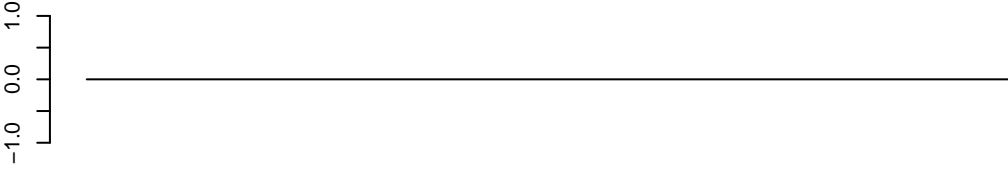


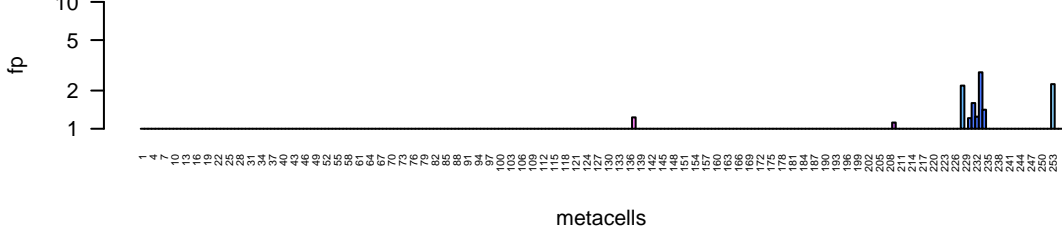
gamma_aminobutyric_acid_type_B_receptor_subunit_2
Tadh | no data



gamma_aminobutyric_acid_type_B_receptor_subunit_2
TrH2 | no data

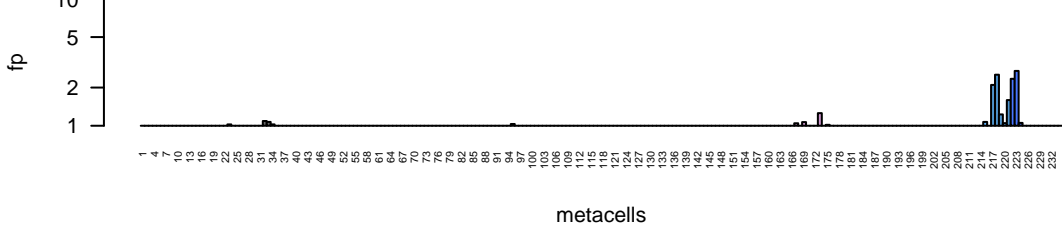


Hhon OG_7236
Hhon_g07333.t1
gamma_aminobutyric_acid_type_B_receptor_subunit_2

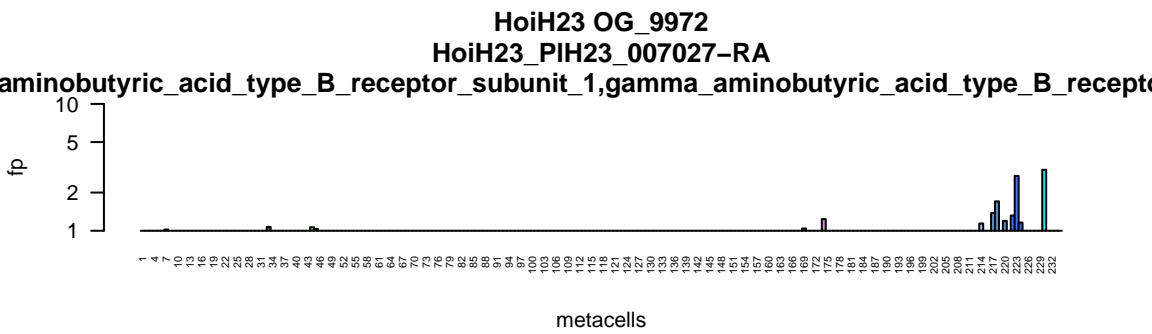
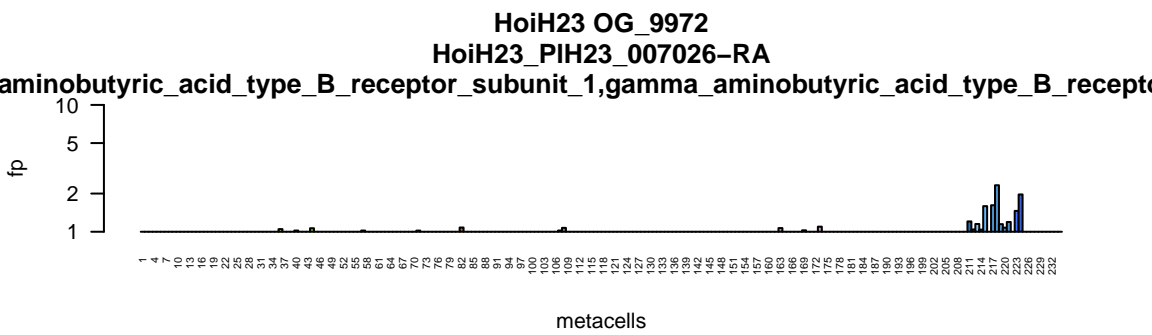
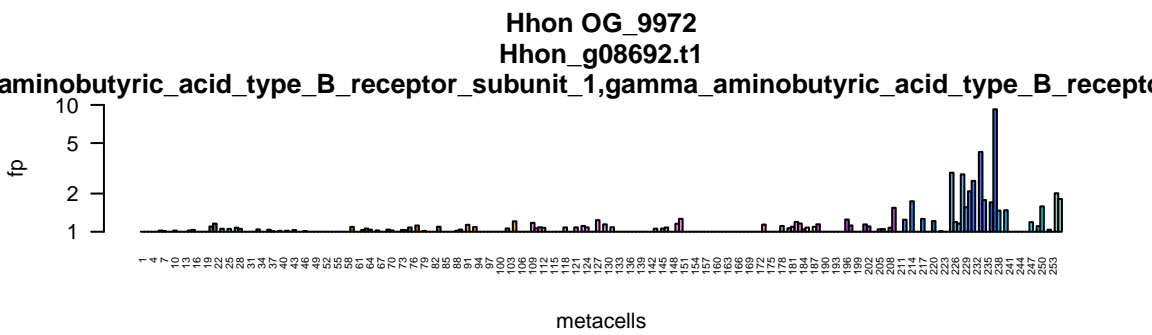
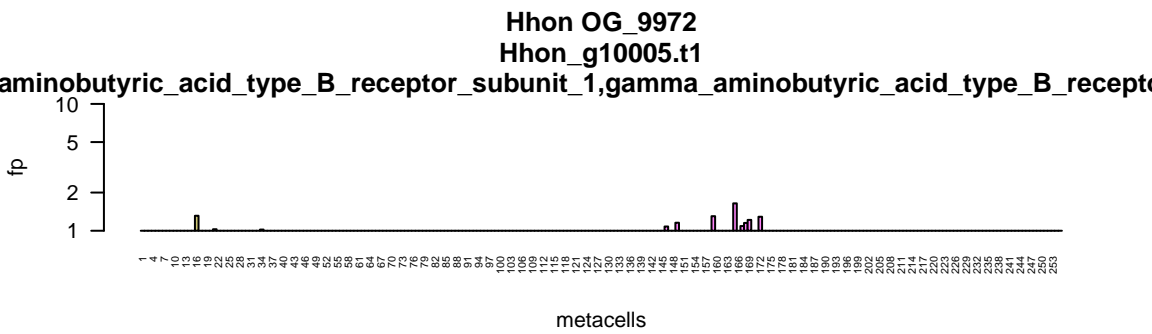
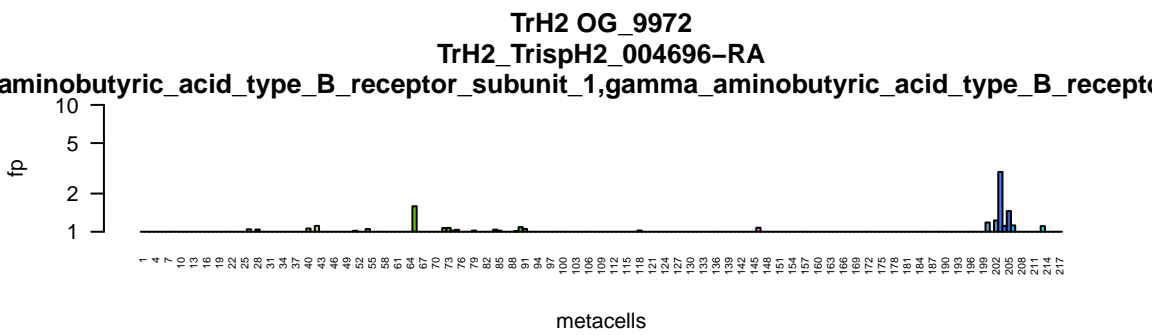
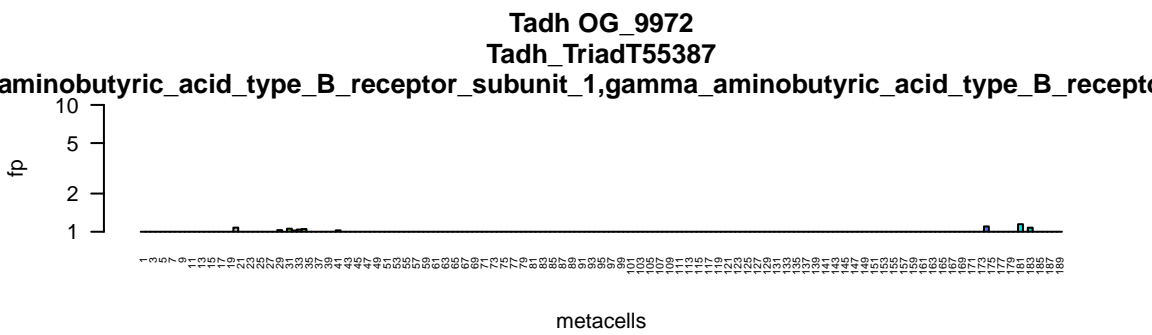
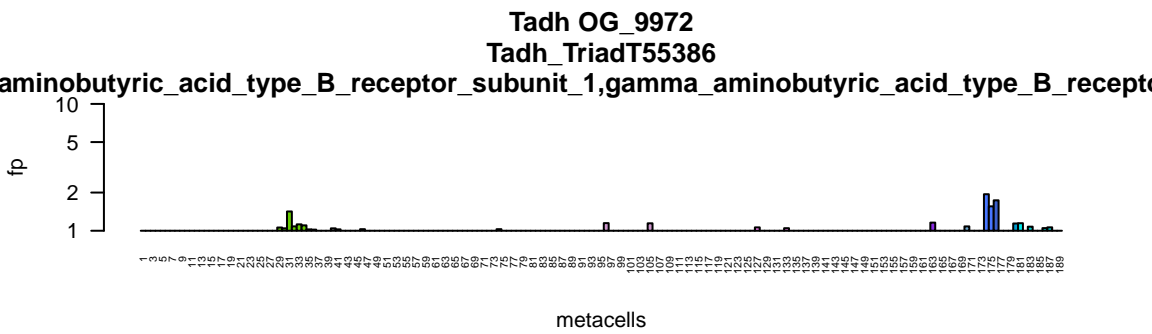


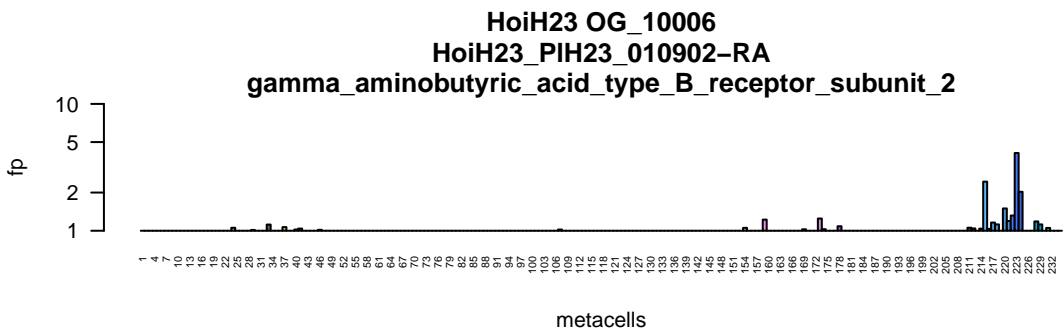
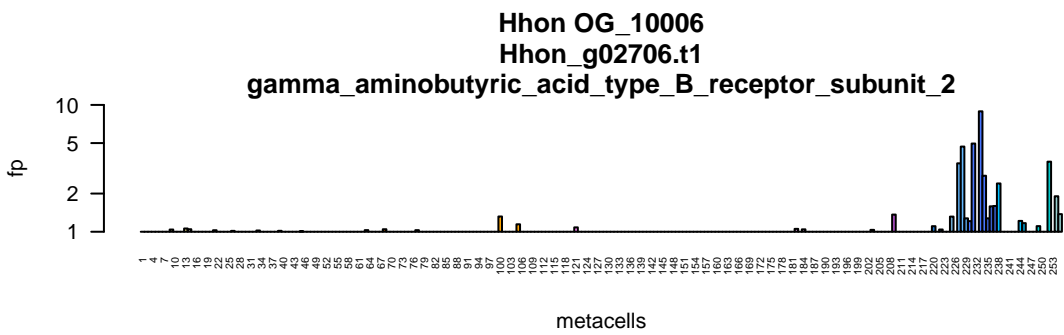
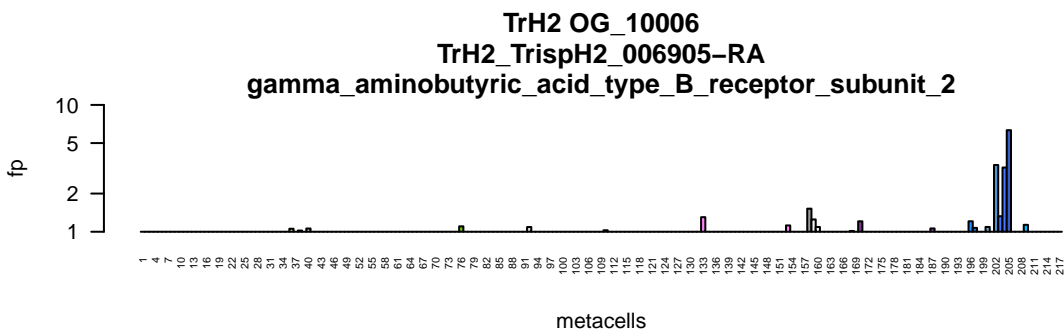
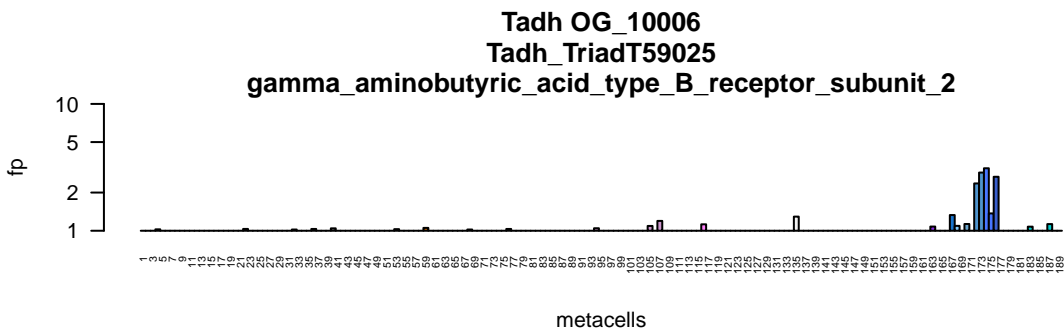
metacells

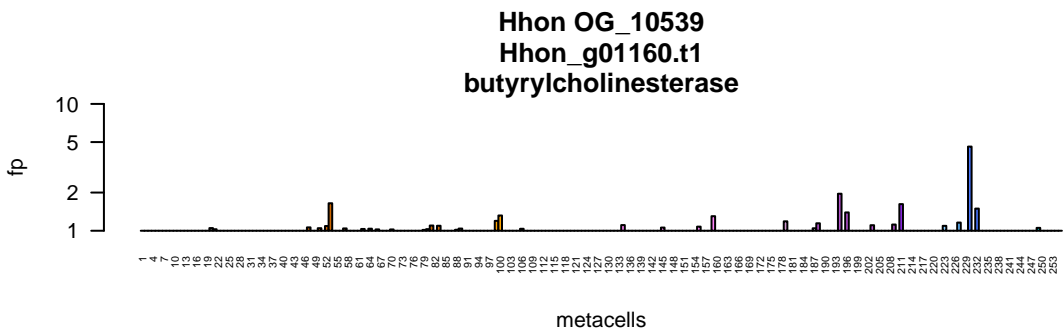
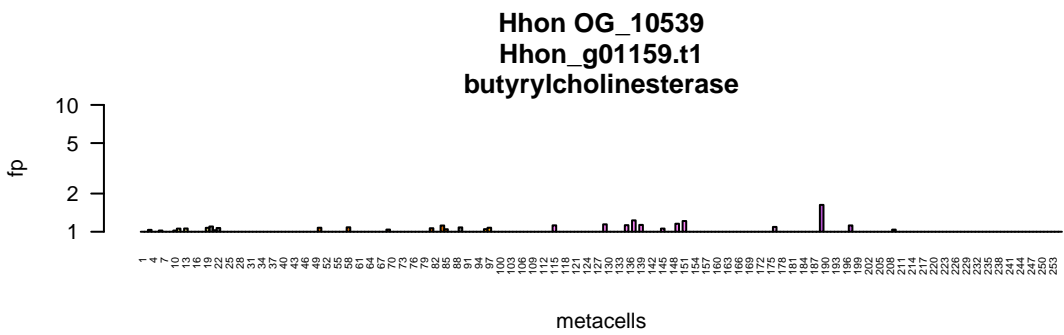
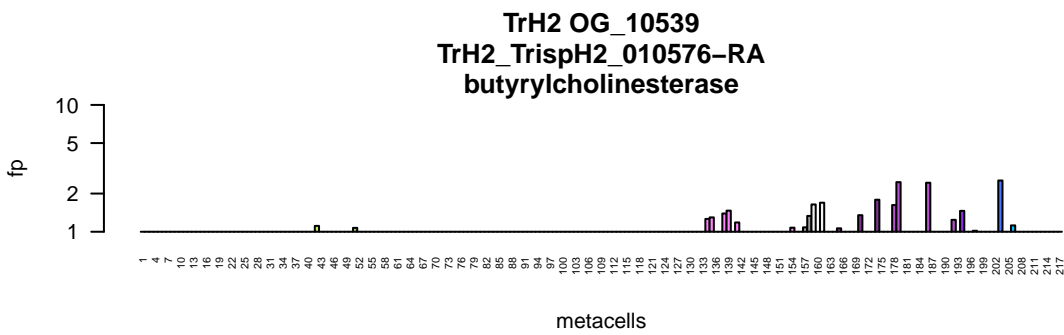
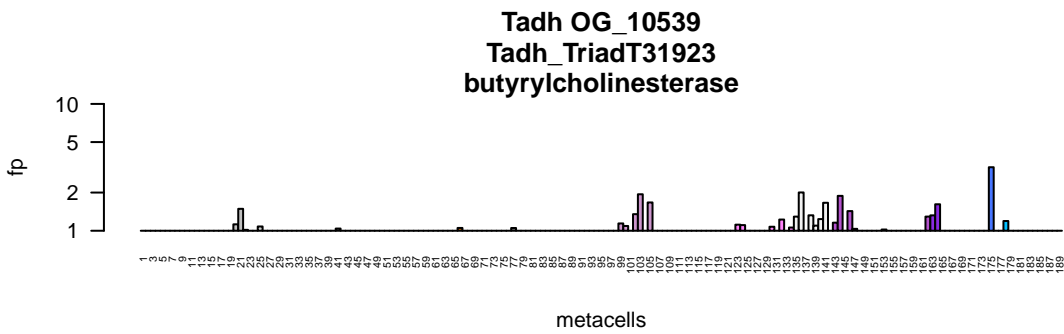
HoiH23 OG_7236
HoiH23_PIH23_003984-RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2

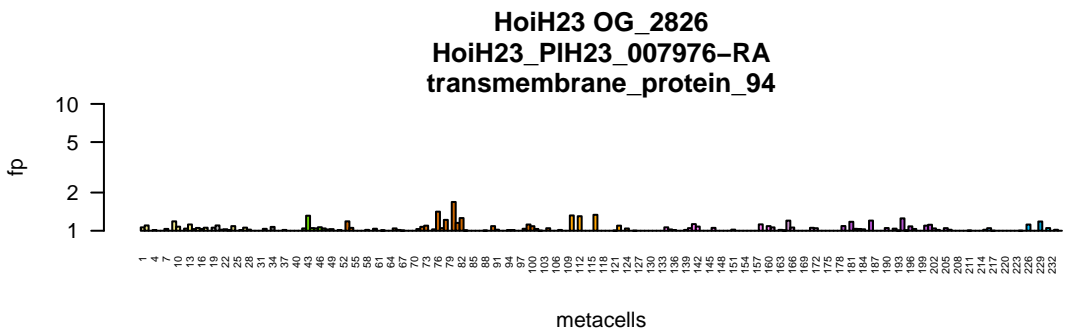
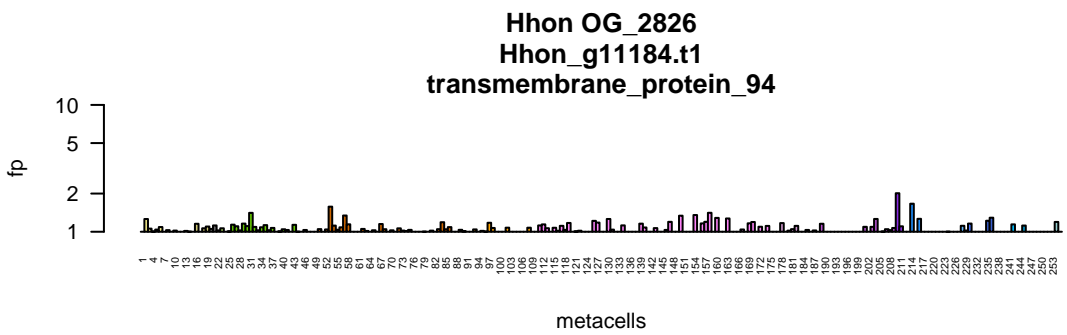
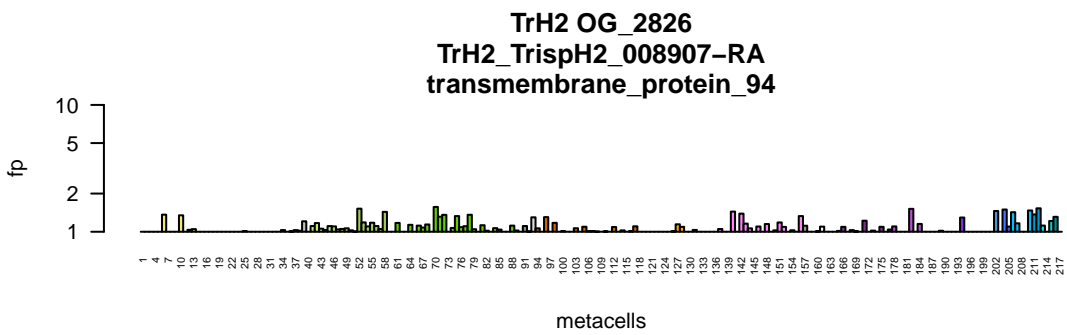
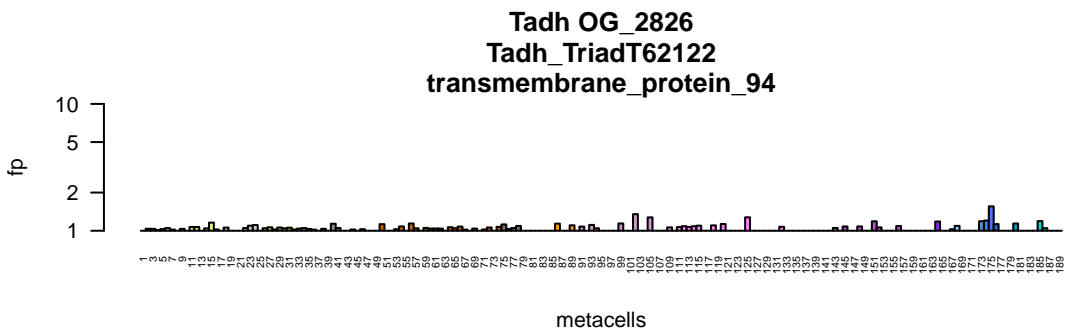


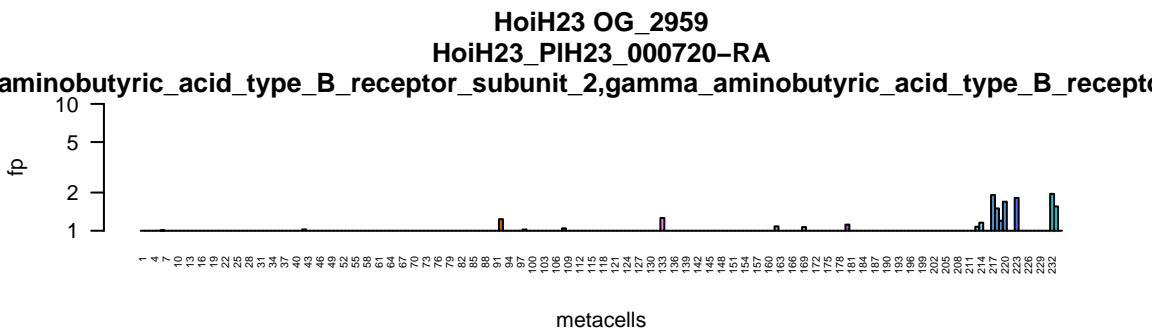
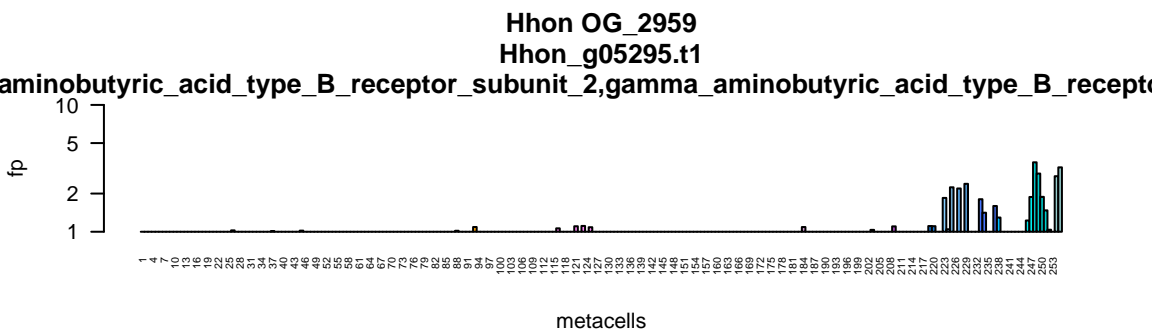
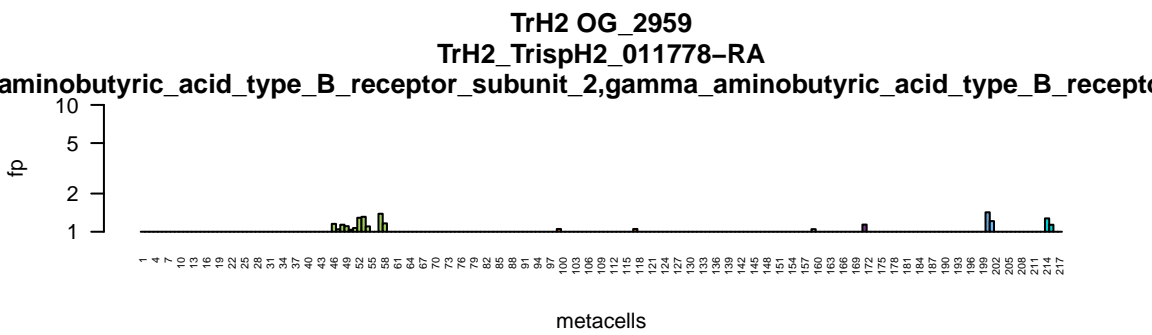
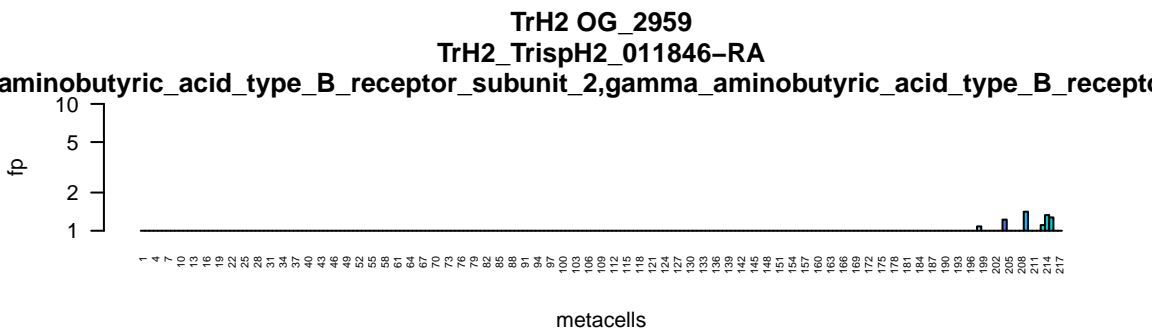
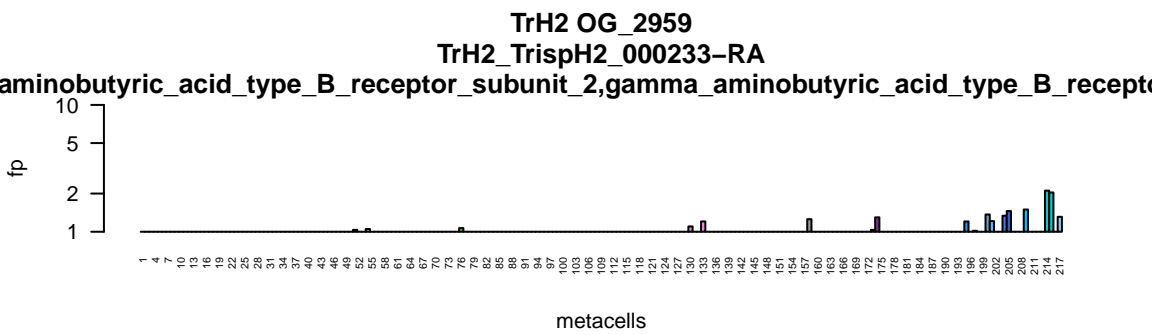
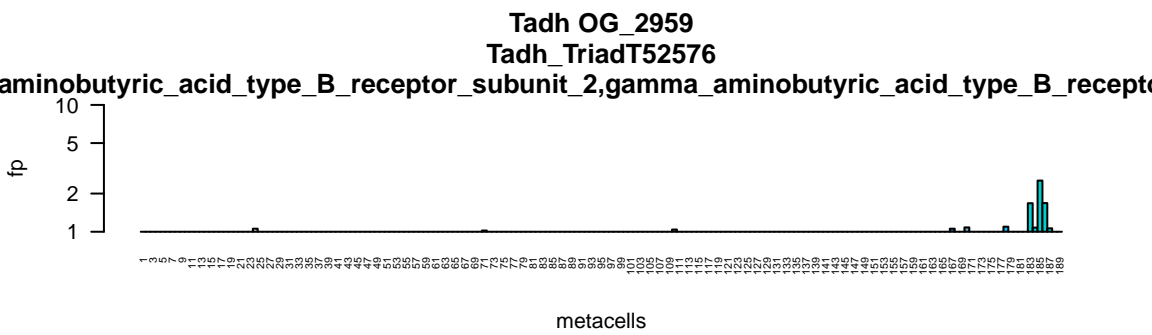
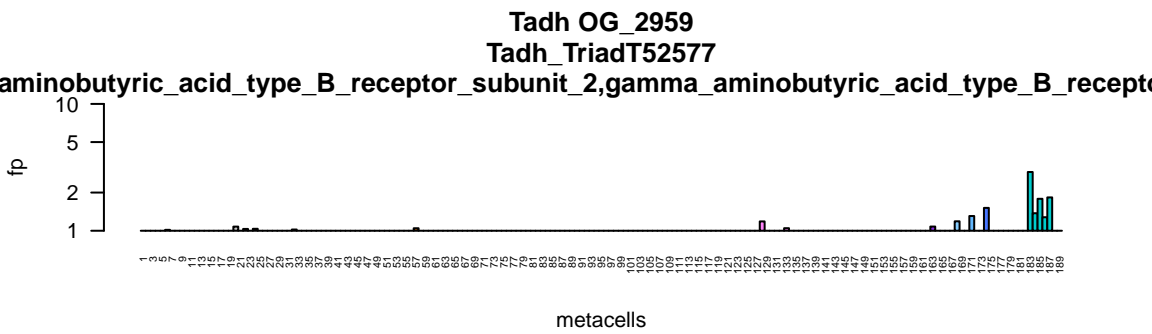
metacells

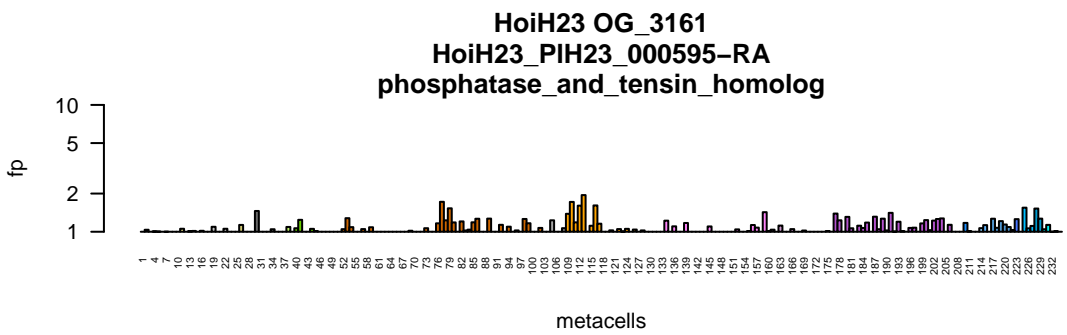
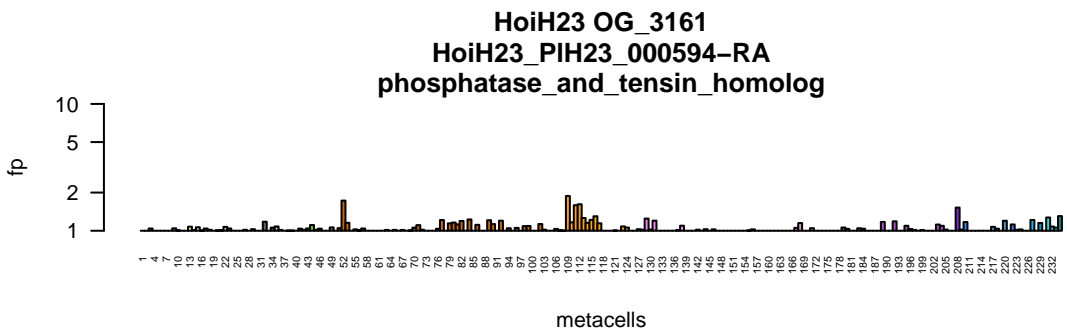
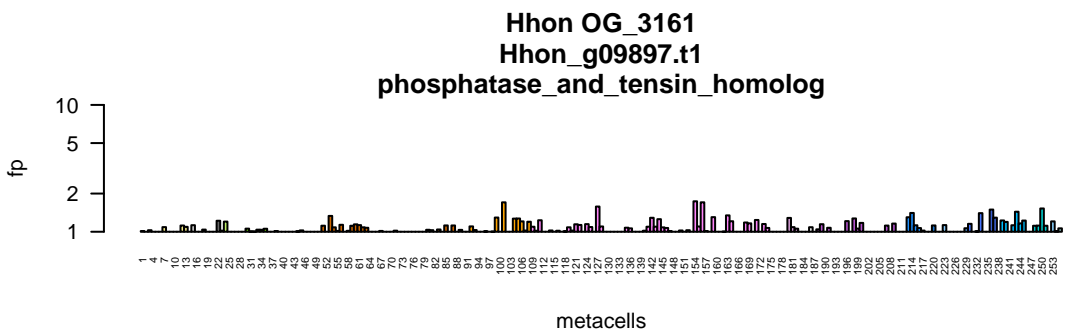
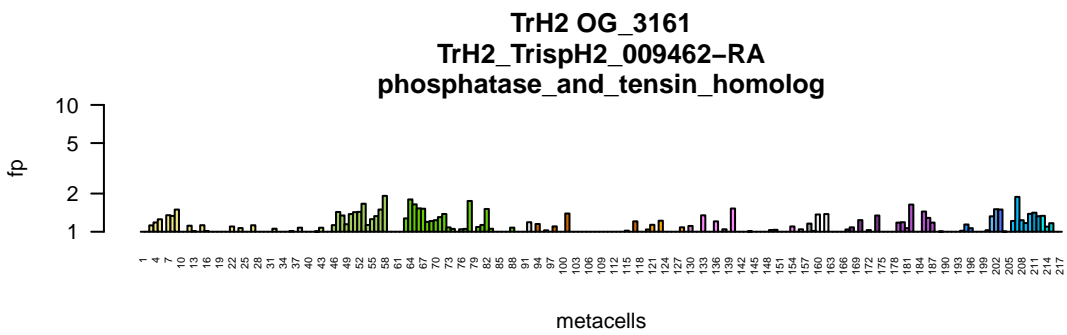
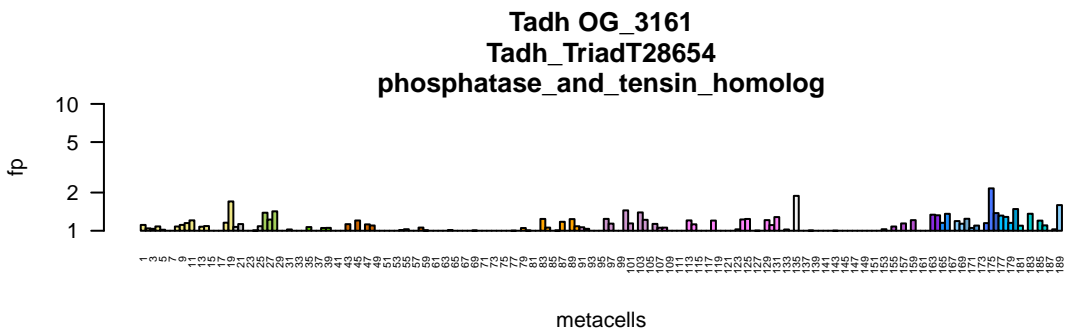


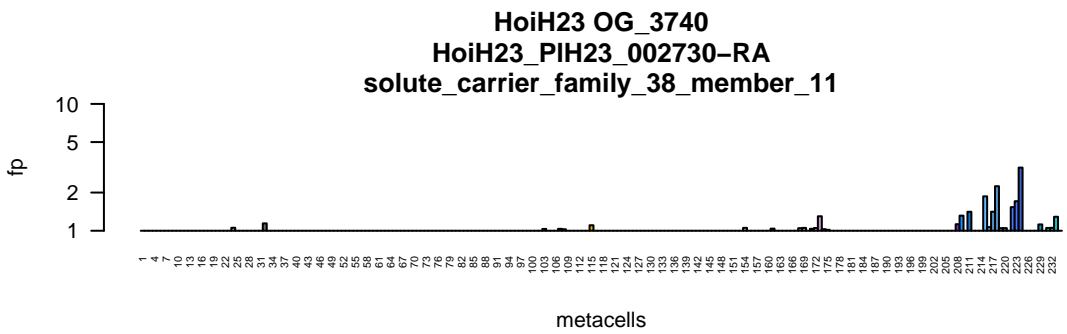
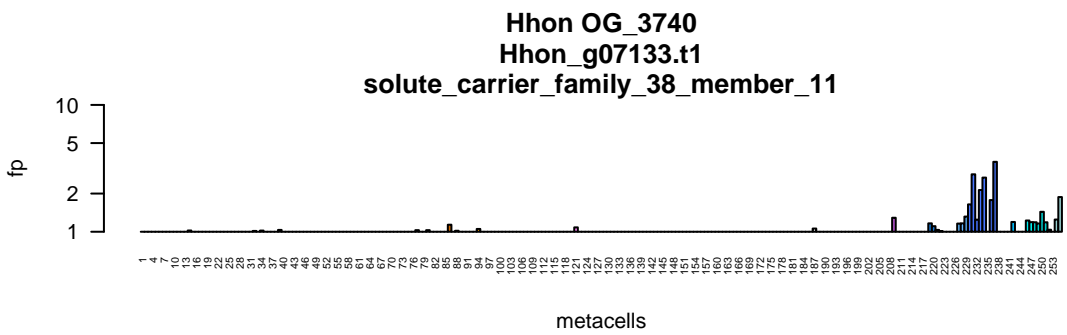
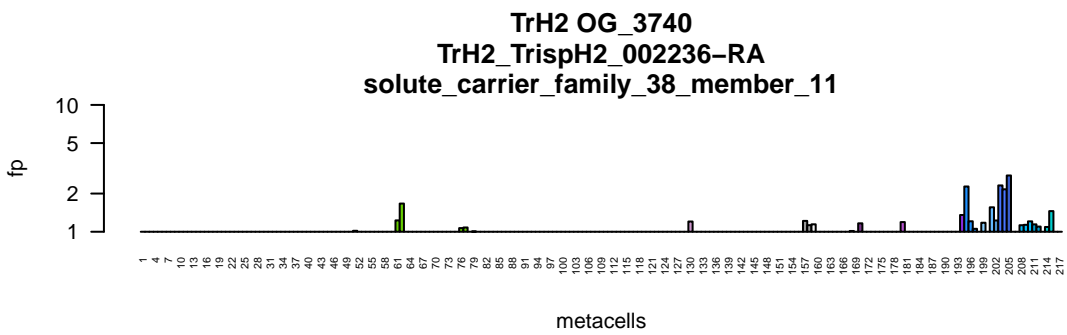
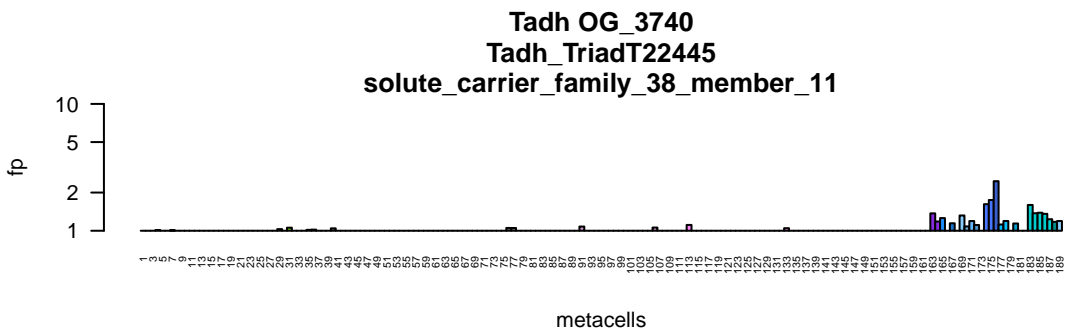


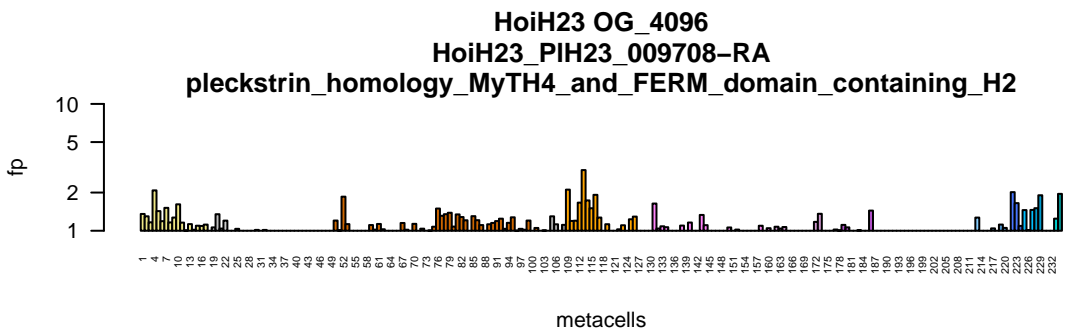
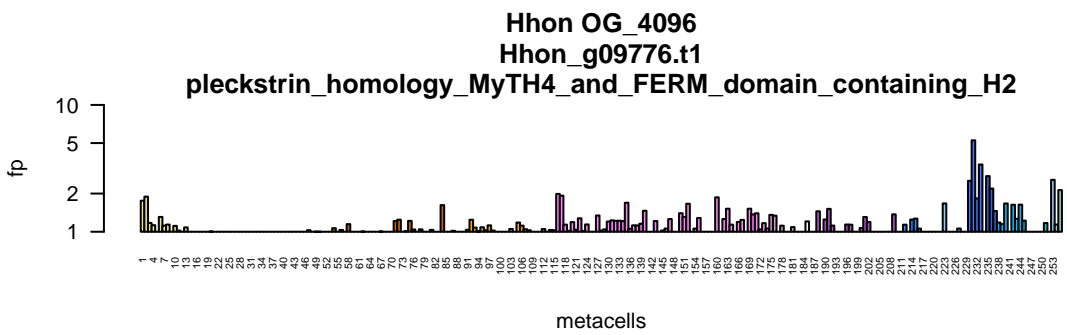
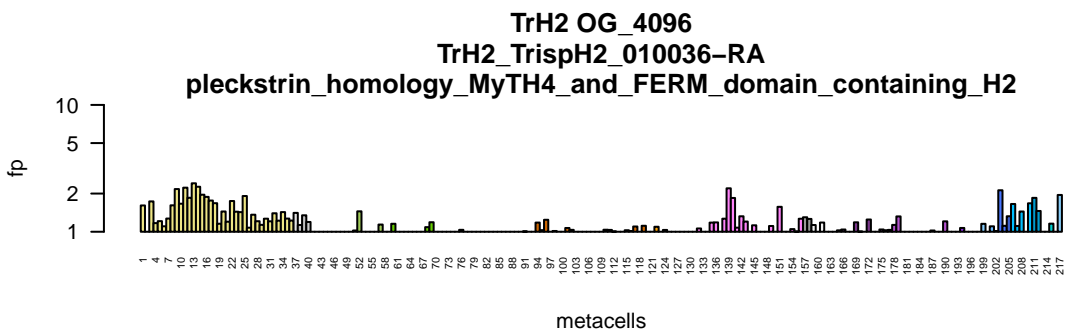
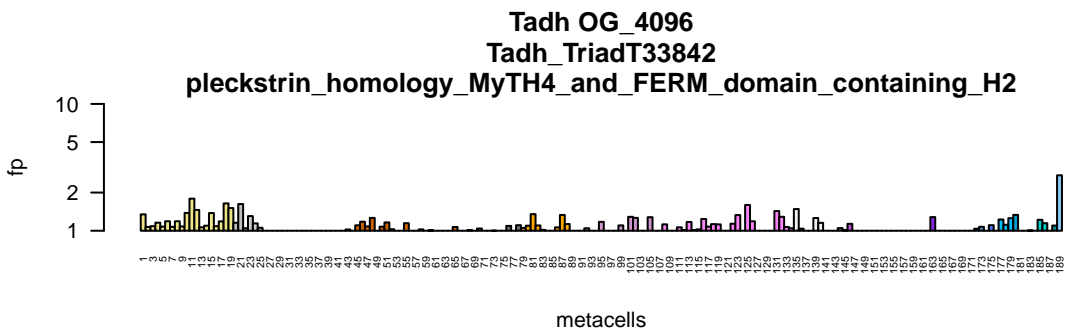












metacells

metacell	fp
1	0
4	0
7	0
13	0
16	0
19	0
25	0
28	0
31	0
34	0
37	0
40	0
43	0
46	0
49	0
52	1
55	1
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	1
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	1
130	0
133	0
136	0
139	0
142	0
145	1
148	1
151	0
154	1
157	0
160	1
163	0
166	0
169	0
172	0
175	1
178	0
181	0
184	0
187	0
190	0
193	0
196	0
199	2
202	2
205	2
208	1
211	1
214	0
217	2

metacell	fp
1	0
4	0
7	0
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	0
34	0
37	0
40	0
43	0
46	0
49	0
52	0
55	0
58	0
61	0
64	0
67	0
70	0
73	0
76	0
79	0
82	0
85	0
88	0
91	0
94	0
97	0
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	0
130	0
133	0
136	0
139	0
142	0
145	0
148	0
151	0
154	0
157	0
160	0
163	0
166	0
169	0
172	0
175	0
178	0
181	0
184	0
187	0
190	0
193	0
196	0
199	1
200	2
201	1
202	0
205	0
208	0
211	0
214	0
217	0

metacell	fp
1	0
4	0
7	0
13	0
16	1
19	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	1
76	0
79	0
82	0
85	0
88	0
91	0
94	0
97	0
100	1
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
157	0
160	0
163	0
166	0
169	0
172	0
175	0
178	0
181	0
184	0
187	0
190	0
193	1
196	1
199	2
202	3
205	1
206	1
209	1
211	1
214	1
217	1

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

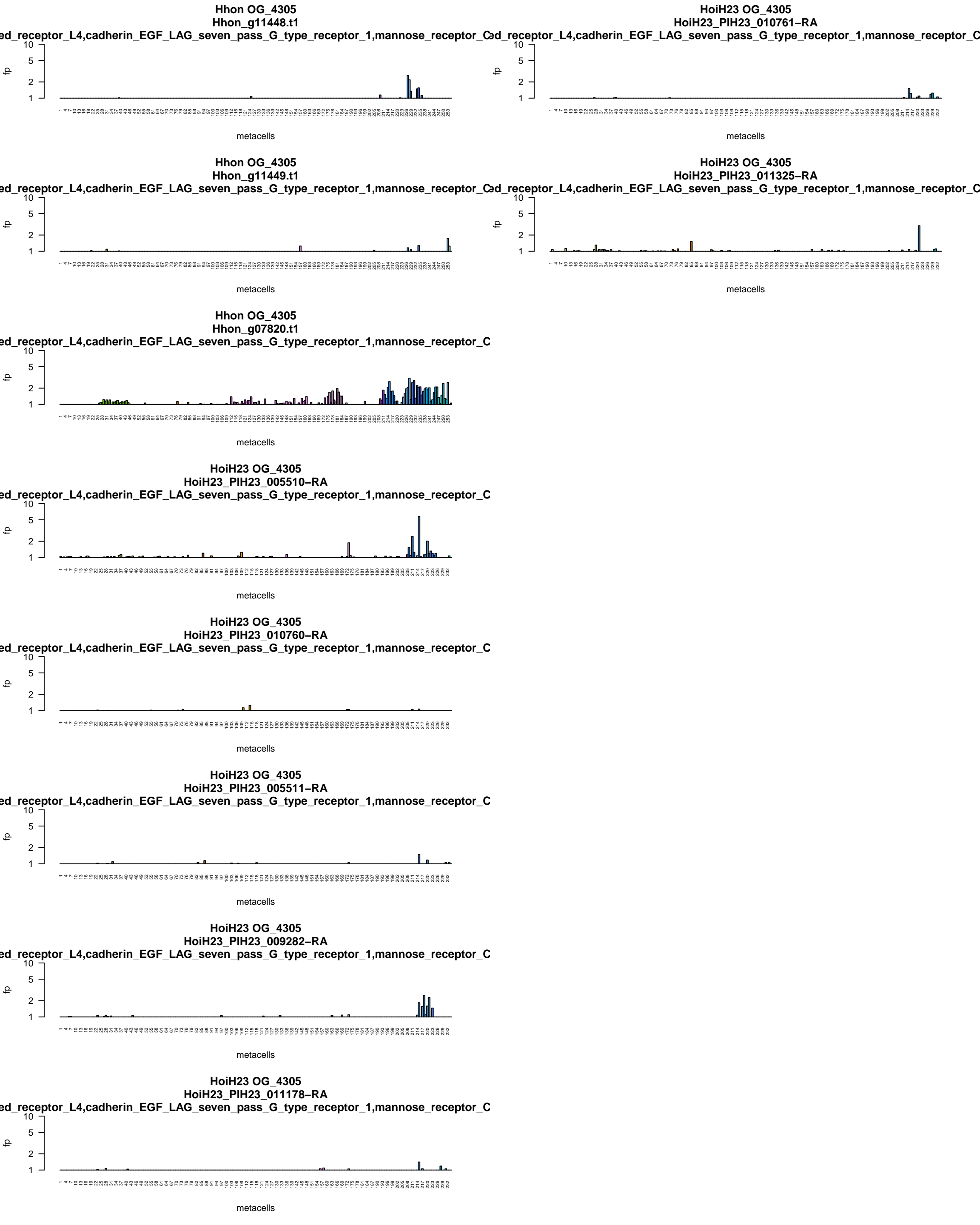
metacell	fp
1	0
4	0
7	0
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	0
34	0
37	0
40	0
43	0
46	0
49	0
52	1
55	0
58	0
61	0
64	0
67	0
70	0
73	0
76	0
79	0
82	0
85	0
88	0
91	0
94	1
97	0
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	0
130	0
133	0
136	0
139	0
142	0
145	0
148	0
151	0
154	0
157	1
160	0
163	0
166	0
169	0
172	0
175	0
178	0
181	0
184	0
187	0
190	0
193	0
196	0
199	1
202	2
205	0
208	0
211	0
214	0
217	1

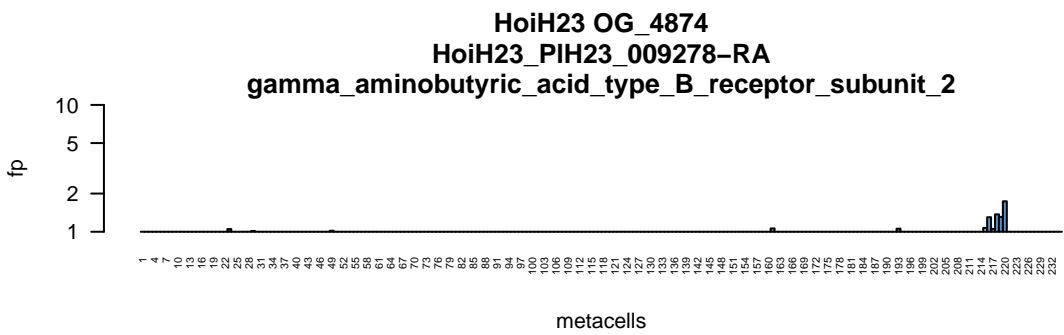
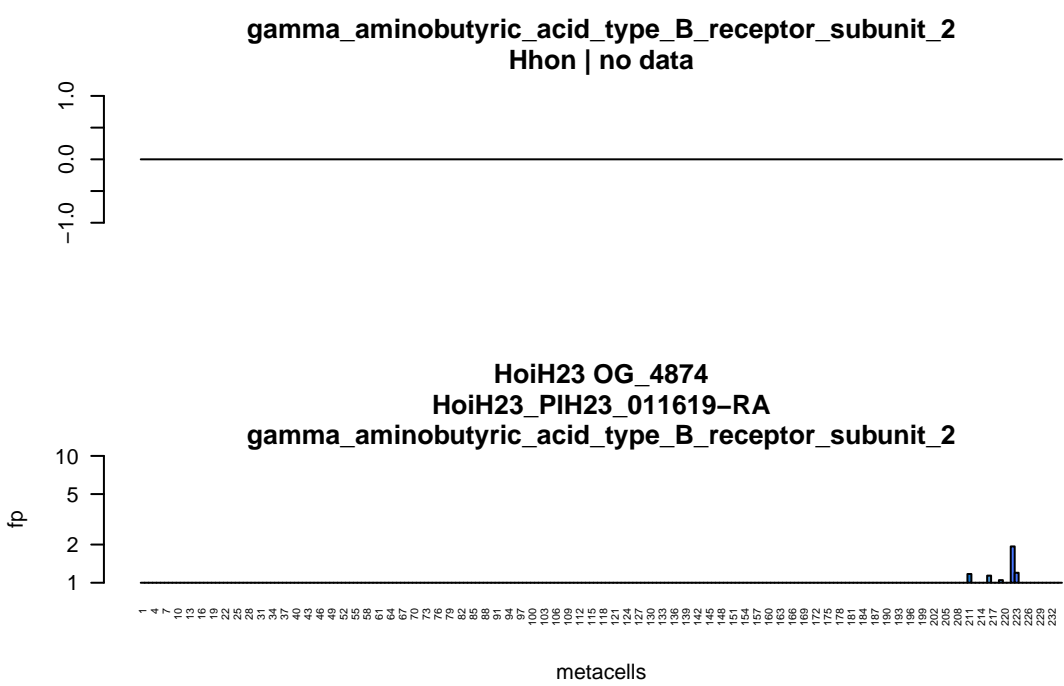
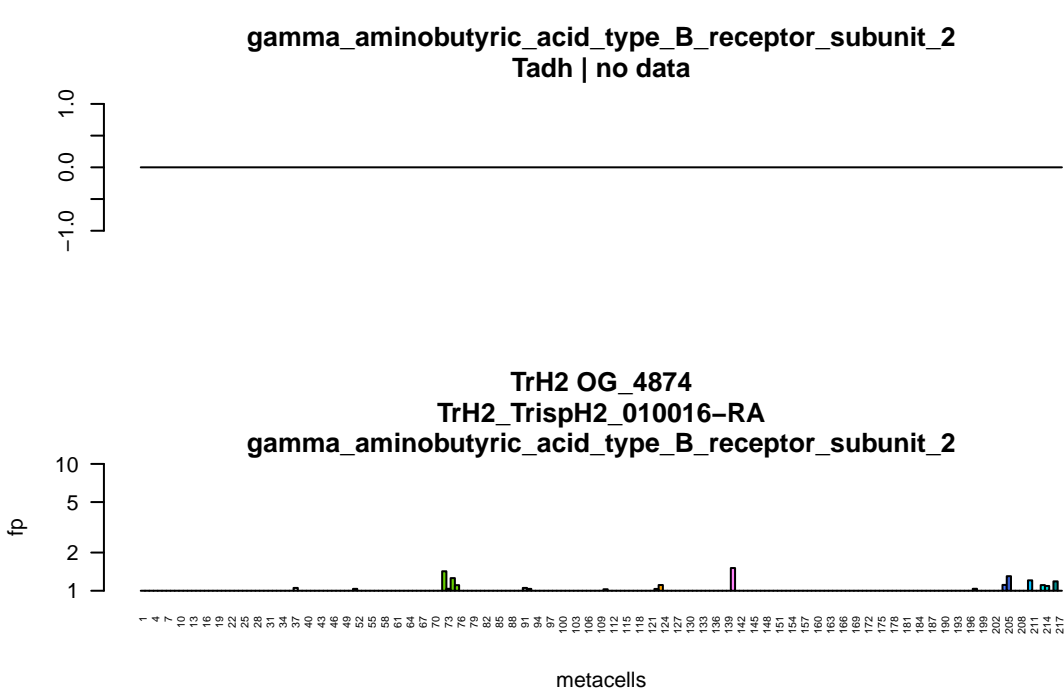
A bar chart showing the frequency of metacells (x-axis) across different frequency bins (y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 253. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows a distribution of metacells across frequency bins, with a peak around 250 metacells.

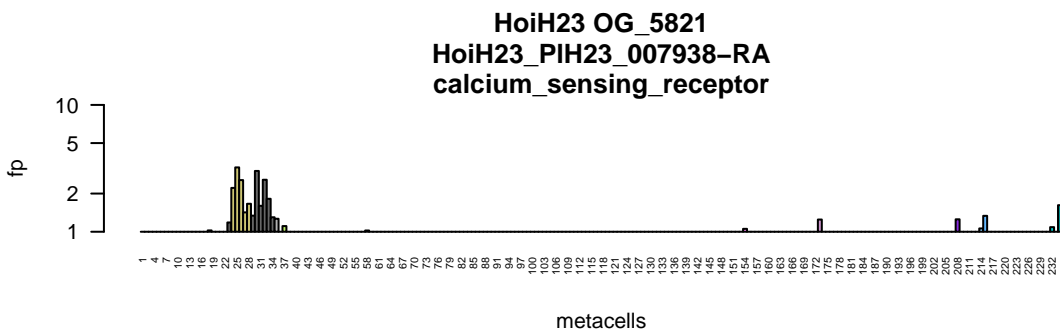
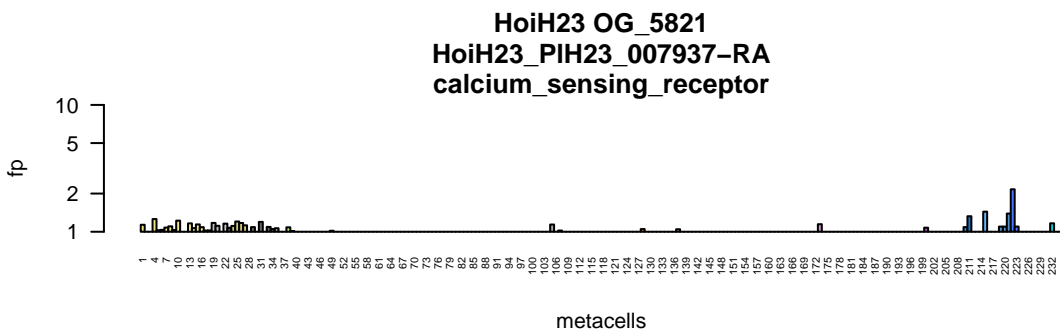
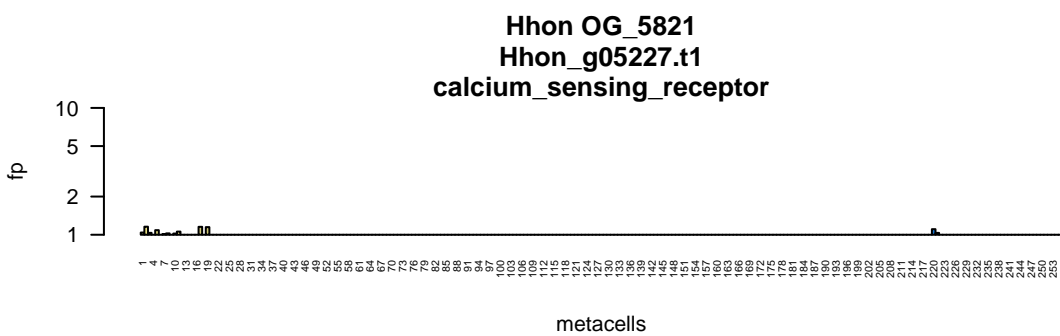
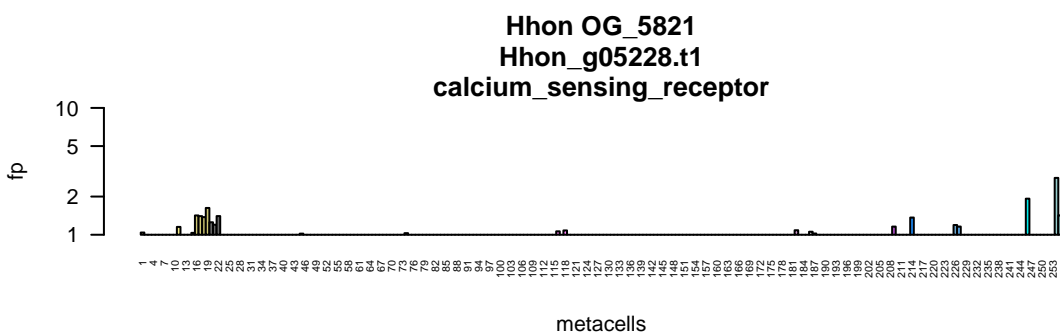
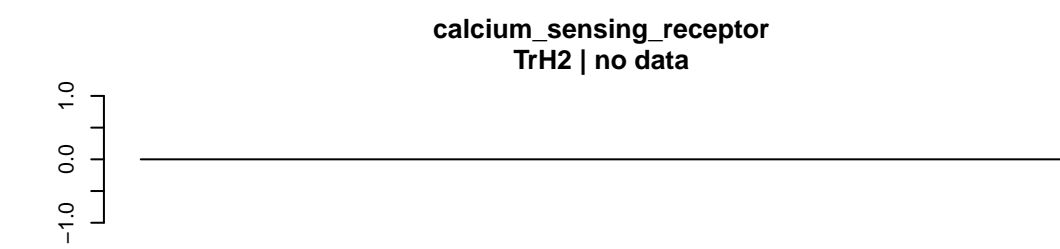
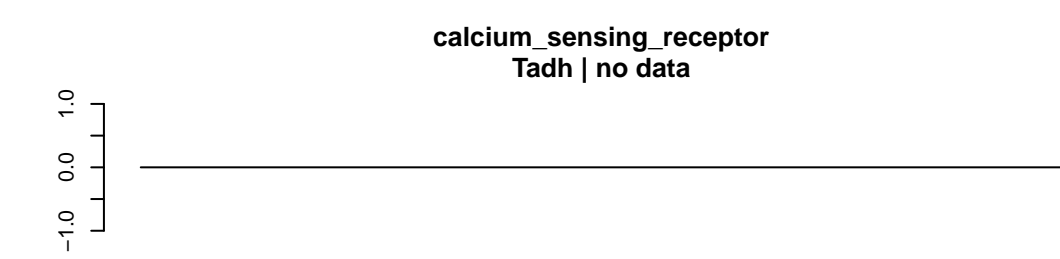
A bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 253. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a few outliers reaching up to 3. The bars are colored in a gradient from dark blue to light blue.

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists indices from 1 to 253. Most metacells have 0 false positives, but some have 1 or 2. Metacells 229 and 253 have the highest number of false positives, with 2 each.

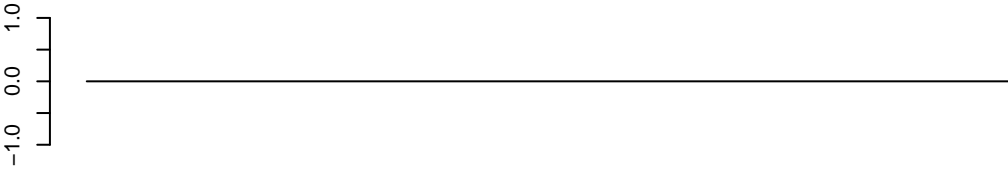
metacell	fp
1	0
4	0
10	0
13	0
19	0
22	0
28	0
31	0
37	0
40	0
46	0
49	0
55	0
61	0
64	0
70	0
73	0
76	0
82	0
85	0
88	0
91	0
97	0
100	0
106	0
109	0
115	0
118	0
127	0
133	0
136	0
142	0
145	0
151	0
154	0
160	0
163	0
169	0
172	0
178	0
184	0
187	0
191	0
193	0
195	0
200	0
202	0
205	0
214	0
219	0
220	0
223	0
229	2
232	0
238	0
241	0
247	0
250	0
253	2







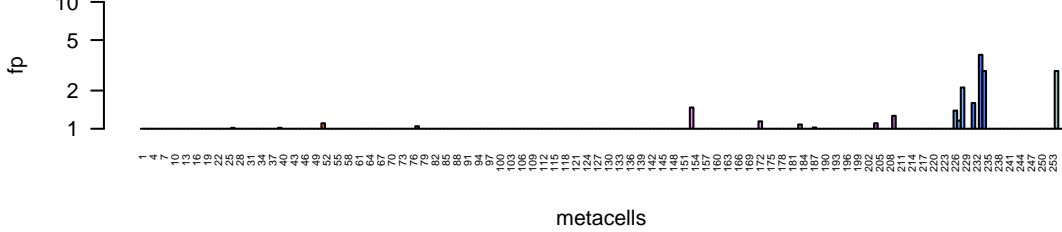
gamma_aminobutyric_acid_type_B_receptor_subunit_2
Tadh | no data



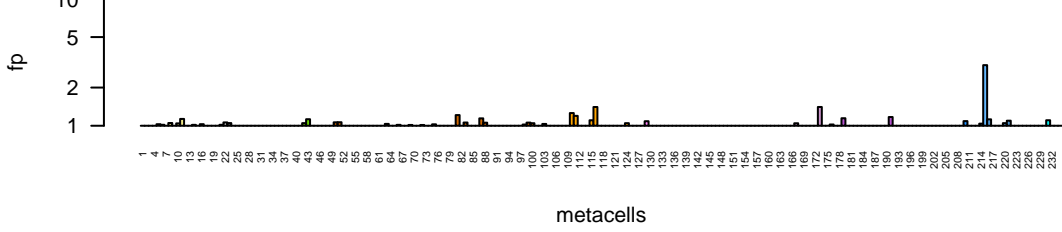
gamma_aminobutyric_acid_type_B_receptor_subunit_2
TrH2 | no data

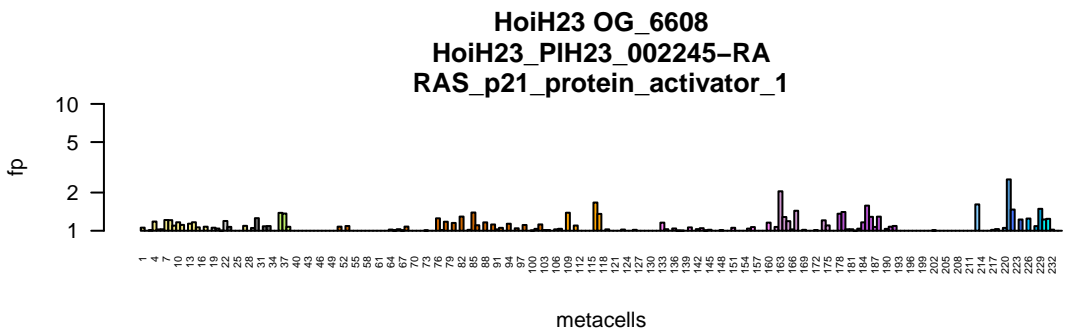
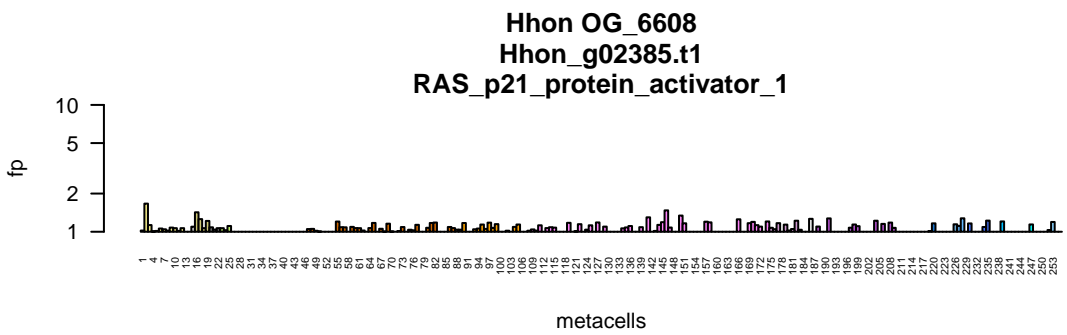
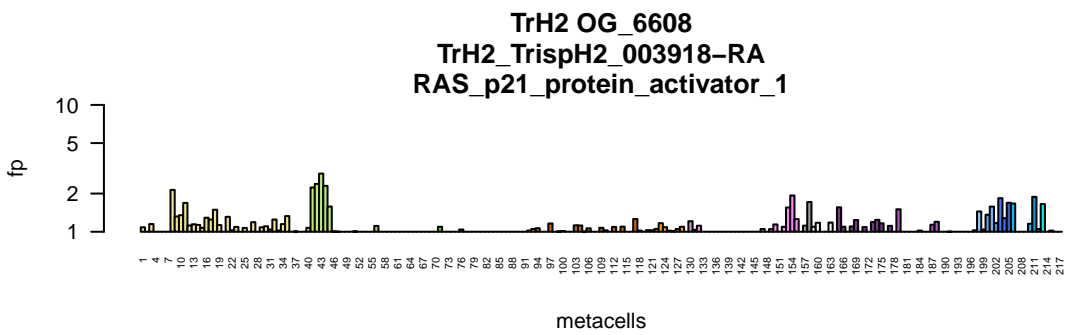
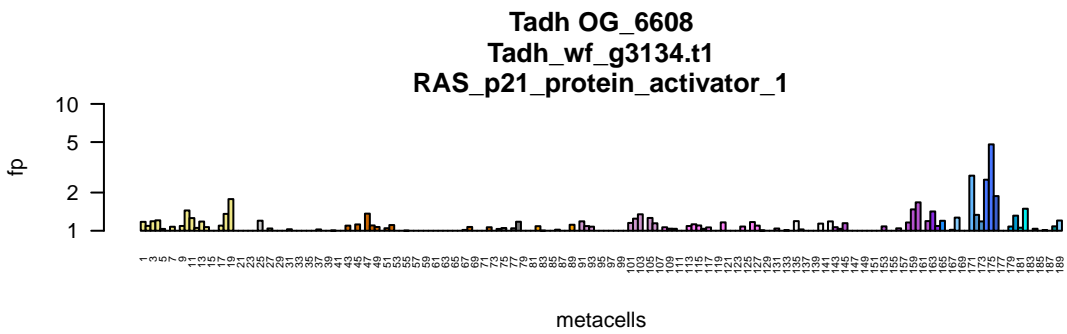


Hhon OG_6010
Hhon_g01250.t1
gamma_aminobutyric_acid_type_B_receptor_subunit_2



HoiH23 OG_6010
HoiH23_PIH23_000005-RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2

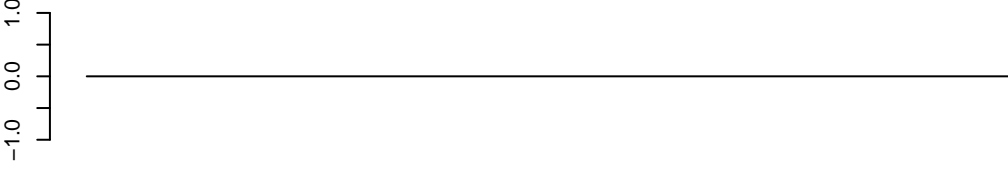




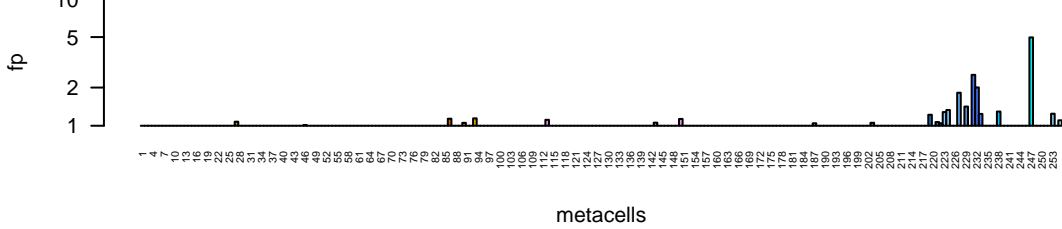
gamma_aminobutyric_acid_type_B_receptor_subunit_1
Tadh | no data



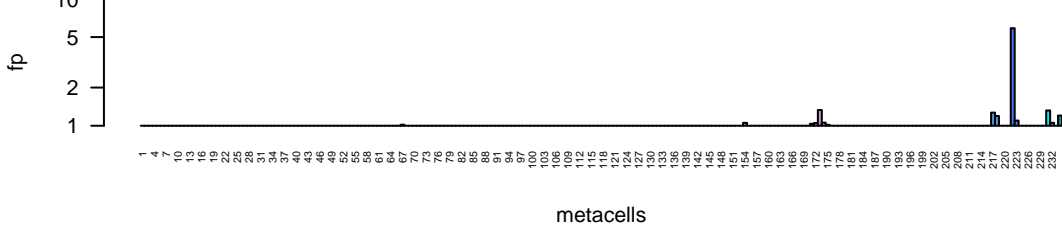
gamma_aminobutyric_acid_type_B_receptor_subunit_1
TrH2 | no data

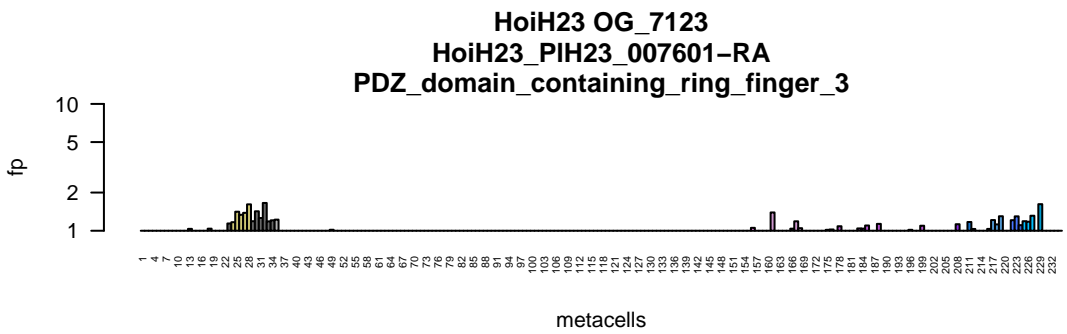
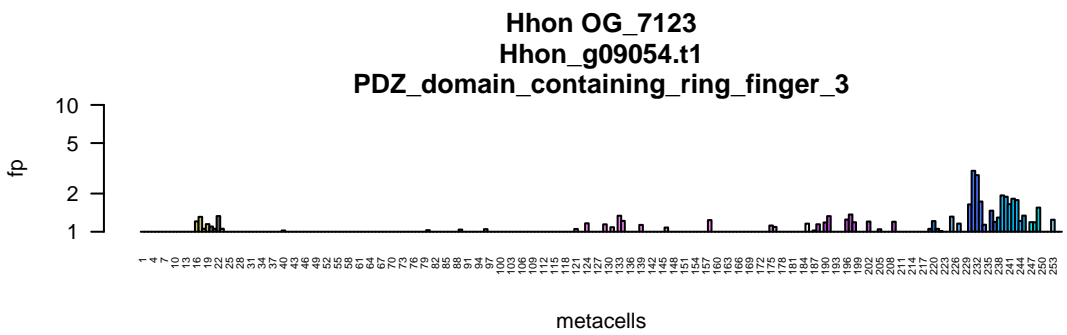
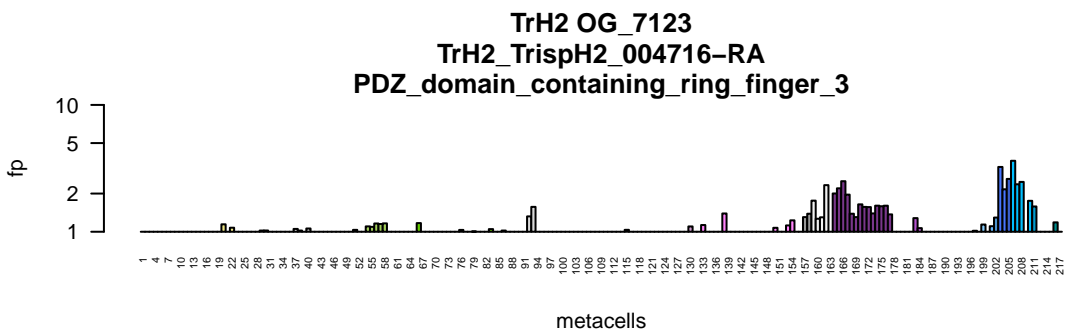
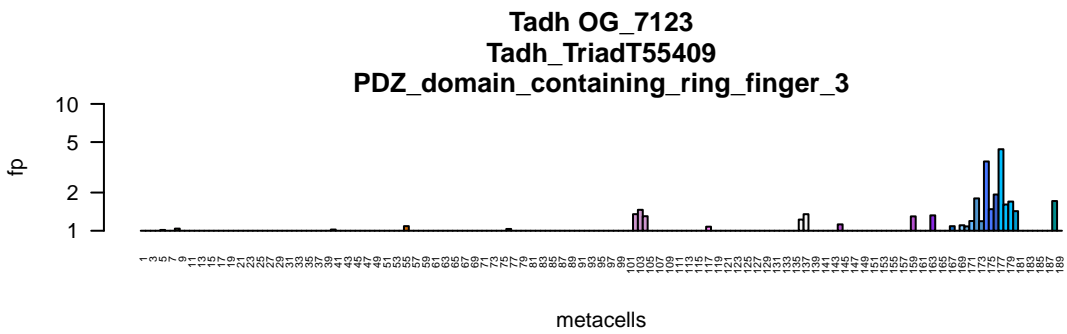


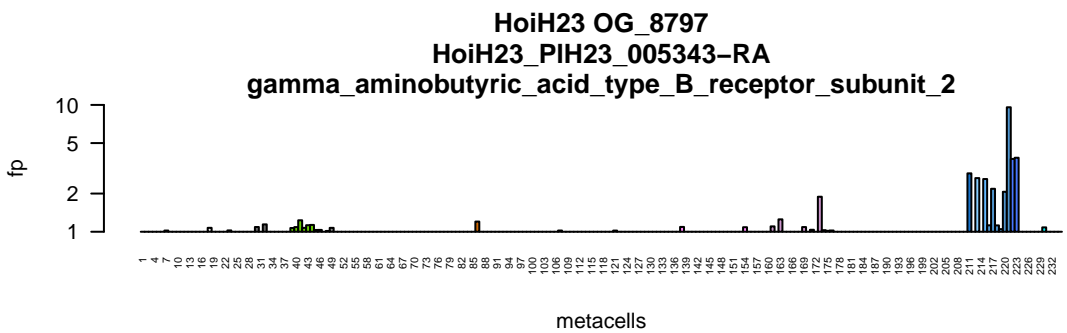
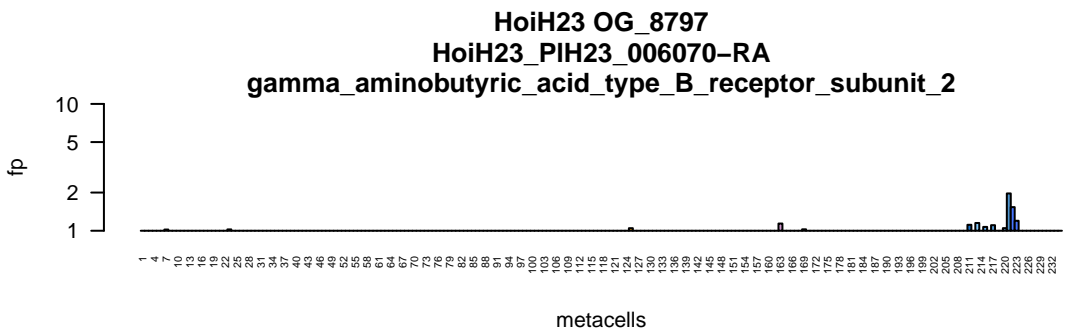
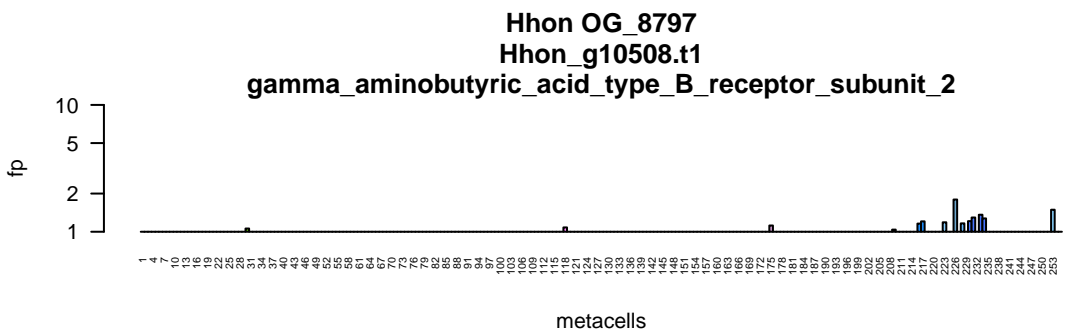
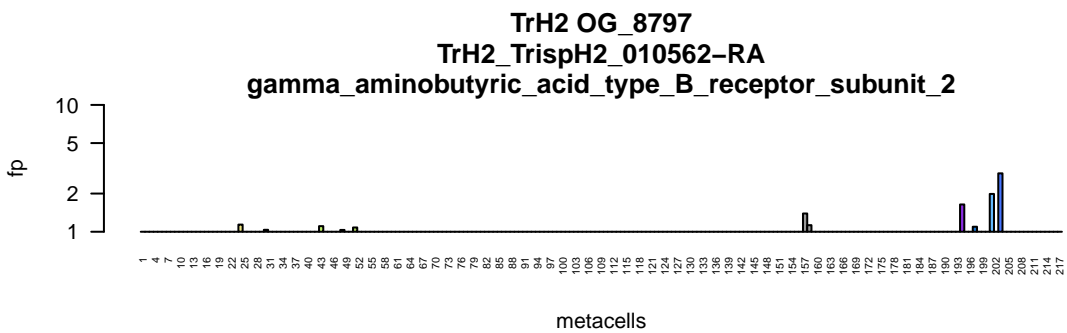
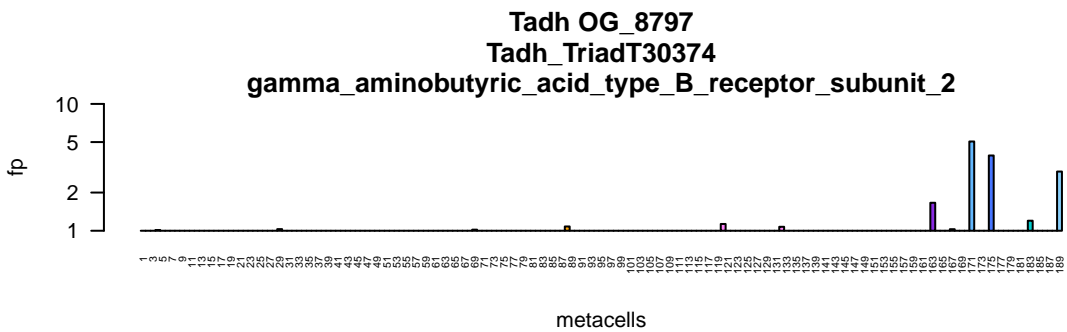
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gamma_aminobutyric_acid_type_B_receptor_subunit_1



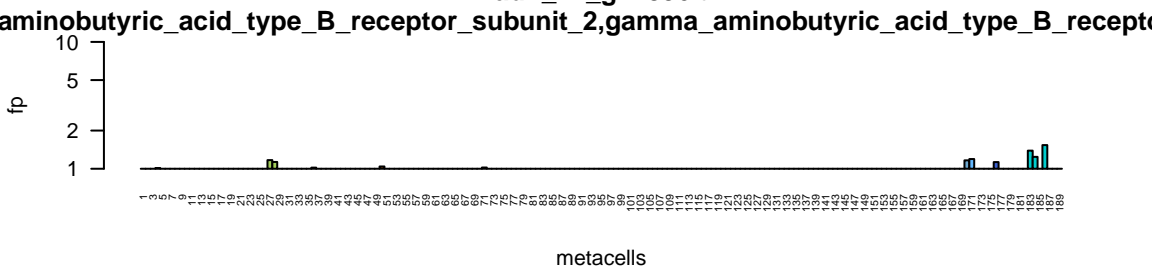
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HoiH23_PIH23_008905-RA
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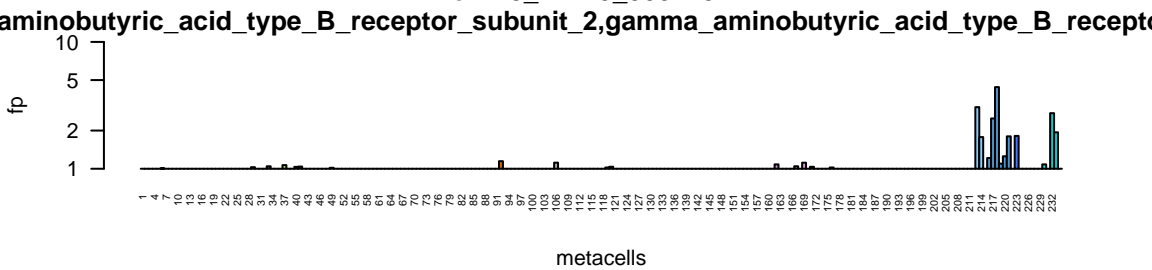




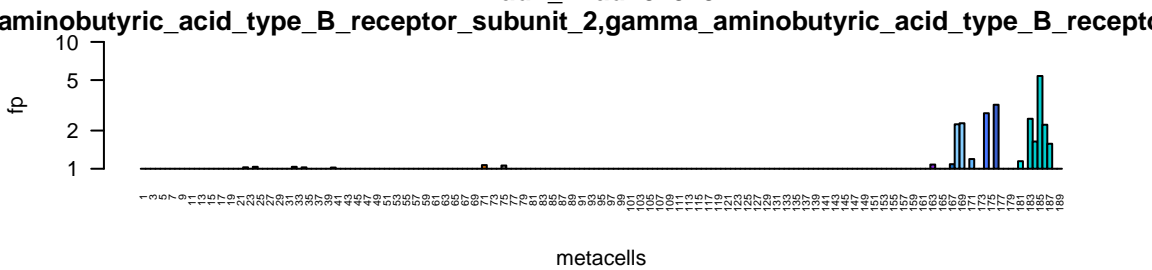
Tadh OG_8960
tadh_wf_g11896.t1



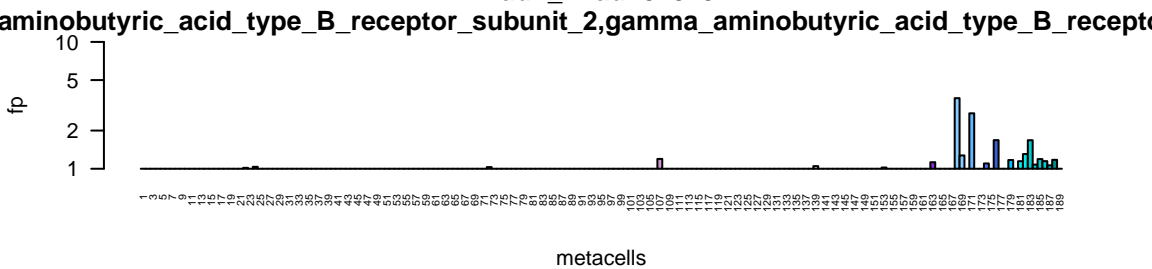
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HoiH23_PIH23_000719-RA



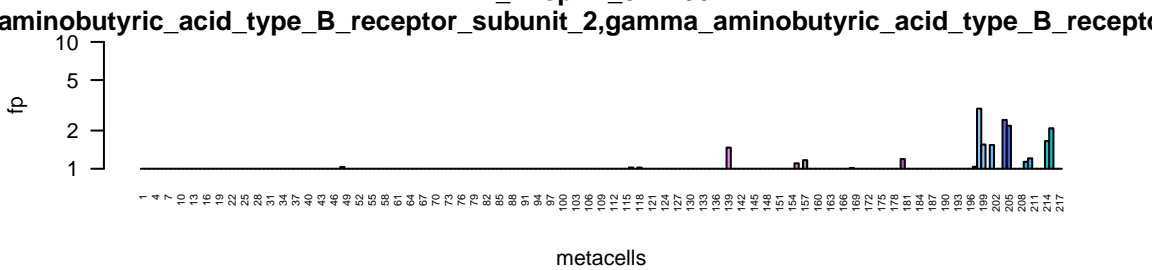
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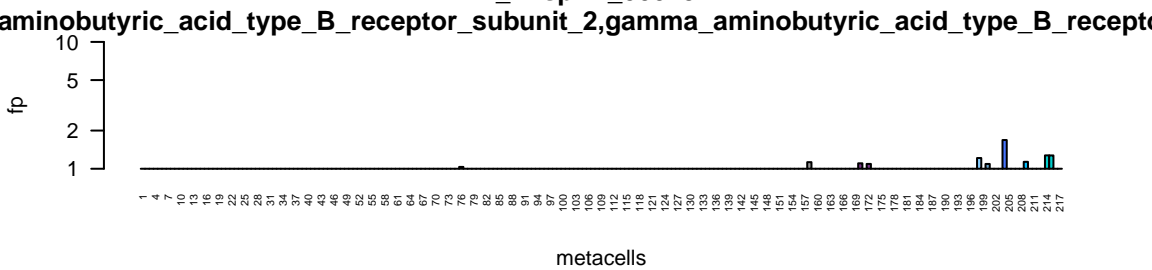
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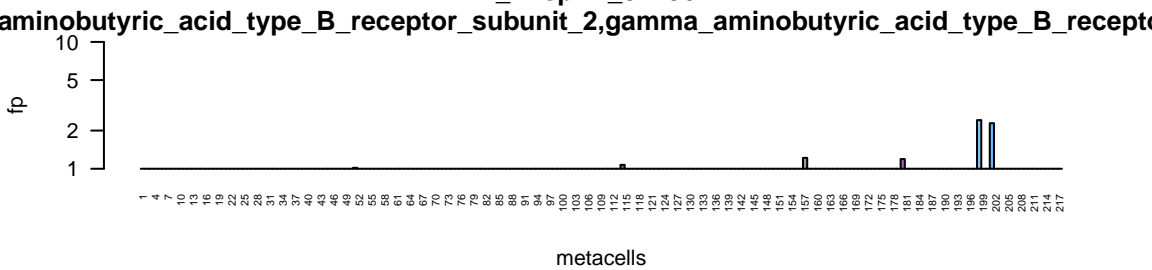
TrH2 OG_8960
TrH2_TrispH2_012100-RA



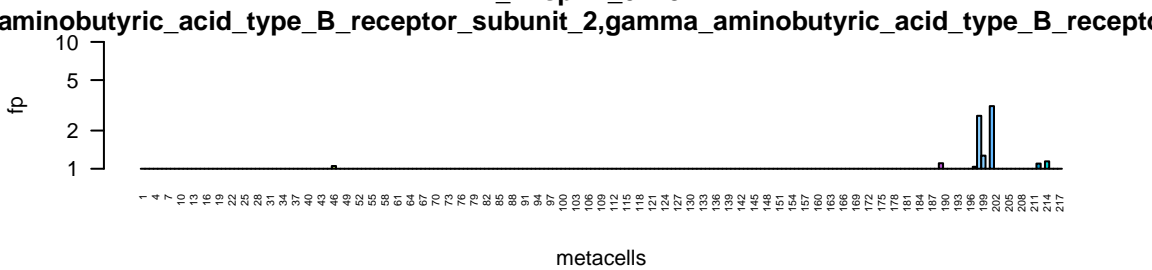
TrH2 OG_8960
TrH2_TrispH2_000234-RA



TrH2 OG_8960
TrH2_TrispH2_011964-RA



TrH2 OG_8960
TrH2_TrispH2_011847-RA



Hhon OG_8960
Hhon_g05296.t1

