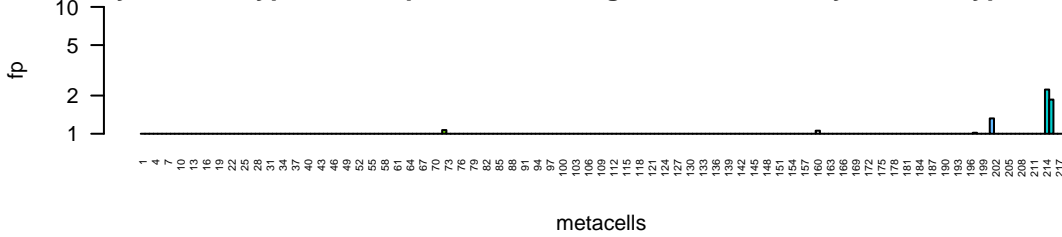


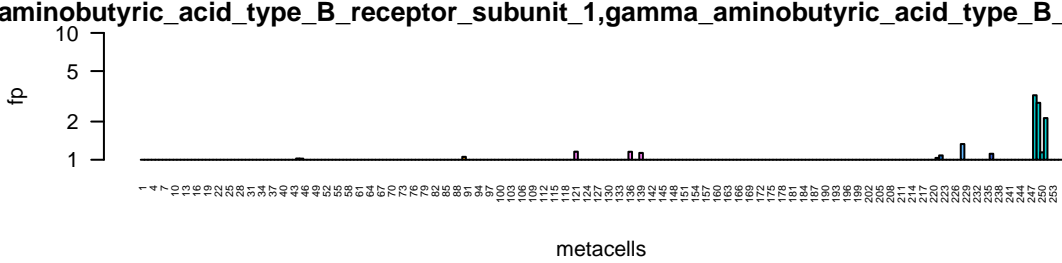
aminobutyric_acid_type_B_receptor_subunit_1,gamma_aminobutyric_acid_type_B_recept
Tadh | no data



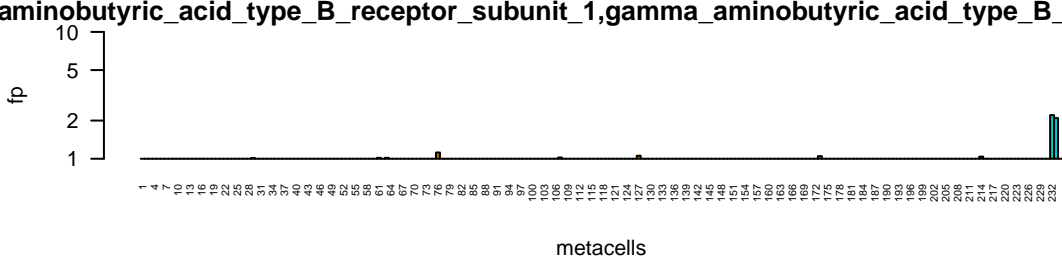
TrH2 OG_8921
TrH2_TrispH2_006332-RA
aminobutyric_acid_type_B_receptor_subunit_1,gamma_aminobutyric_acid_type_B_recept

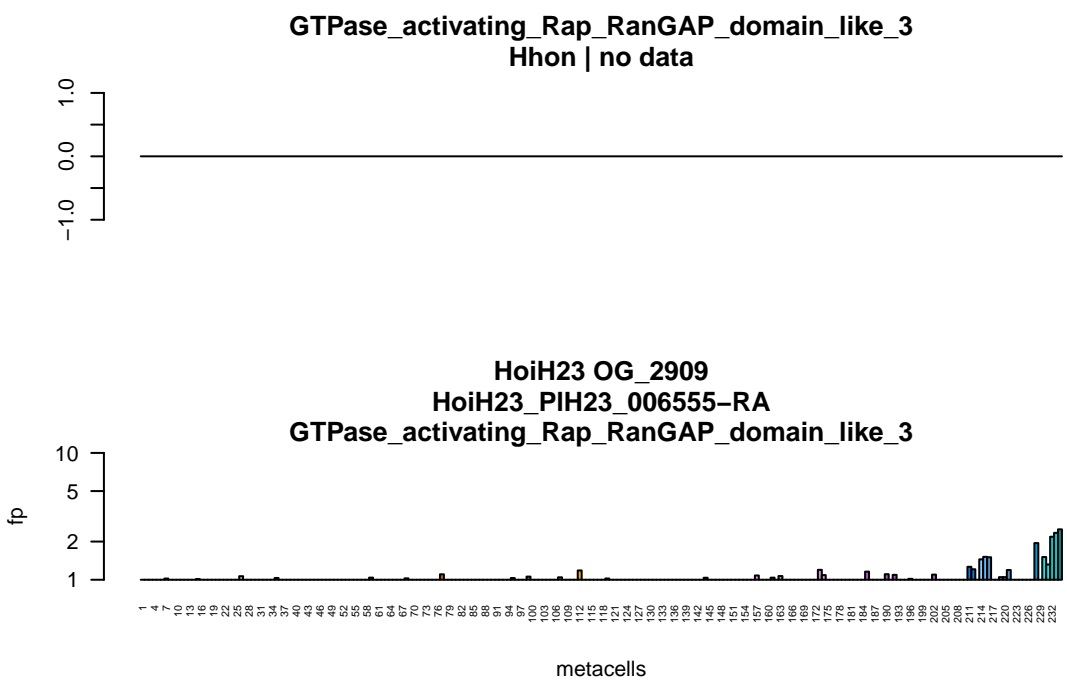
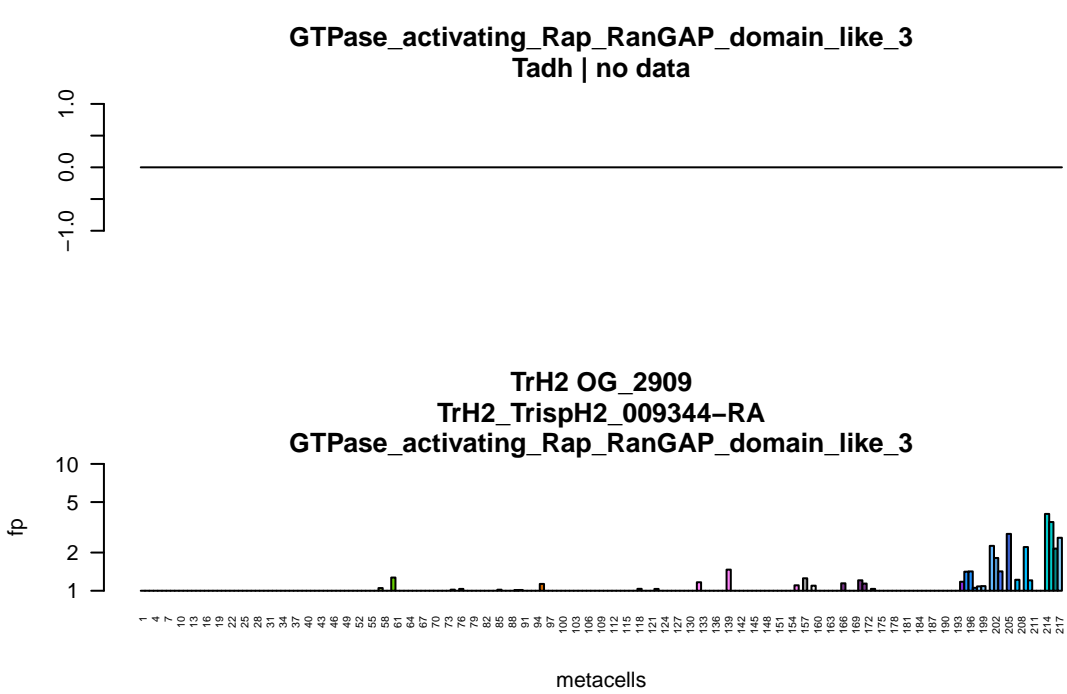


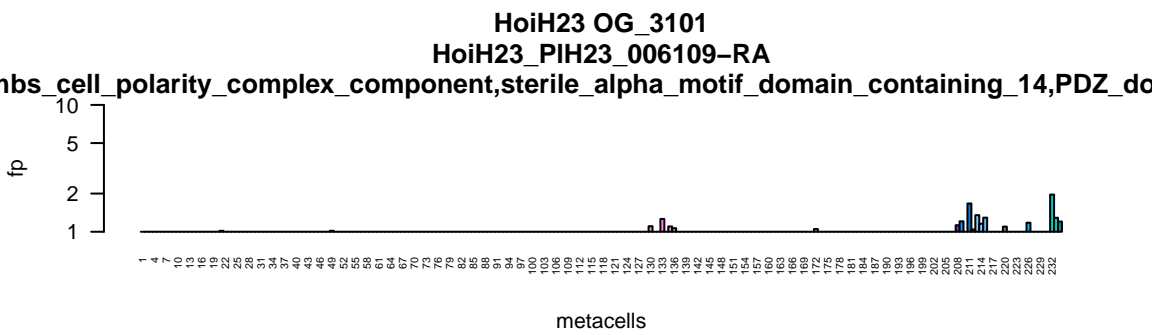
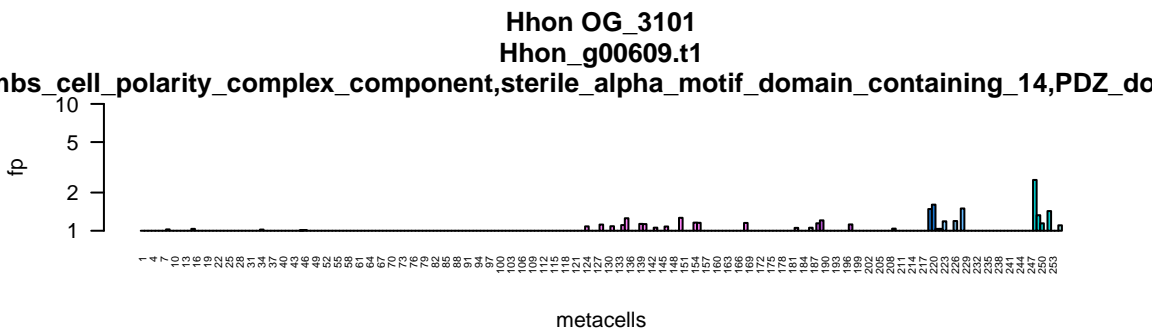
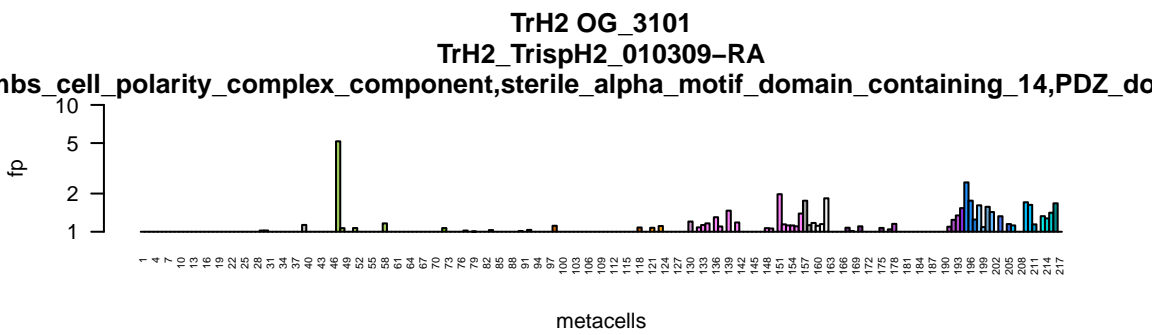
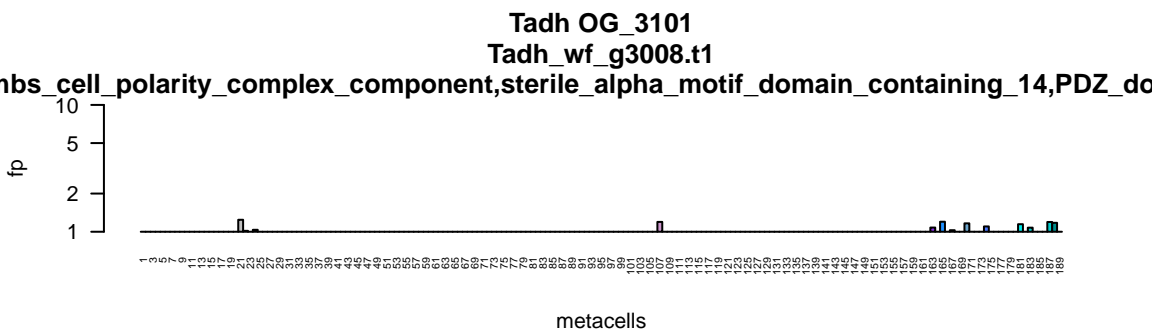
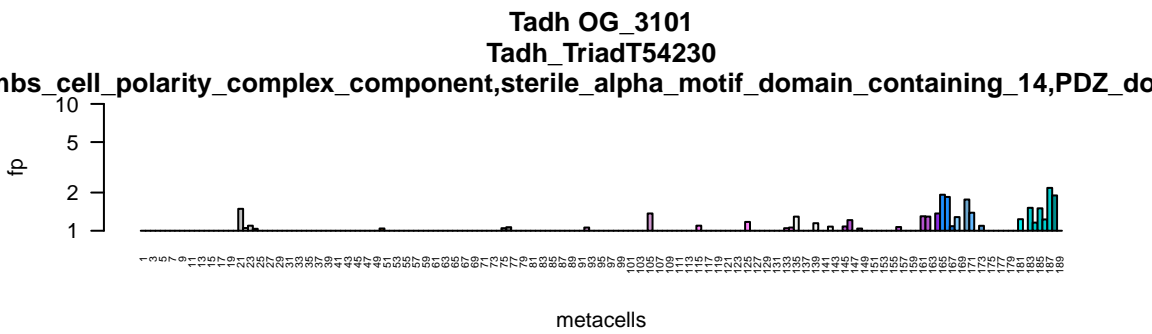
Hhon OG_8921
Hhon_g05728.t1
aminobutyric_acid_type_B_receptor_subunit_1,gamma_aminobutyric_acid_type_B_recept

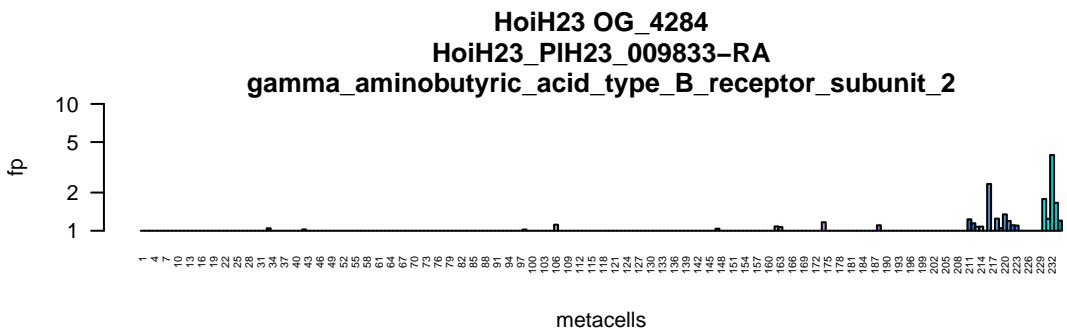
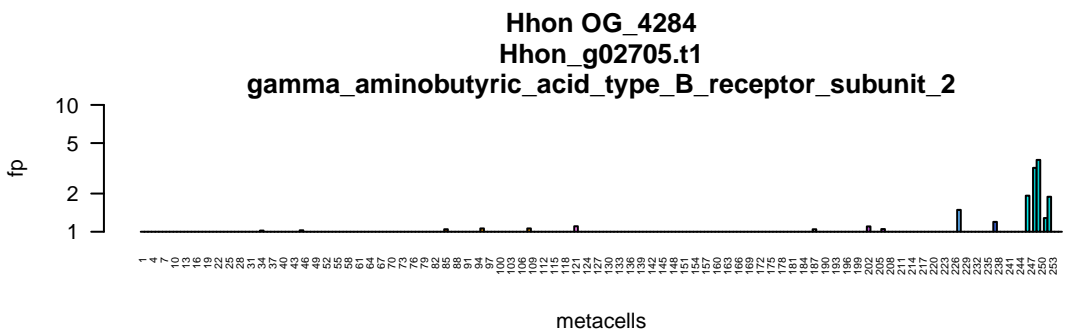
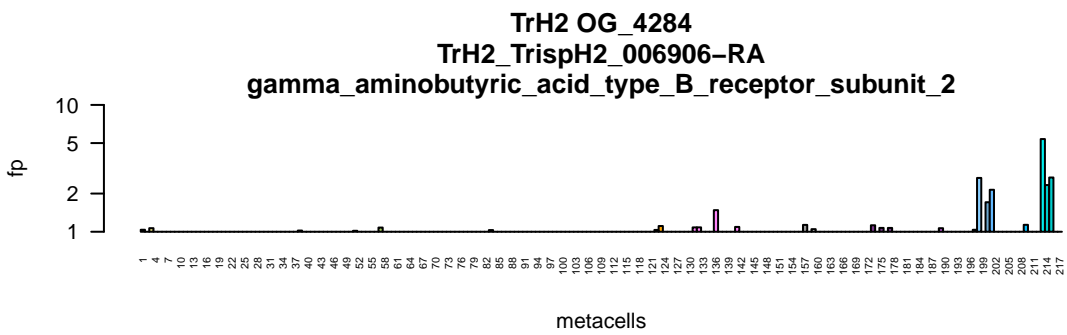
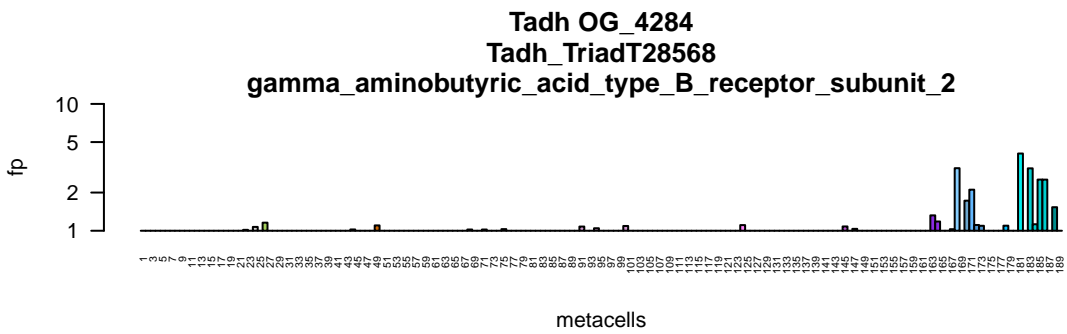


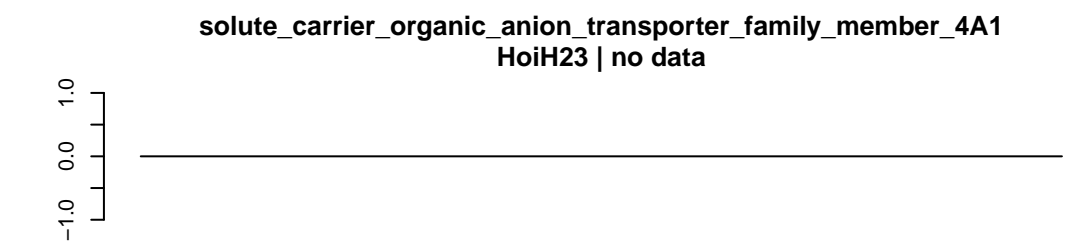
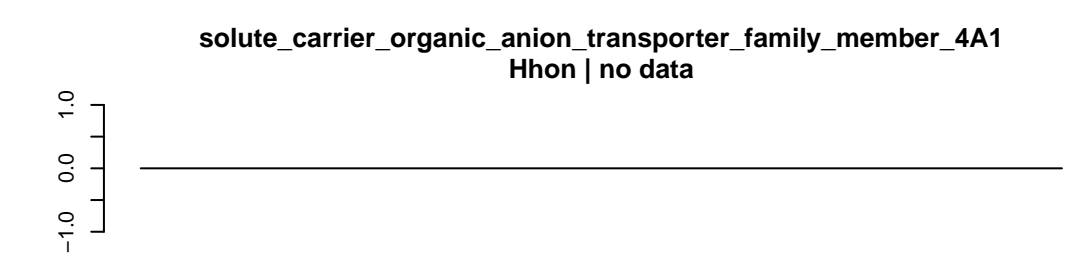
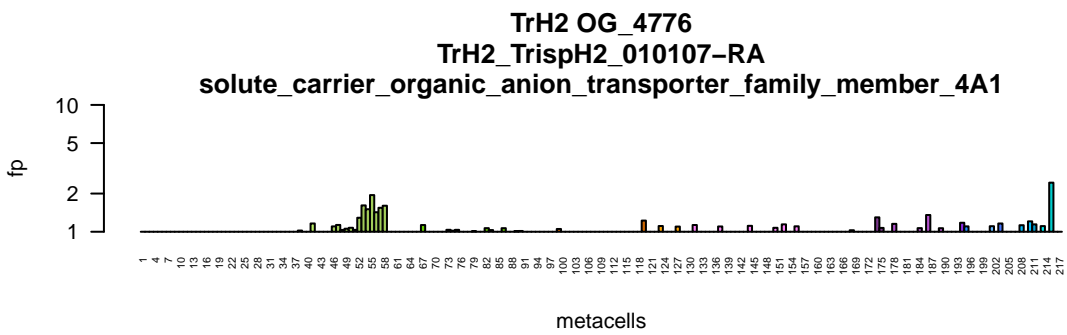
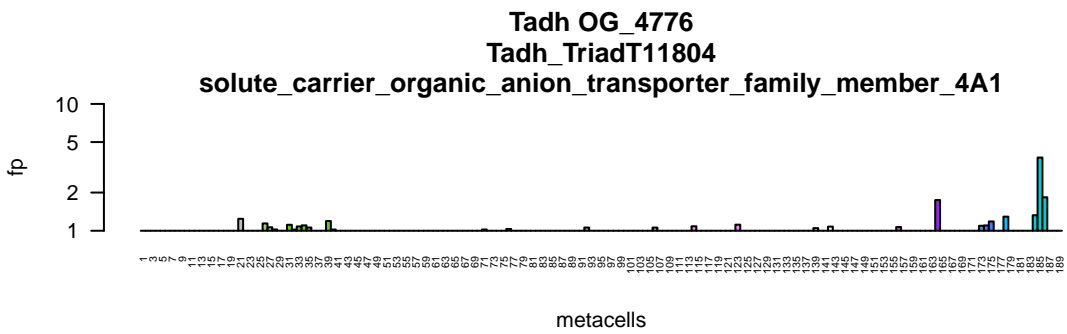
HoiH23 OG_8921
HoiH23_PIH23_002276-RA
aminobutyric_acid_type_B_receptor_subunit_1,gamma_aminobutyric_acid_type_B_recept

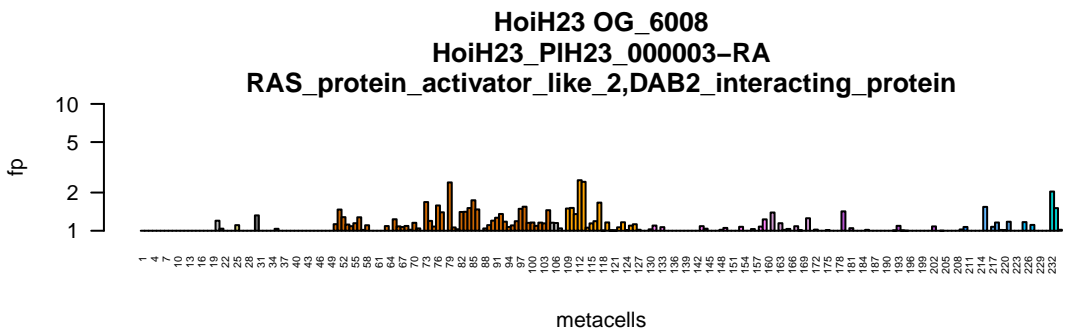
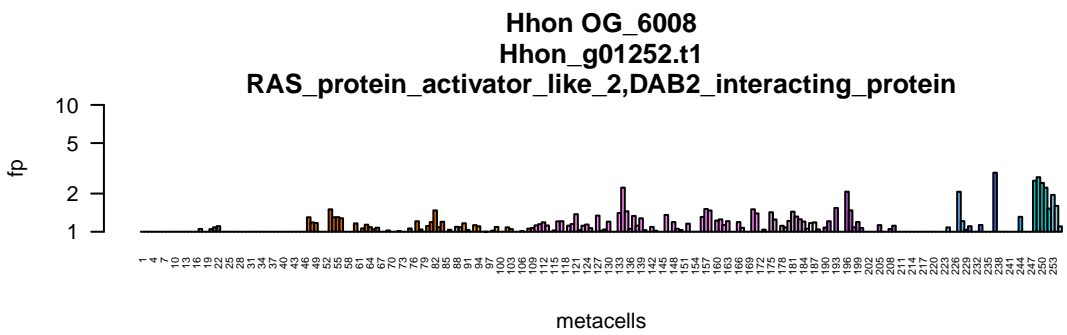
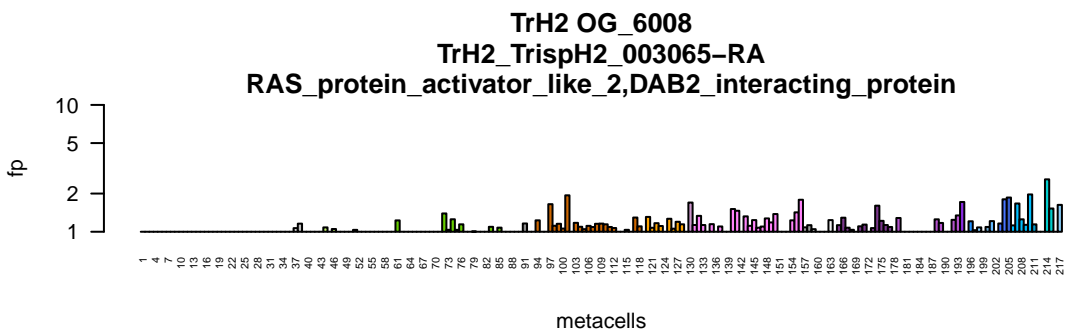
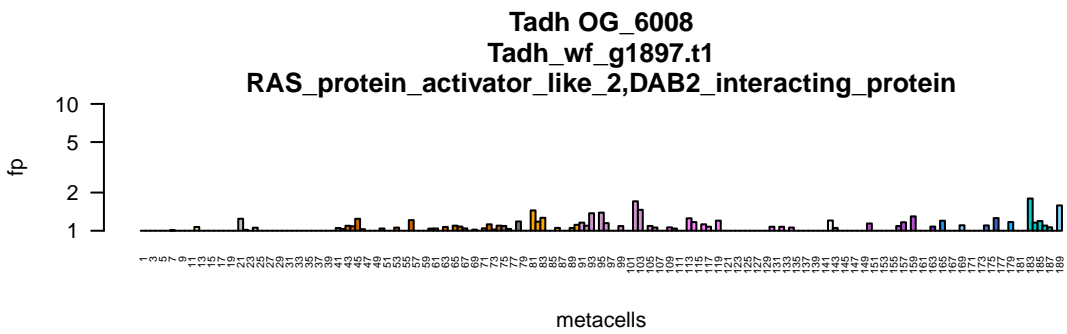


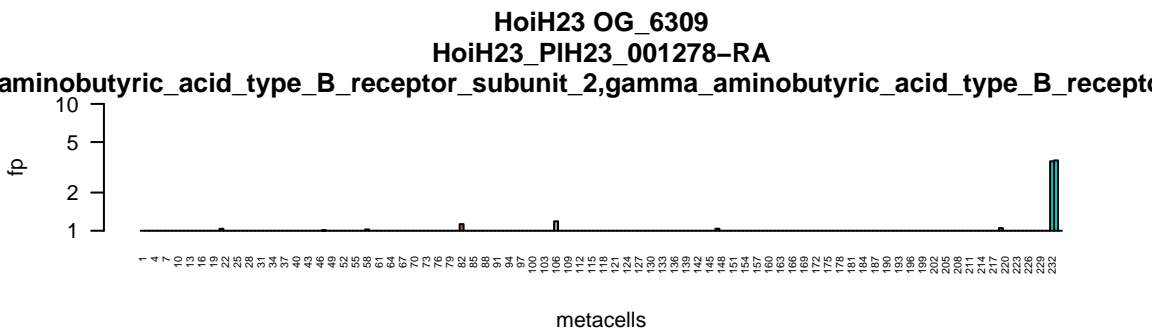
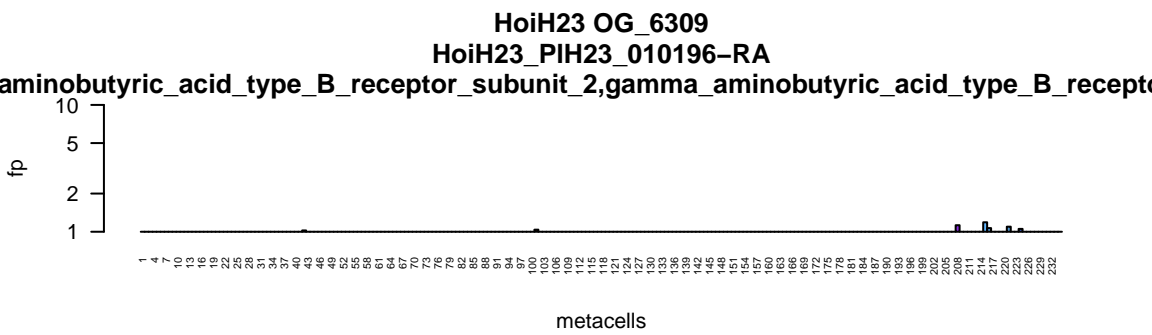
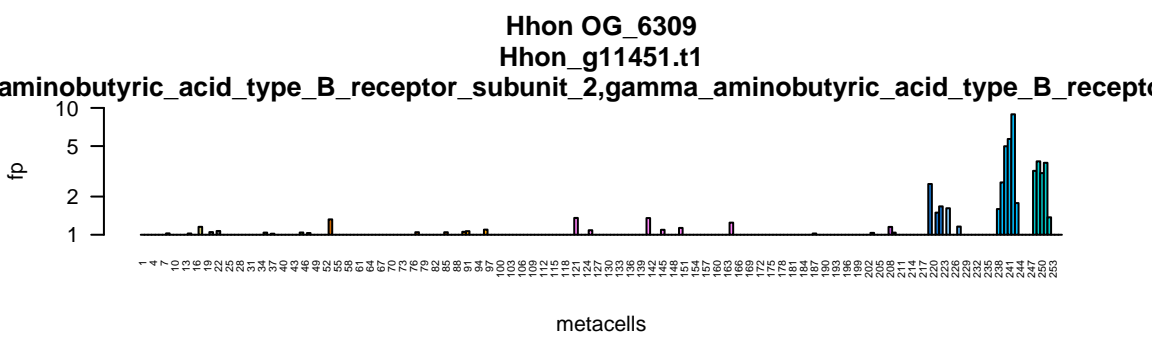
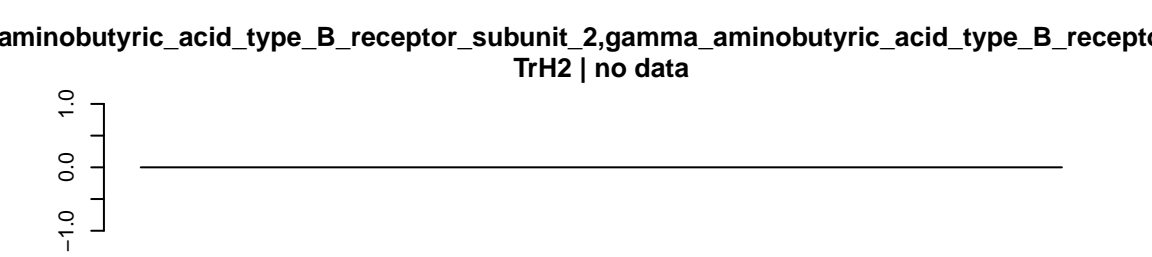
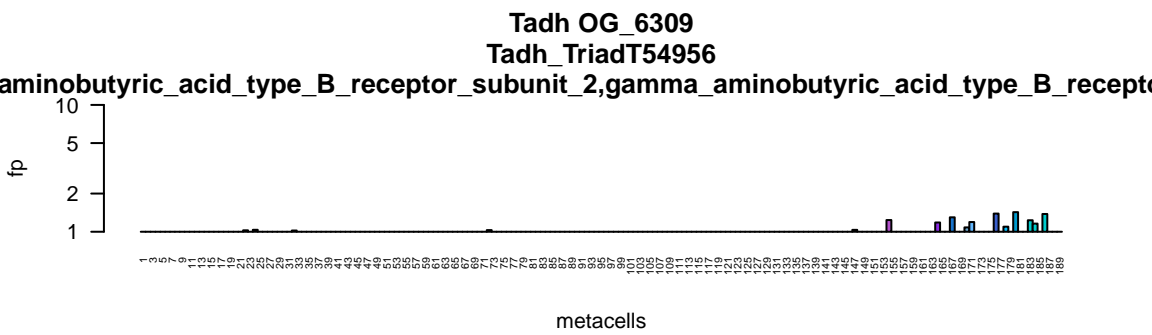
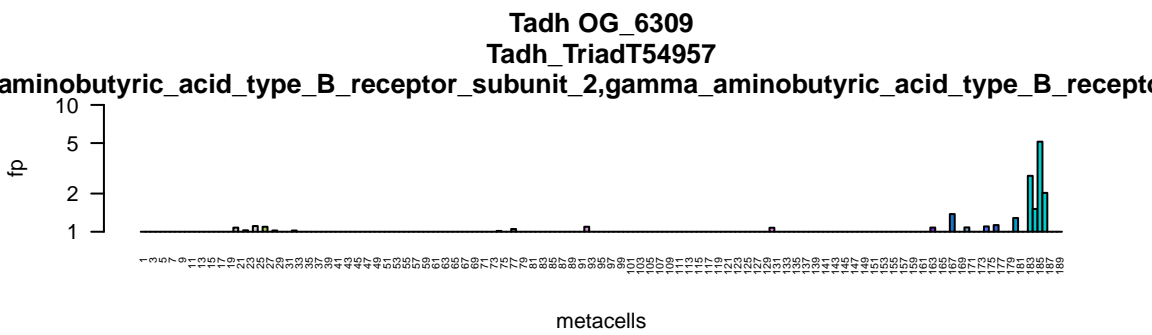
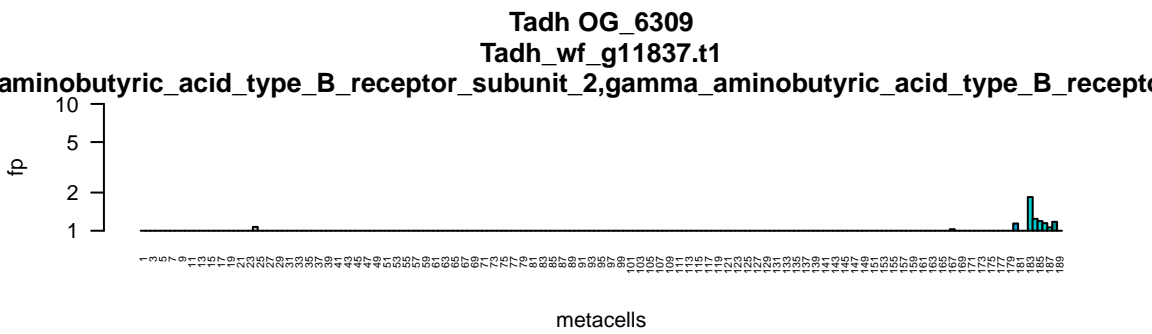


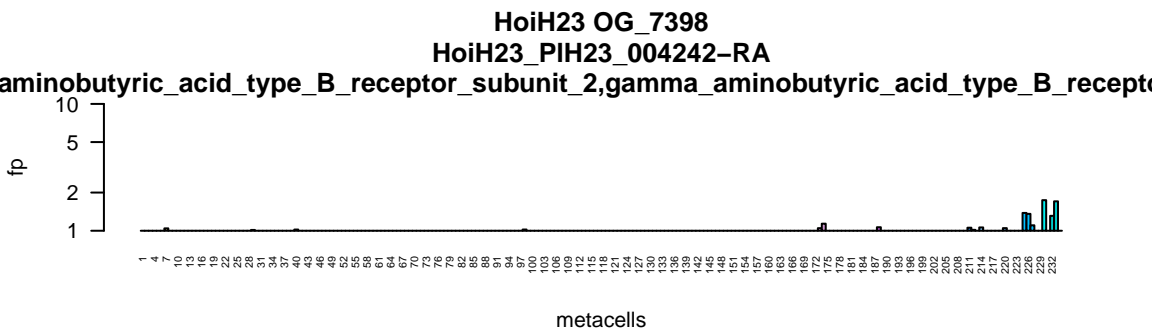
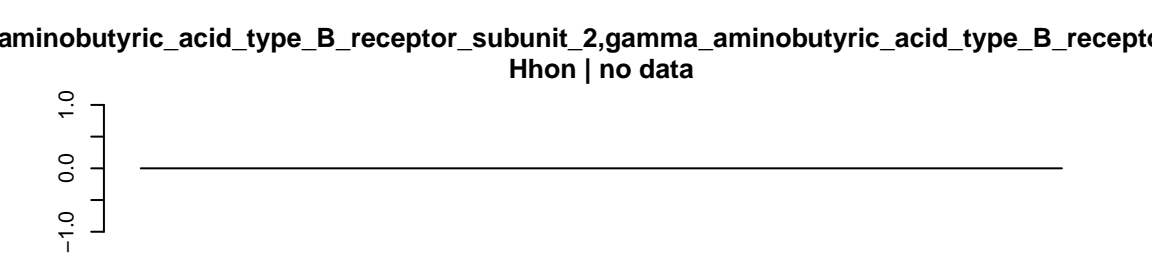
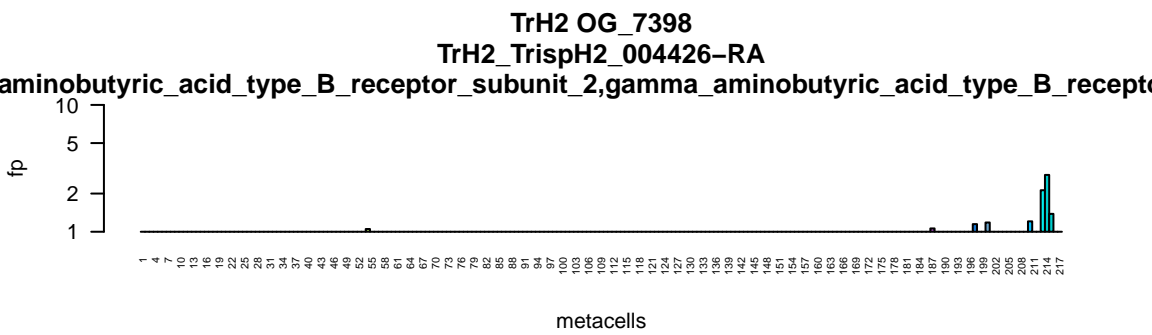
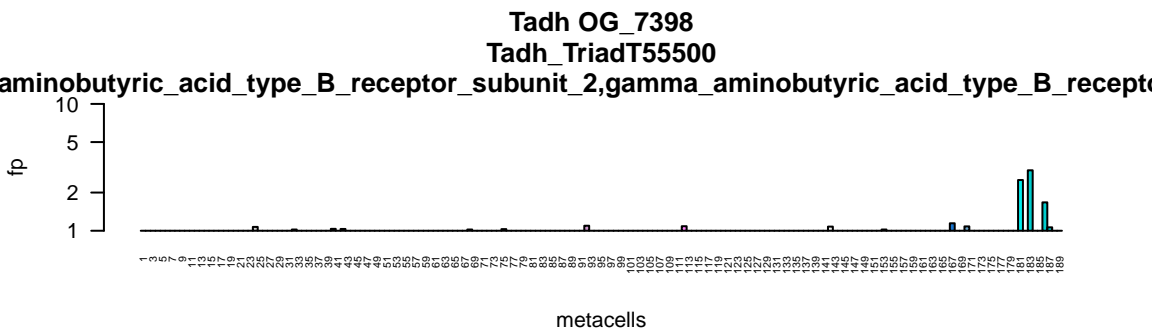


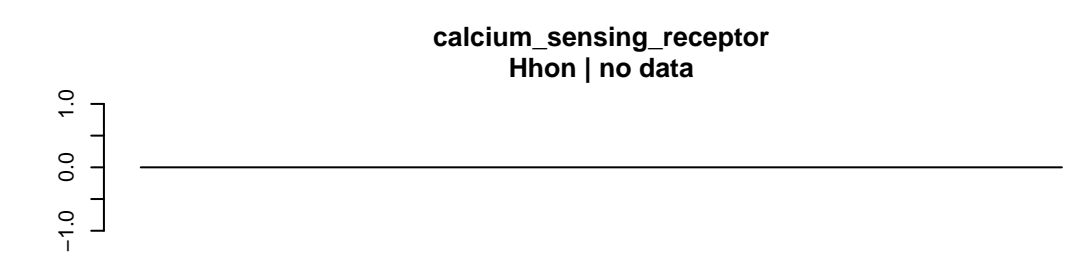
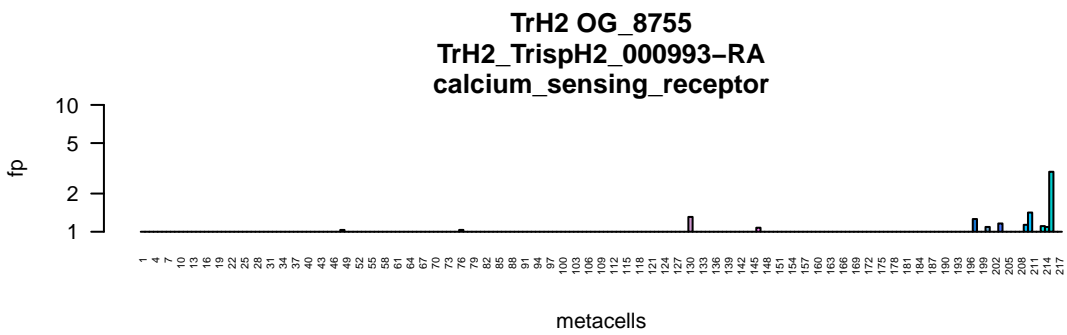
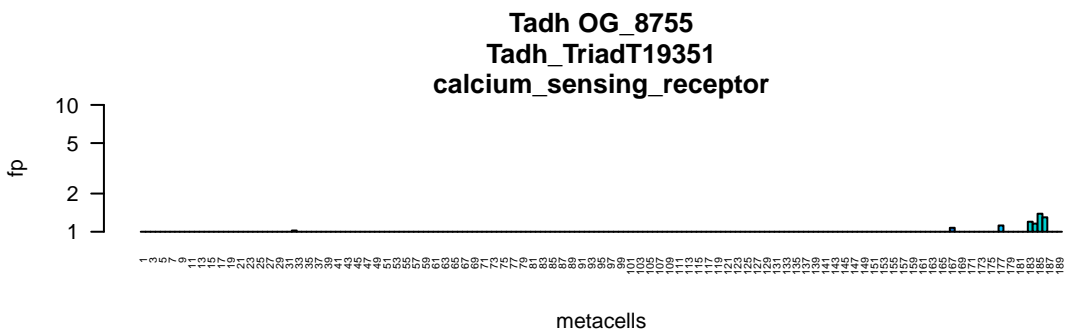
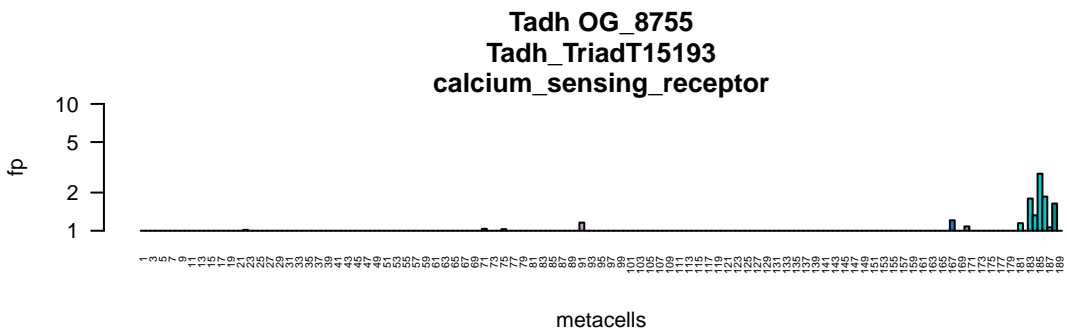


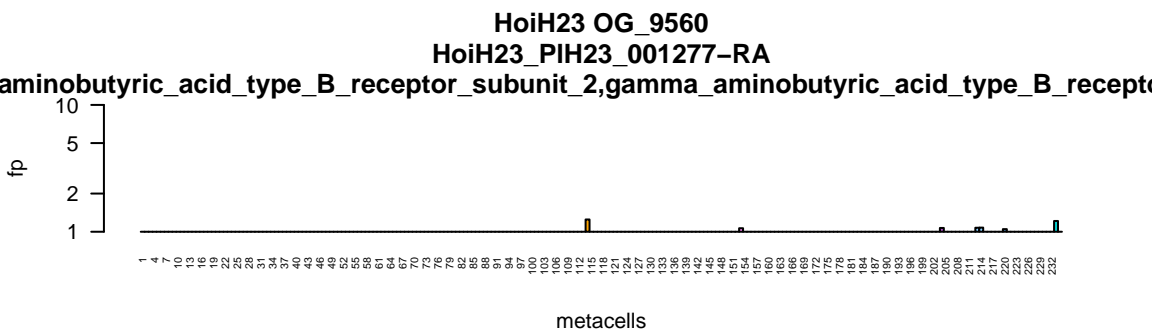
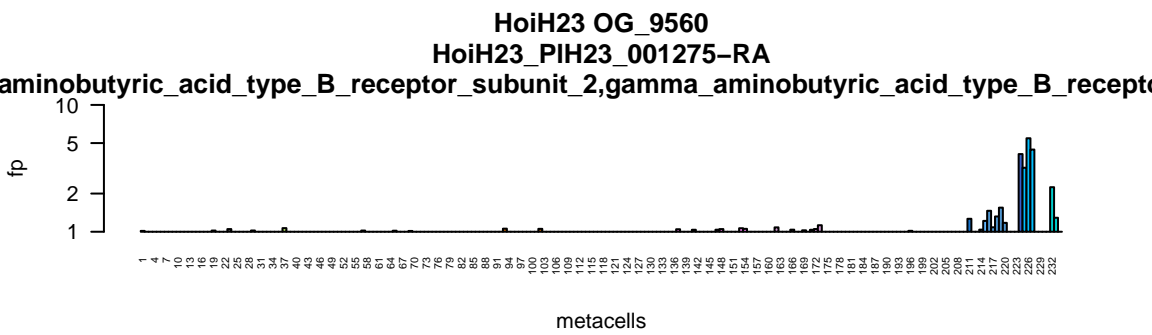
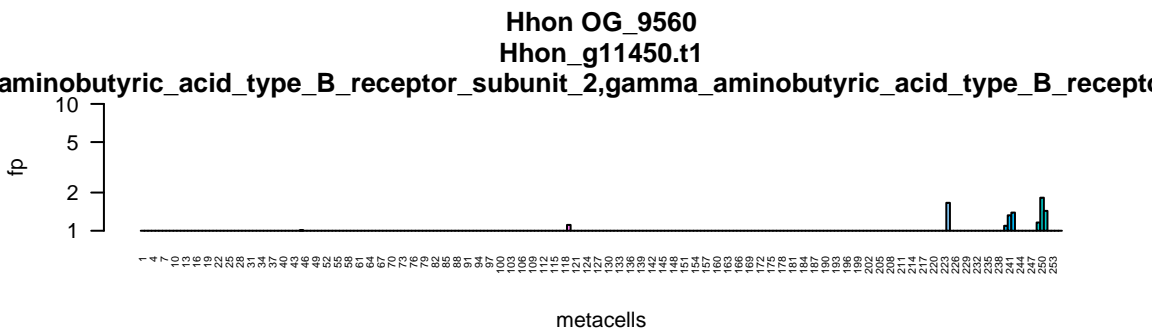
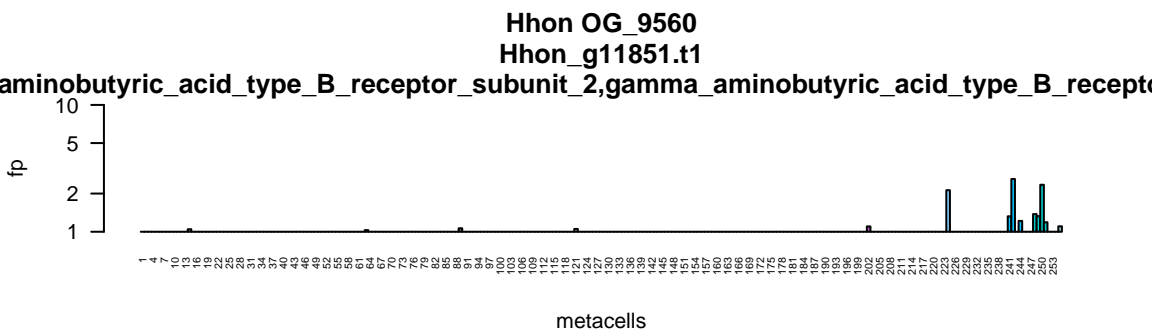
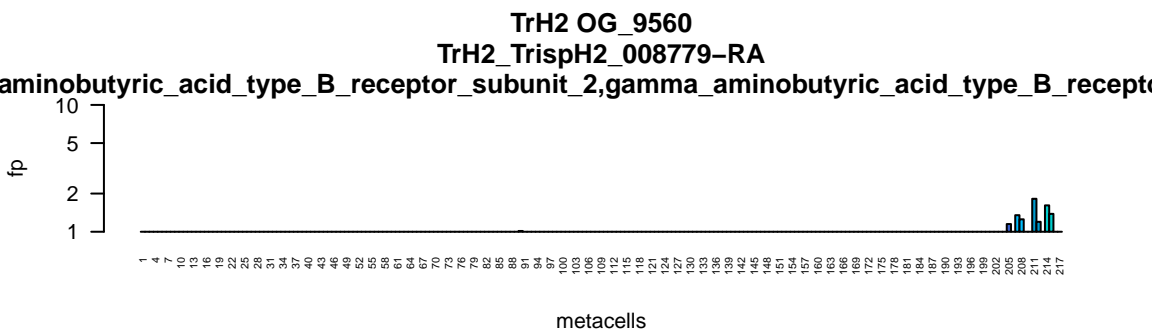
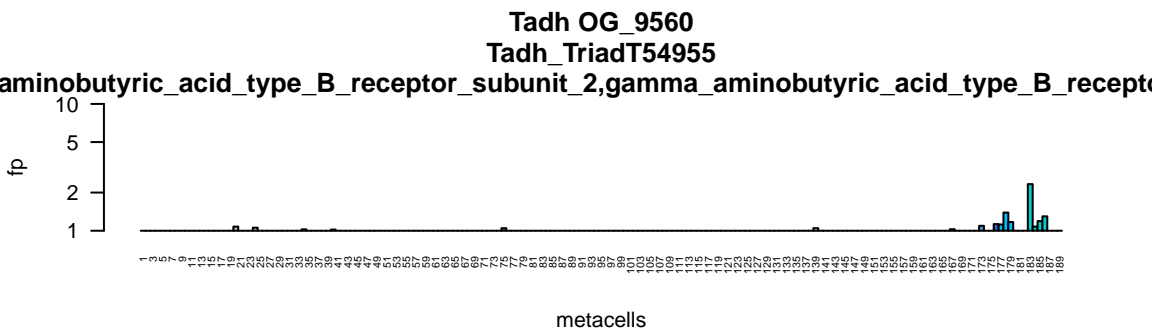




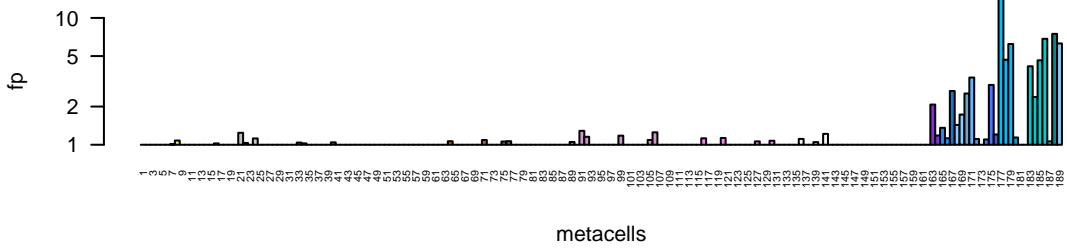




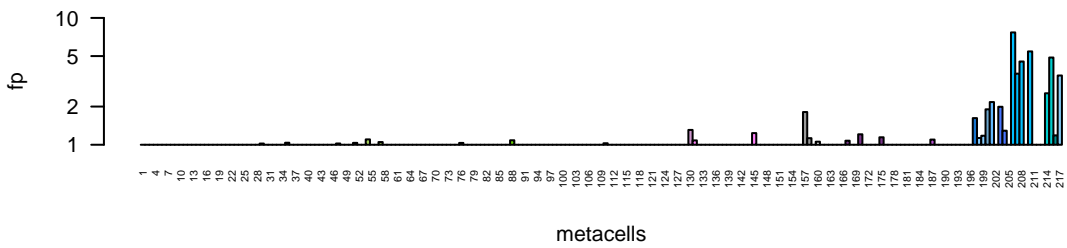




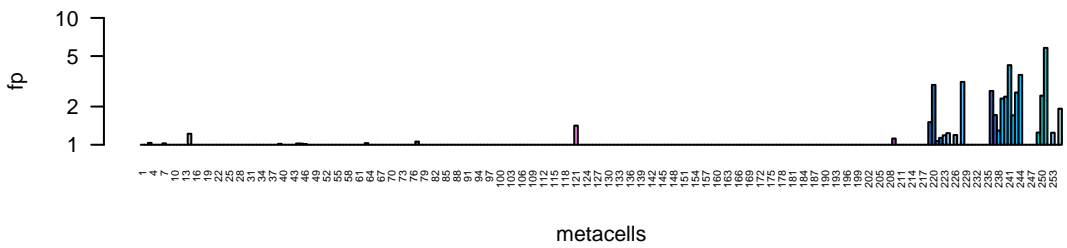
Tadh OG_1478
Tadh_wf_g4010.t1



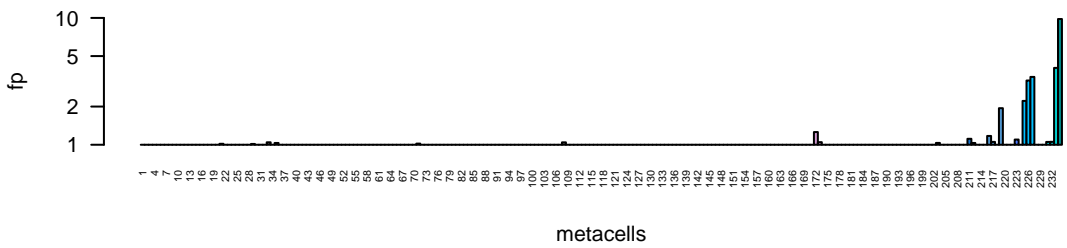
TrH2 OG_1478
TrH2_TrispH2_001729-RA

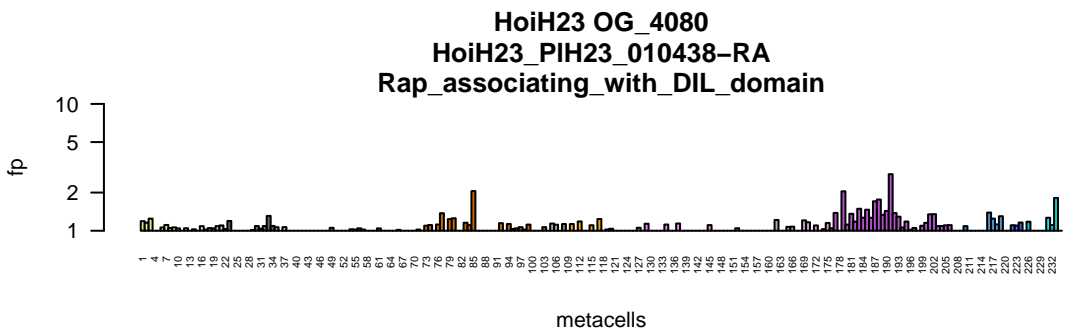
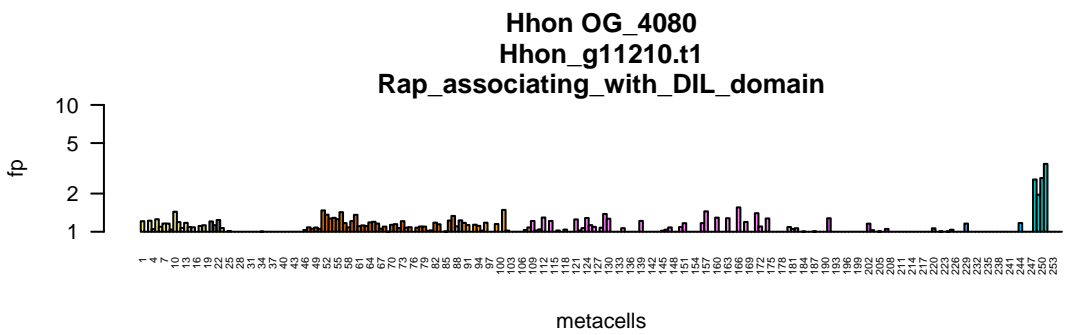
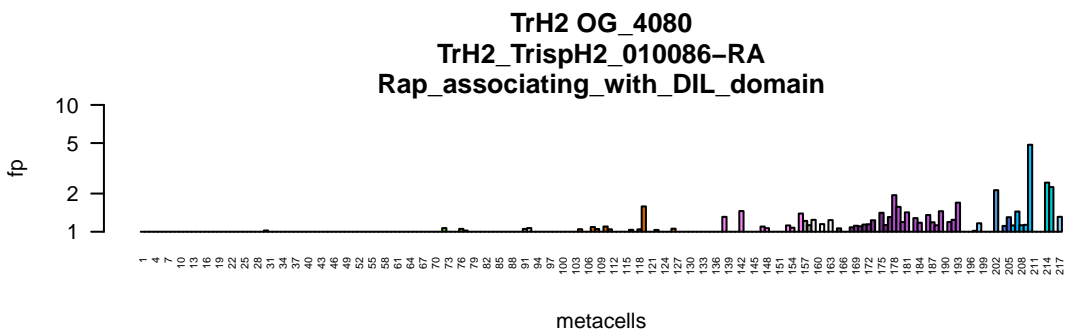
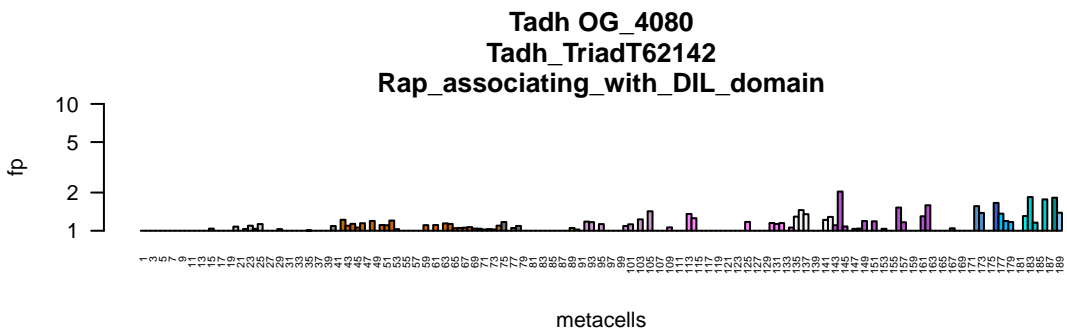


Hhon OG_1478
Hhon_g05248.t1



HoiH23 OG_1478
HoiH23_PIH23_010817-RA





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

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

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

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

metacell	fp
1	0
4	0
7	0
10	0
13	0
16	0
19	0
22	0
25	0
28	0
31	0
34	0
37	0
40	0
43	0
46	0
49	0
52	1
55	0
58	0
61	0
64	0
67	0
70	0
73	0
76	0
79	0
82	0
85	0
88	0
91	0
94	0
97	0
100	0
103	0
106	0
109	0
112	0
115	0
118	0
121	0
124	0
127	0
130	0
133	1
136	0
139	0
142	0
145	0
148	0
151	0
154	1
155	1
156	0
159	0
162	0
165	0
168	0
171	0
174	0
177	0
180	0
183	0
186	0
189	1
192	0
195	0
198	0
201	2
202	2
203	1
204	1
205	1
206	1
207	1
208	1
209	1
210	1
211	1
214	1
217	1

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacell	fp
1	1
4	1
13	1
16	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	1.1
77	1
79	1
82	1
85	1
88	1
94	1
97	1
100	1.3
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1.1
133	1.2
136	1
139	1
142	1
145	1
148	1
151	1
154	1.3
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	1.8
199	1.5
202	2.2
205	1.5
208	1.2
211	1.2
214	1.2
217	1.5

A bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists indices from 1 to 217. Most metacells have 1 false positive, but metacells 202 and 214 have 2 false positives.

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

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

fp

metacells

Hhon OG_4305
Hhon_g09074.t1

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

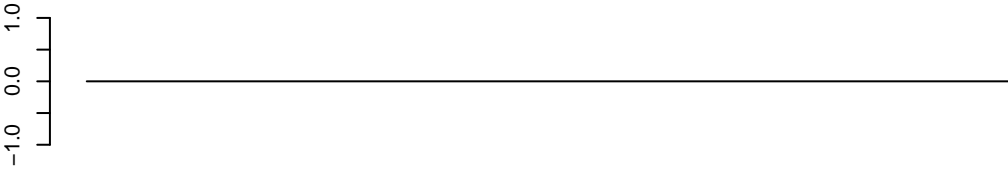
fp

metacells

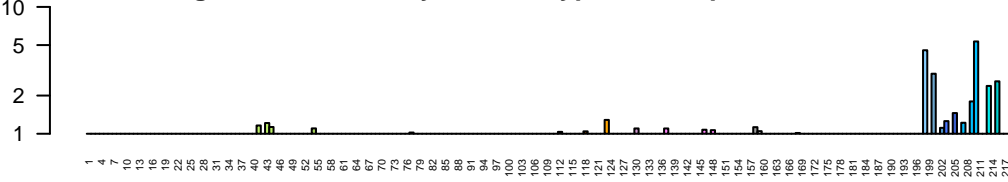
A bar chart showing the frequency of metacells (x-axis) versus the frequency of pairs (fp, y-axis). The x-axis is labeled 'metacells' and ranges from 1 to 232. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a few outliers reaching up 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
52	1
55	1
58	1
64	1
67	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

gamma_aminobutyric_acid_type_B_receptor_subunit_2
Tadh | no data

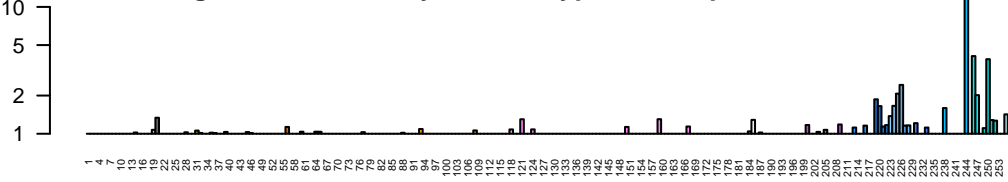


TrH2 OG_4636
TrH2_TrispH2_003379-RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2



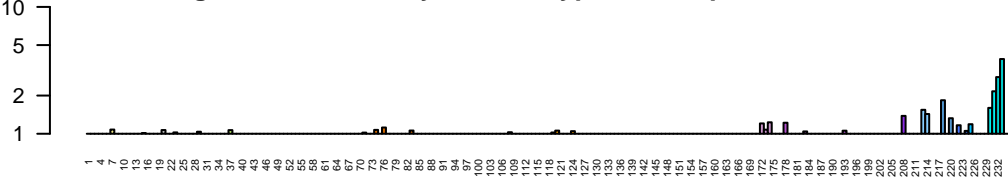
metacells

Hhon OG_4636
Hhon_g07295.t1
gamma_aminobutyric_acid_type_B_receptor_subunit_2

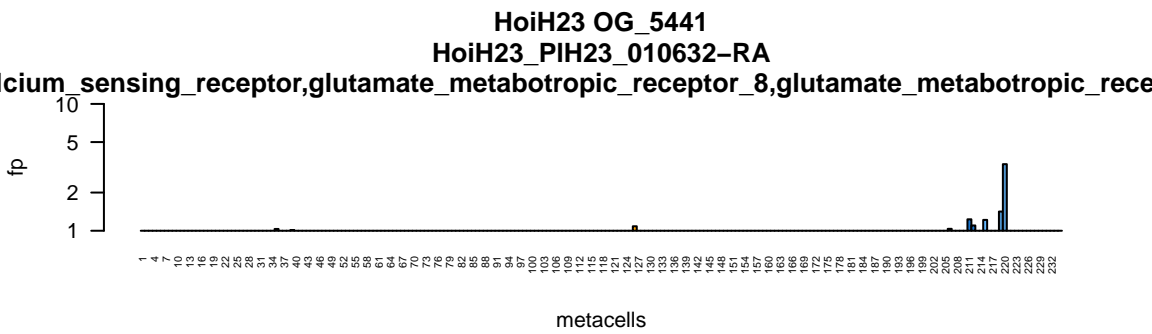
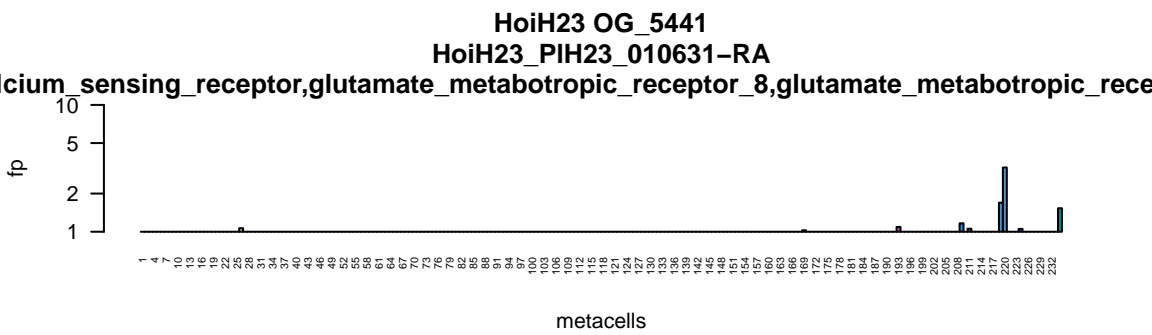
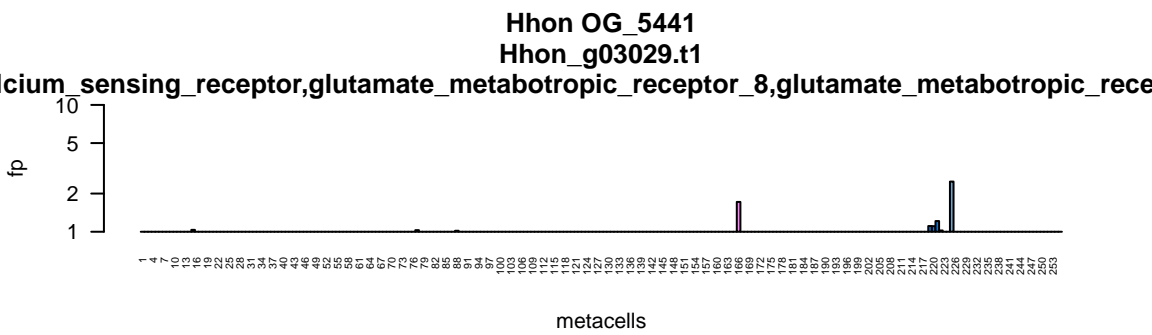
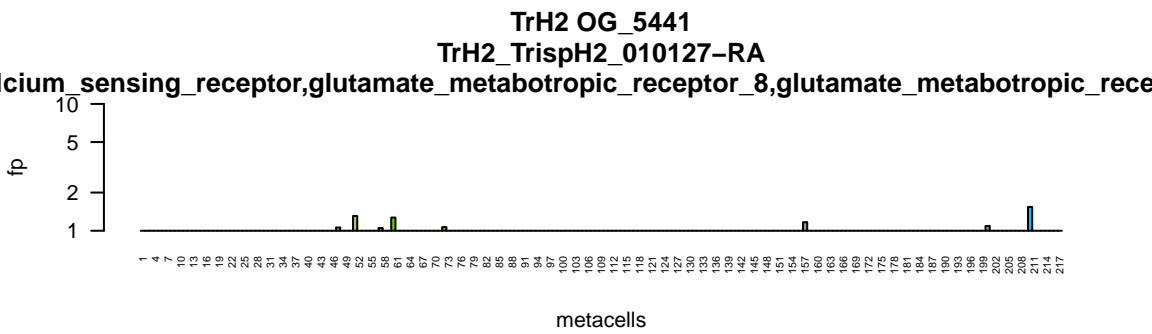
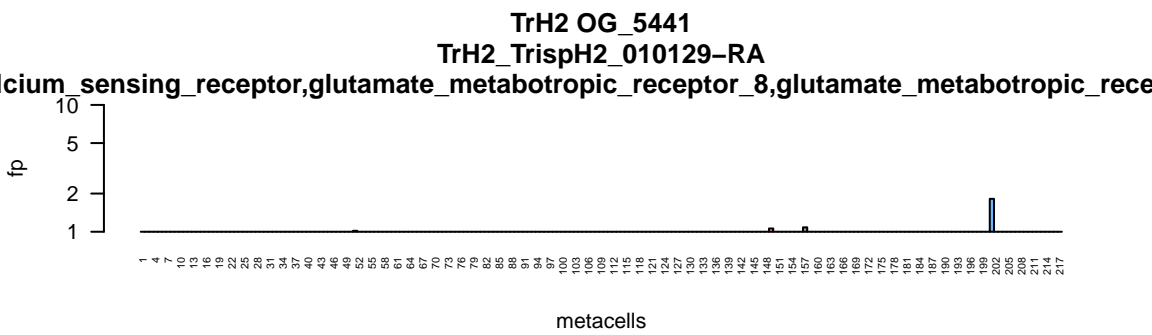
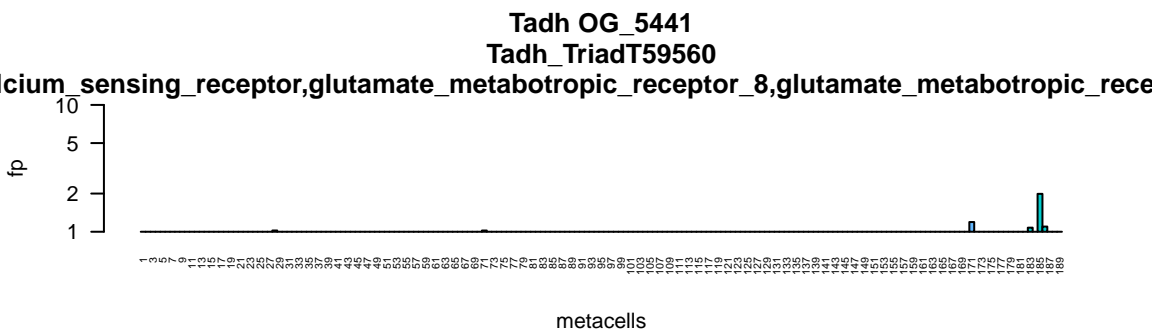
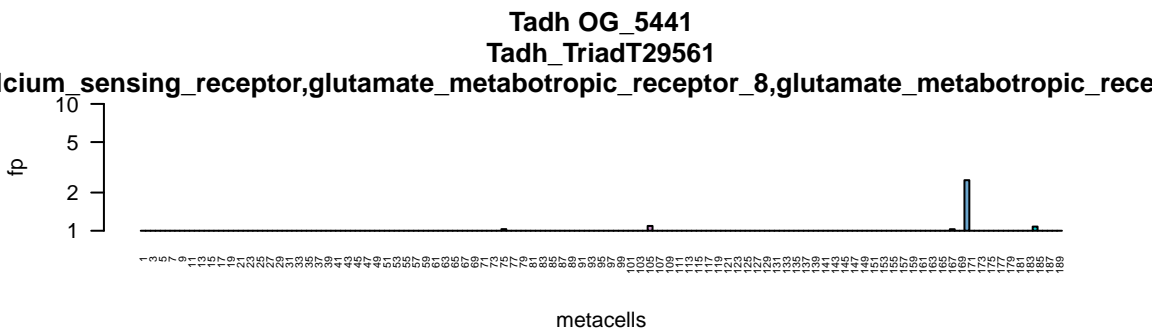


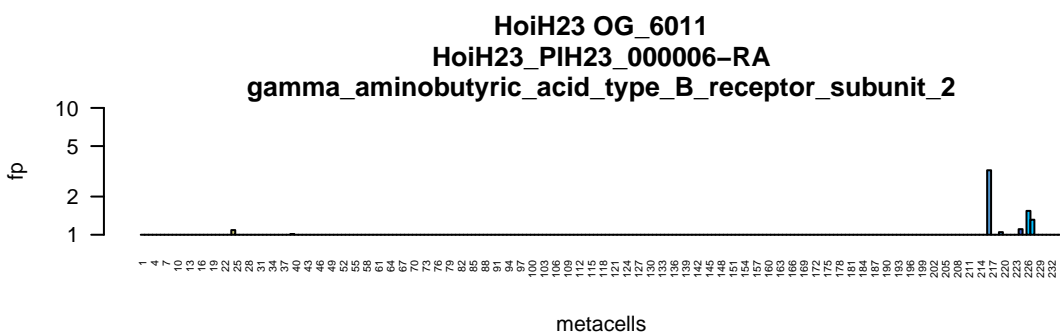
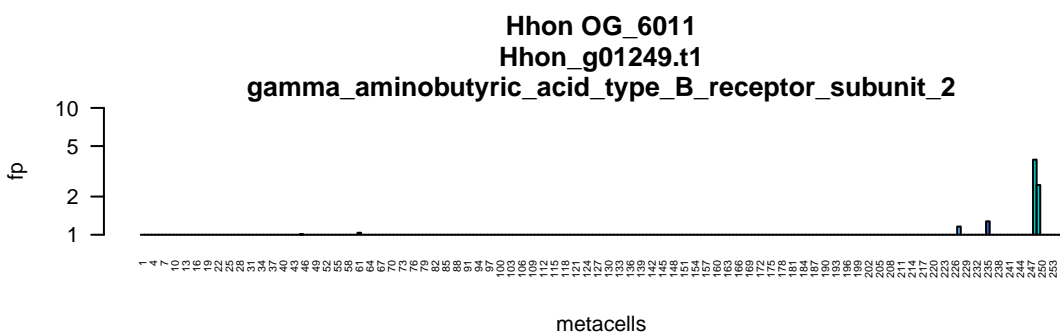
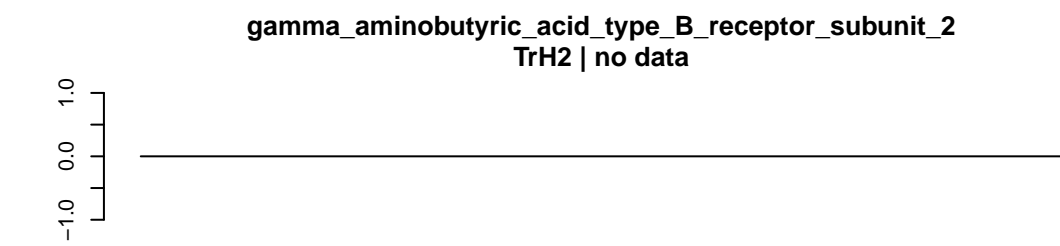
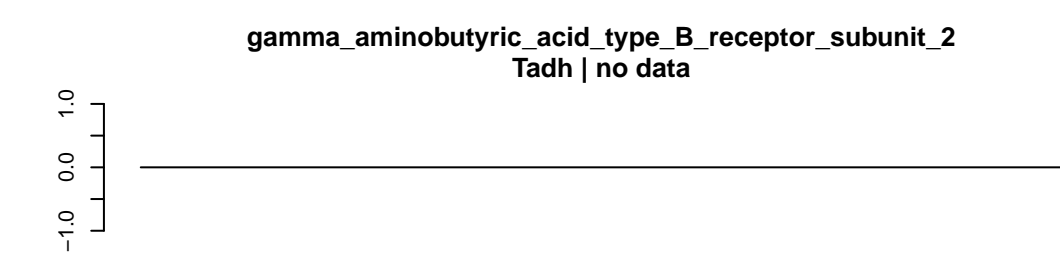
metacells

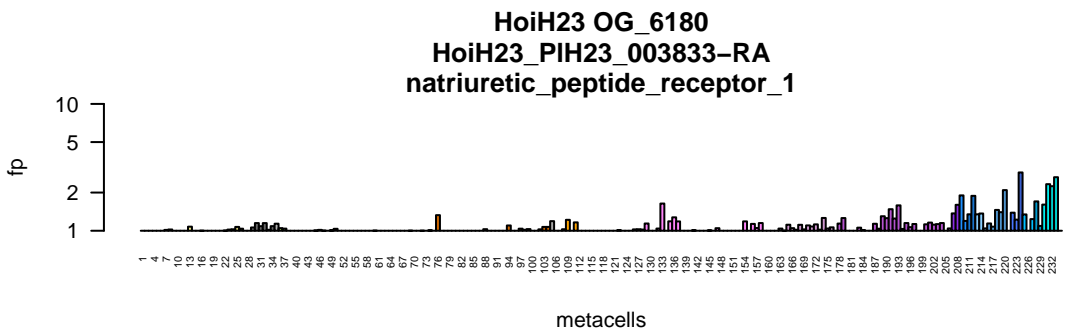
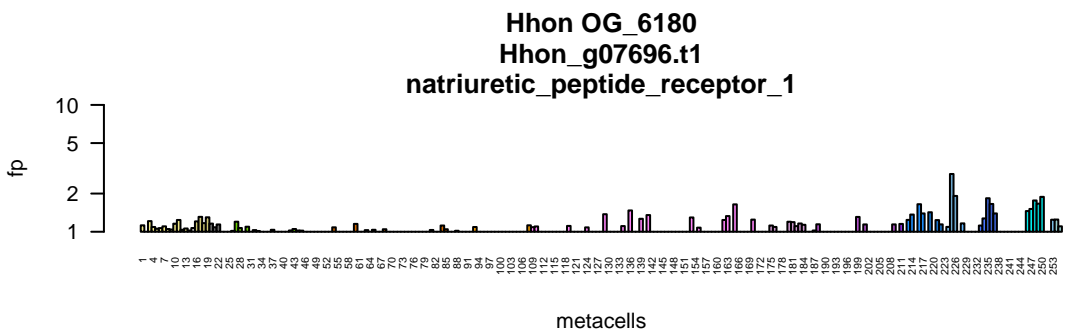
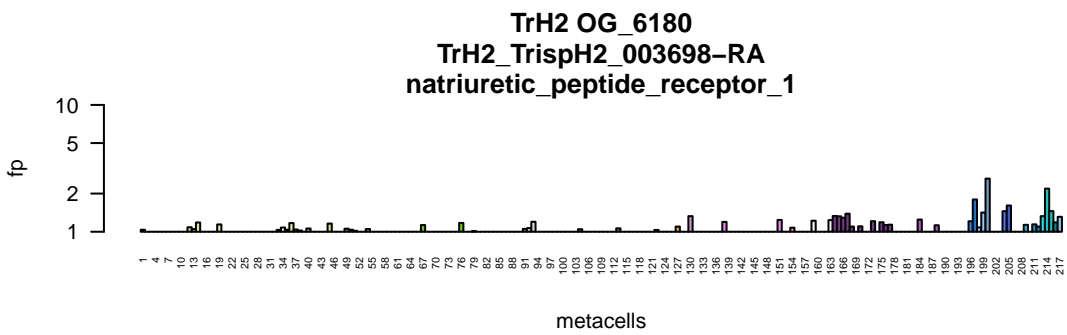
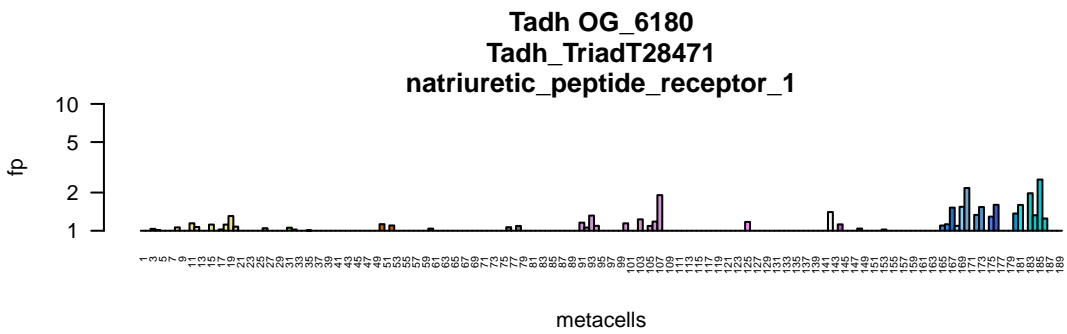
HoiH23 OG_4636
HoiH23_PIH23_010039-RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2



metacells



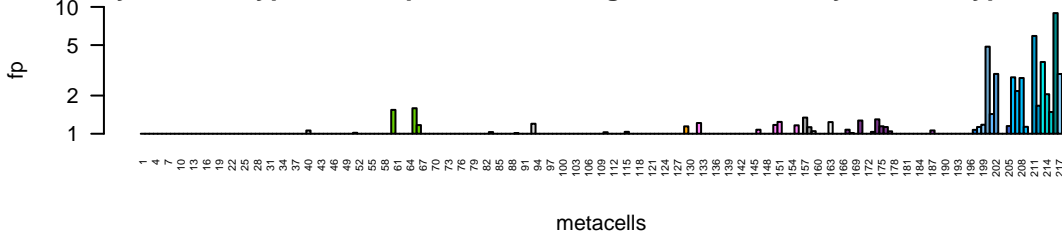




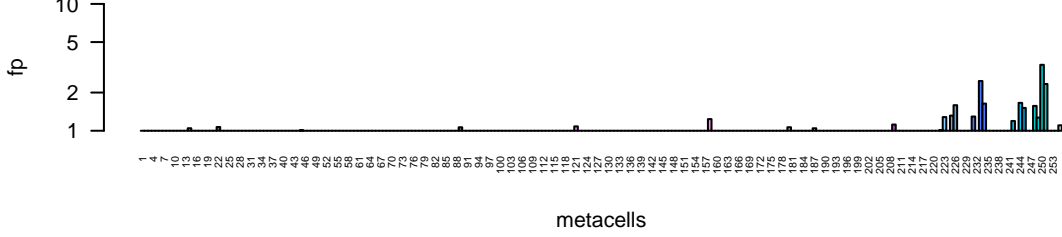
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recepto
Tadh | no data



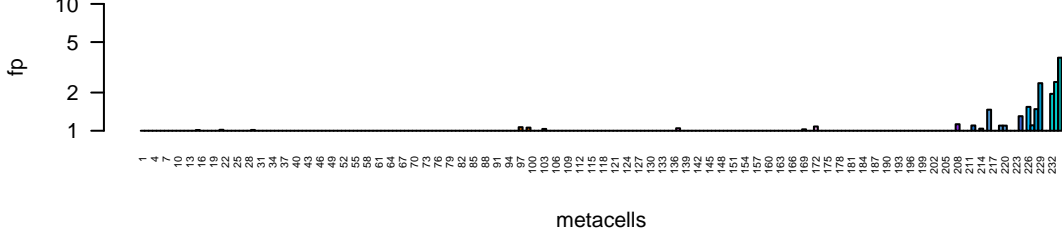
TrH2 OG_7186
TrH2_TrispH2_010138-RA
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recepto



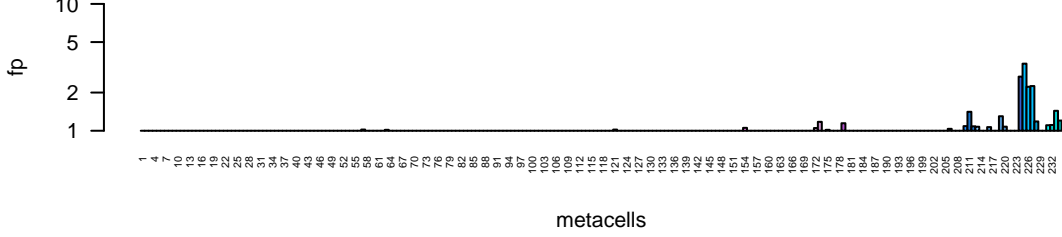
Hhon OG_7186
Hhon_g01977.t1
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recepto



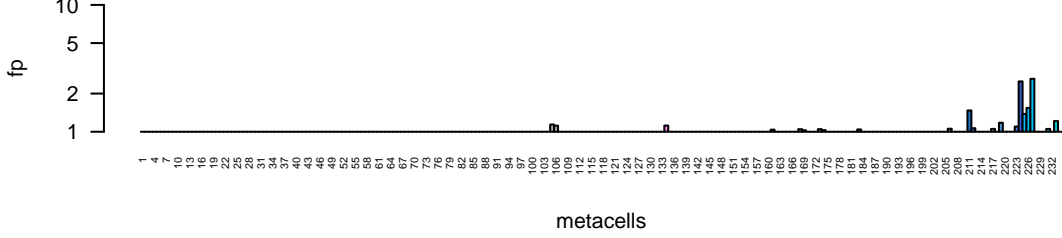
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HoiH23_PIH23_011144-RA
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recepto



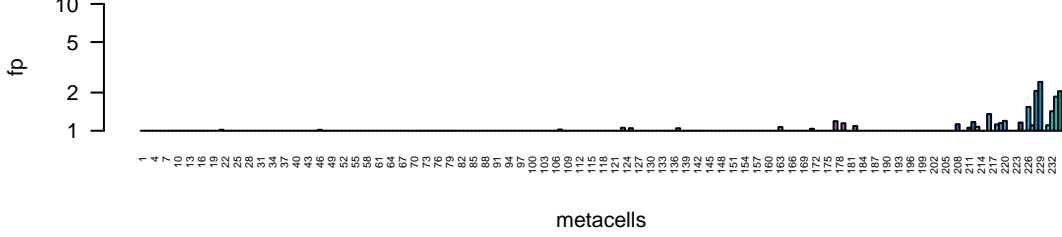
HoiH23 OG_7186
HoiH23_PIH23_011313-RA
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recepto

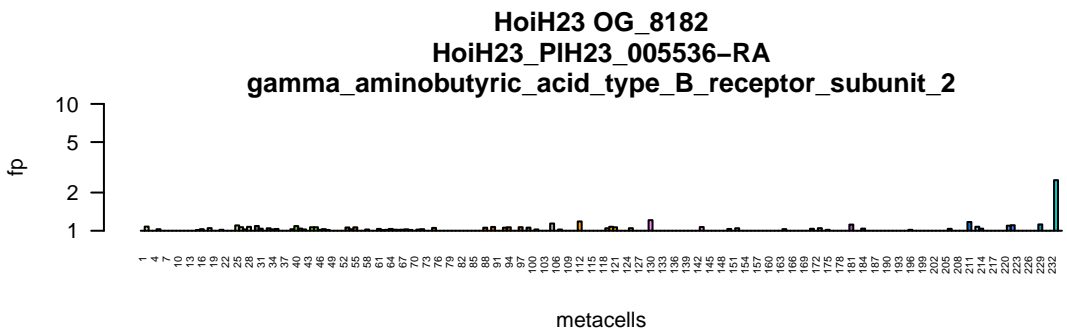
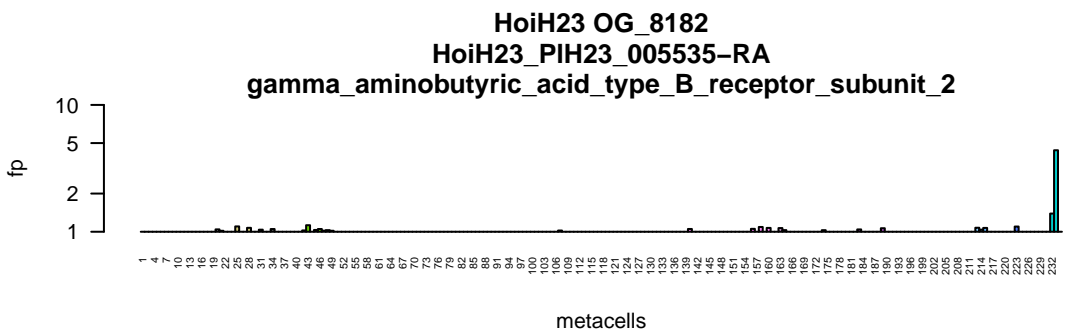
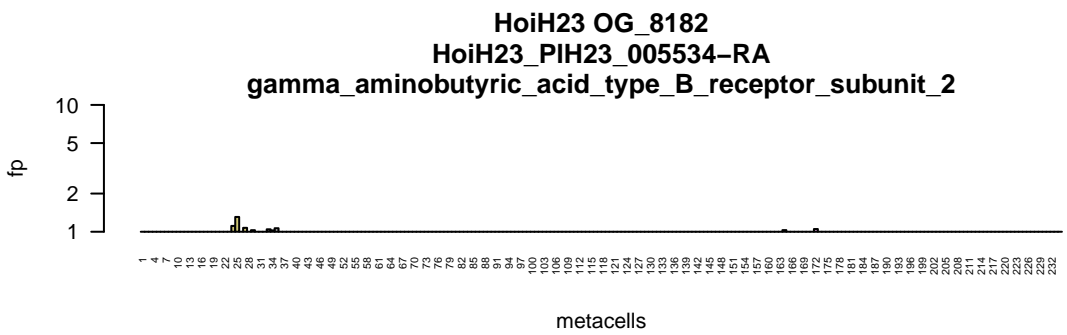
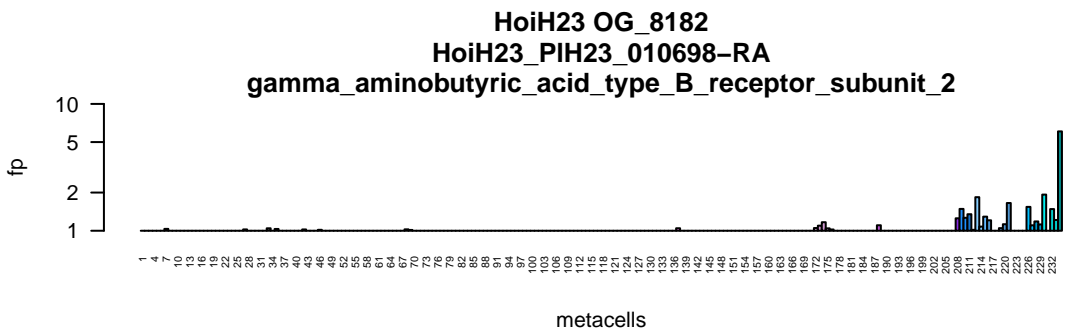
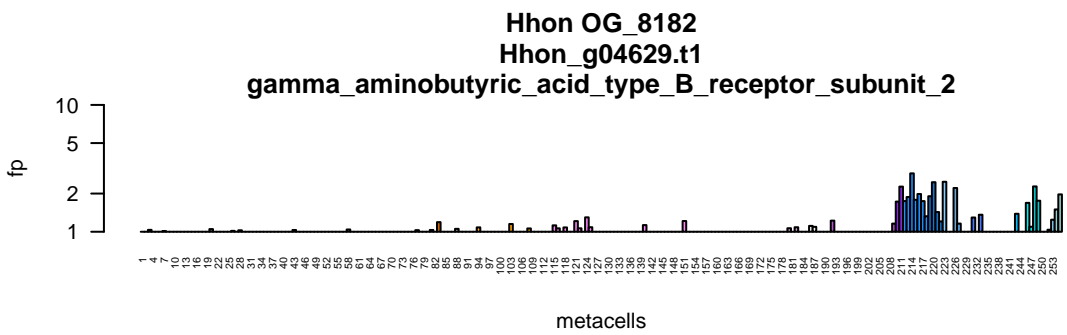
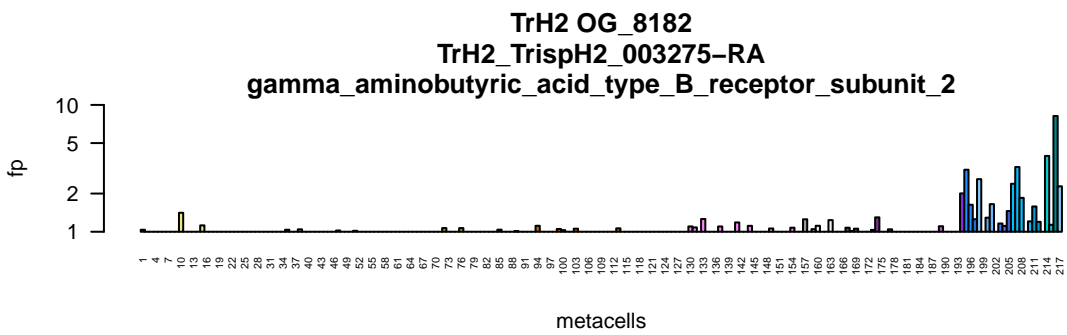
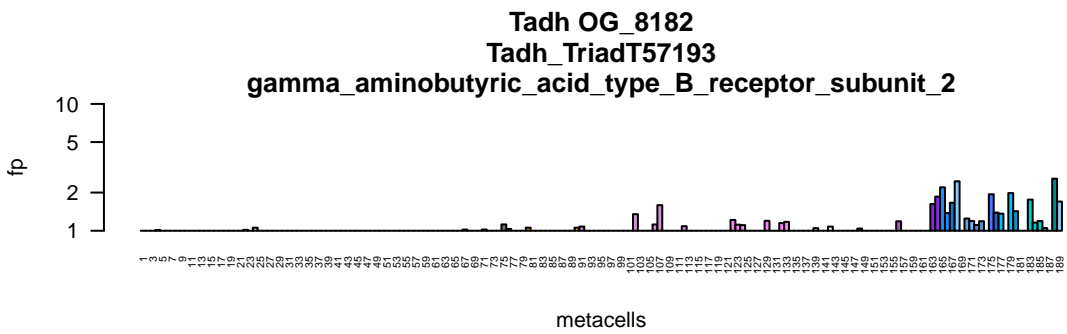


HoiH23 OG_7186
HoiH23_PIH23_004027-RA
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recepto

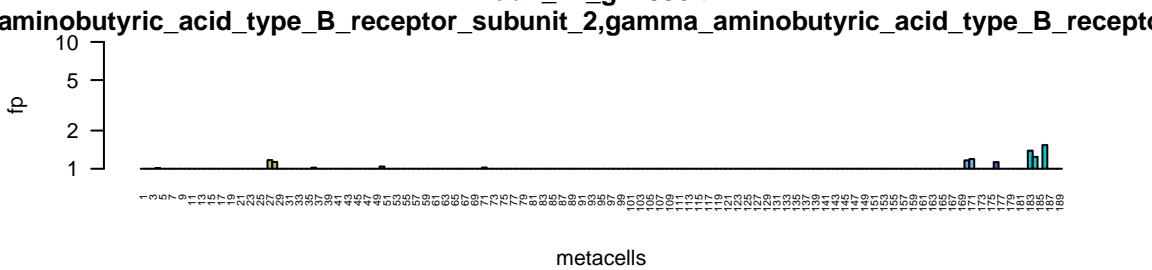


HoiH23 OG_7186
HoiH23_PIH23_011312-RA
aminobutyric_acid_type_B_receptor_subunit_2,gamma_aminobutyric_acid_type_B_recepto

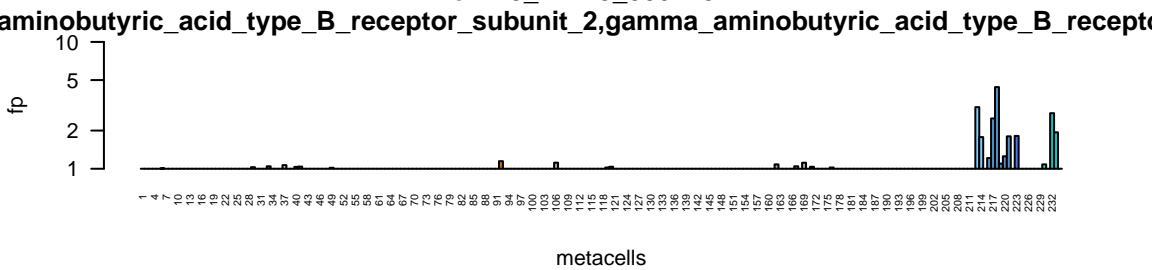




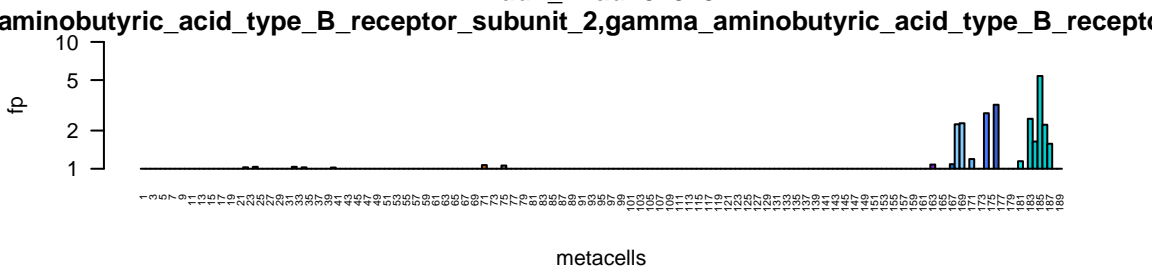
Tadh OG_8960
tadh_wf_g11896.t1



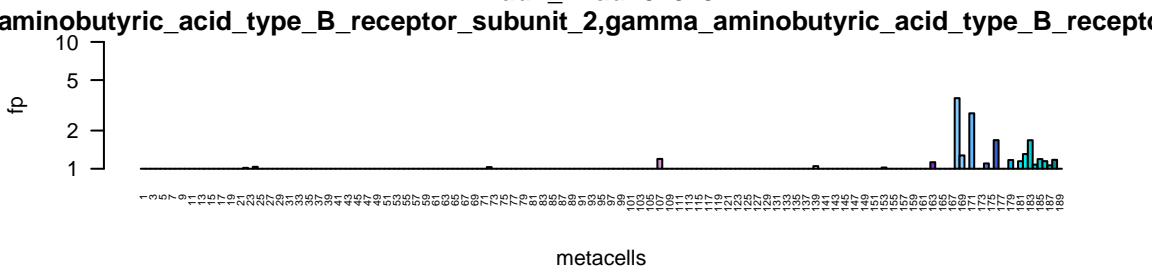
HoiH23 OG_8960
HoiH23_PIH23_000719-RA



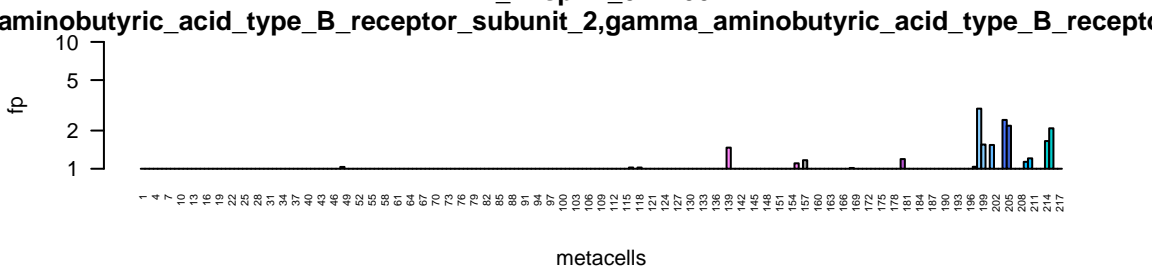
Tadh OG_8960
Tadh_TriadT52578



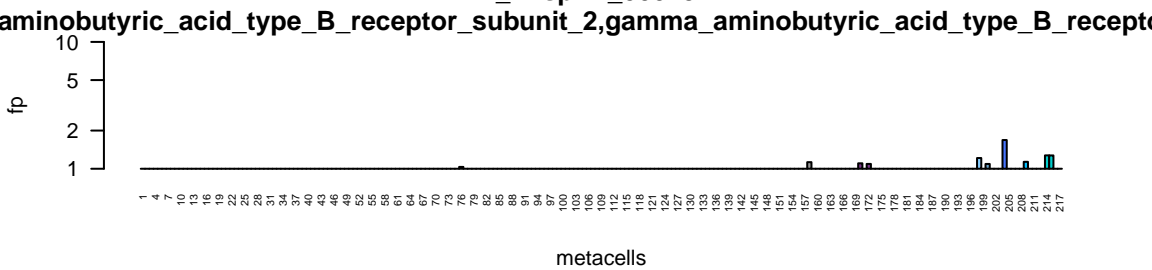
Tadh OG_8960
Tadh_TriadT52579



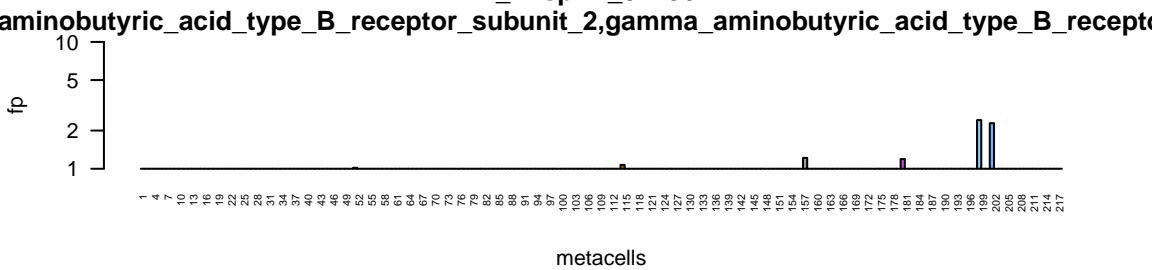
TrH2 OG_8960
TrH2_TrispH2_012100-RA



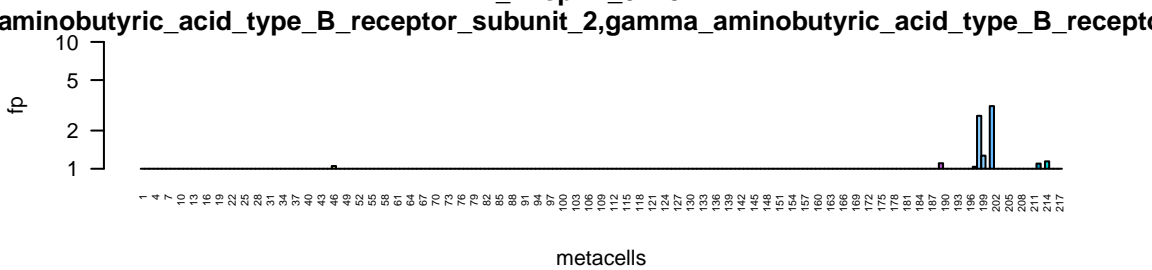
TrH2 OG_8960
TrH2_TrispH2_000234-RA



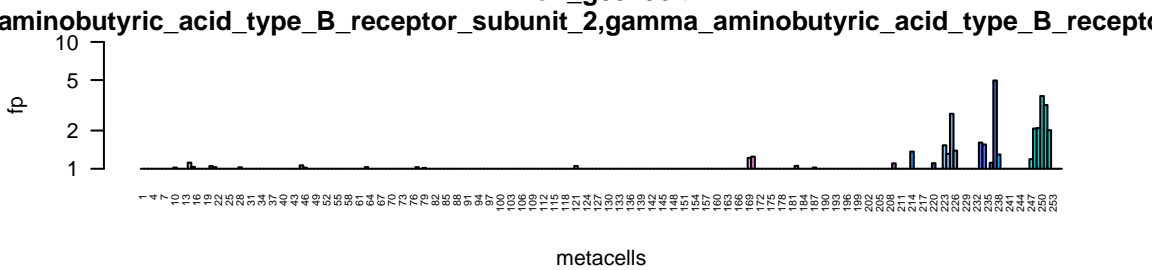
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TrH2_TrispH2_011964-RA

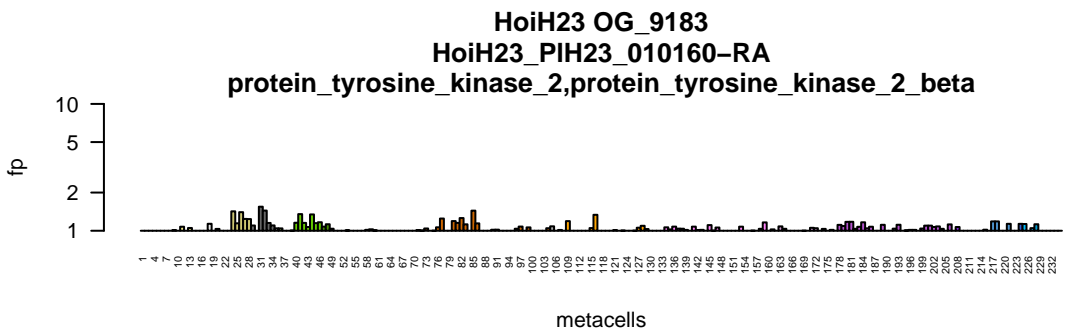
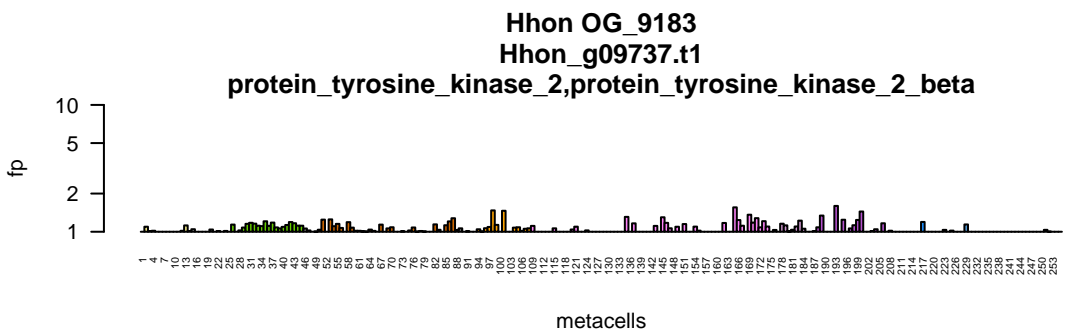
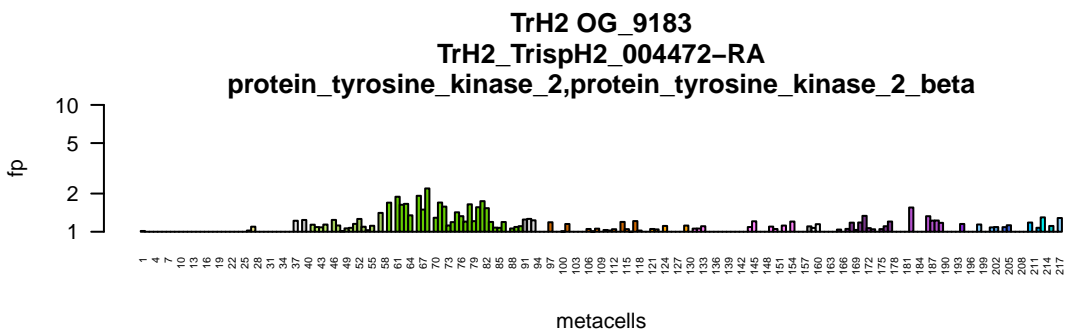
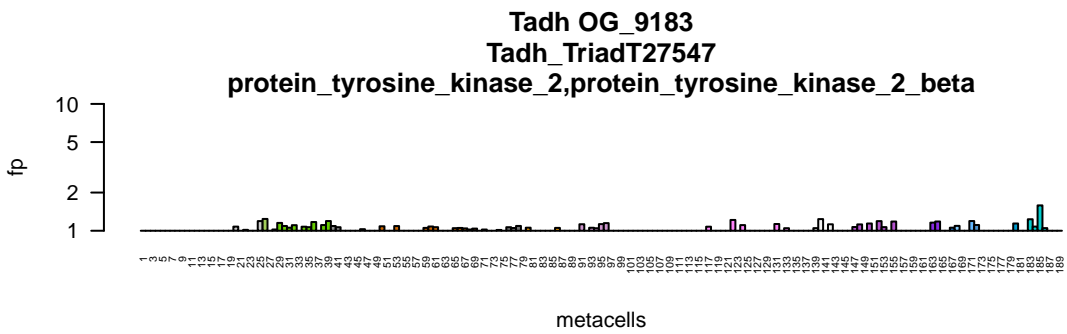


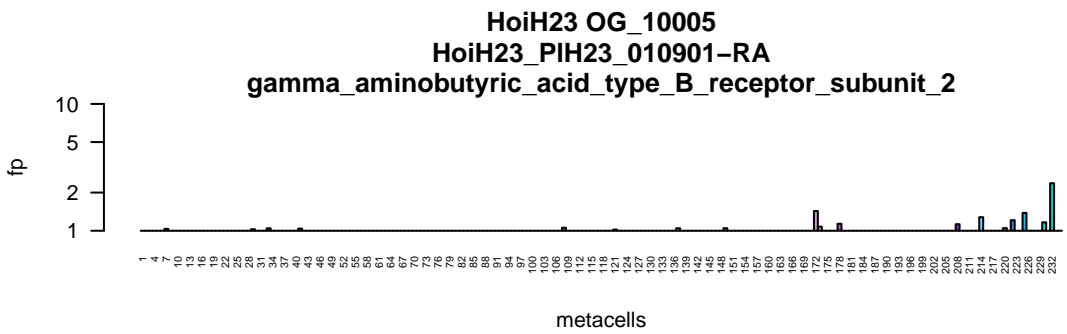
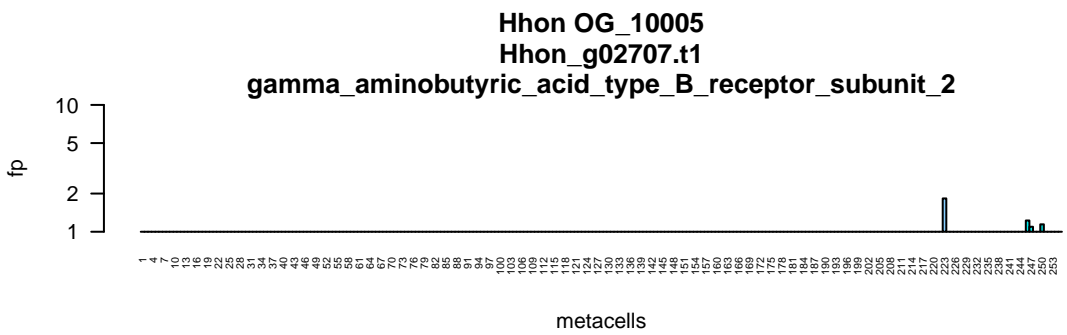
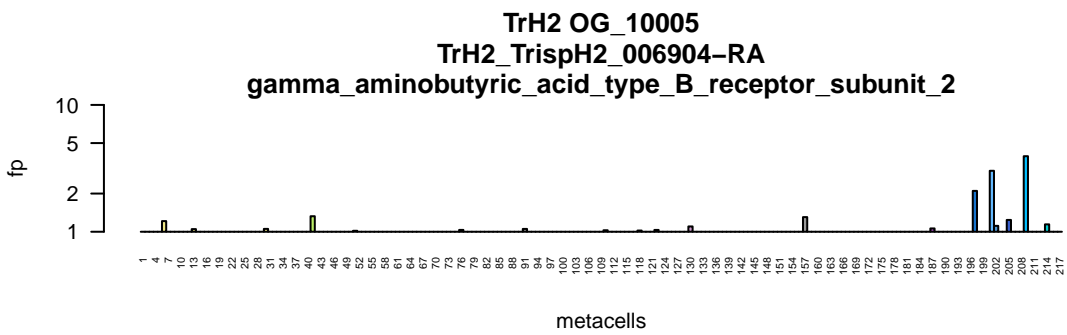
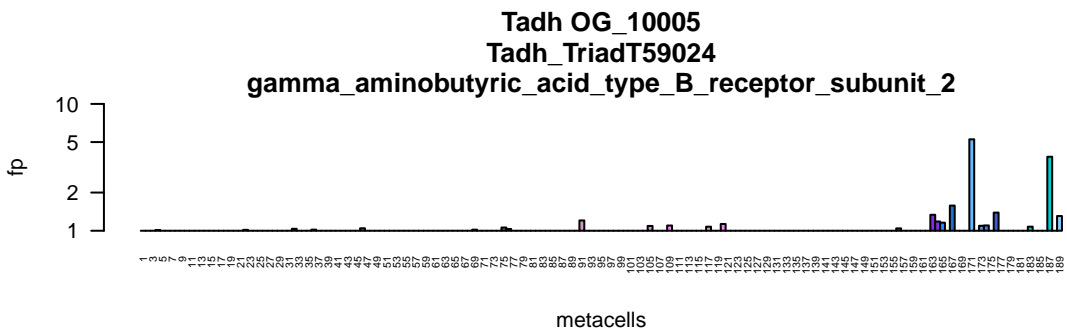
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TrH2_TrispH2_011847-RA

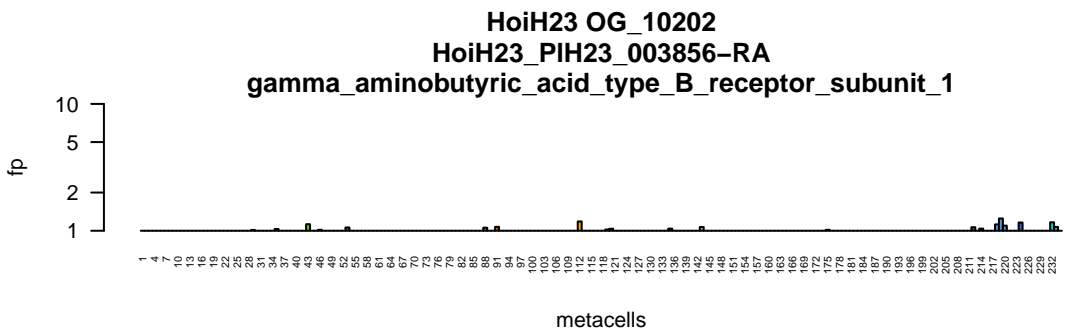
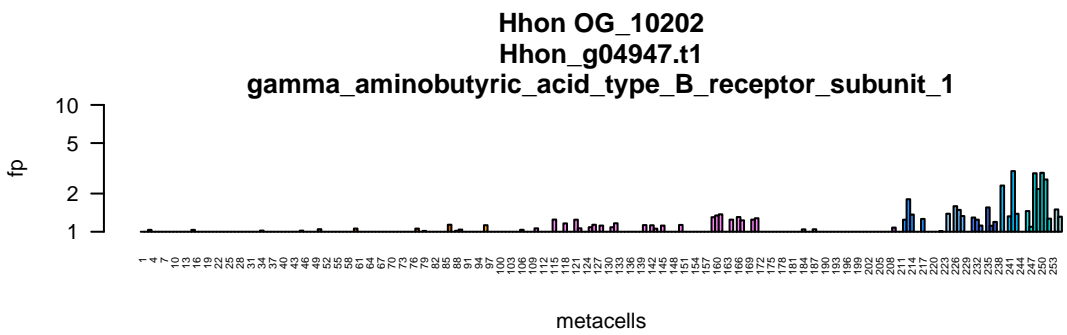
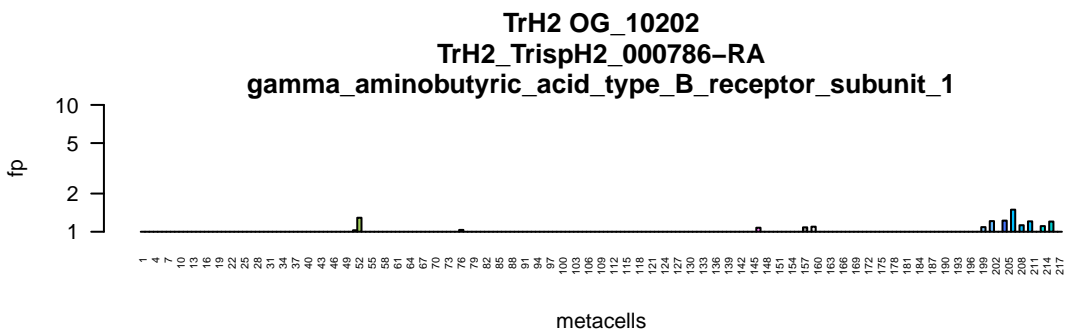
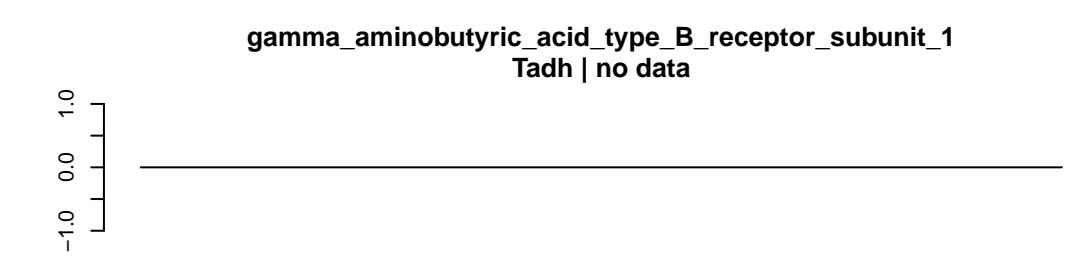


Hhon OG_8960
Hhon_g05296.t1

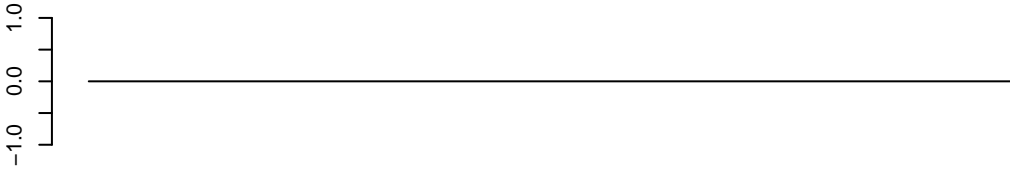




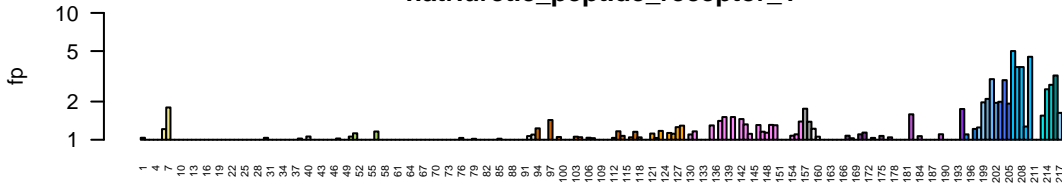




natriuretic_peptide_receptor_1
Tadh | no data

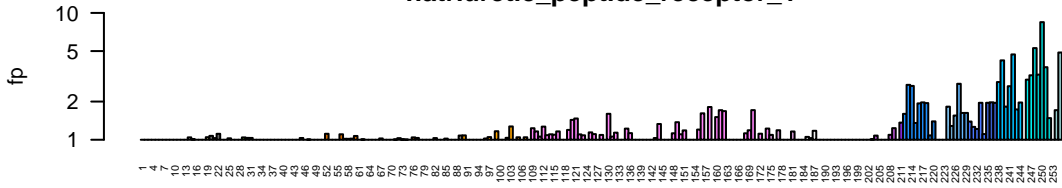


TrH2 OG_10203
TrH2_TrispH2_000784-RA
natriuretic_peptide_receptor_1



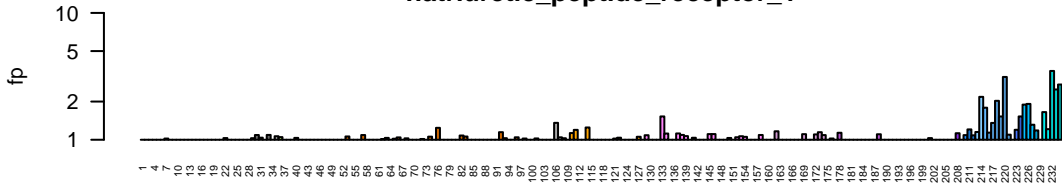
metacells

Hhon OG_10203
Hhon_g04948.t1
natriuretic_peptide_receptor_1



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HoiH23 OG_10203
HoiH23_PIH23_003854-RA
natriuretic_peptide_receptor_1



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