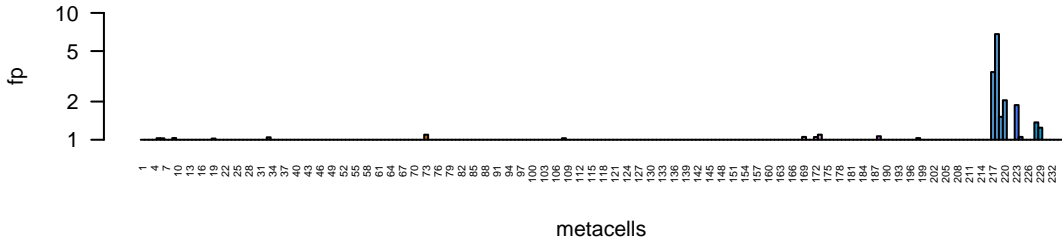
TrH2 | no data



Hhon OG\_6709  
Hhon\_g01989.t1



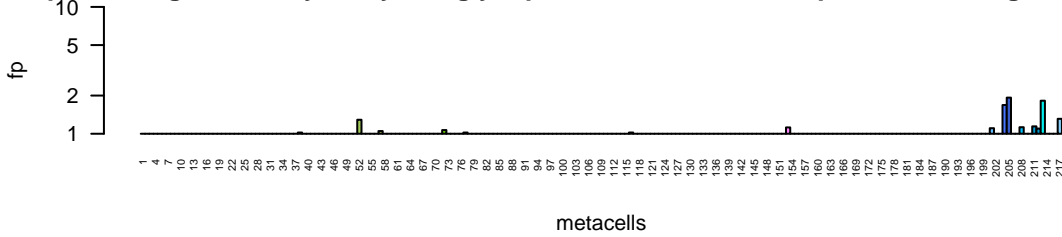
HoiH23 OG\_6709  
HoiH23\_PIH23\_003664-RA



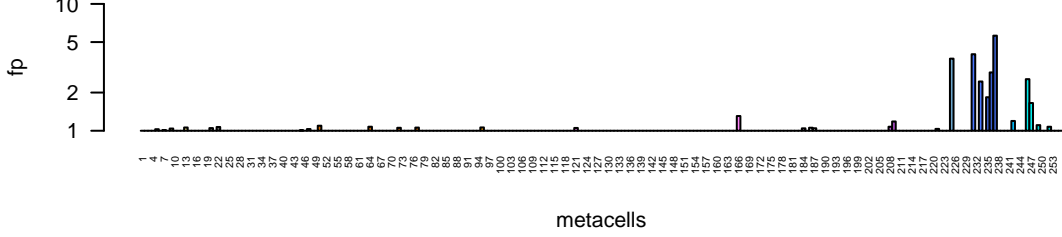
receptor\_1,oligodendrocyte\_myelin\_glycoprotein,leucine\_rich\_repeat\_containing\_40,relaxir  
Tadh | no data



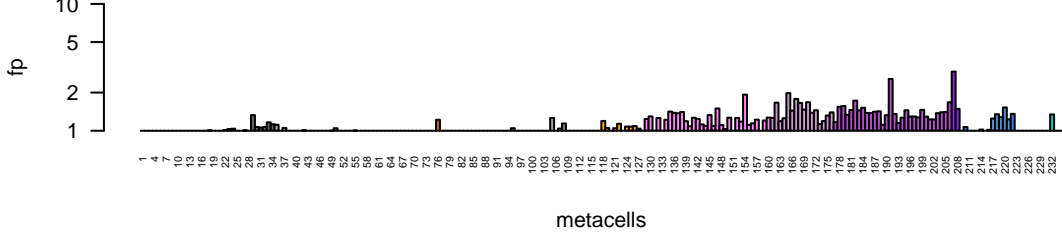
TrH2 OG\_6800  
TrH2\_TrispH2\_006274-RA  
receptor\_1,oligodendrocyte\_myelin\_glycoprotein,leucine\_rich\_repeat\_containing\_40,relaxir



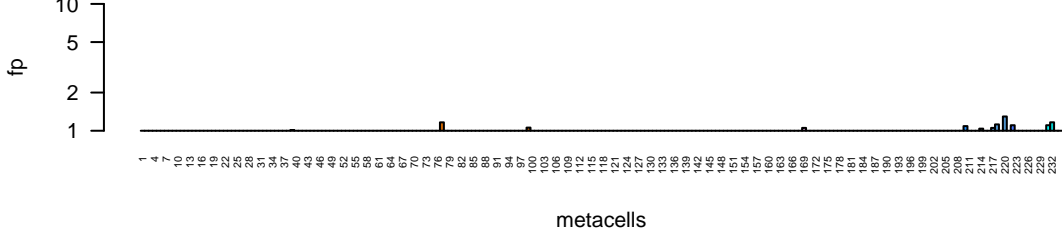
Hhon OG\_6800  
Hhon\_g04761.t1  
receptor\_1,oligodendrocyte\_myelin\_glycoprotein,leucine\_rich\_repeat\_containing\_40,relaxir



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

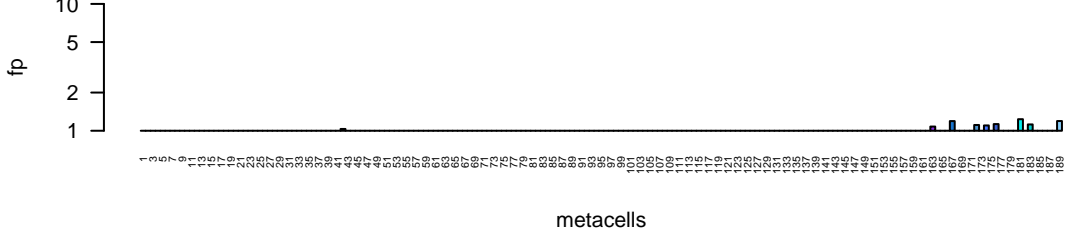




Tadh OG\_7054

Tadh\_TriadT59349

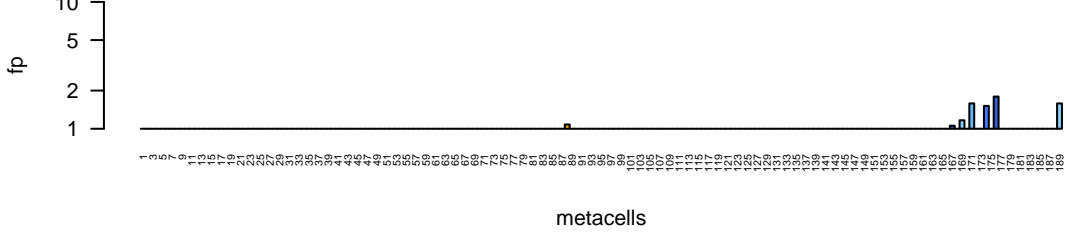
adhesion\_G\_protein\_coupled\_receptor\_D1,adhesion\_G\_protein\_coupled\_receptor\_L3



Tadh OG\_7054

Tadh\_TriadT59345

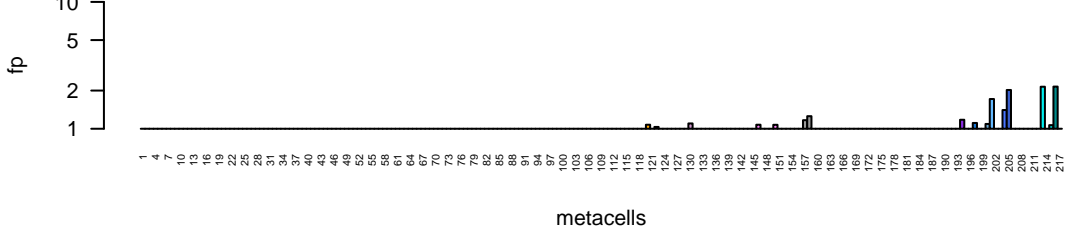
adhesion\_G\_protein\_coupled\_receptor\_D1,adhesion\_G\_protein\_coupled\_receptor\_L3



TrH2 OG\_7054

TrH2\_TrispH2\_011841-RA

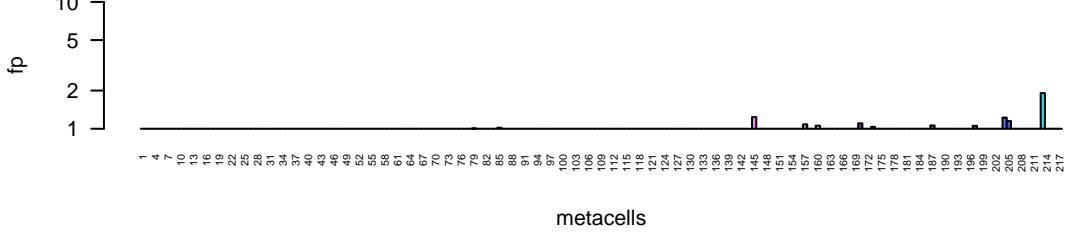
adhesion\_G\_protein\_coupled\_receptor\_D1,adhesion\_G\_protein\_coupled\_receptor\_L3



TrH2 OG\_7054

TrH2\_TrispH2\_010220-RA

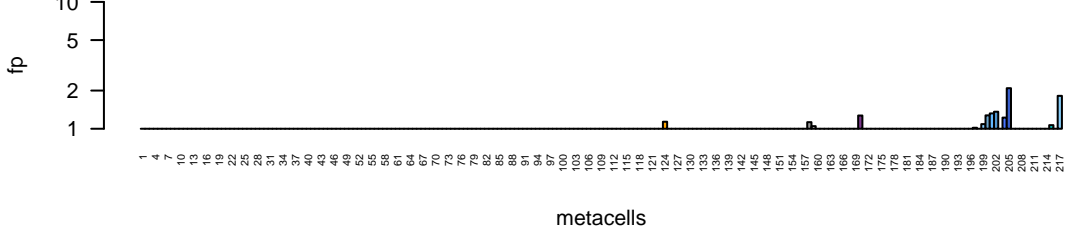
adhesion\_G\_protein\_coupled\_receptor\_D1,adhesion\_G\_protein\_coupled\_receptor\_L3



TrH2 OG\_7054

TrH2\_TrispH2\_010222-RA

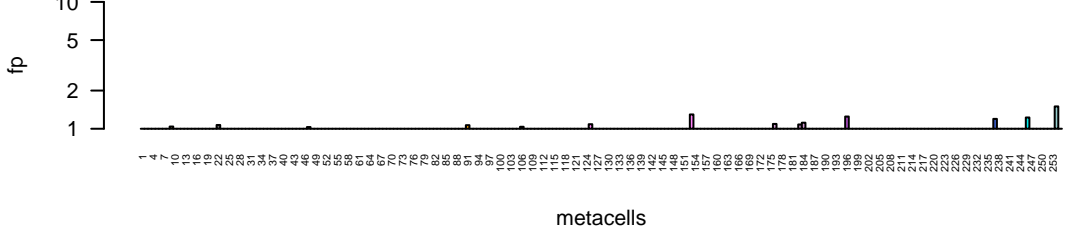
adhesion\_G\_protein\_coupled\_receptor\_D1,adhesion\_G\_protein\_coupled\_receptor\_L3



Hhon OG\_7054

Hhon\_g10892.t1

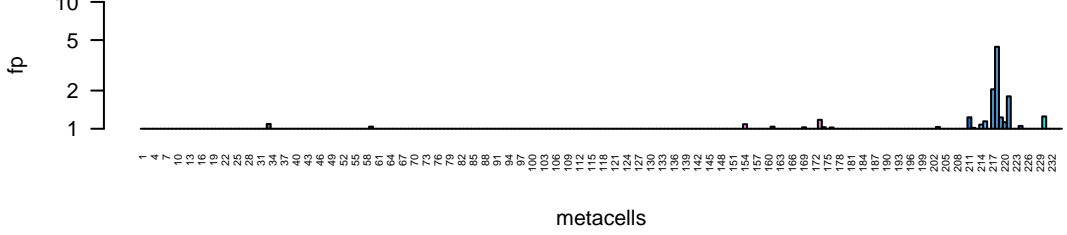
adhesion\_G\_protein\_coupled\_receptor\_D1,adhesion\_G\_protein\_coupled\_receptor\_L3



HoiH23 OG\_7054

HoiH23\_PIH23\_010401-RA

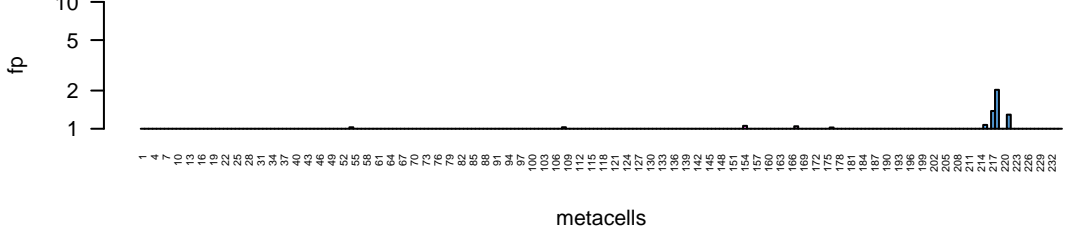
adhesion\_G\_protein\_coupled\_receptor\_D1,adhesion\_G\_protein\_coupled\_receptor\_L3

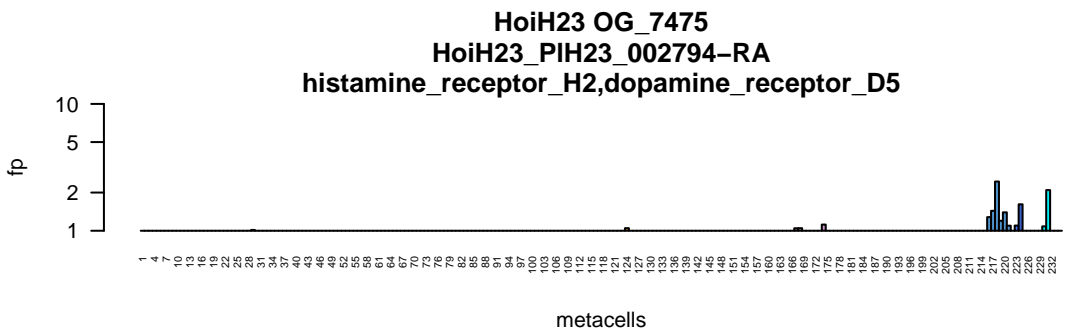
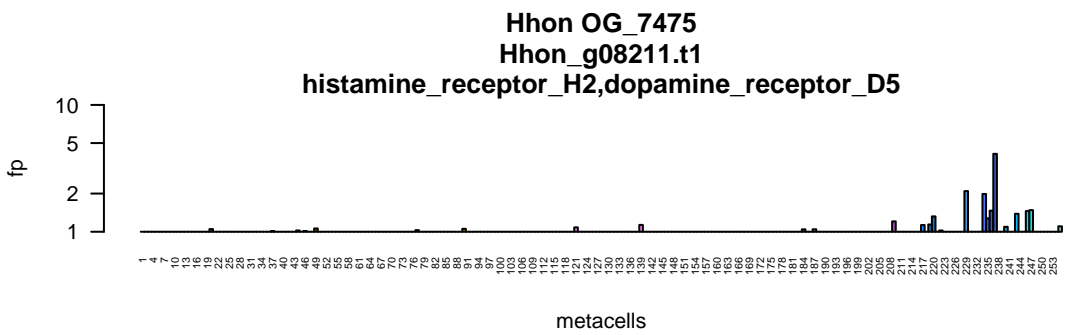
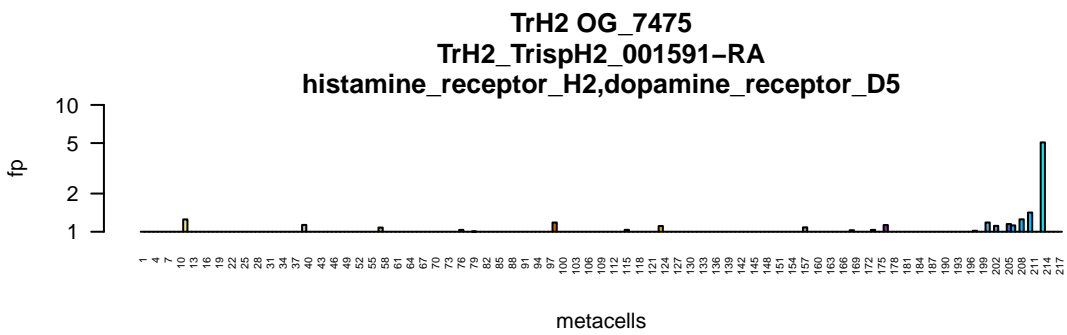


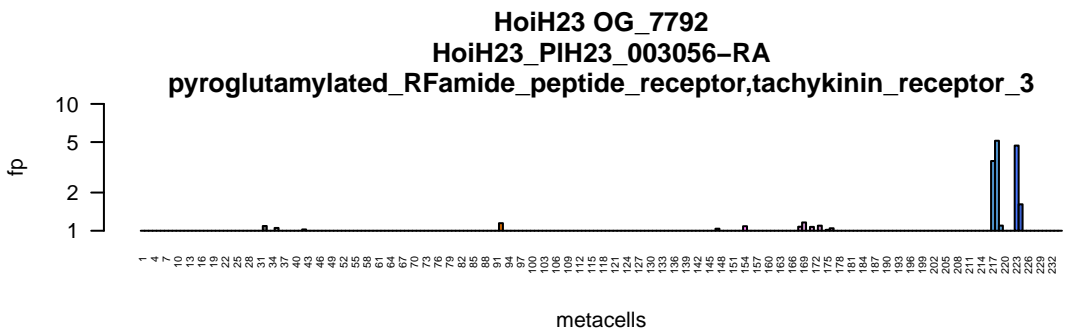
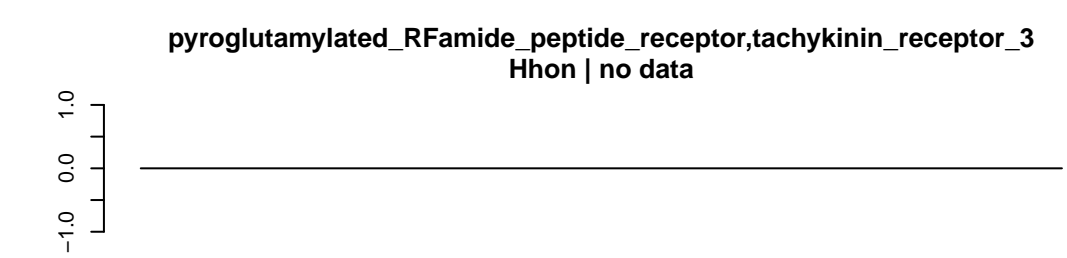
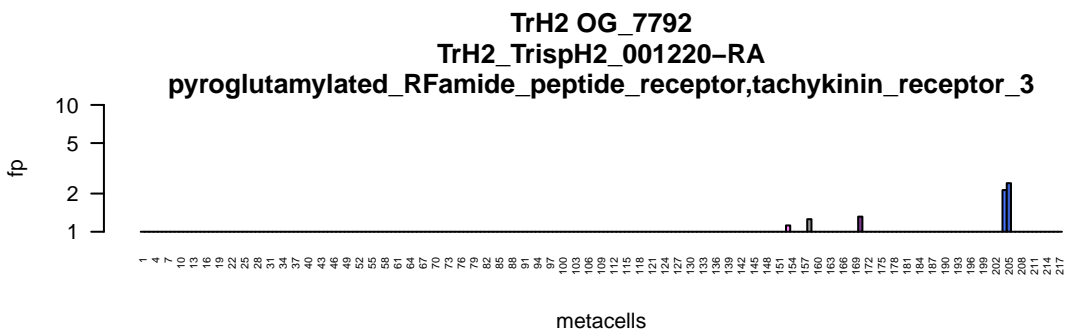
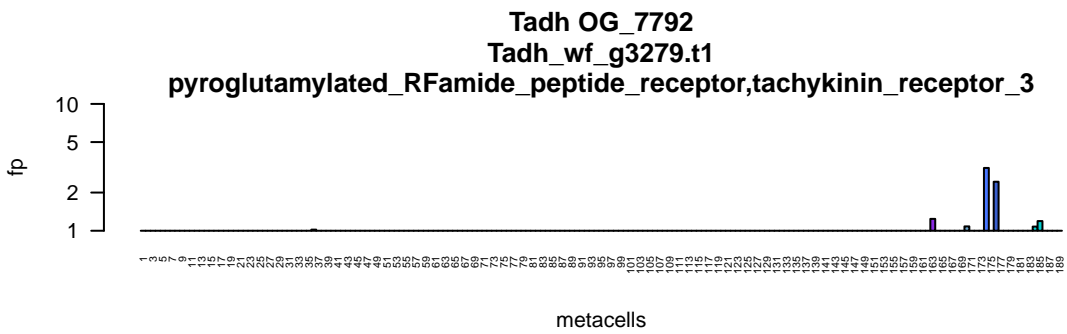
HoiH23 OG\_7054

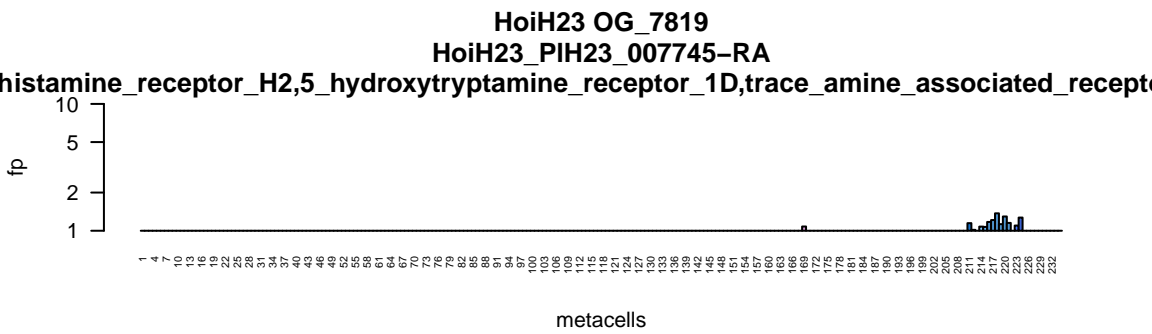
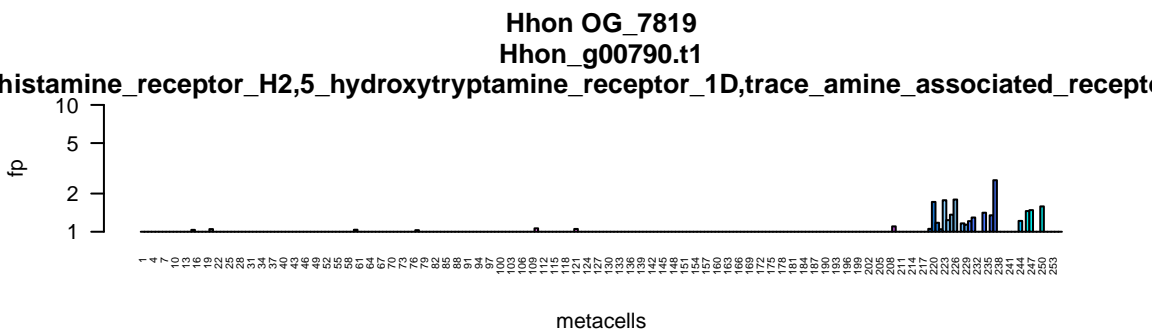
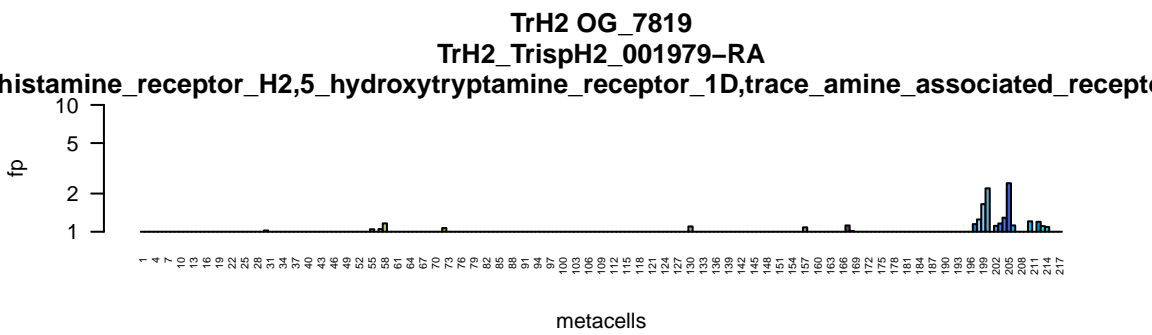
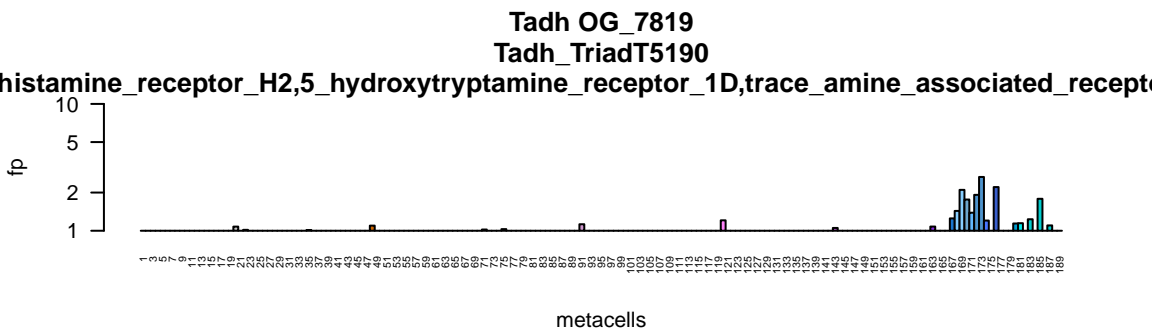
HoiH23\_PIH23\_006966-RA

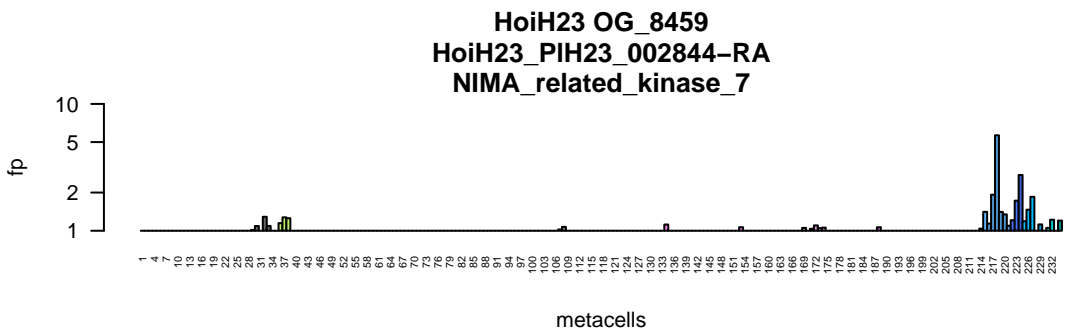
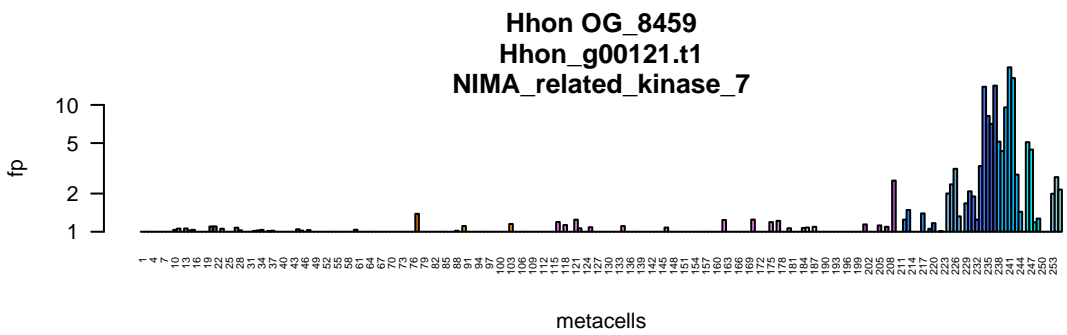
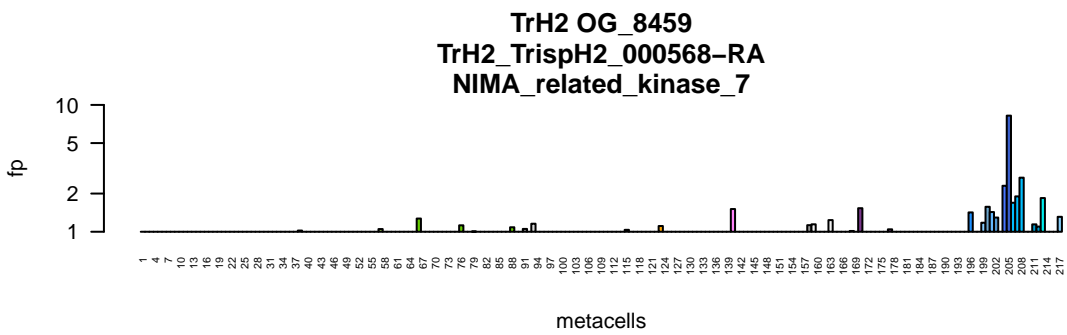
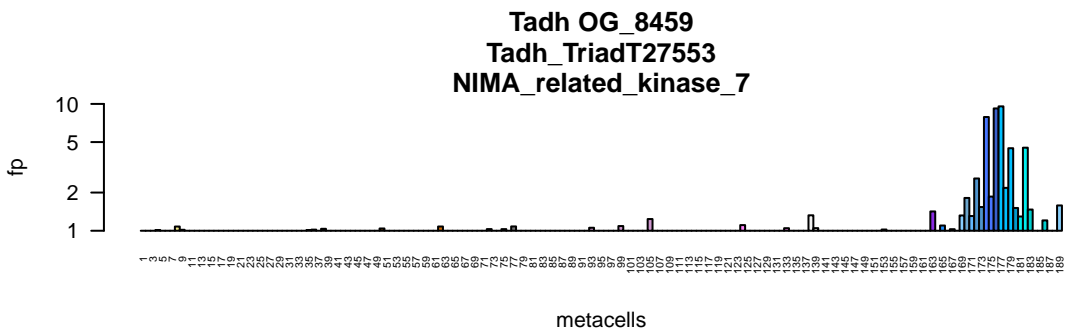
adhesion\_G\_protein\_coupled\_receptor\_D1,adhesion\_G\_protein\_coupled\_receptor\_L3





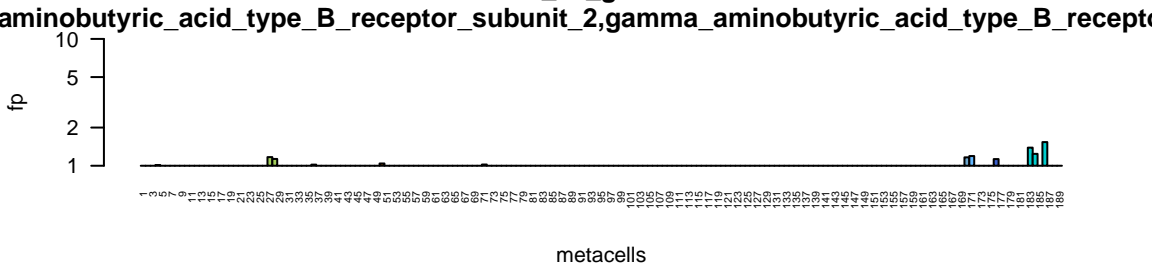






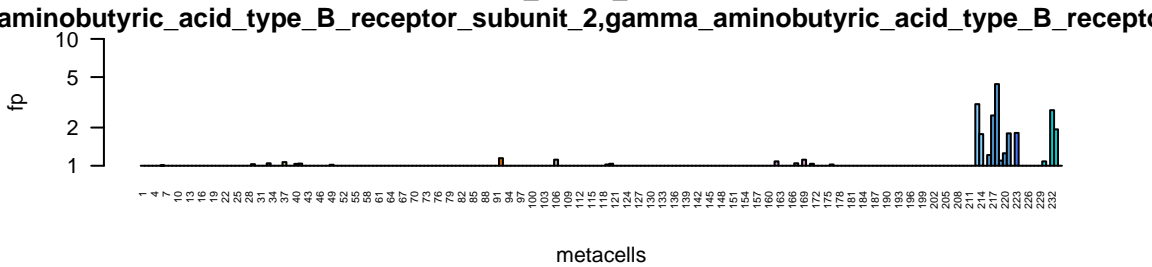
**Tadh OG\_8960**

**Tadh\_wf\_g11896.t1**



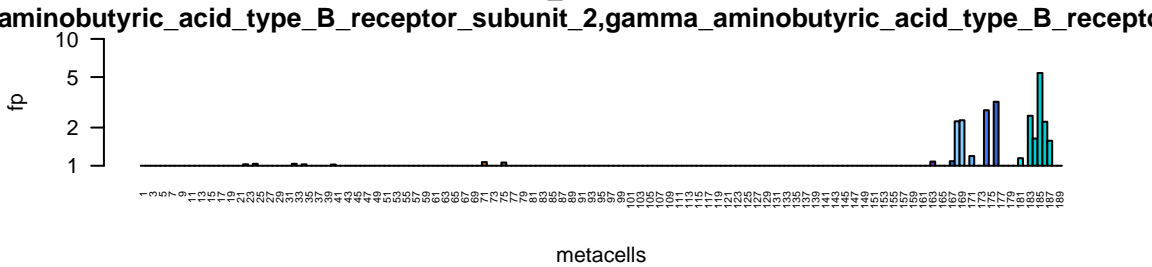
**HoiH23 OG\_8960**

**HoiH23\_PIH23\_000719-RA**



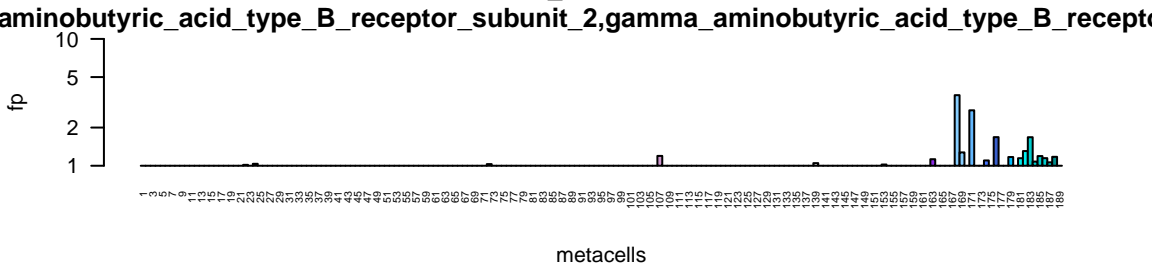
**Tadh OG\_8960**

**Tadh\_TriadT52578**



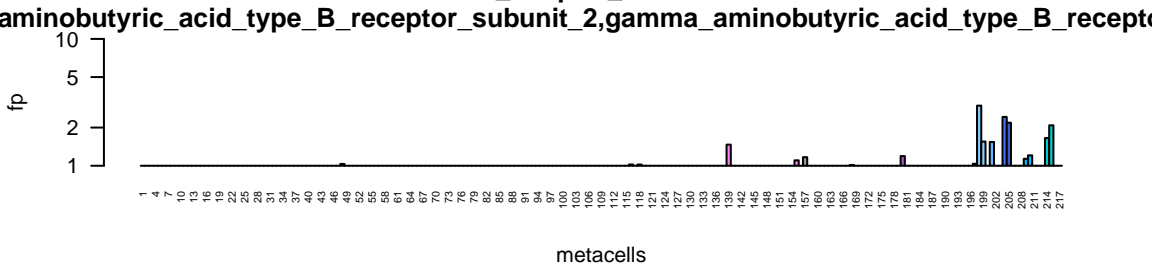
**Tadh OG\_8960**

**Tadh\_TriadT52579**



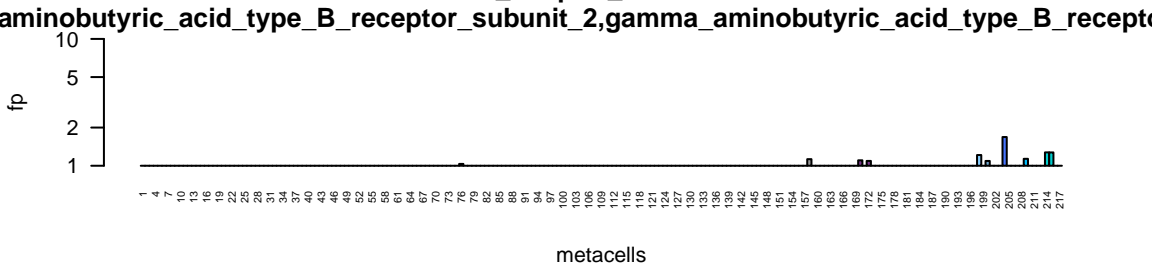
**TrH2 OG\_8960**

**TrH2\_TrispH2\_012100-RA**



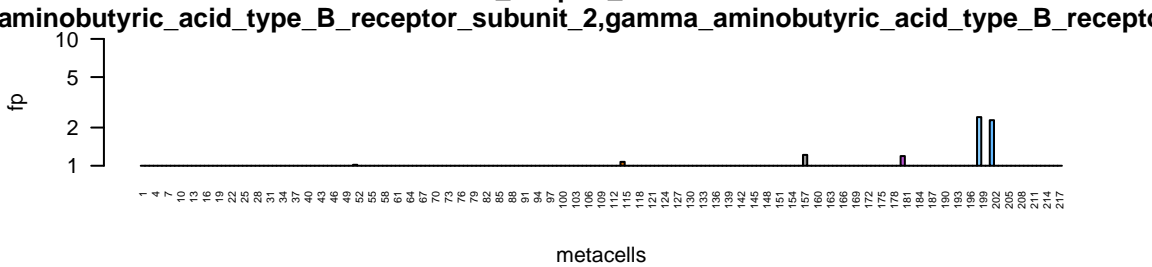
**TrH2 OG\_8960**

**TrH2\_TrispH2\_000234-RA**



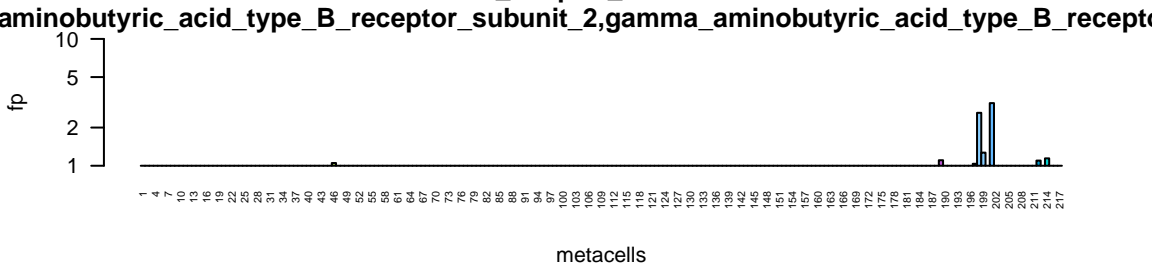
**TrH2 OG\_8960**

**TrH2\_TrispH2\_011964-RA**



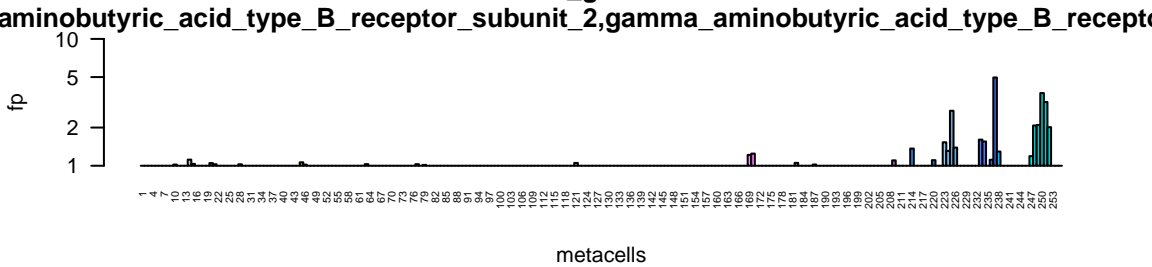
**TrH2 OG\_8960**

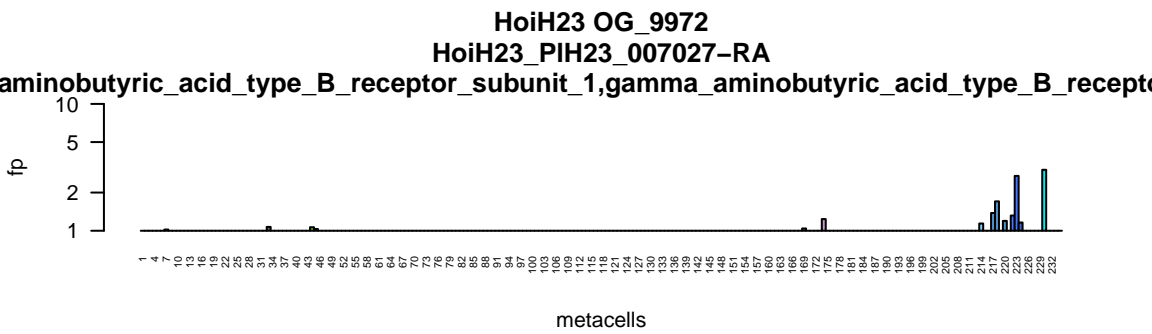
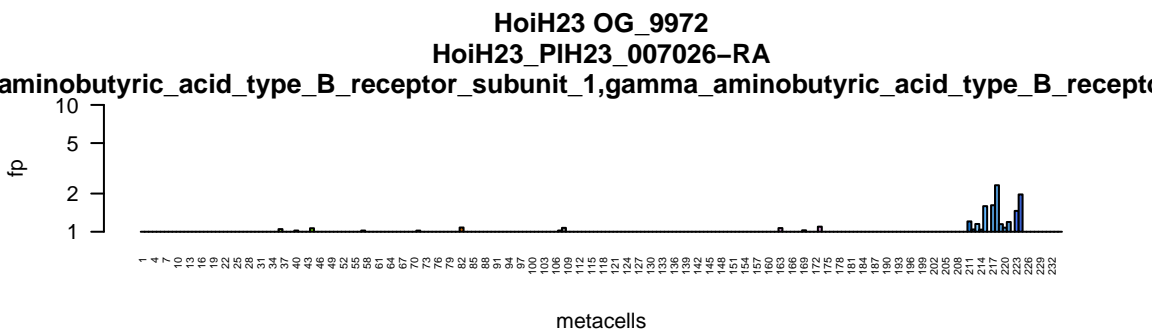
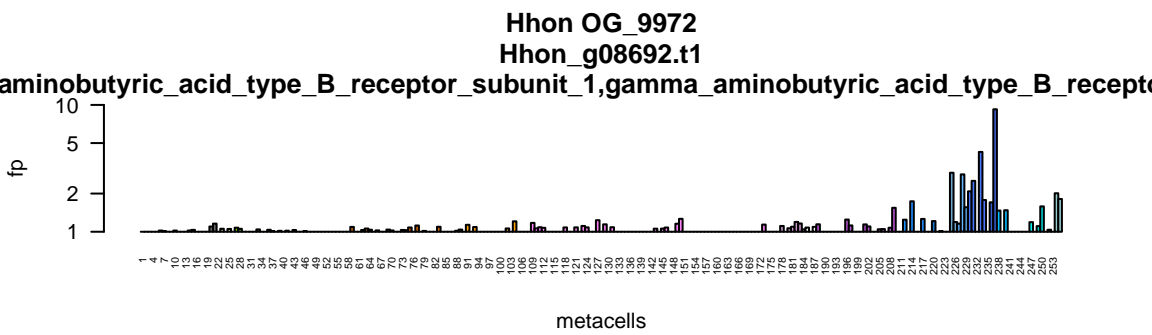
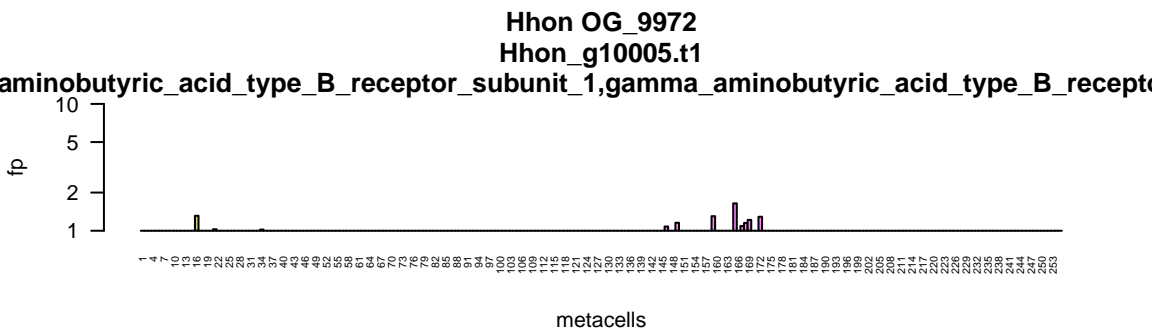
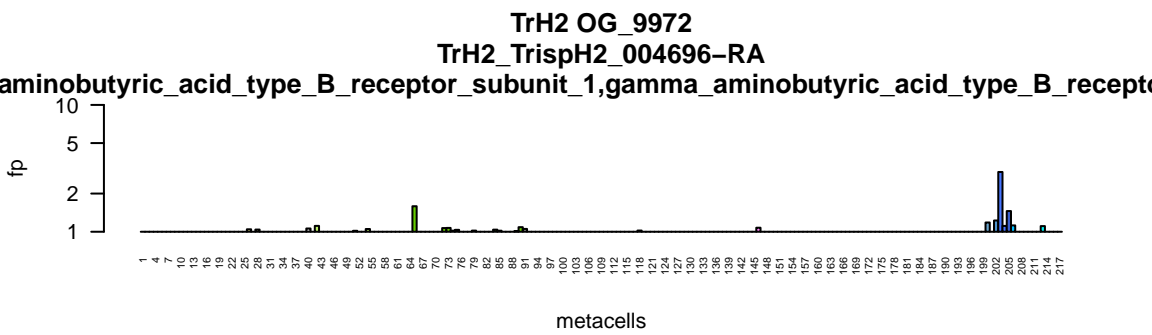
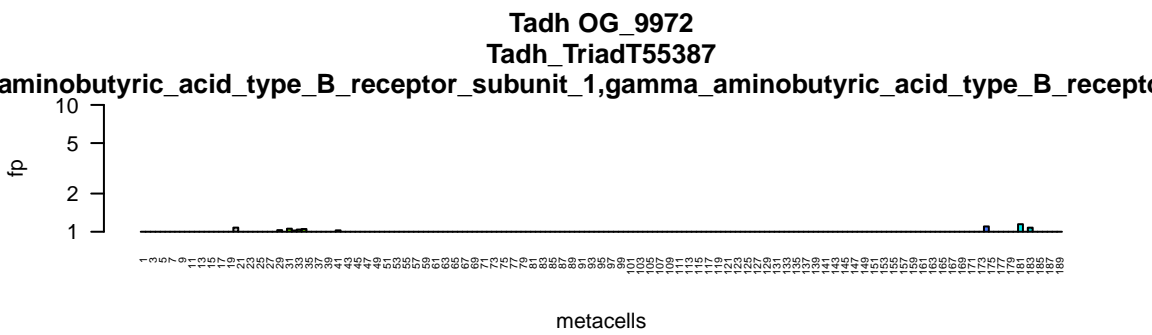
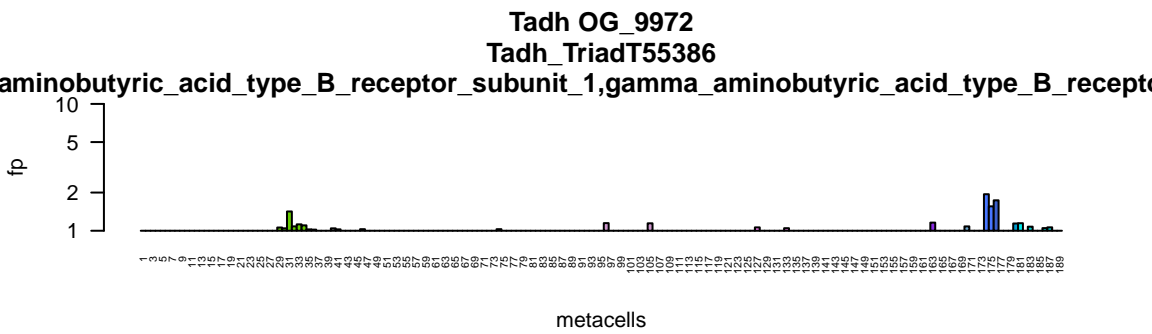
TrH2\_TrispH2\_011847-RA

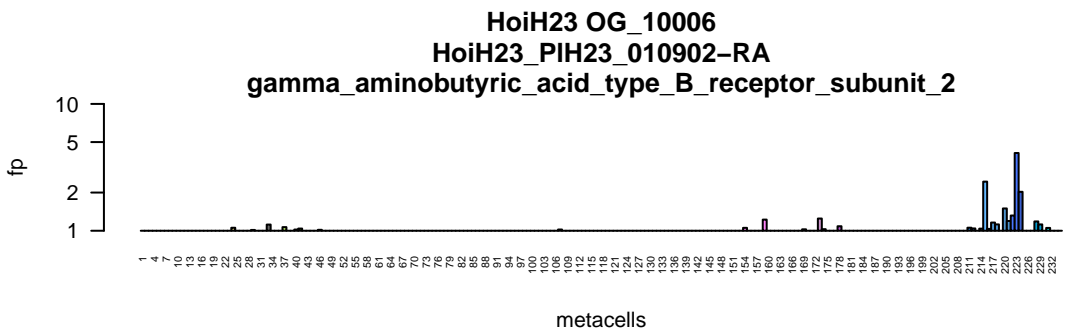
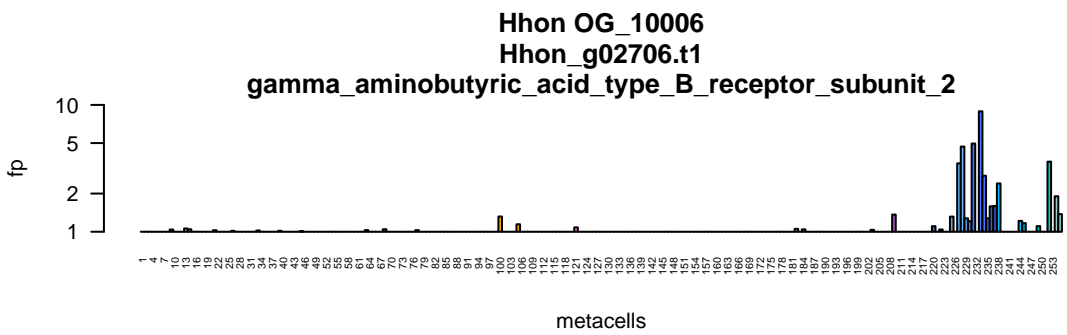
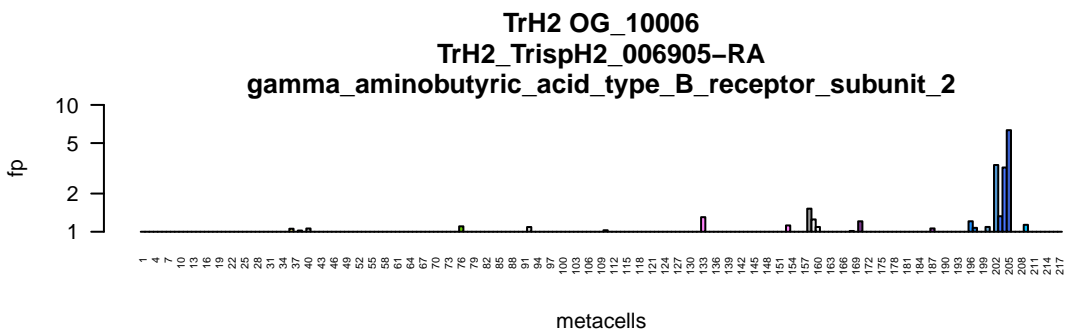
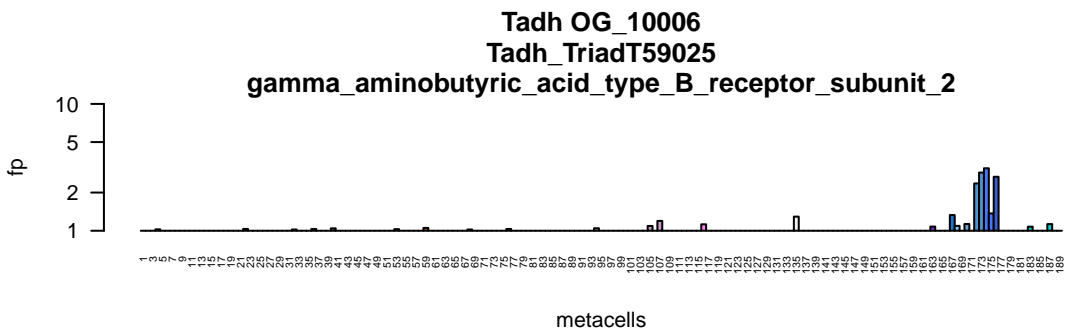


**Hhon OG\_8960**

Hhon\_g05296.t1

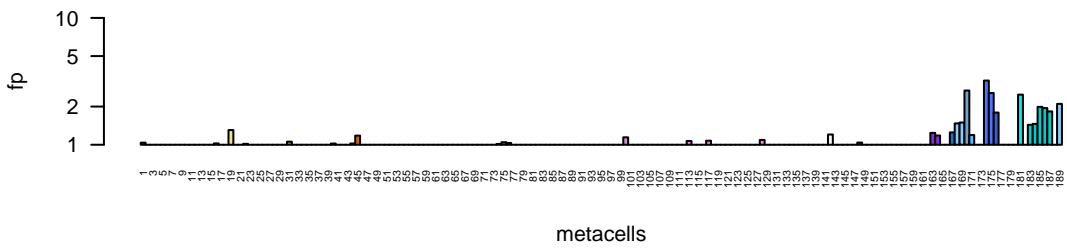




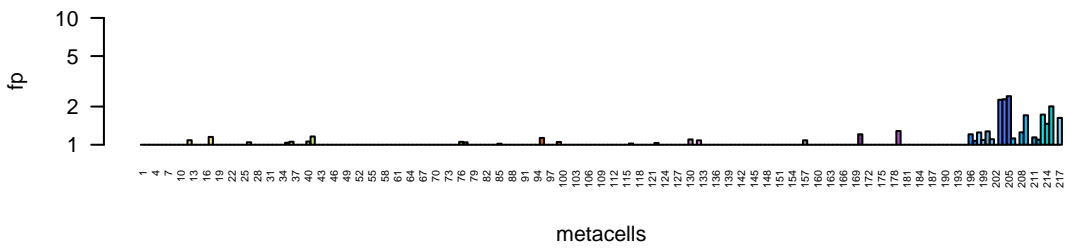




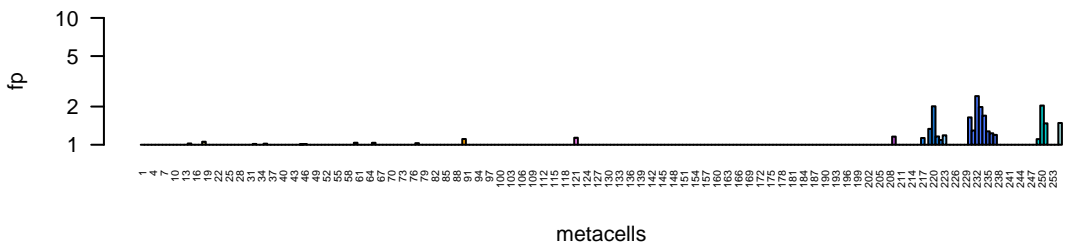
Tadh OG\_1238  
Tadh\_wf\_g1259.t1



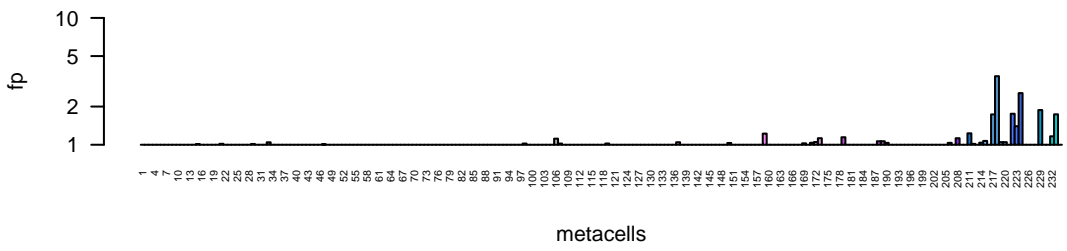
TrH2 OG\_1238  
TrH2\_TrispH2\_000180-RA

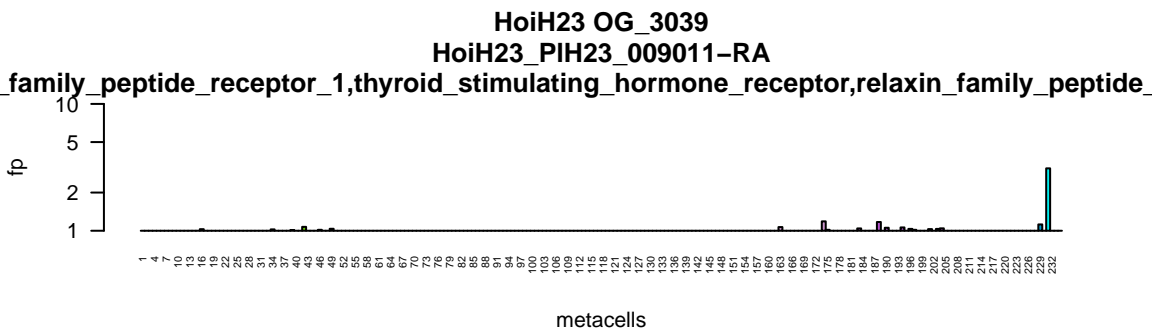
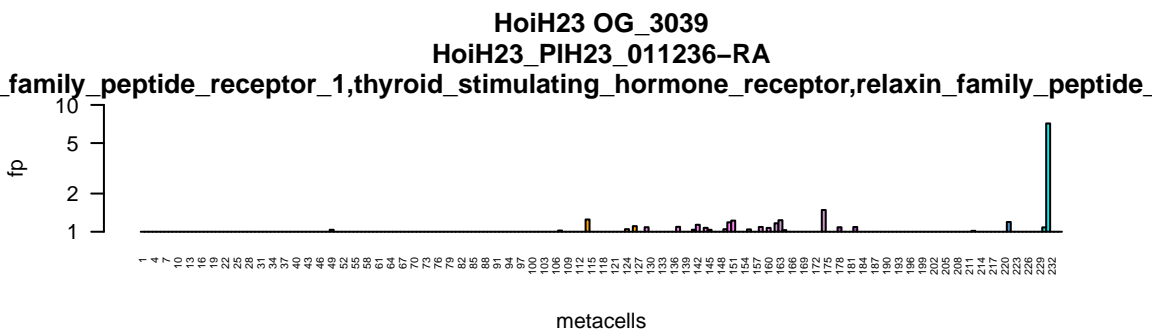
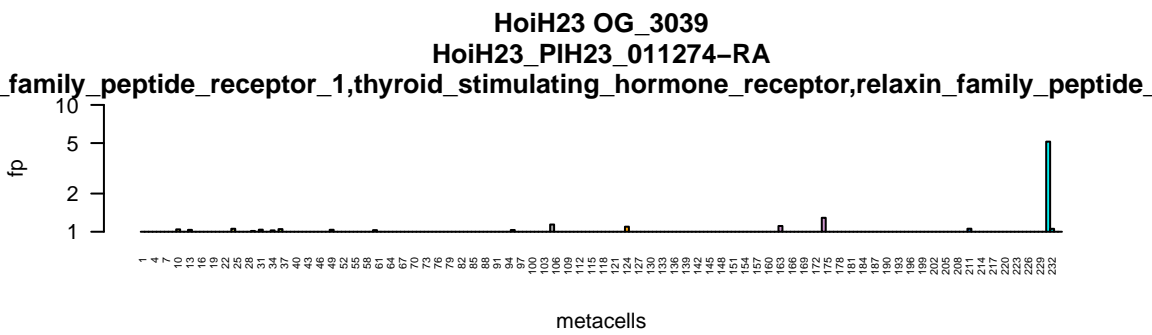
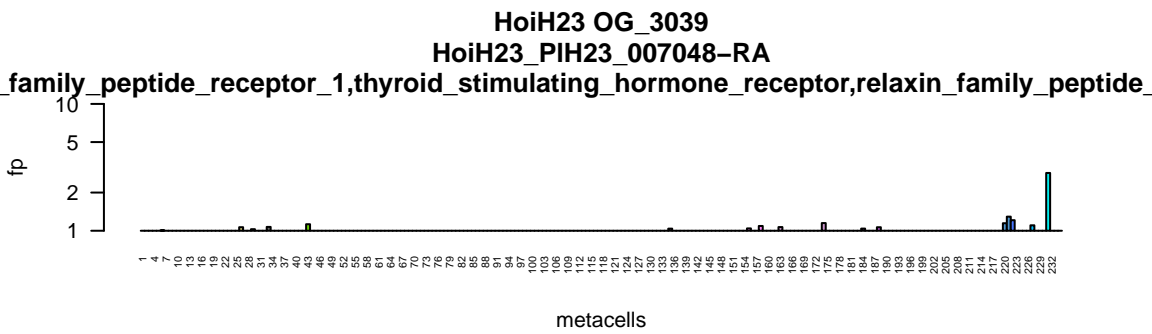
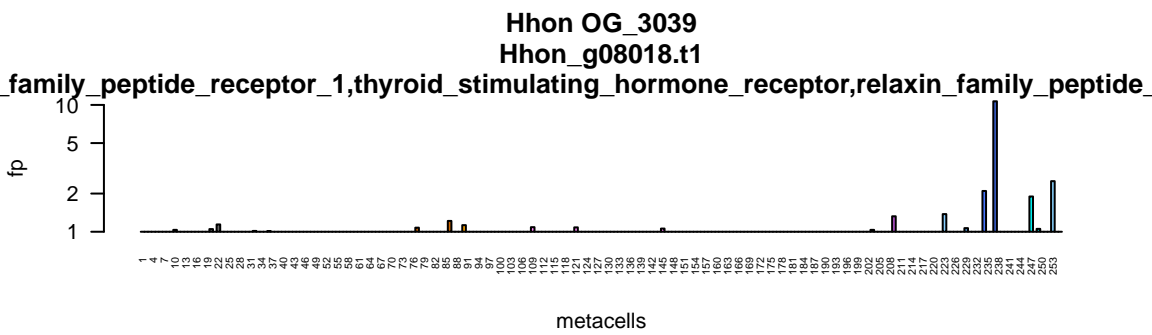
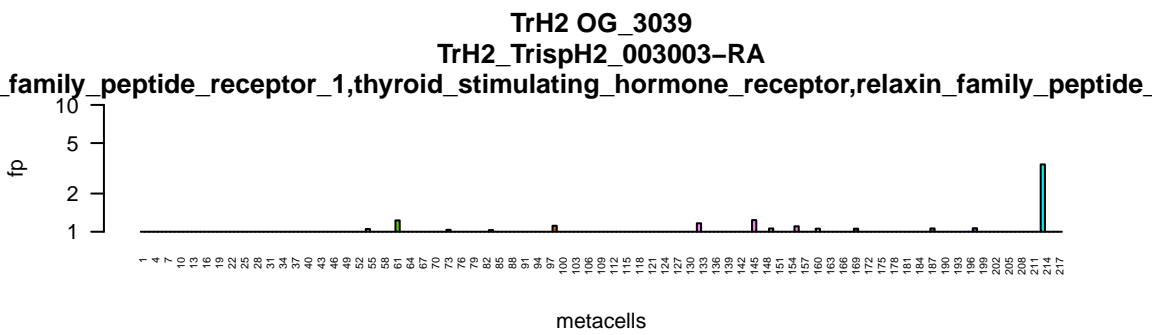
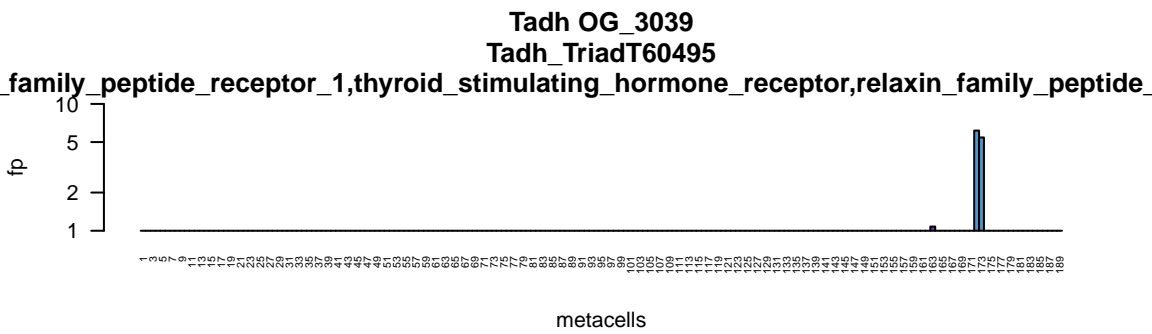


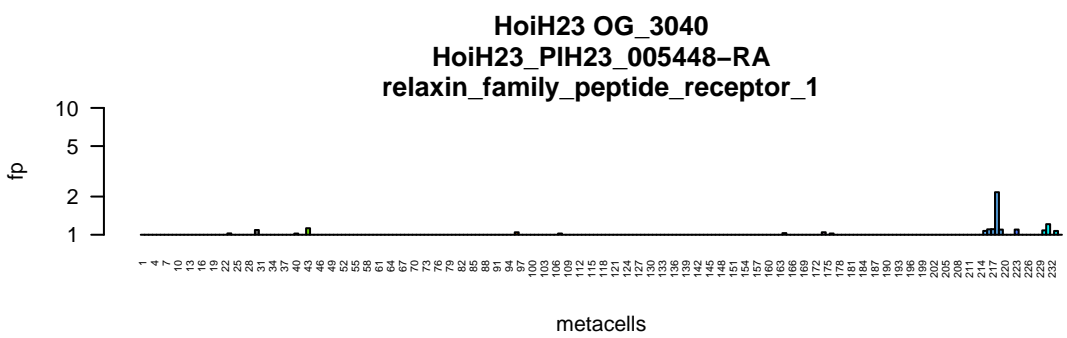
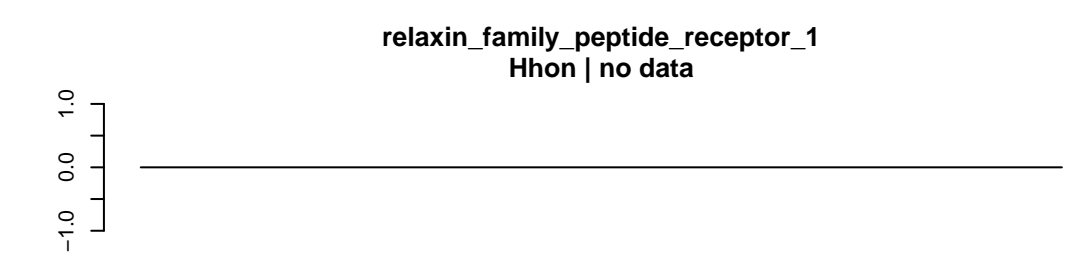
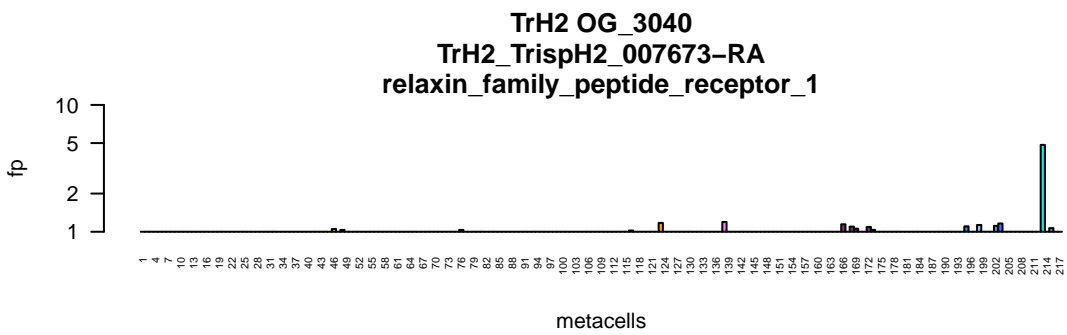
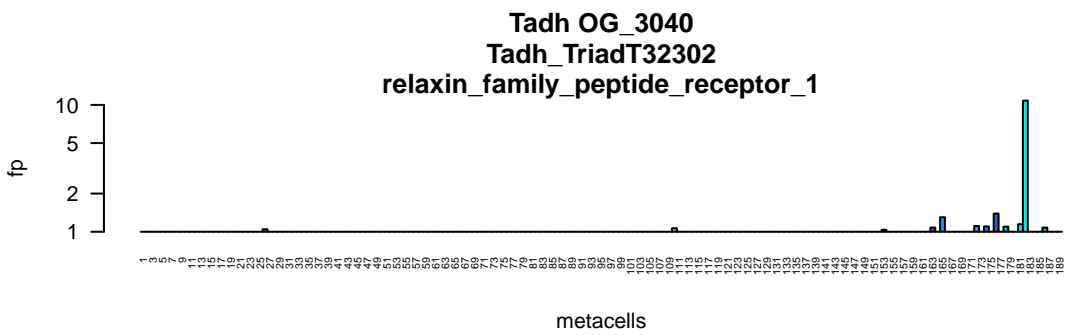
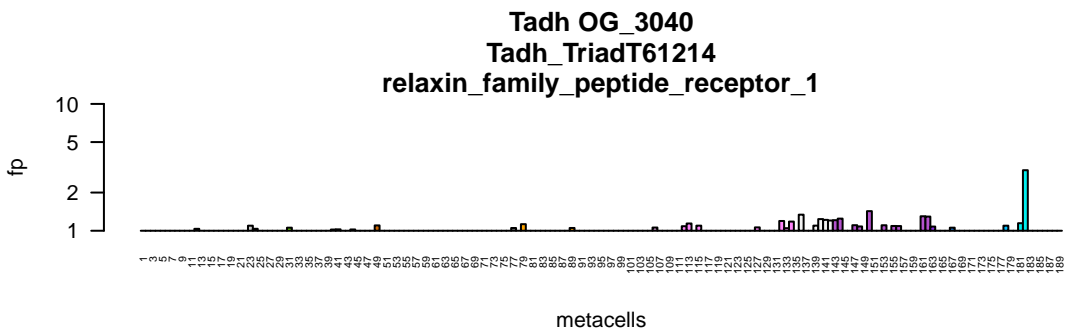
Hhon OG\_1238  
Hhon\_g04472.t1

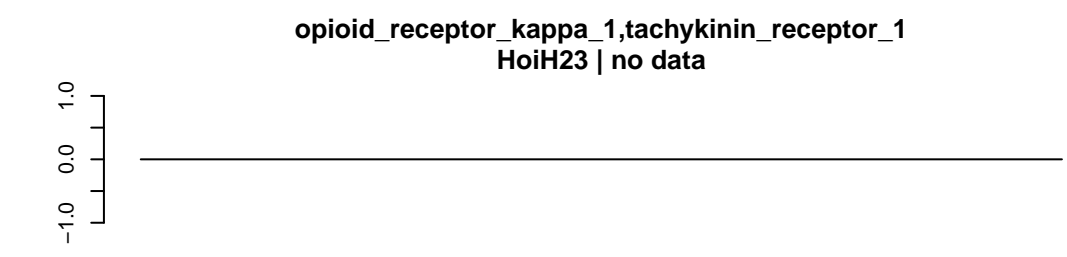
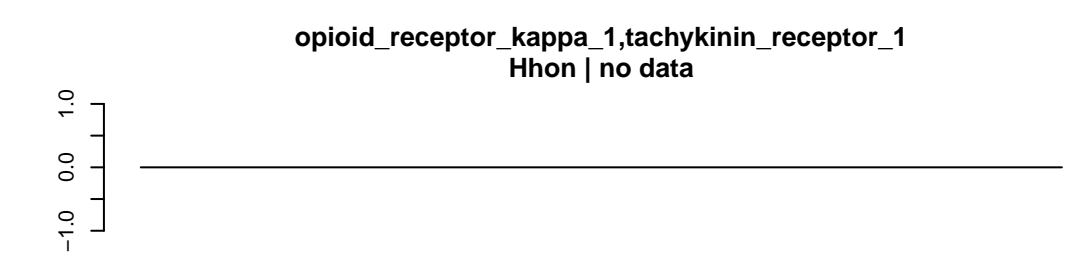
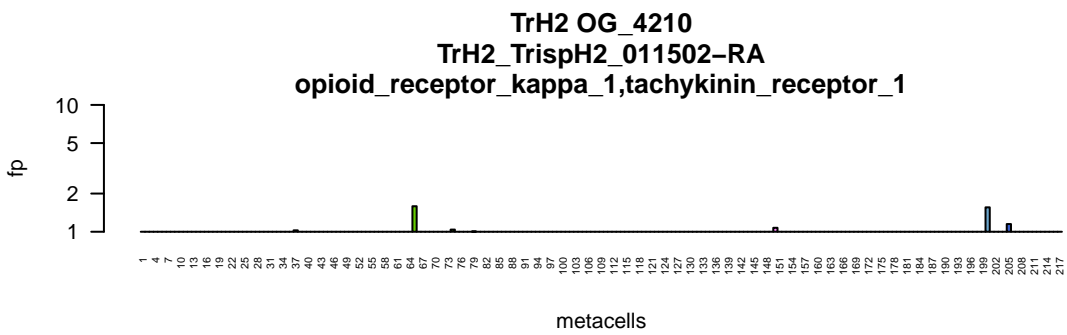
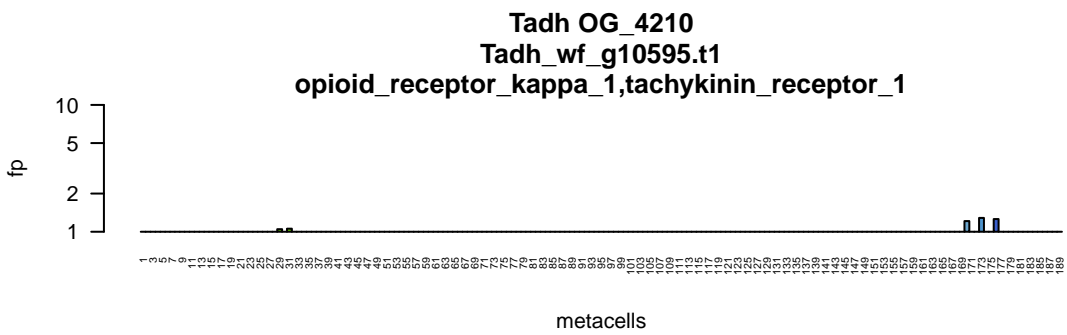
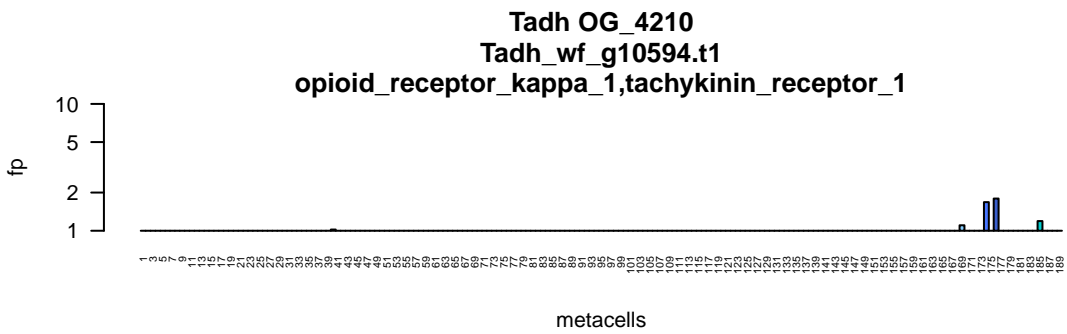


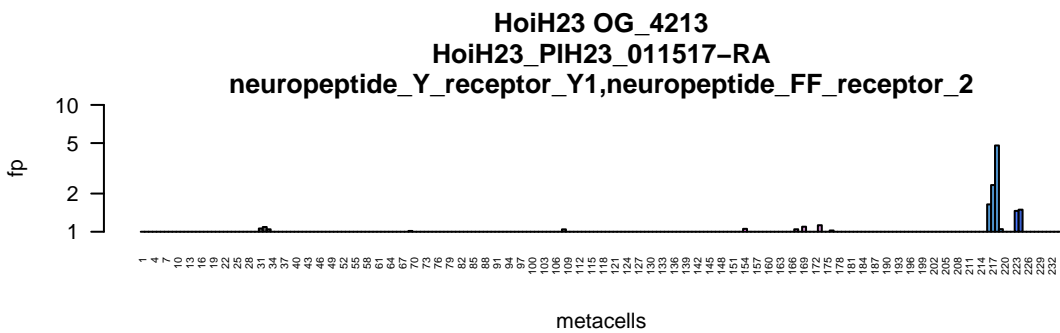
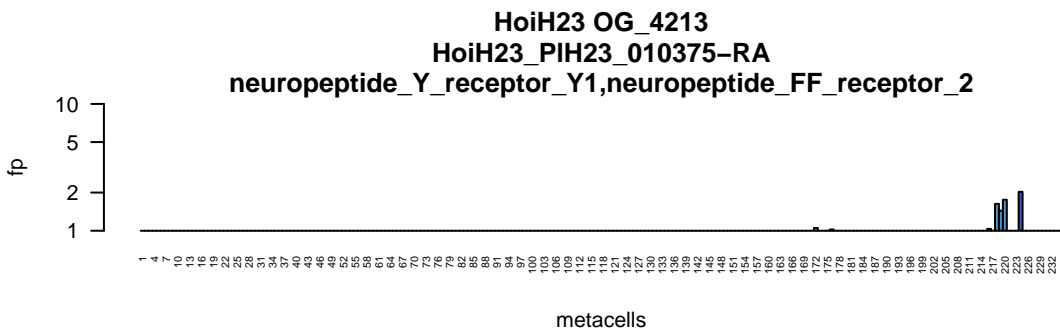
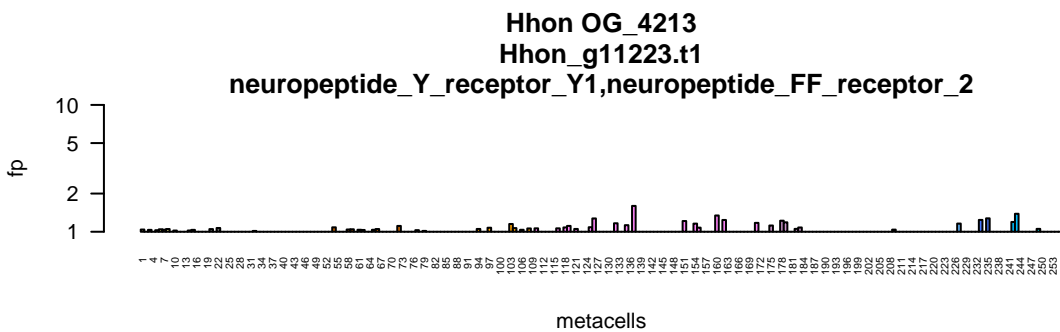
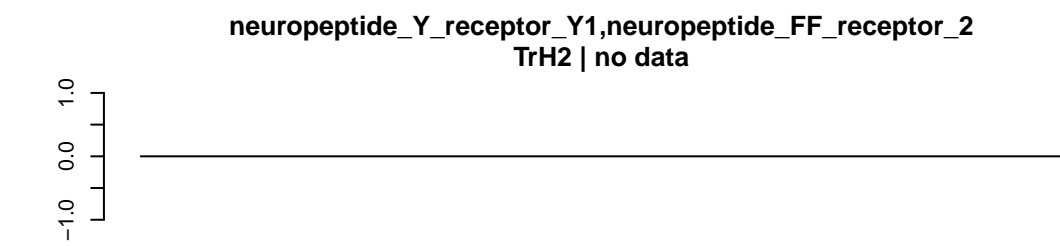
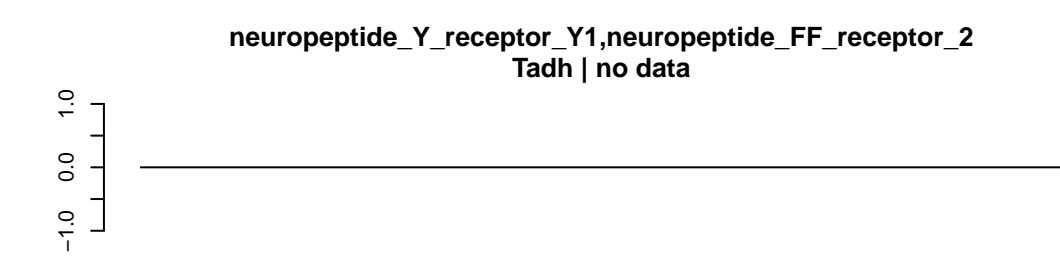
HoiH23 OG\_1238  
HoiH23\_PIH23\_000771-RA

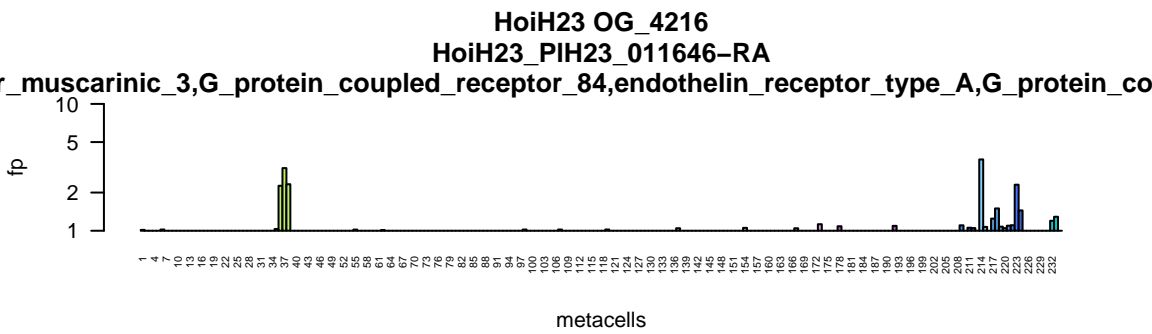
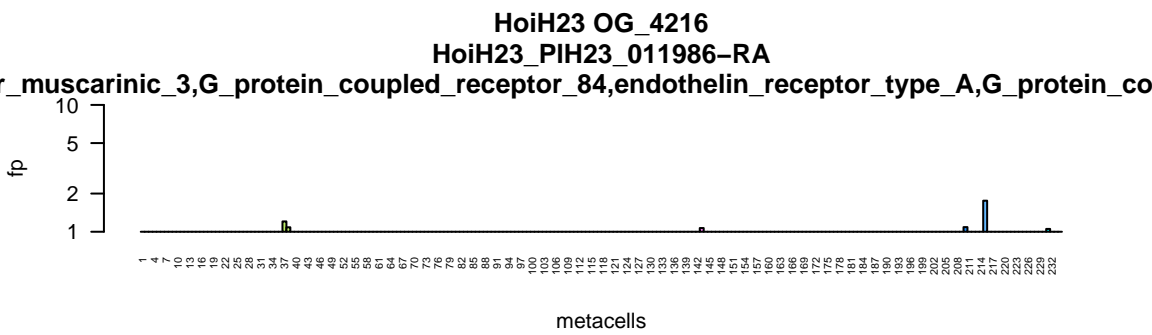
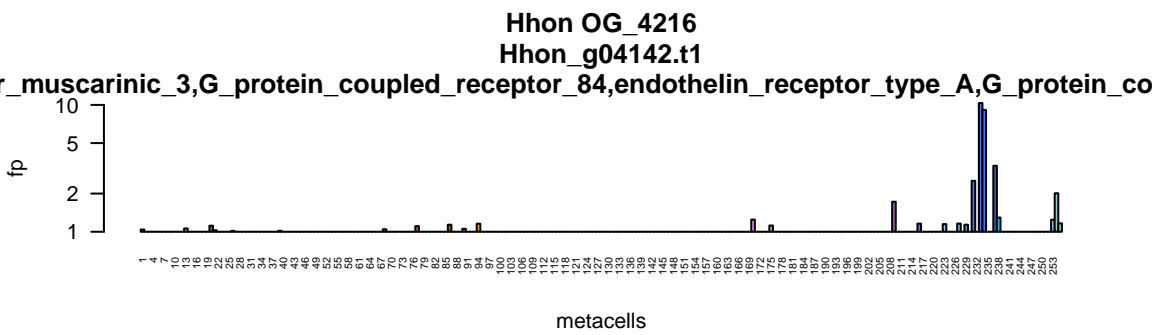
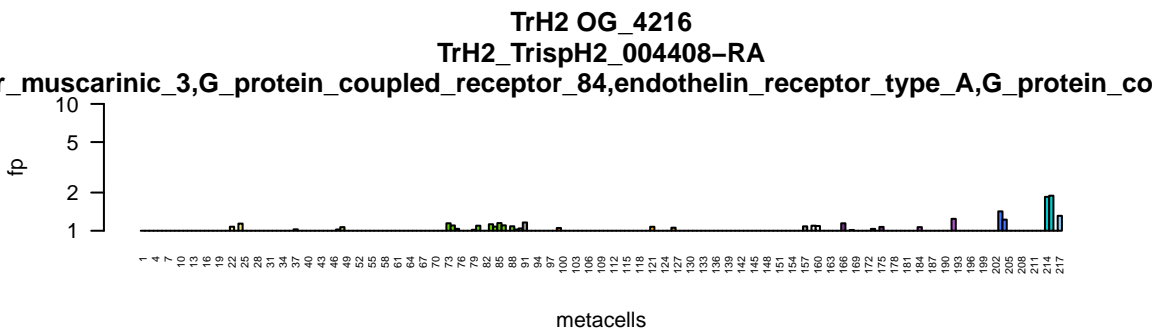
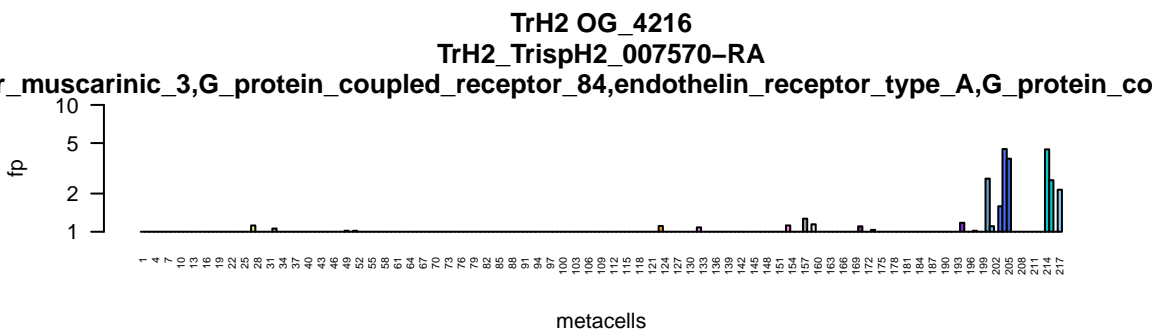
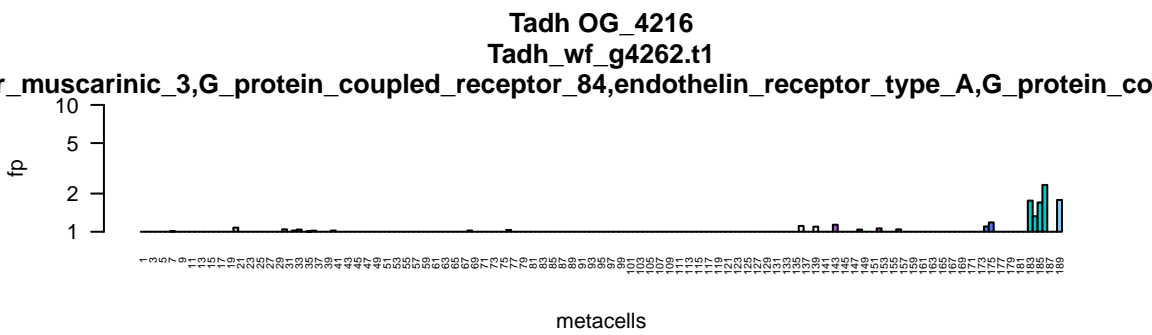
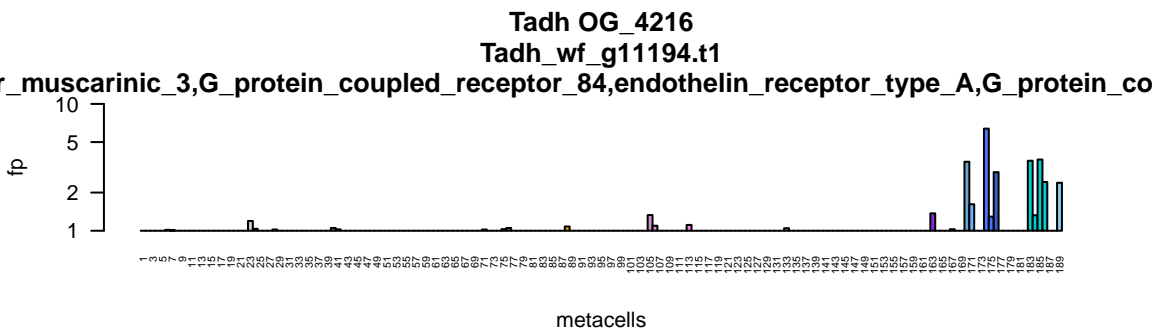


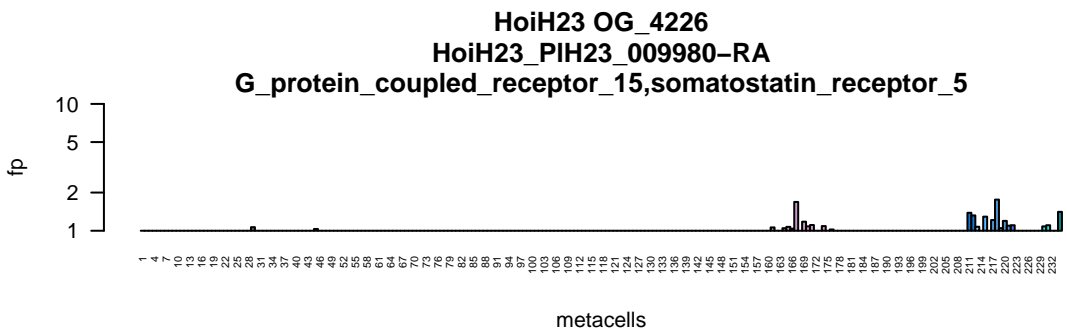
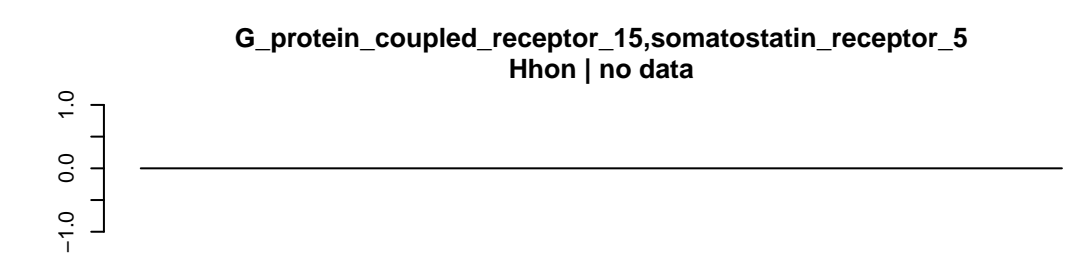
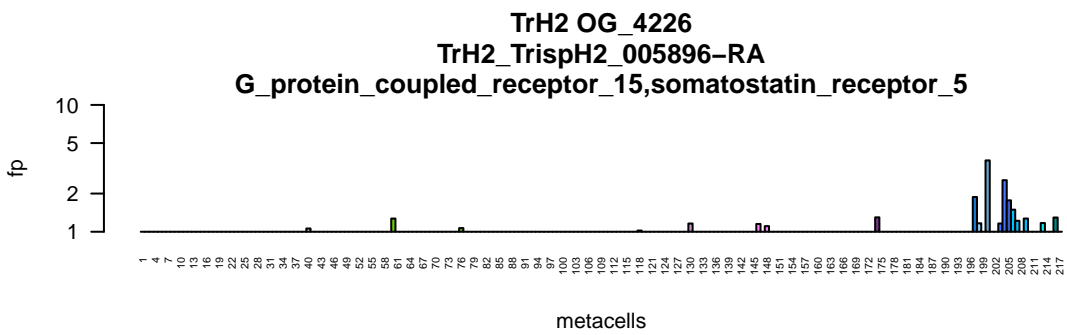
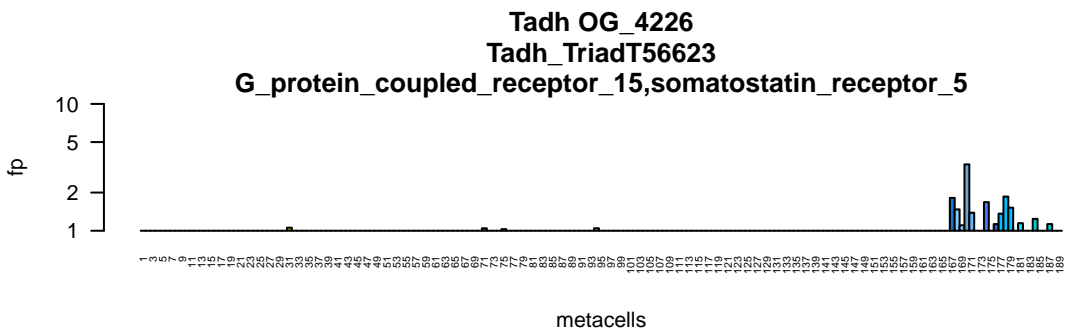












A bar chart showing the frequency of metacells (x-axis) across different frequency bins (y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows a distribution of metacells across frequency bins, with a notable peak at metacell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

metacells	fp
1	1
4	1
7	1
13	1
16	1
19	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
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	4
205	1
208	1
211	1
214	1
217	1

ed\_receptor\_L4,cadherin\_EGF\_LAG\_seven\_pass\_G\_type\_receptor\_1,mannose\_receptor\_C

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

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	1
55	0
58	0
61	0
64	0
67	0
70	0
73	0
76	0
79	0
82	0
85	0
88	0
91	0
94	0
97	0
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	0
130	0
133	1
136	0
139	0
142	0
145	0
148	0
151	0
154	1
155	1
156	0
159	0
162	0
165	0
168	0
171	0
174	0
177	0
180	0
183	0
186	0
189	1
192	0
195	0
198	0
201	2
202	2
203	1
204	1
205	1
206	1
207	1
208	1
209	1
210	1
211	1
214	1
217	1

ed\_receptor\_L4,cadherin\_EGF\_LAG\_seven\_pass\_G\_type\_receptor\_1,mannose\_receptor\_C

metacell	fp
1	1
4	1
13	1
16	2
17	1
19	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
52	1
55	1
58	1
61	1
67	1
70	1
73	1
77	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	2
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
133	2
136	1
139	1
142	1
145	1
148	1
151	1
154	2
157	1
160	2
163	1
166	1
169	1
172	1
175	1
179	1
181	1
184	1
187	1
190	1
193	1.5
196	1.5
199	3
202	3
205	3
208	1.5
211	1.5
214	1.5
217	1.5

metacells

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 indices from 1 to 217. Most metacells have 1 false positive, but metacells 202 and 214 have 2 false positives.

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

ed\_receptor\_L4,cadherin\_EGF\_LAG\_seven\_pass\_G\_type\_receptor\_1,mannose\_receptor\_C

fp

metacells

Bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis lists metacells from 1 to 253. The y-axis shows fp values from 1 to 10. Most metacells have a frequency of 1, with a few having higher frequencies (e.g., metacell 220 has a frequency of 2).

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 253. 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
19	1
22	1
28	1
31	1
37	1
43	1
46	1
55	1
61	1
64	1
70	1
73	1
79	1
88	1
91	1
97	1
100	1
109	1
115	1
119	1
127	1
133	1
136	1
146	1
151	1
154	1
160	1
163	1
169	1
175	1
184	1
187	1
193	1
196	1
202	1
205	1
211	1
214	1
220	1
223	1
229	1
232	1
238	1
241	1
247	1
250	1
253	2

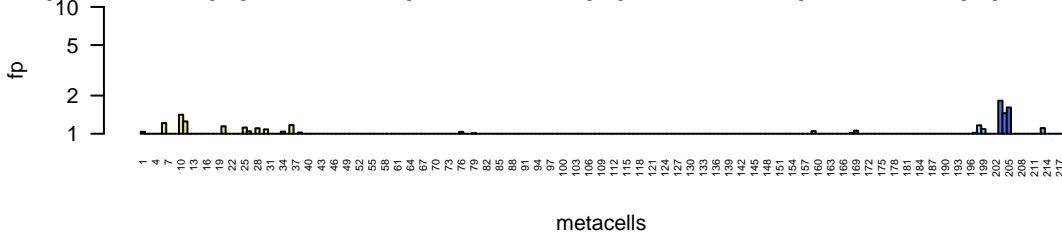


A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 232. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a notable peak at metacell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

receptor\_1,neuropeptide\_Y\_receptor\_Y1,neuropeptide\_FF\_receptor\_2,neuropeptide\_FF\_rec  
Tadh | no data



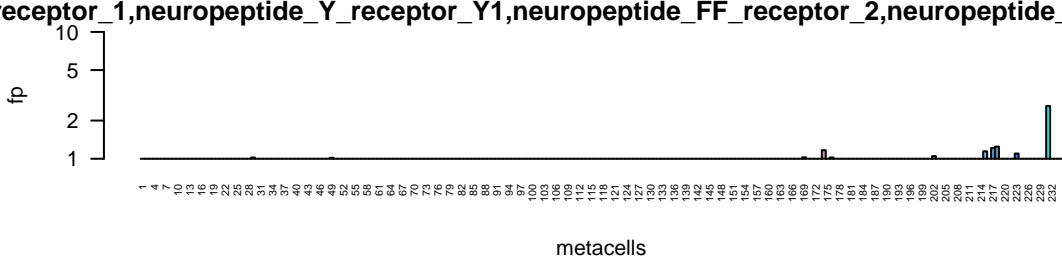
TrH2 OG\_4548  
TrH2\_TrispH2\_007473-RA  
receptor\_1,neuropeptide\_Y\_receptor\_Y1,neuropeptide\_FF\_receptor\_2,neuropeptide\_FF\_rec



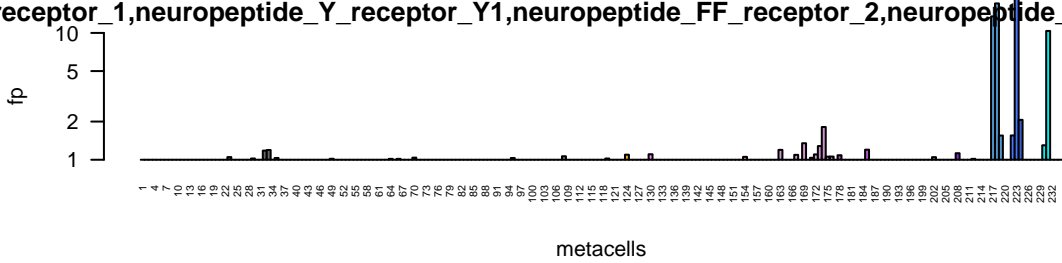
receptor\_1,neuropeptide\_Y\_receptor\_Y1,neuropeptide\_FF\_receptor\_2,neuropeptide\_FF\_rec  
Hhon | no data



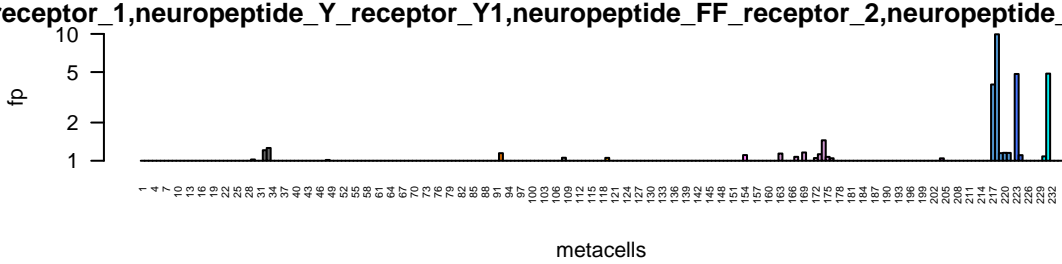
HoiH23 OG\_4548  
HoiH23\_PIH23\_006484-RA  
receptor\_1,neuropeptide\_Y\_receptor\_Y1,neuropeptide\_FF\_receptor\_2,neuropeptide\_FF\_rec



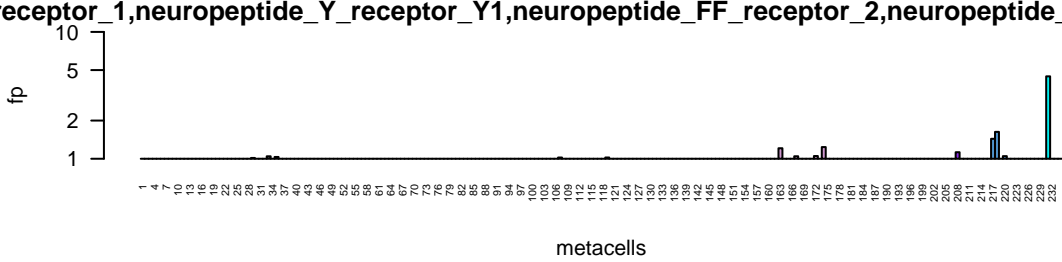
HoiH23 OG\_4548  
HoiH23\_PIH23\_011478-RA  
receptor\_1,neuropeptide\_Y\_receptor\_Y1,neuropeptide\_FF\_receptor\_2,neuropeptide\_FF\_rec



HoiH23 OG\_4548  
HoiH23\_PIH23\_011479-RA  
receptor\_1,neuropeptide\_Y\_receptor\_Y1,neuropeptide\_FF\_receptor\_2,neuropeptide\_FF\_rec



HoiH23 OG\_4548  
HoiH23\_PIH23\_011481-RA  
receptor\_1,neuropeptide\_Y\_receptor\_Y1,neuropeptide\_FF\_receptor\_2,neuropeptide\_FF\_rec



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

fp

metacells

**Tadh OG\_4762**  
**Tadh\_wf\_g8464.t1**  
**hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chen**

metacells	fp
1	1.0
2	1.0
3	1.0
4	1.0
5	1.0
6	1.0
7	1.0
8	1.0
9	1.0
10	1.0
11	1.0
12	1.0
13	1.0
14	1.0
15	1.0
16	1.0
17	1.0
18	1.0
19	1.0
20	1.0
21	1.0
22	1.0
23	1.0
24	1.0
25	1.0
26	1.0
27	1.0
28	1.0
29	1.0
30	1.0
31	1.0
32	1.0
33	1.0
34	1.0
35	1.0
36	1.0
37	1.0
38	1.0
39	1.0
40	1.0
41	1.0
42	1.0
43	1.0
44	1.0
45	1.0
46	1.0
47	1.0
48	1.0
49	1.0
50	1.0
51	1.0
52	1.0
53	1.0
54	1.0
55	1.0
56	1.0
57	1.0
58	1.0
59	1.0
60	1.0
61	1.0
62	1.0
63	1.0
64	1.0
65	1.0
66	1.0
67	1.0
68	1.0
69	1.0
70	1.0
71	1.0
72	1.0
73	1.0
74	1.0
75	1.0
76	1.0
77	1.0
78	1.0
79	1.0
80	1.0
81	1.0
82	1.0
83	1.0
84	1.0
85	1.0
86	1.0
87	1.0
88	1.0
89	1.0
90	1.0
91	1.0
92	1.0
93	1.0
94	1.0
95	1.0
96	1.0
97	1.0
98	1.0
99	1.0
100	1.0
101	1.0
102	1.0
103	1.0
104	1.0
105	1.0
106	1.0
107	1.0
108	1.0
109	1.0
110	1.0
111	1.0
112	1.0
113	1.0
114	1.0
115	1.0
116	1.0
117	1.0
118	1.0
119	1.0
120	1.0
121	1.0
122	1.0
123	1.0
124	1.0
125	1.0
126	1.0
127	1.0
128	1.0
129	1.0
130	1.0
131	1.0
132	1.0
133	1.0
134	1.0
135	1.0
136	1.0
137	1.0
138	1.0
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140	1.0
141	1.0
142	1.0
143	1.0
144	1.0
145	1.0
146	1.0
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152	1.0
153	1.0
154	1.0
155	1.0
156	1.0
157	1.0
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165	1.0
166	1.0
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168	1.0
169	1.0
170	1.0
171	1.0
172	1.0
173	1.0
174	1.0
175	1.0
176	1.0
177	1.0
178	1.0
179	1.0
180	1.0
181	1.0
182	1.0
183	1.0
184	1.0
185	1.0
186	1.0
187	1.0
188	1.0
189	1.0
190	1.0
191	1.0
192	1.0
193	1.0
194	1.0
195	1.0
196	1.0
197	1.0
198	1.0
199	1.0
200	1.0
201	1.0

TrH2 OG\_4762  
TrH2\_TrispH2\_010271-RA

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

fp

metacells

Tadh OG\_4762  
Tadh\_TriadT59534  
hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chen

fp

metacells

Hhon OG\_4762  
Hhon\_g10761.t1

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

fp

metacells

TrH2 OG\_4762  
TrH2\_TrispH2\_010634-RA  
hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chen

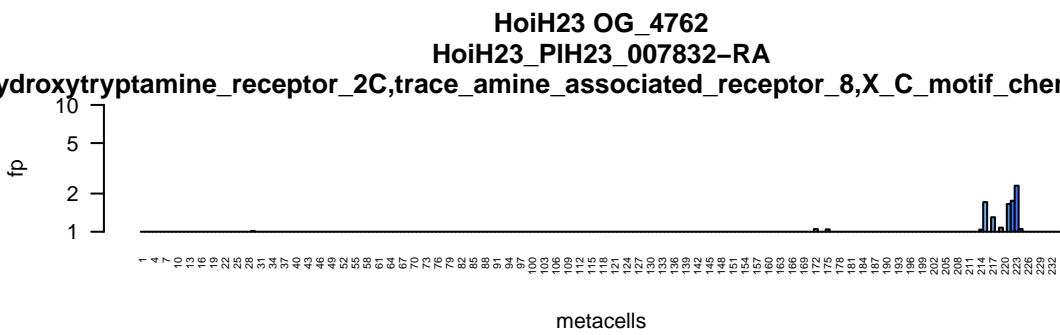
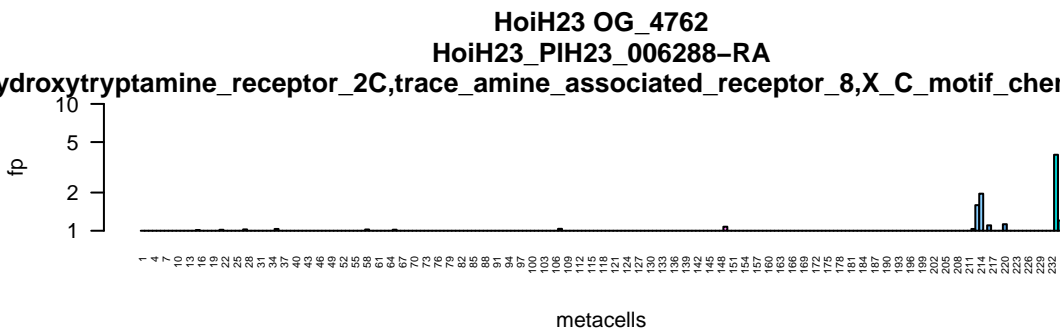
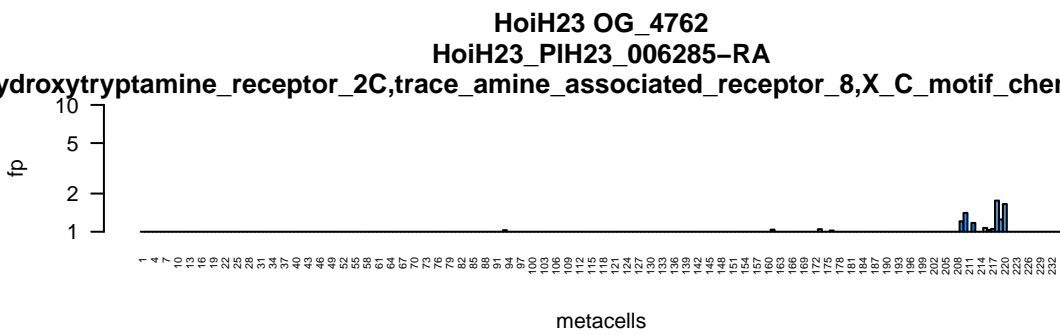
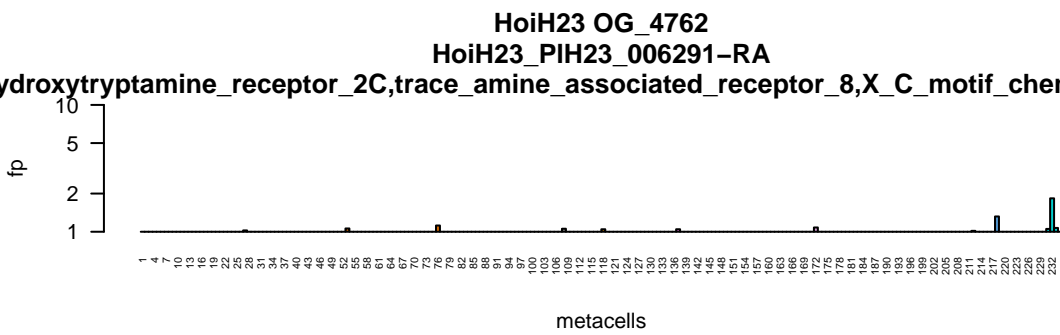
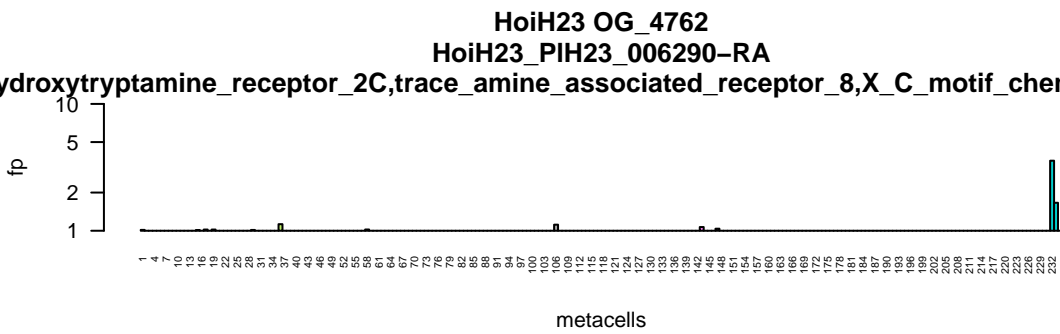
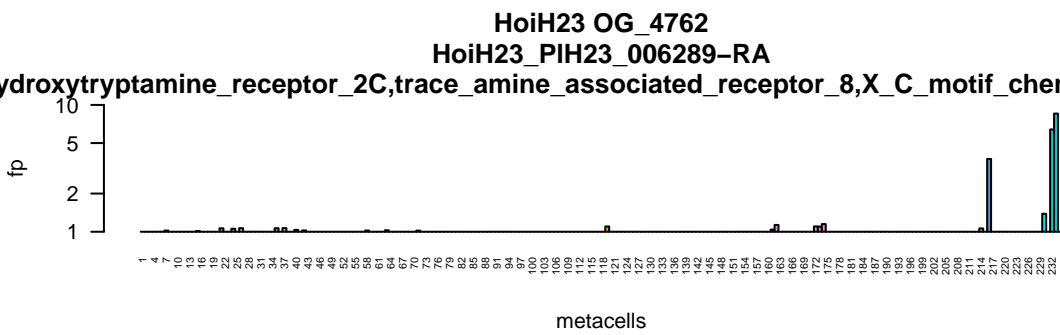
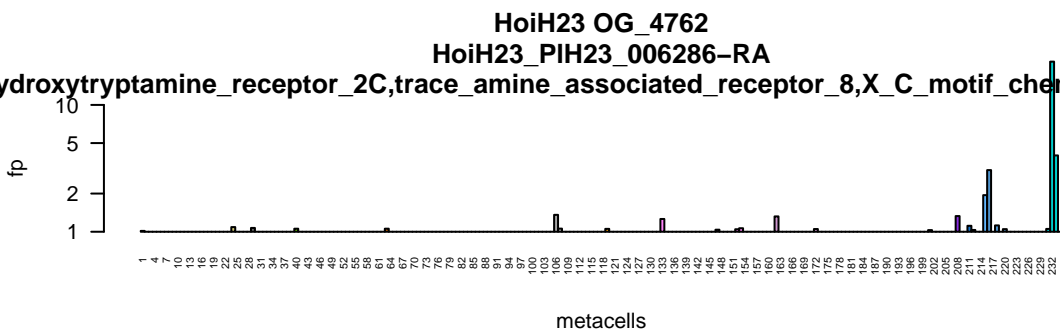
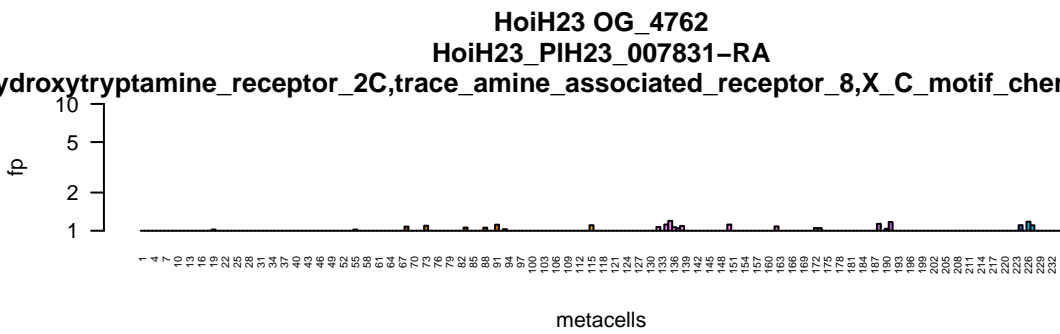
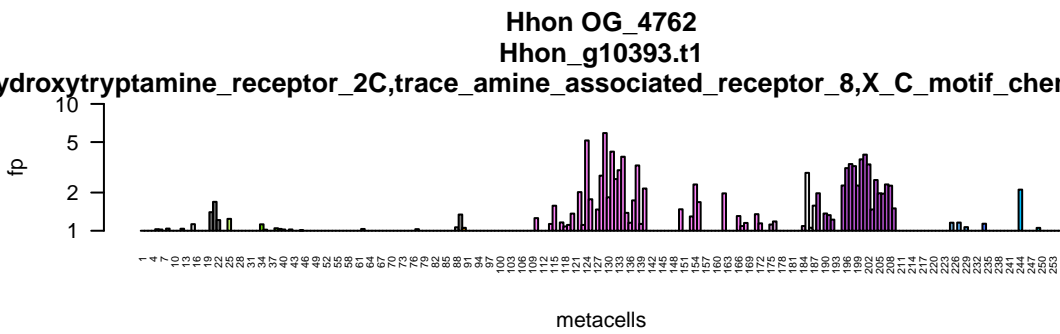
metacells	fp
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10	1
13	1
16	1
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22	1
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28	1
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40	1
43	1
46	1
52	1
55	1
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61	1
64	1
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157	1
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169	1
172	1
175	1
178	1
181	1
184	1
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211	1.1
214	1.8
217	2.2

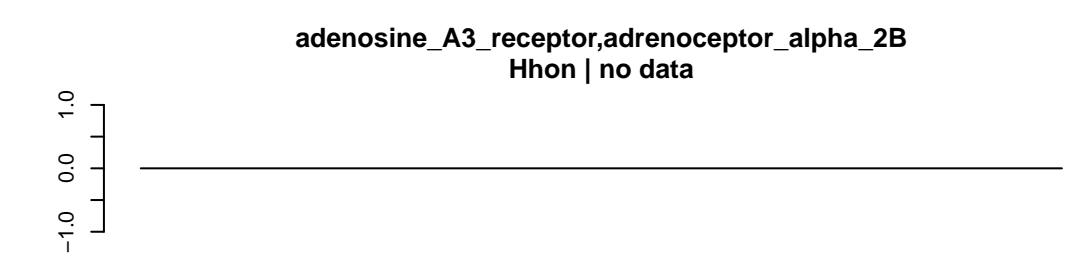
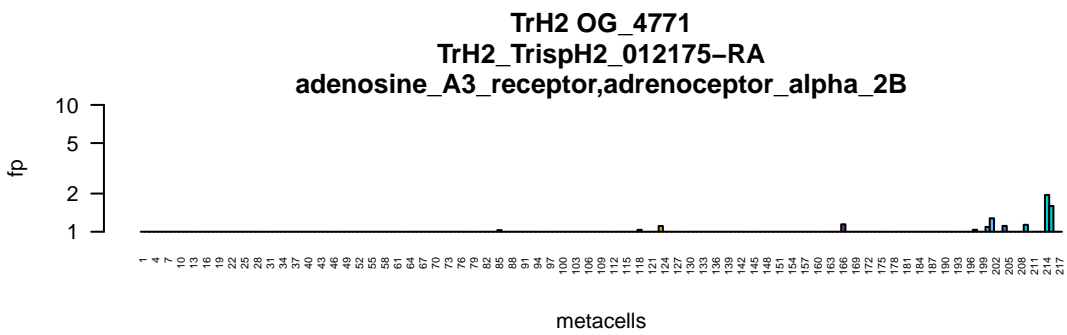
**Hhon OG\_4762**  
**Hhon\_g01703.t1**

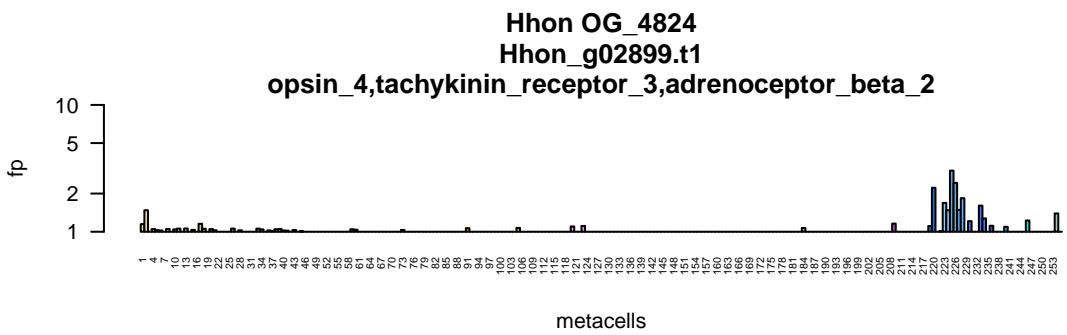
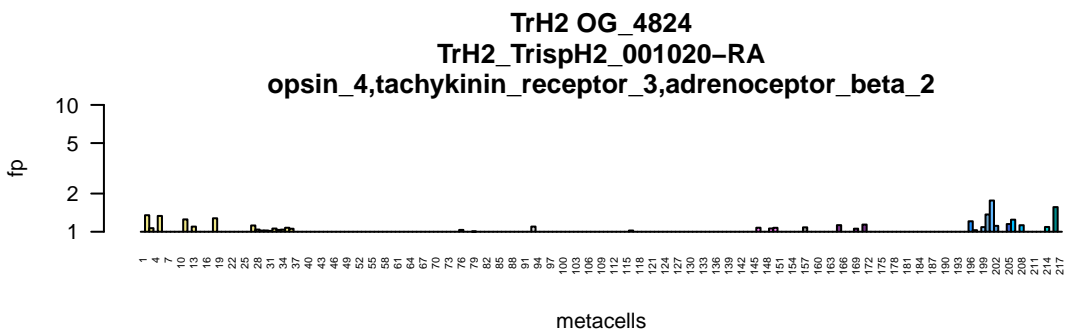
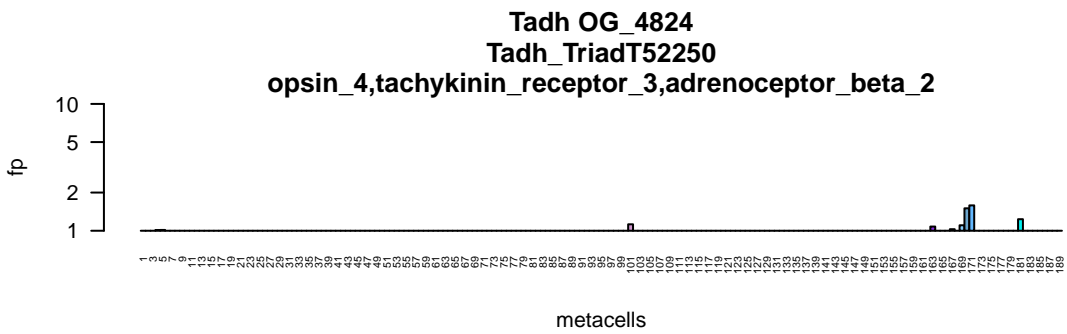
hydroxytryptamine\_receptor\_2C,trace\_amine\_associated\_receptor\_8,X\_C\_motif\_chen

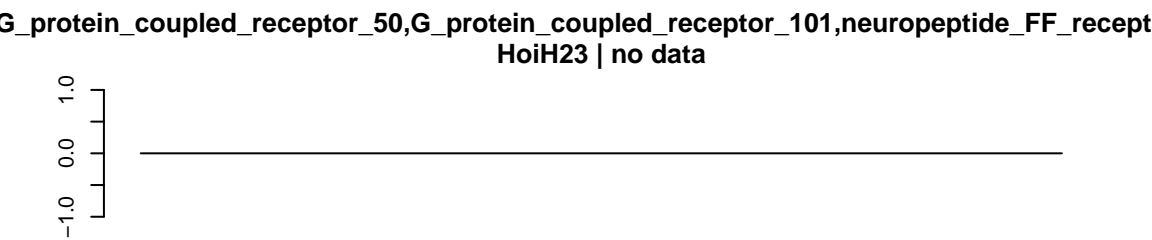
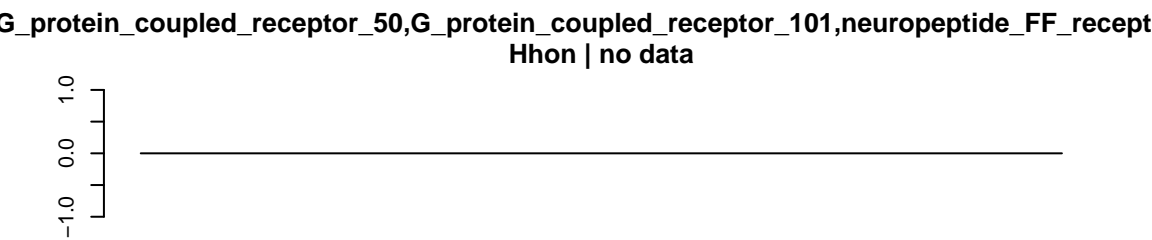
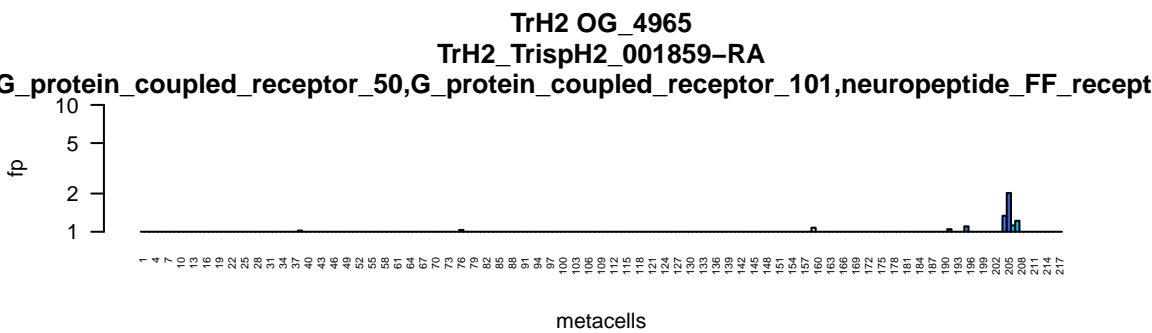
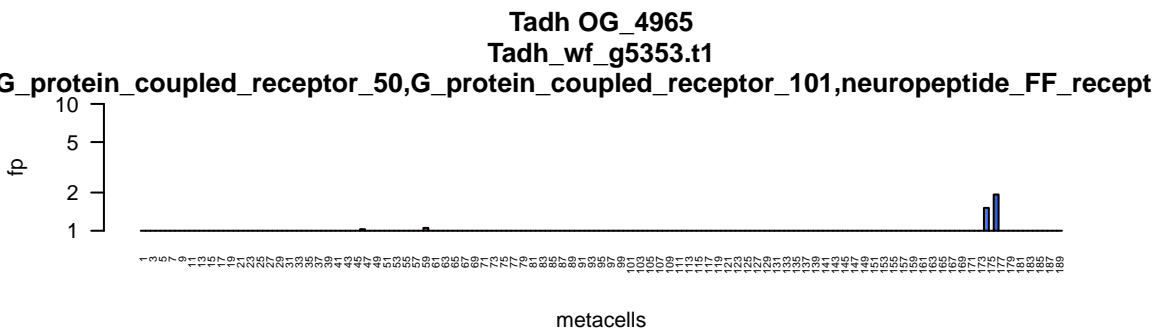
fp

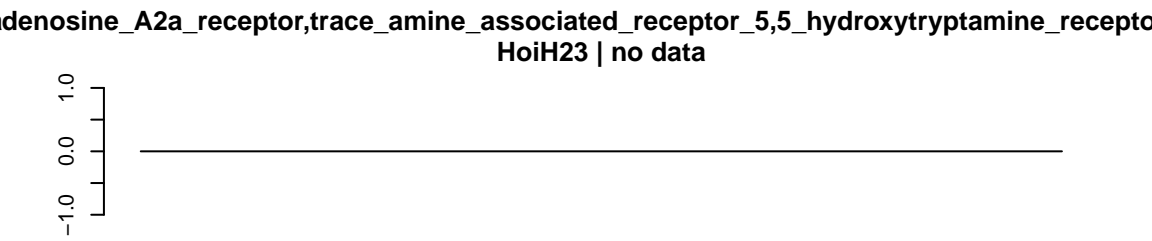
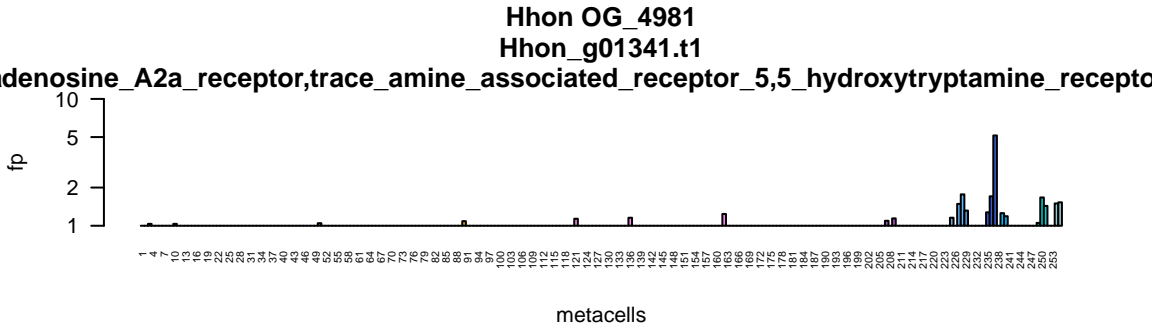
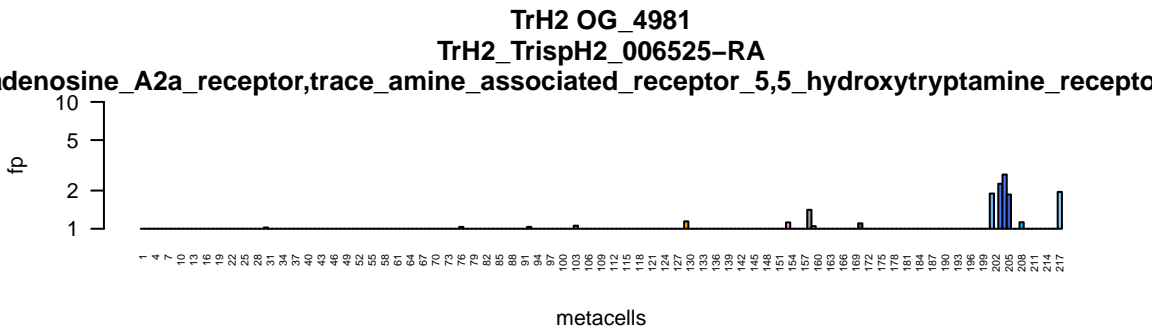
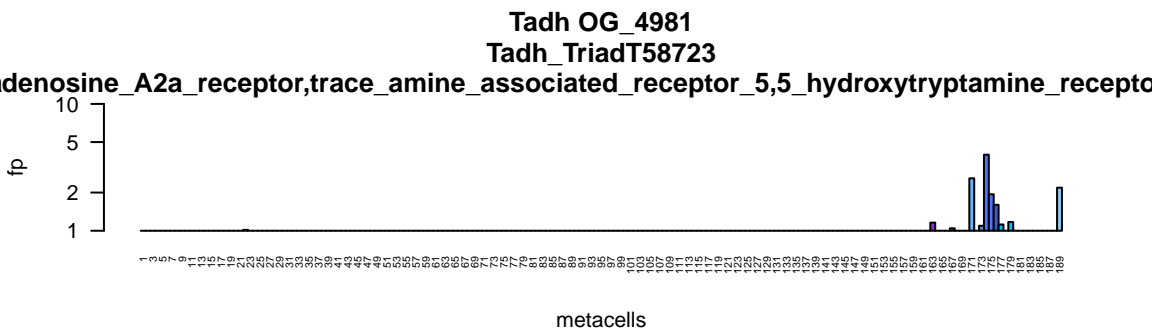
metacells



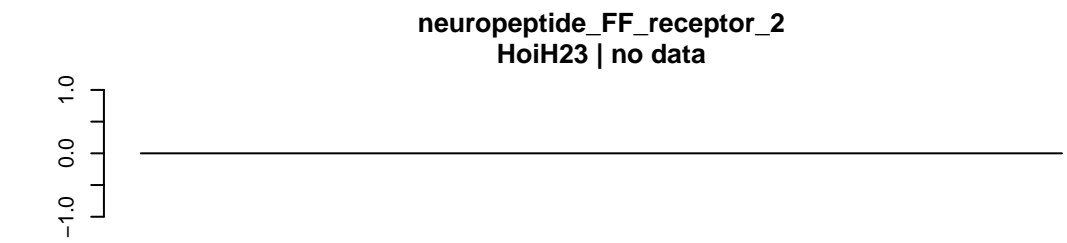
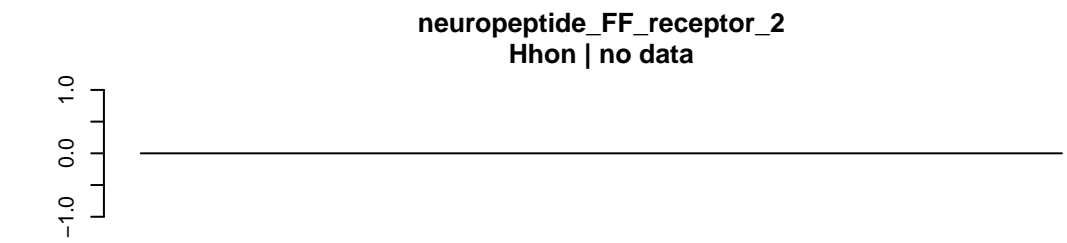
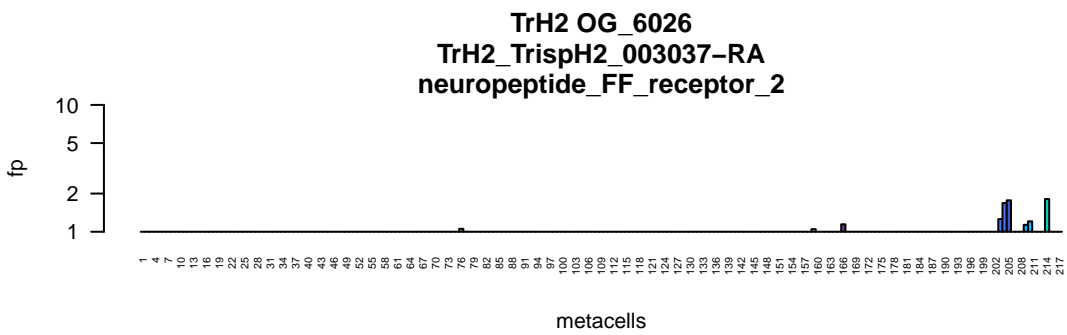
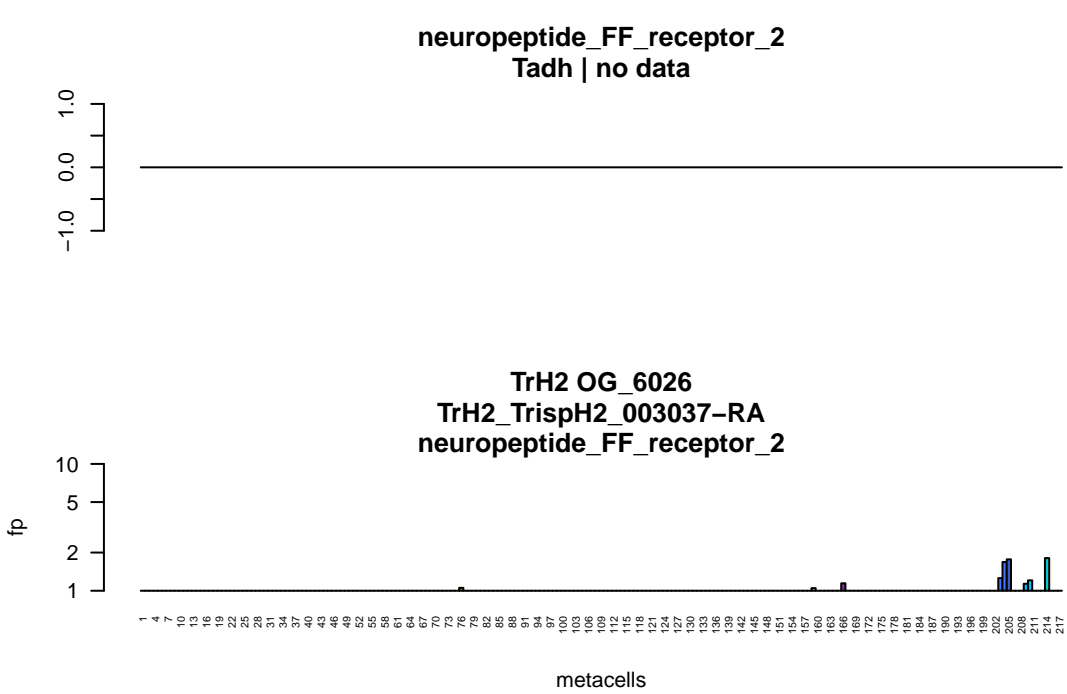


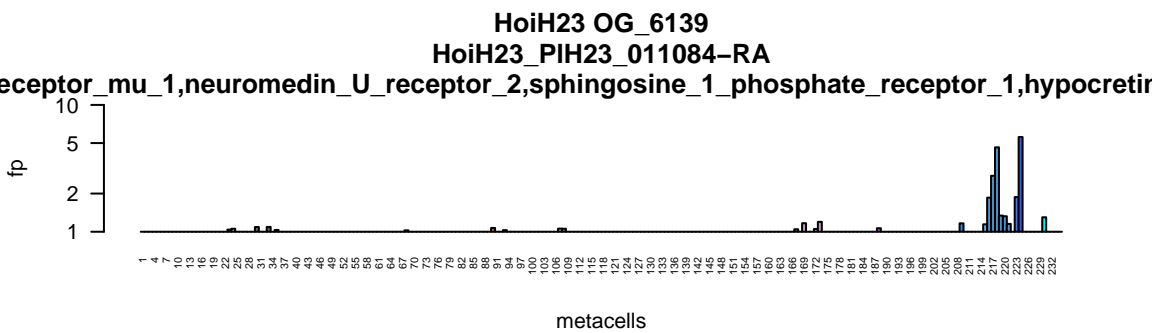
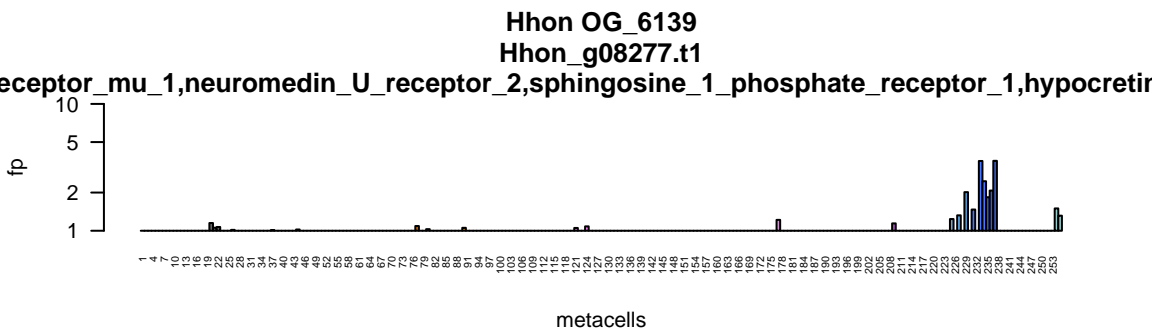
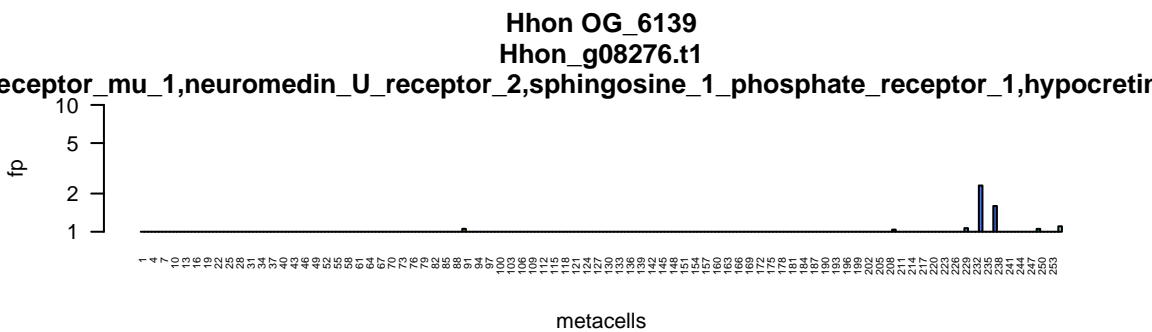
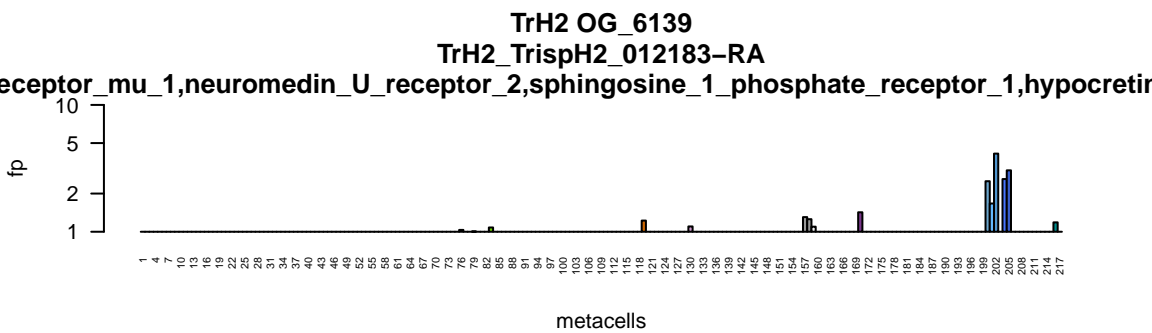
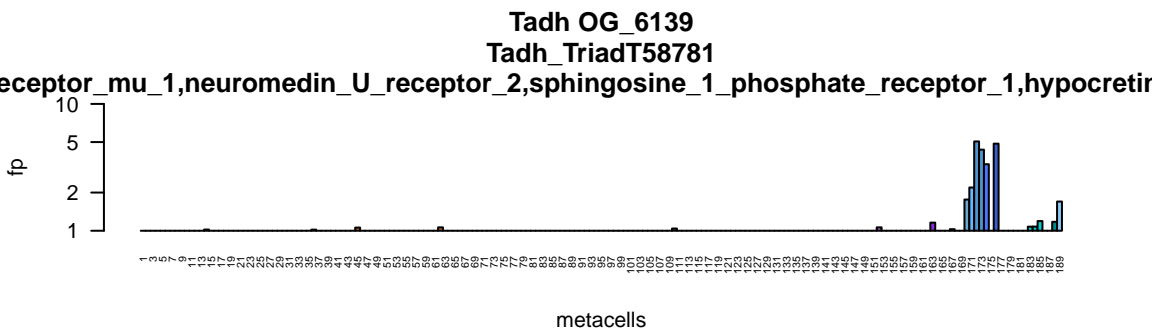


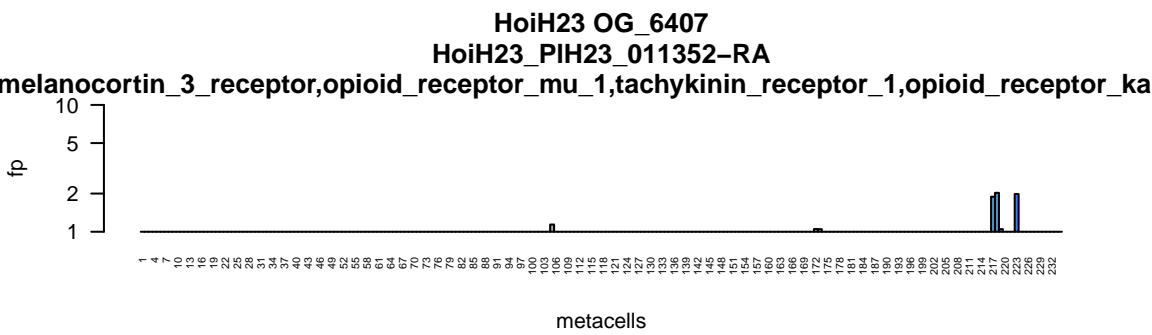
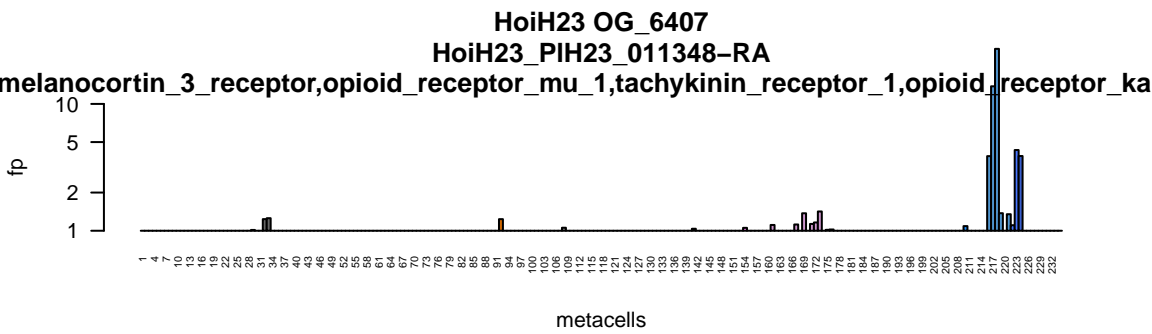
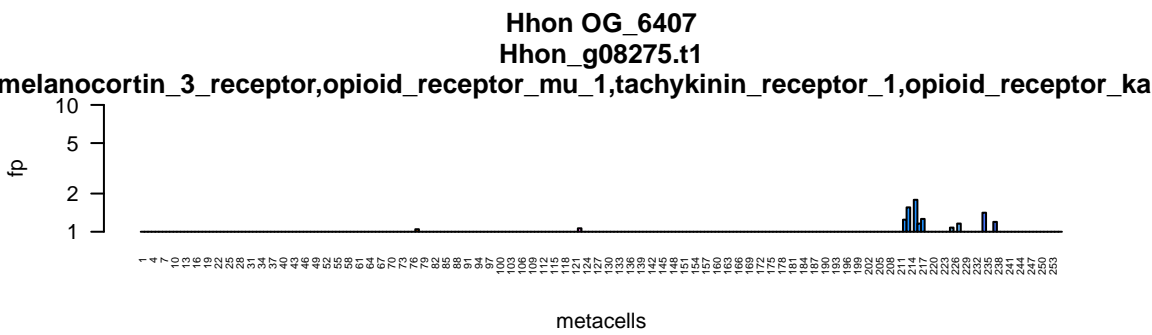
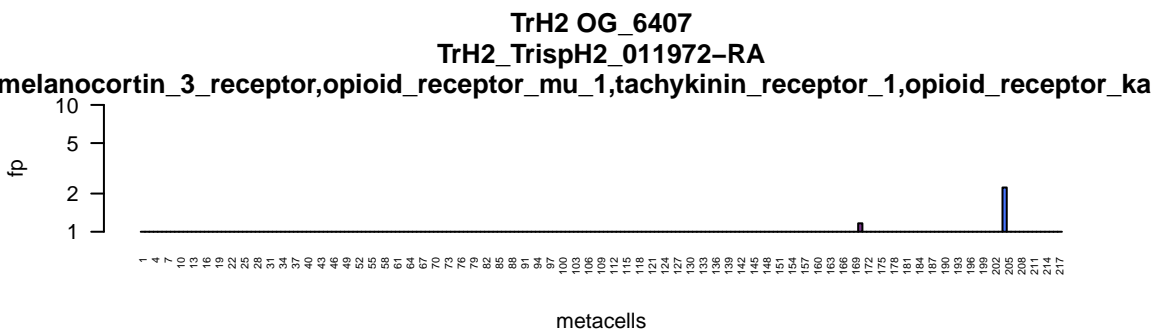
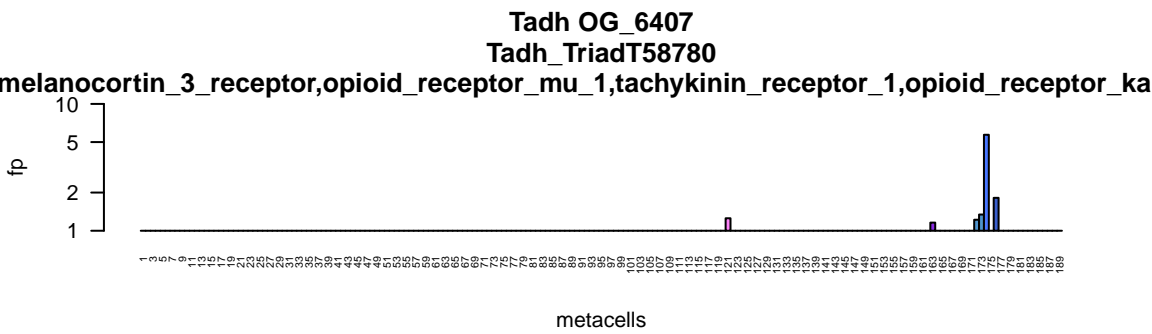












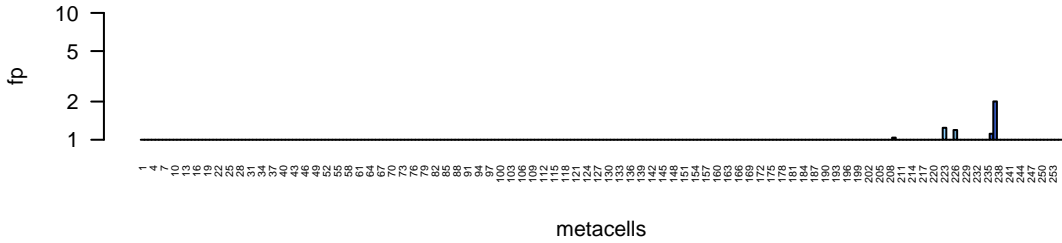
Tadh | no data



TrH2 | no data

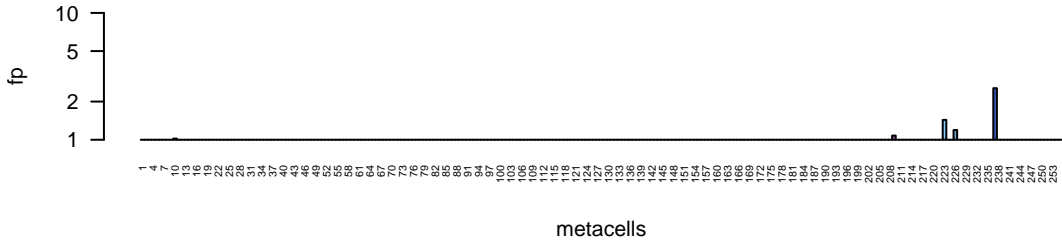


Hhon OG\_6424  
Hhon\_g11706.t1



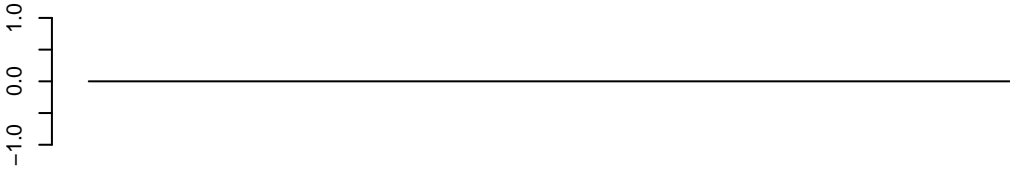
metacells

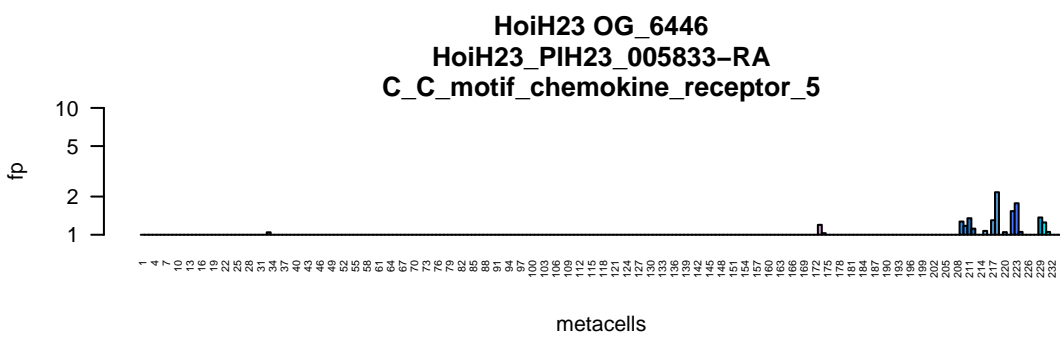
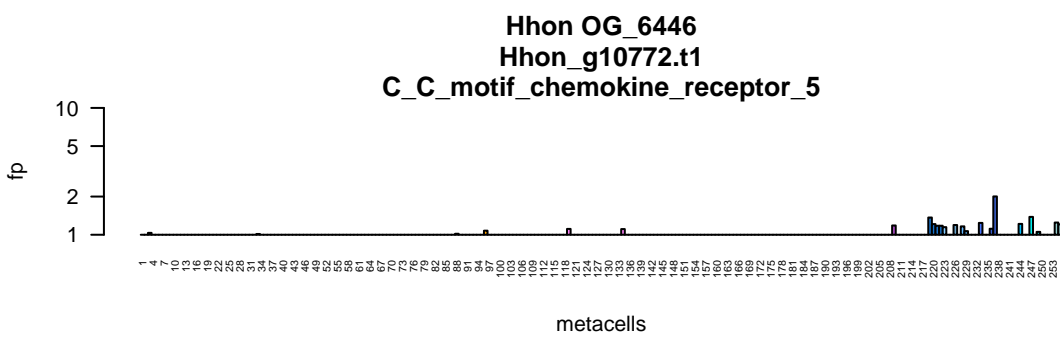
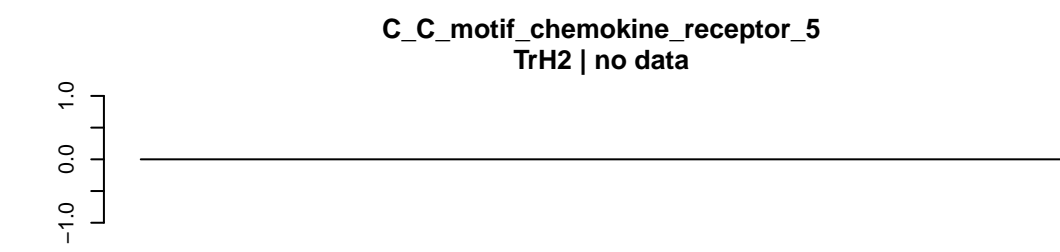
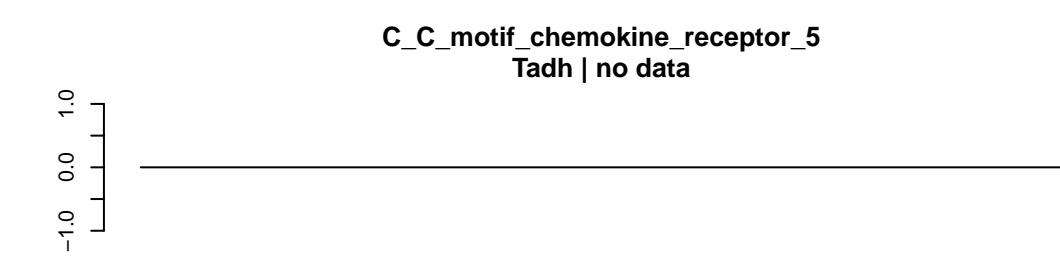
Hhon OG\_6424  
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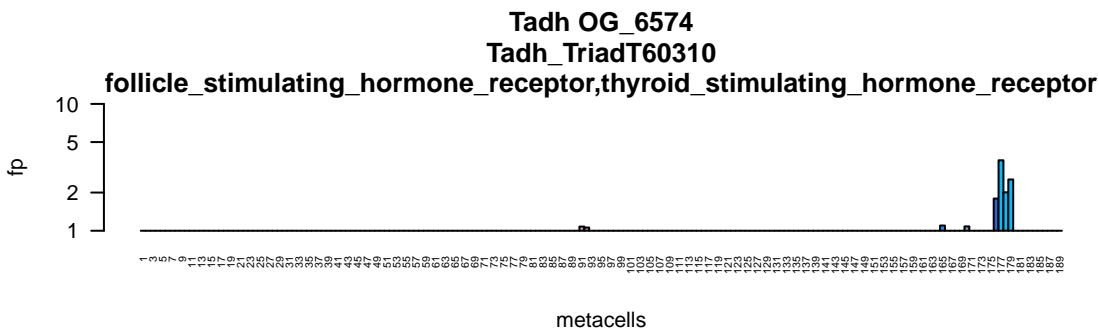


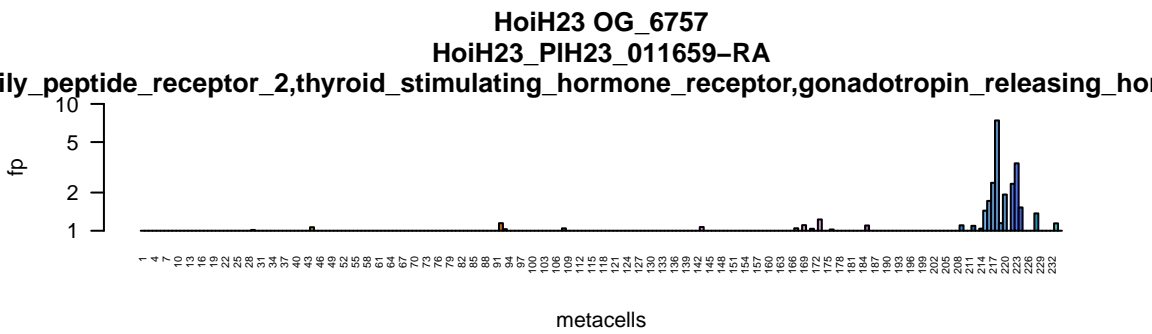
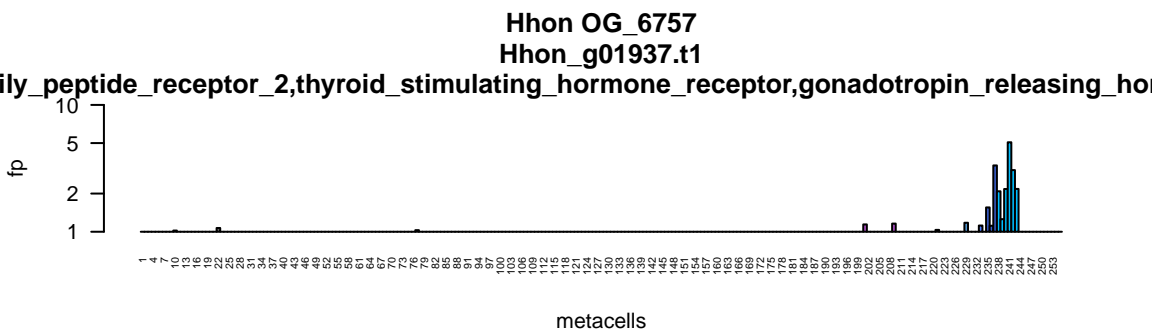
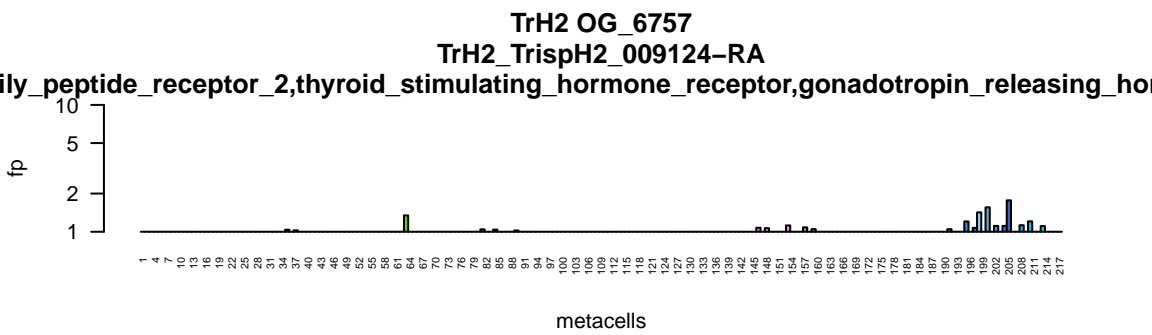
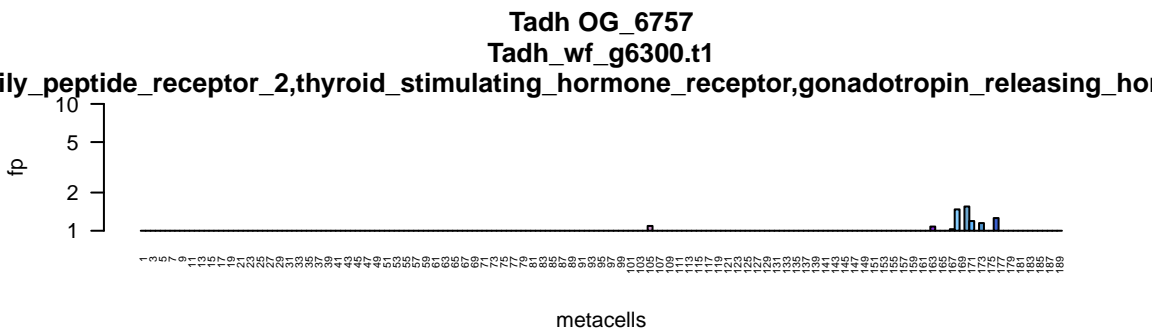
metacells

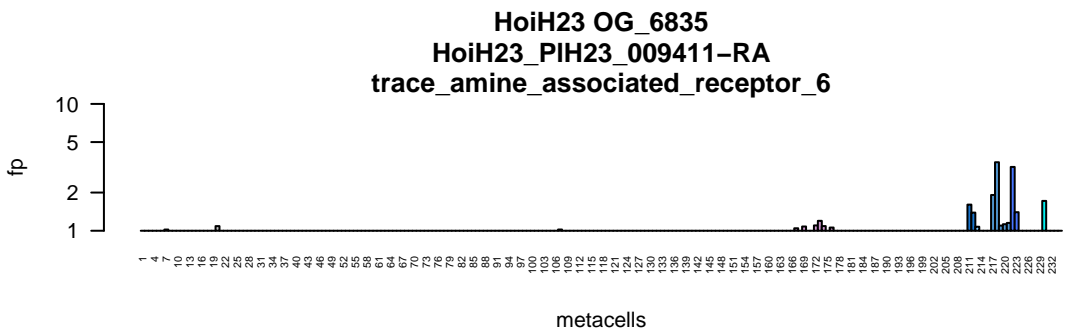
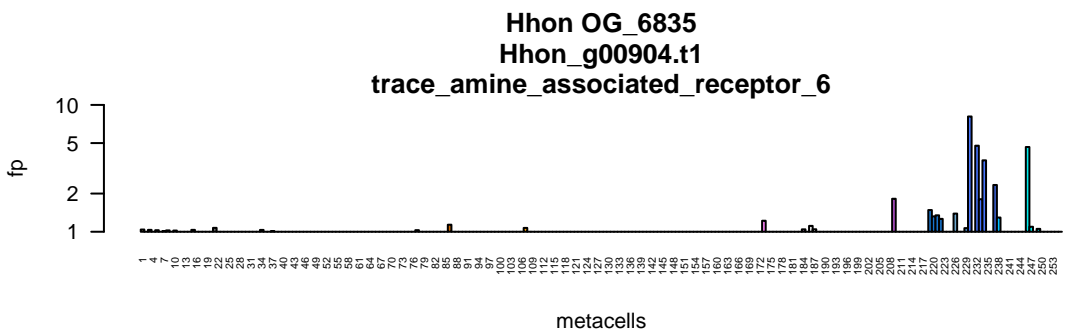
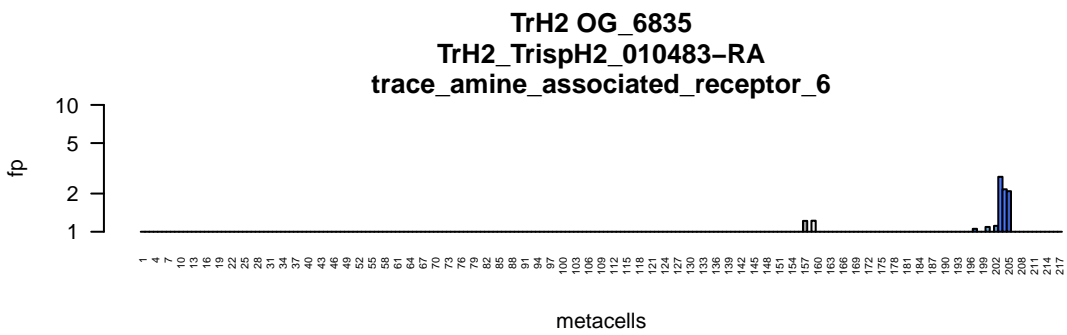
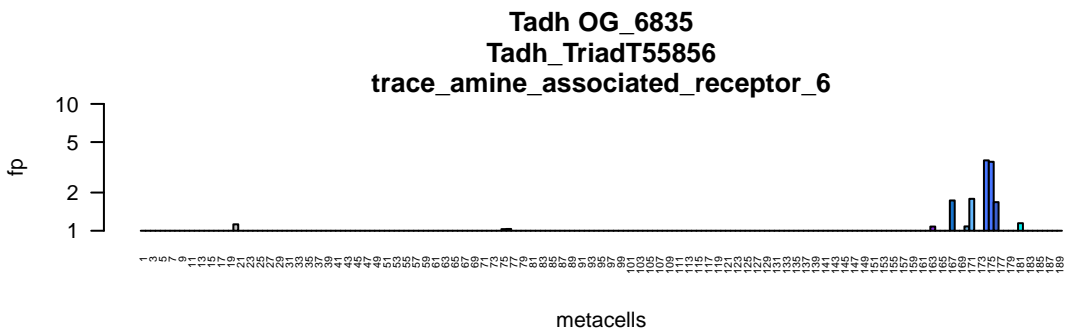
HoiH23 | no data



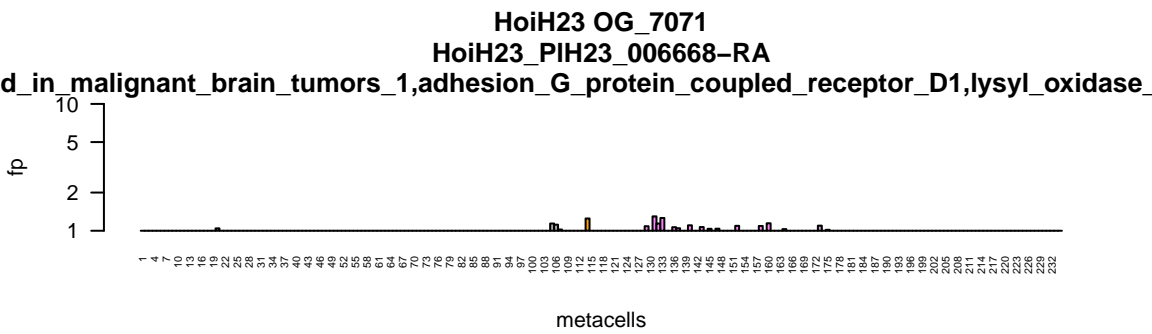
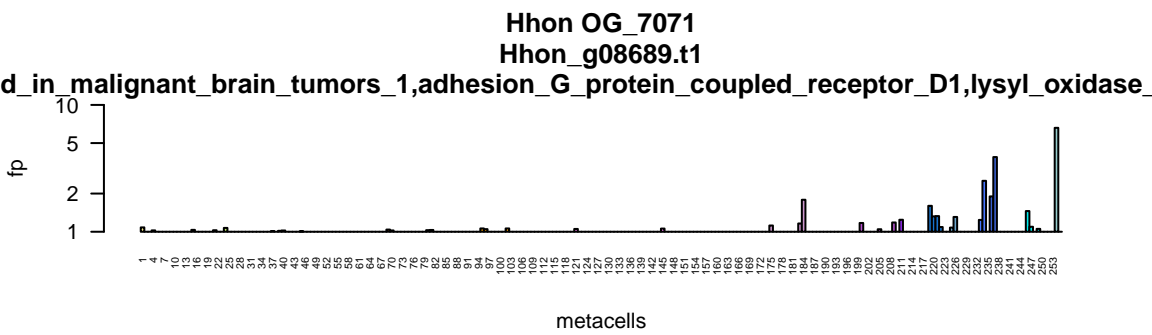
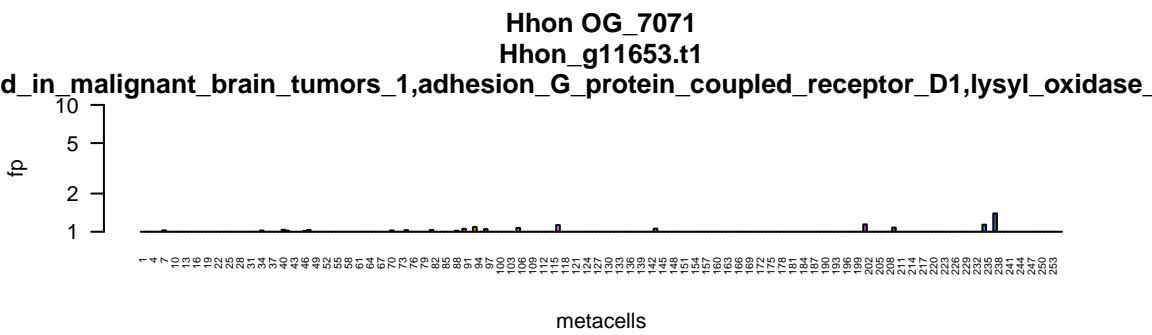
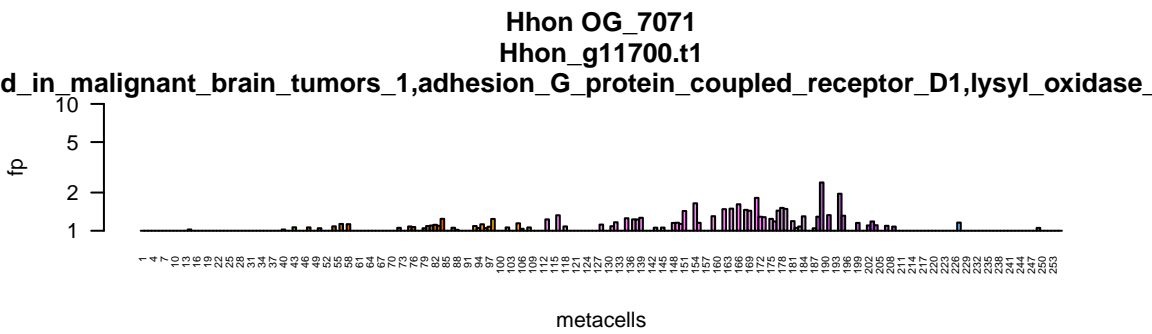
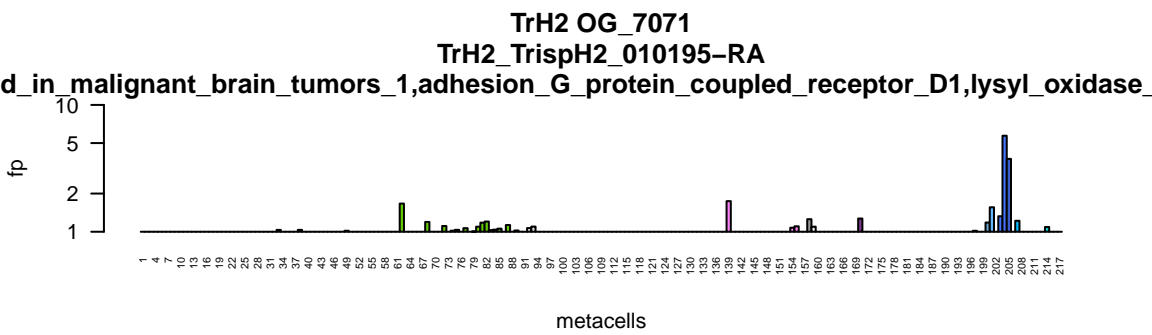
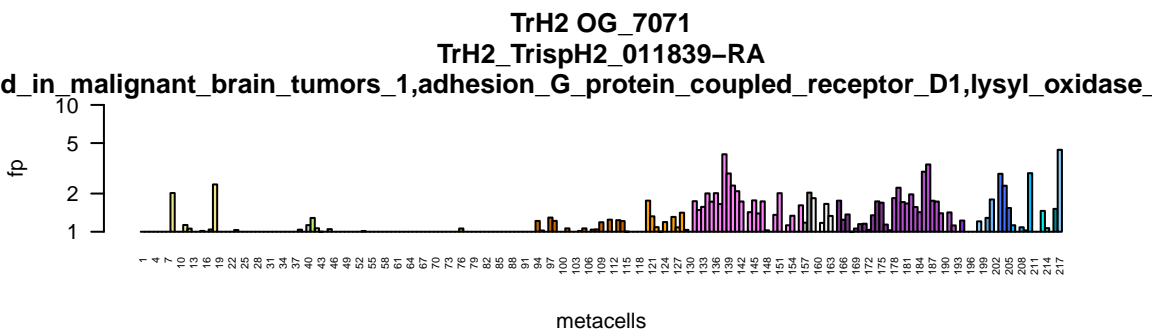
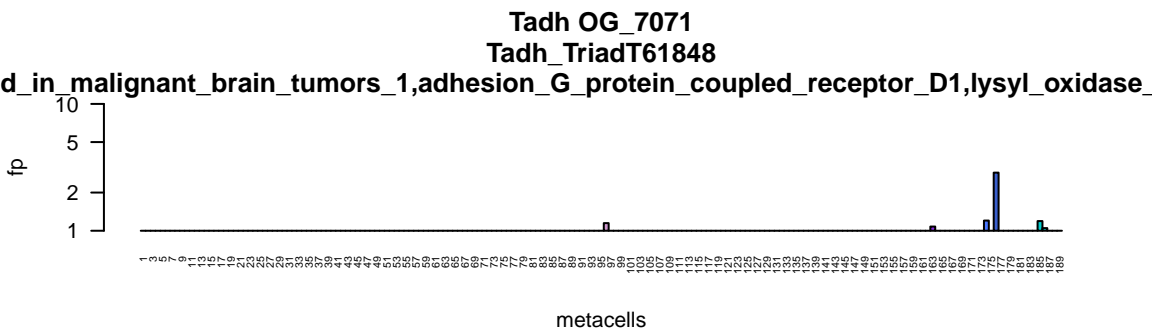








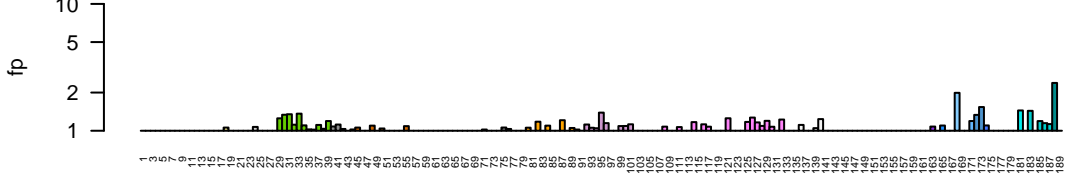




Tadh OG\_7573

Tadh\_TriadT26040

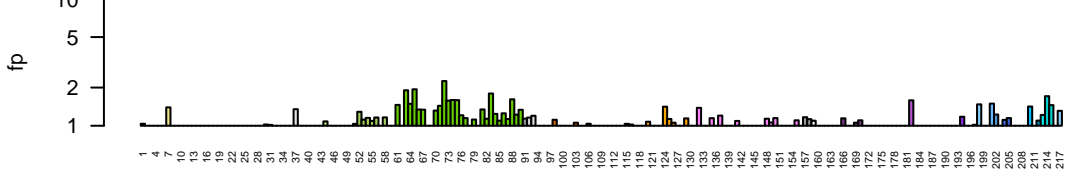
5\_hydroxytryptamine\_receptor\_1B,histamine\_receptor\_H2,opsin\_4,adrenoceptor\_alpha\_



TrH2 OG\_7573

TrH2\_TrispH2\_008979-RA

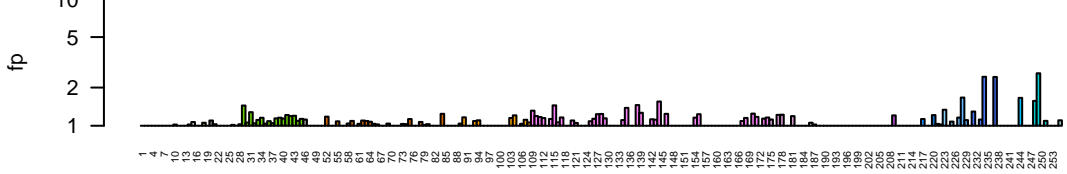
5\_hydroxytryptamine\_receptor\_1B,histamine\_receptor\_H2,opsin\_4,adrenoceptor\_alpha\_



Hhon OG\_7573

Hhon\_g05380.t1

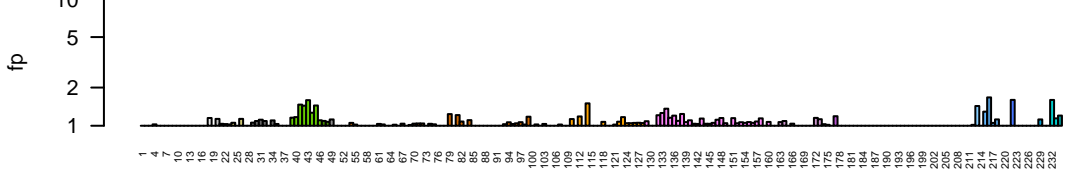
5\_hydroxytryptamine\_receptor\_1B,histamine\_receptor\_H2,opsin\_4,adrenoceptor\_alpha\_



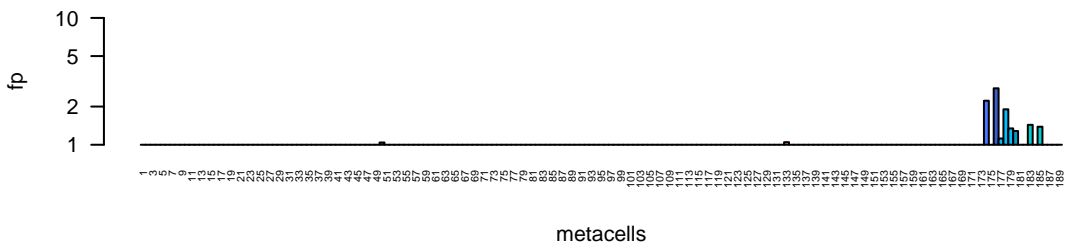
HoiH23 OG\_7573

HoiH23\_PIH23\_000307-RA

5\_hydroxytryptamine\_receptor\_1B,histamine\_receptor\_H2,opsin\_4,adrenoceptor\_alpha\_

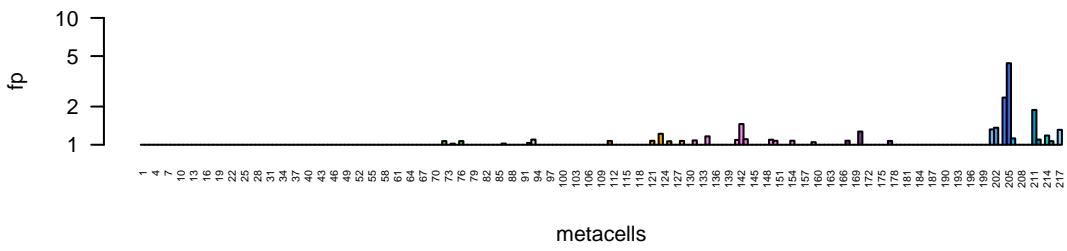


Tadh OG\_8094  
Tadh\_wf\_g6497.t1



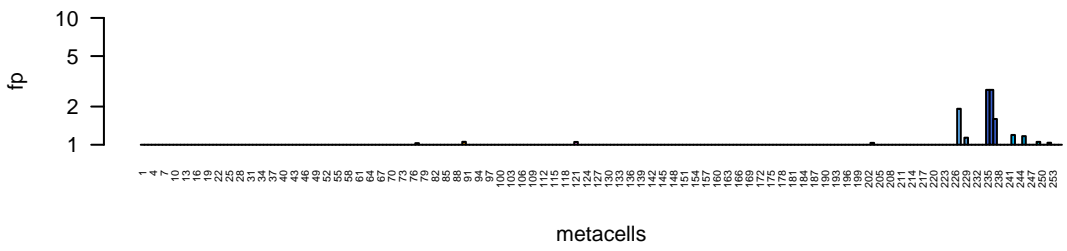
metacells

TrH2 OG\_8094  
TrH2\_TrispH2\_011737-RA



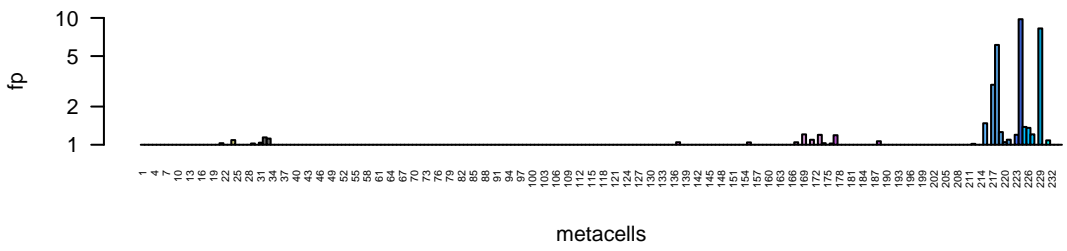
metacells

Hhon OG\_8094  
Hhon\_g08752.t1



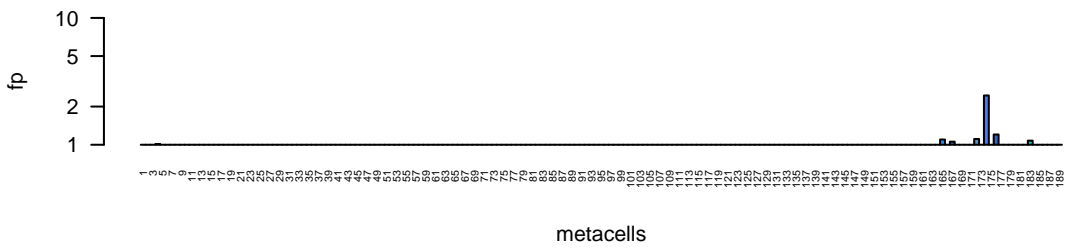
metacells

HoiH23 OG\_8094  
HoiH23\_PIH23\_007147-RA

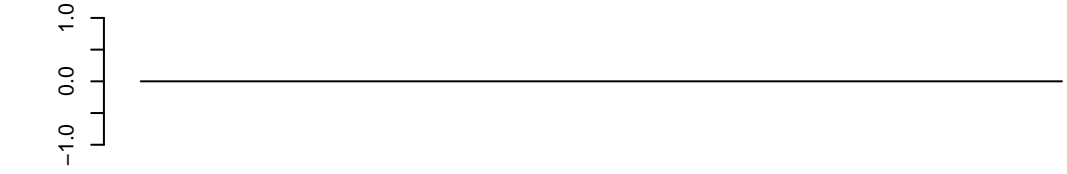


metacells

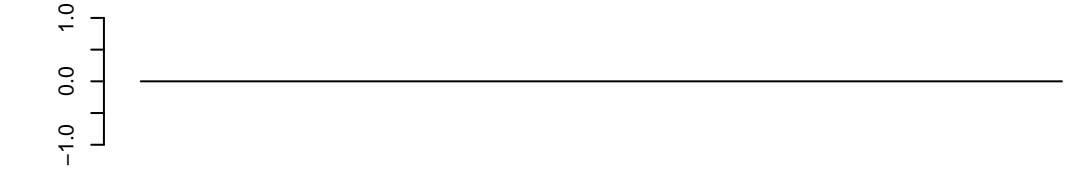
Tadh OG\_8095  
Tadh\_TriadT57627



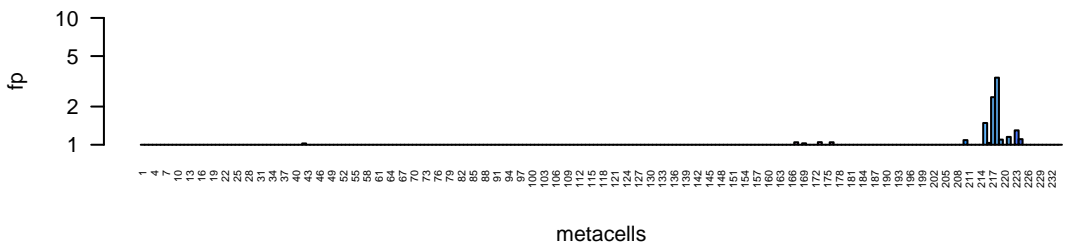
TrH2 | no data



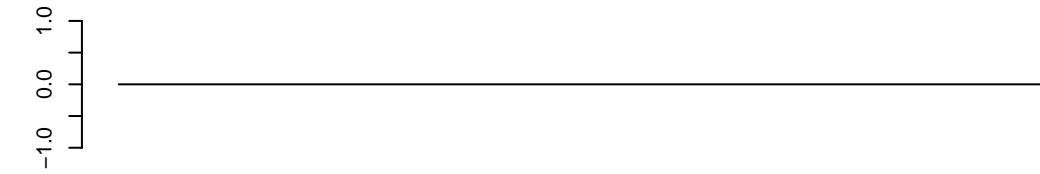
Hhon | no data



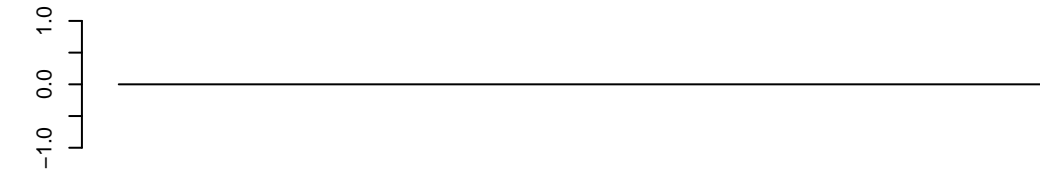
HoiH23 OG\_8095  
HoiH23\_PIH23\_007148-RA



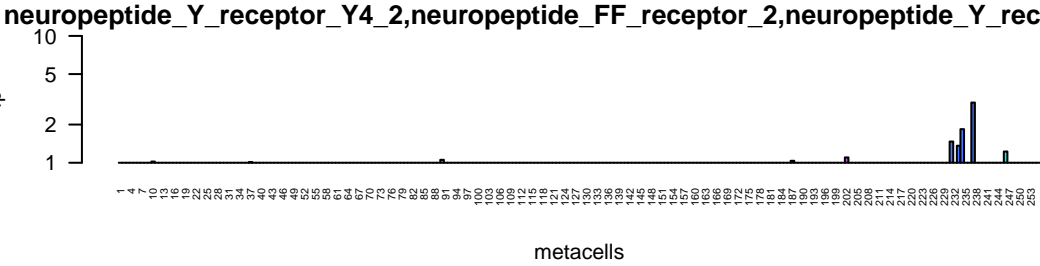
neuropeptide\_Y\_receptor\_Y4\_2,neuropeptide\_FF\_receptor\_2,neuropeptide\_Y\_receptor\_\  
Tadh | no data



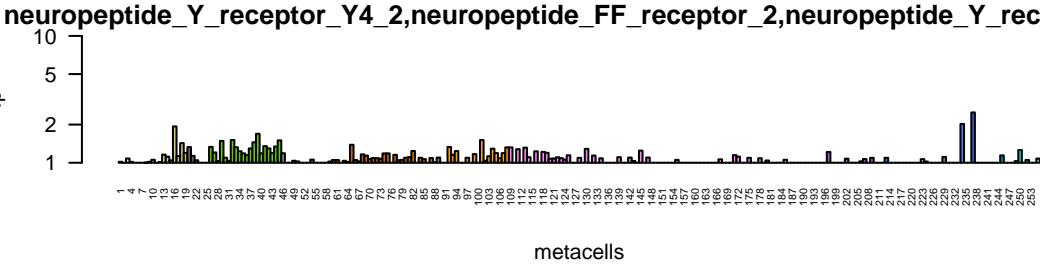
neuropeptide\_Y\_receptor\_Y4\_2,neuropeptide\_FF\_receptor\_2,neuropeptide\_Y\_receptor\_\  
TrH2 | no data



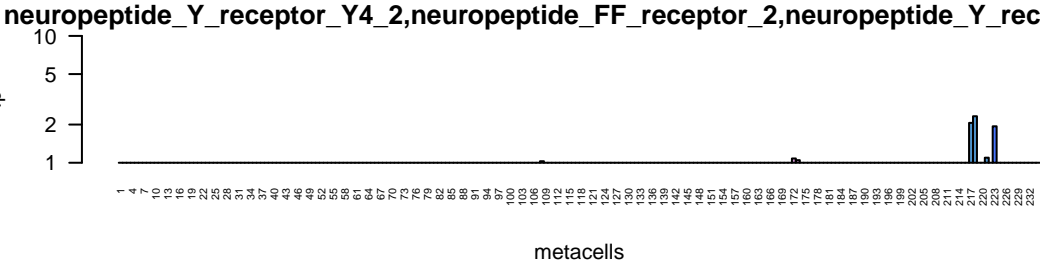
Hhon OG\_8289  
Hhon\_g07093.t1  
neuropeptide\_Y\_receptor\_Y4\_2,neuropeptide\_FF\_receptor\_2,neuropeptide\_Y\_receptor\_\  
metacells

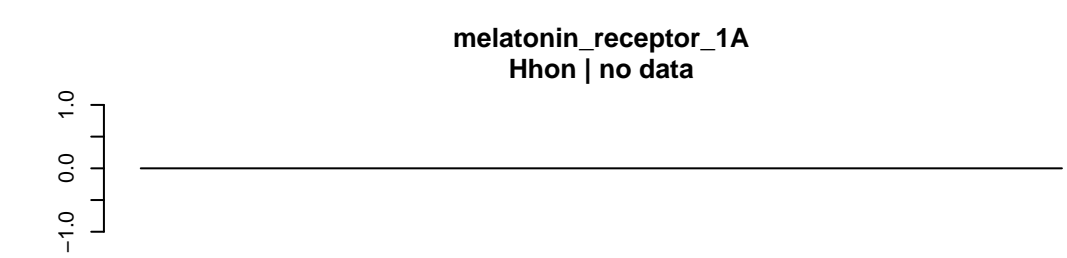
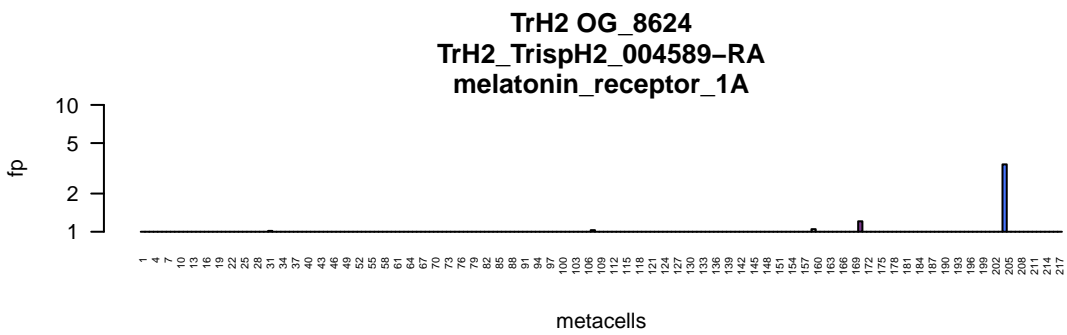
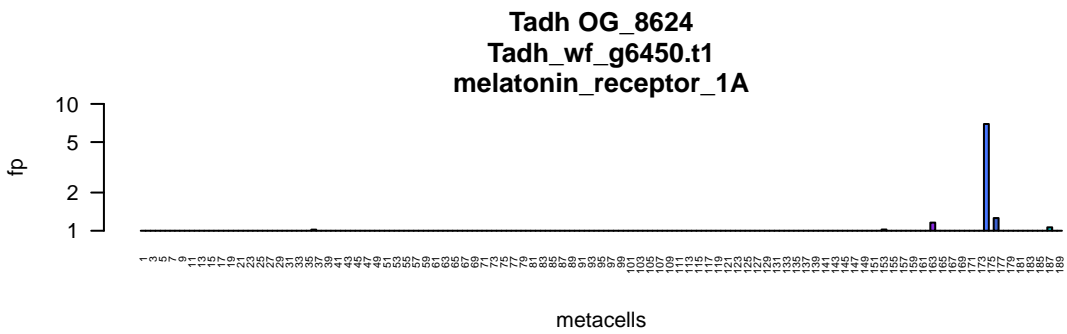


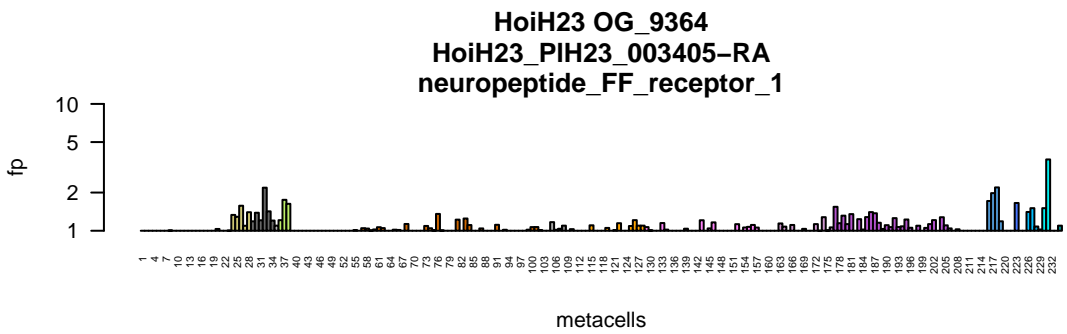
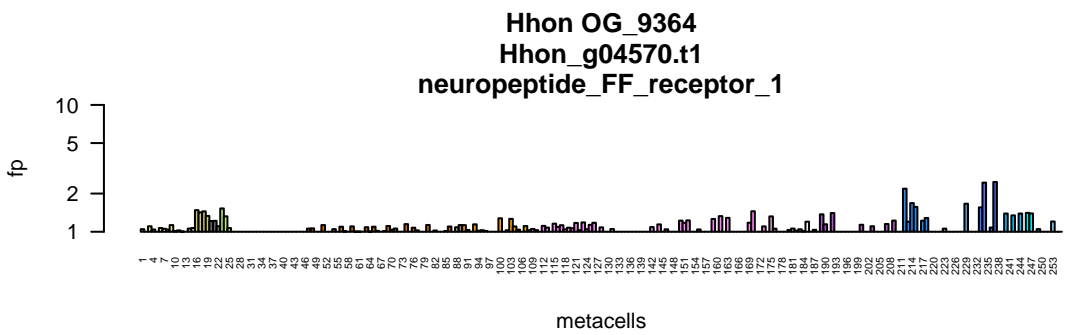
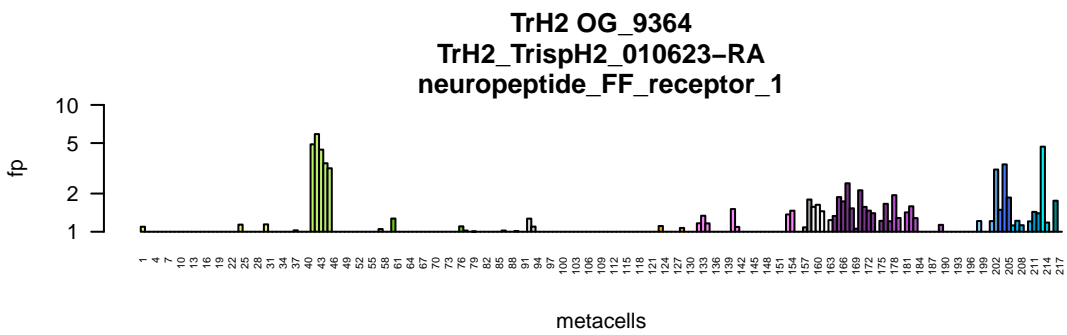
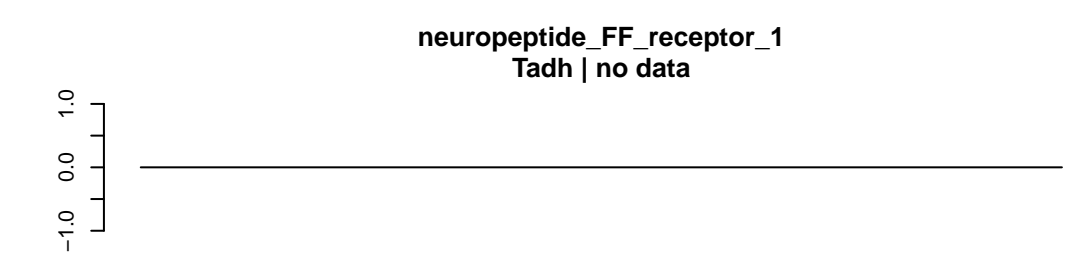
Hhon OG\_8289  
Hhon\_g11435.t1  
neuropeptide\_Y\_receptor\_Y4\_2,neuropeptide\_FF\_receptor\_2,neuropeptide\_Y\_receptor\_\  
metacells



HoiH23 OG\_8289  
HoiH23\_PIH23\_002693-RA  
neuropeptide\_Y\_receptor\_Y4\_2,neuropeptide\_FF\_receptor\_2,neuropeptide\_Y\_receptor\_\  
metacells



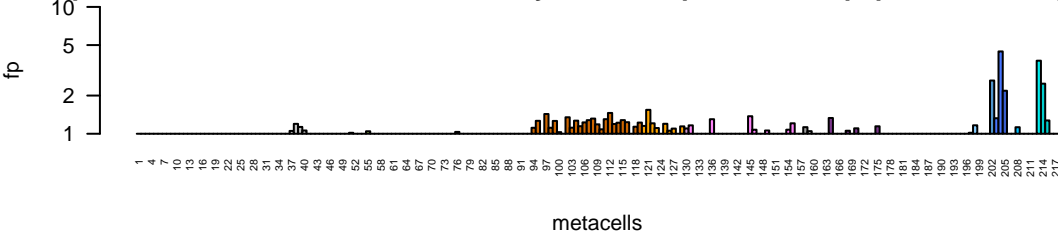




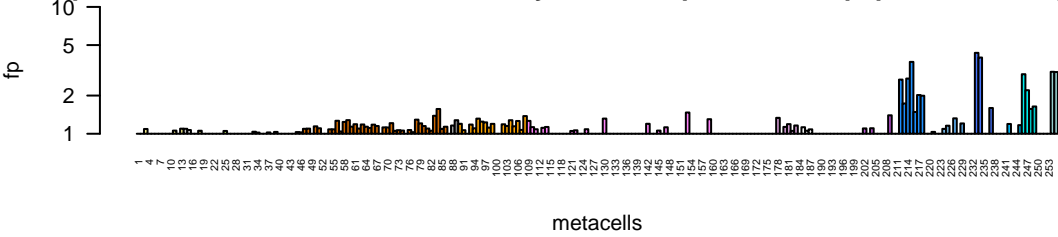
aprataxin\_and\_PNKP\_like\_factor,tachykinin\_receptor\_2,neuropeptide\_FF\_receptor\_1  
Tadh | no data



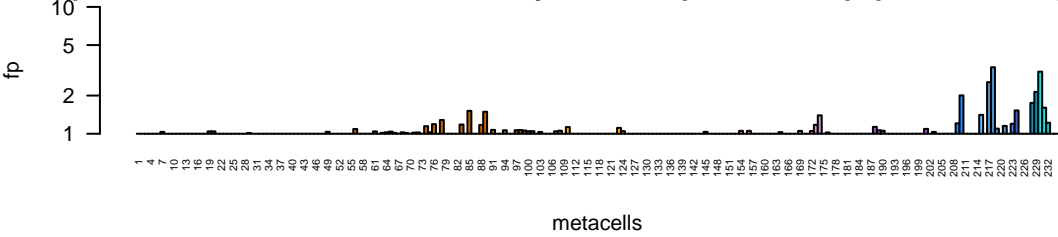
TrH2 OG\_9368  
TrH2\_TrispH2\_010619-RA  
aprataxin\_and\_PNKP\_like\_factor,tachykinin\_receptor\_2,neuropeptide\_FF\_receptor\_1



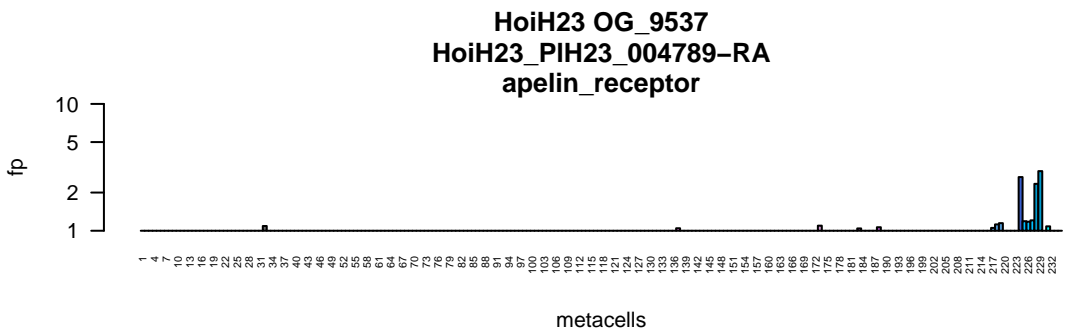
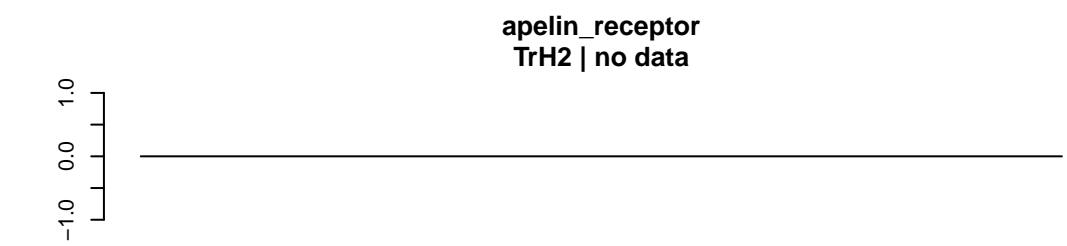
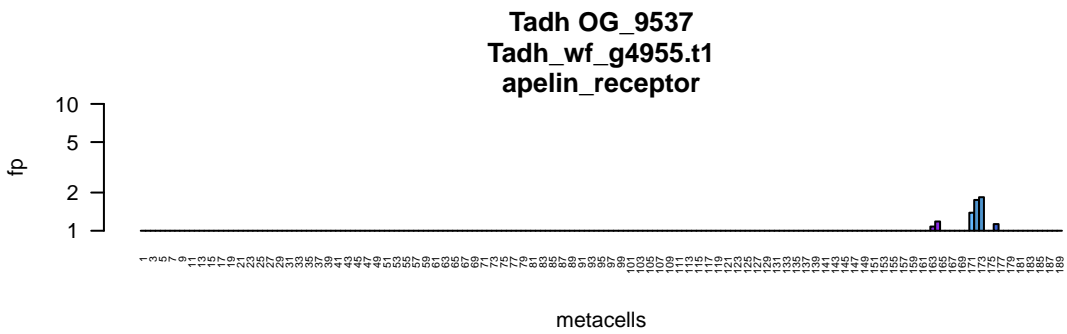
Hhon OG\_9368  
Hhon\_g04575.t1  
aprataxin\_and\_PNKP\_like\_factor,tachykinin\_receptor\_2,neuropeptide\_FF\_receptor\_1



HoiH23 OG\_9368  
HoiH23\_PIH23\_003409-RA  
aprataxin\_and\_PNKP\_like\_factor,tachykinin\_receptor\_2,neuropeptide\_FF\_receptor\_1



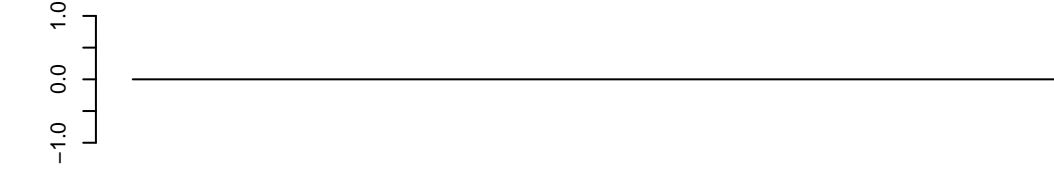




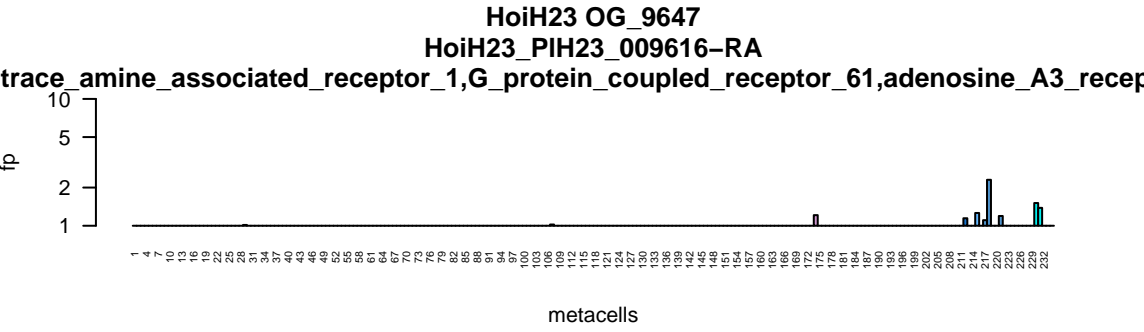
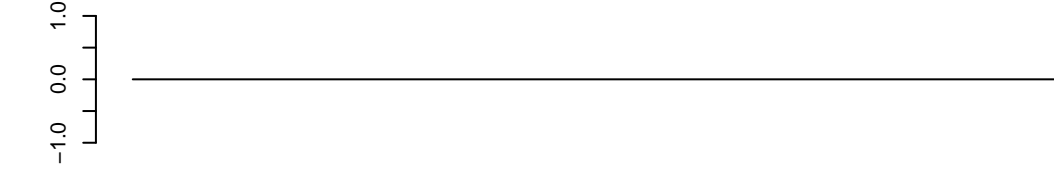
trace\_amine\_associated\_receptor\_1,G\_protein\_coupled\_receptor\_61,adenosine\_A3\_receptor  
Tadh | no data



trace\_amine\_associated\_receptor\_1,G\_protein\_coupled\_receptor\_61,adenosine\_A3\_receptor  
TrH2 | no data



trace\_amine\_associated\_receptor\_1,G\_protein\_coupled\_receptor\_61,adenosine\_A3\_receptor  
Hhon | no data



Tadh | no data

TrH2 | no data

Hhon OG\_10393  
Hhon\_g01936.t1

HoiH23 | no data

