

Two bar charts showing the frequency of metacells for different cell types across two conditions: TadH OG_4214 and TrH2 OG_4214. The y-axis is 'fp' (frequency) on a log scale from 1 to 10. The x-axis is 'metacells' with indices 1 to 217. The left chart is for TadH OG_4214 and the right chart is for TrH2 OG_4214. Both charts show a similar distribution of metacells, with a peak around metacell 173.

Tadh OG_4214
Tadh_TriadT30372
 subtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

TrH2 OG_4214
TrH2_TripH2_010333-RA
 subtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

fp

metacells

TadH OG_4214

TrH2 OG_4214

metacells

Tad

Tadh OG_4214
Tadh_TriadT60080

Hh

Hhon OG_4214
Hhon_g06801.t1

ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

fp

metacells

TadH OG_4214
TadH_TriadT60077
 subtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

Hhon OG_4214
Hhon_g10731.t1
 subtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

fp

metacells

Tadh OG_4214

Hhon OG_4214

fp

10

5

2

1

1 3 7 11 13 17 21 23 27 31 33 37 41 43 47 51 53 55 59 61 63 65 67 71 73 75 79 81 83 85 87 89 91 93 95 97 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189

metacells

Hhon OG_4214

Hhon_g10727.t1

fp

10

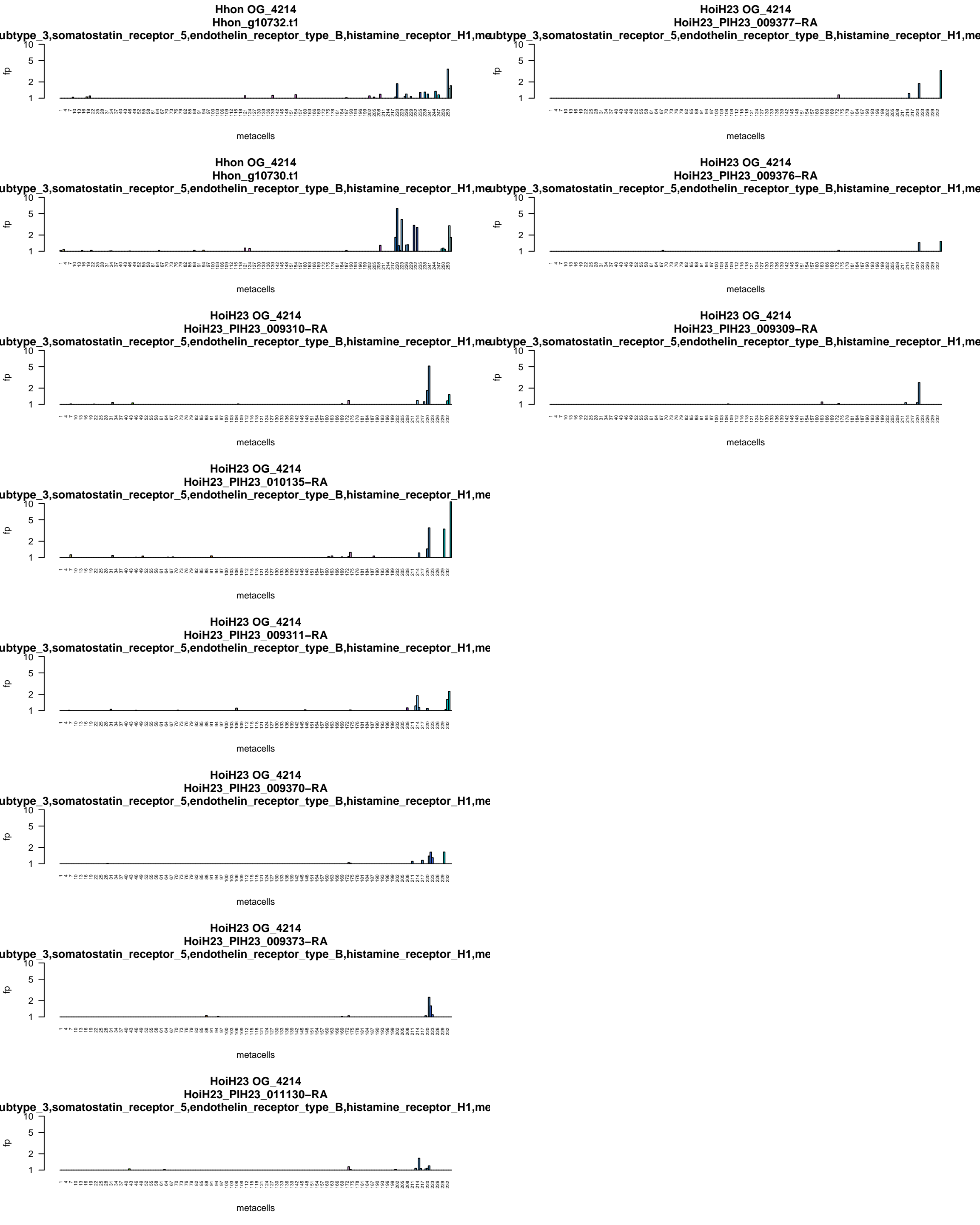
5

2

1

1 4 10 13 19 22 28 31 37 43 49 52 55 59 61 63 65 67 69 71 73 75 77 79 81 83 85 87 89 91 93 95 97 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189

metacells



HoiH23 OG_4214

HoiH23_PIH23_010135-RA

ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

HoiH23 OG_4214

HoiH23_PIH23_009311-RA

ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

HoiH23 OG_4214

HoiH23_PIH23_009370-RA

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HoiH23 OG_4214

HoiH23_PIH23_009373-RA

ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

HoiH23 OG_4214

HoiH23_PIH23_011130-RA

ubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,meubtype_3,somatostatin_receptor_5,endothelin_receptor_type_B,histamine_receptor_H1,me

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

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

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

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
158	0
161	0
164	0
167	0
170	0
173	0
176	0
179	0
182	0
185	0
188	0
191	0
194	0
197	0
200	1
201	1
202	2
203	2
205	0
208	0
211	0
214	1
217	0

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacell	fp
1	1
4	1
13	1
16	1.2
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	1.2
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	1.2
157	1
160	1
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	2.0
202	2.2
205	1.5
208	1.5
211	1.2
214	1.2
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.

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

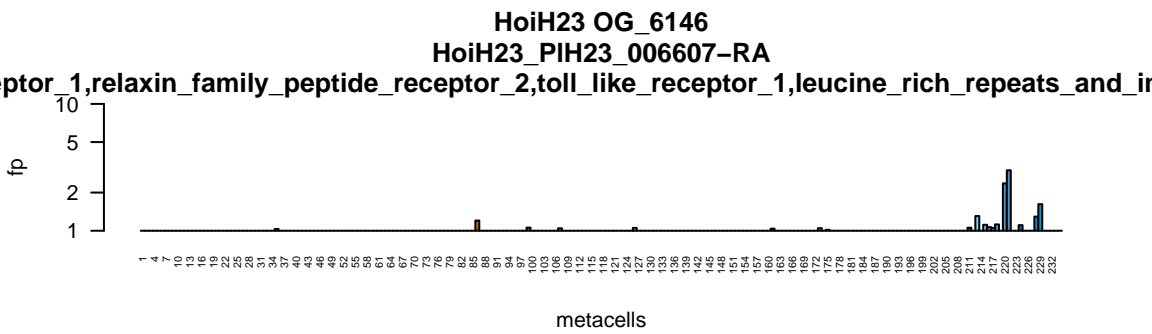
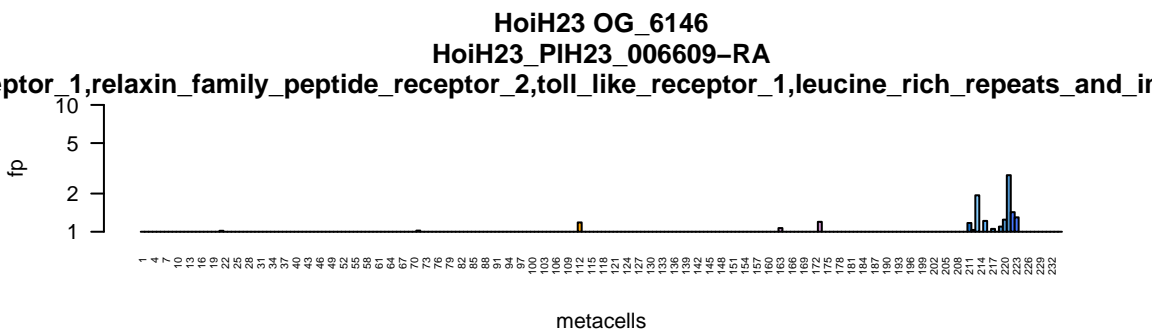
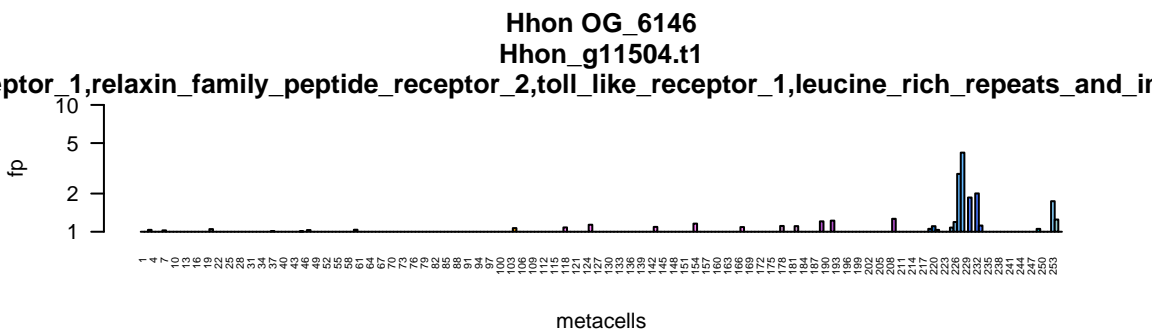
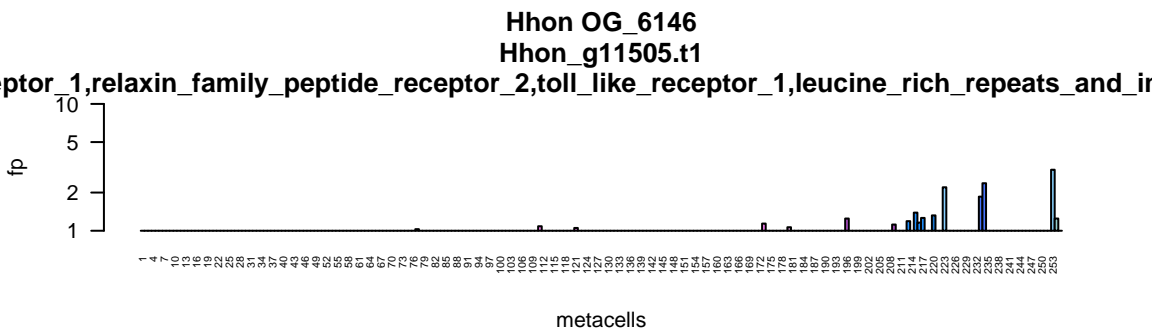
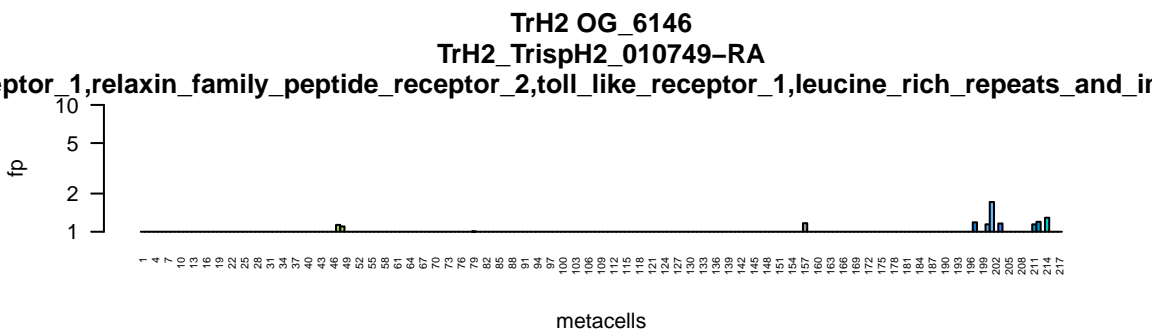
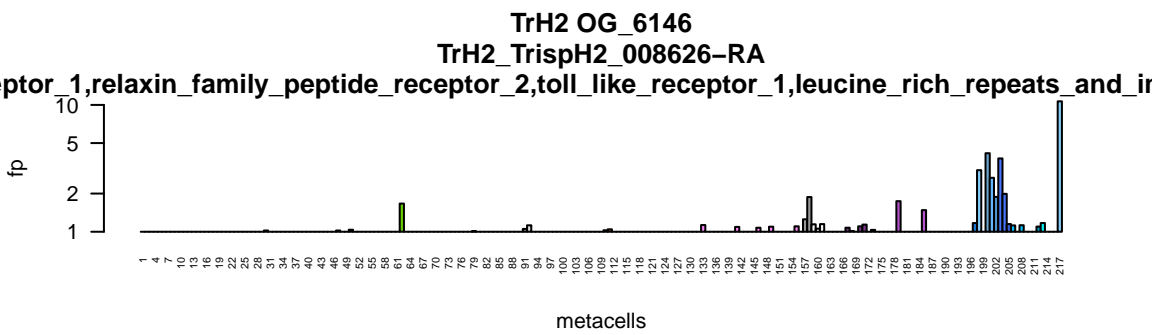
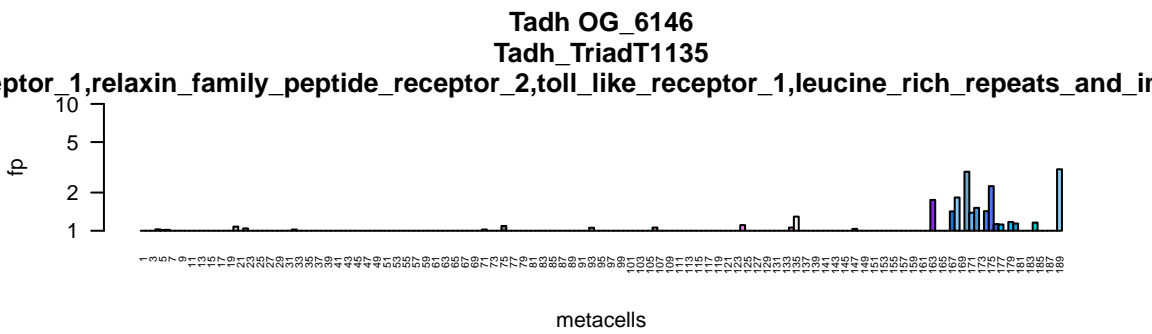
A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 253. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a few outliers reaching up to 2.

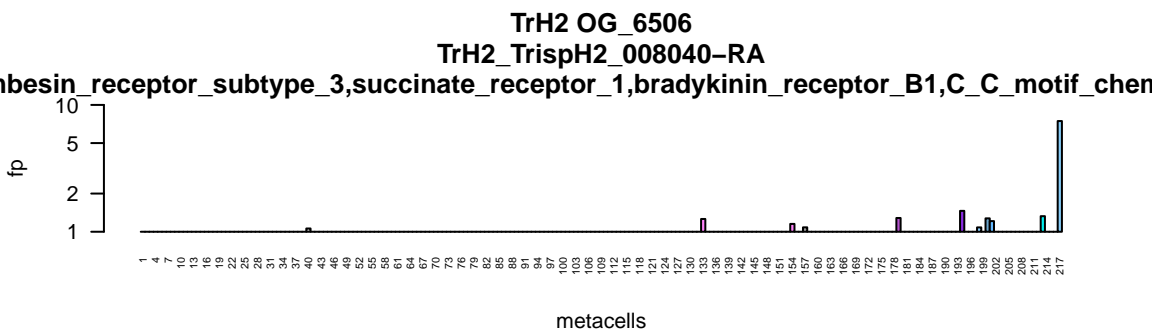
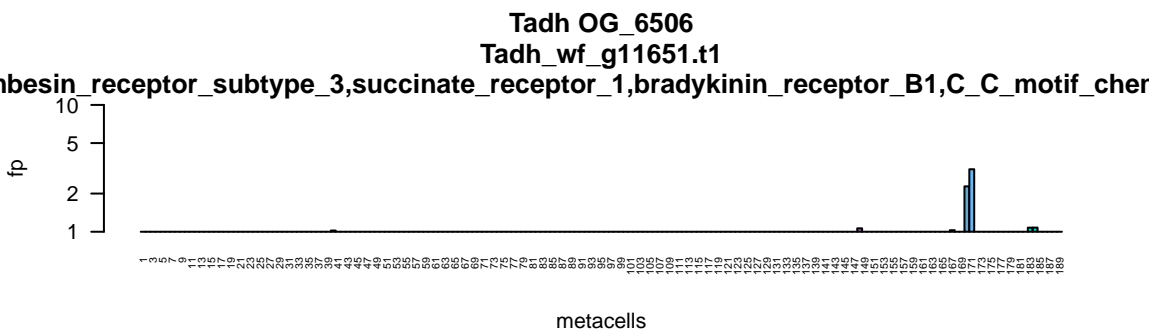
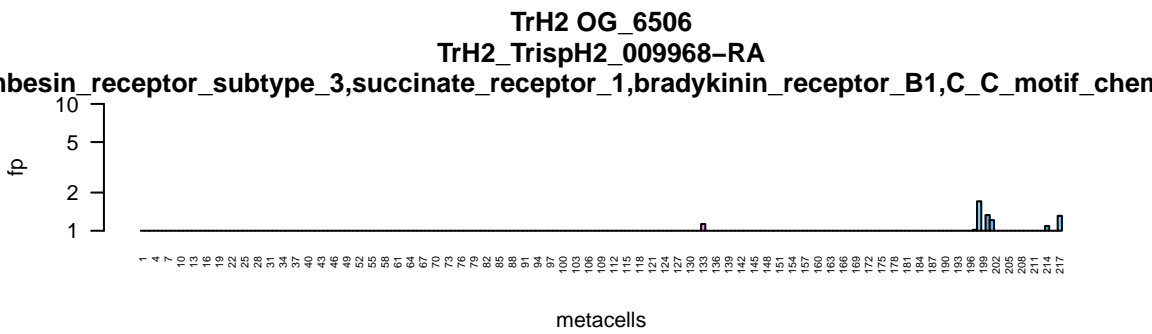
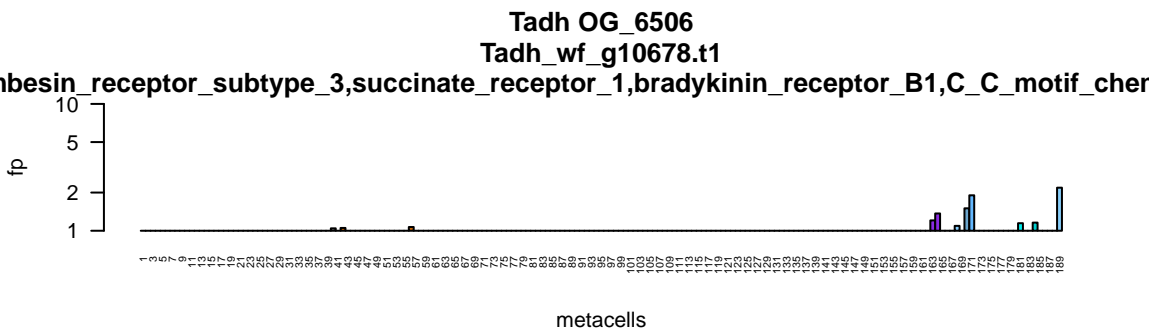
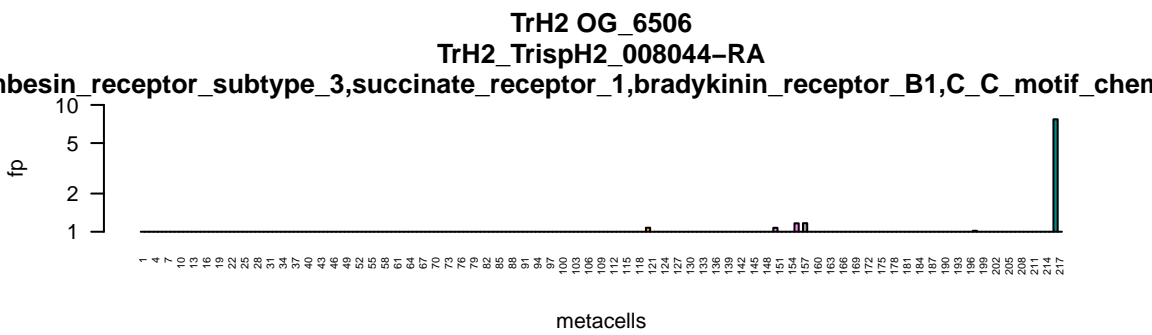
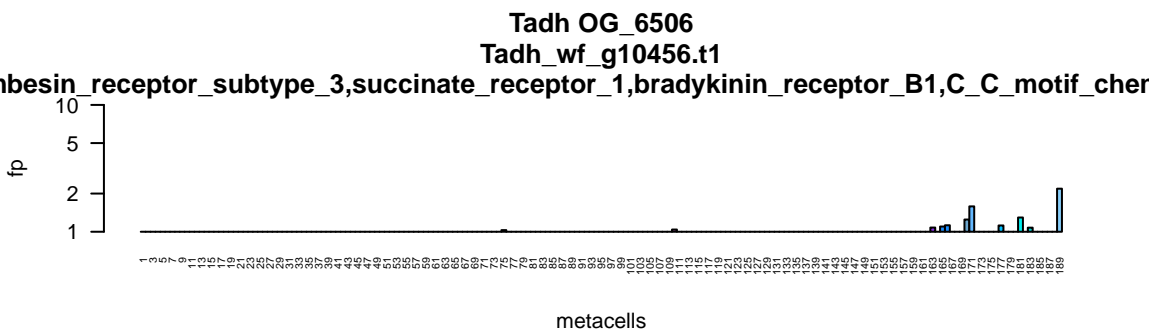
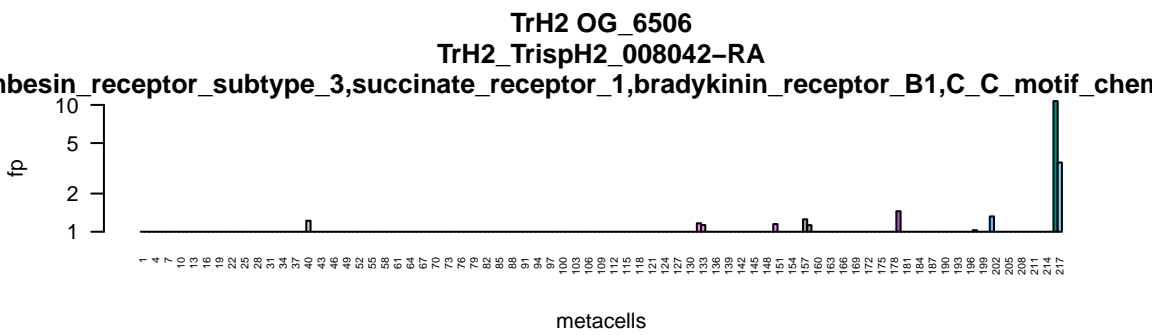
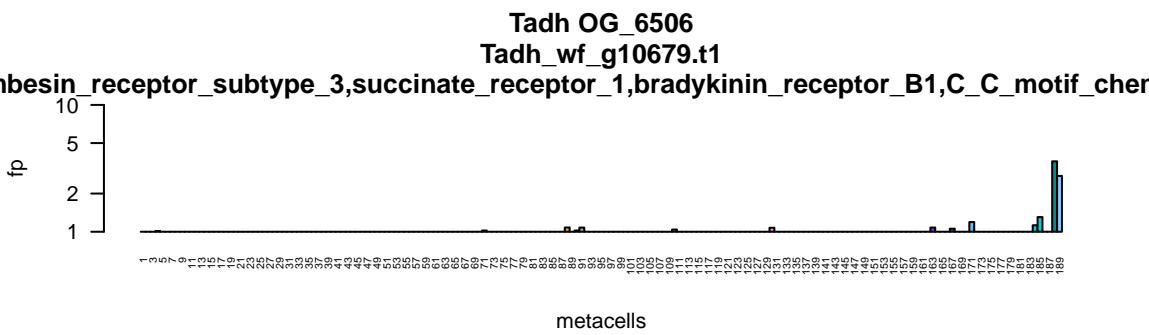
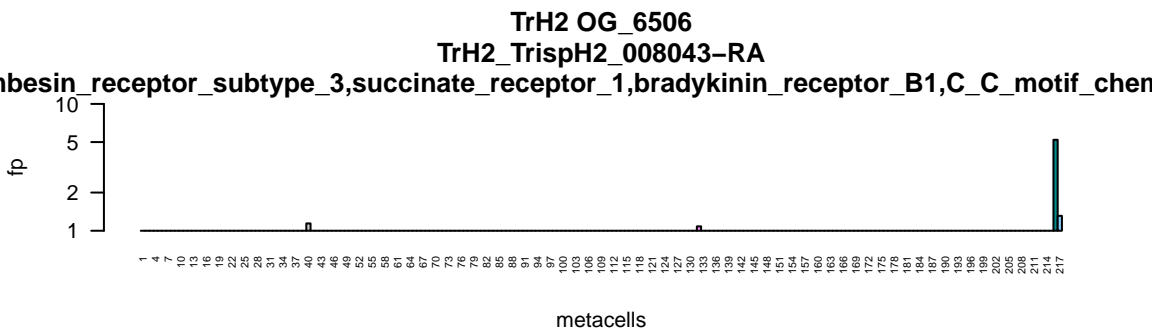
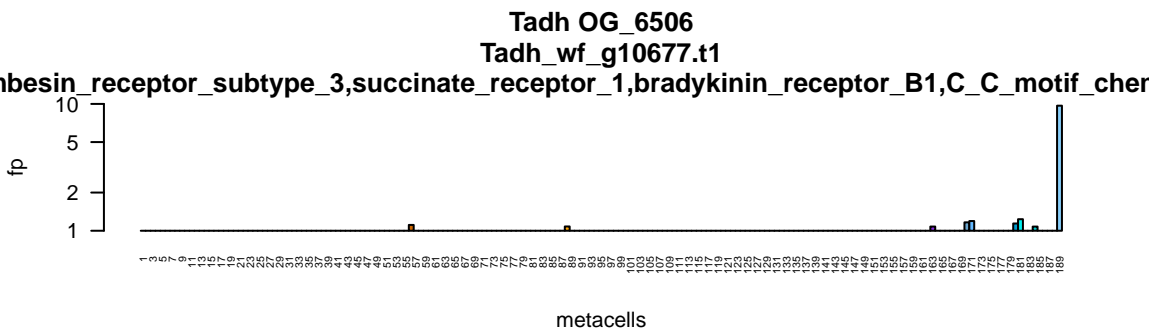
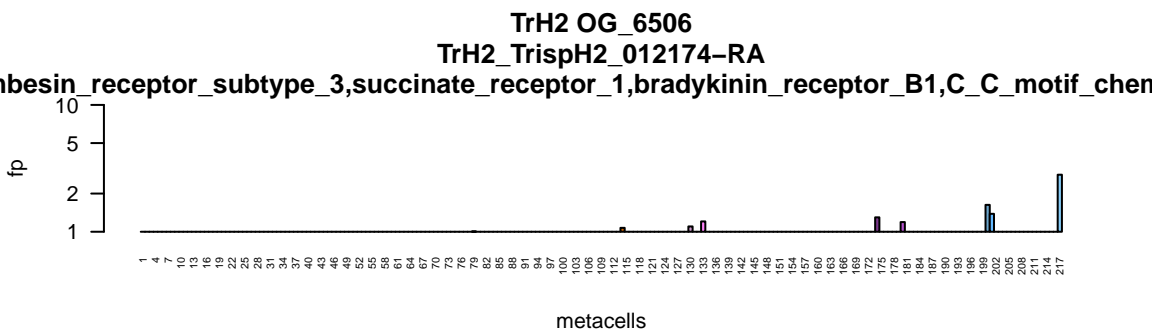
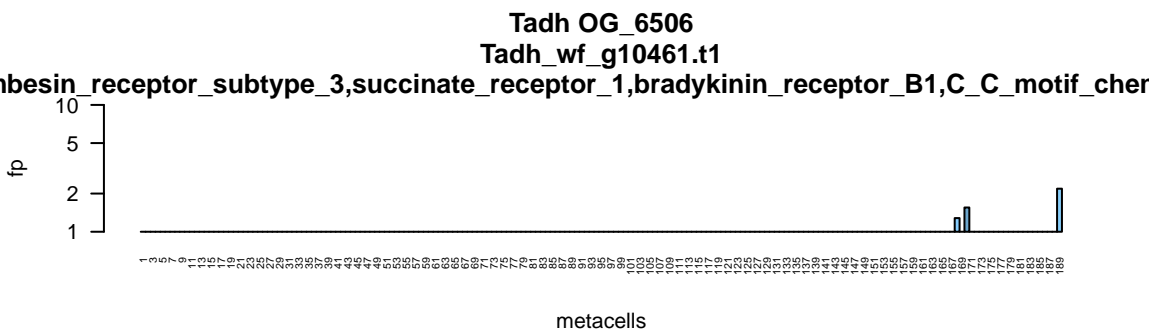
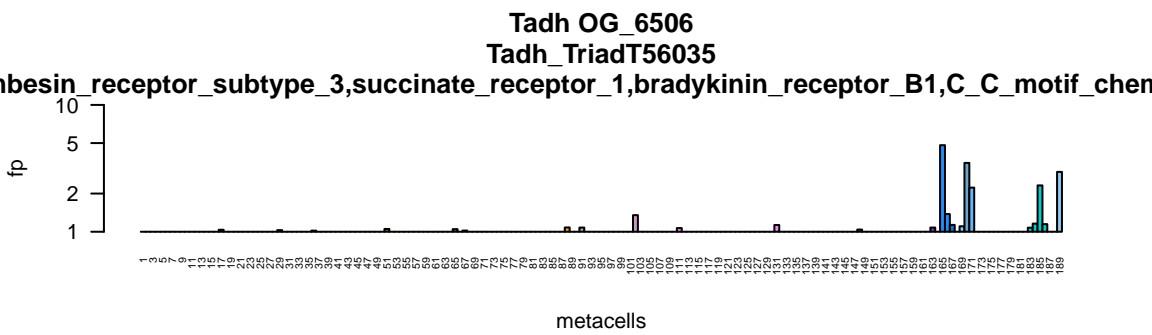
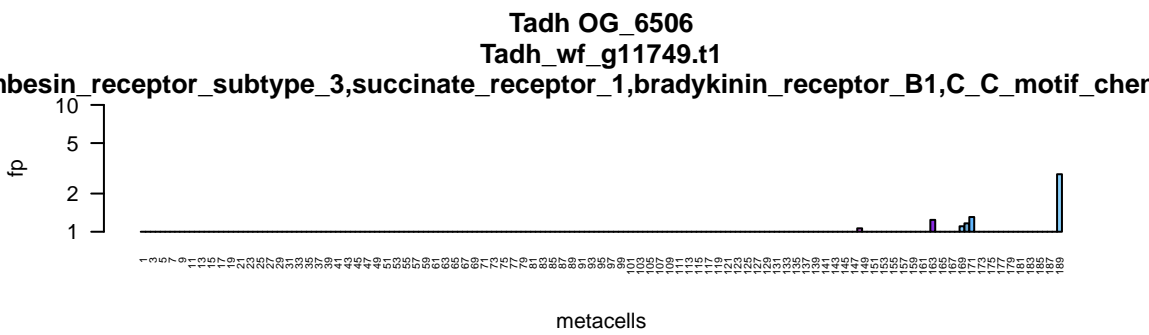
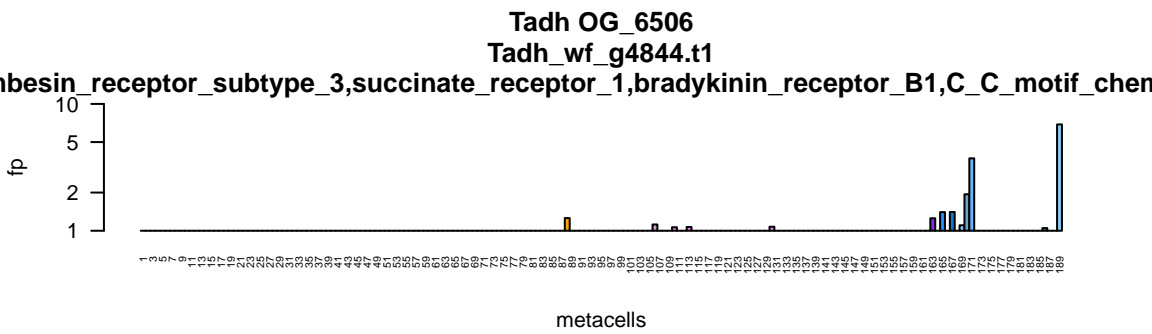
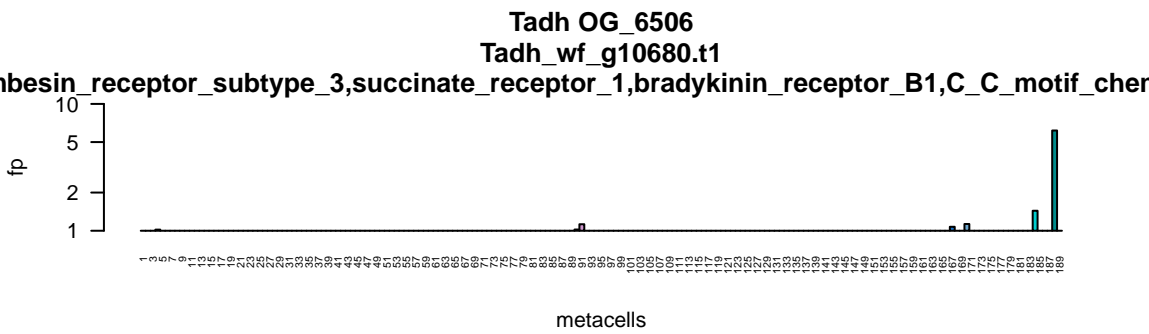
metacells	fp
1	1
4	1
10	1
13	1
19	1
22	1
28	1
31	1
37	1
43	1
46	1
55	1
61	1
64	1
70	1
73	1
79	1
88	1
91	1
97	1
100	1
109	1
115	1
119	1
127	1
133	1
136	1
146	1
151	1
154	1
160	1
163	1
169	1
175	1
184	1
187	1
193	1
196	1
202	1
205	1
211	1
214	1
220	1
223	1
229	1
232	1
238	1
241	1
247	1
250	1
253	2

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

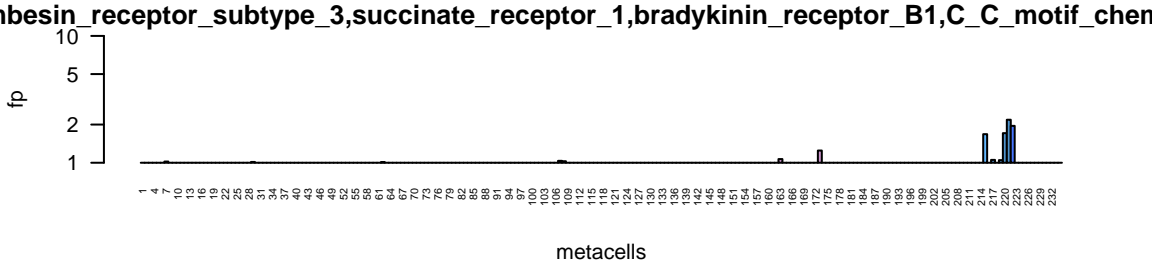
A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 232. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a few outliers reaching up to 4.

metacells	fp
1	1
4	1
10	1
13	1
16	1
22	1
25	1
31	1
34	1
37	1
43	1
46	1
55	1
68	1
69	1
73	1
76	1
82	1
85	4
88	1
94	1
97	1
103	1
106	1
109	1
115	1
118	1
124	1
127	1
130	1
136	1
139	1
142	1
145	1
148	1
151	1
157	1
160	1
163	1
166	1
169	1
172	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
220	4
223	1
229	1
232	1

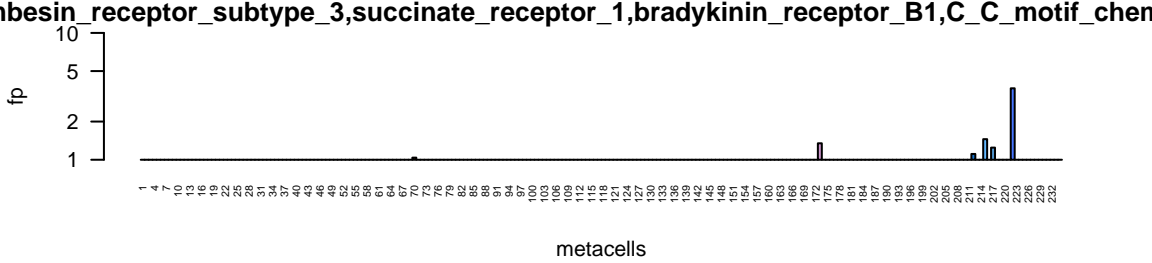


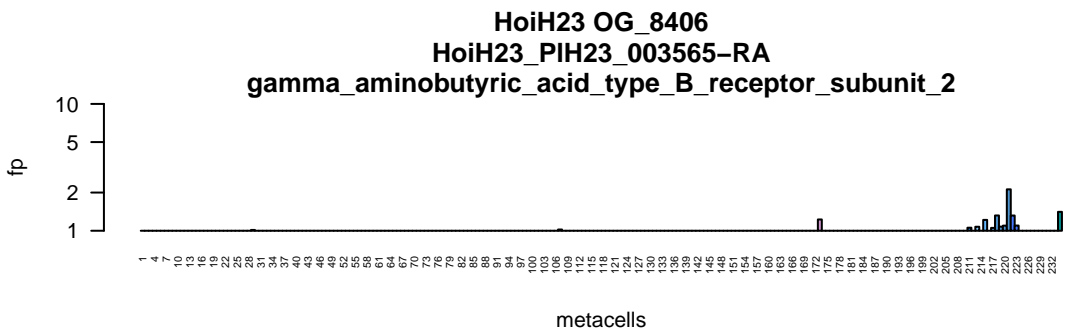
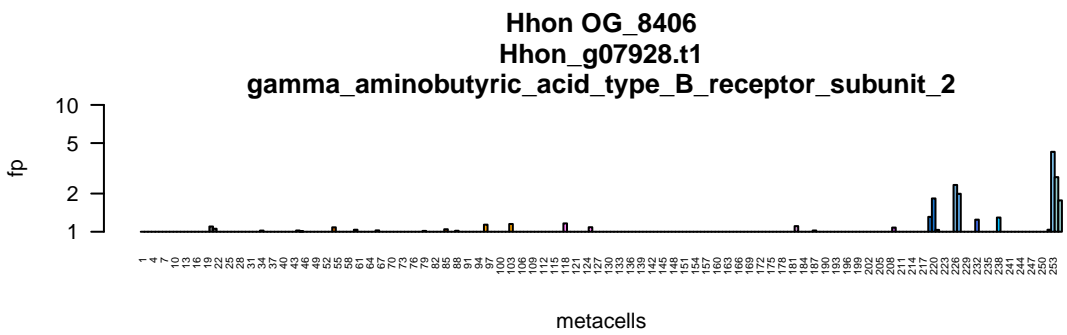
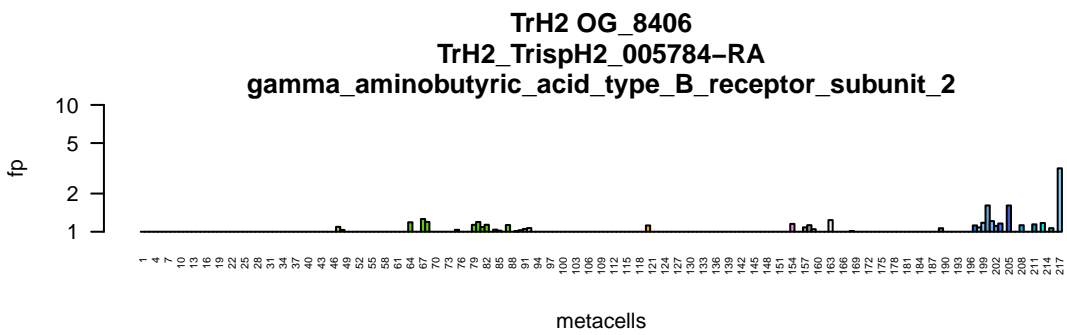
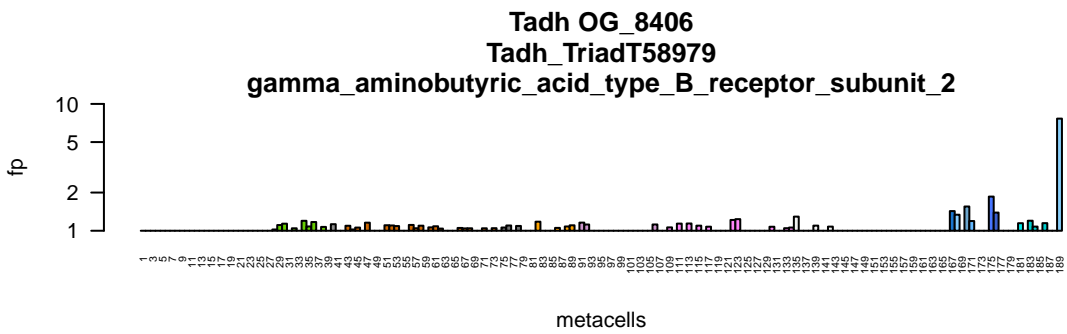


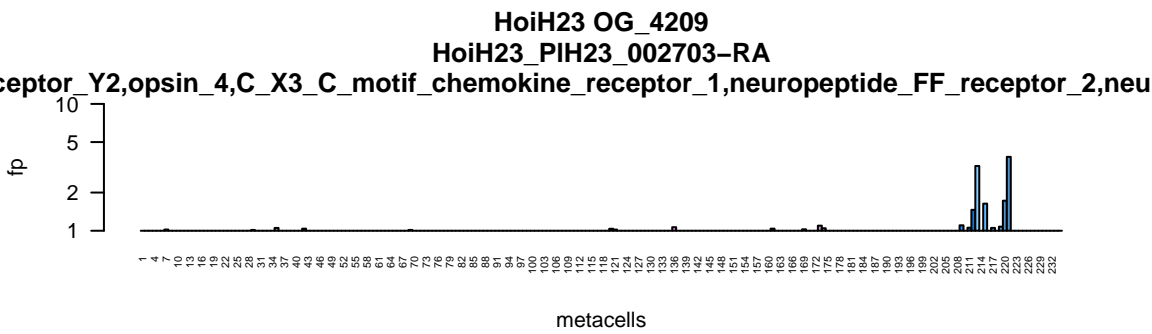
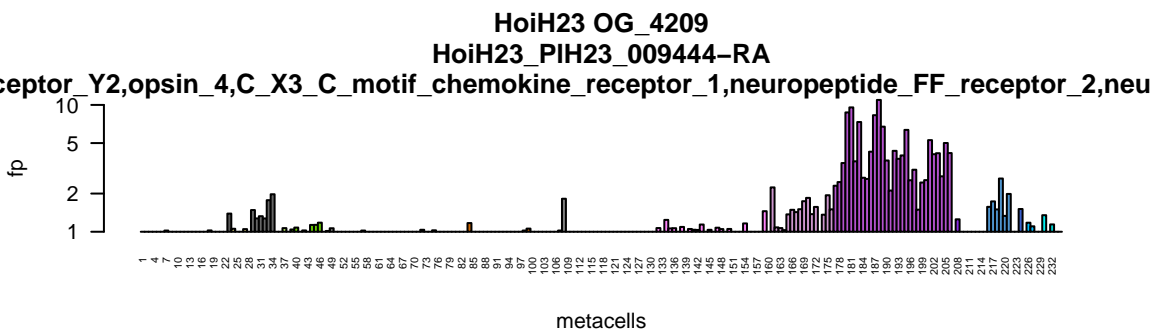
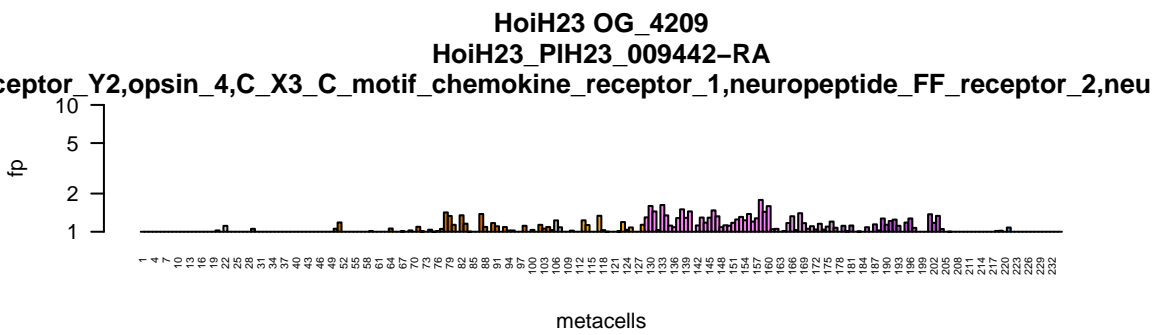
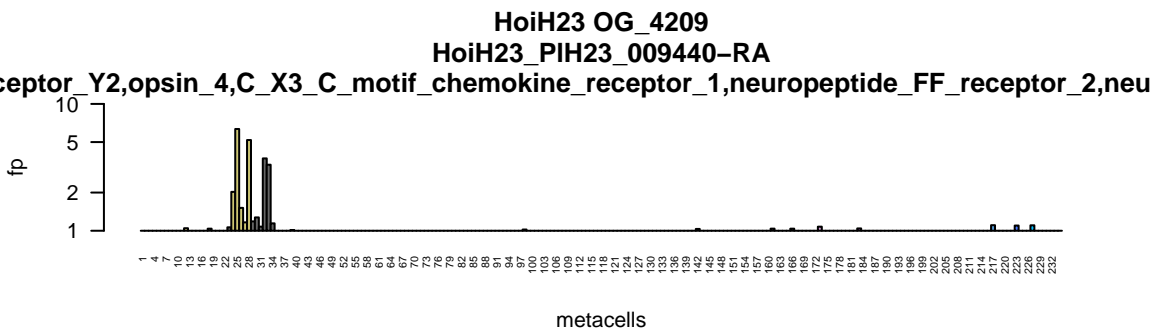
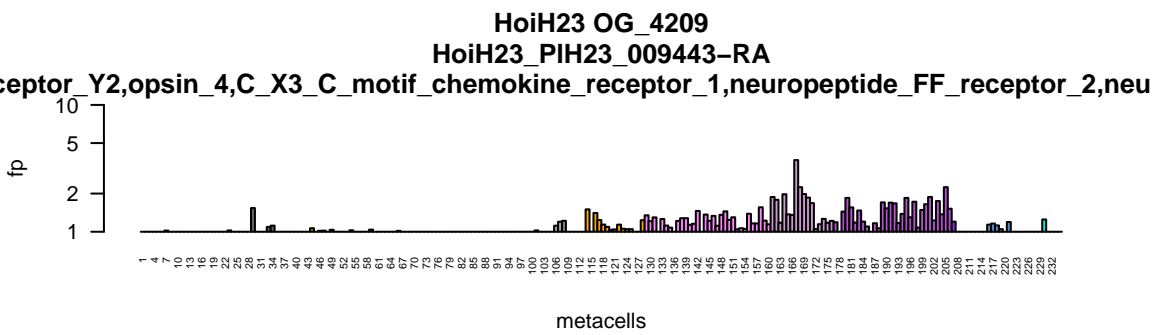
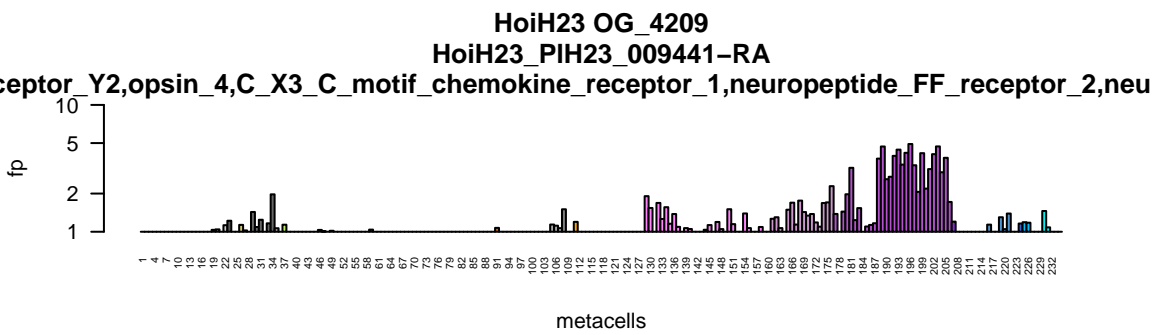
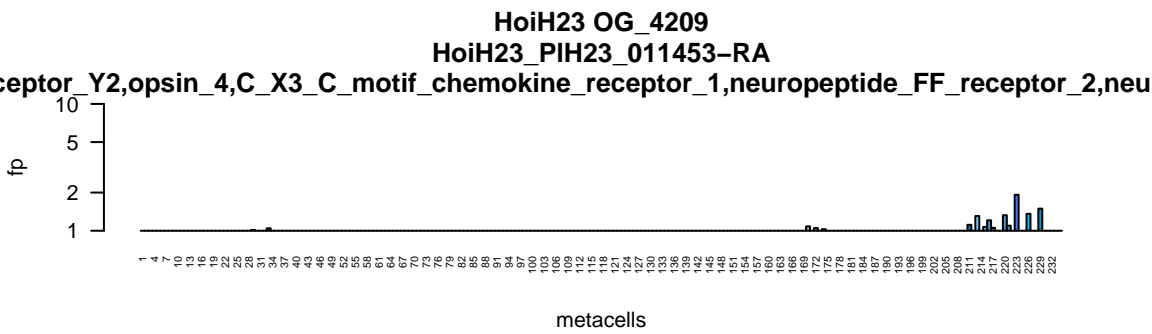
HoiH23 OG_6506
HoiH23_PIH23_008211-RA

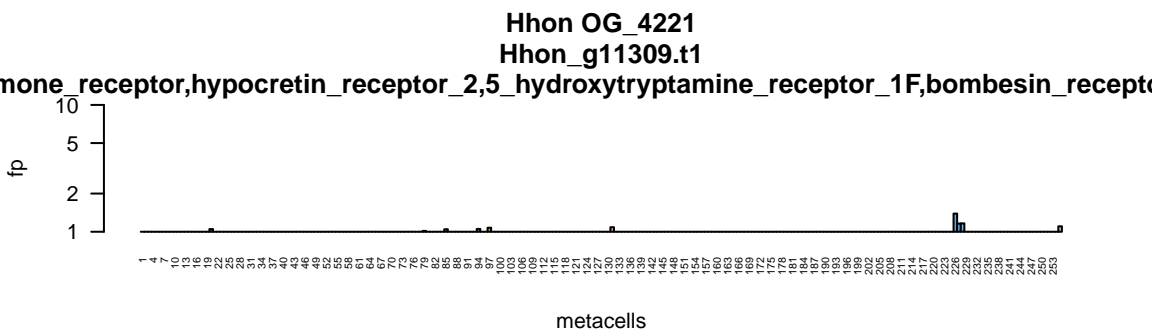
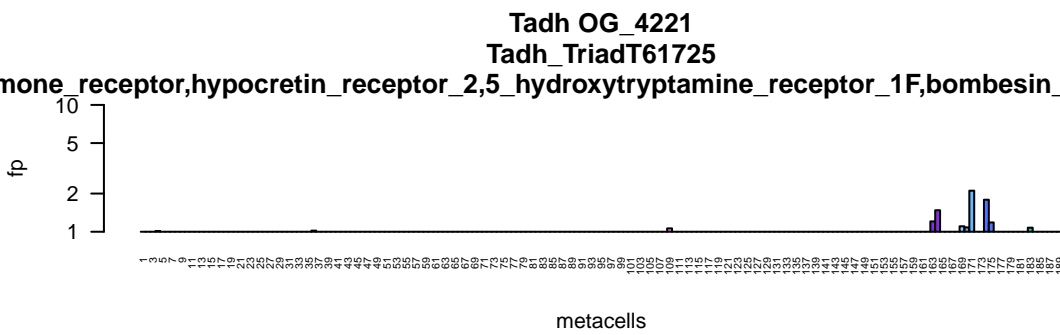
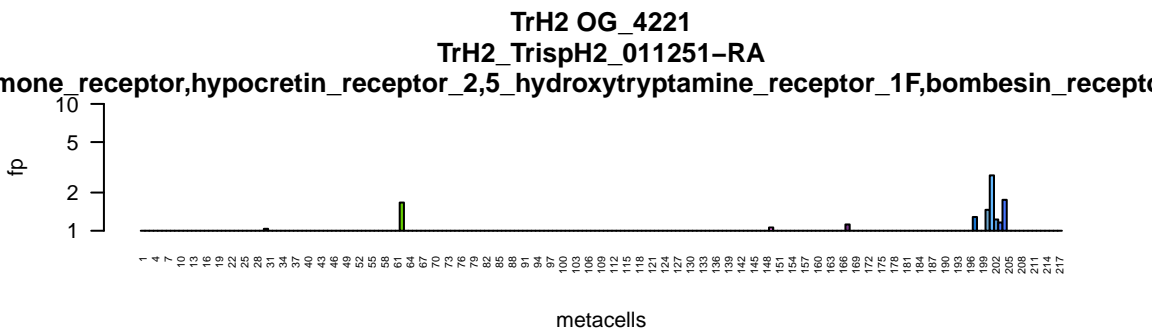
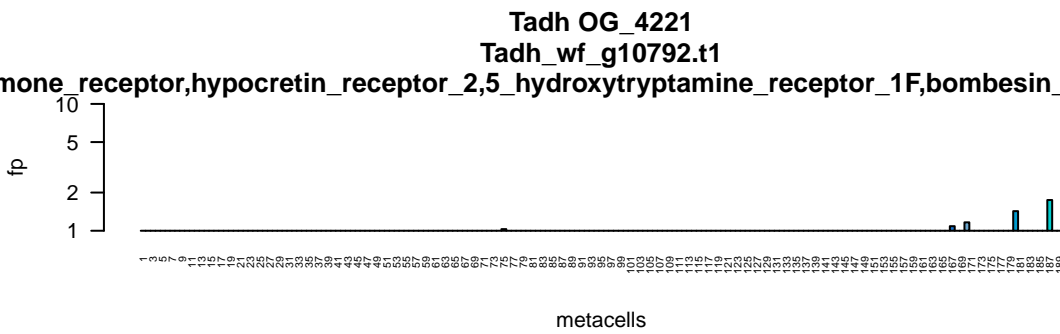
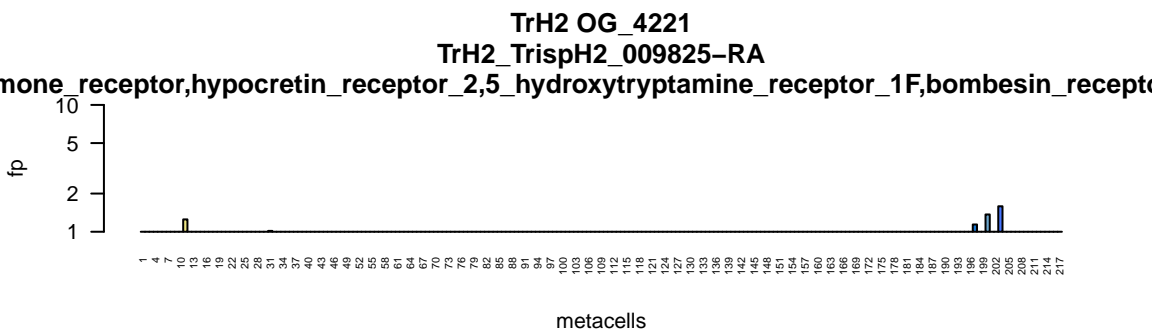
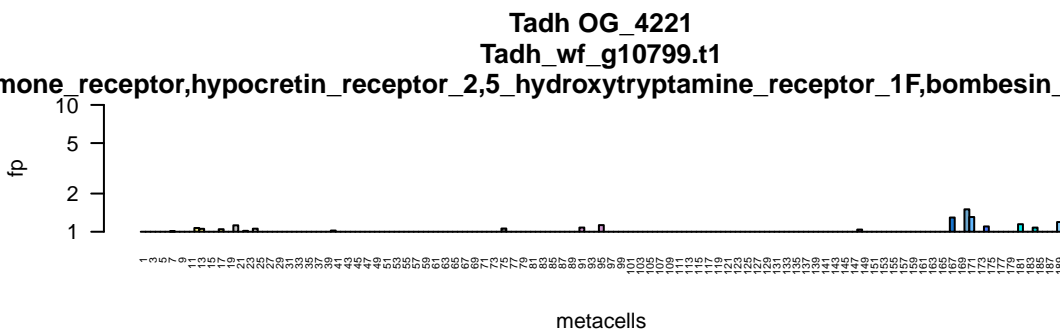
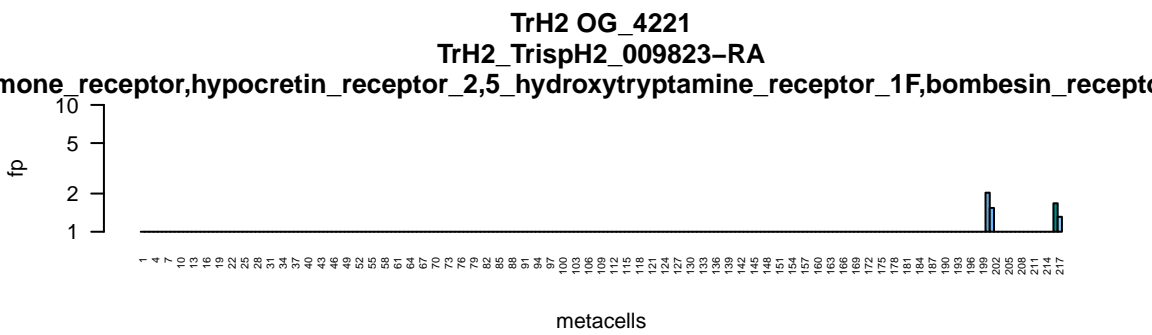
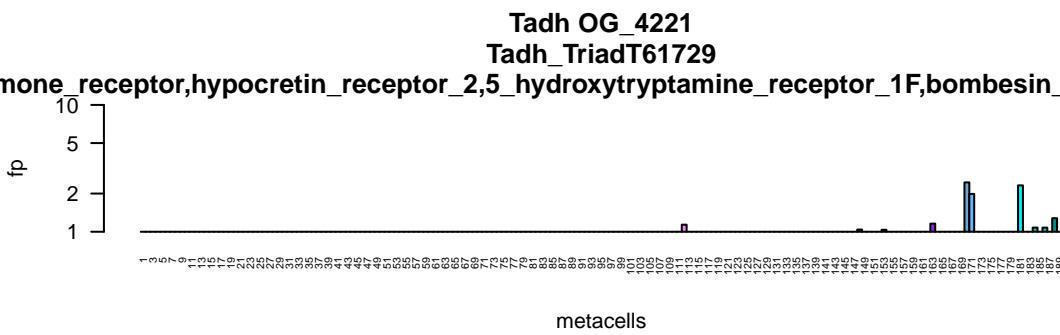
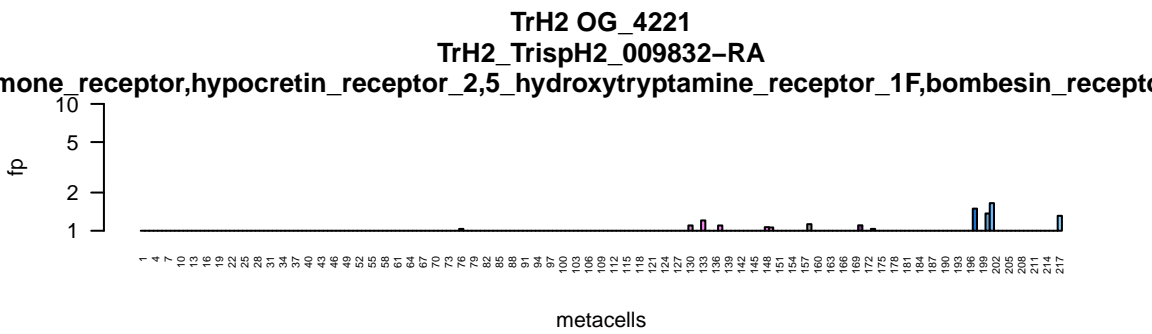
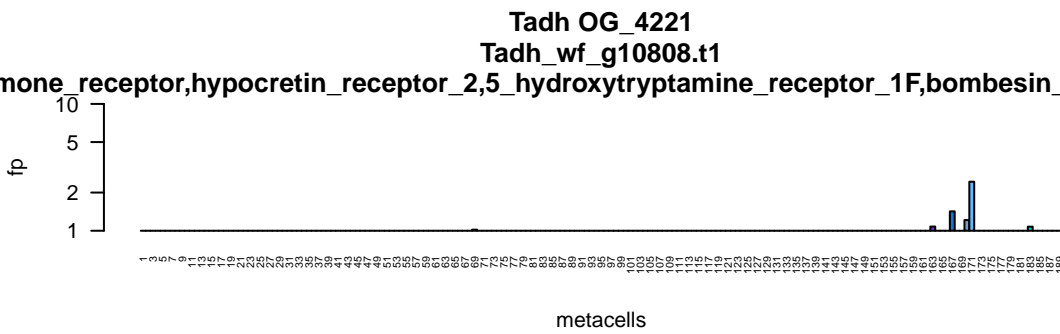
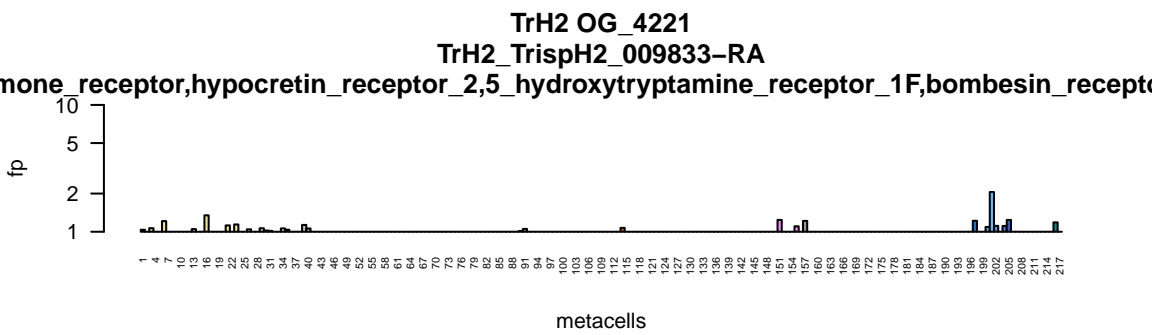
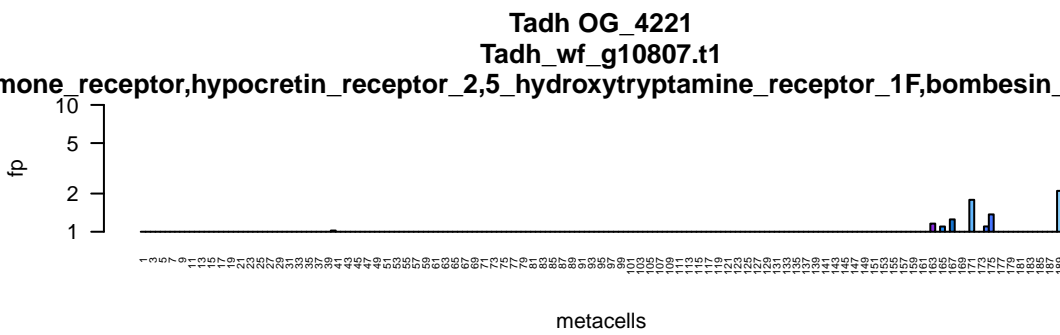
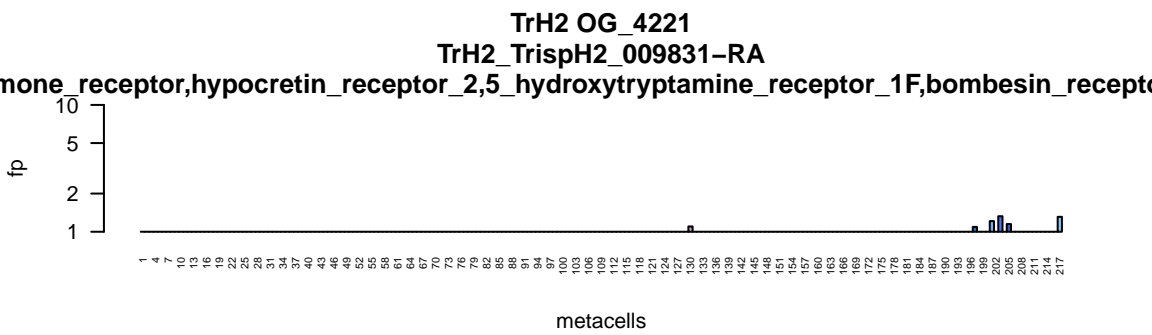
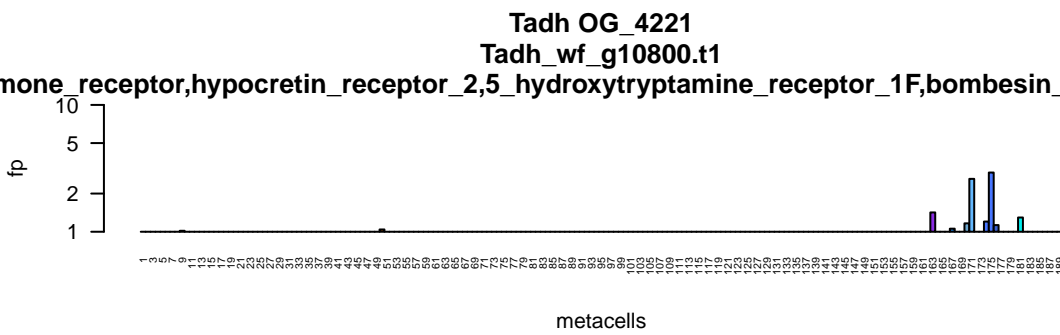
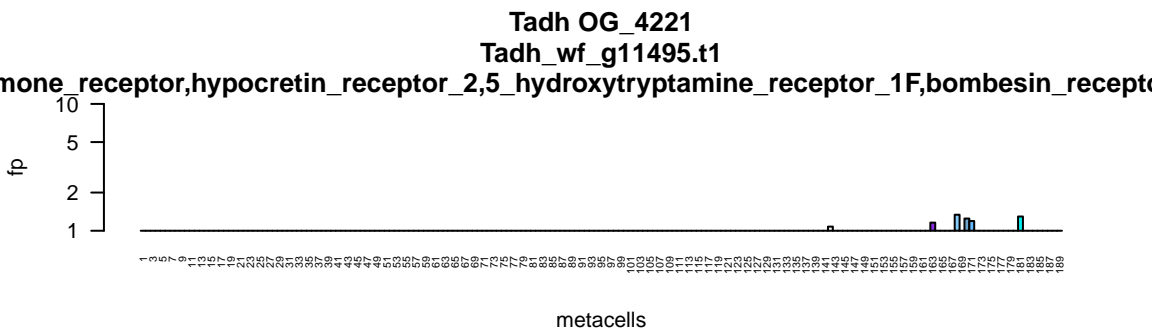
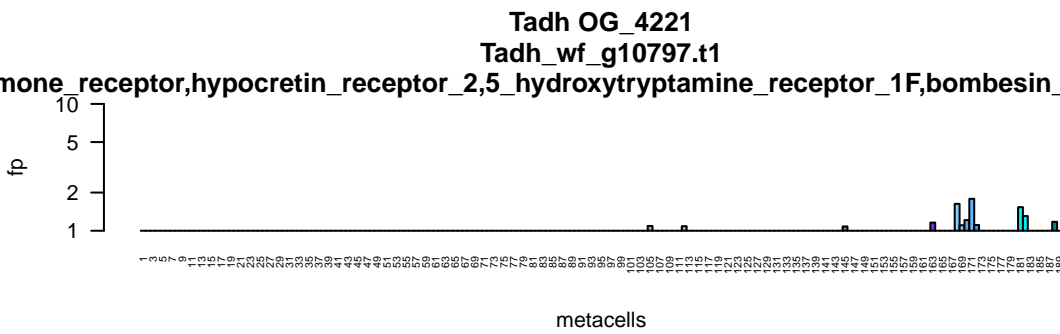


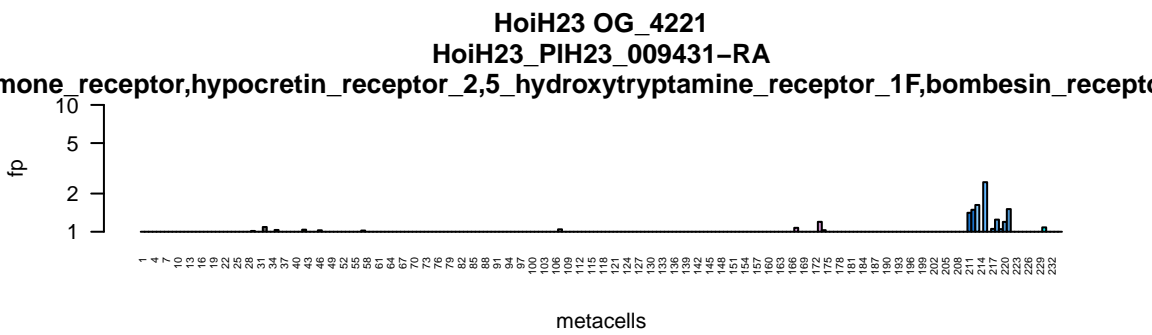
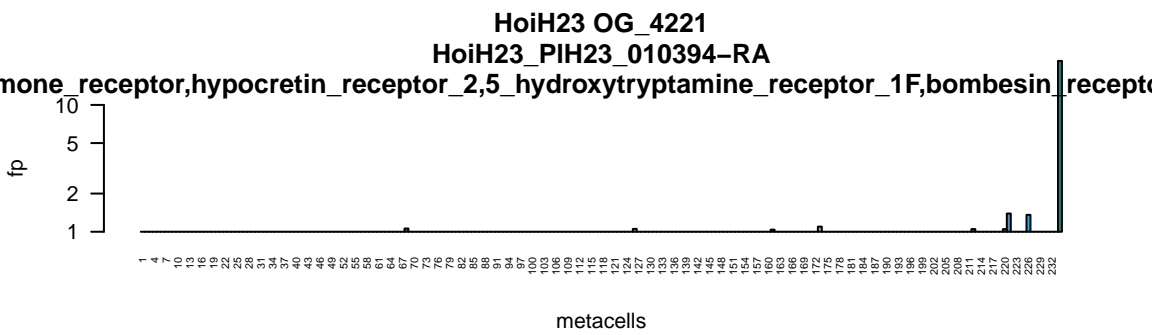
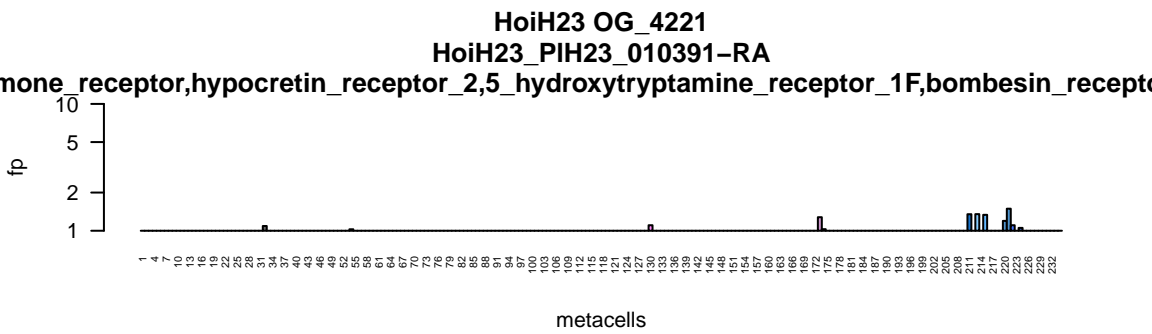
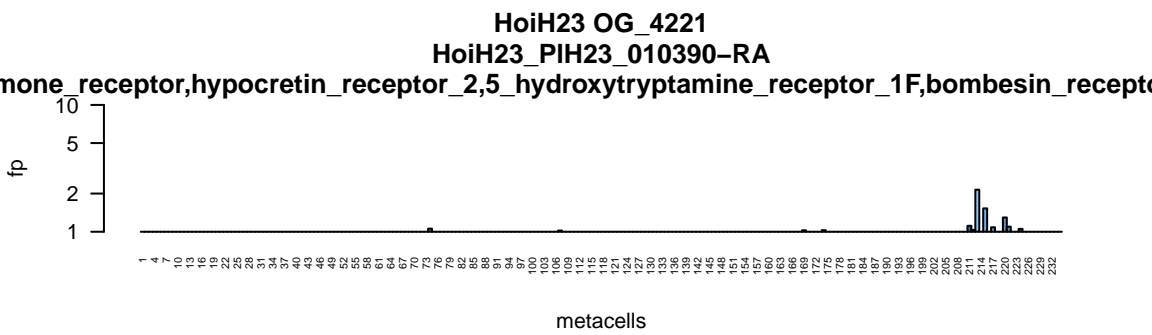
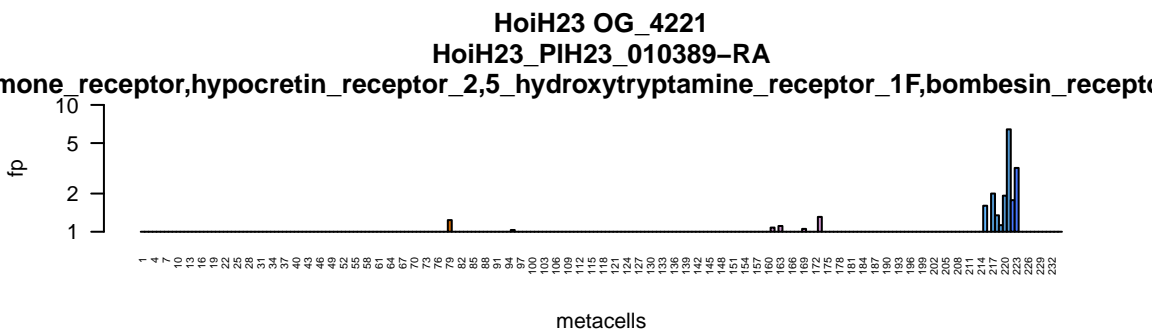
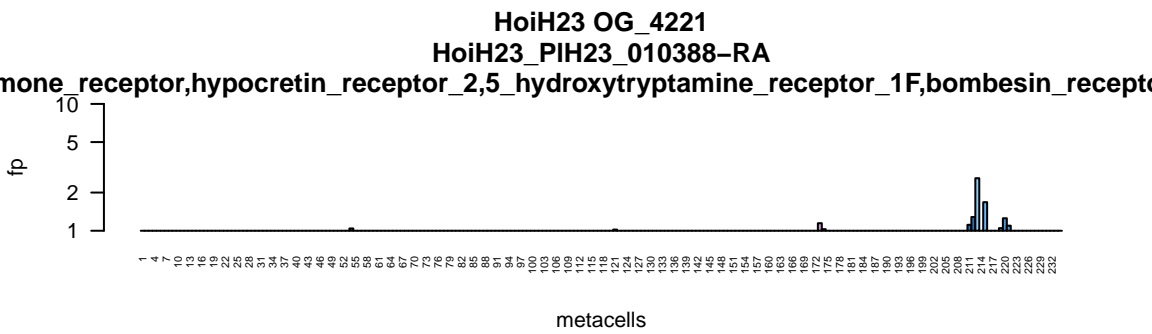
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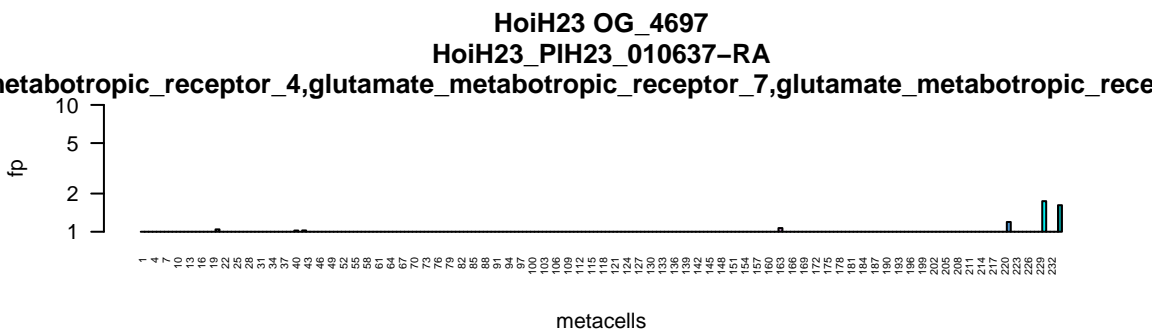
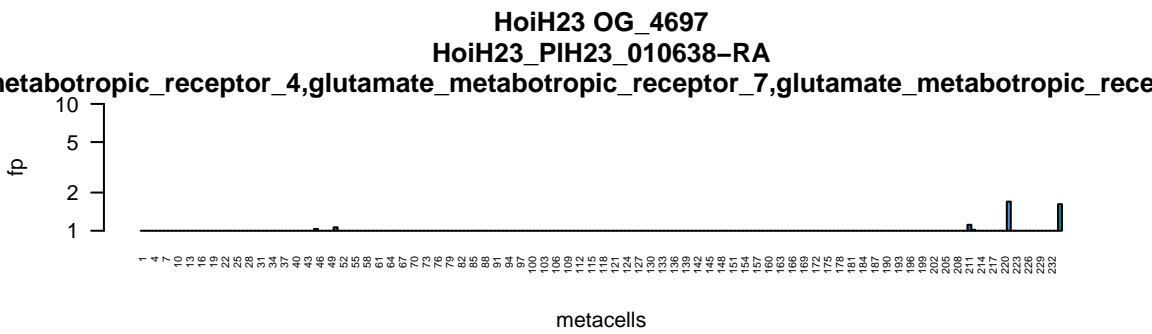
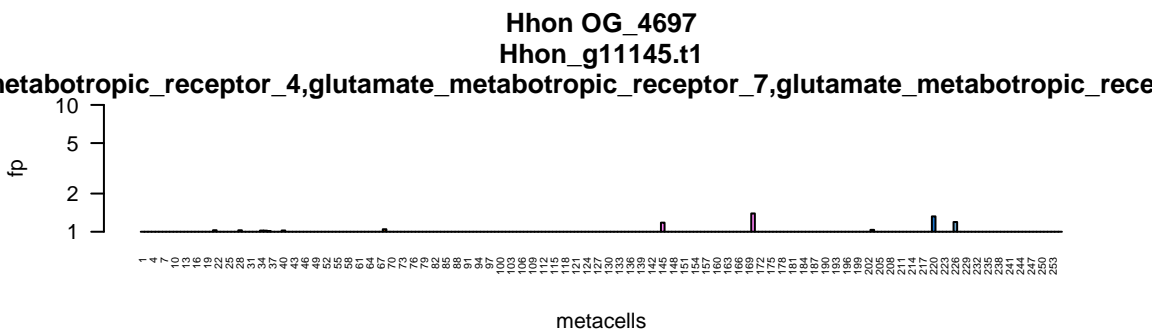
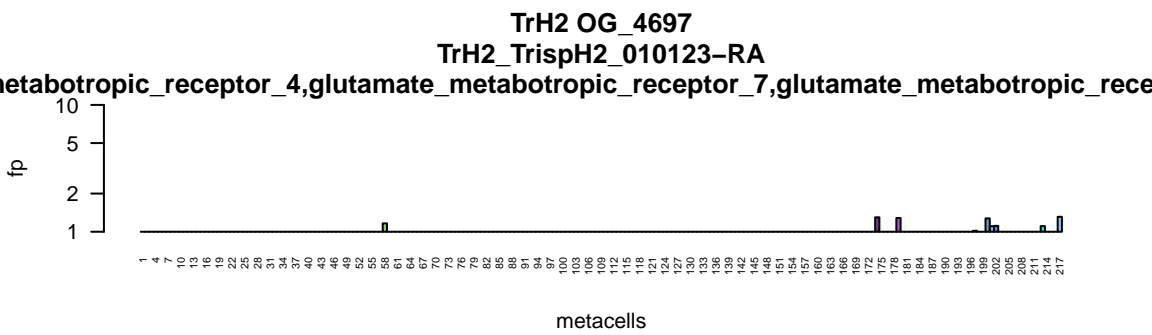
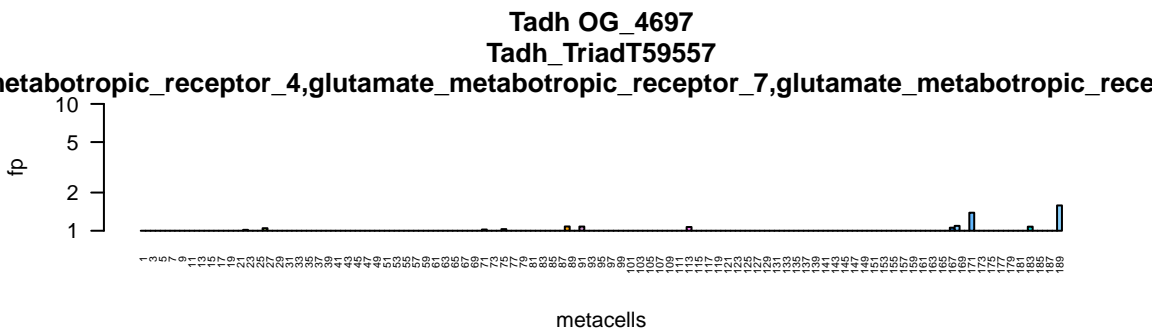


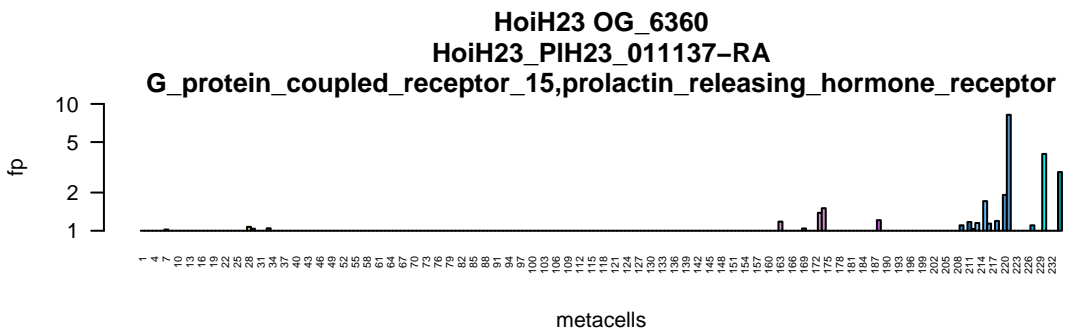
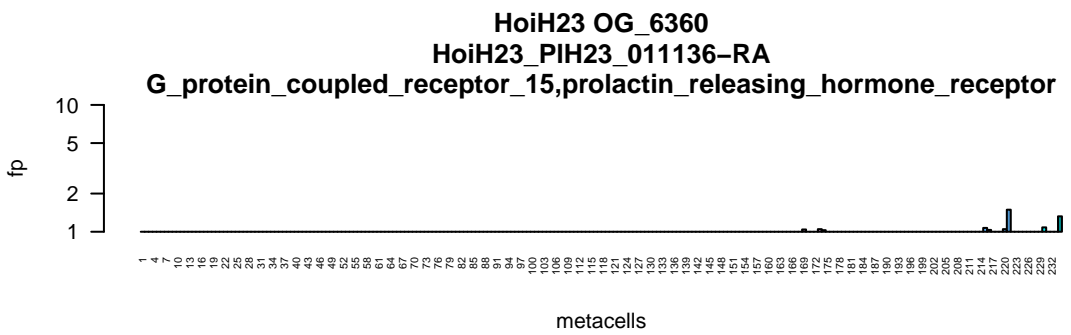
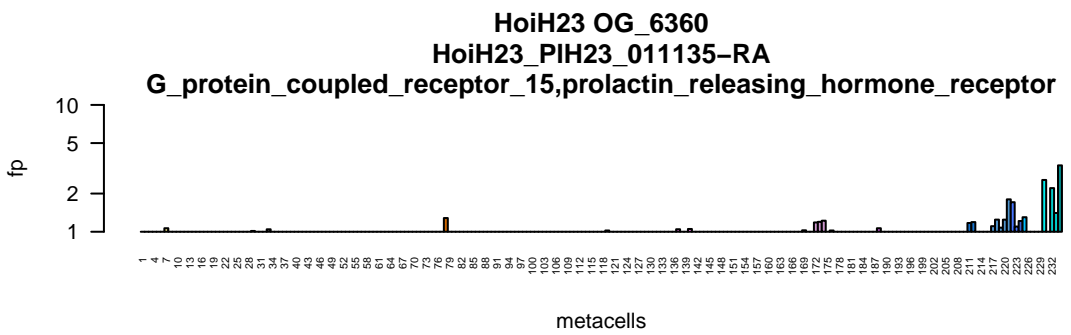
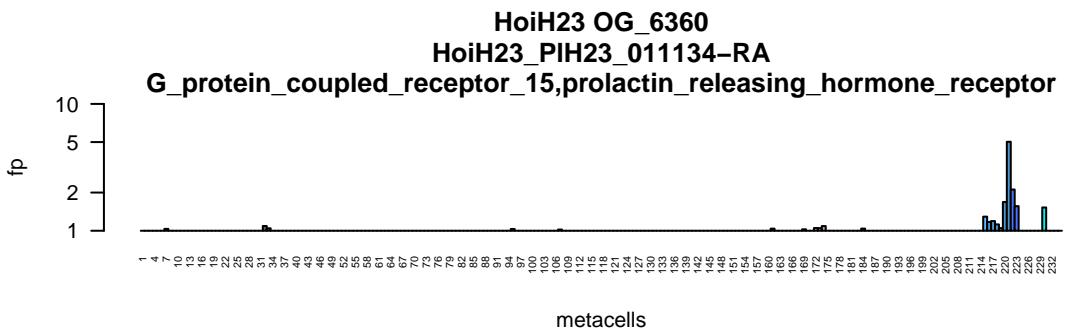
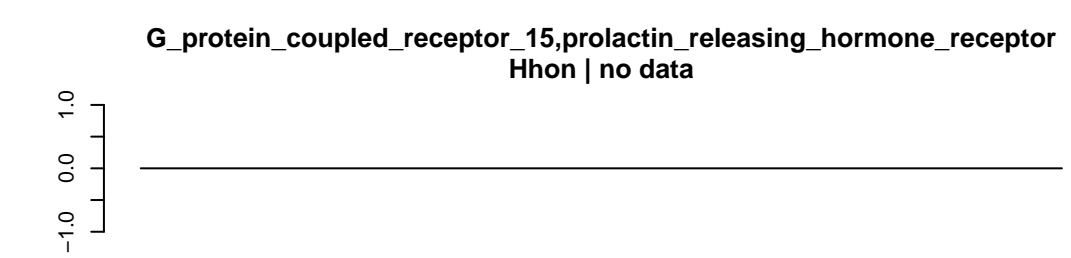
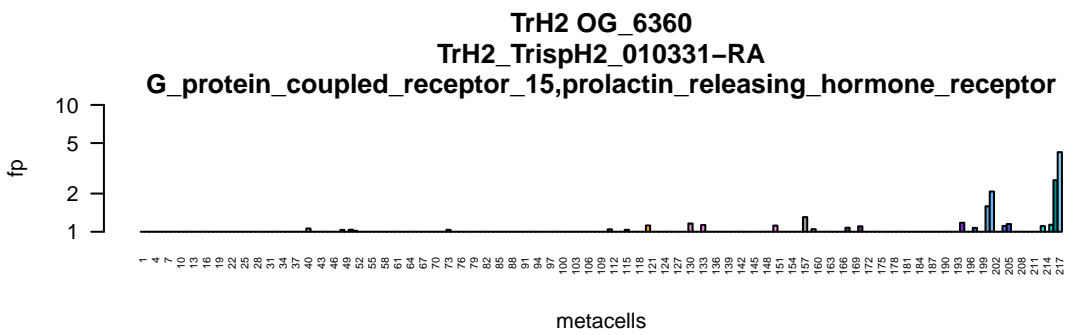
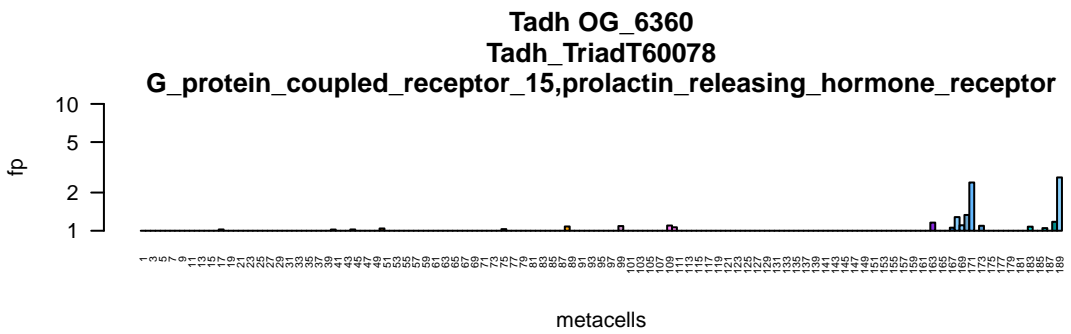


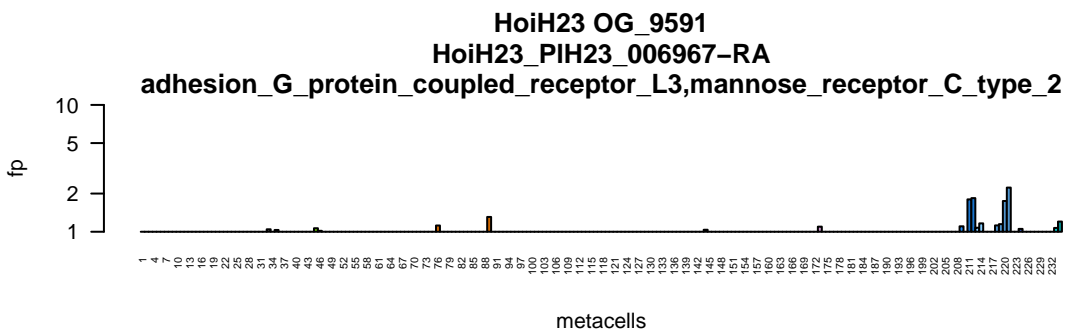
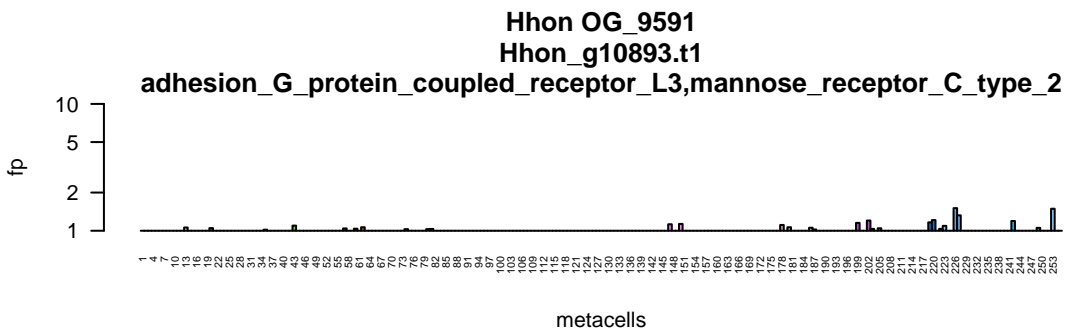
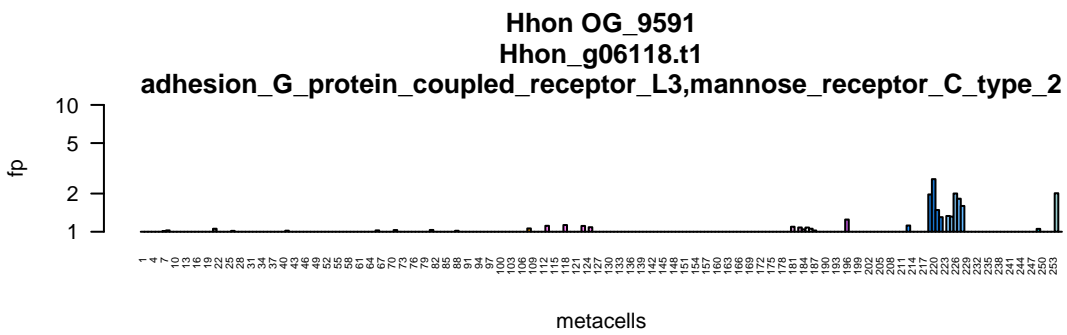
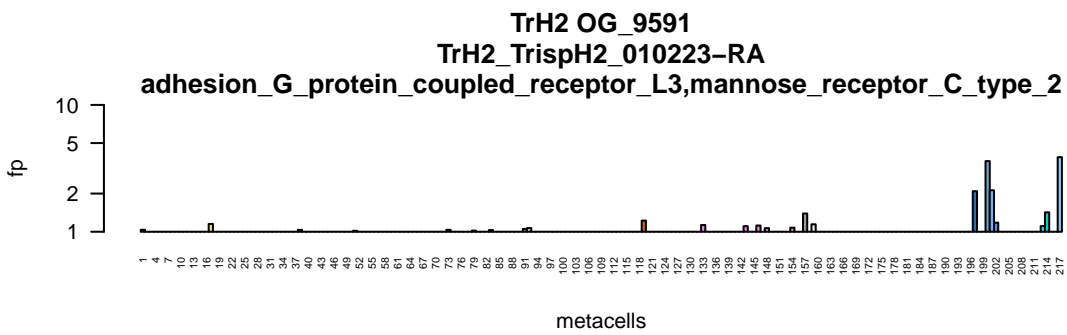
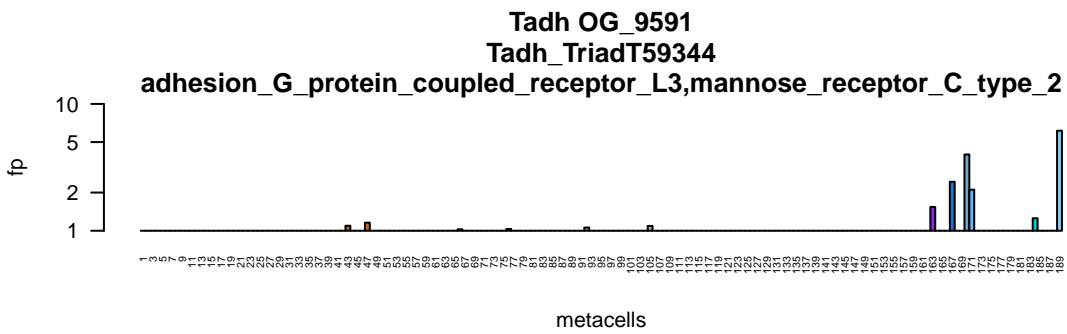


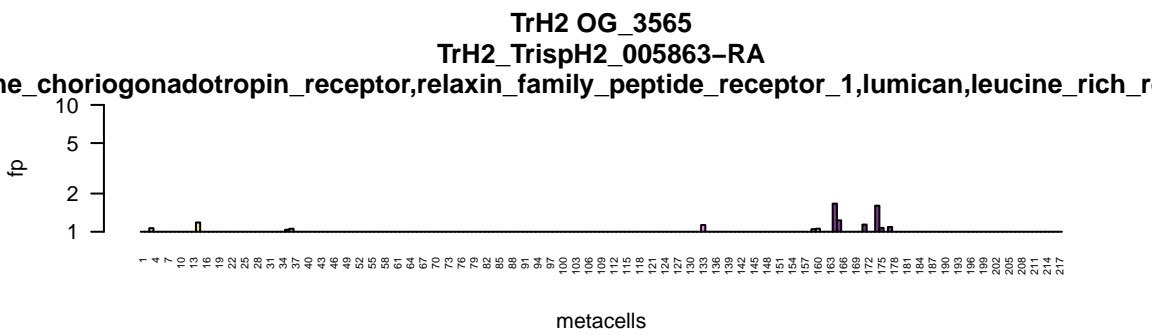
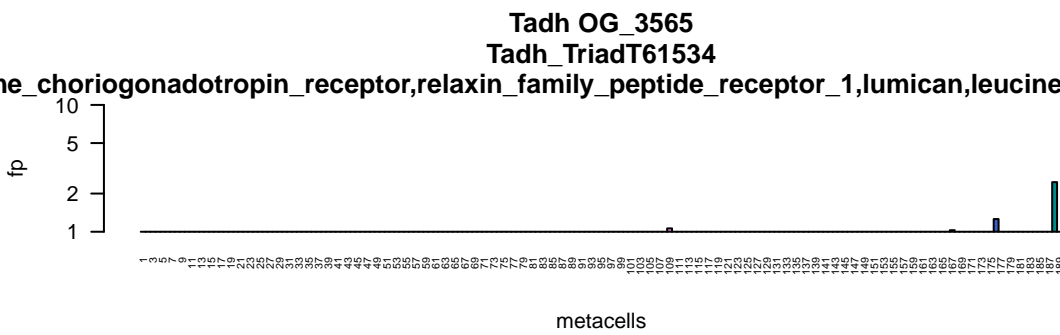
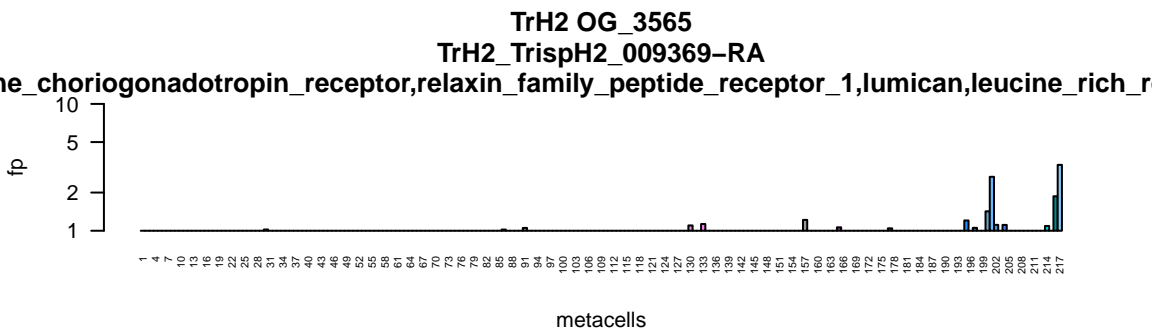
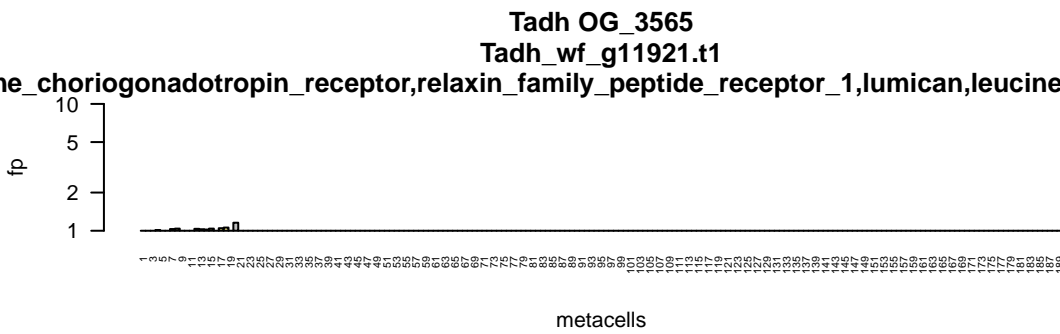
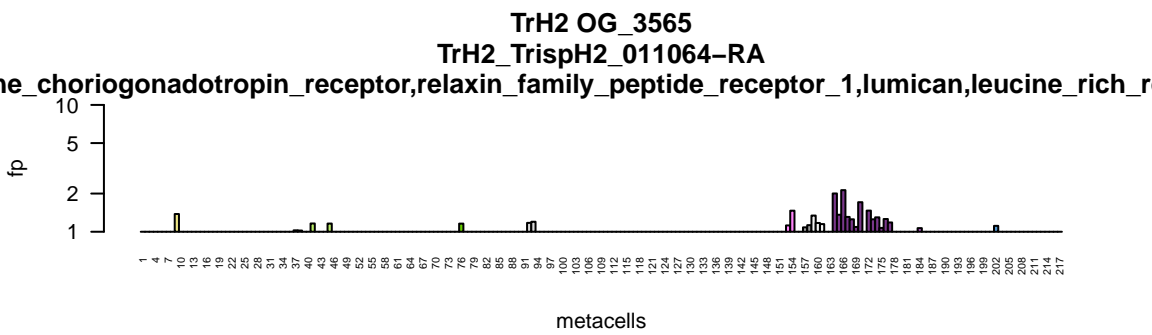
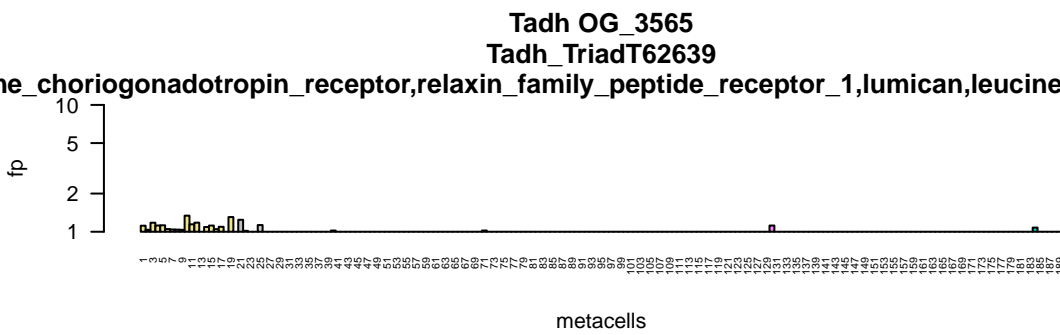
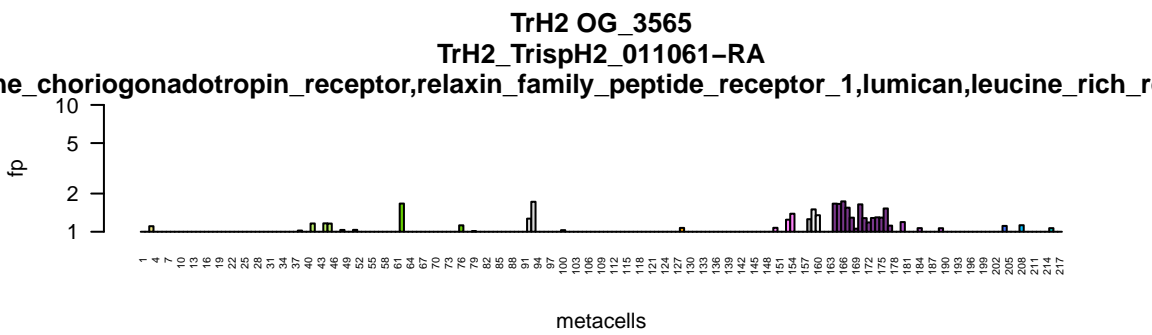
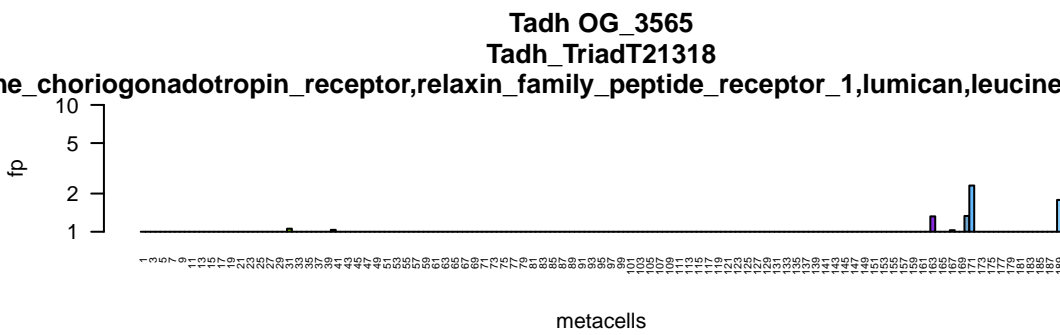
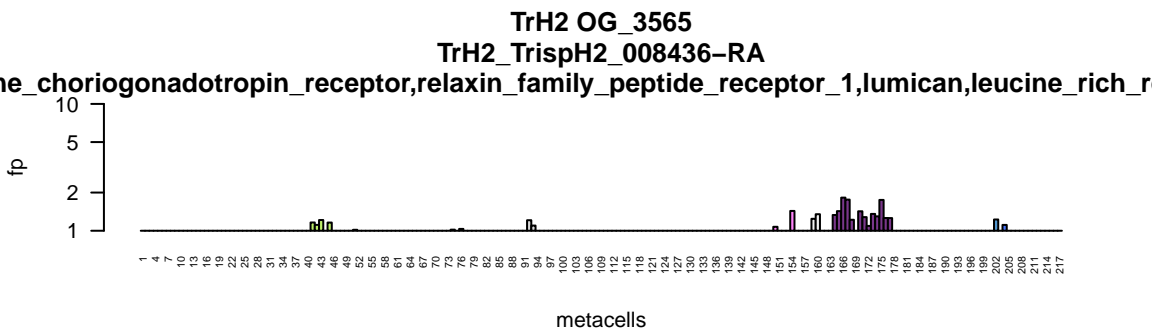
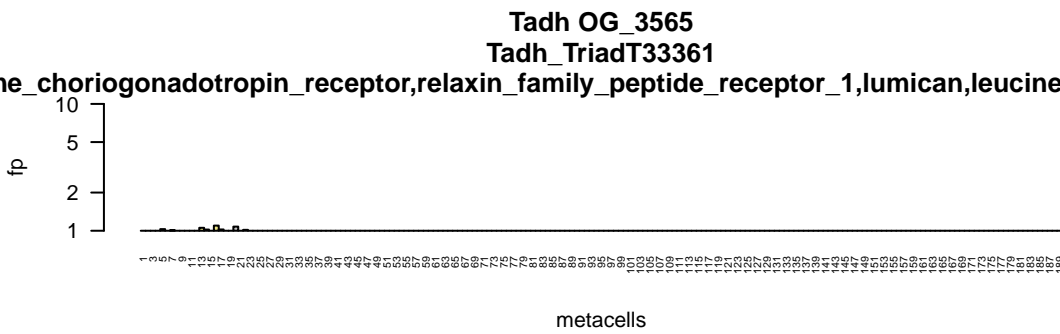
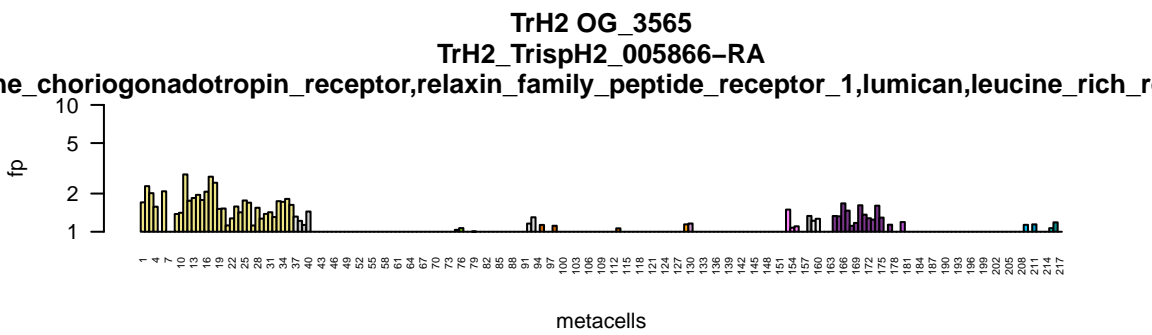
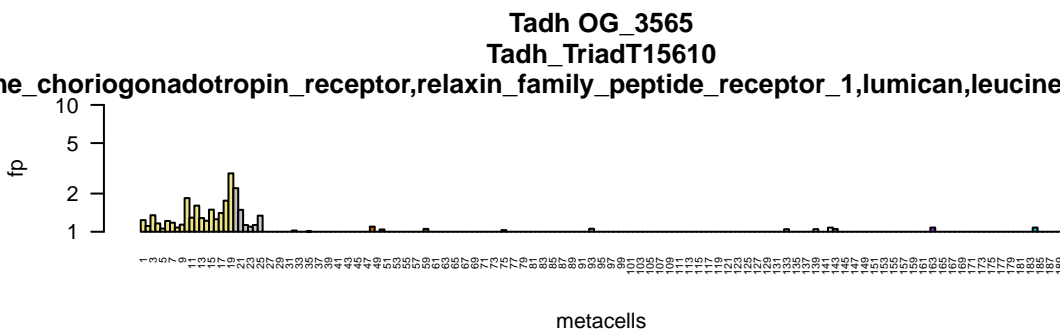
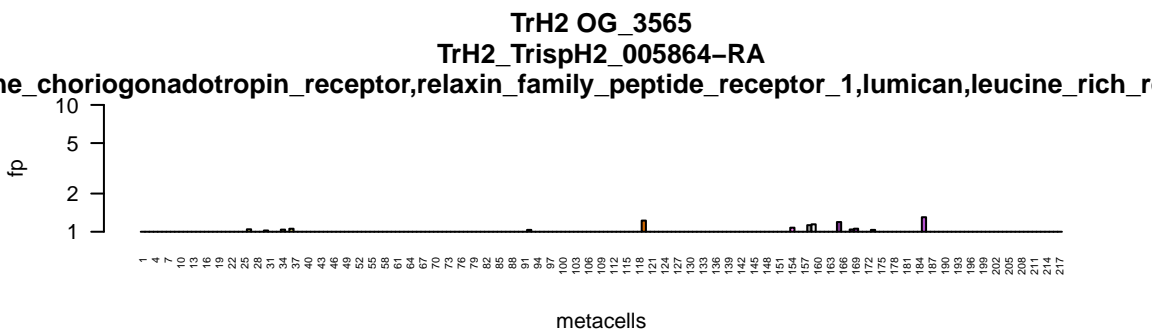
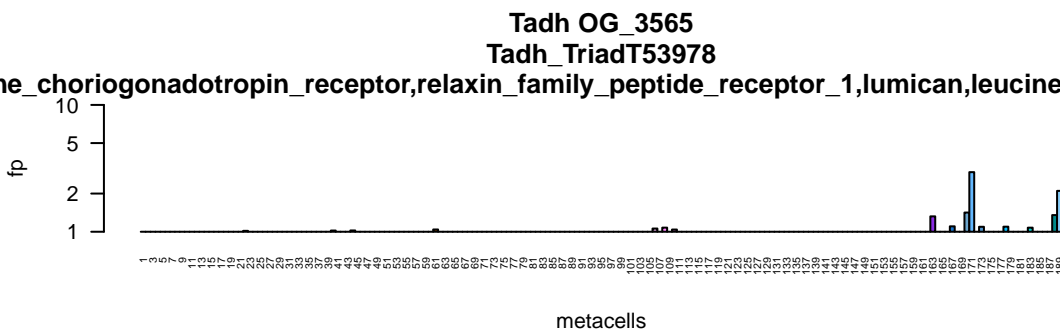
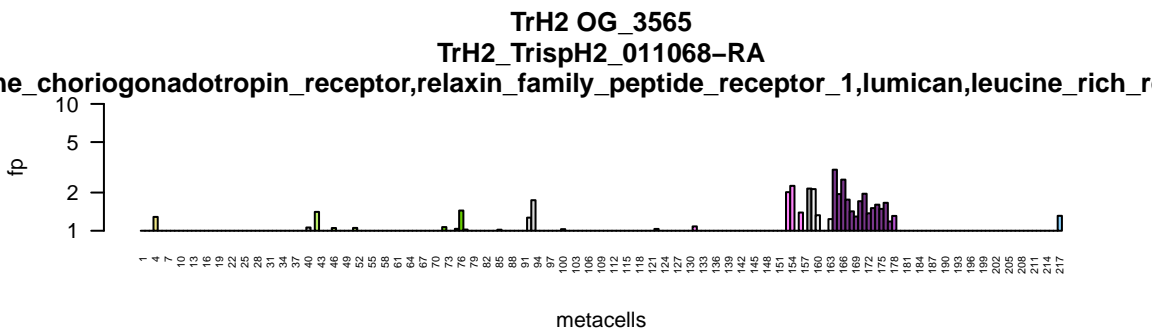
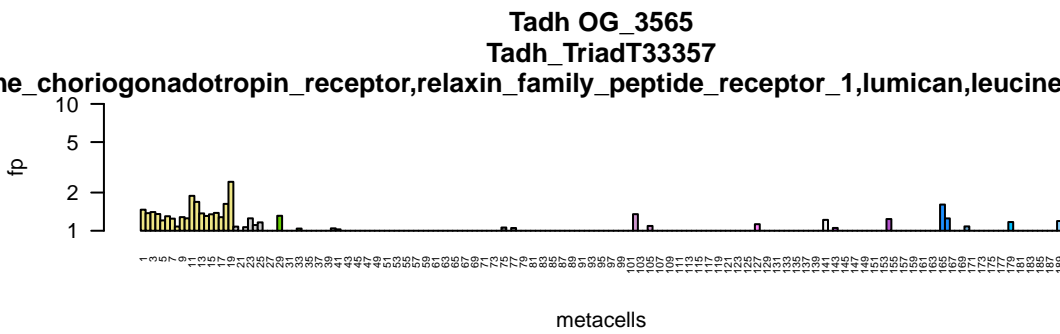


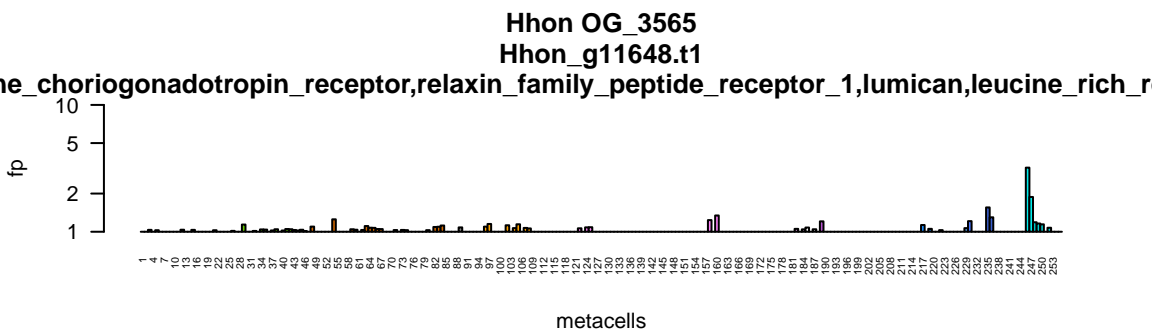
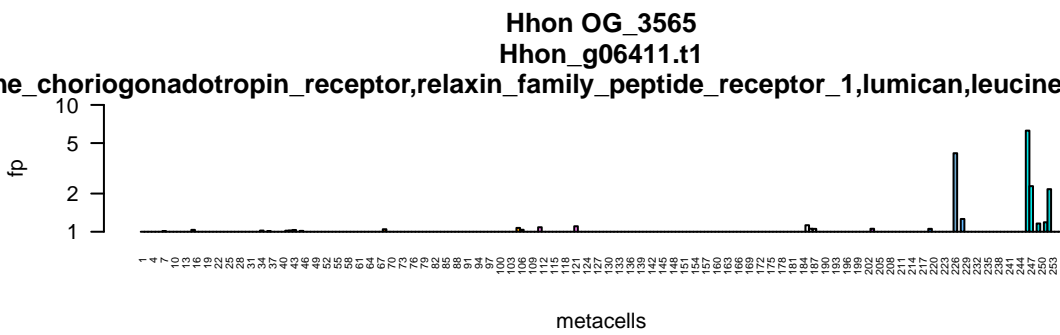
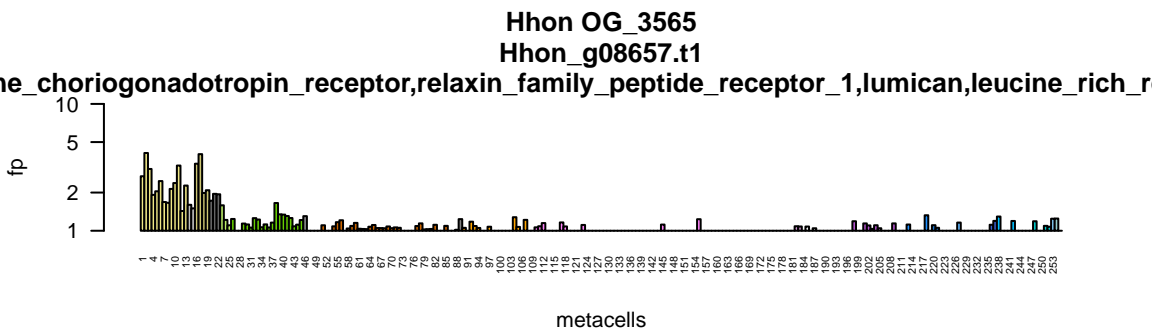
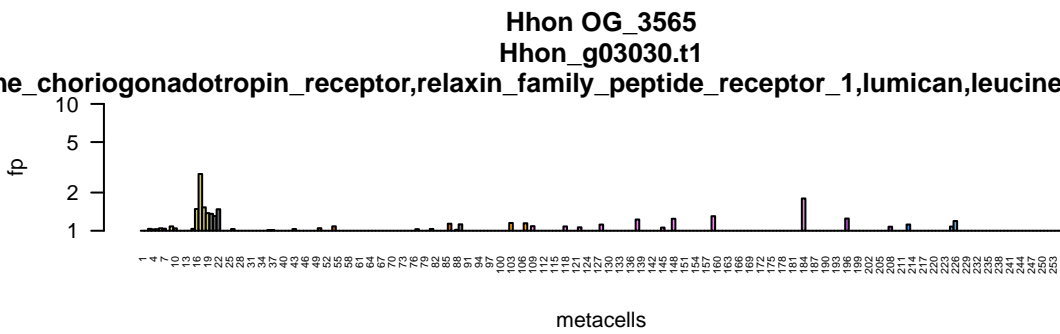
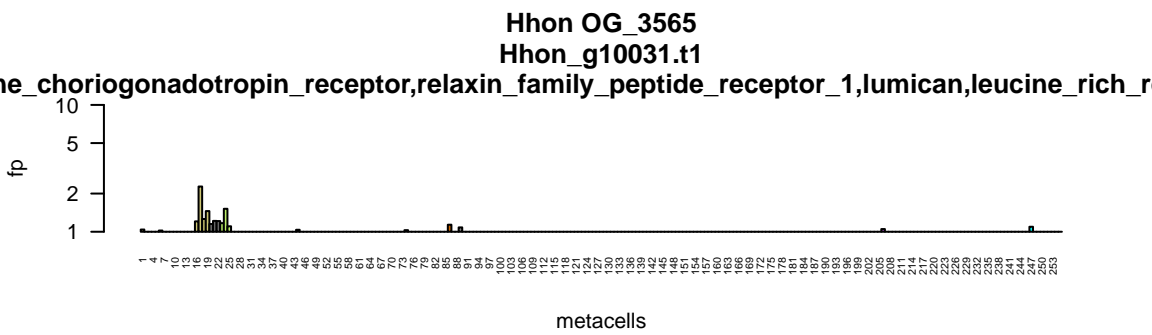
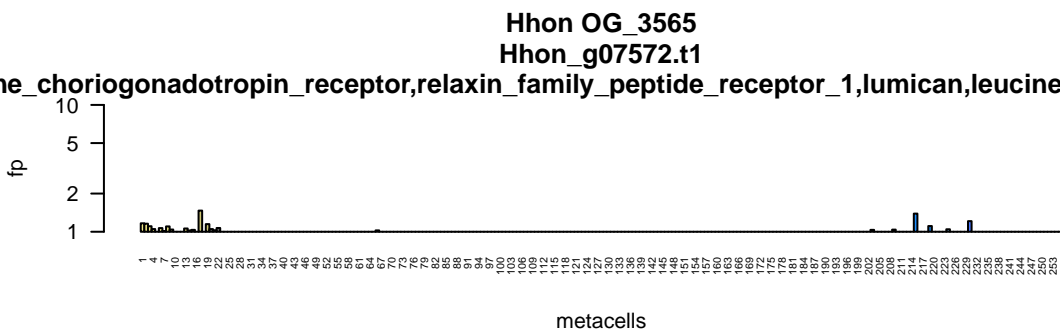
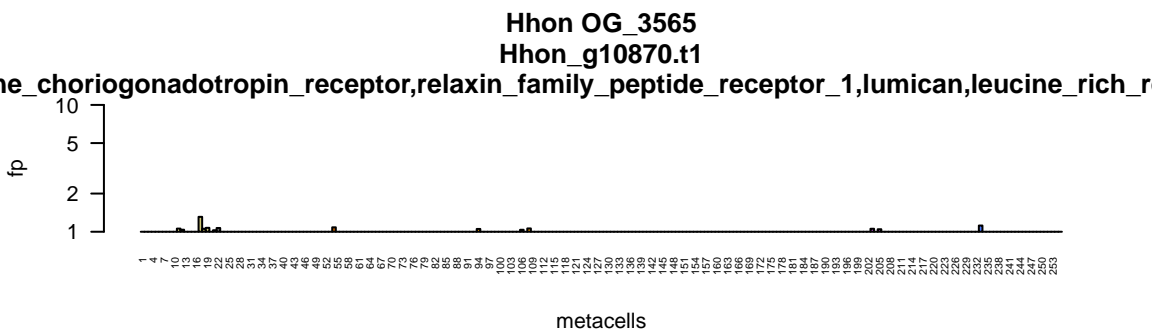
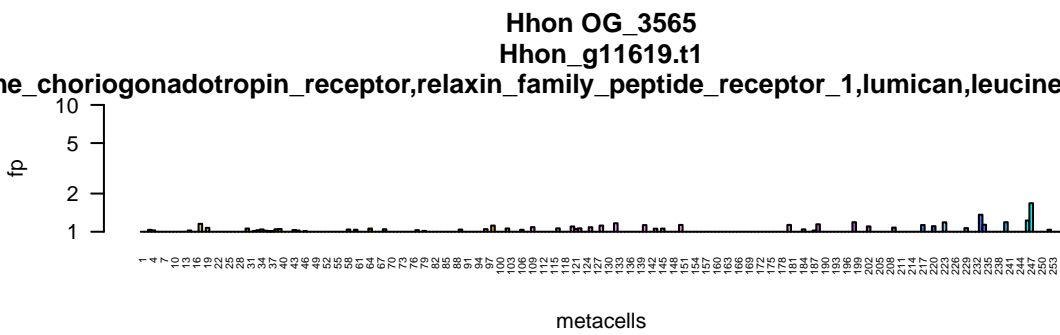
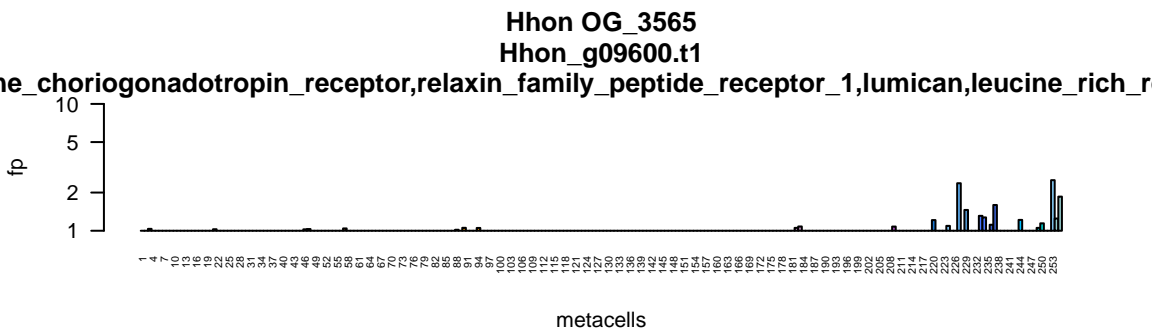
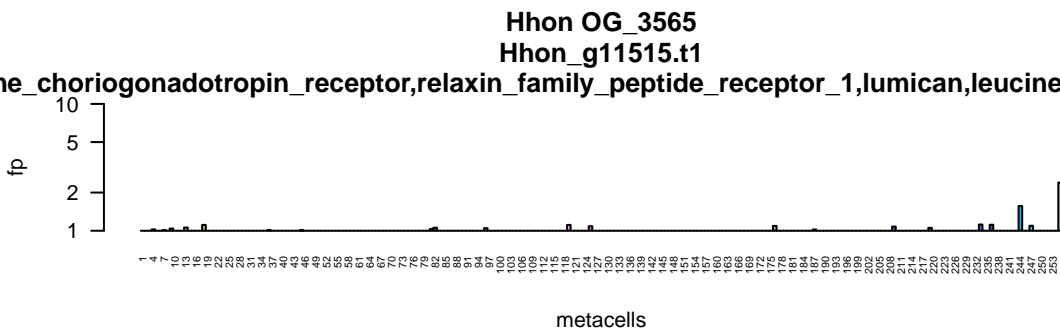
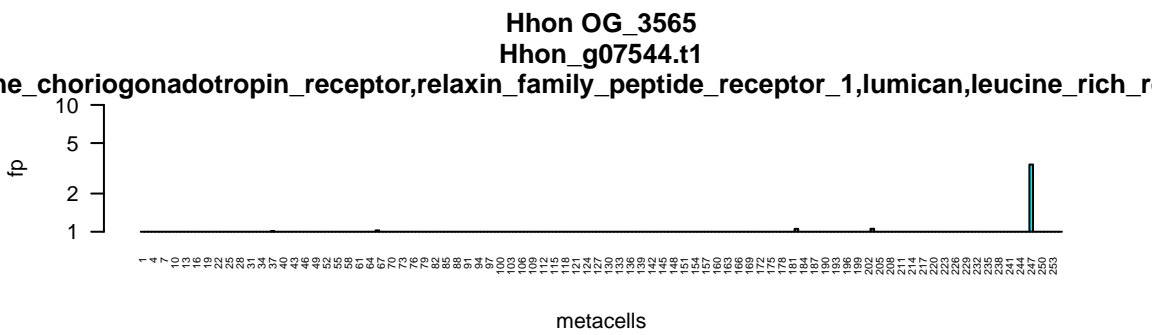
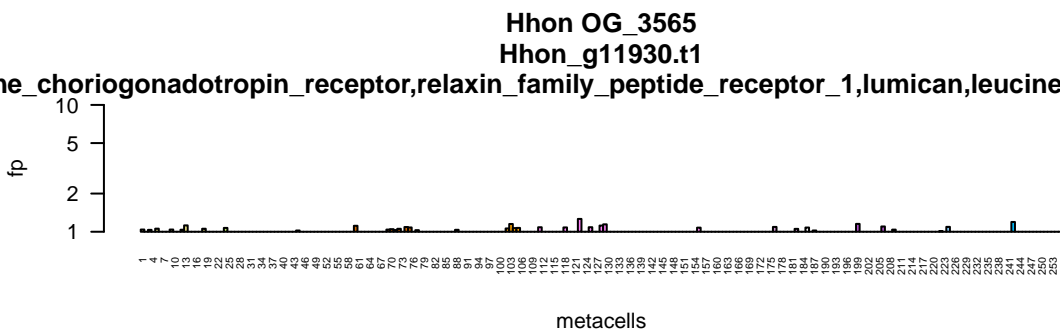
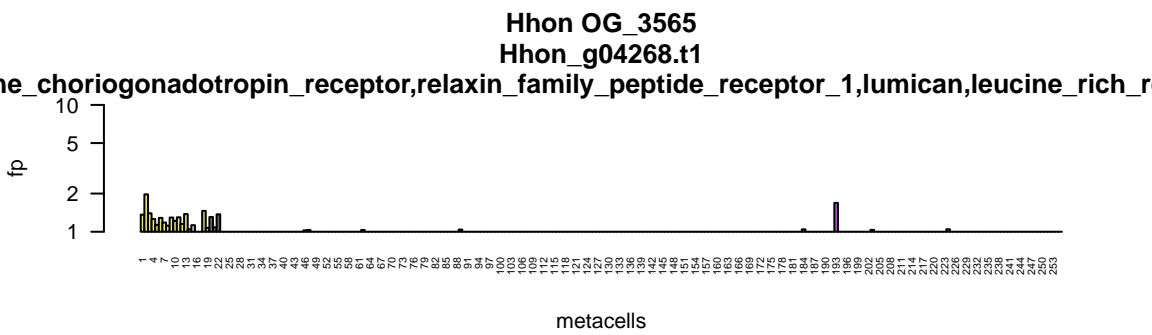
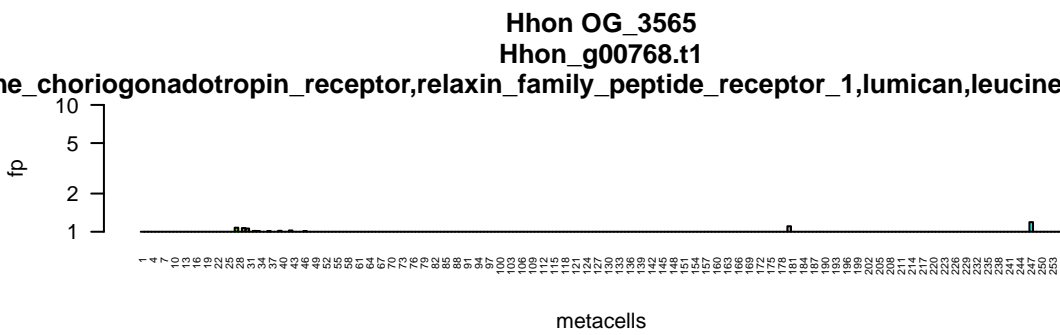
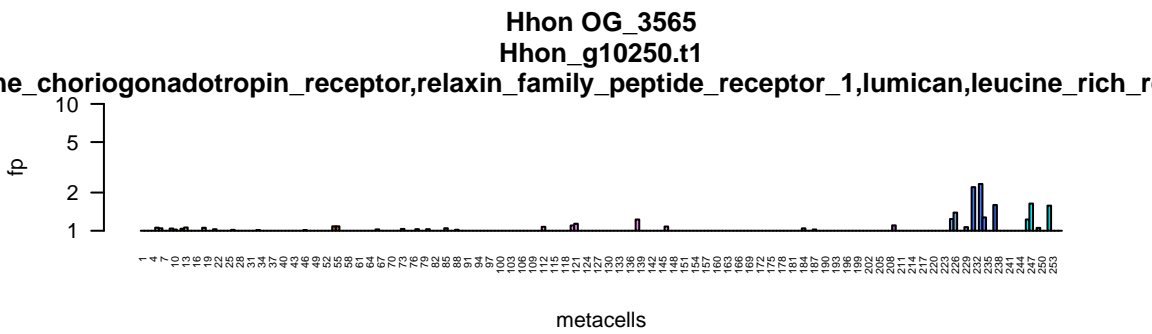
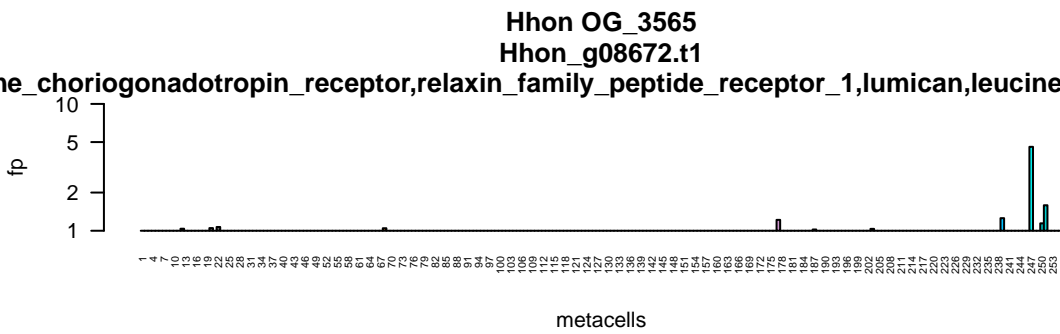


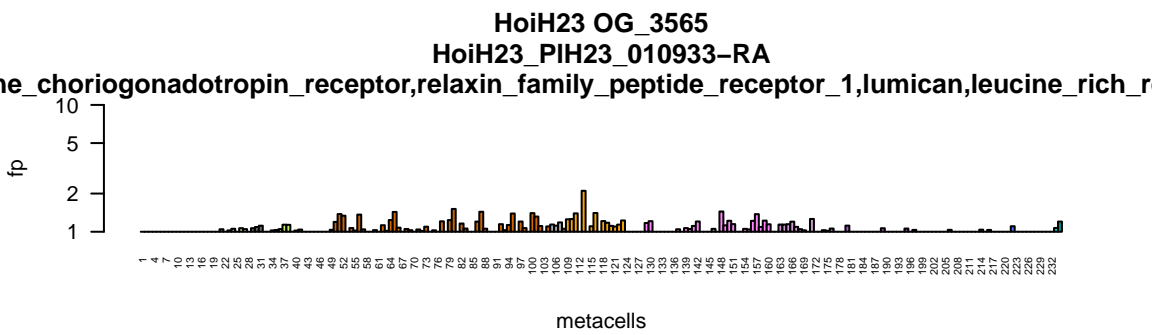
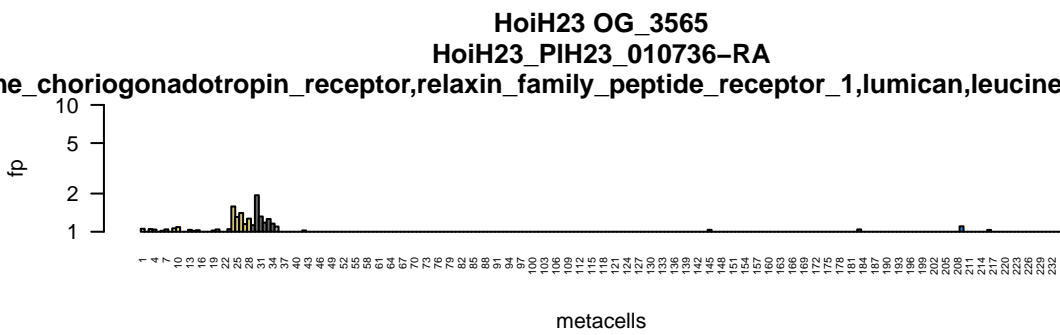
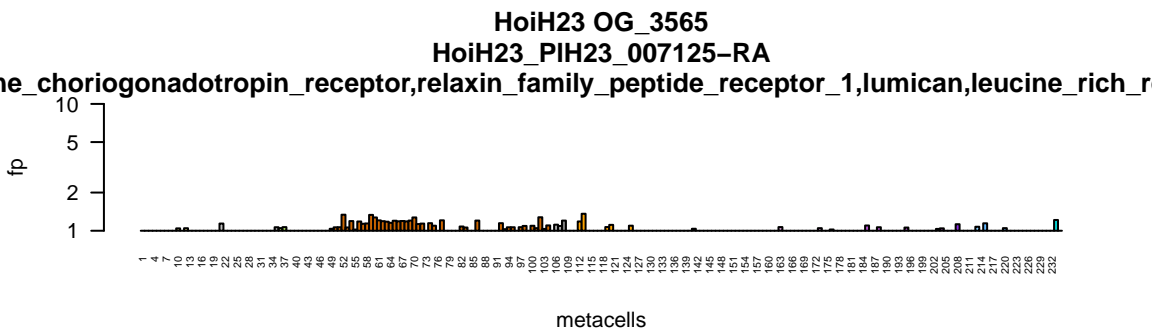
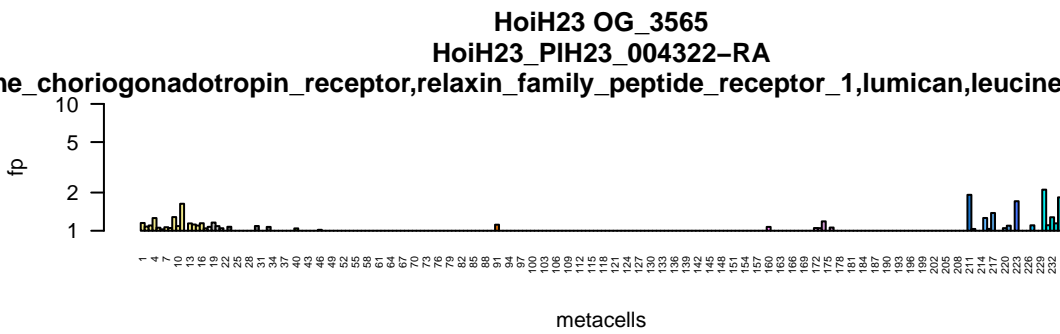
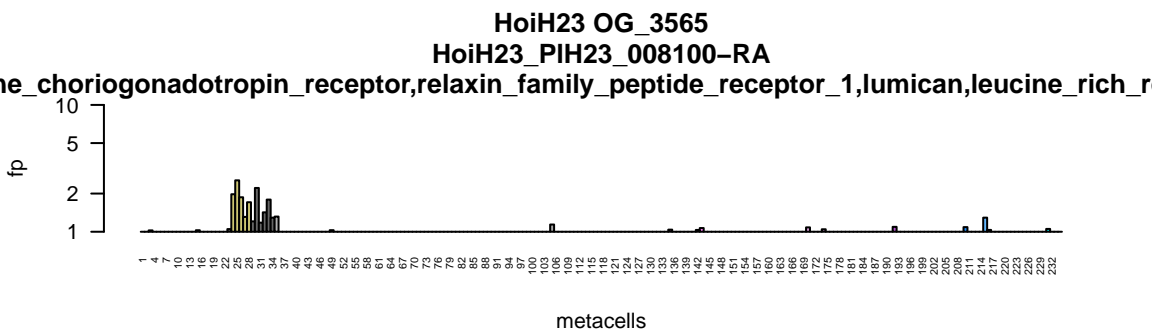
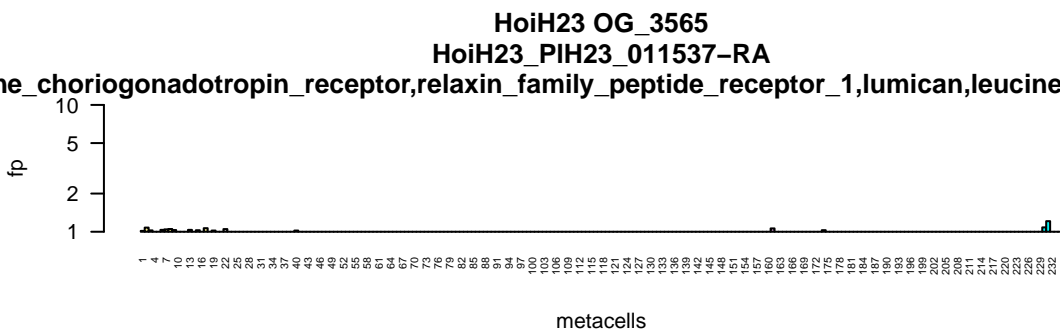
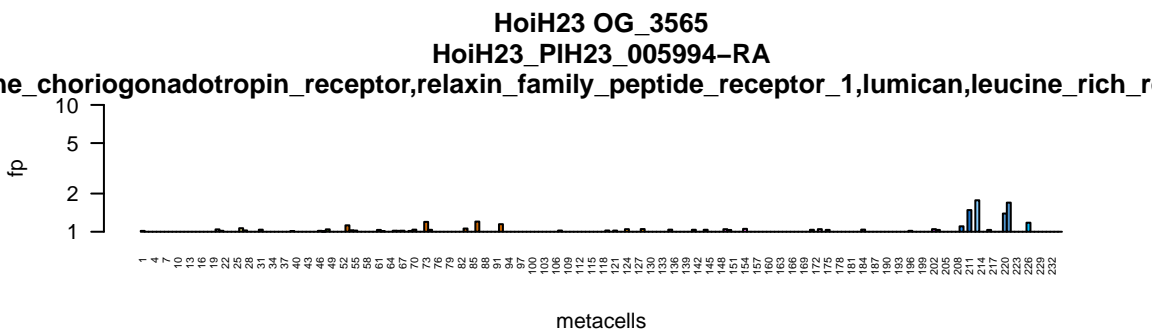
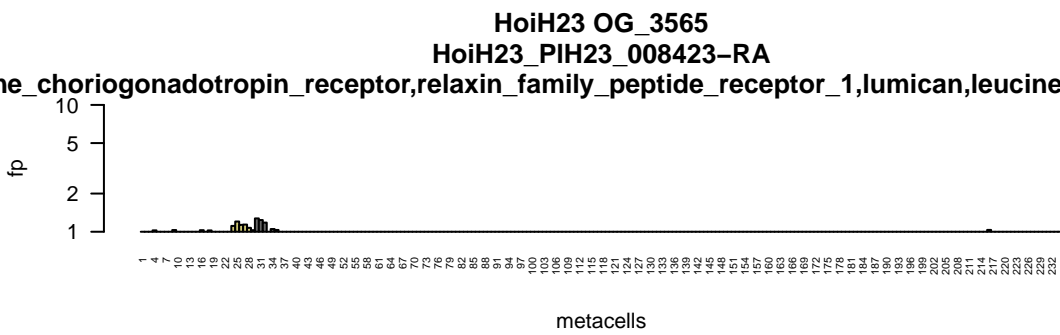
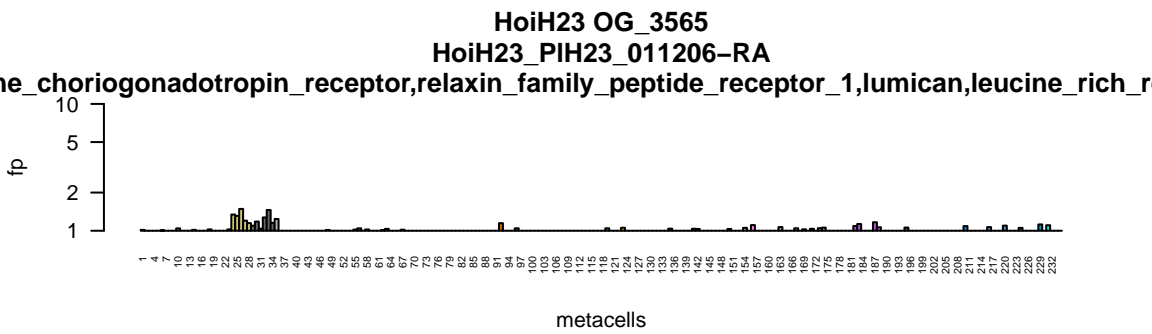
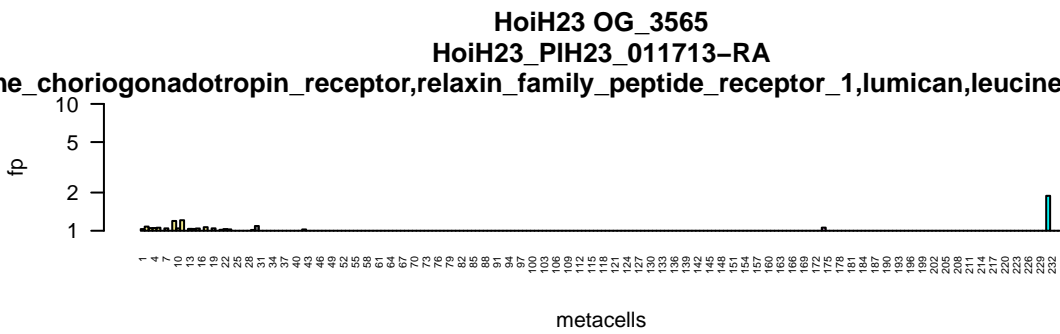
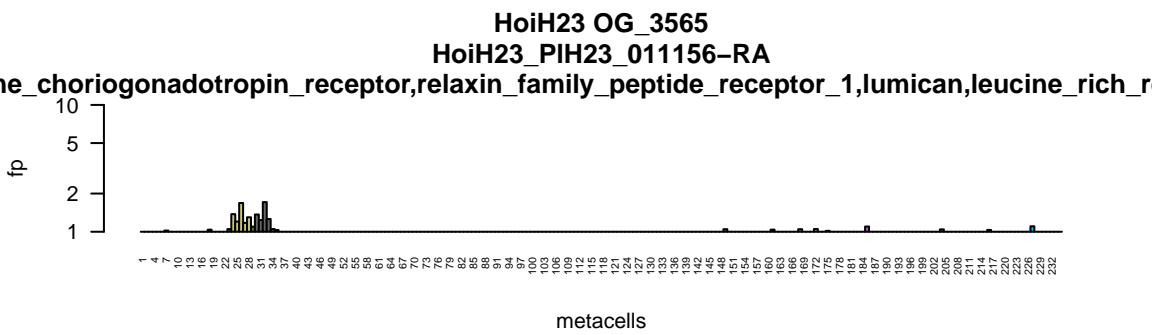
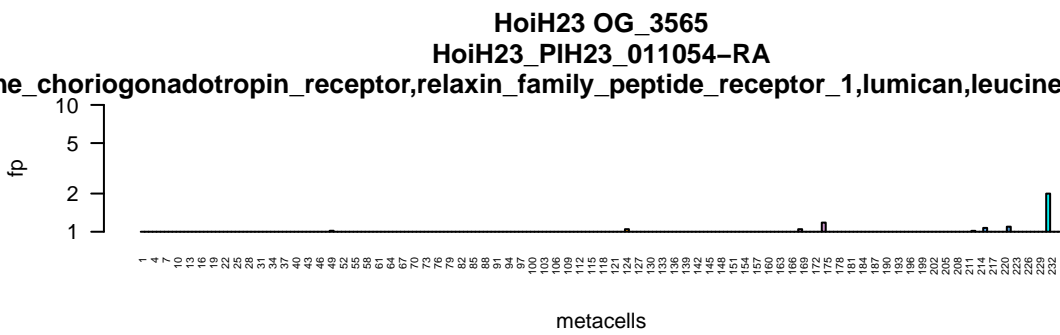
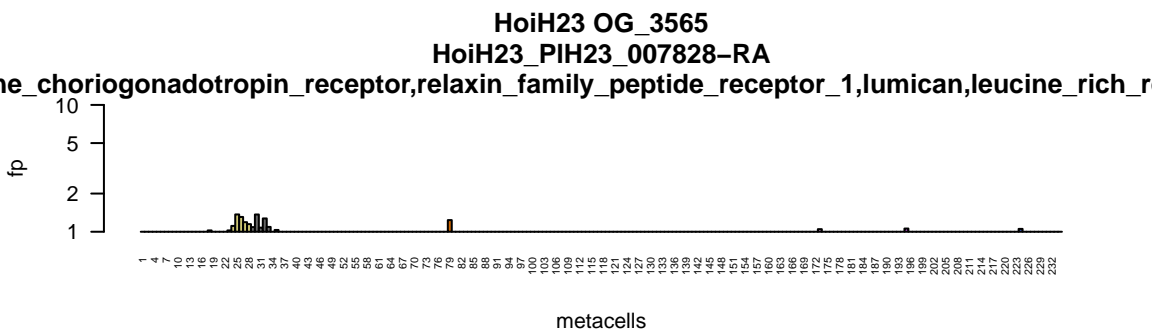
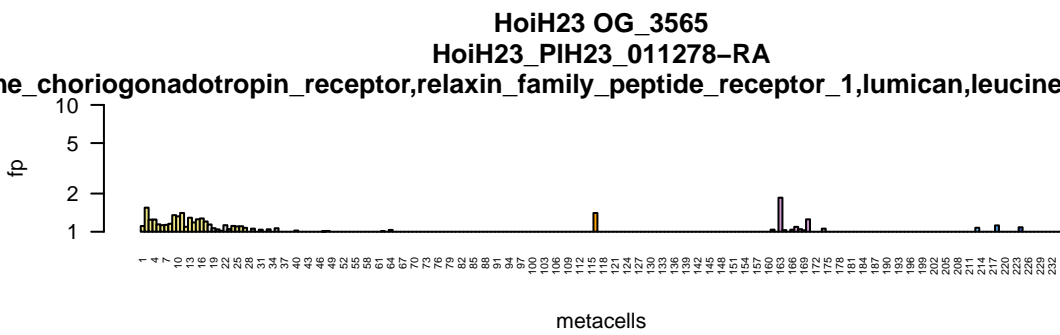
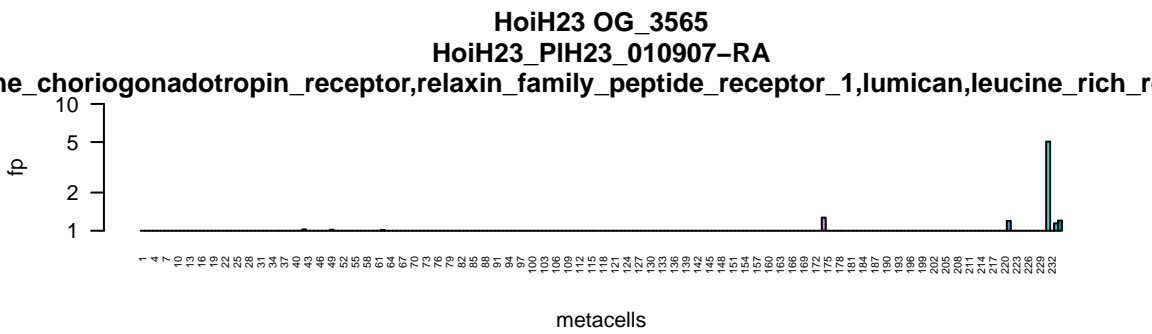
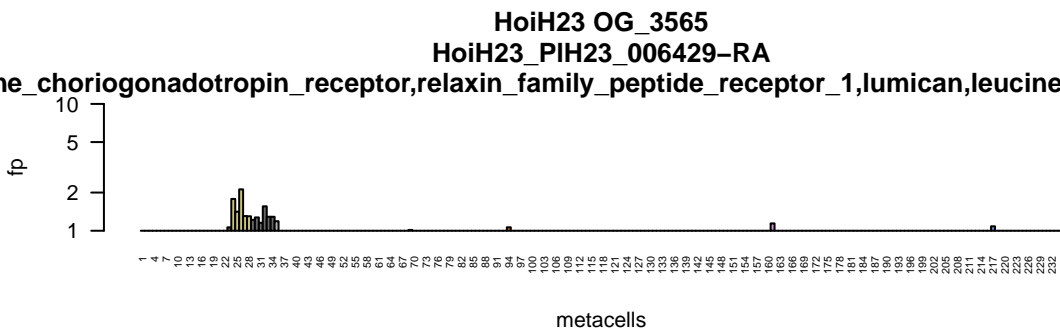


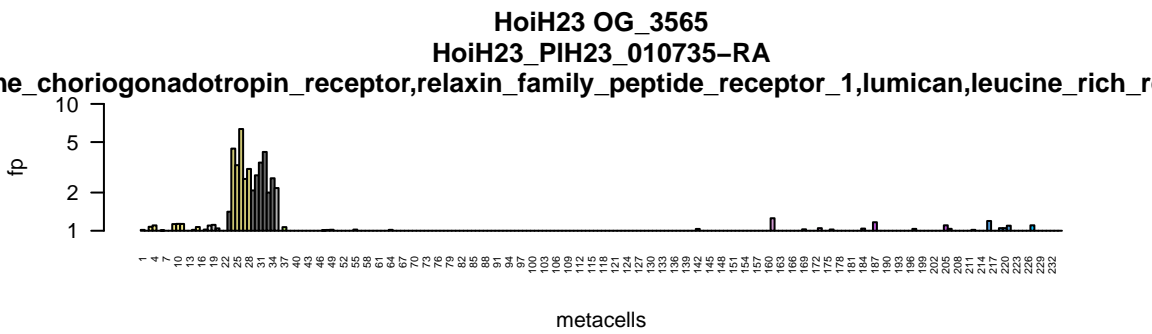
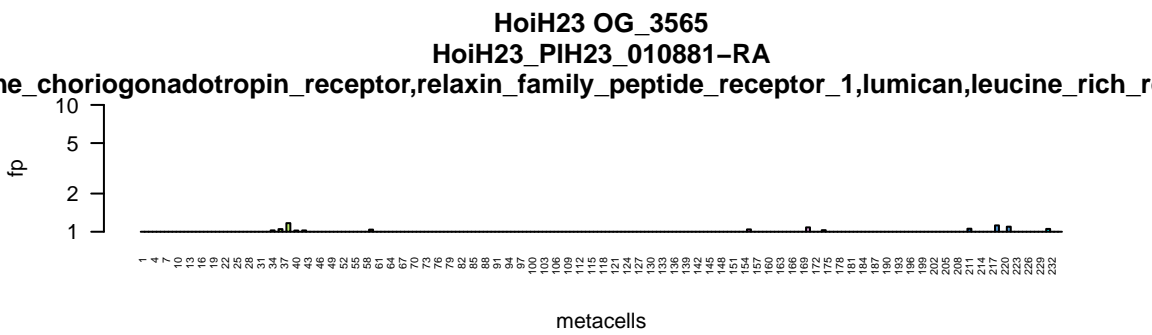
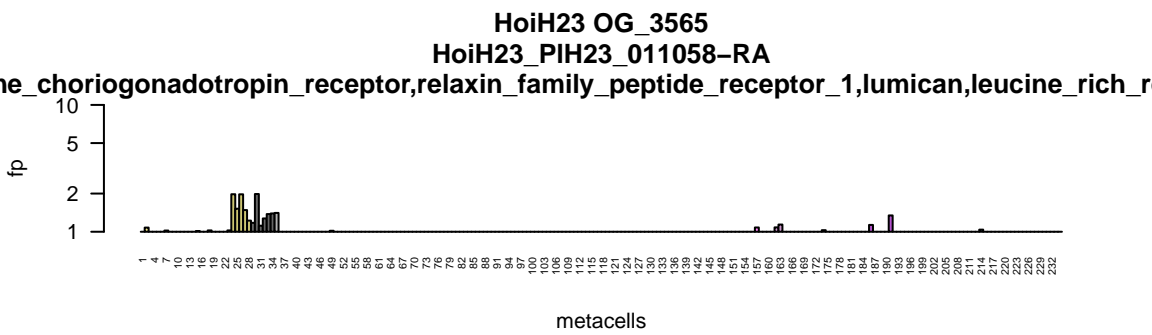
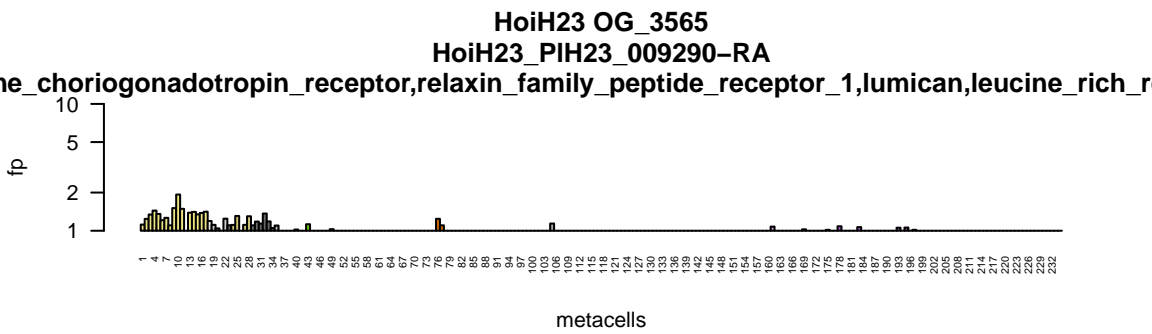


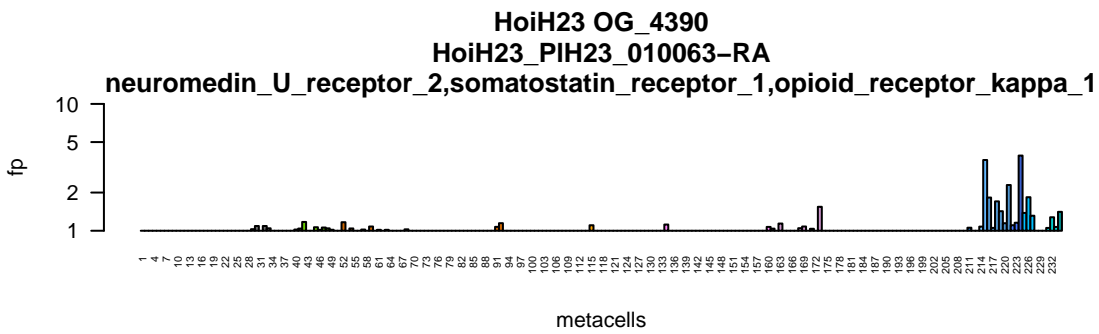
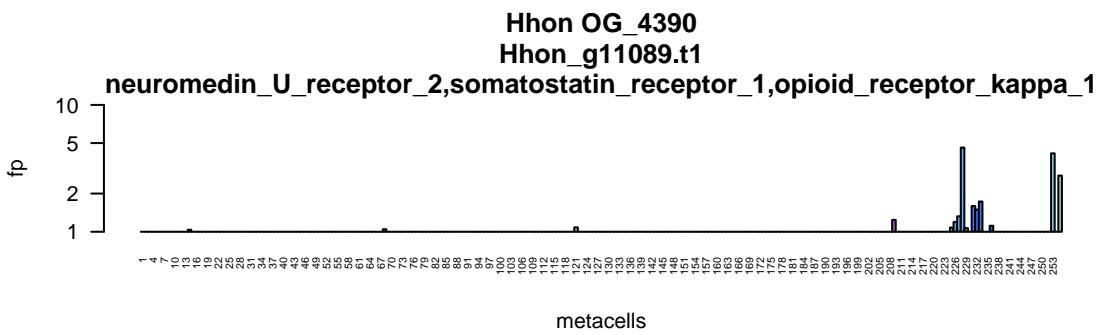
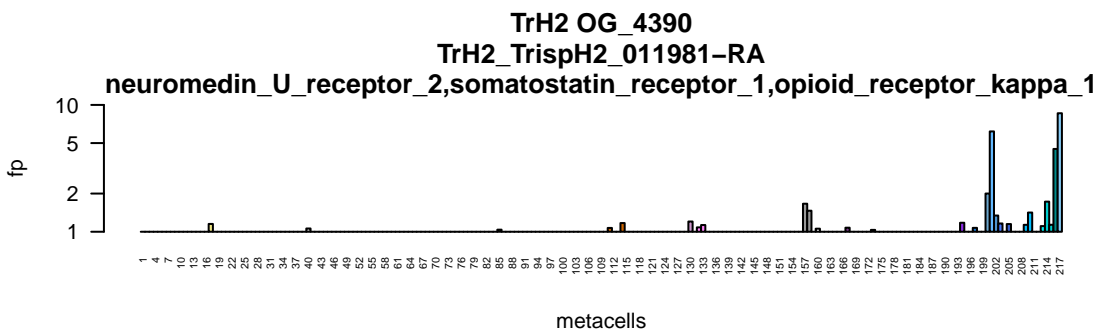
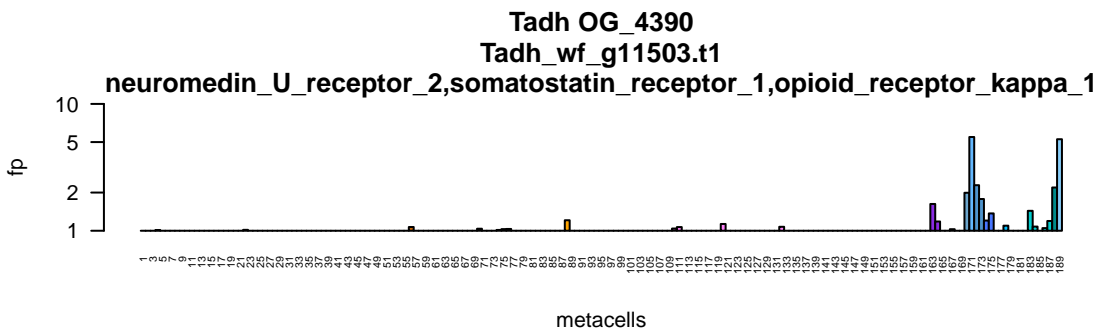




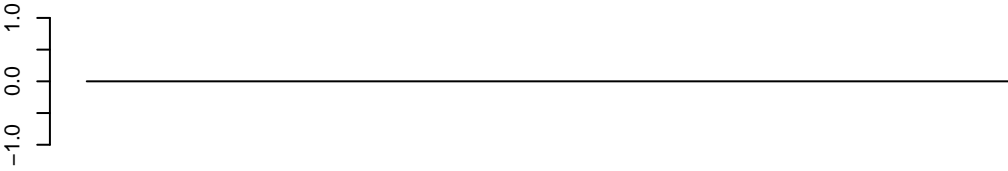




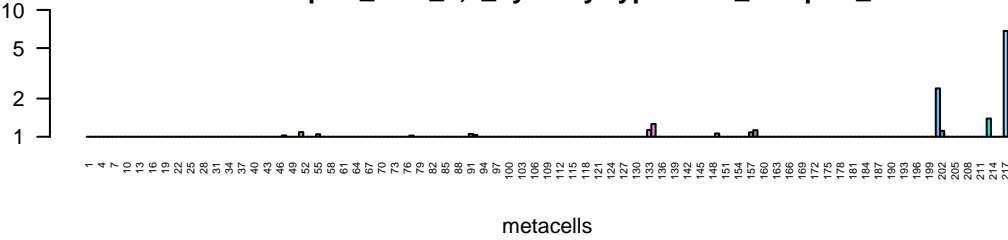




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Tadh | no data



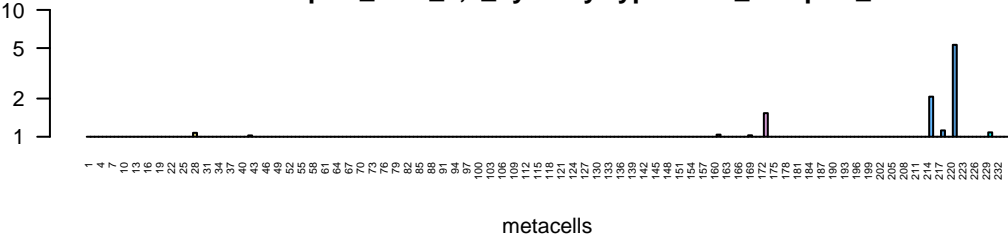
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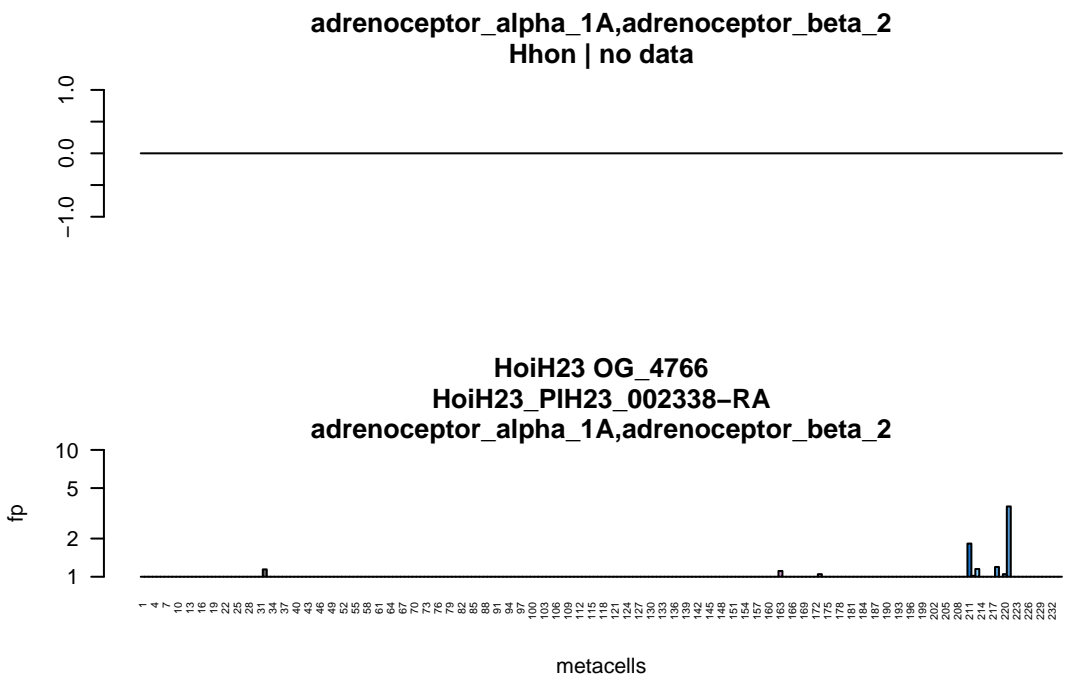
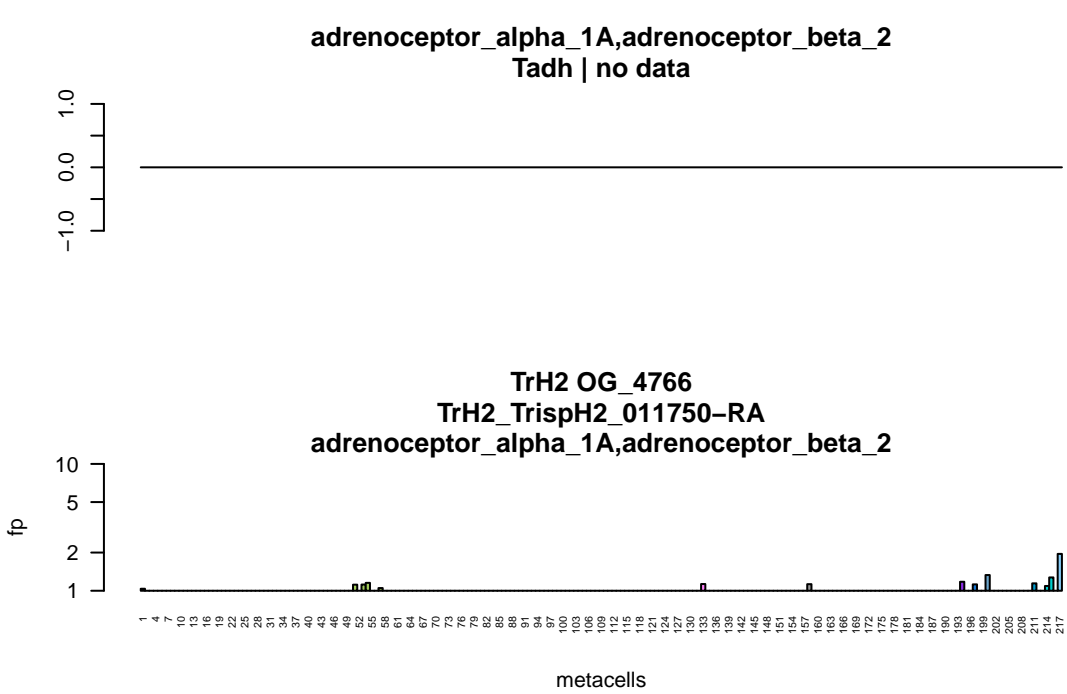


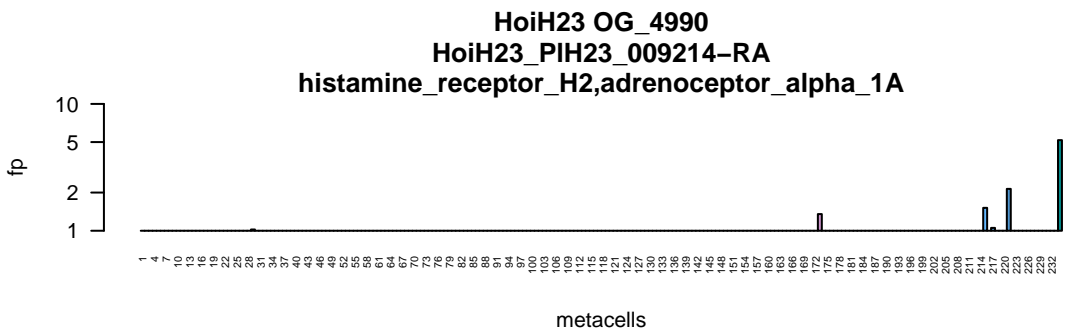
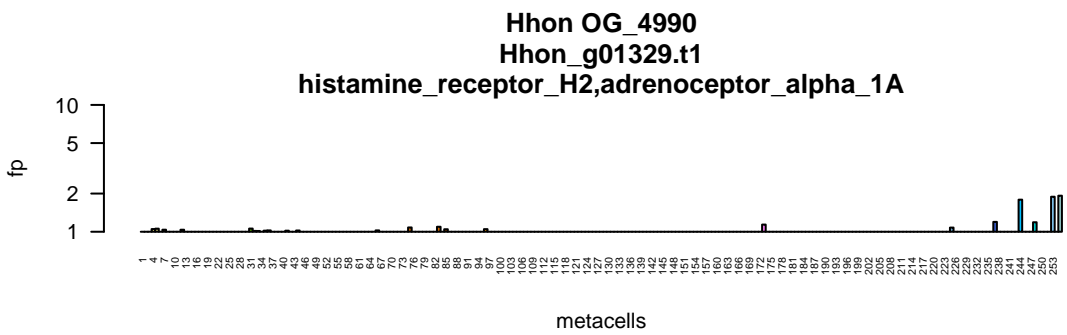
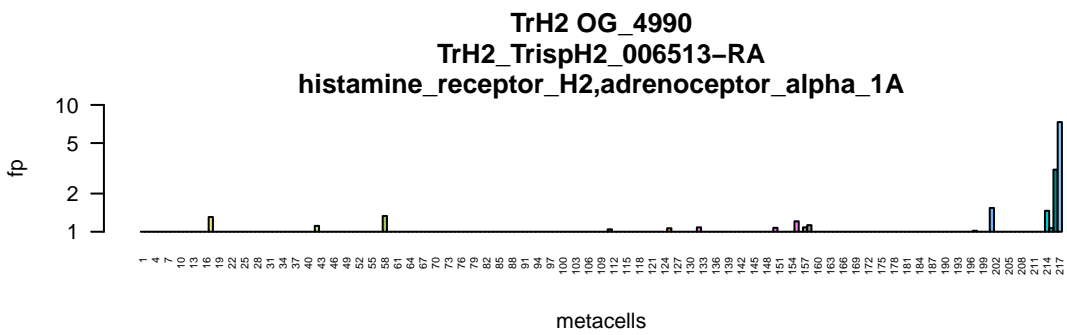
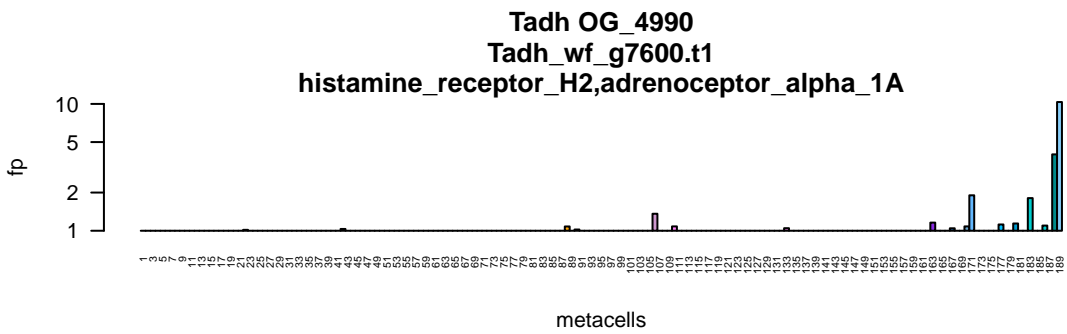
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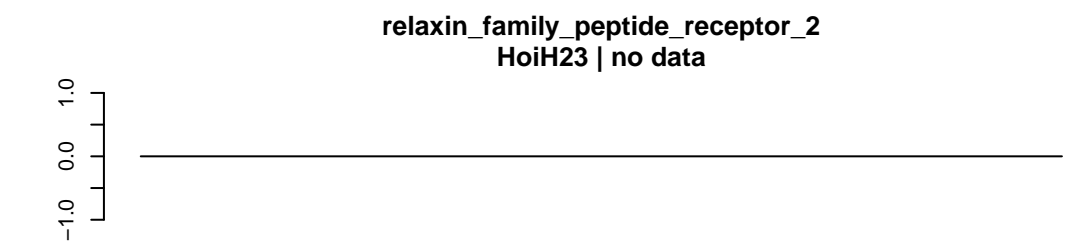
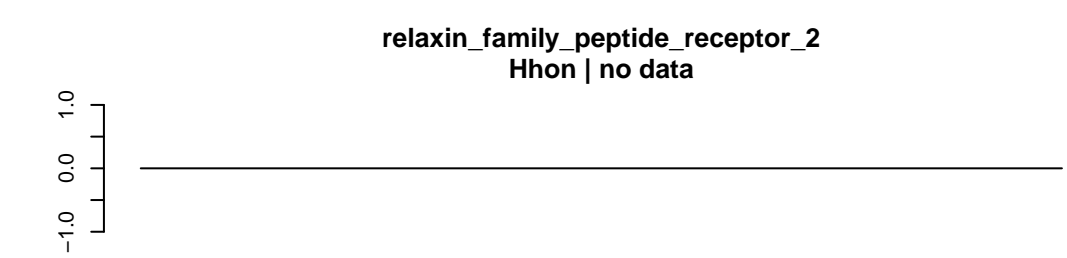
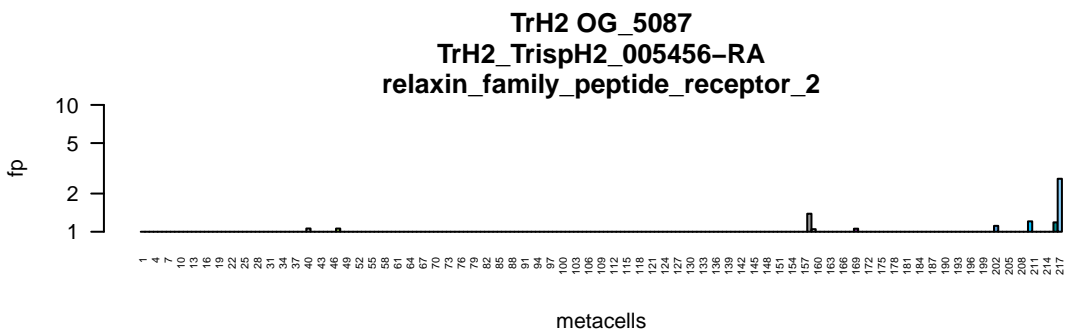
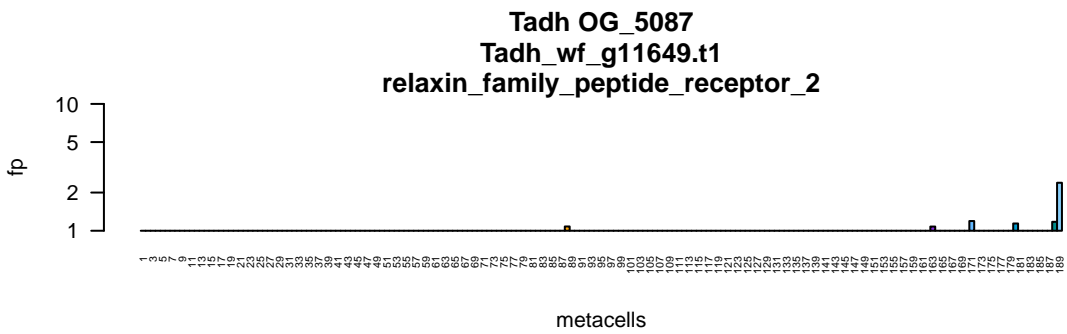


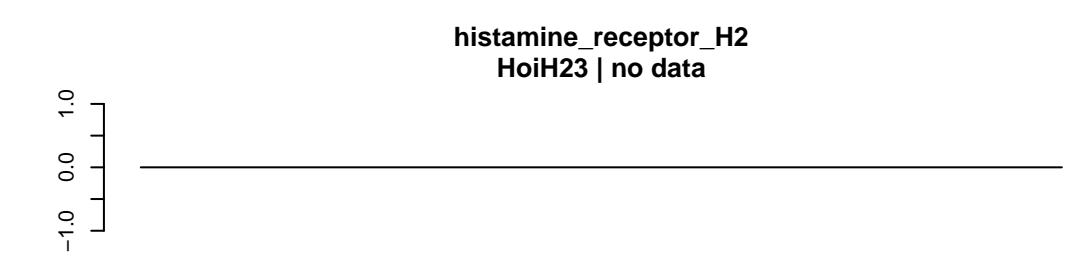
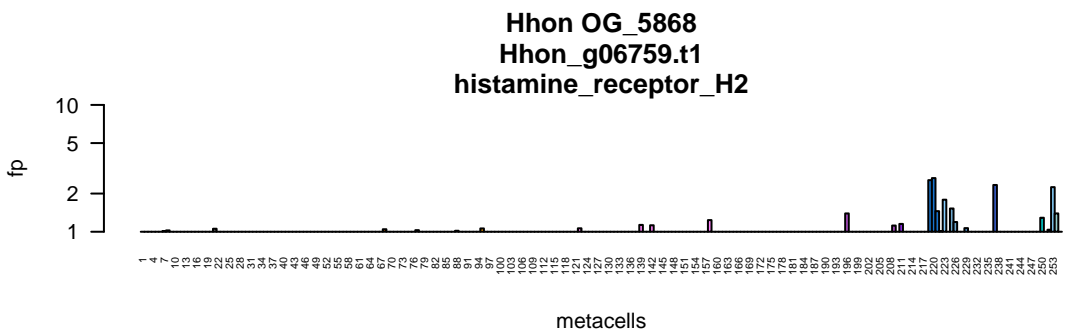
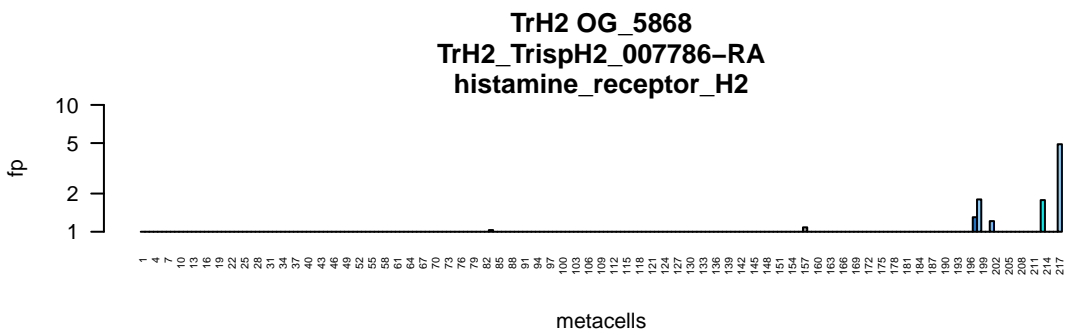
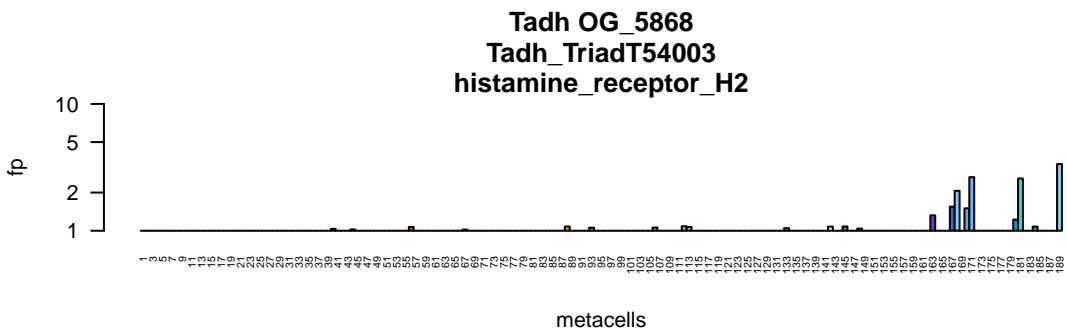
HoiH23 OG_4563
HoiH23_PIH23_009618-RA
adrenoceptor_beta_2,5_hydroxytryptamine_receptor_2A



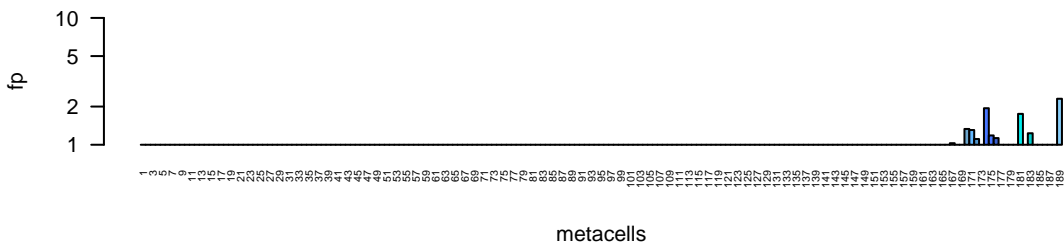




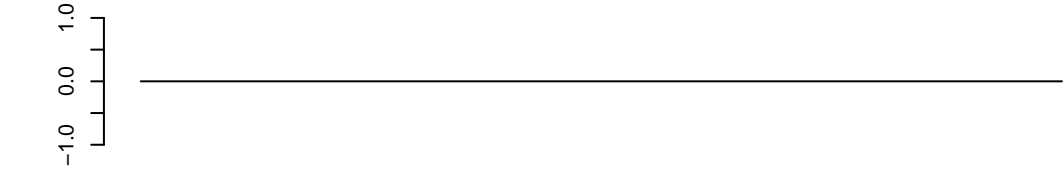




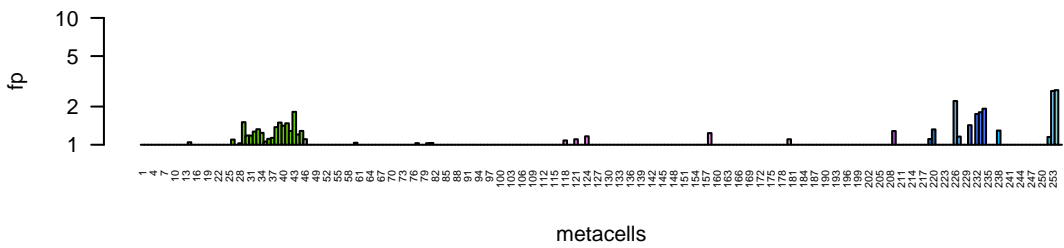
Tadh OG_6481
Tadh_wf_g9558.t1



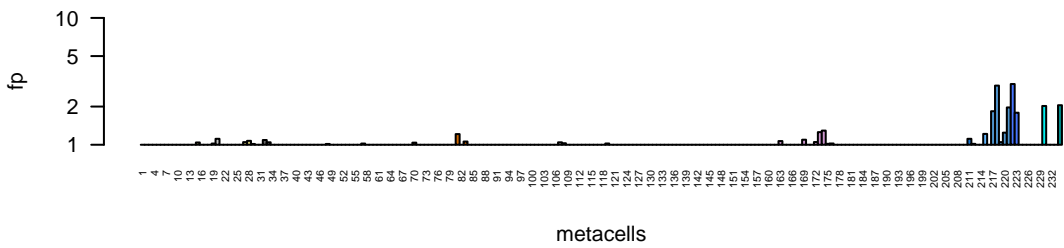
TrH2 | no data

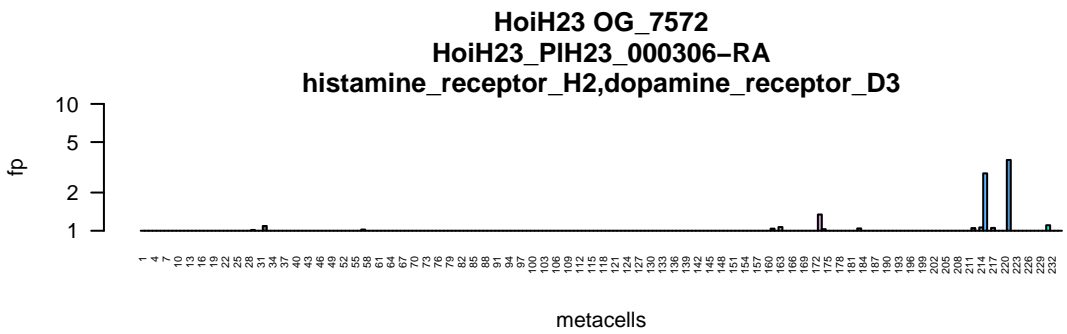
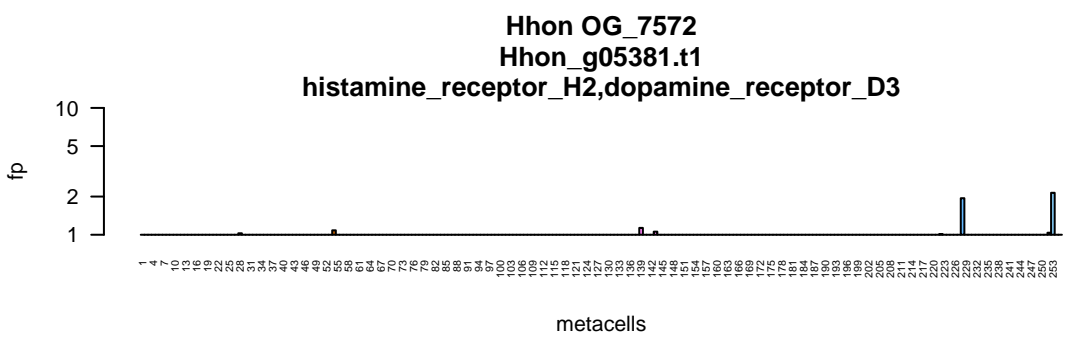
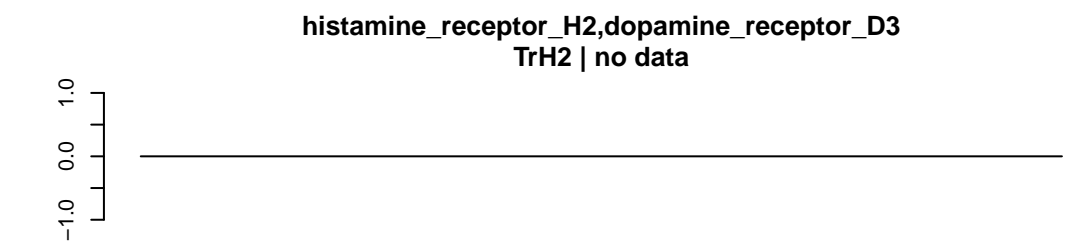
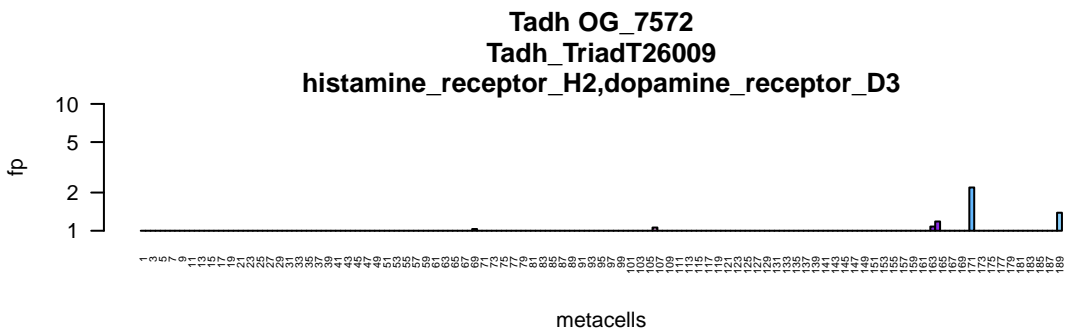


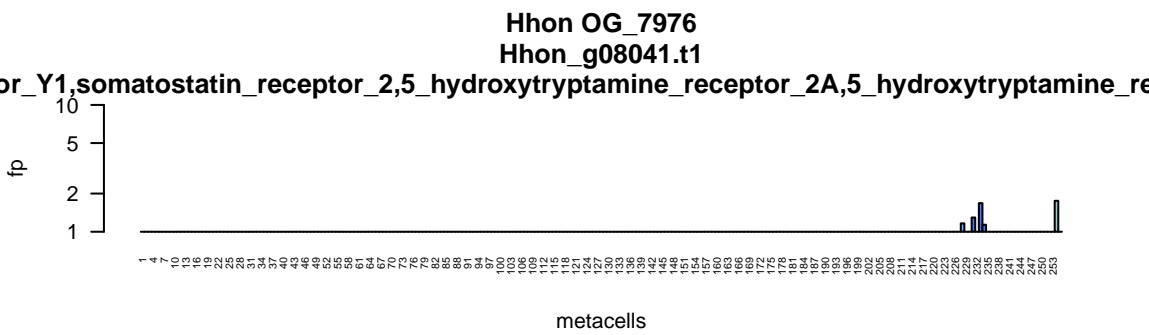
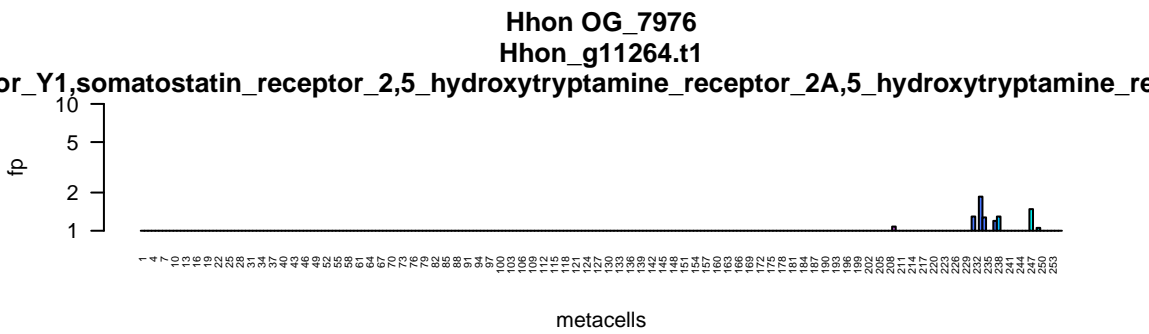
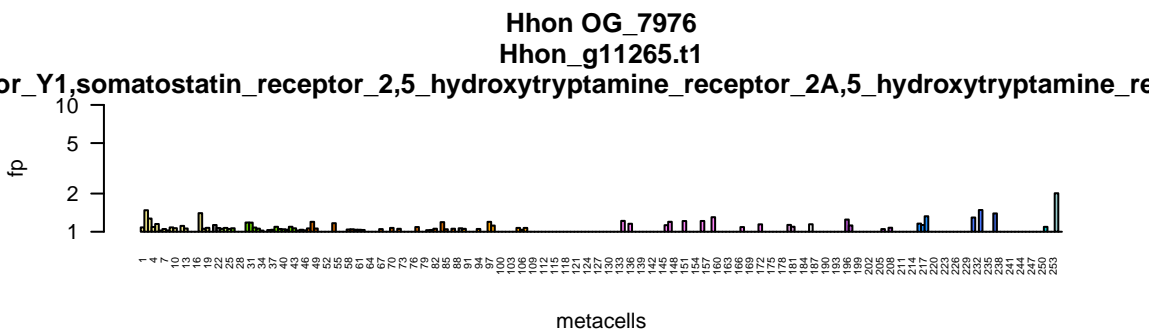
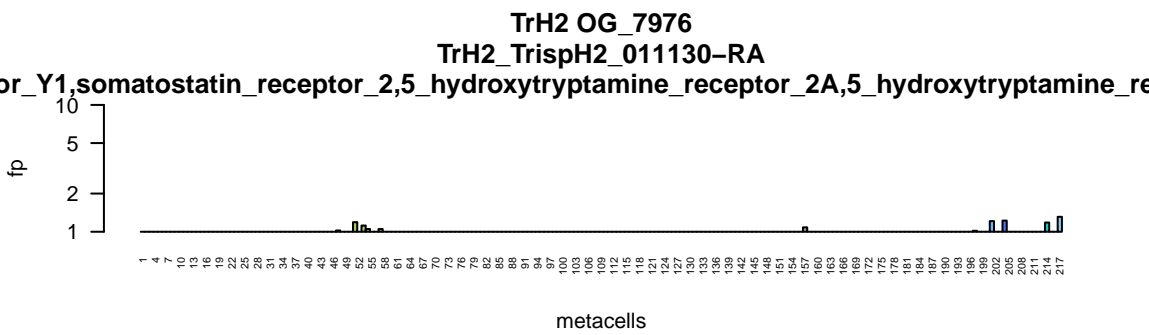
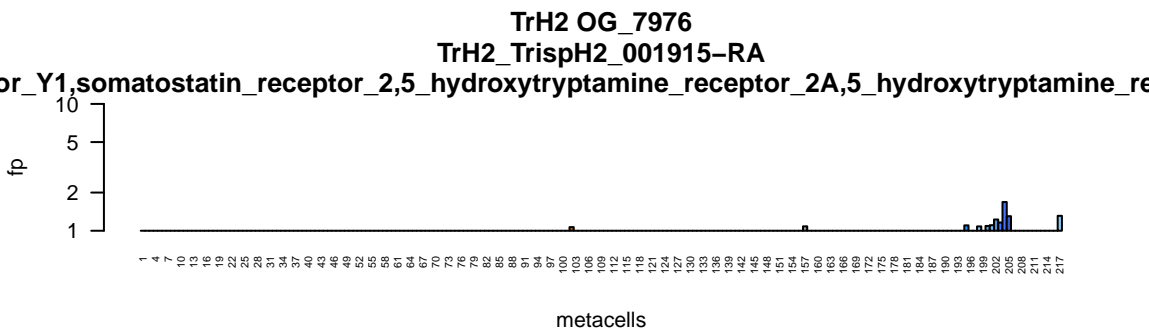
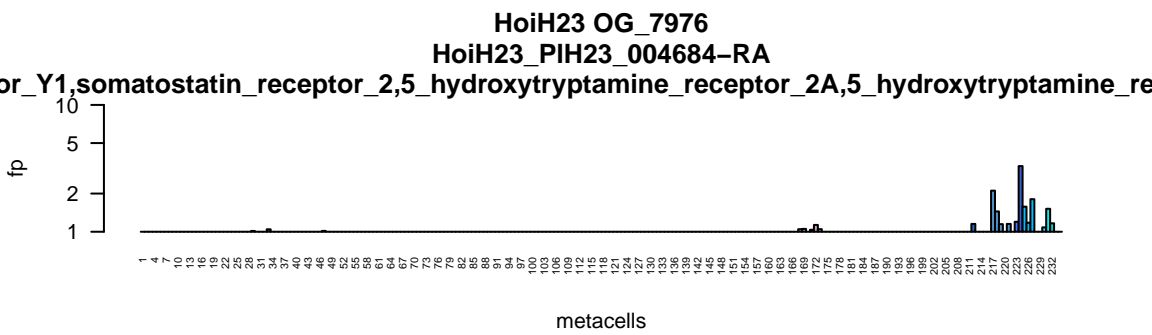
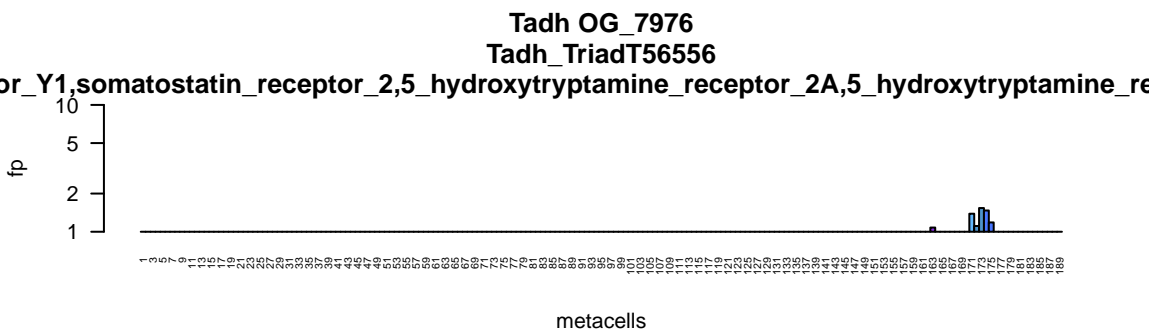
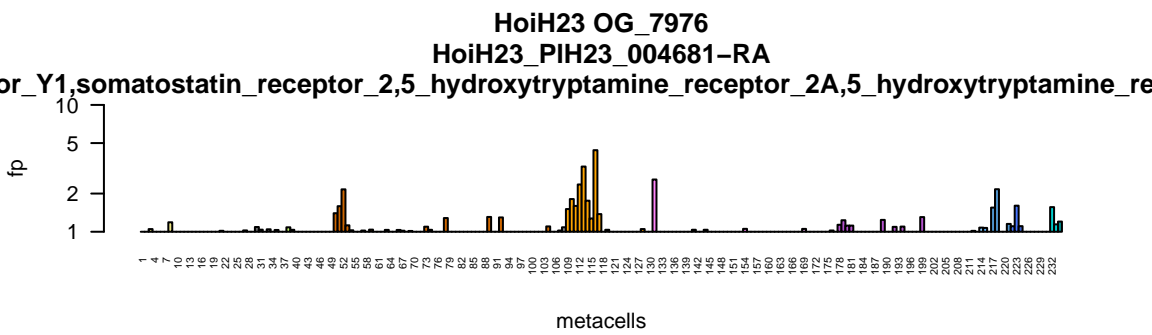
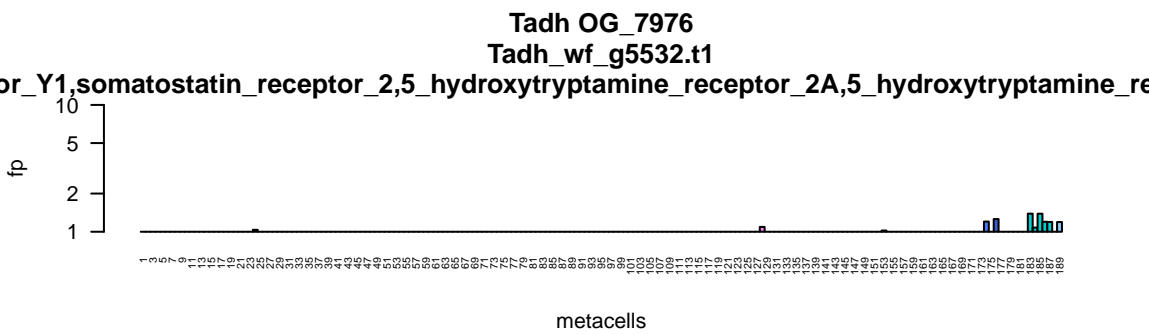
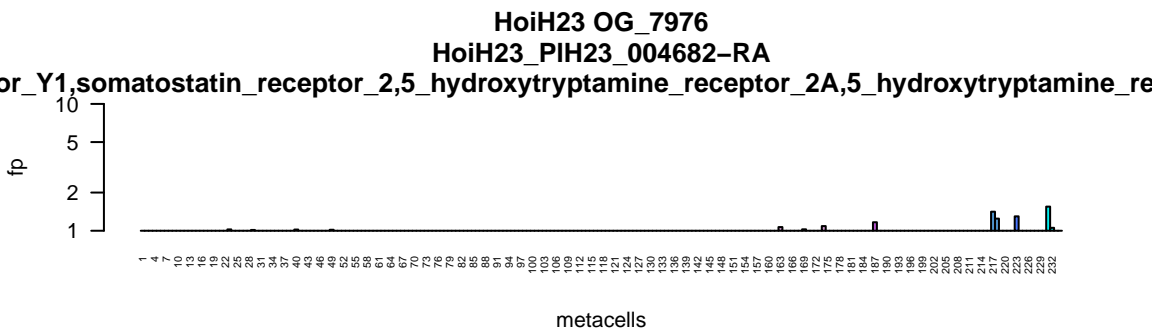
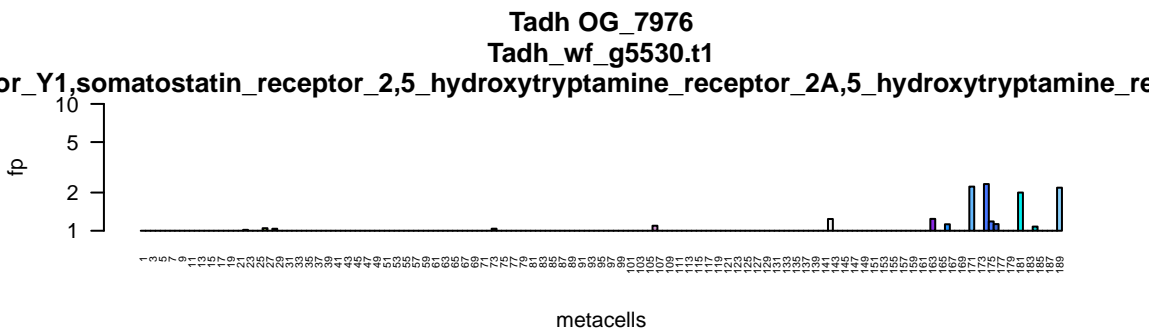
Hhon OG_6481
Hhon_g10968.t1

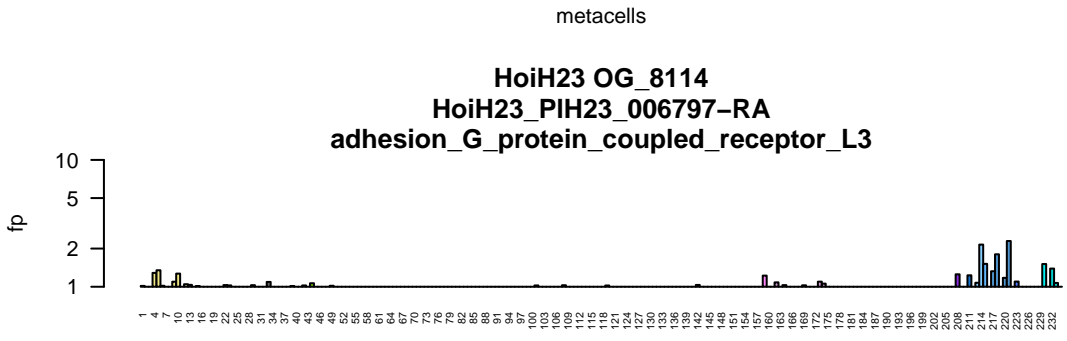
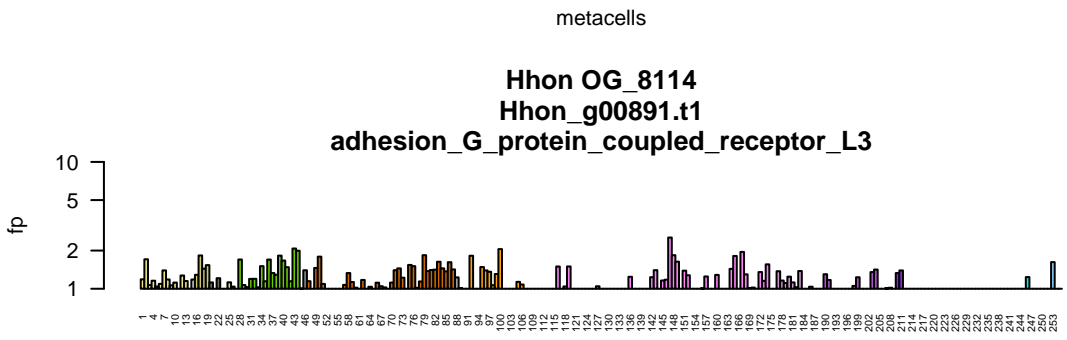
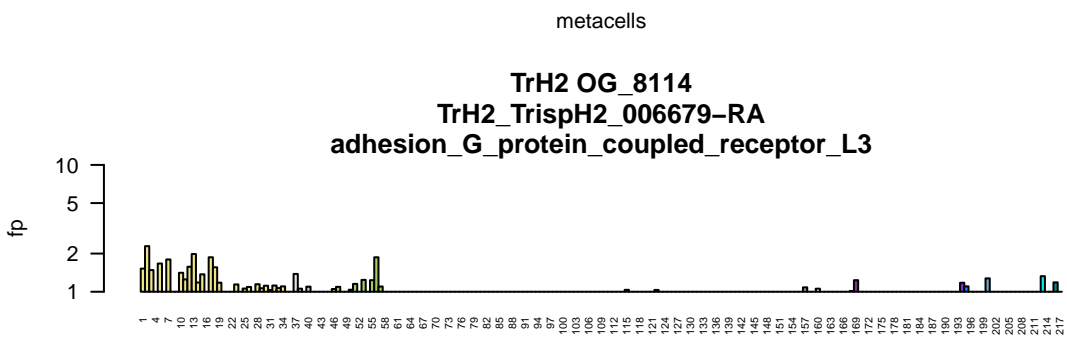
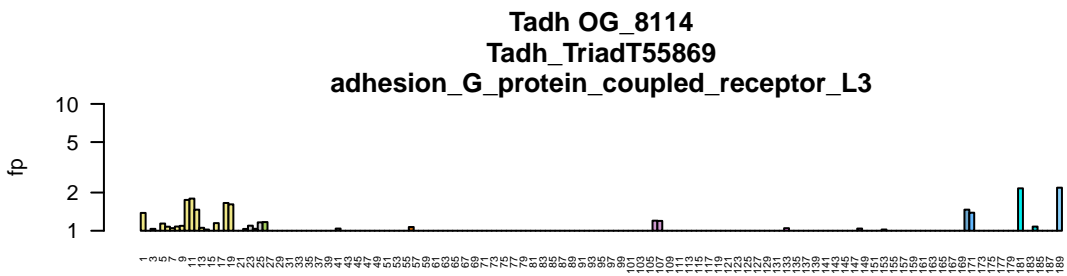


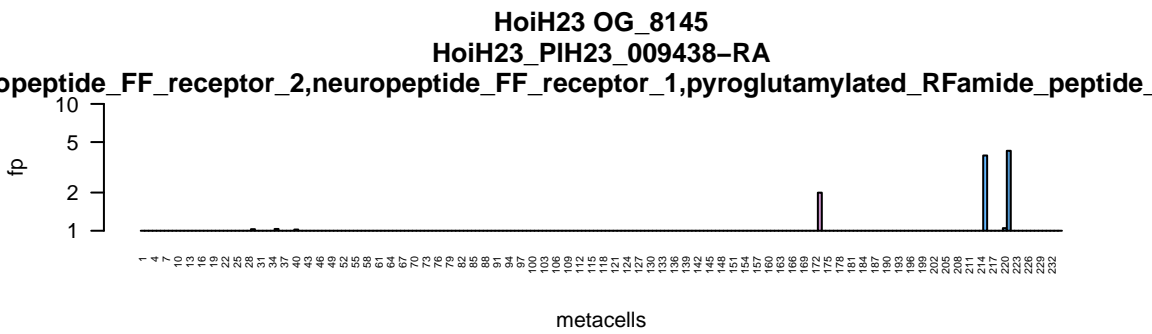
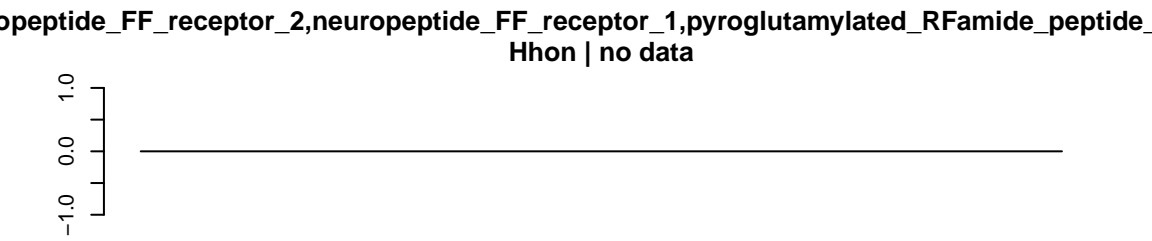
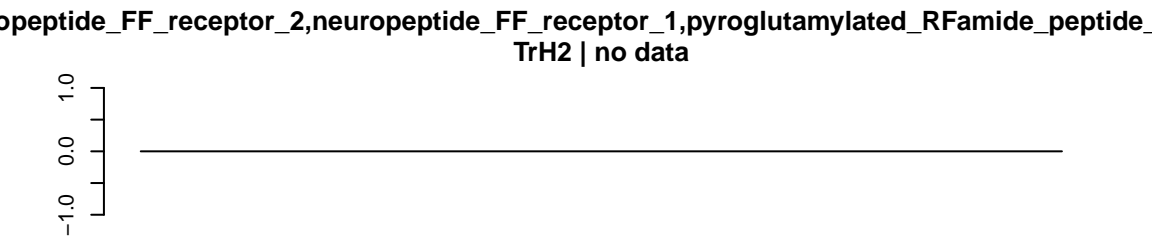
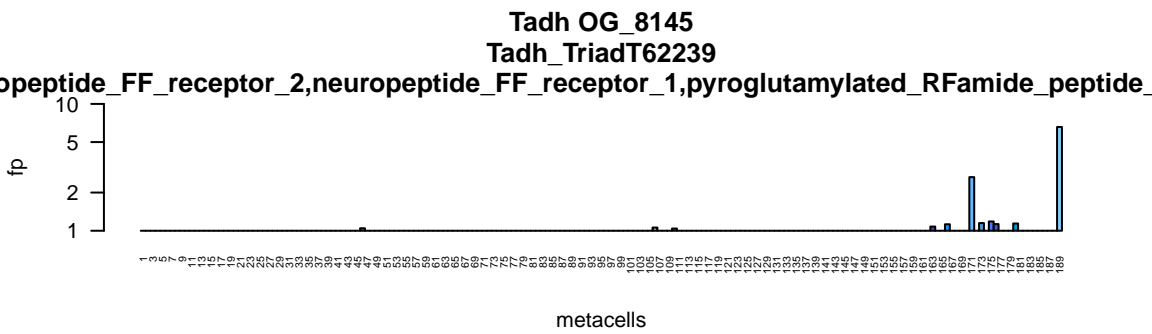
HoiH23 OG_6481
HoiH23_PIH23_011005-RA

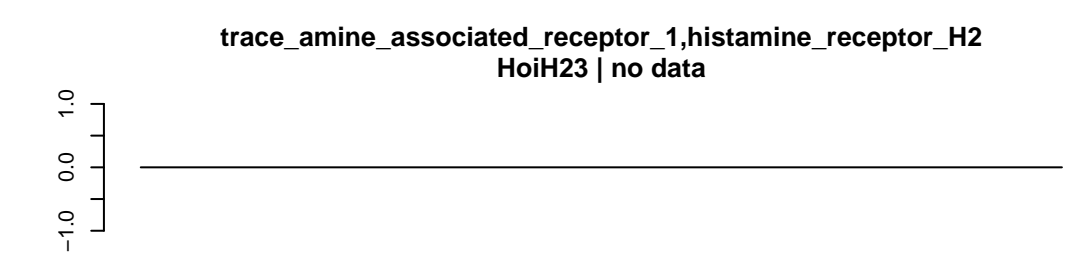
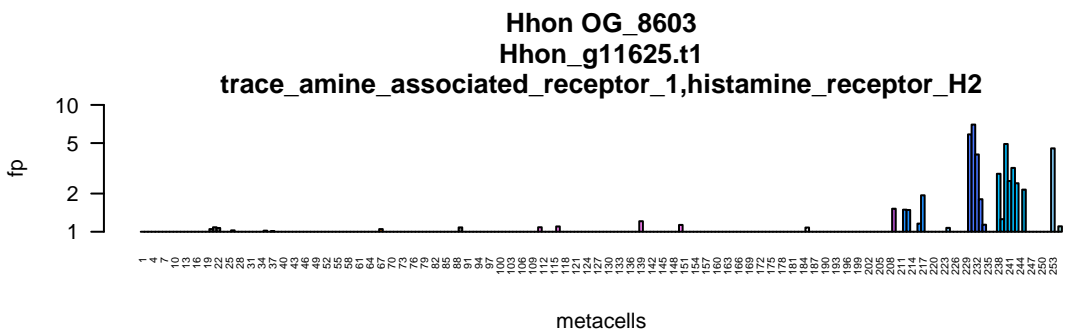
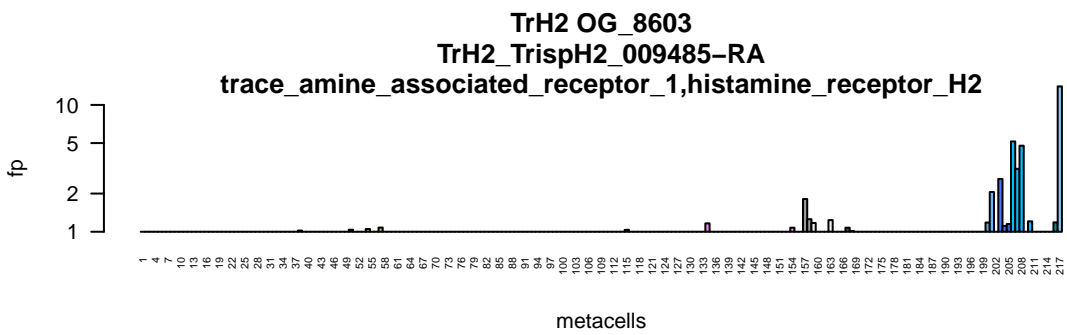
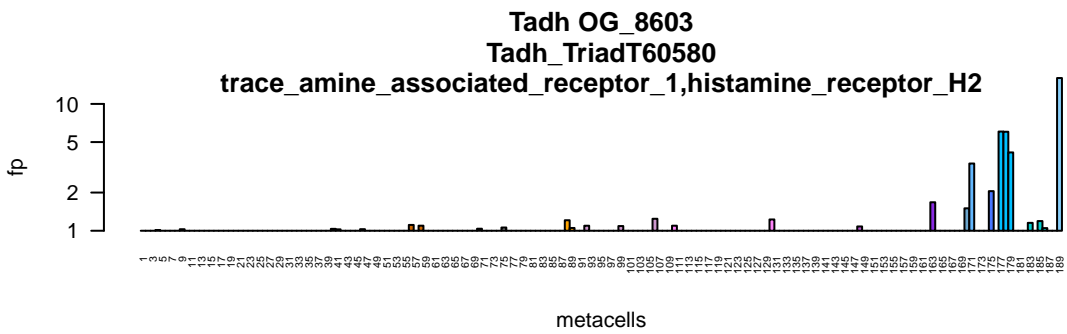


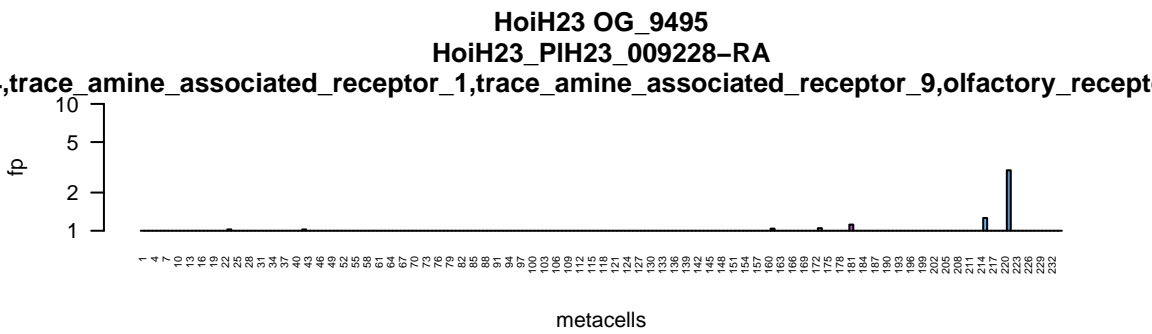
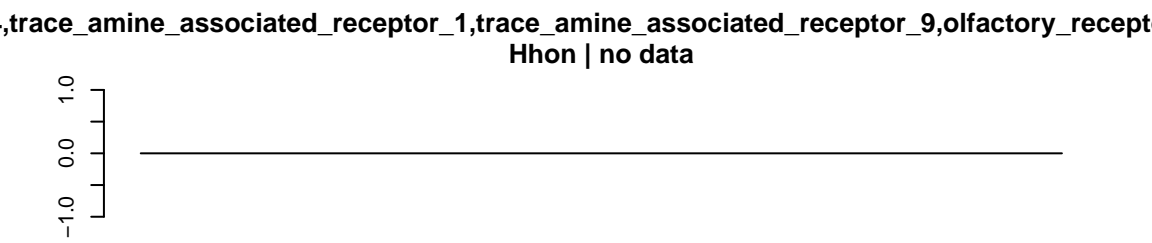
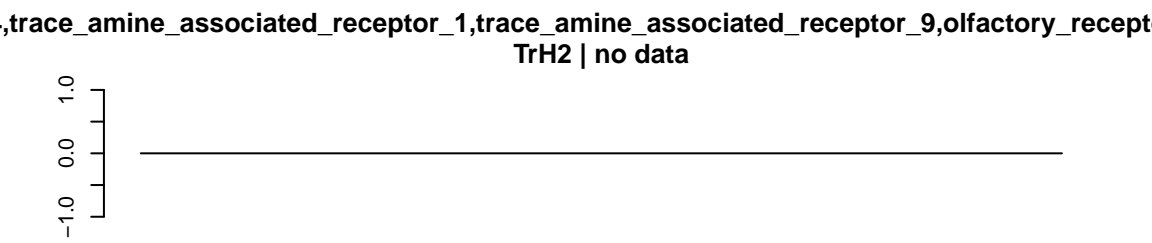
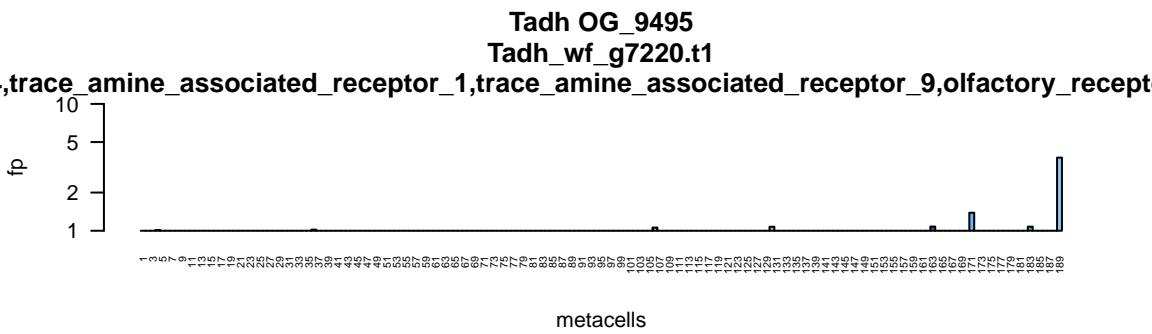


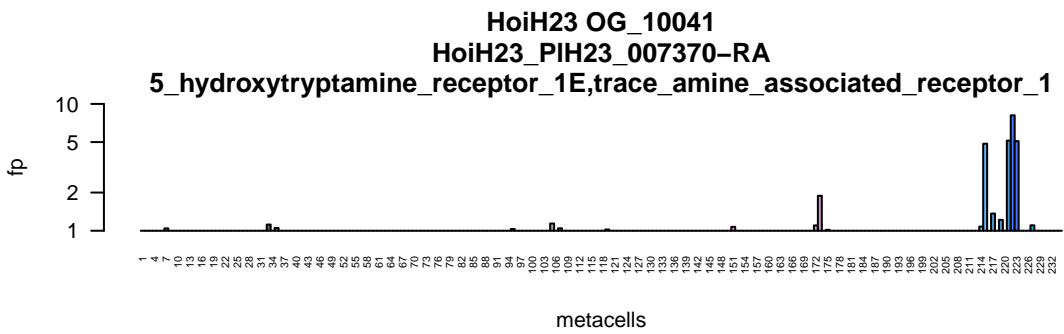
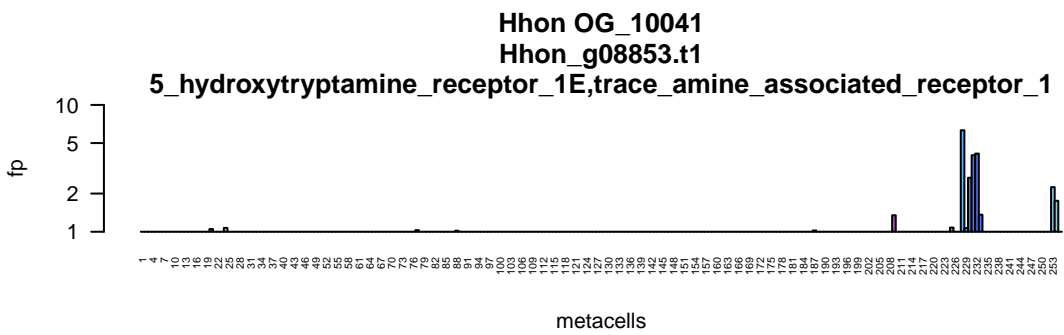
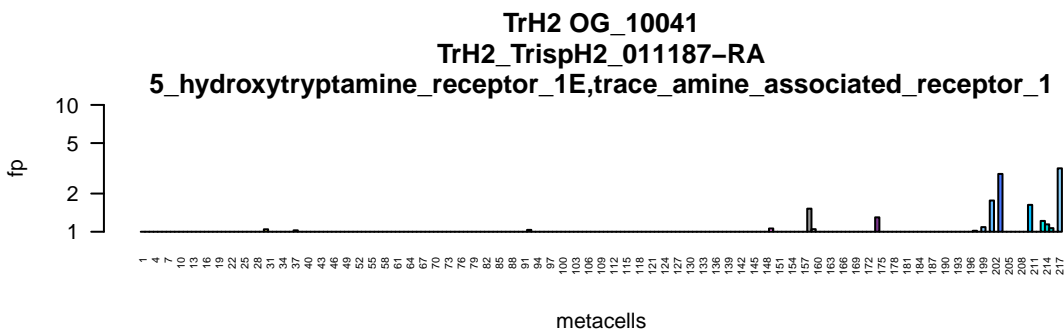
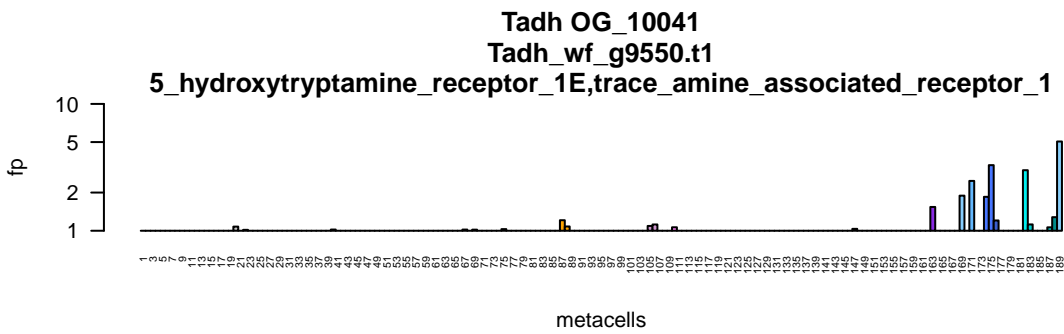




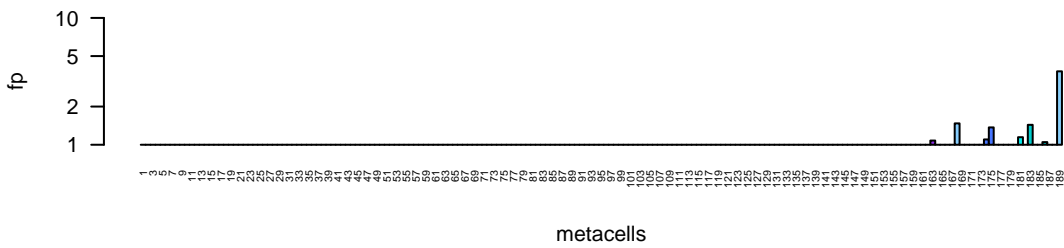




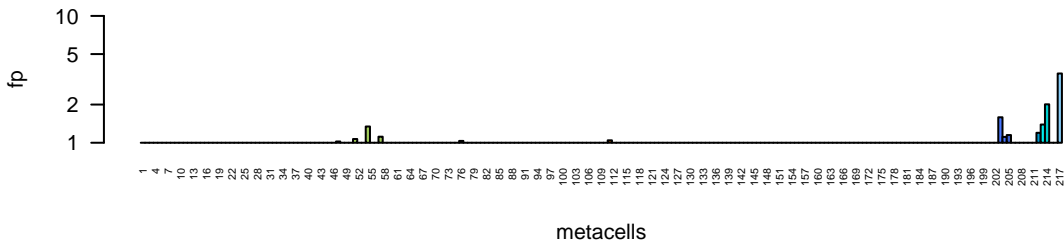




Tadh OG_10104
Tadh_TriadT60048



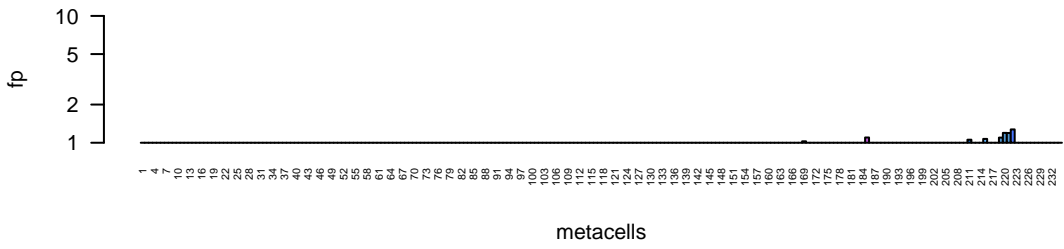
TrH2 OG_10104
TrH2_TrispH2_011300-RA

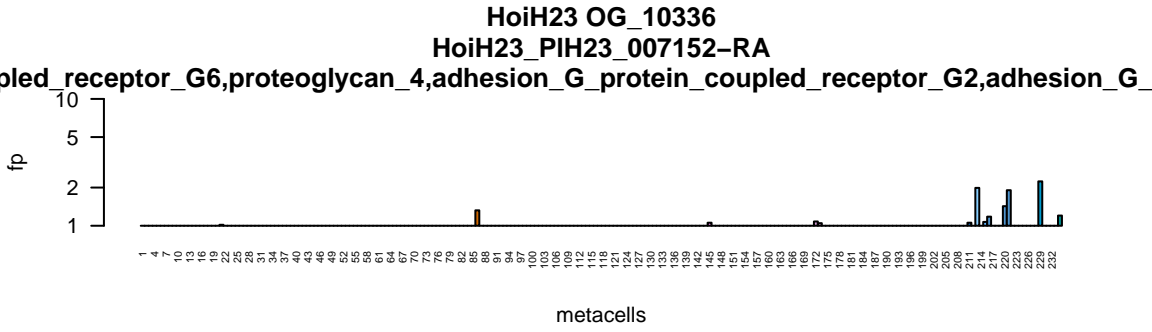
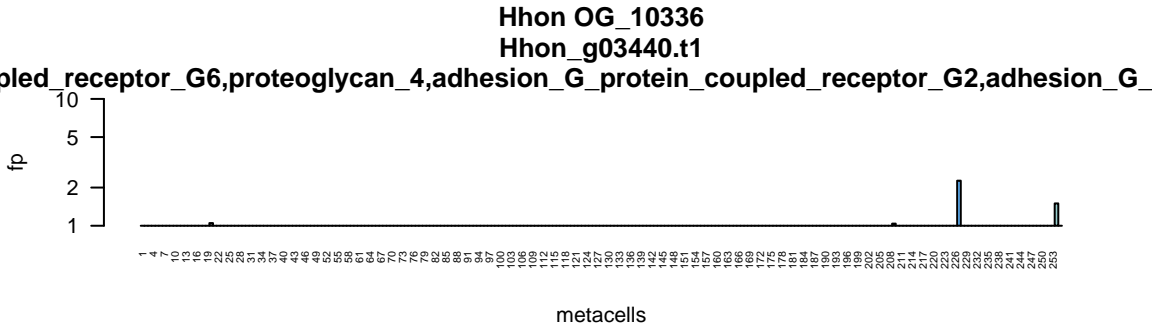
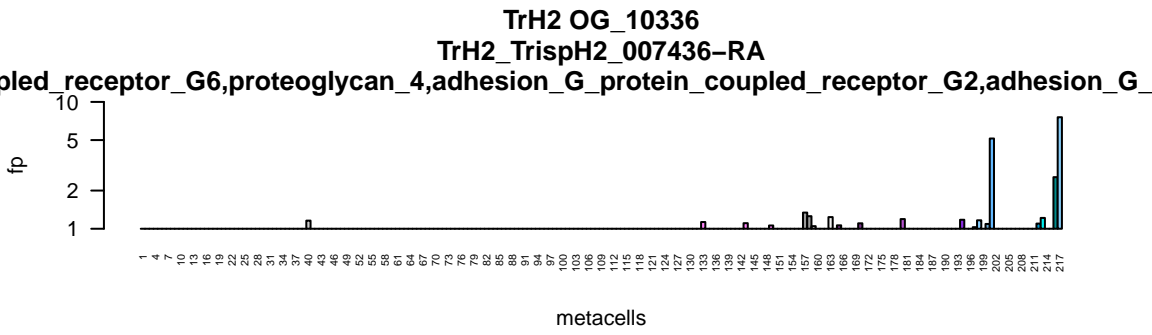
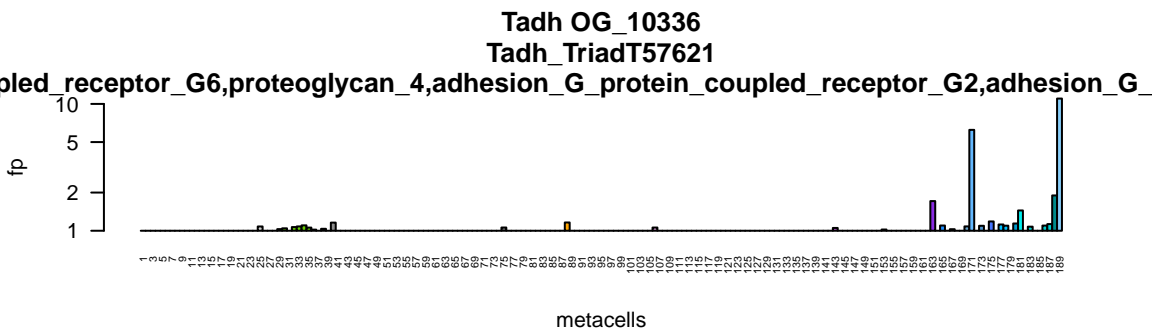


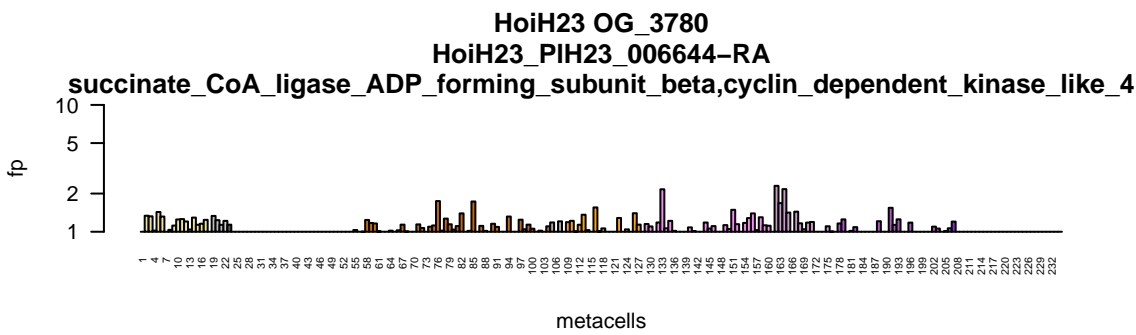
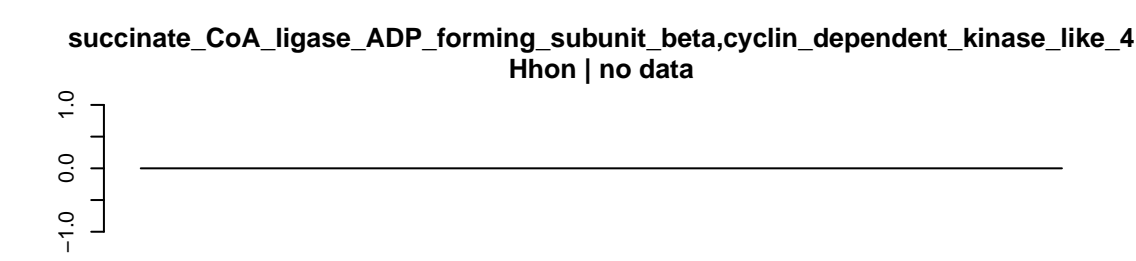
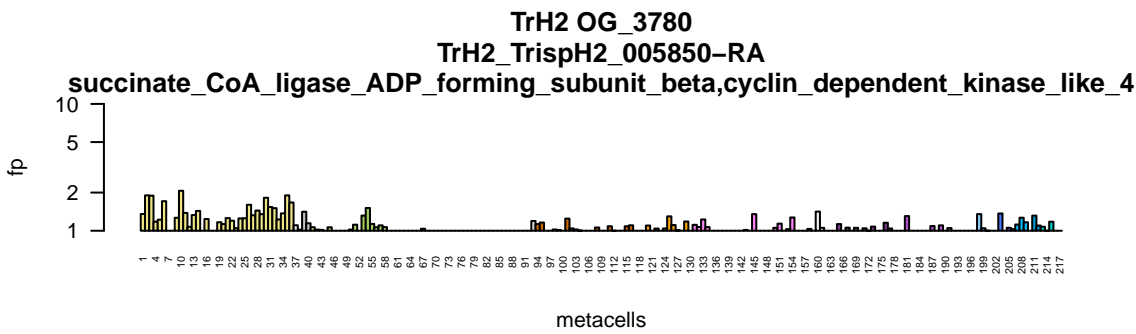
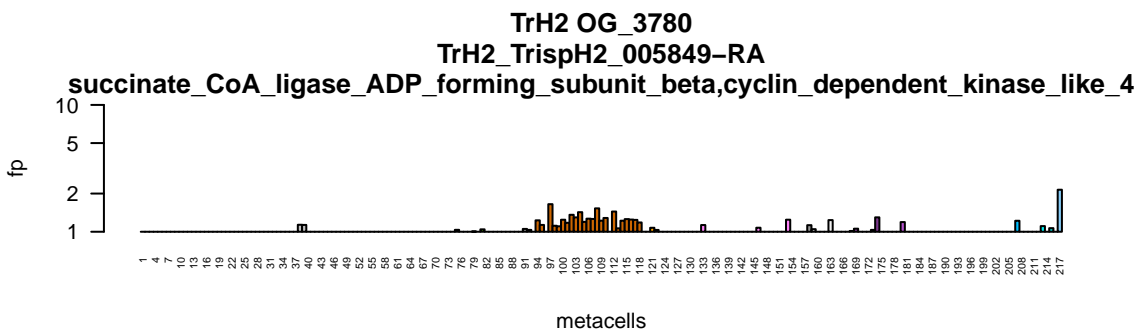
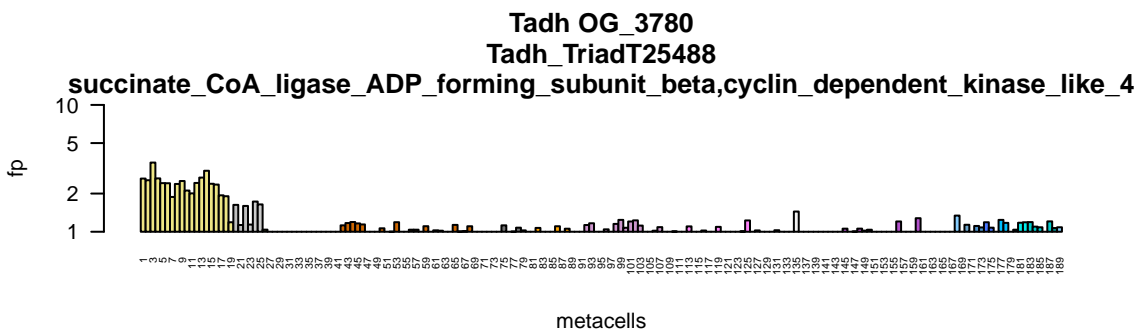
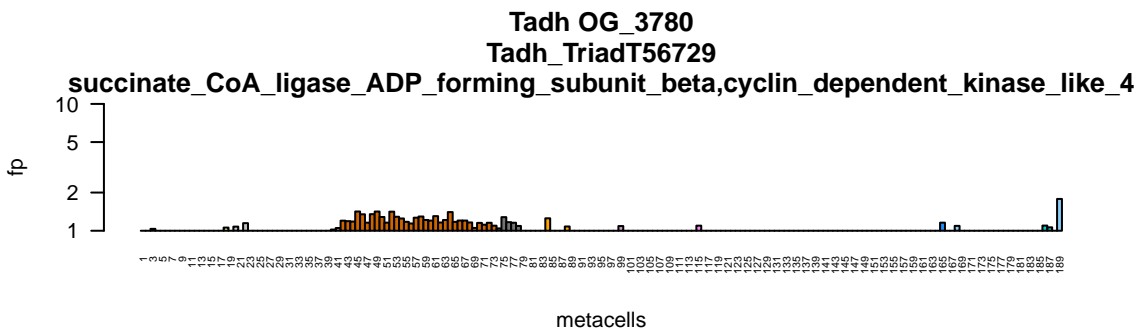
Hhon | no data

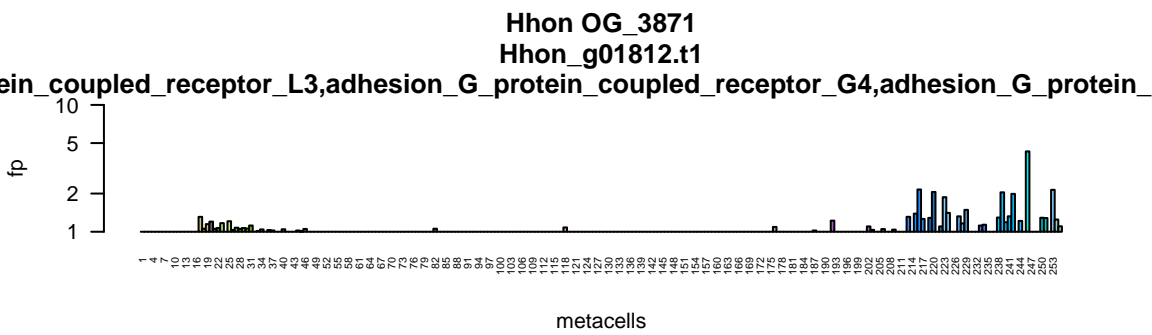
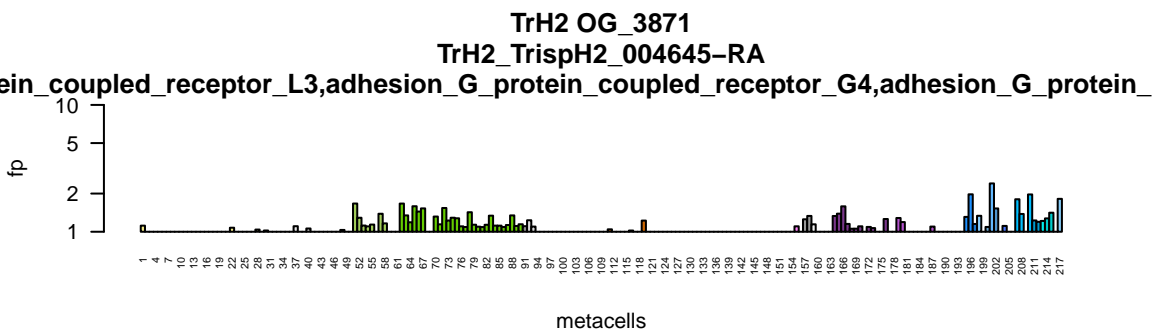
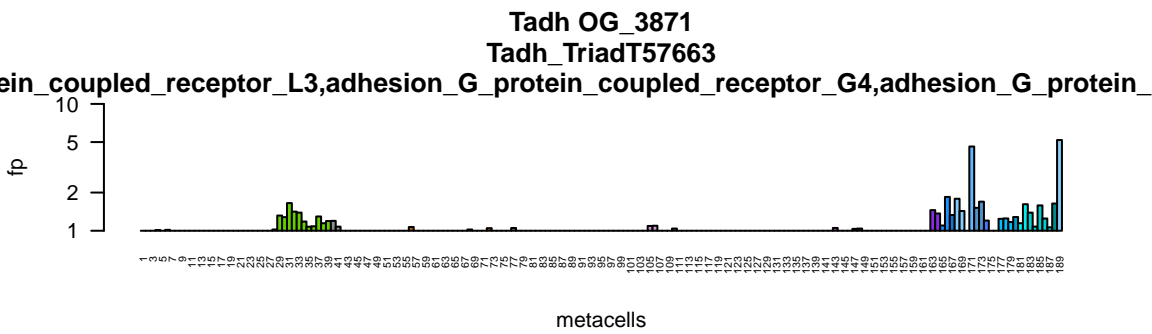


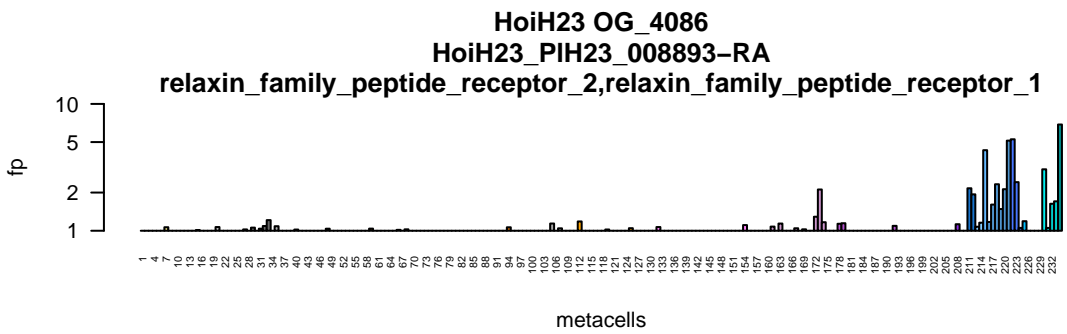
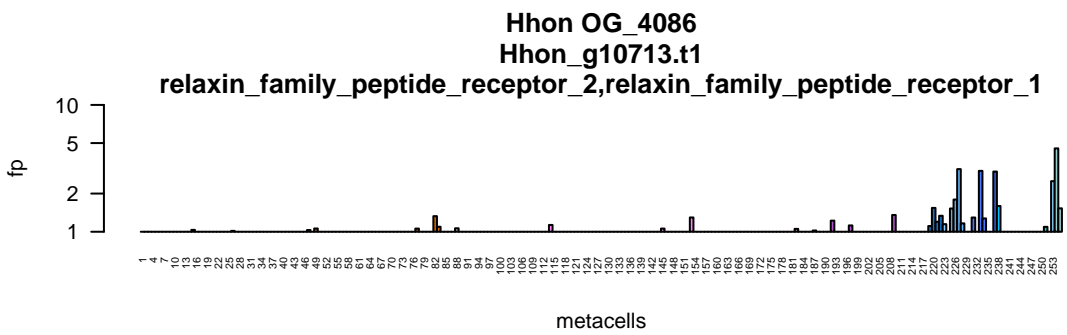
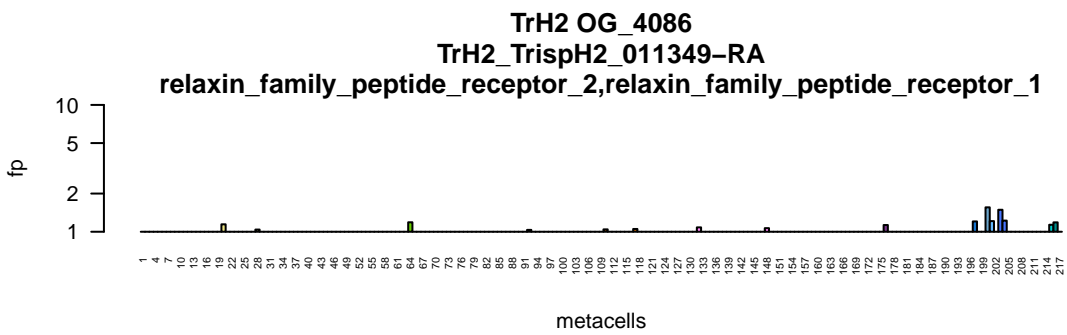
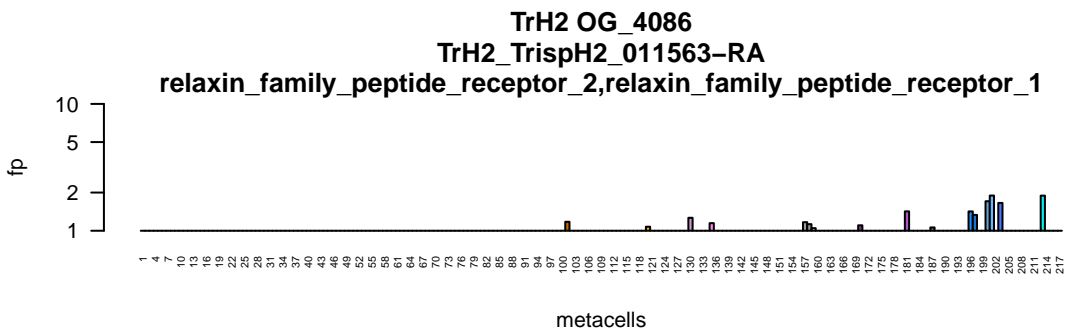
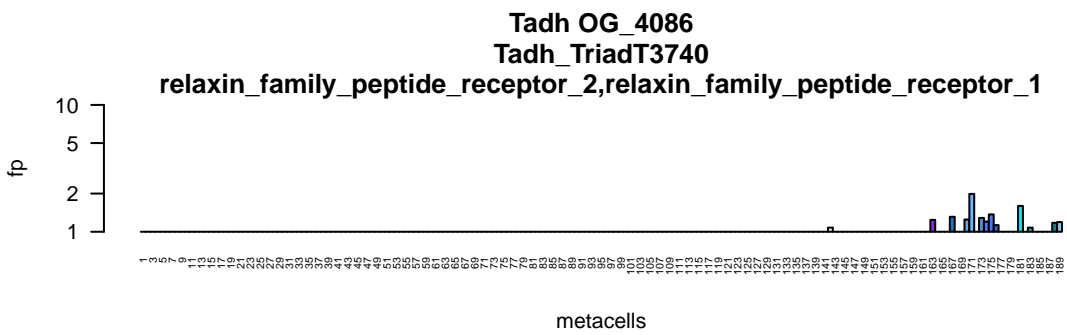
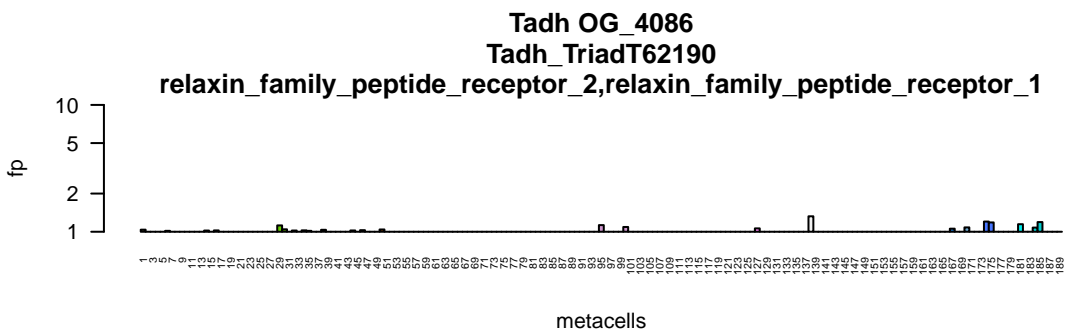
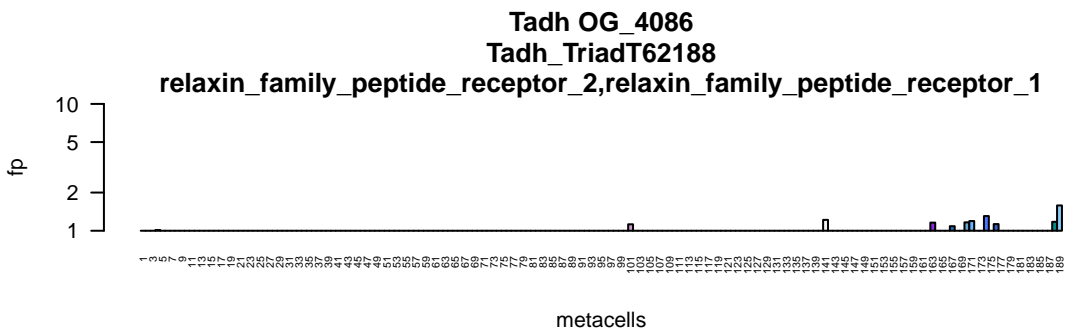
HoiH23 OG_10104
HoiH23_PIH23_006092-RA

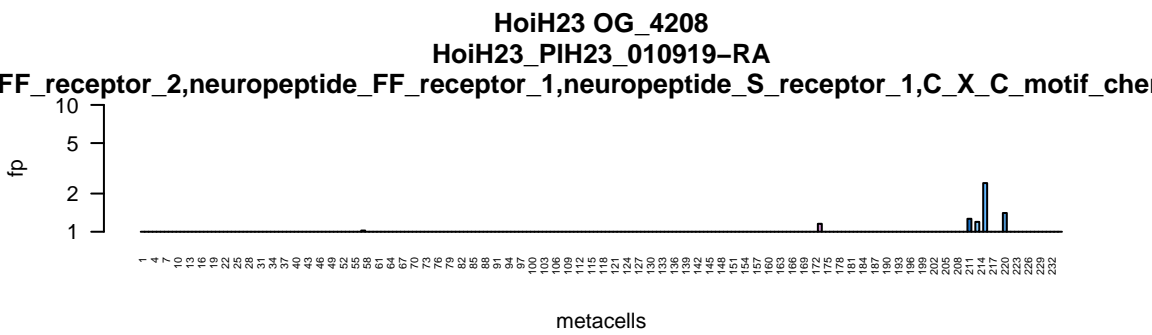
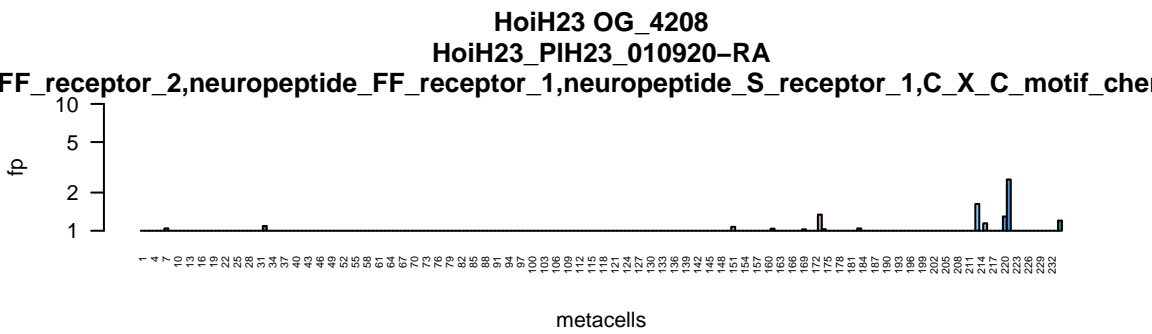
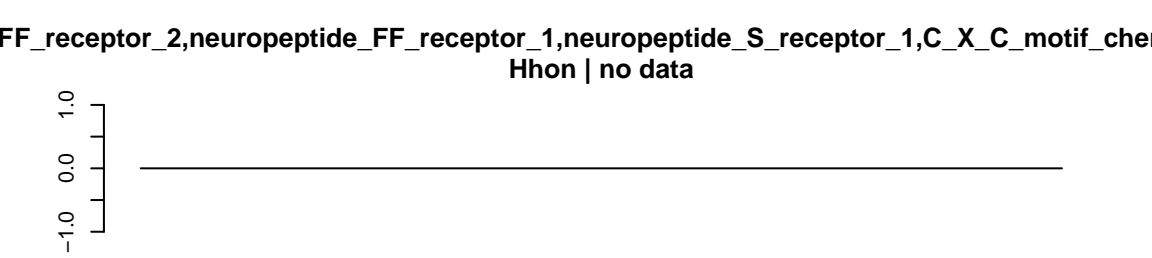
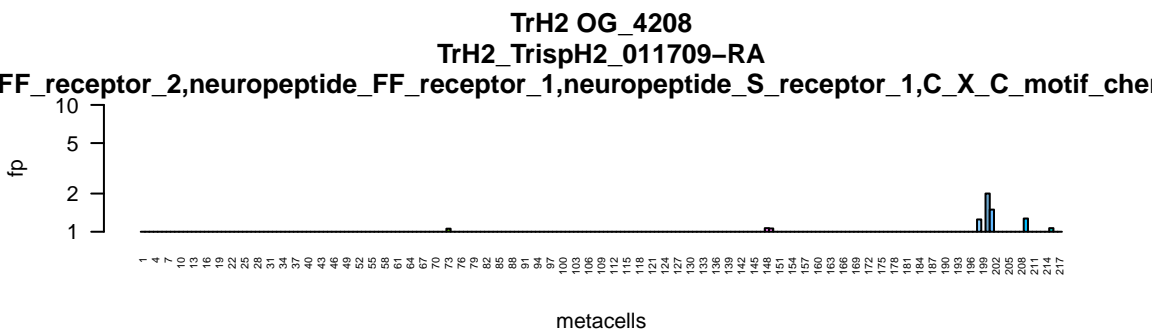
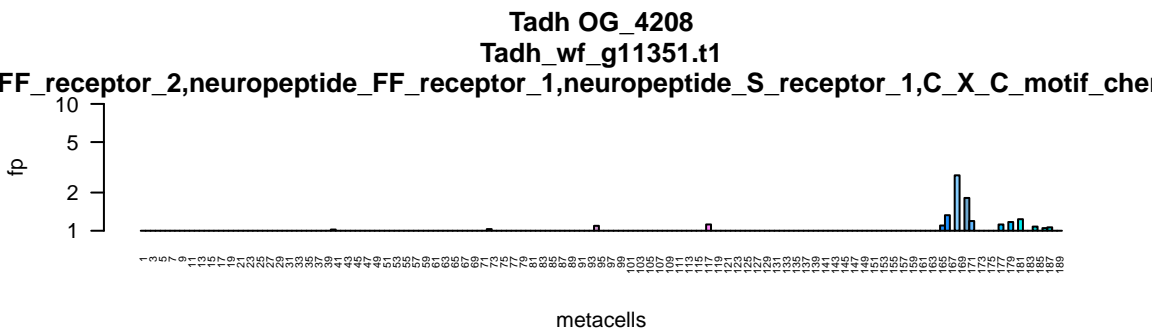












Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists 100 metacells. Most metacells have 0 false positives, but a few have 1, and a small cluster of metacells (173-187) has 2, 3, or 4 false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

metacell	fp
1	0
2	0
3	0
4	0
5	0
6	0
7	0
8	0
9	0
10	0
11	0
12	0
13	0
14	0
15	0
16	0
17	0
18	0
19	0
20	0
21	0
22	0
23	0
24	0
25	0
26	0
27	0
28	0
29	0
30	0
31	0
32	0
33	0
34	0
35	0
36	0
37	0
38	0
39	0
40	0
41	0
42	0
43	0
44	0
45	0
46	0
47	0
48	0
49	0
50	0
51	0
52	0
53	0
54	0
55	0
56	0
57	0
58	0
59	0
60	0
61	0
62	0
63	0
64	0
65	0
66	0
67	0
68	0
69	0
70	0
71	0
72	0
73	0
74	0
75	0
76	0
77	0
78	0
79	0
80	0
81	0
82	0
83	0
84	0
85	0
86	0
87	0
88	0
89	0
90	0
91	0
92	0
93	0
94	0
95	0
96	0
97	0
98	0
99	0
100	0
101	0
102	0
103	0
104	0
105	0
106	0
107	0
108	0
109	0
110	0
111	0
112	0
113	0
114	0
115	0
116	0
117	0
118	0
119	0
120	0
121	0
122	0
123	0
124	0
125	0
126	0
127	0
128	0
129	0
130	0
131	0
132	0
133	0
134	0
135	0
136	0
137	0
138	0
139	0
140	0
141	0
142	0
143	0
144	0
145	0
146	0
147	0
148	0
149	0
150	0
151	0
152	0
153	0
154	0
155	0
156	0
157	0
158	0
159	0
160	0
161	0
162	0
163	0
164	0
165	0
166	0
167	0
168	0
169	0
170	0
171	0
172	0
173	1
174	2
175	2
176	3
177	4
178	3
179	2
180	1
181	1
182	1
183	1
184	1
185	1
186	1
187	1
188	1
189	1
190	1
191	1
192	1
193	1
194	1
195	1
196	1
197	1
198	1
199	1
200	1

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
163	2
167	3
171	2
173	4
177	5
185	2
187	2

Bar chart showing the frequency of metacells. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists 100 metacells. Most metacells have a frequency of 1, with a few having frequencies of 2 or 3.

metacell	fp
1	1
2	1
3	1
4	1
5	1
6	1
7	1
8	1
9	1
10	1
11	1
12	1
13	1
14	1
15	1
16	1
17	1
18	1
19	1
20	1
21	1
22	1
23	1
24	1
25	1
26	1
27	1
28	1
29	1
30	1
31	1
32	1
33	1
34	1
35	1
36	1
37	1
38	1
39	1
40	1
41	1
42	1
43	1
44	1
45	1
46	1
47	1
48	1
49	1
50	1
51	1
52	1
53	1
54	1
55	1
56	1
57	1
58	1
59	1
60	1
61	1
62	1
63	1
64	1
65	1
66	1
67	1
68	1
69	1
70	1
71	1
72	1
73	1
74	1
75	1
76	1
77	1
78	1
79	1
80	1
81	1
82	1
83	1
84	1
85	1
86	1
87	1
88	1
89	1
90	1
91	1
92	1
93	1
94	1
95	1
96	1
97	1
98	1
99	1
100	1

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists 48 metacells. Most metacells have a false positive count of 1, with a few having 2 or 3. Metacells 199, 202, 203, 208, and 214 show higher counts, up to 4.

metacell	fp
1	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	2
199	2
202	1
203	4
206	2
208	3
211	2
214	1

Bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and the y-axis is labeled 'fp'. The chart shows a very high frequency for metacell 1 (fp=10) and a few other metacells with low frequencies (fp=1 or 2).

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

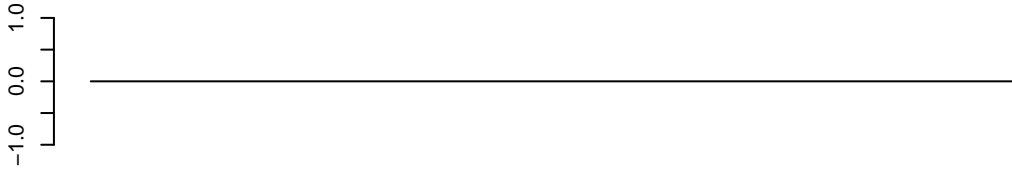
metacell	fp
1	1
4	1
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	0
34	0
37	0
40	0
43	0
46	0
49	0
52	0
55	0
58	0
61	0
64	0
67	0
70	0
73	0
76	1
78	1
79	1
82	1
85	0
88	0
91	1
94	1
97	0
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	0
130	0
133	0
136	0
139	0
142	0
145	1
148	1
151	0
154	1
157	1
160	1
163	1
166	0
169	1
172	1
175	1
178	0
181	0
184	0
187	0
190	0
193	0
196	0
199	1
202	4
203	5
208	3
211	1
214	1

metacell	fp
1	1
2	1
3	1
4	1
5	1
6	1
7	1
8	1
9	1
10	1
11	1
12	1
13	1
14	1
15	1
16	1
17	1
18	1
19	1
20	1
21	1
22	1
23	1
24	1
25	1
26	1
27	1
28	1
29	1
30	1
31	1
32	1
33	1
34	1
35	1
36	1
37	1
38	1
39	1
40	1
41	1
42	1
43	1
44	1
45	1
46	1
47	1
48	1
49	1
50	1
51	1
52	1
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55	1
56	1
57	1
58	1
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60	1
61	1
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75	1
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78	1
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80	1
81	1
82	1
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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
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103	1
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106	1
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113	1
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117	1
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119	1
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122	1
123	1
124	1
125	1
126	1
127	1
128	1
129	1
130	1
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141	1
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150	1
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152	1
153	1
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157	1
158	1
159	1
160	1
161	1
162	1
163	1
164	1
165	1
166	1
167	1
168	1
169	1
170	1
171	1
172	1
173	1
174	1
175	1
176	1
177	1
178	1
179	1
180	1
181	1
182	1
183	1
184	1
185	1
186	1
187	1
188	1
189	1
190	1
191	1
192	1
193	1
194	1
195	1
196	1
197	1
198	1
199	1
200	1
201	1
202	1
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208	1
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211	1
212	1
213	1

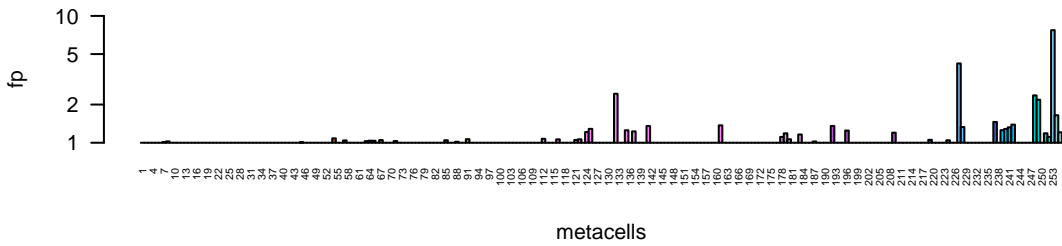
Tadh | no data



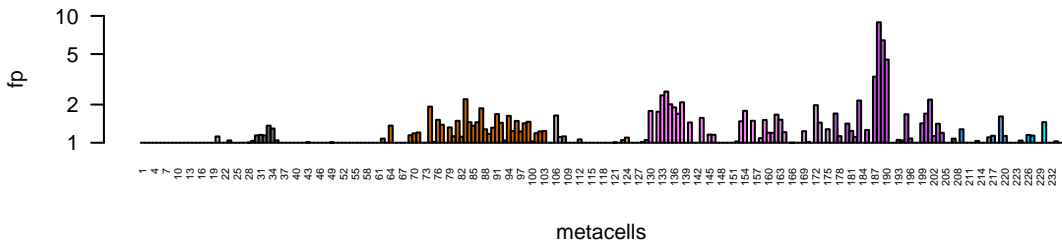
TrH2 | no data



Hhon OG_4260
Hhon_g04196.t1



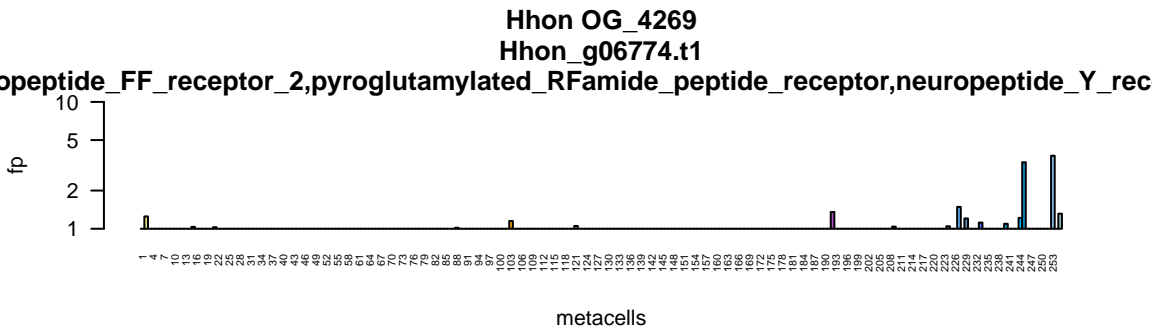
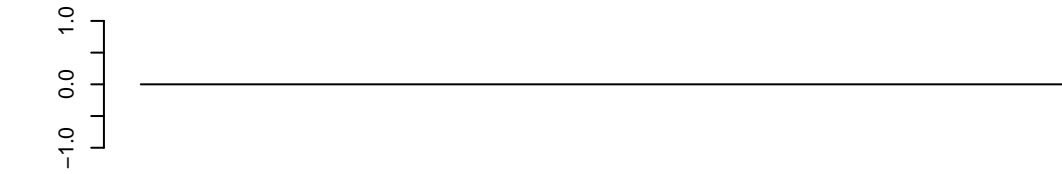
HoiH23 OG_4260
HoiH23_PIH23_009314-RA



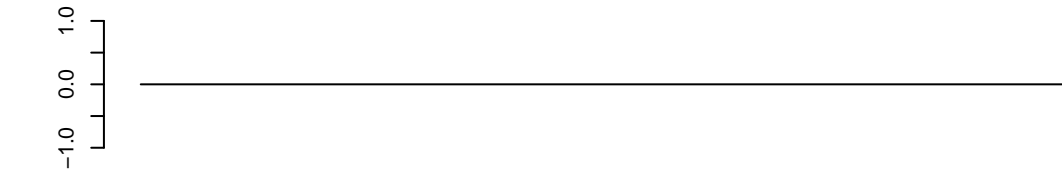
peptide_FF_receptor_2,pyroglutamylated_RFamide_peptide_receptor,neuropeptide_Y_rec
Tadh | no data

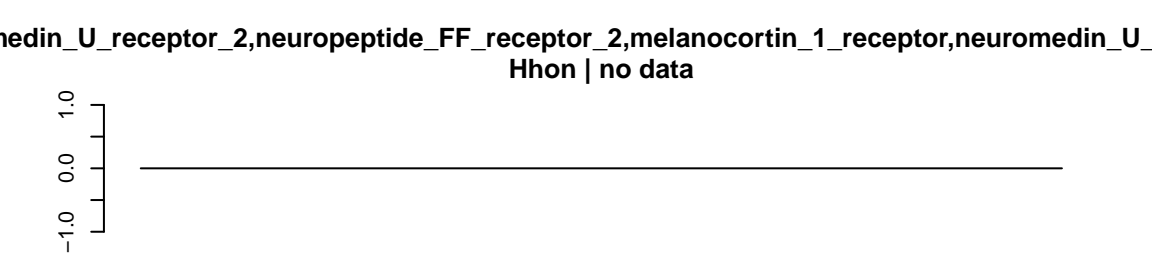
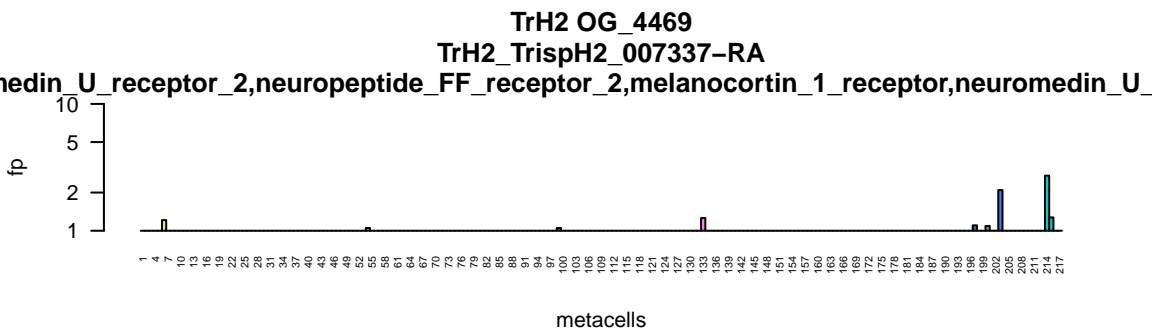
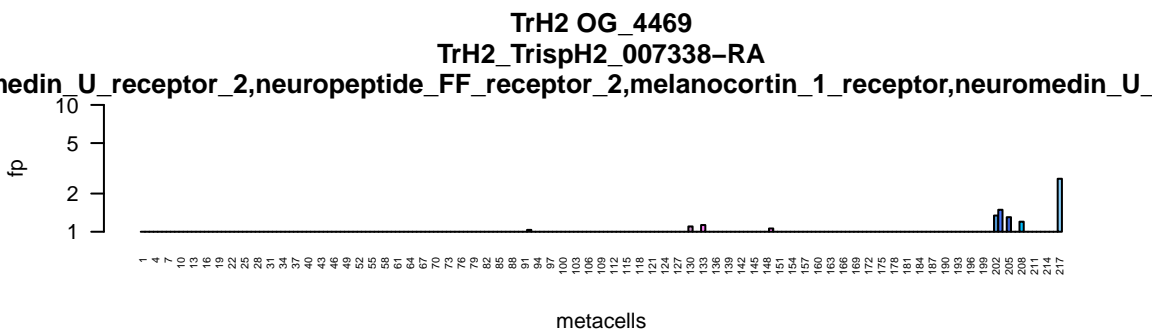
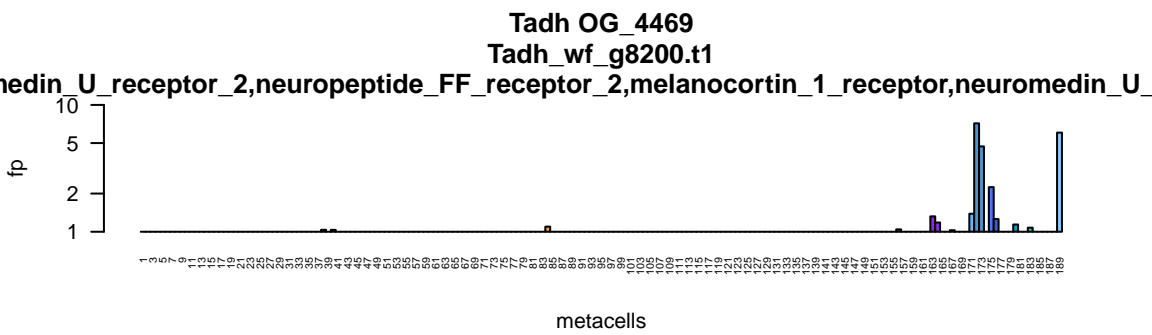
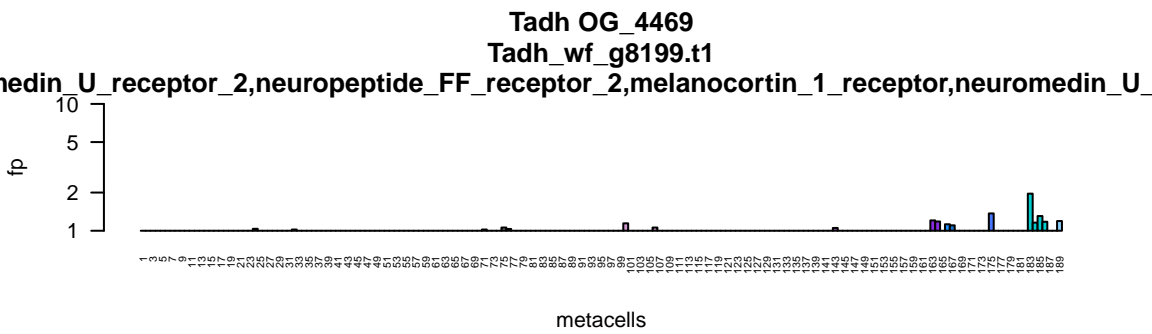


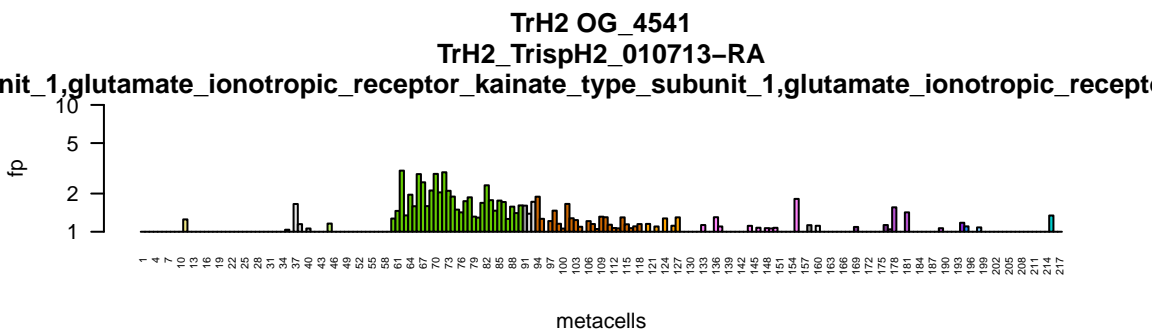
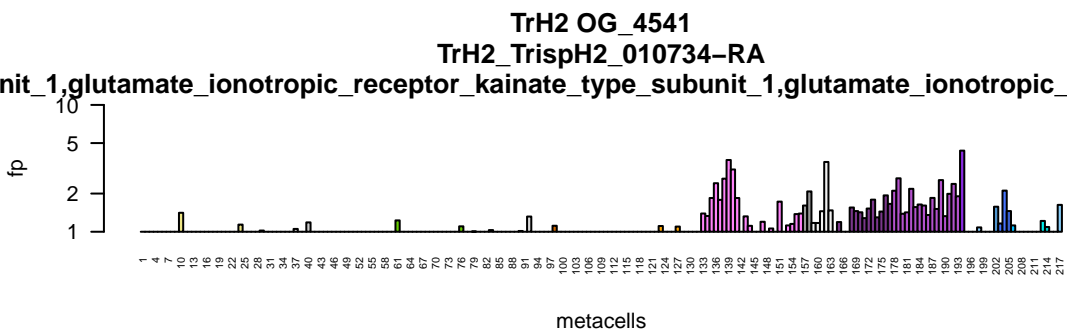
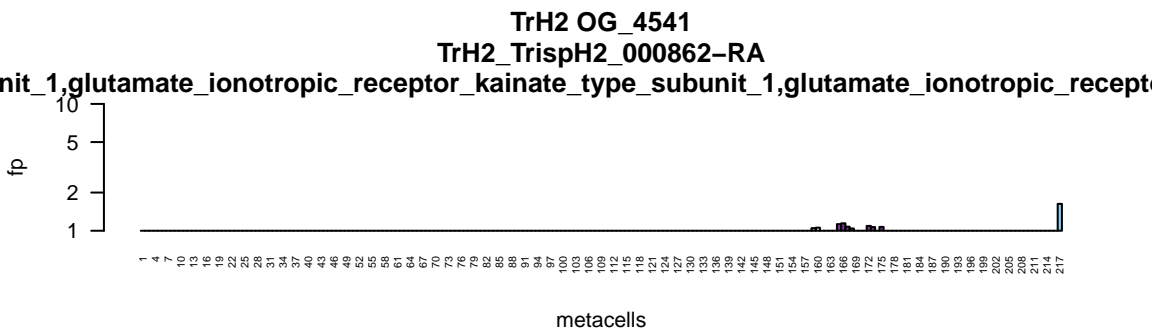
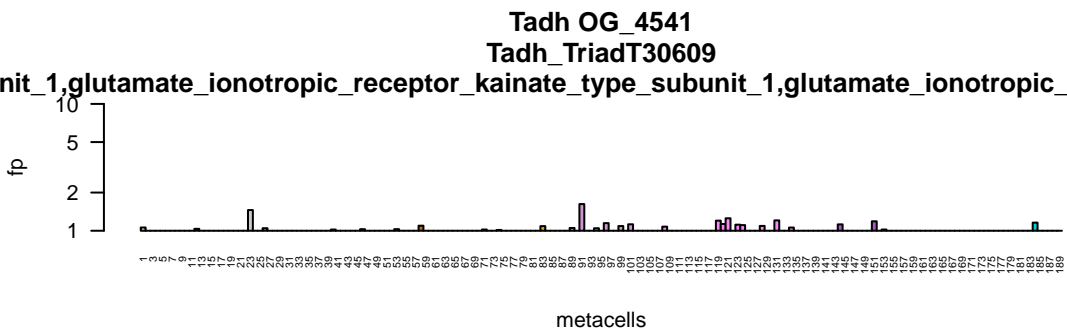
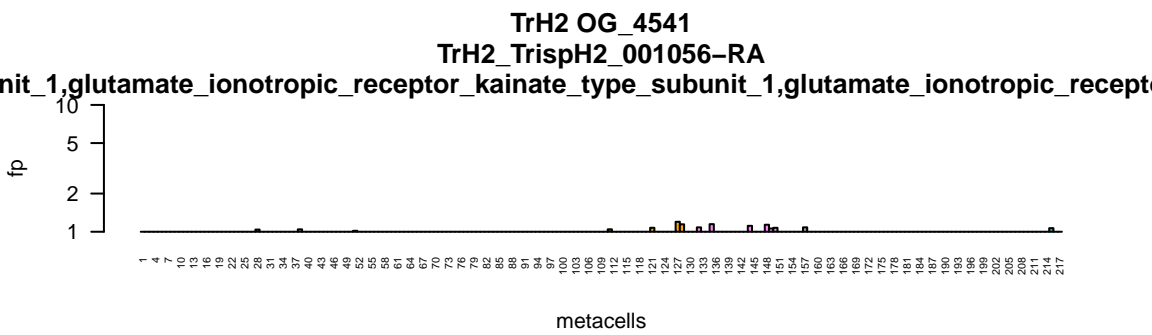
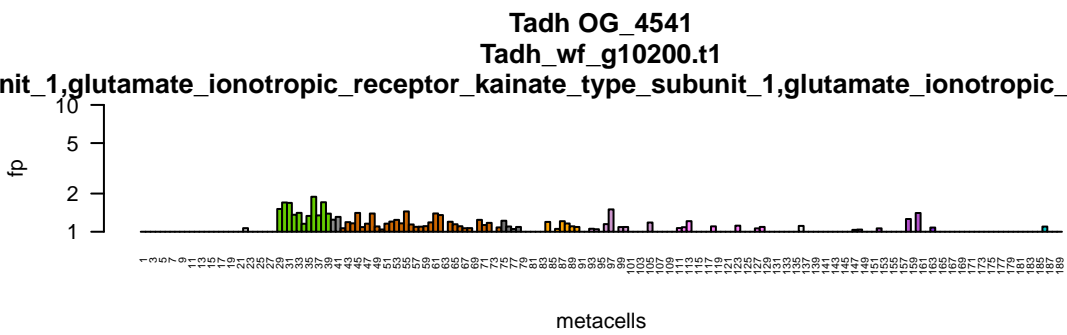
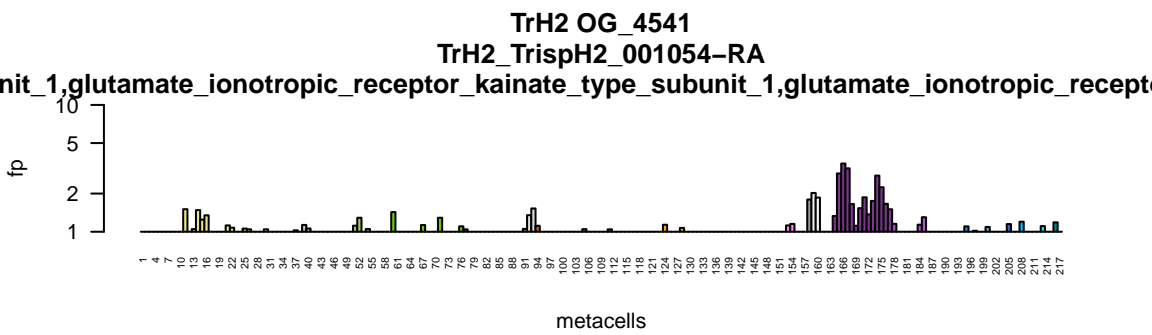
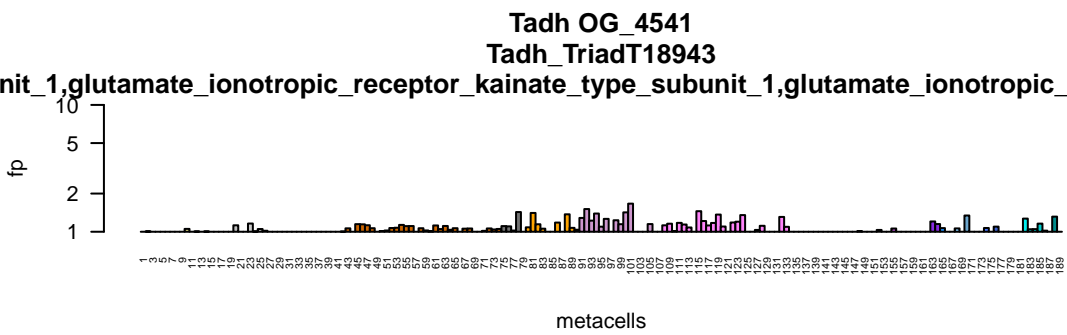
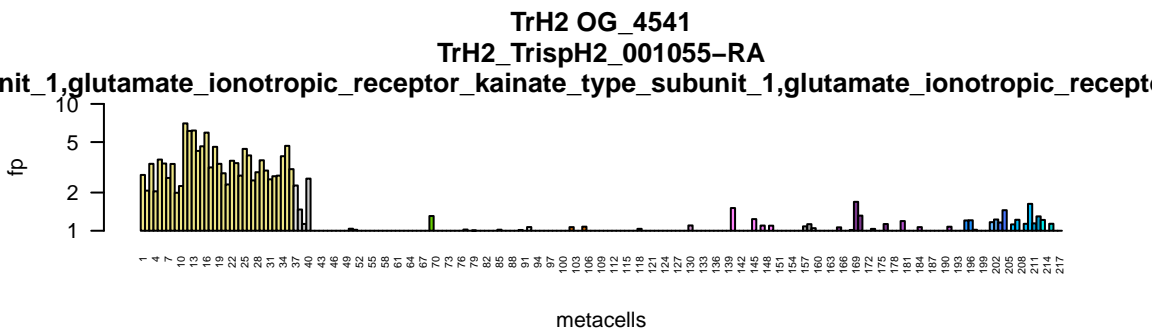
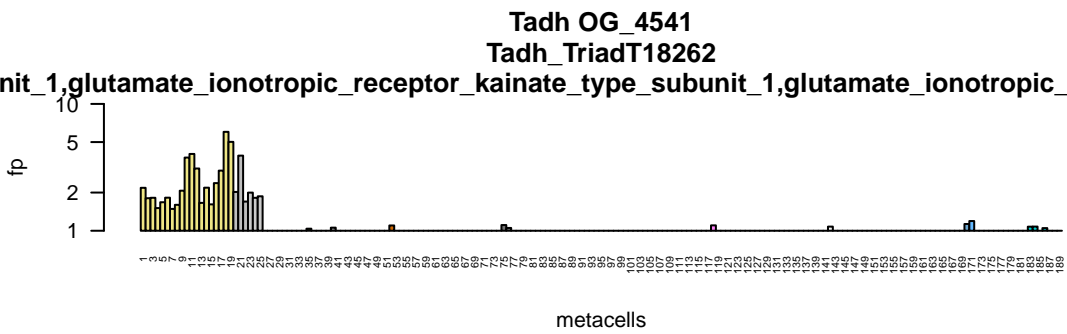
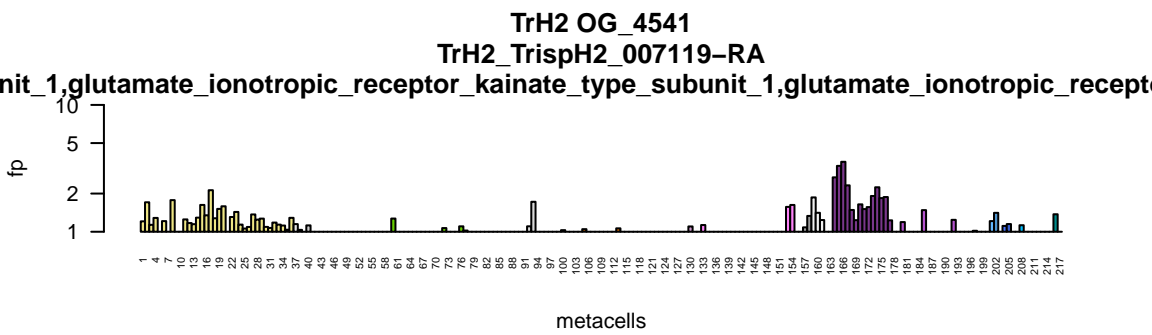
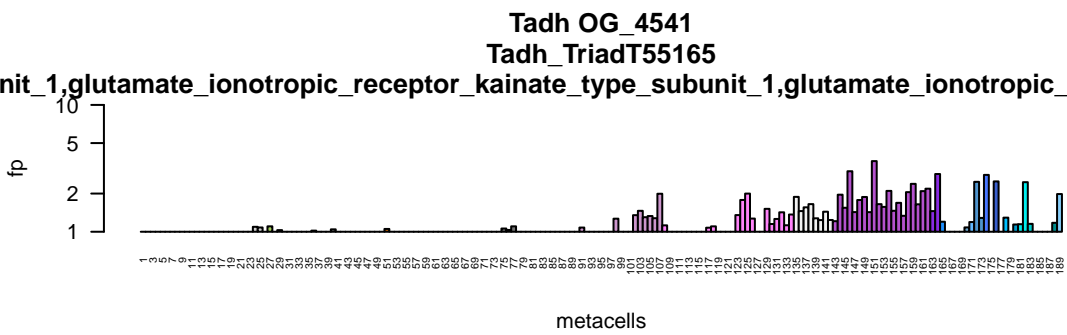
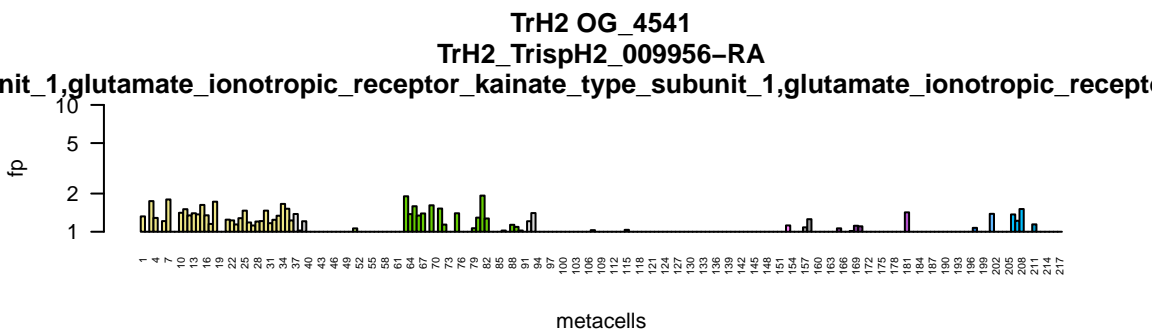
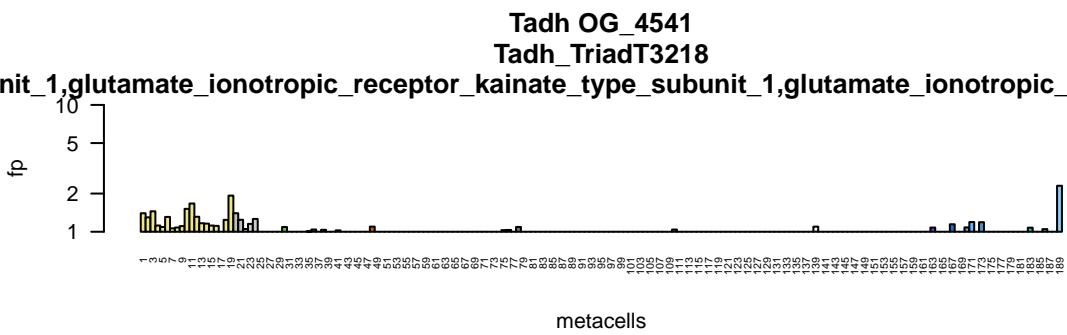
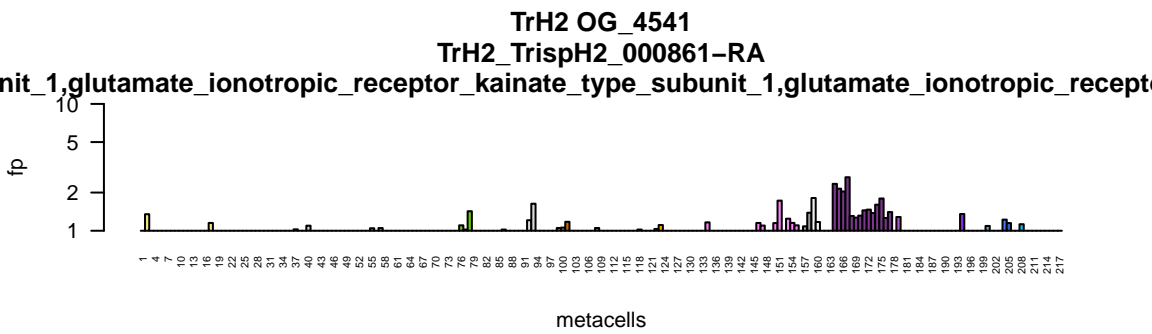
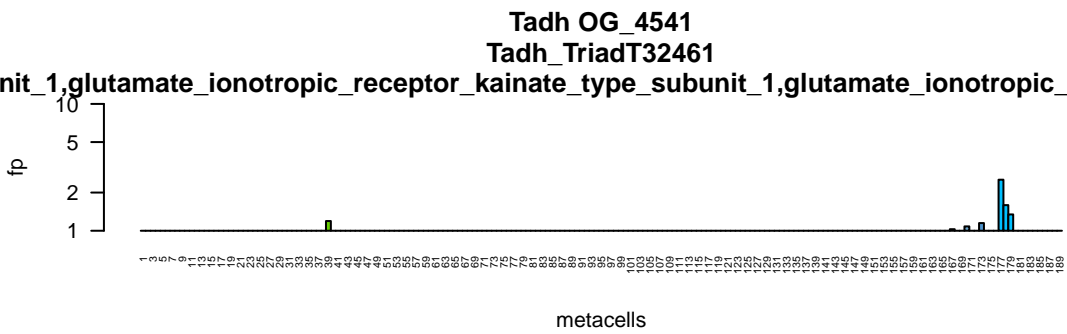
peptide_FF_receptor_2,pyroglutamylated_RFamide_peptide_receptor,neuropeptide_Y_rec
TrH2 | no data

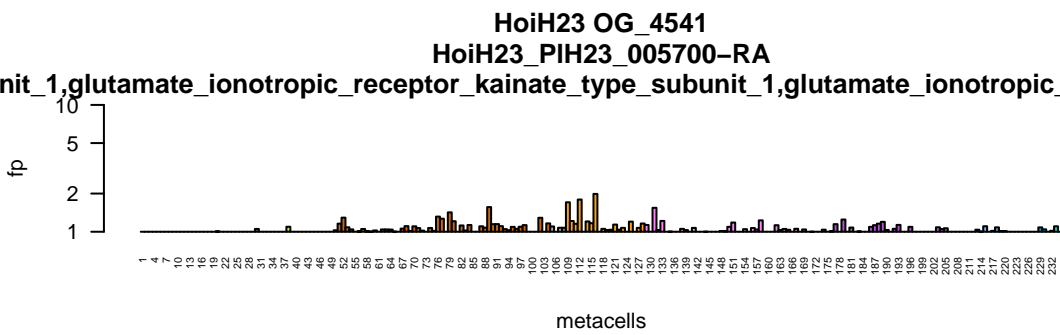
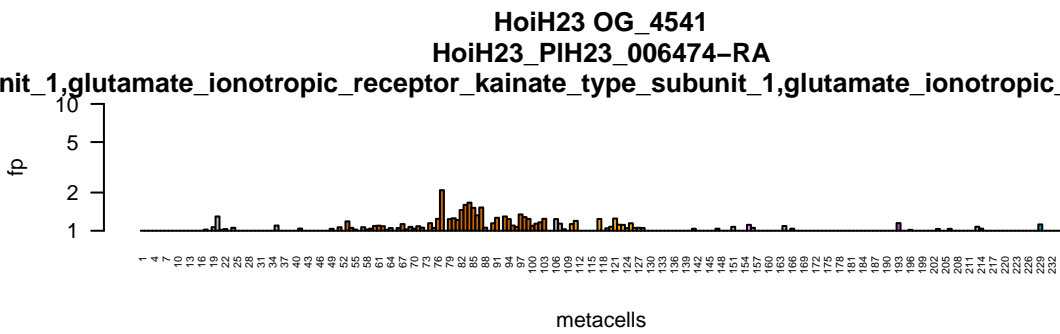
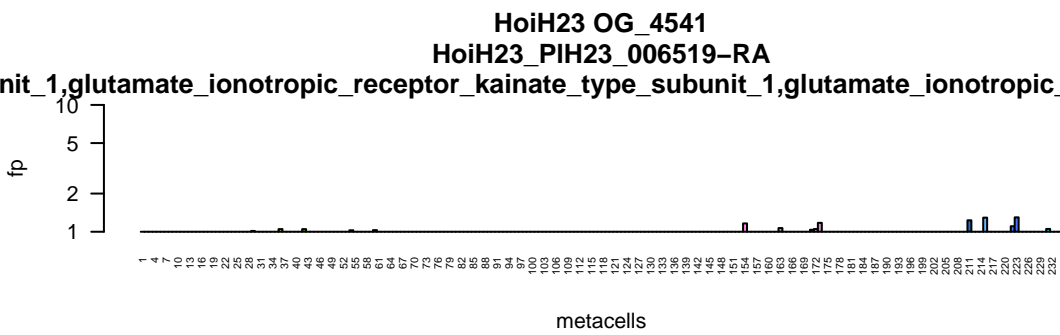
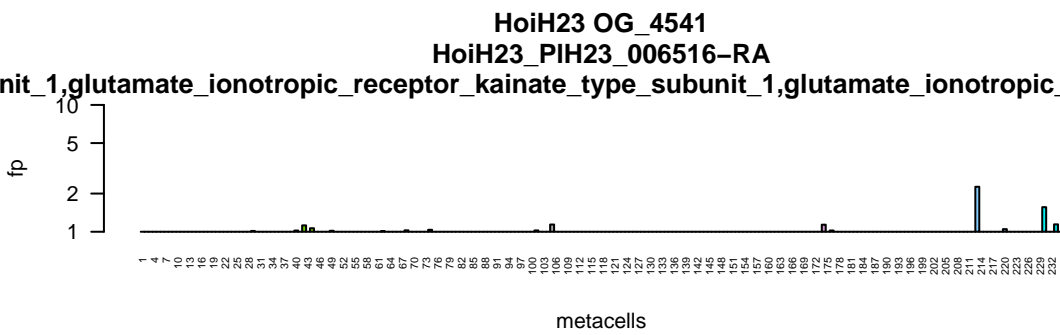
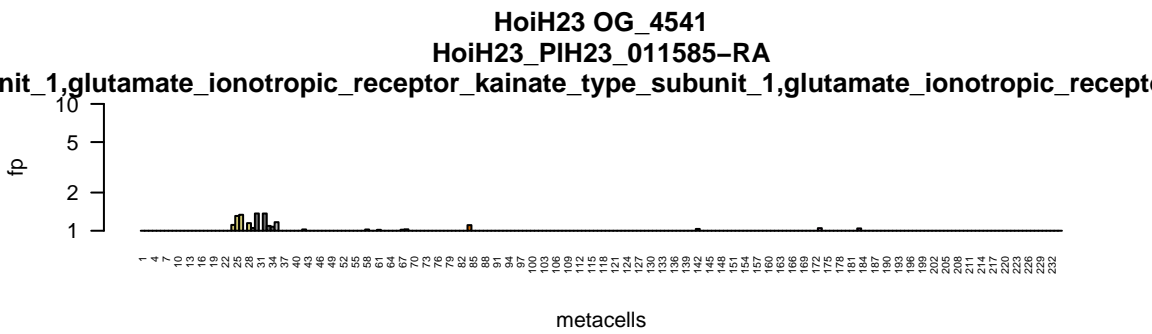
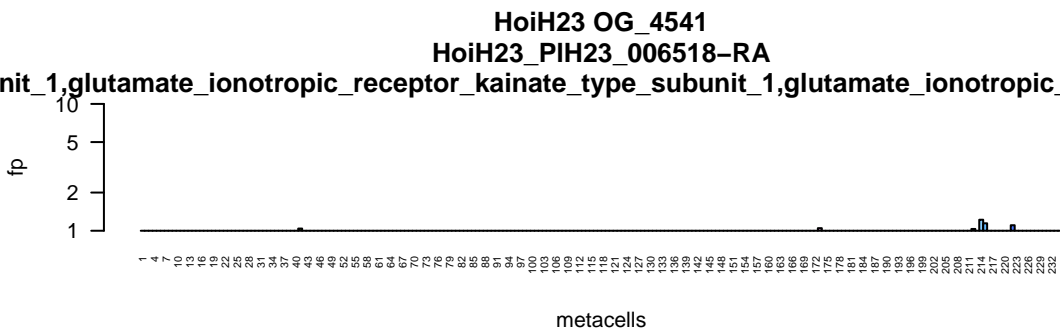
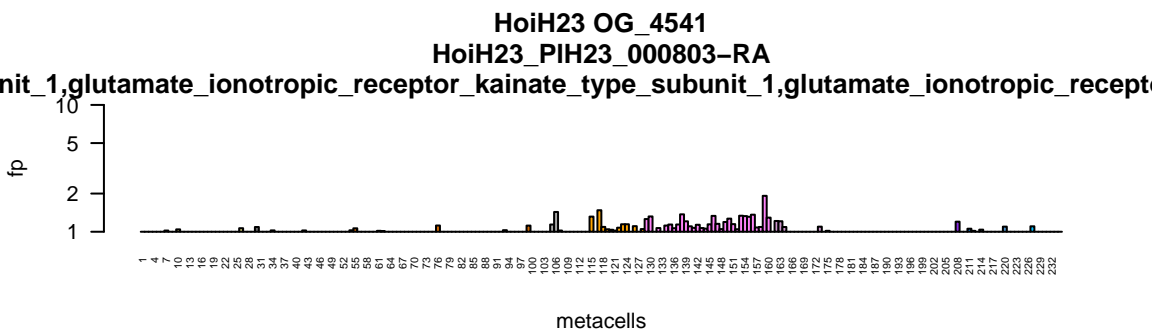
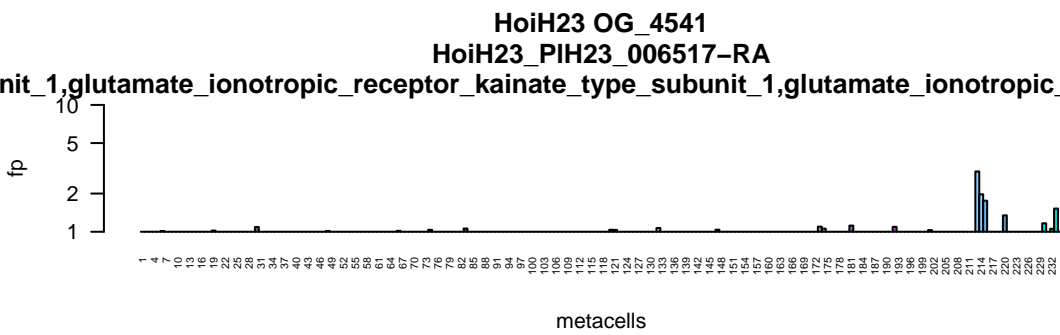
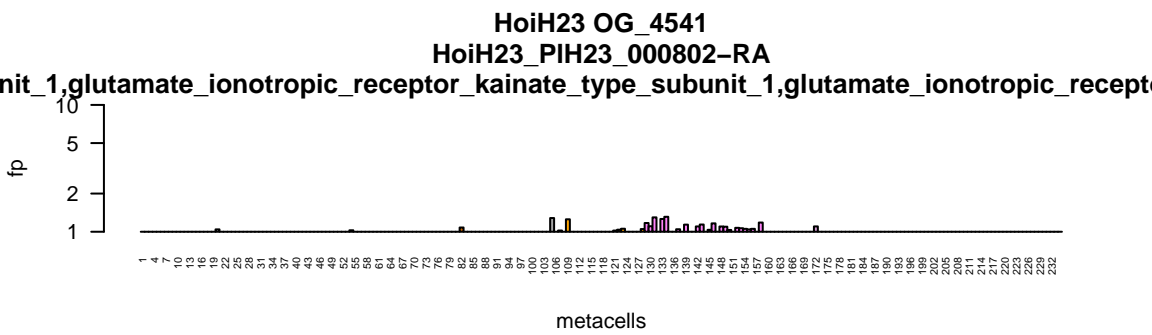
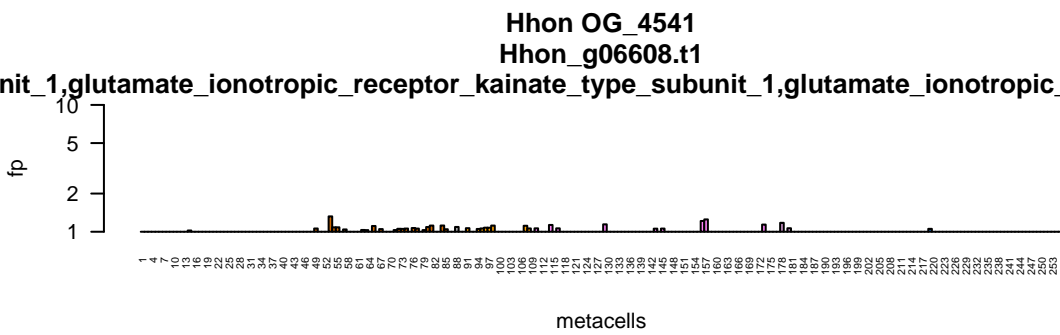
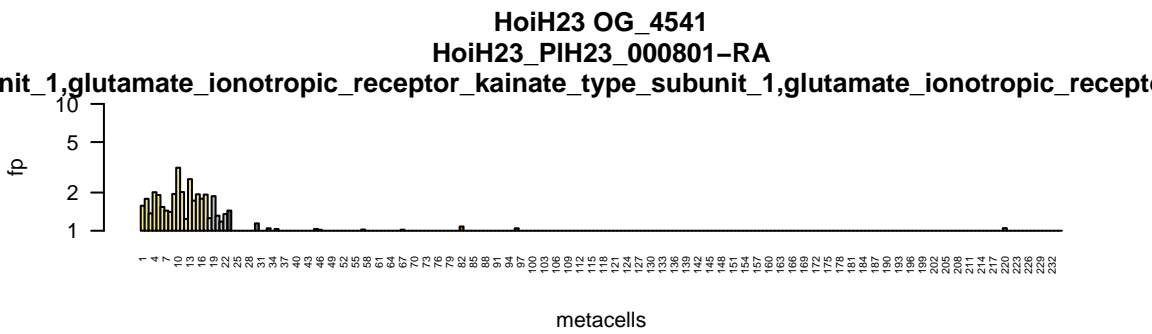
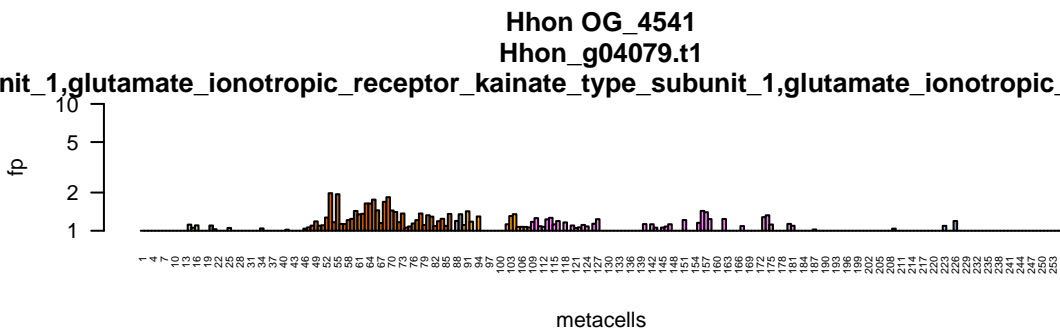


peptide_FF_receptor_2,pyroglutamylated_RFamide_peptide_receptor,neuropeptide_Y_rec
HoiH23 | no data

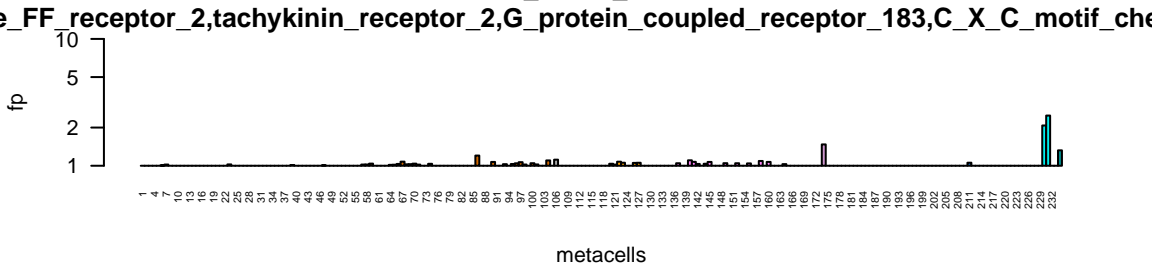


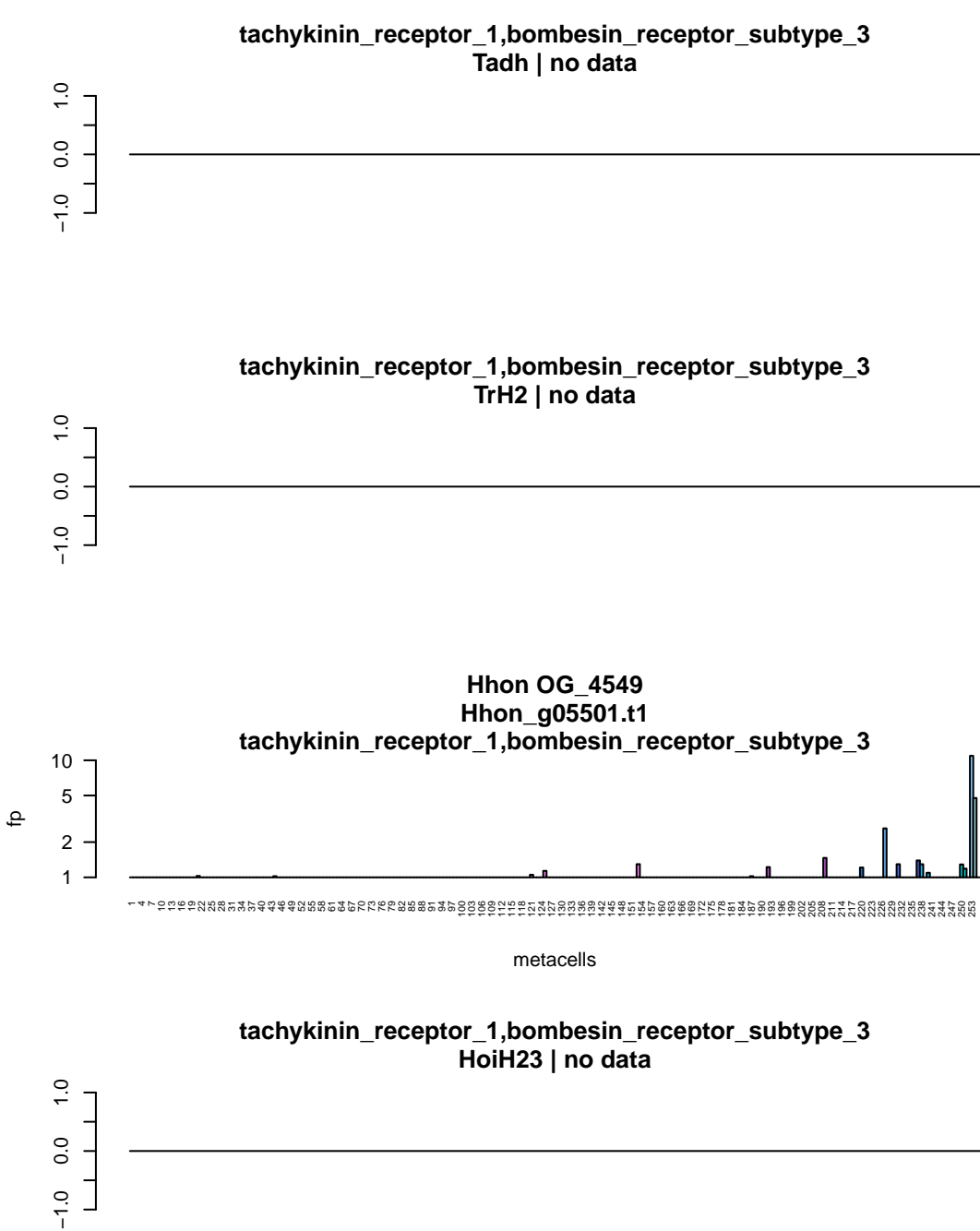


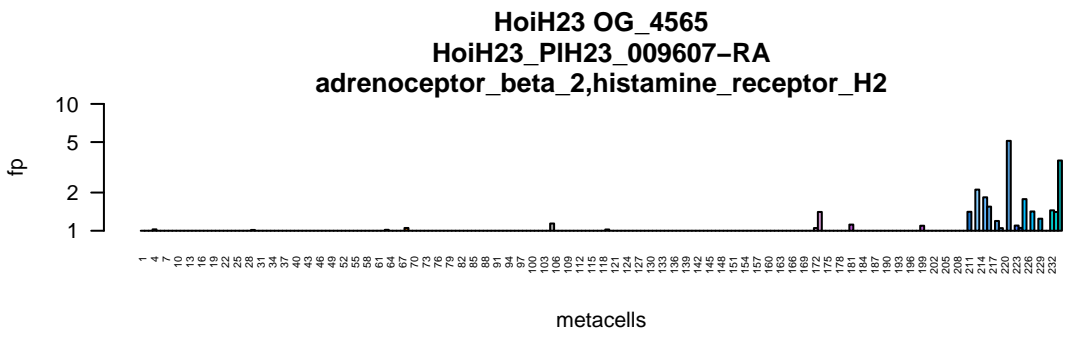
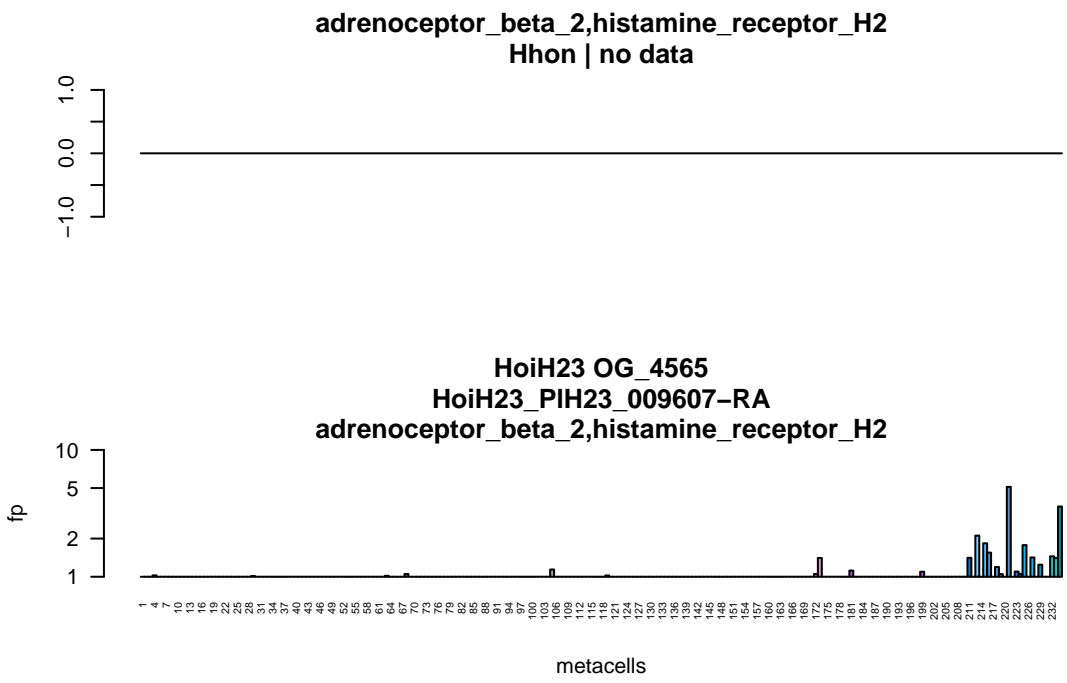
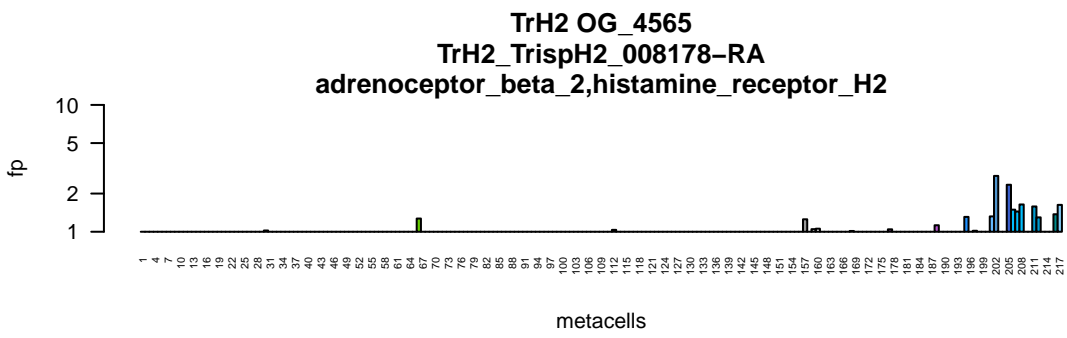
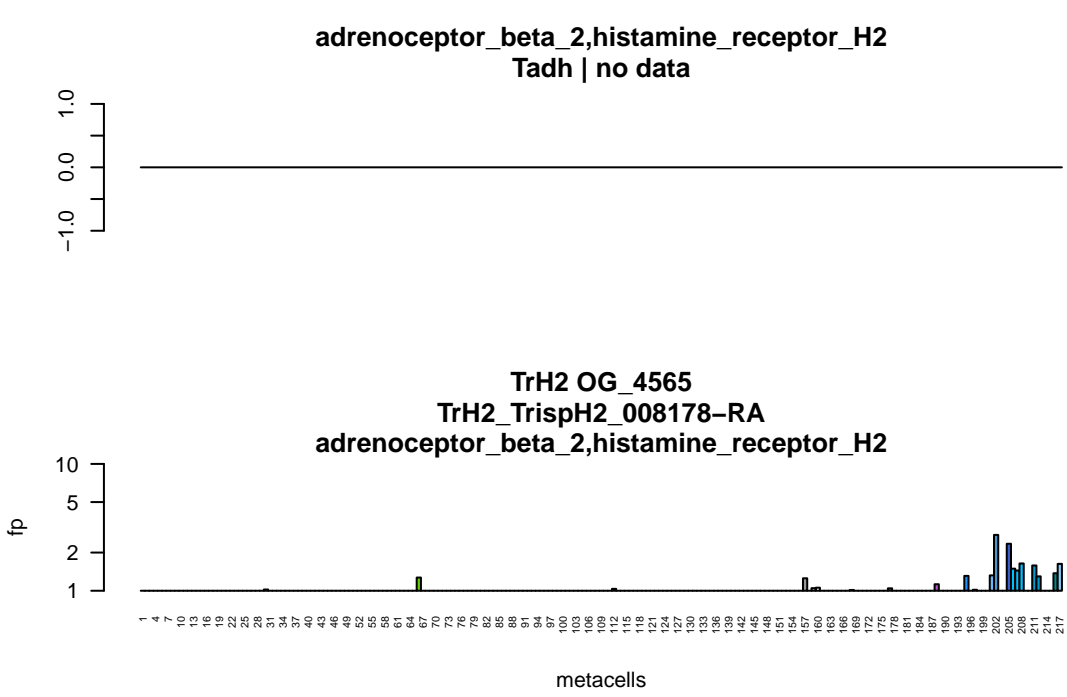




HoiH23 OG_4547
HoiH23_PIH23_002323-RA







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
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
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31	1
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220	1
221	1
222	1
223	1
224	1
225	1
226	1
227	1
228	1
229	1
230	1
231	1
232	

TrH2 OG_4762
TrH2_TrispH2_010634-RA
hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chen

metacells	fp
1	1
4	1
10	1
13	1
16	1
19	1
22	1
25	1
28	1
31	1
37	1
40	1
43	1
46	1
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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	1.2
211	1.1
214	1.8
217	2.2

TrH2 OG_4762
TrH2_TrispH2_010271-RA

hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

metacells

Hhon OG_4762
Hhon_g10761.t1

hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

fp

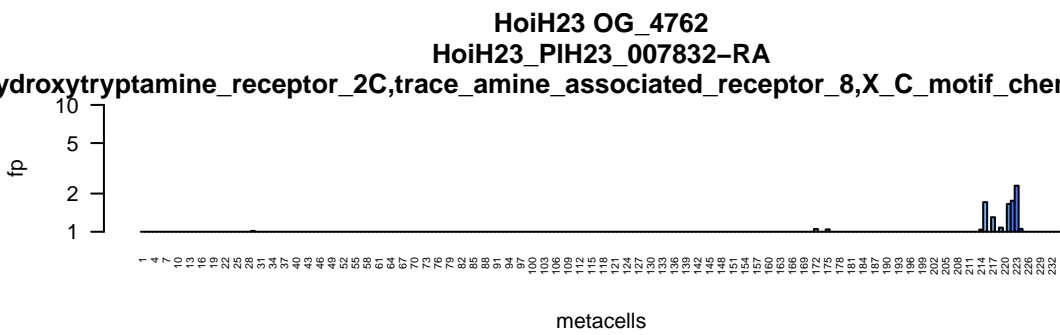
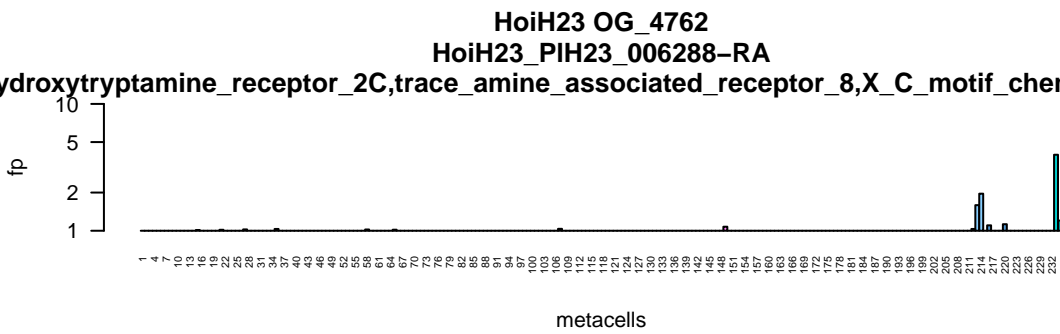
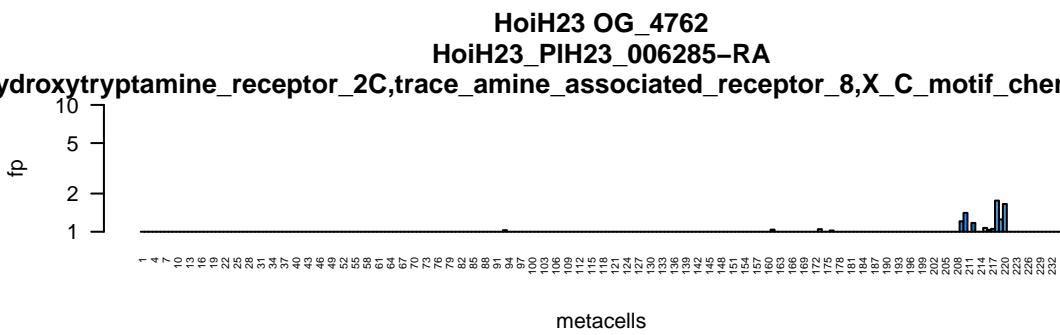
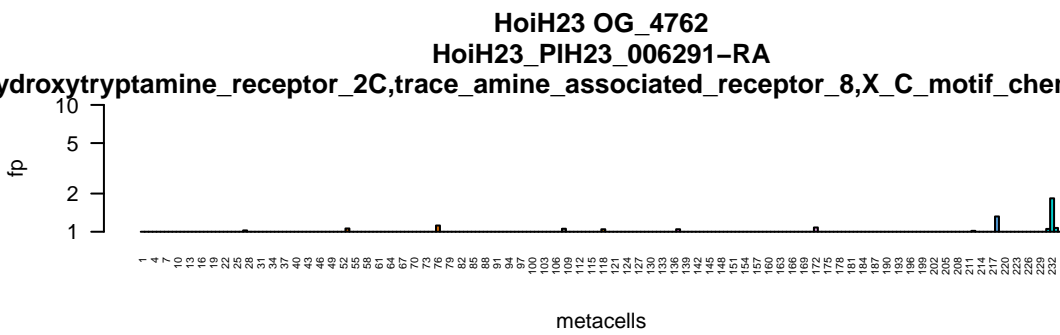
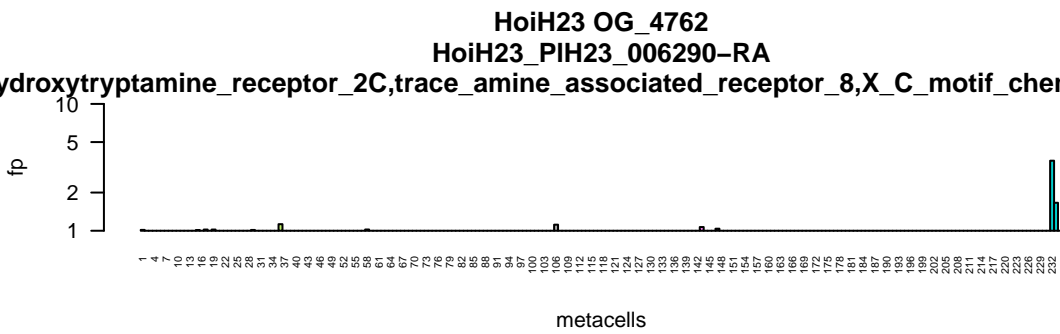
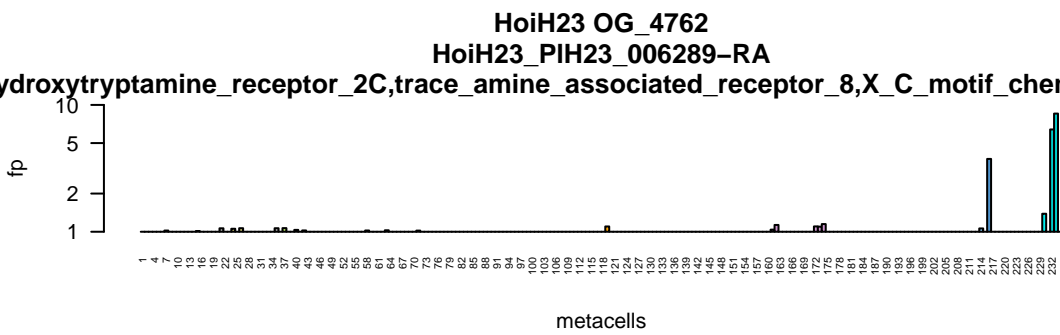
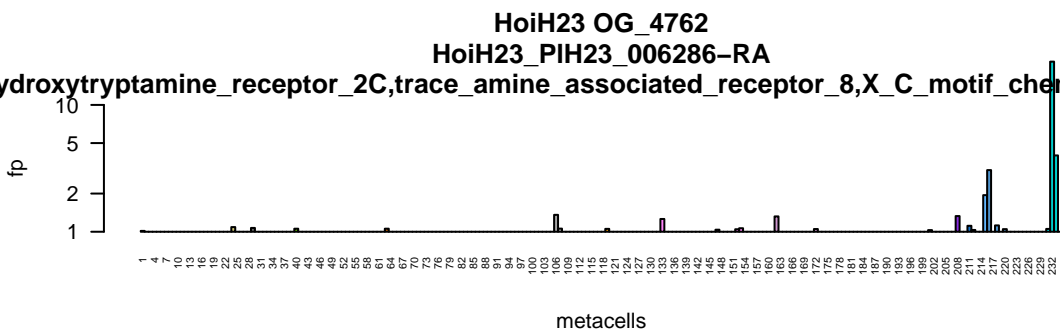
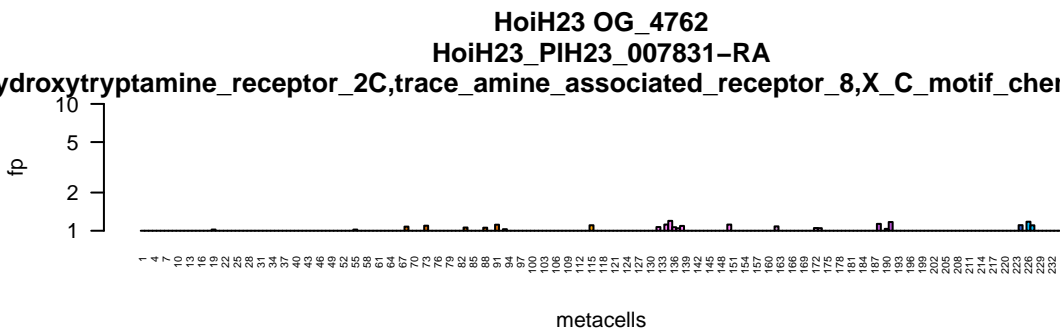
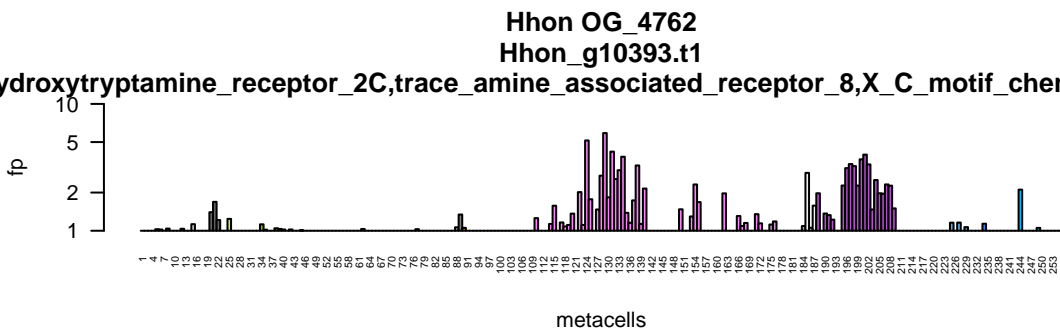
metacells

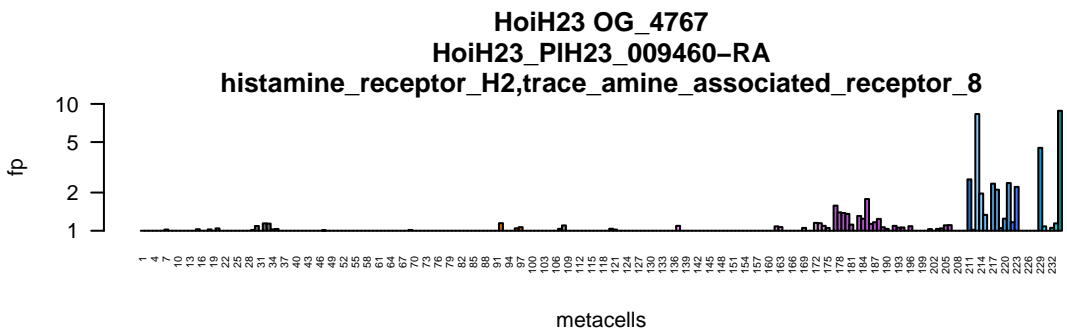
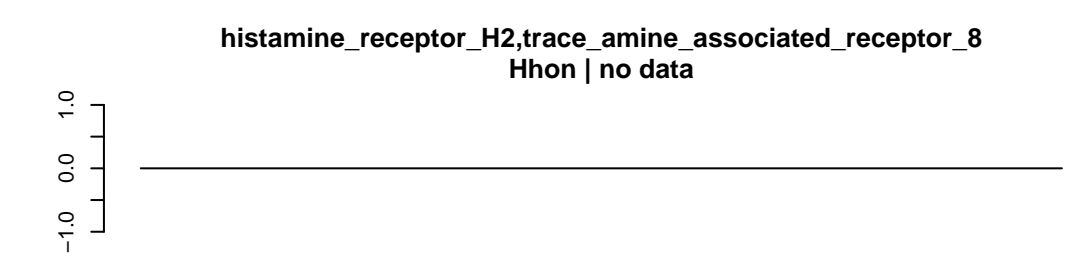
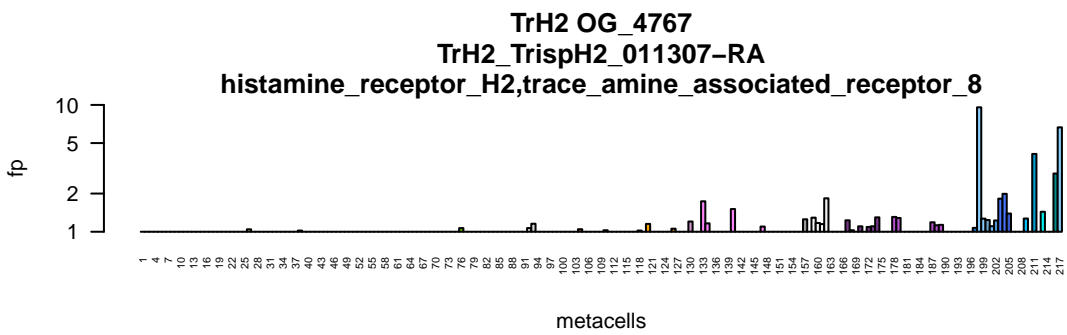
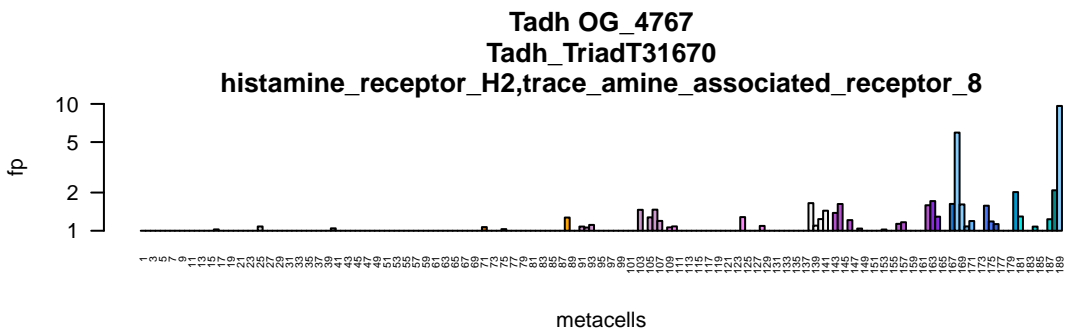
Hhon OG_4762
Hhon_g01703.t1

hydroxytryptamine_receptor_2C,trace_amine_associated_receptor_8,X_C_motif_chemokine

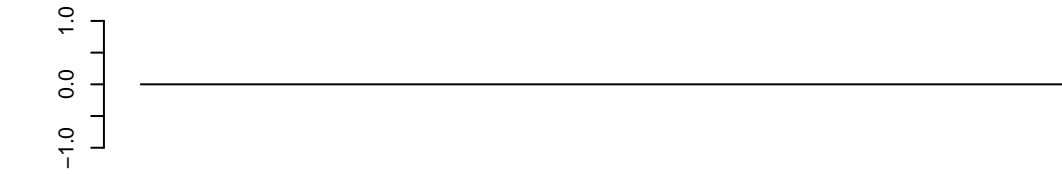
fp

metacells

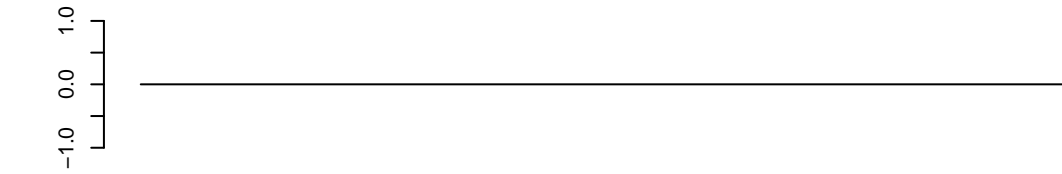




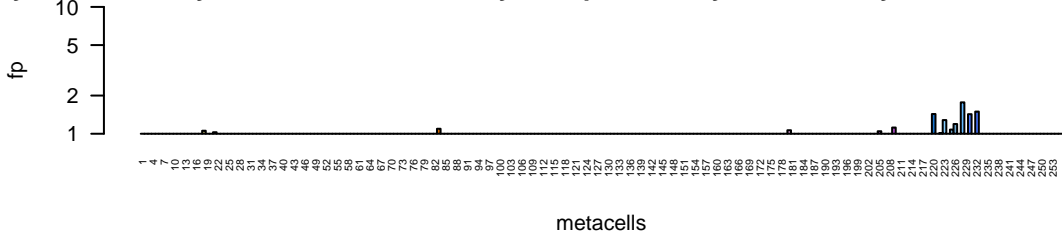
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi
Tadh | no data



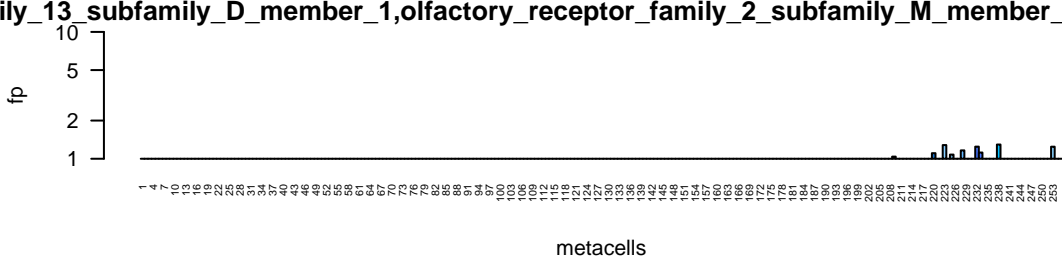
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi
TrH2 | no data



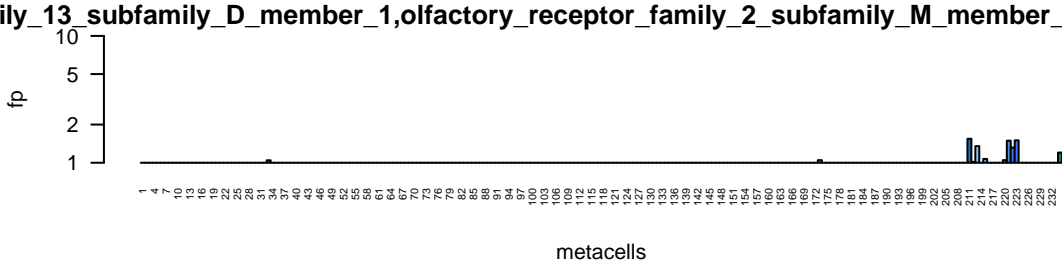
Hhon OG_4788
Hhon_g08016.t1
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi



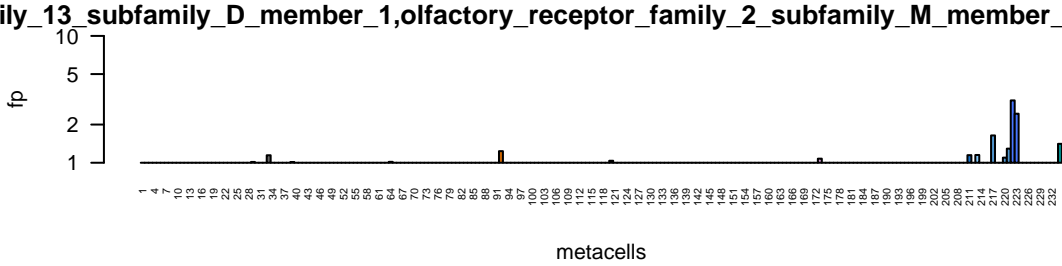
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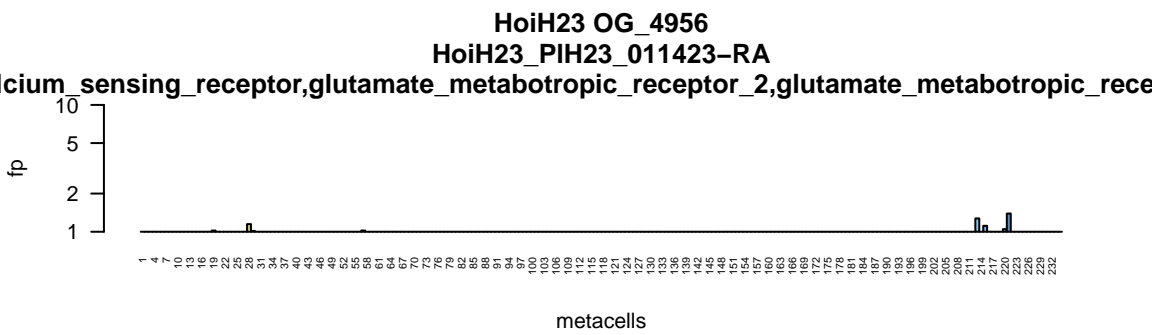
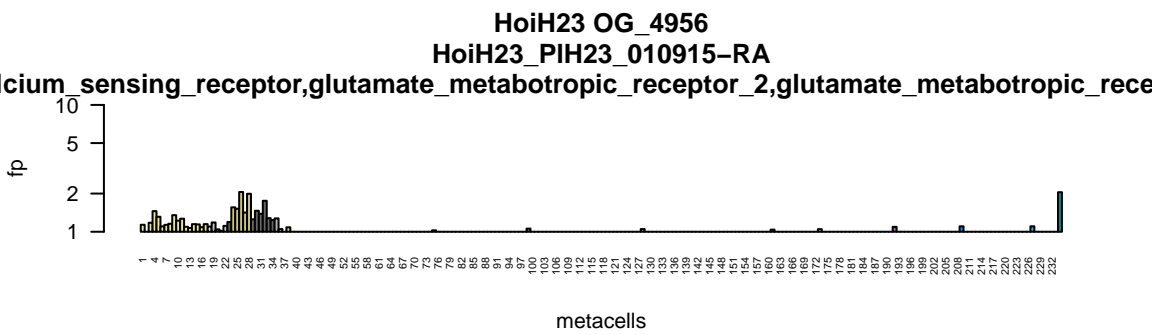
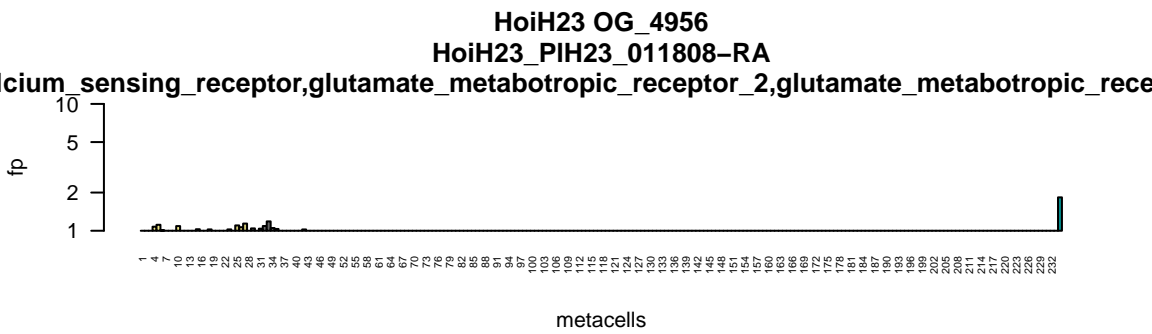
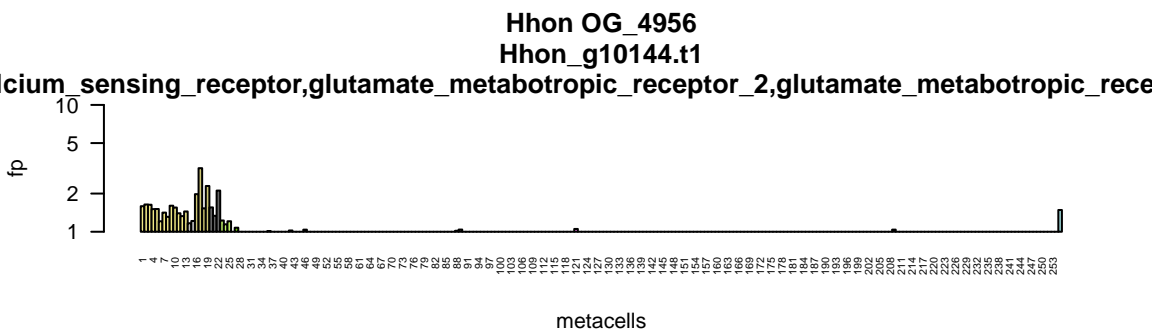
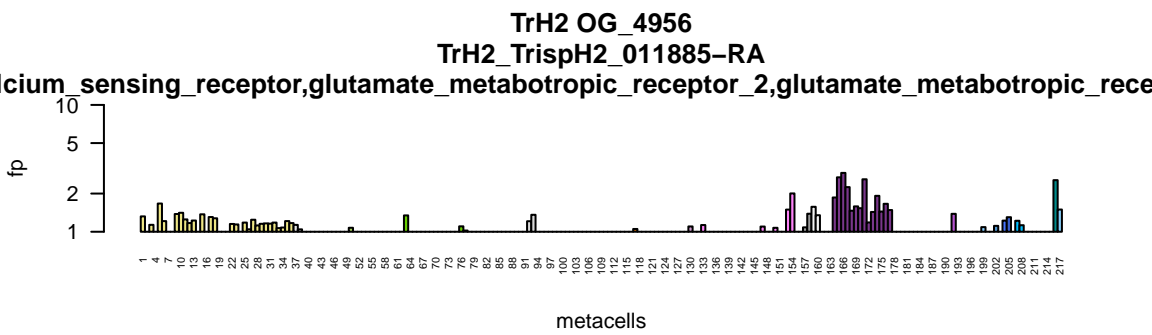
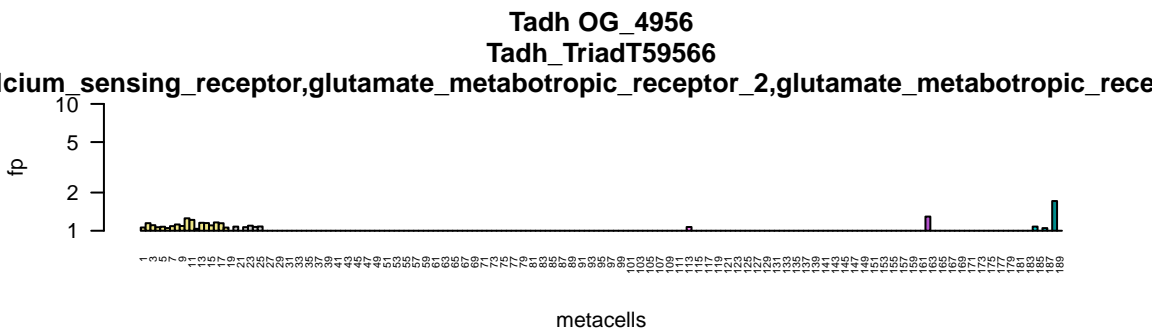


HoiH23 OG_4788
HoiH23_PIH23_005444-RA
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi

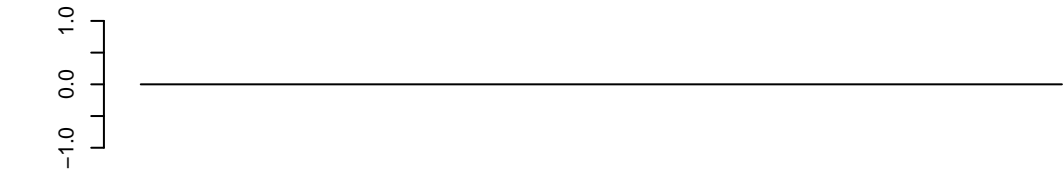


HoiH23 OG_4788
HoiH23_PIH23_005445-RA
ily_13_subfamily_D_member_1,olfactory_receptor_family_2_subfamily_M_member_3,opioi

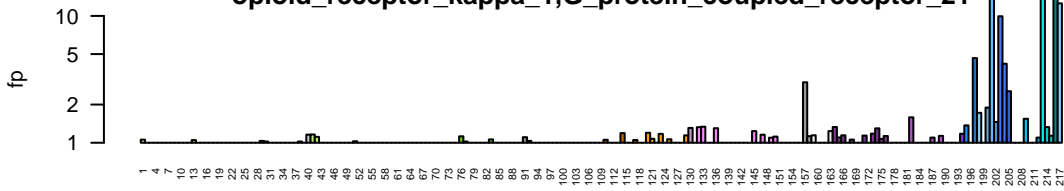




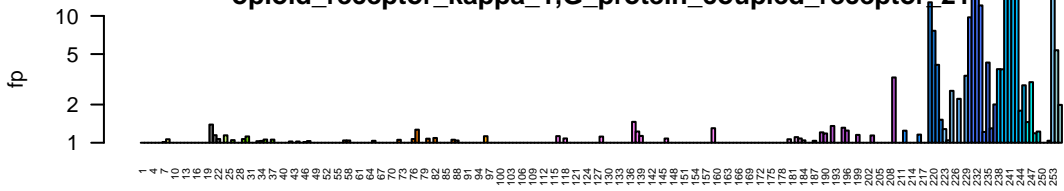
opioid_receptor_kappa_1,G_protein_coupled_receptor_21
Tadh | no data



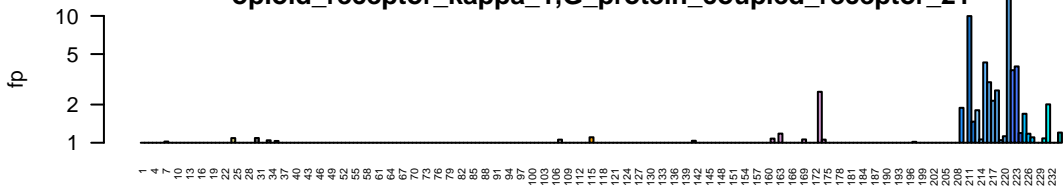
TrH2 OG_5203
TrH2_TrispH2_004138-RA
opioid_receptor_kappa_1,G_protein_coupled_receptor_21



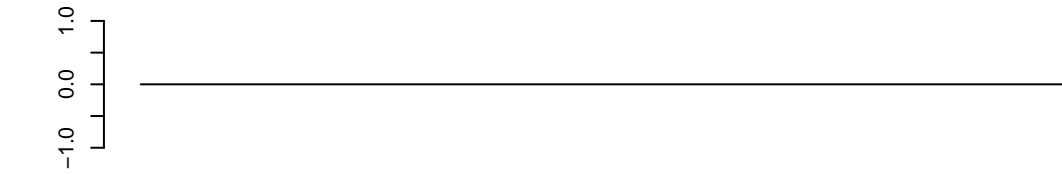
Hhon OG_5203
Hhon_g01680.t1
opioid_receptor_kappa_1,G_protein_coupled_receptor_21



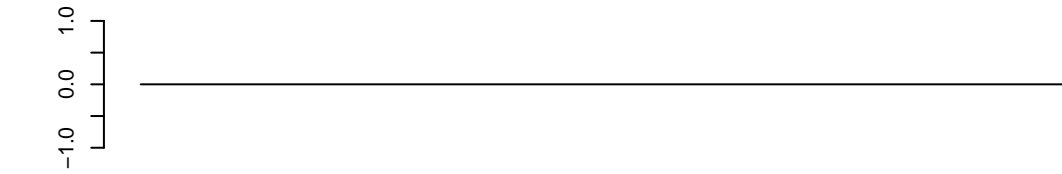
HoiH23 OG_5203
HoiH23_PIH23_003635-RA
opioid_receptor_kappa_1,G_protein_coupled_receptor_21



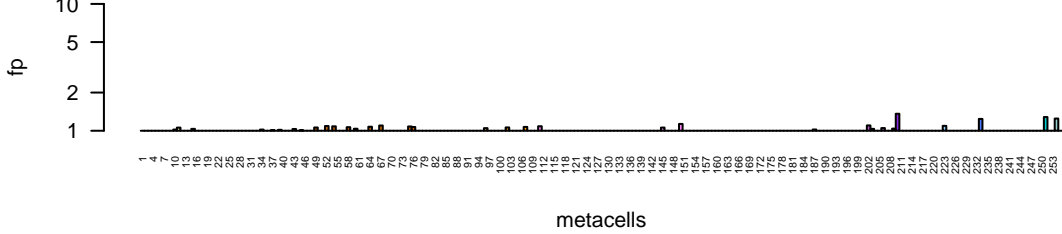
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recept
Tadh | no data



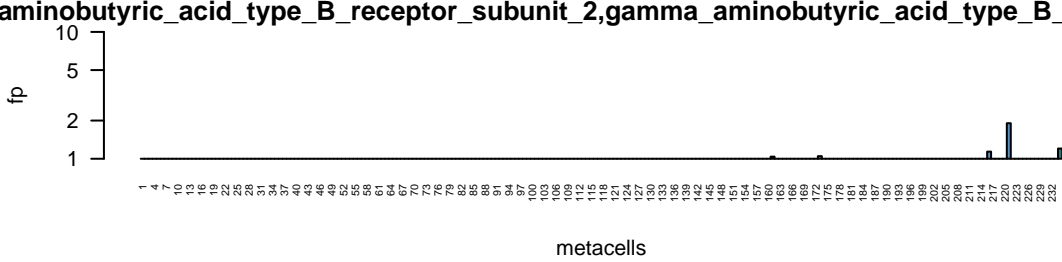
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TrH2 | no data

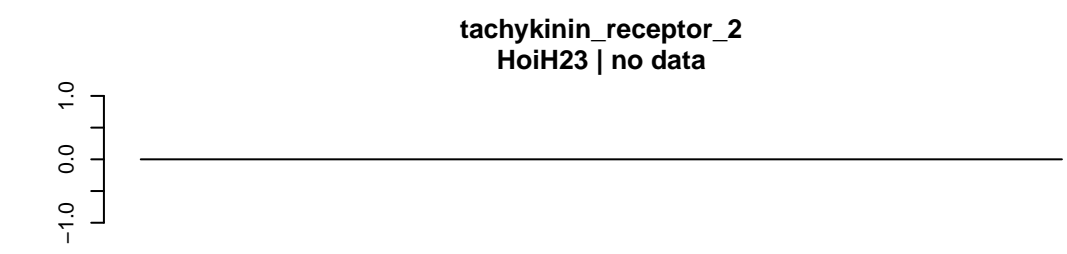
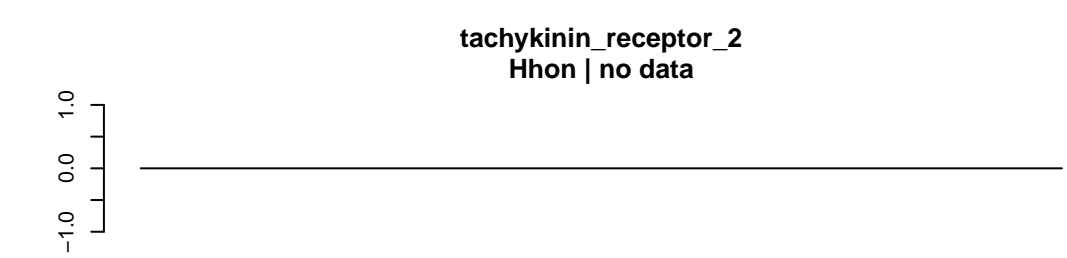
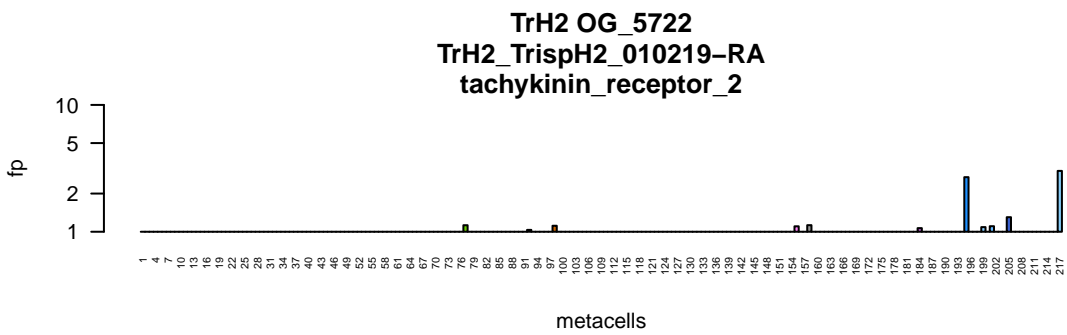
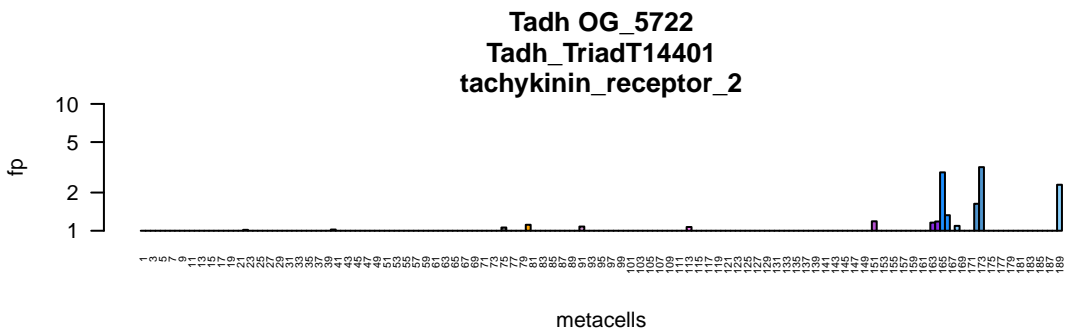


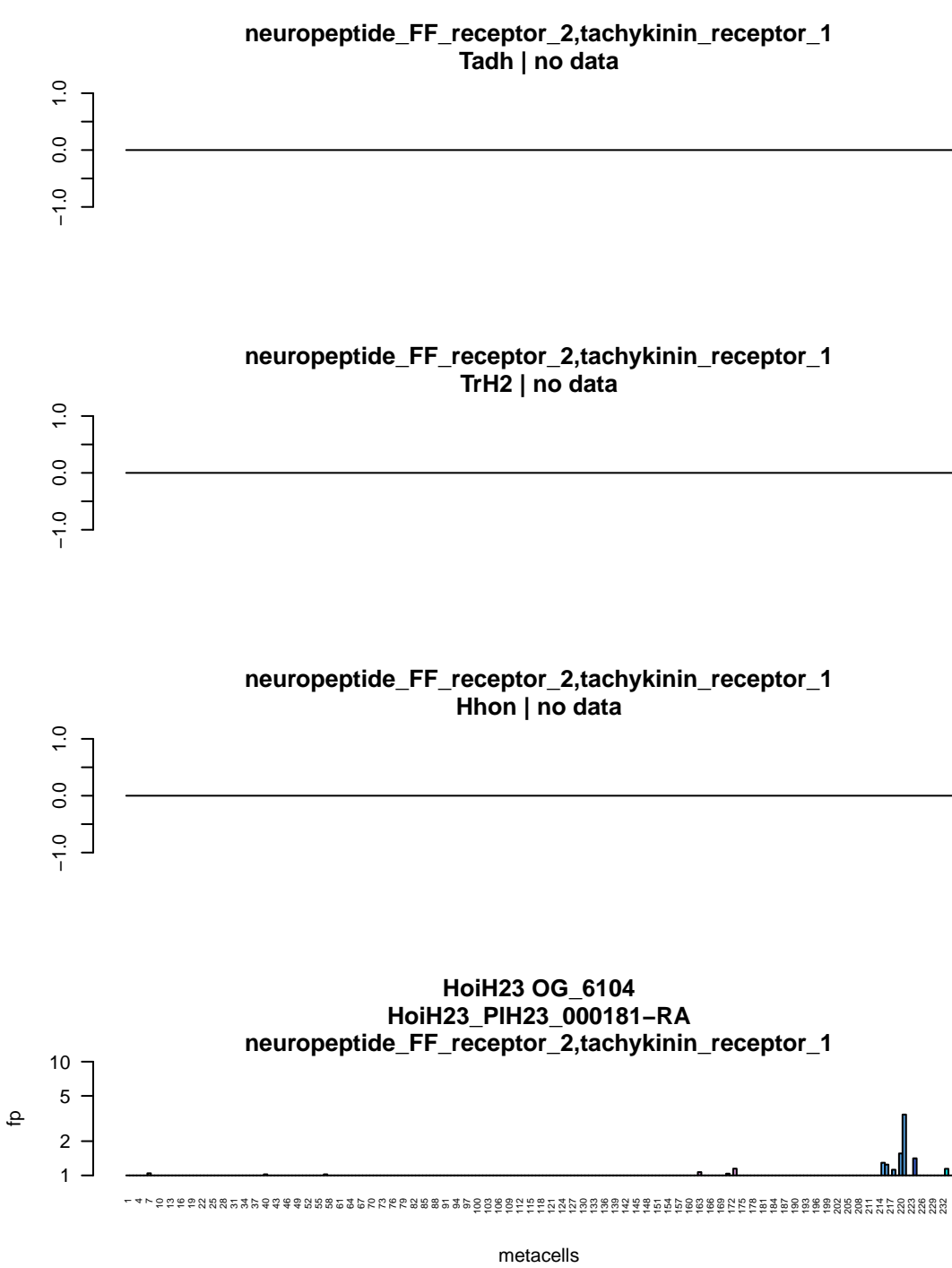
Hhon OG_5437
Hhon_g11390.t1
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recept

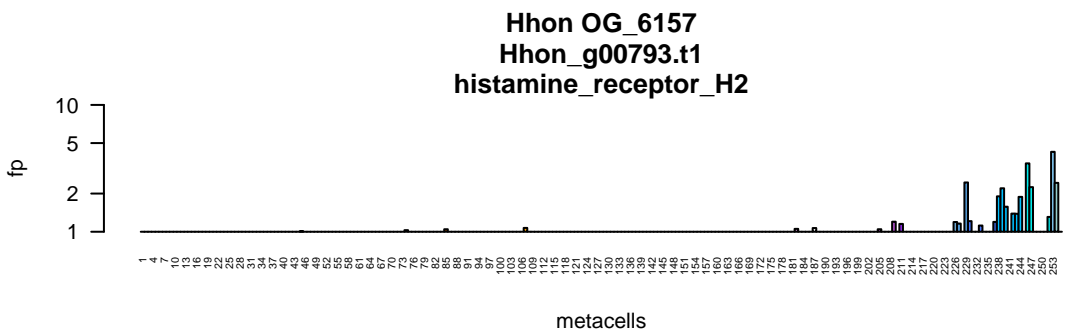
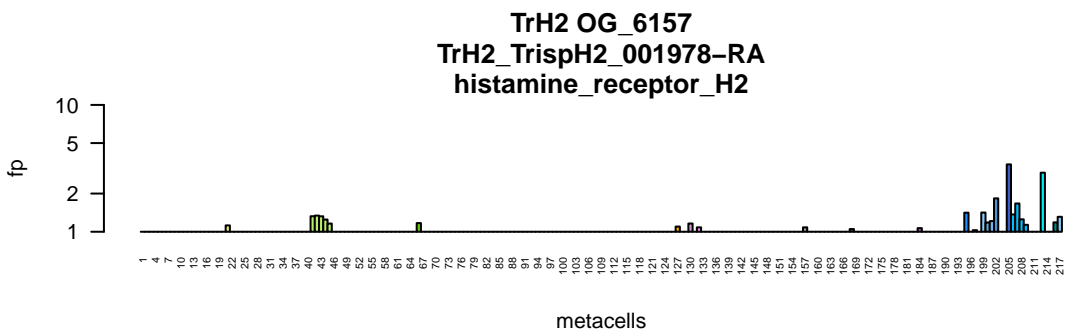
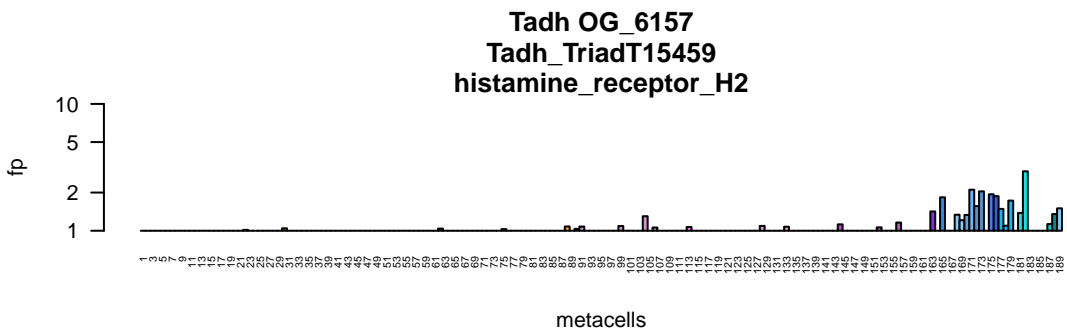


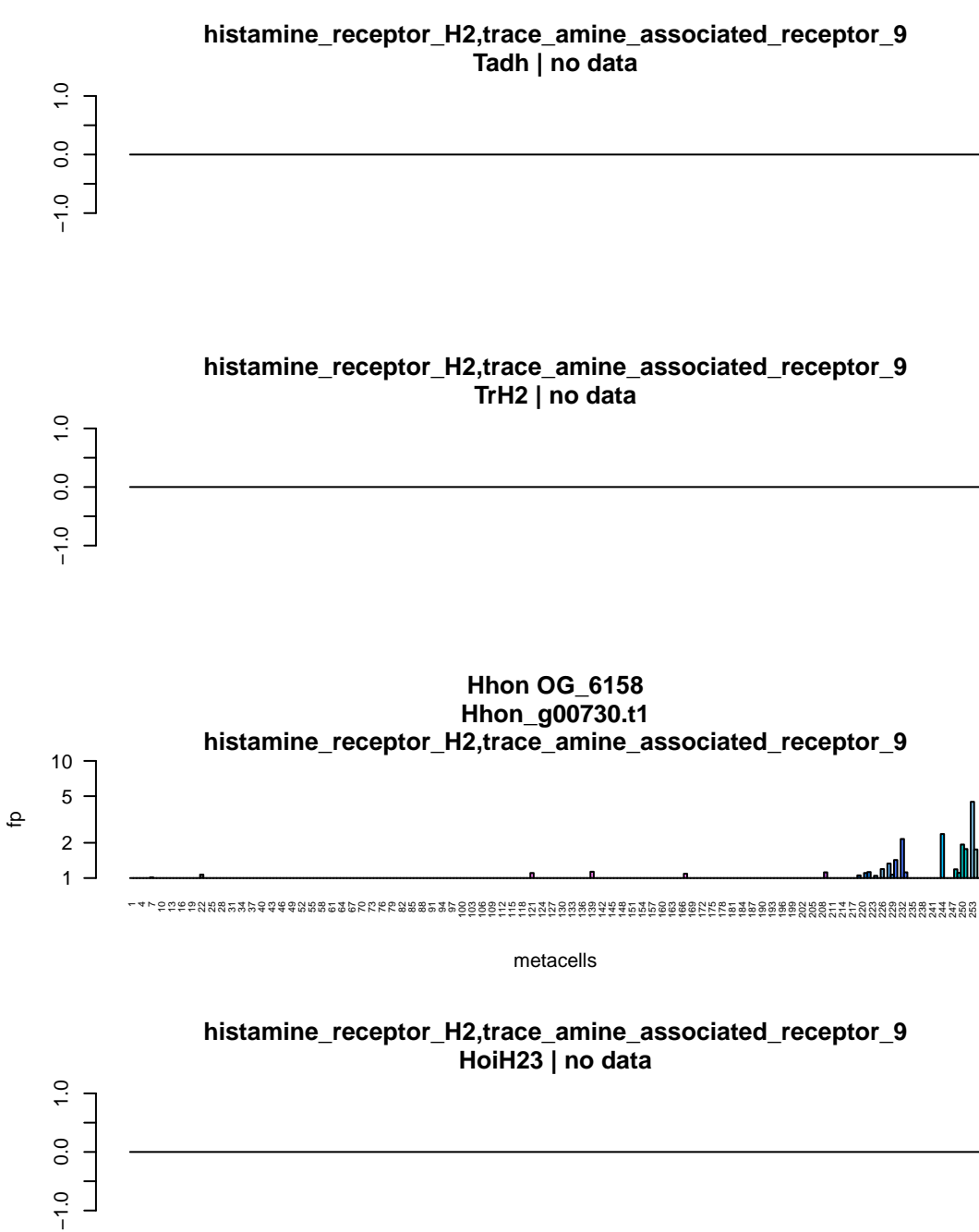
HoiH23 OG_5437
HoiH23_PIH23_011579-RA
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recept



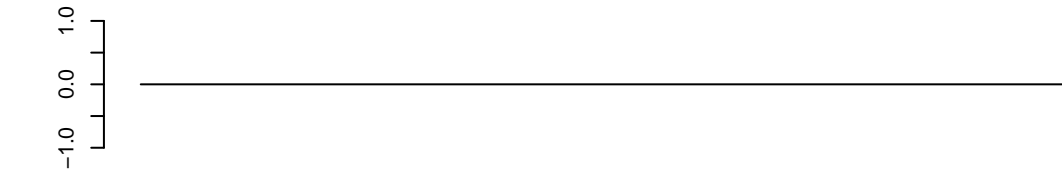




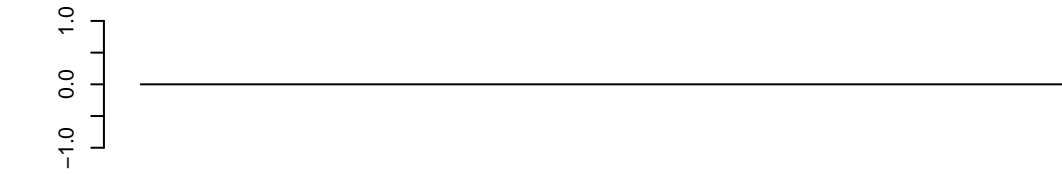




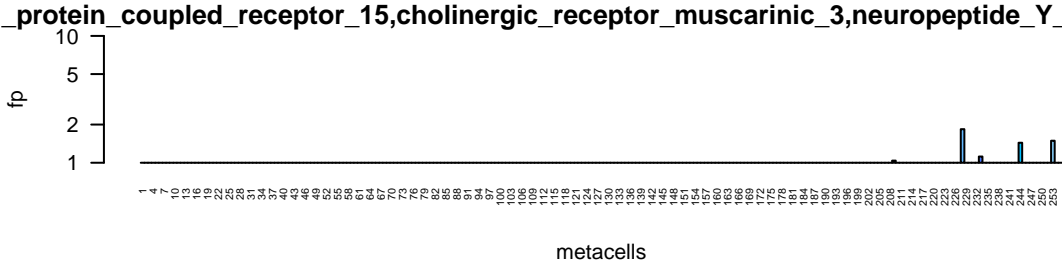
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept
Tadh | no data



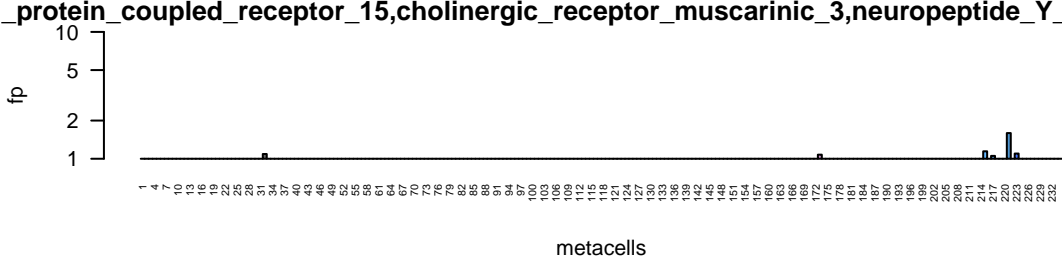
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept
TrH2 | no data

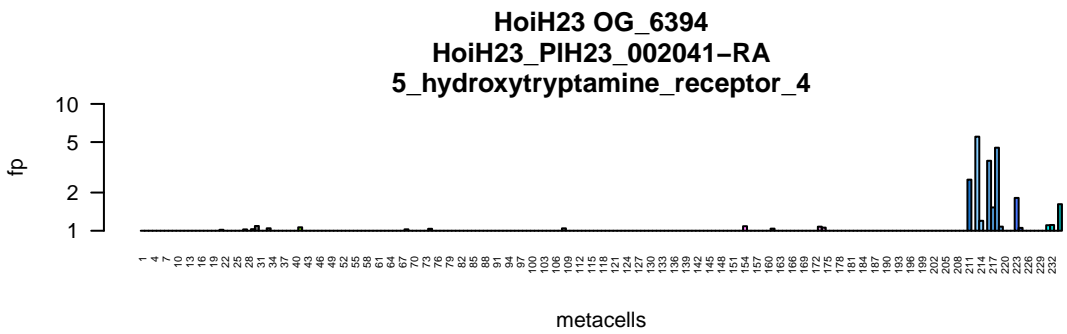
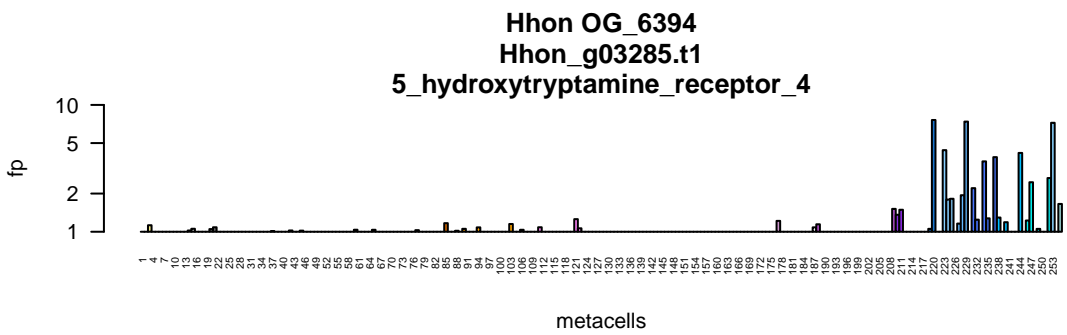
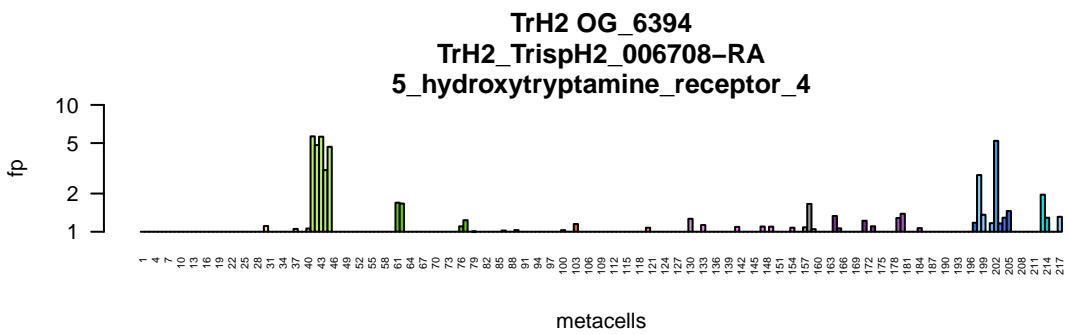
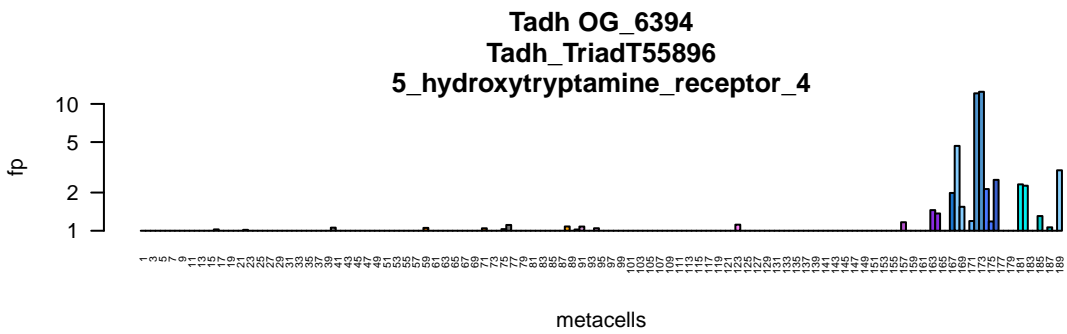


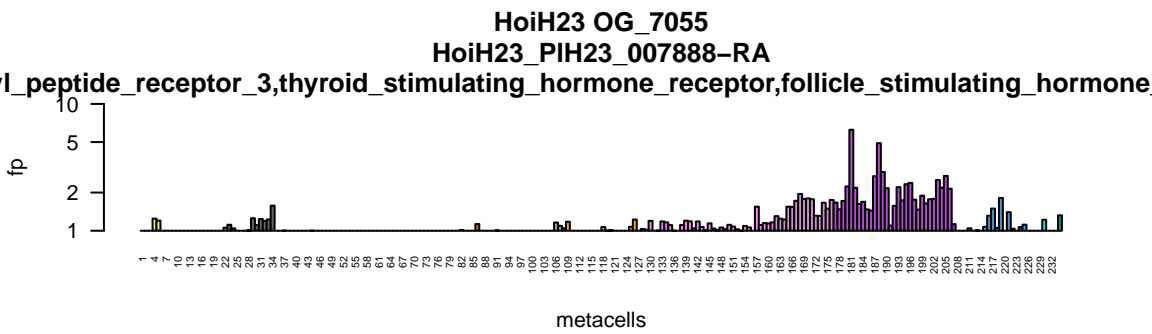
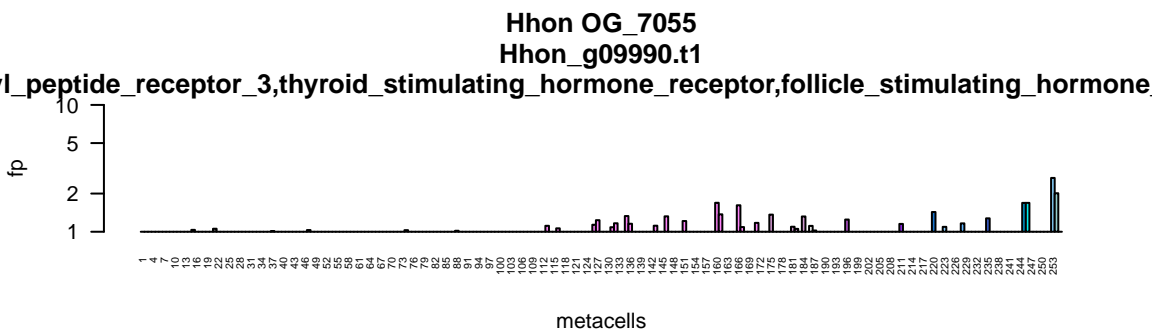
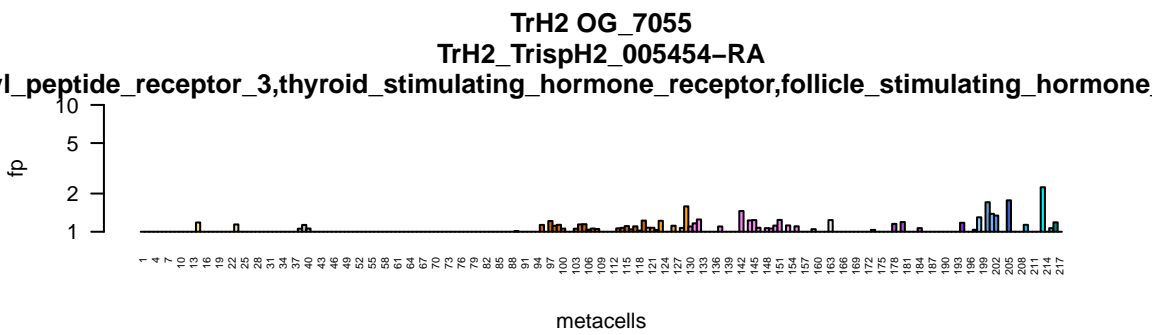
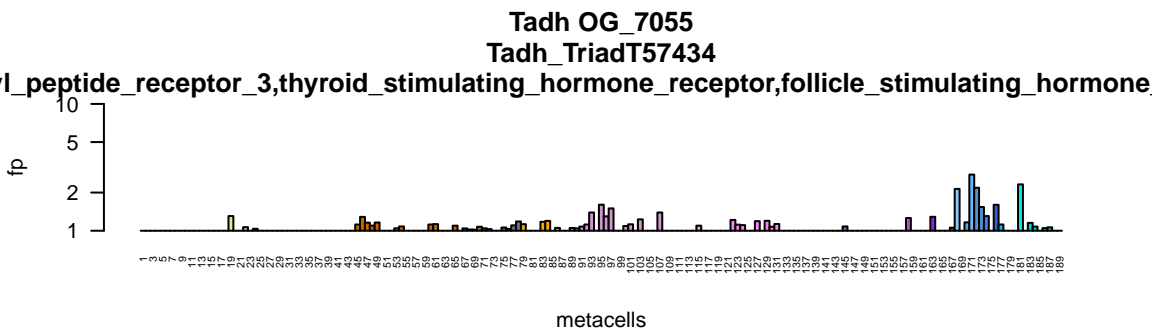
Hhon OG_6355
Hhon_g11199.t1
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept

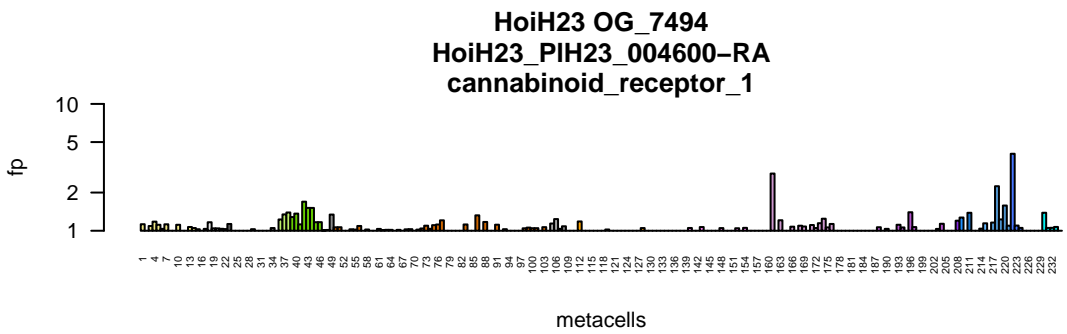
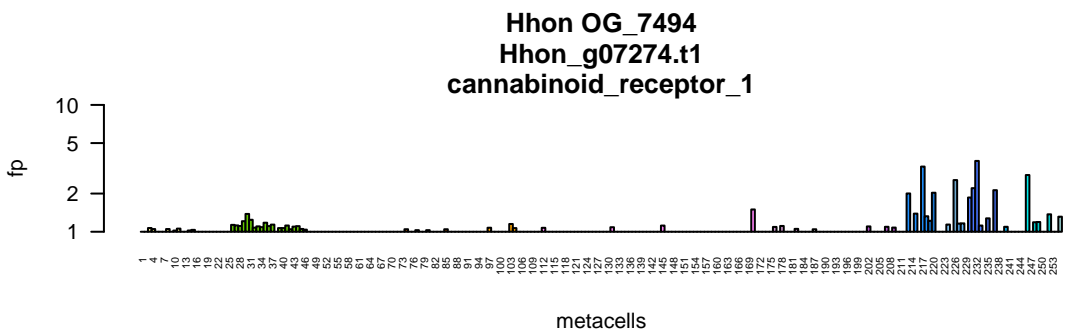
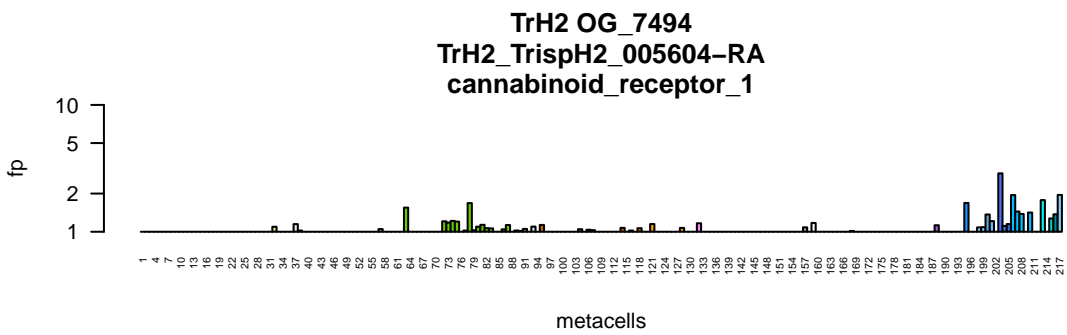
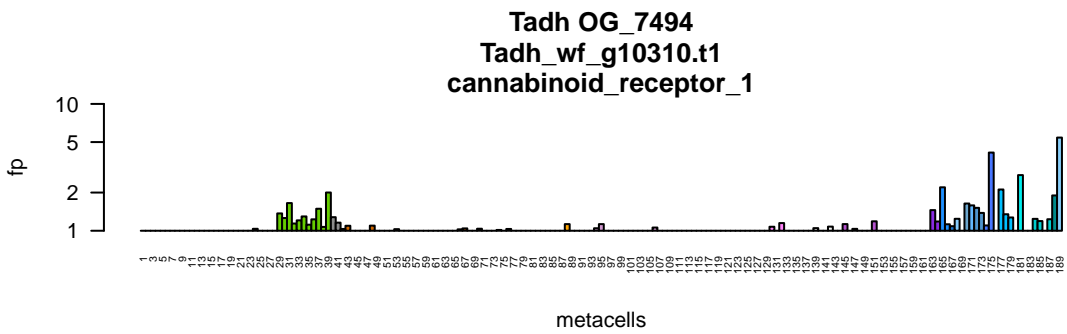


HoiH23 OG_6355
HoiH23_PIH23_012003-RA
_protein_coupled_receptor_15,cholinergic_receptor_muscarinic_3,neuropeptide_Y_recept

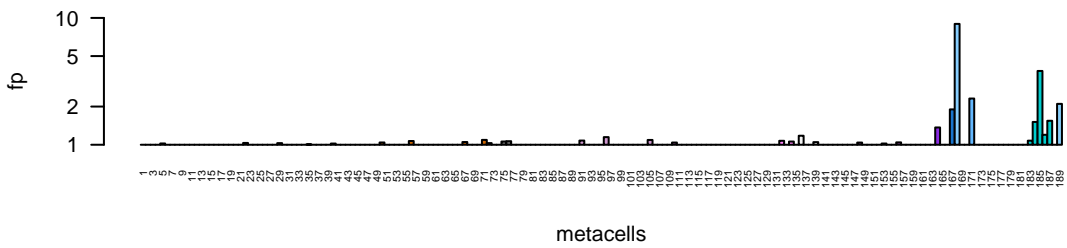




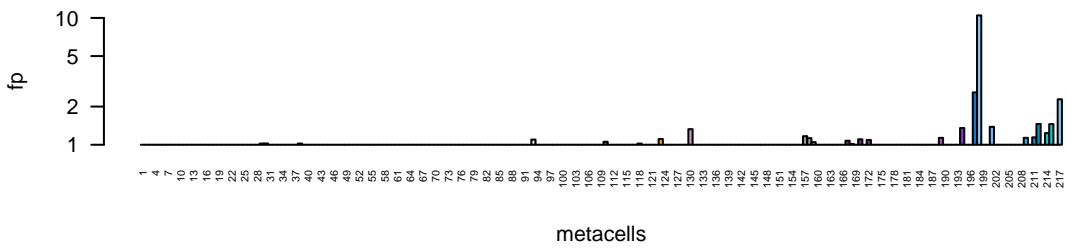




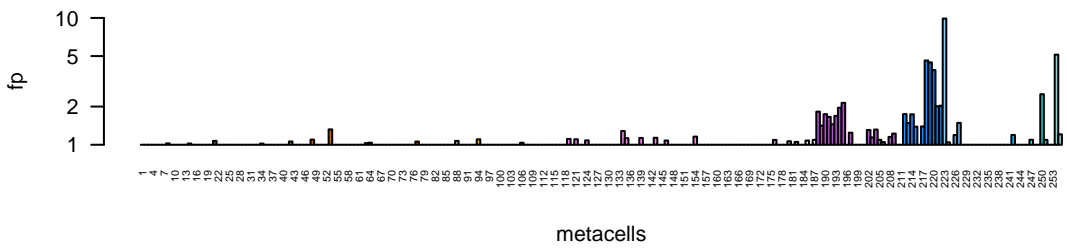
Tadh OG_7537
Tadh_wf_g5232.t1



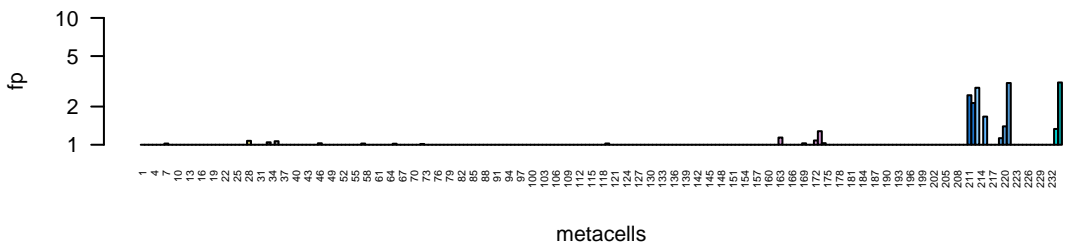
TrH2 OG_7537
TrH2_TrispH2_001160-RA

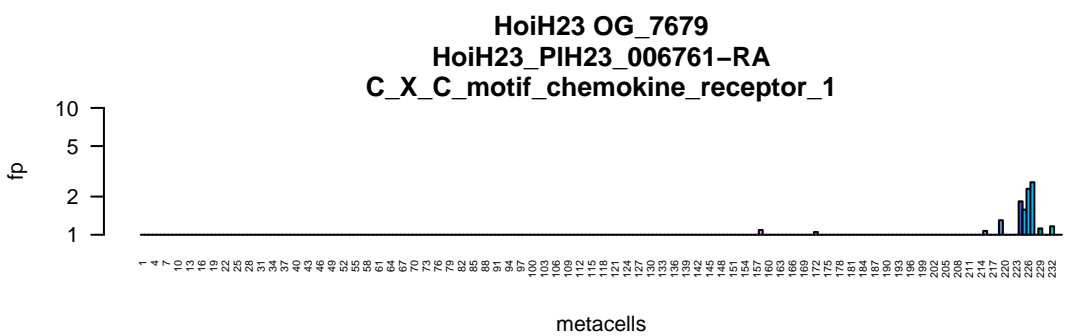
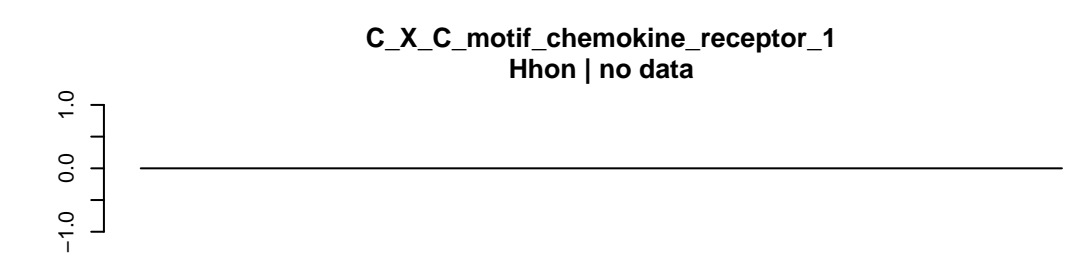
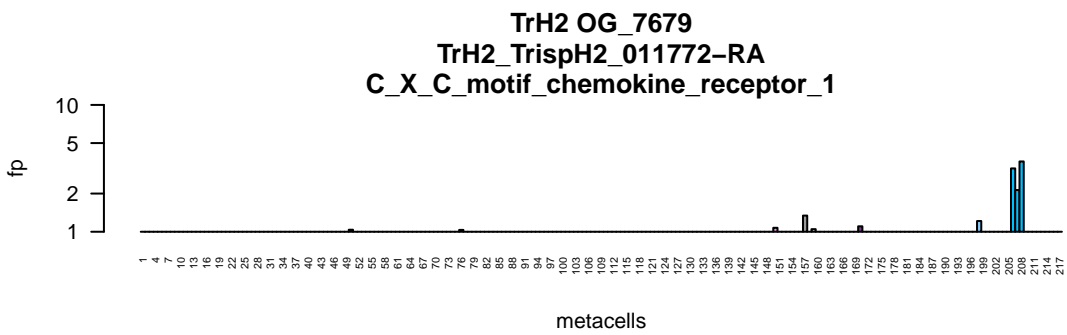
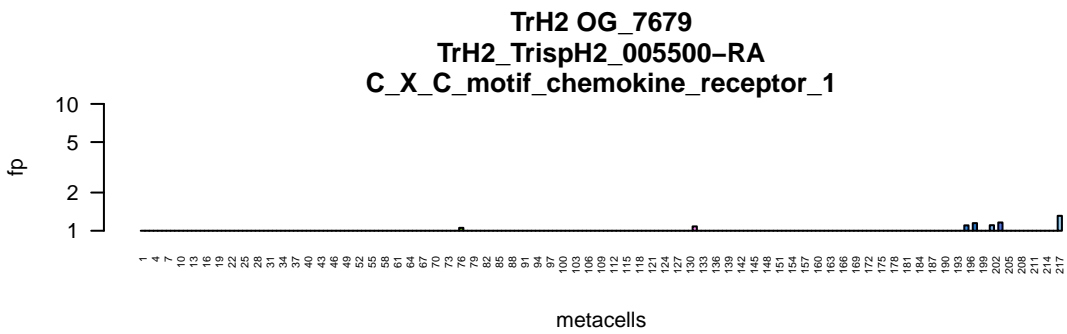
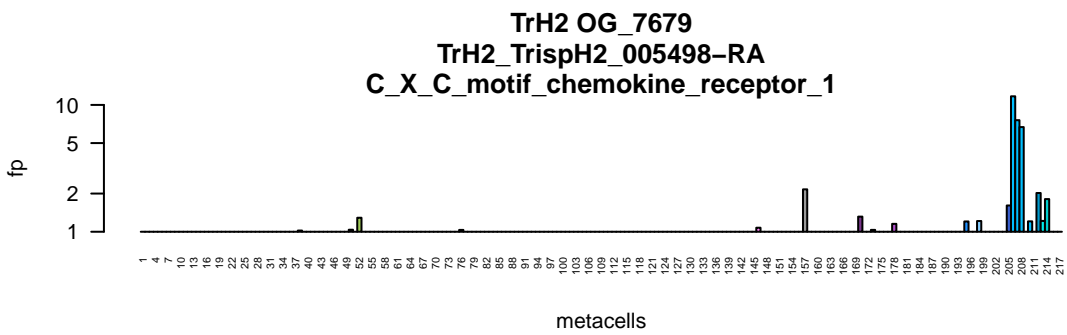
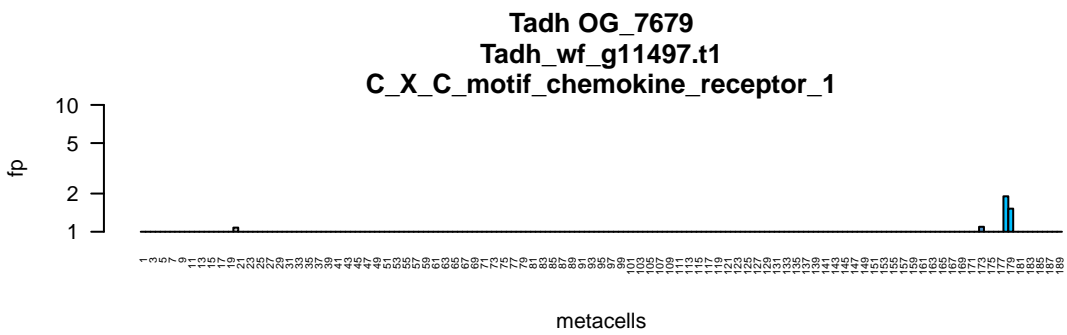
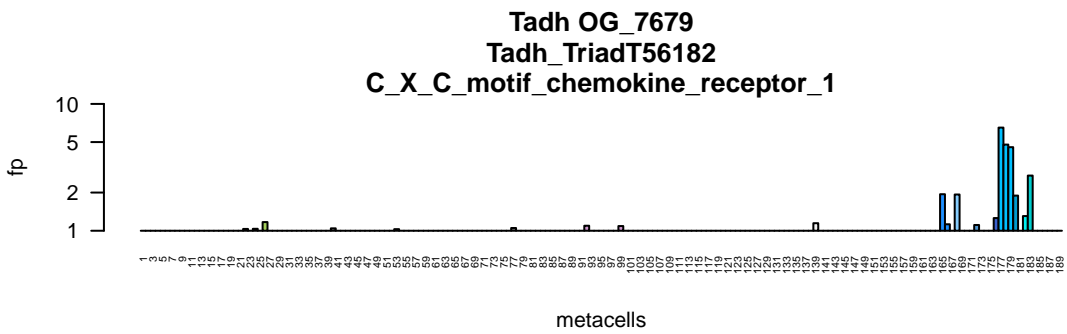


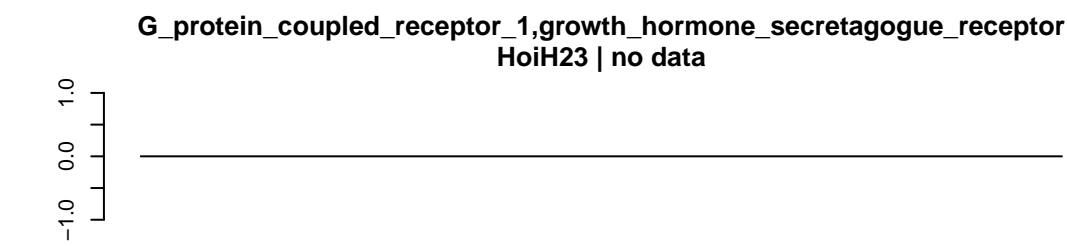
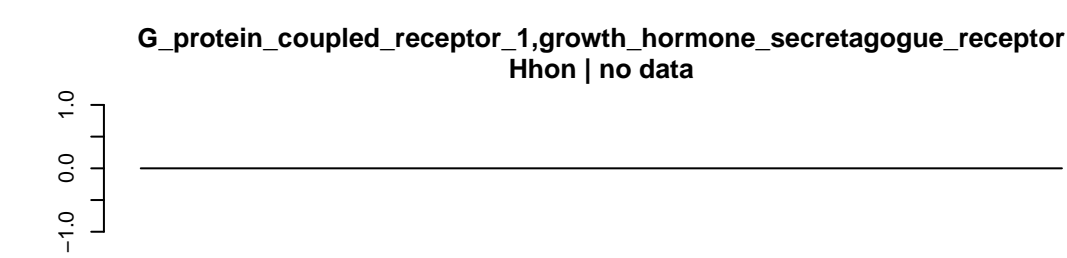
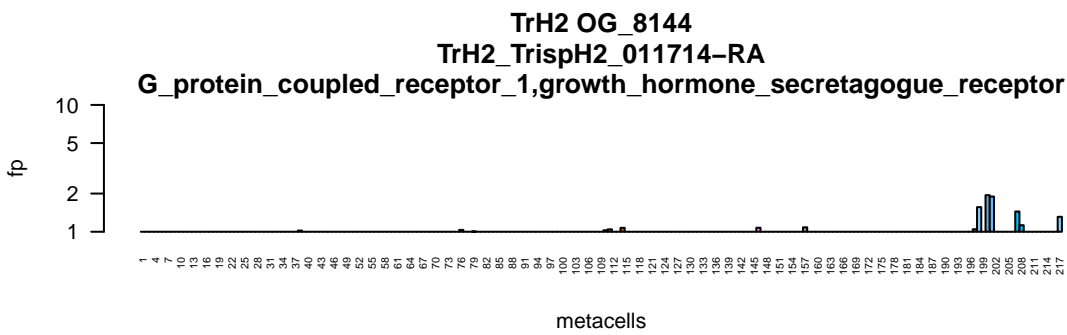
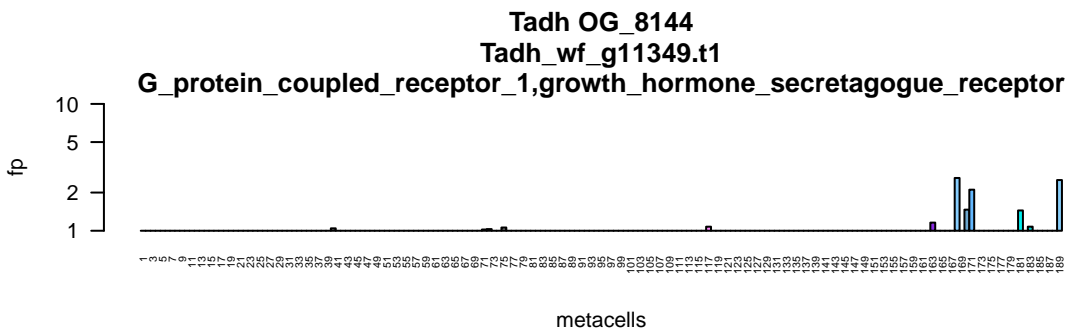
Hhon OG_7537
Hhon_g10209.t1

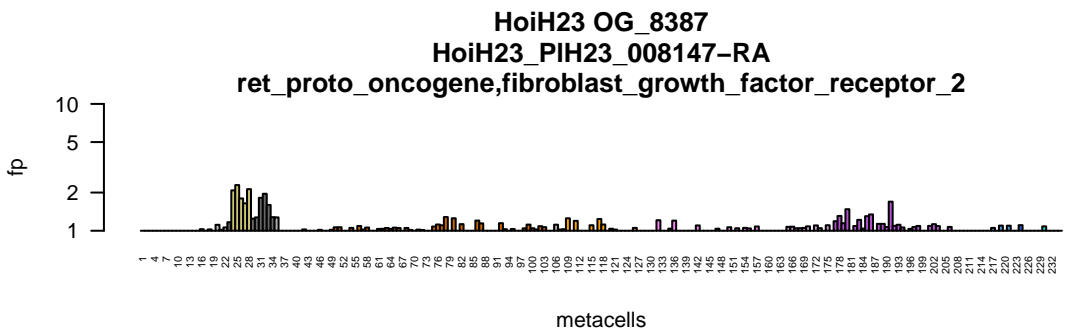
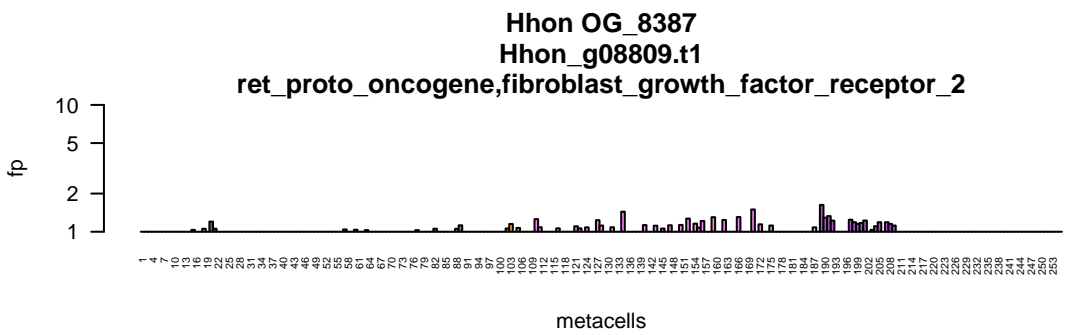
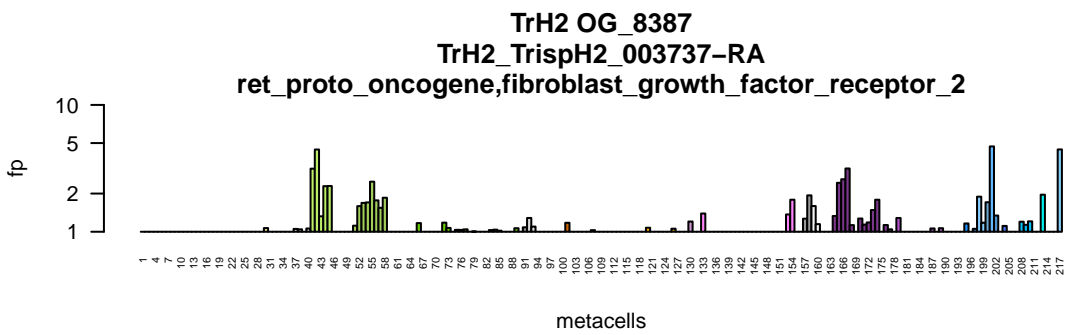
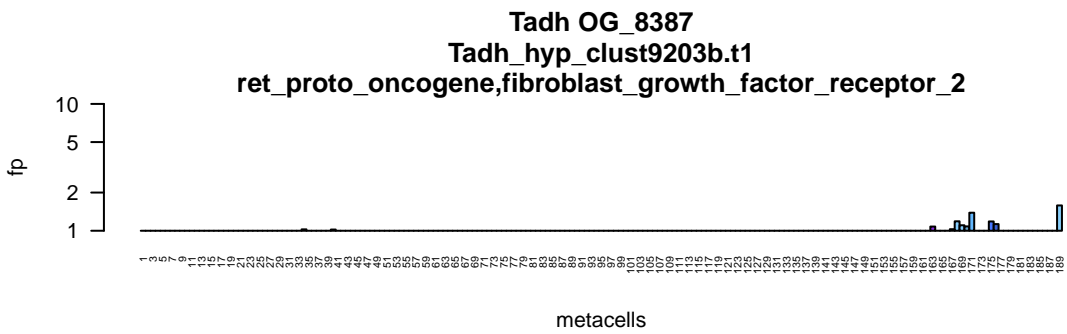


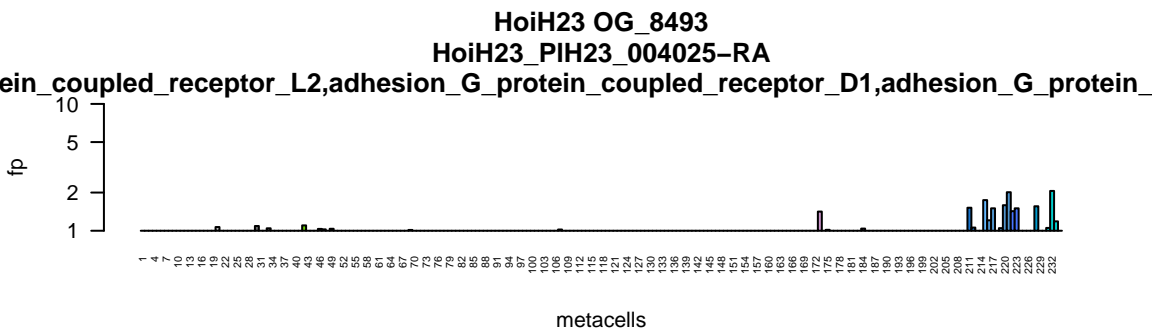
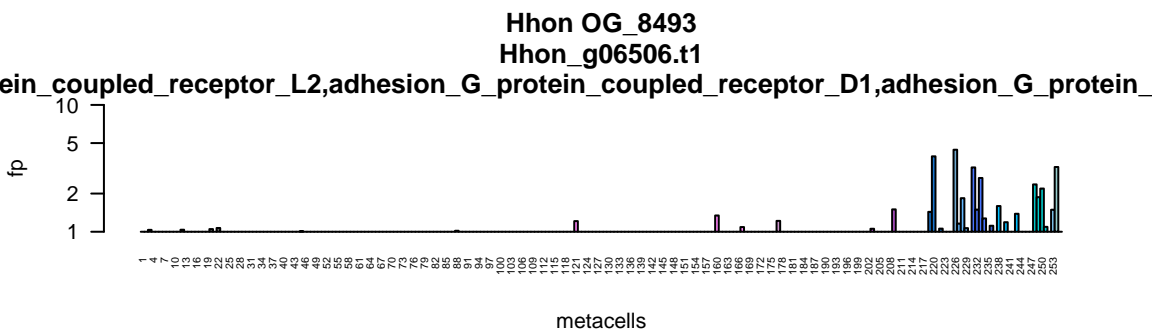
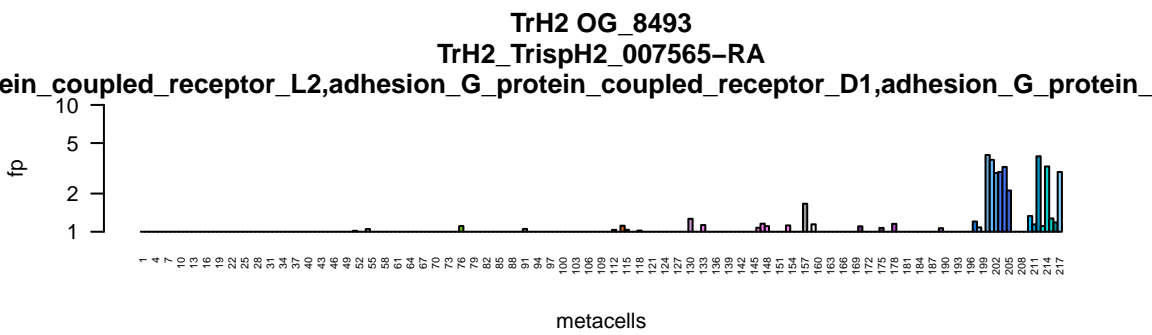
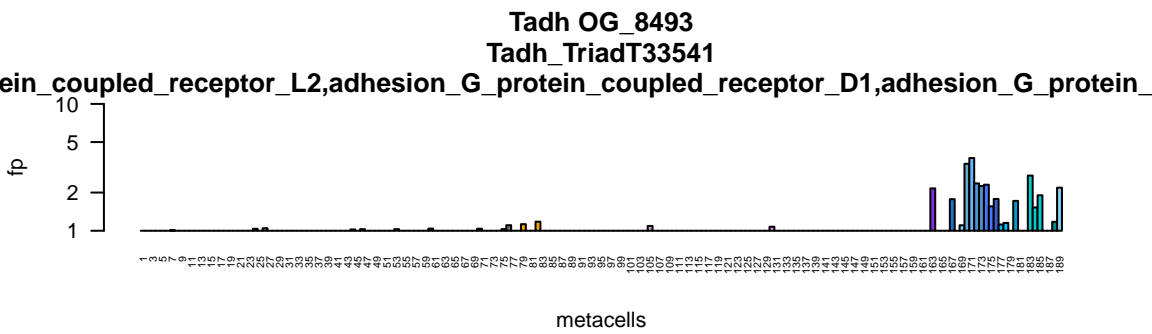
HoiH23 OG_7537
HoiH23_PIH23_000241-RA

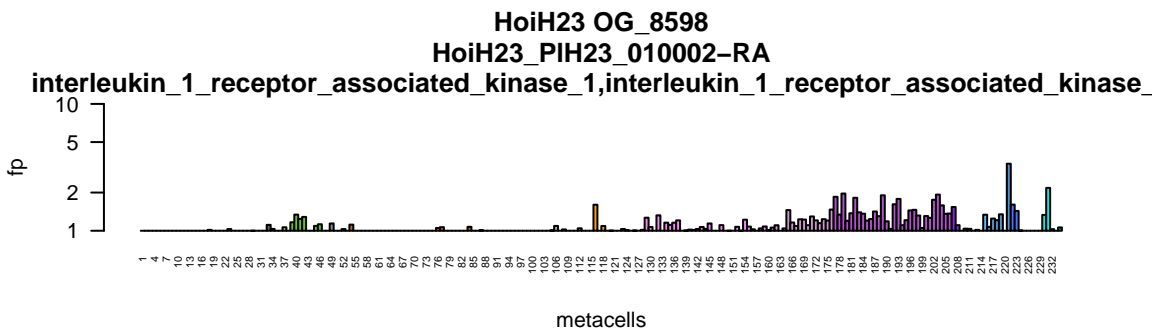
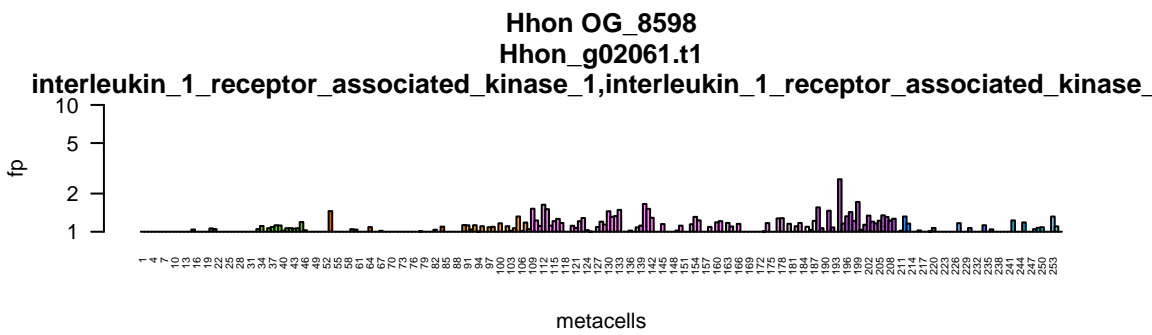
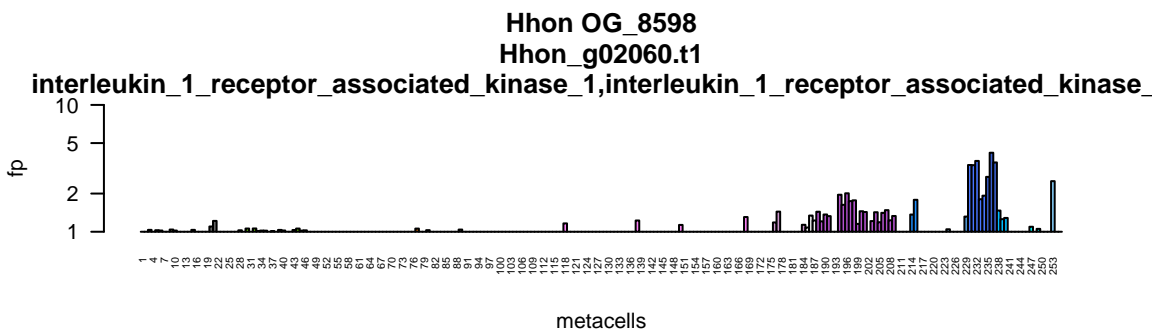
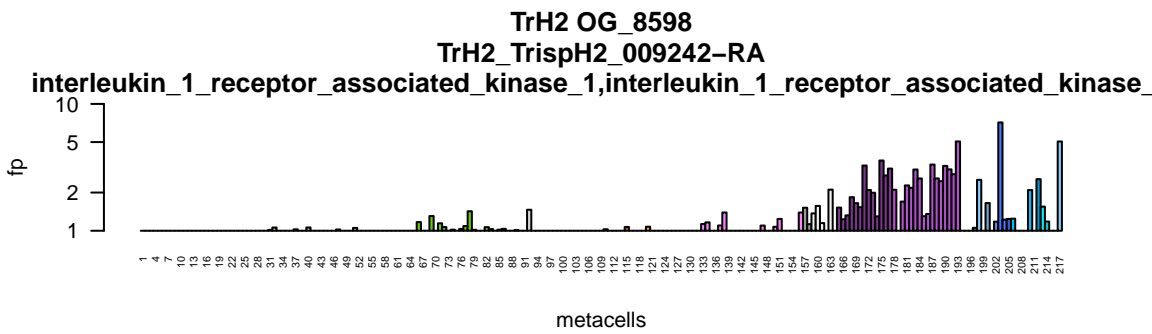
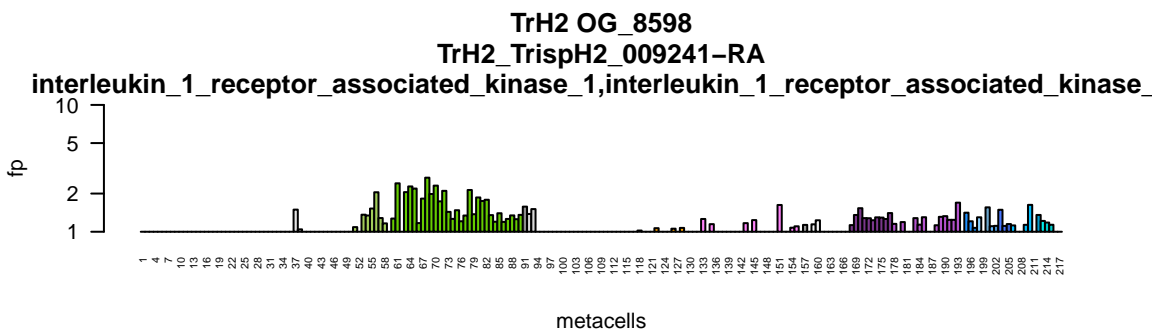
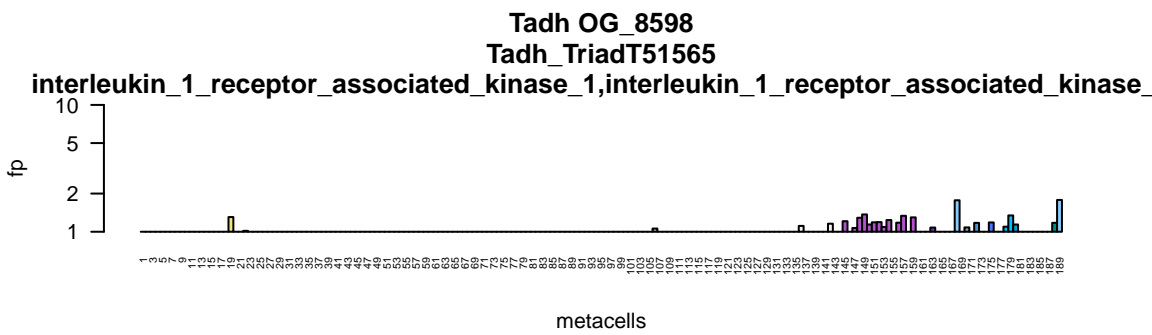
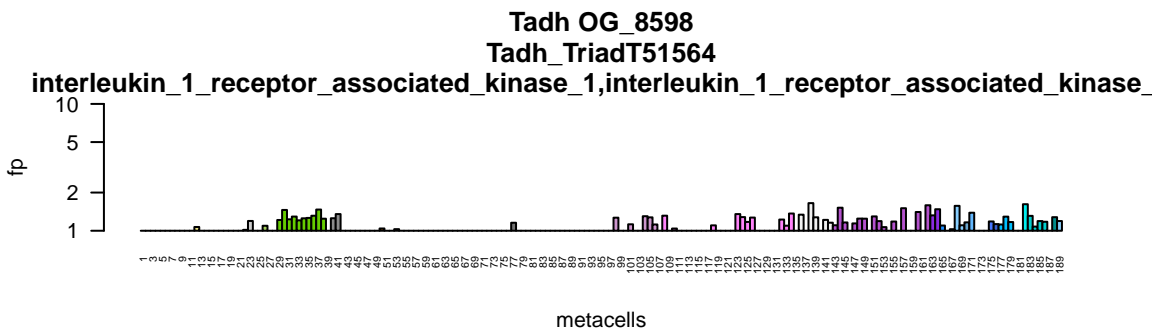




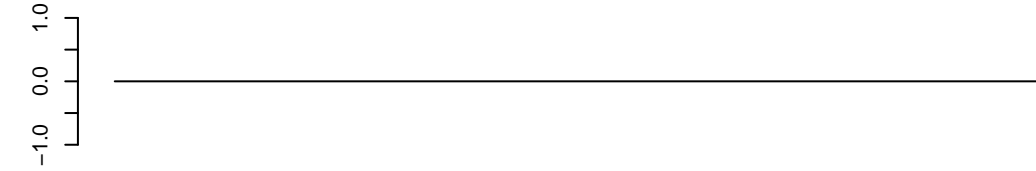




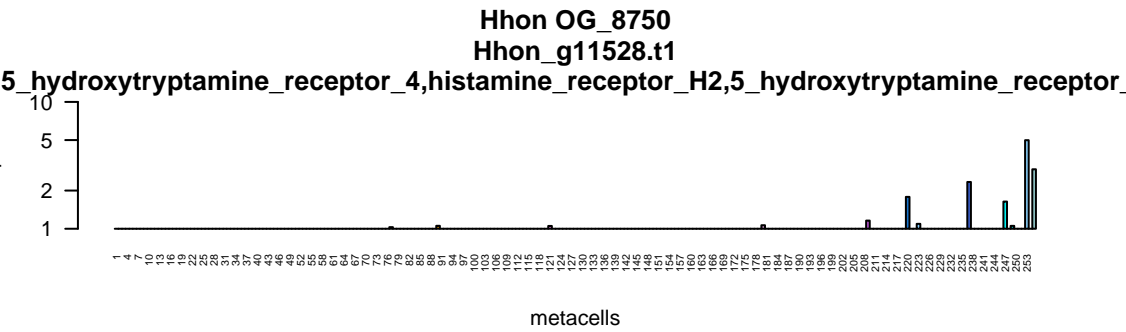
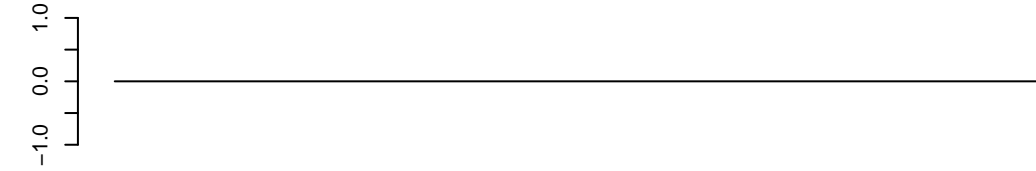




5_hydroxytryptamine_receptor_4,histamine_receptor_H2,5_hydroxytryptamine_receptor_
Tadh | no data

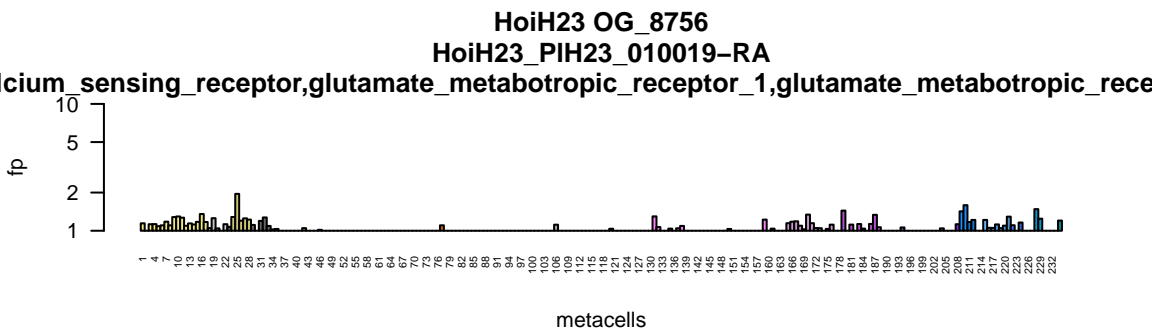
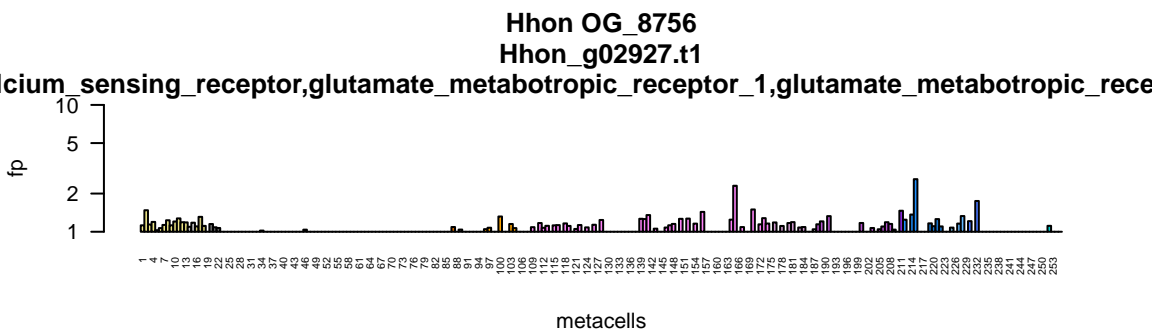
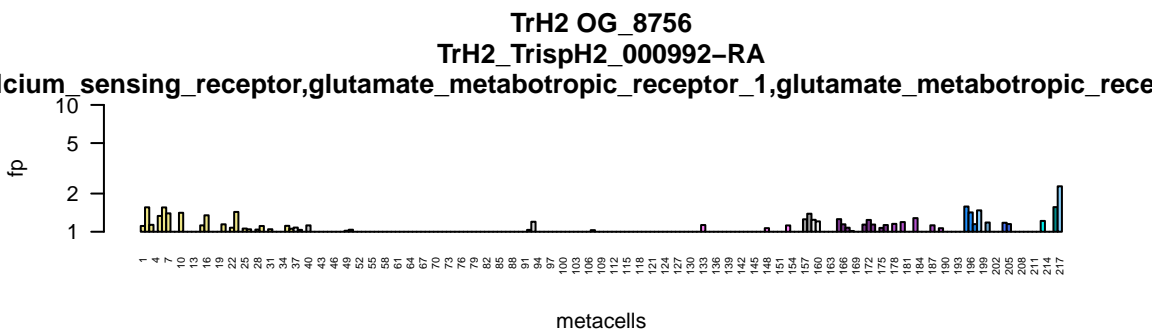
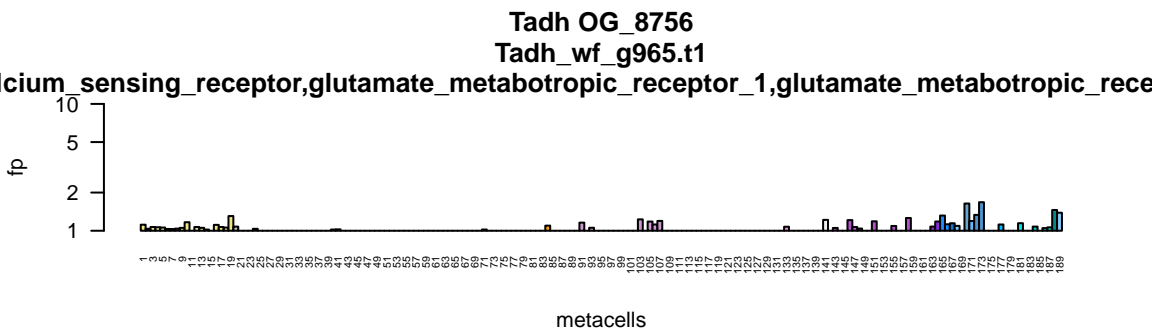


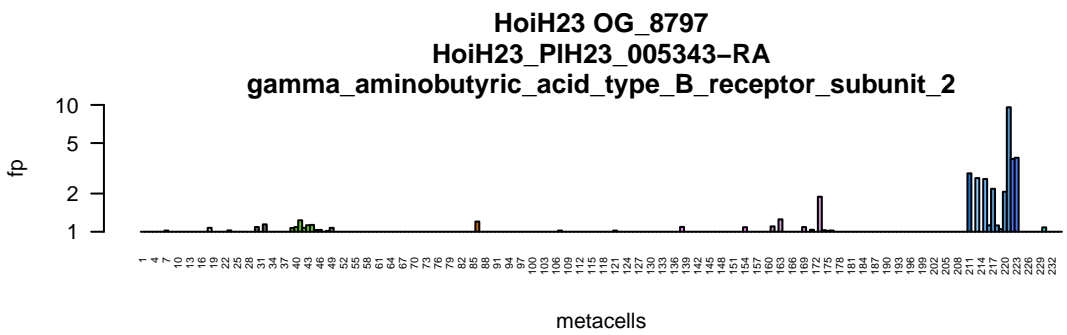
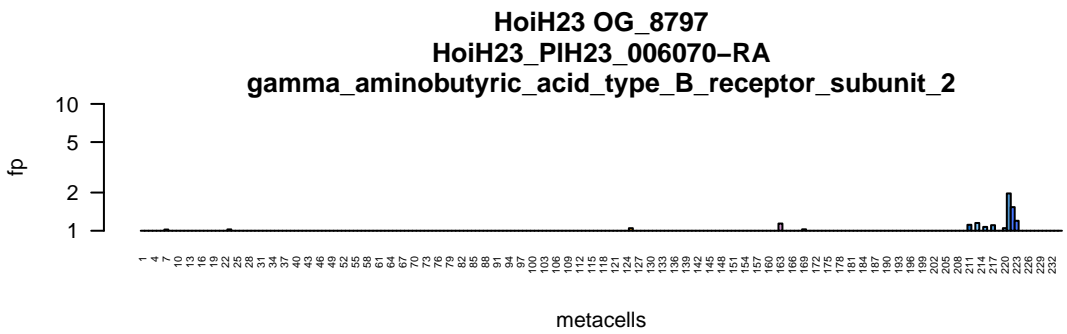
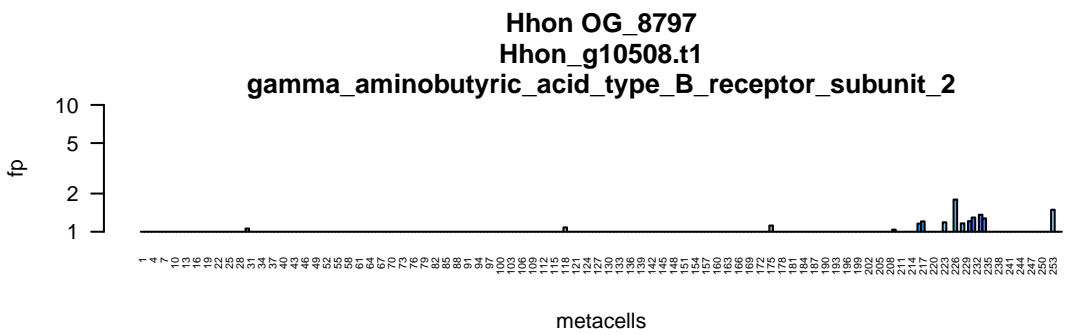
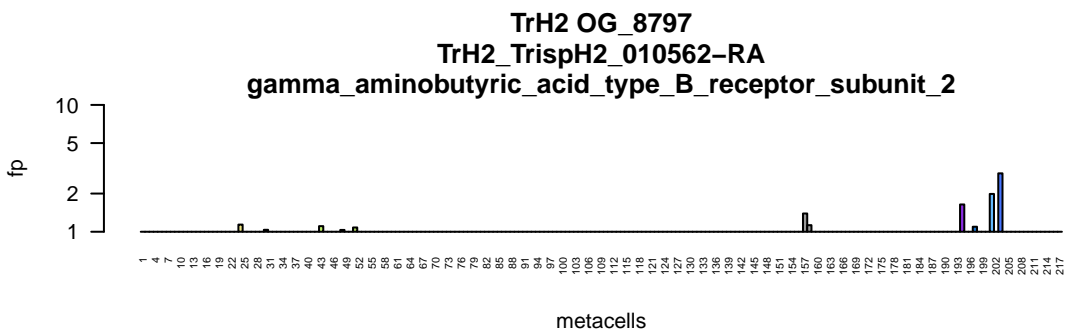
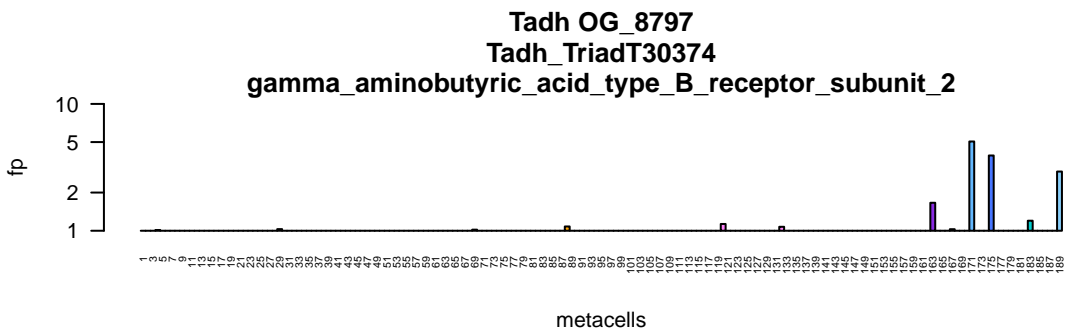
5_hydroxytryptamine_receptor_4,histamine_receptor_H2,5_hydroxytryptamine_receptor_
TrH2 | no data



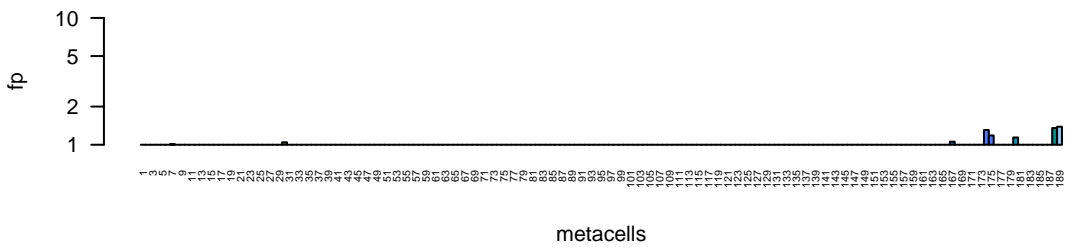
5_hydroxytryptamine_receptor_4,histamine_receptor_H2,5_hydroxytryptamine_receptor_
HoiH23 | no data







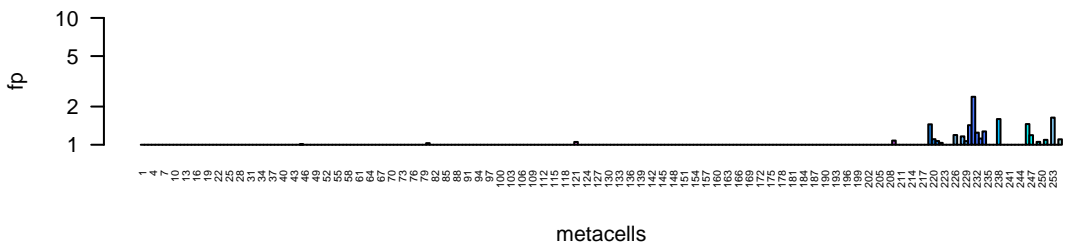
Tadh OG_9610
Tadh_TriadT60281



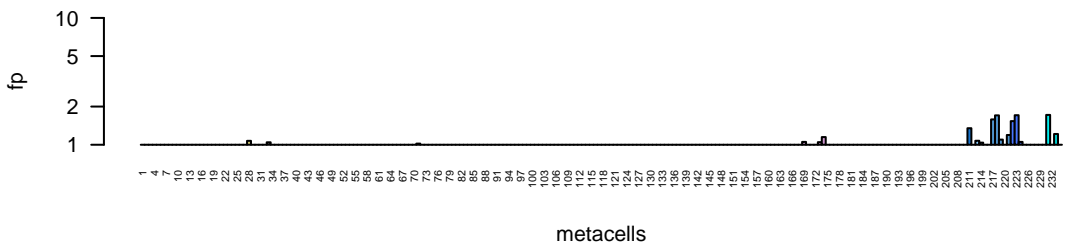
TrH2 | no data

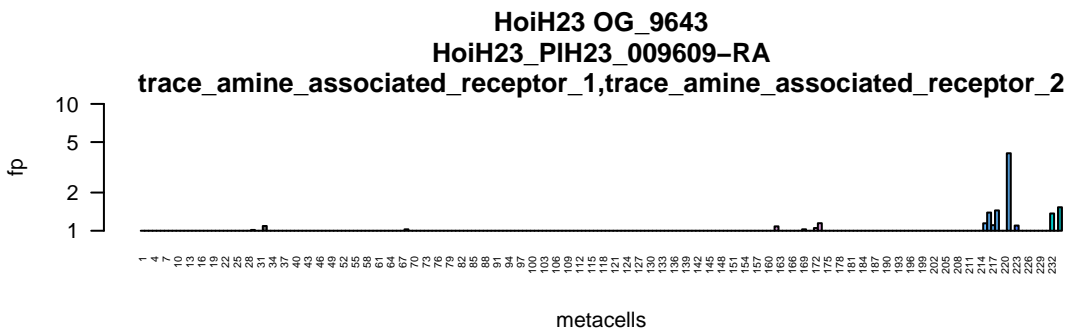
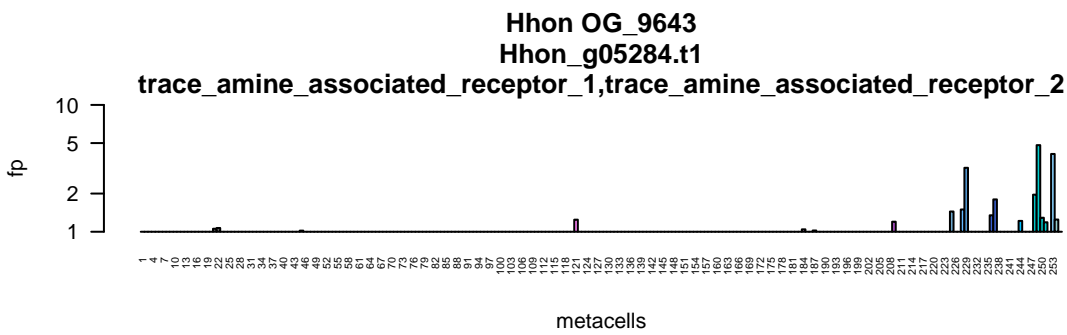
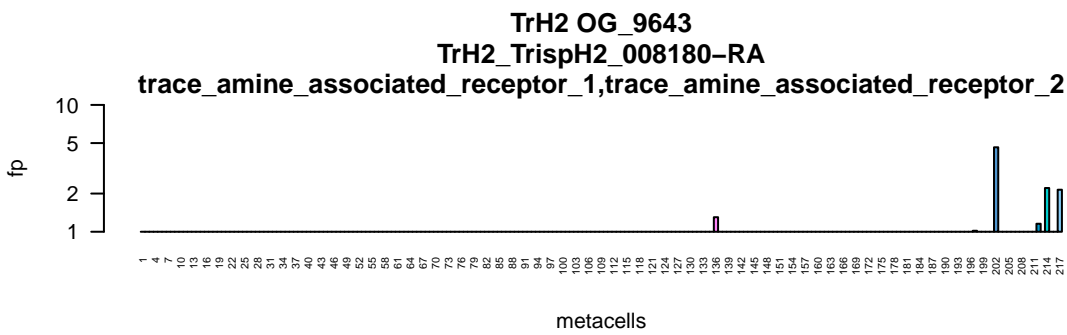
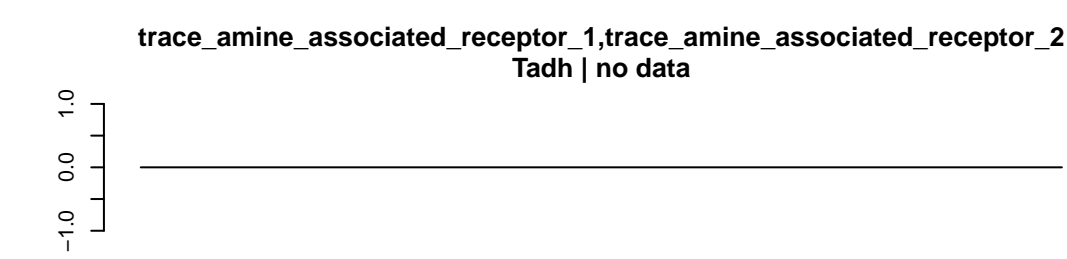


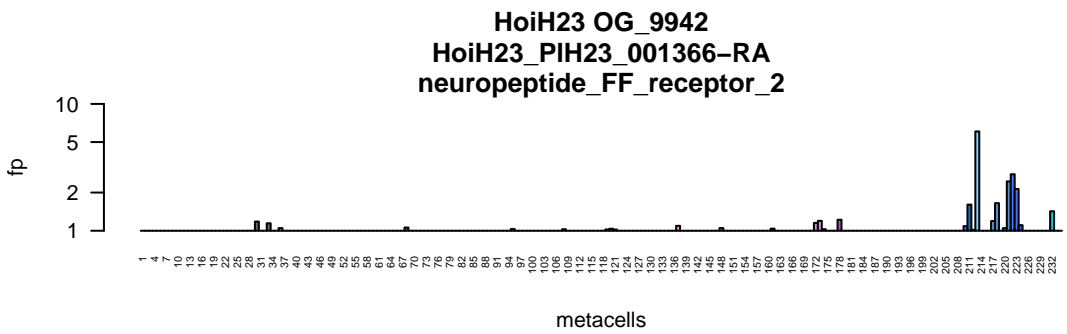
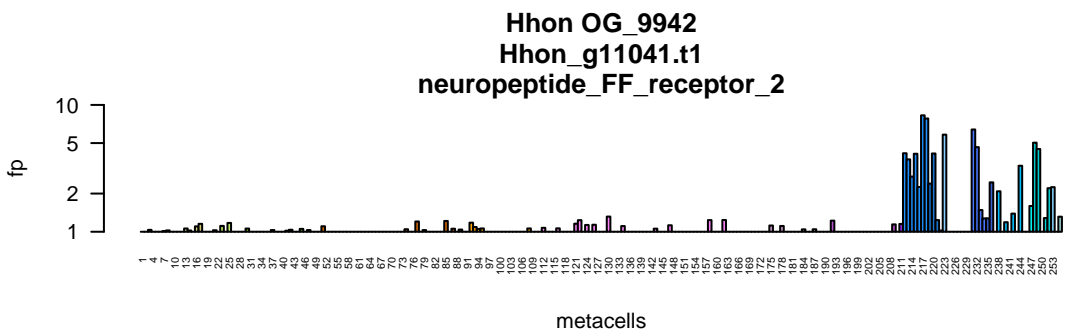
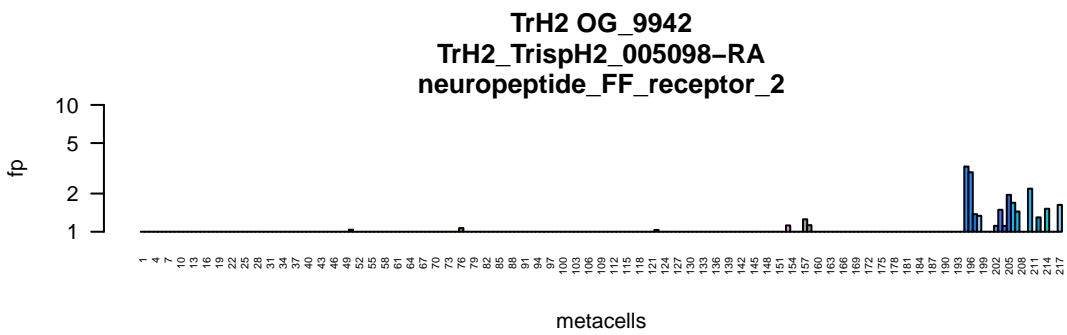
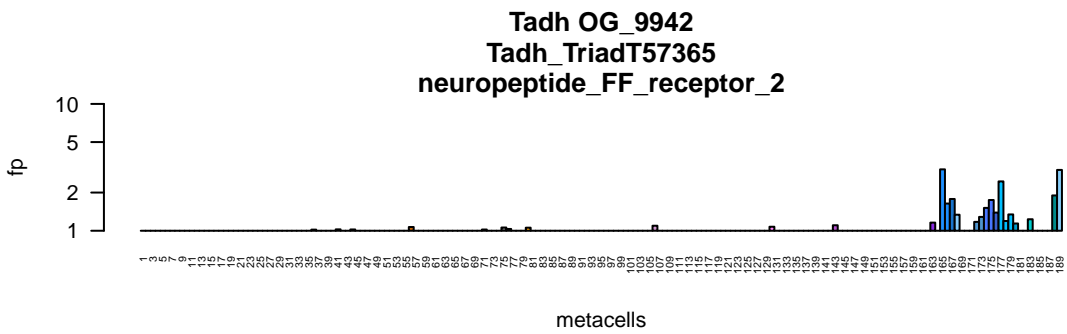
Hhon OG_9610
Hhon_g08104.t1

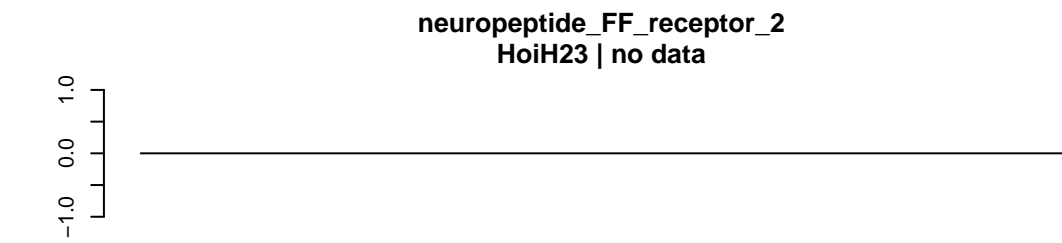
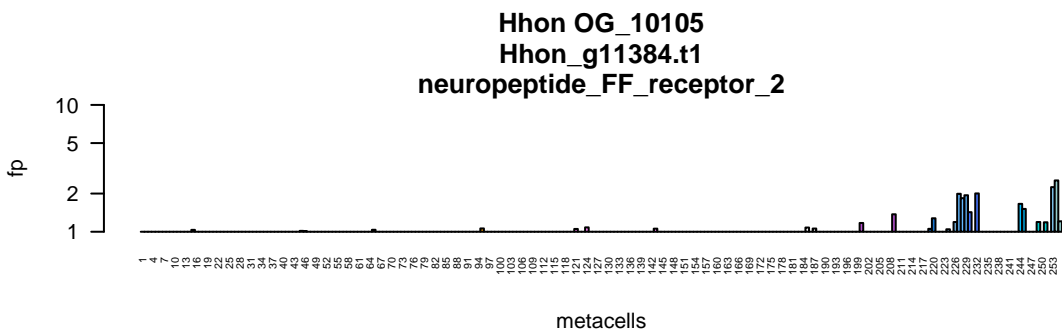
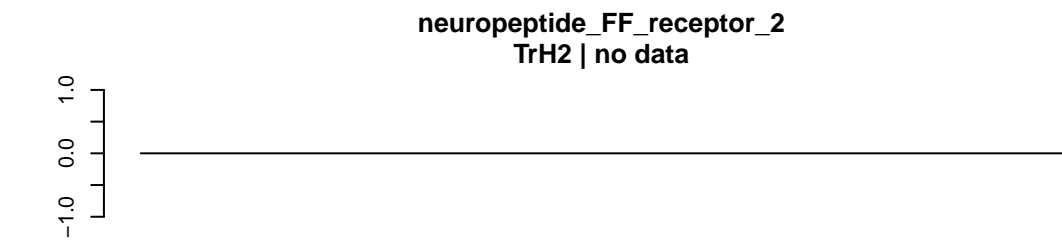
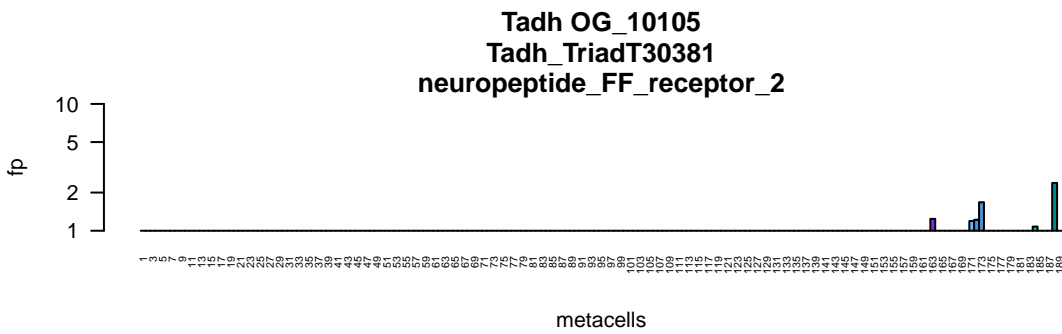


HoiH23 OG_9610
HoiH23_PIH23_005684-RA





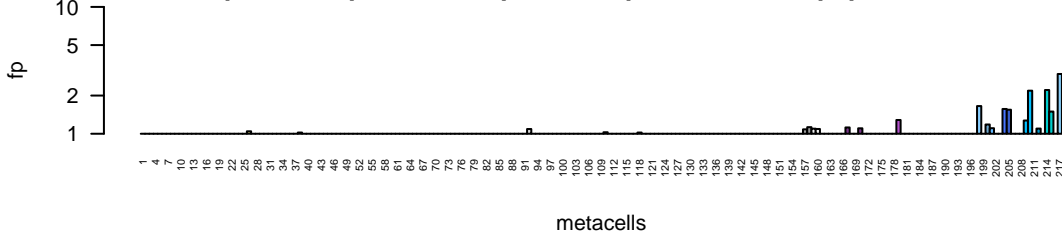




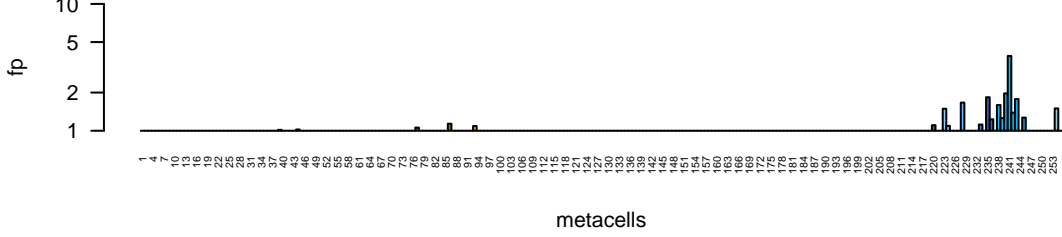
omatostatin_receptor_2,G_protein_coupled_receptor_183,neuropeptides_B_and_W_recept
Tadh | no data



omatostatin_receptor_2,G_protein_coupled_receptor_183,neuropeptides_B_and_W_recept
TrH2 OG_10233
TrH2_TrispH2_006452-RA



omatostatin_receptor_2,G_protein_coupled_receptor_183,neuropeptides_B_and_W_recept
Hhon OG_10233
Hhon_g05148.t1



omatostatin_receptor_2,G_protein_coupled_receptor_183,neuropeptides_B_and_W_recept
HoiH23 OG_10233
HoiH23_PIH23_001231-RA

