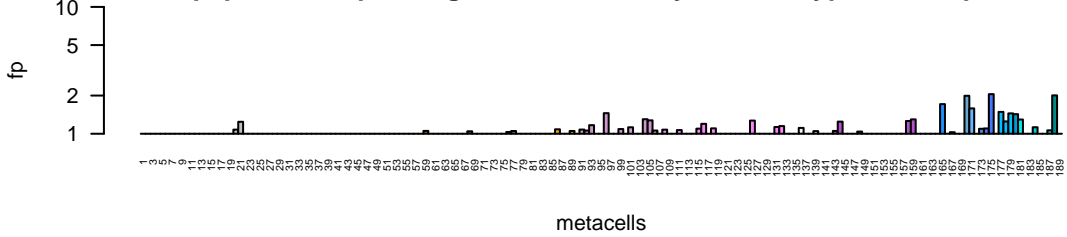


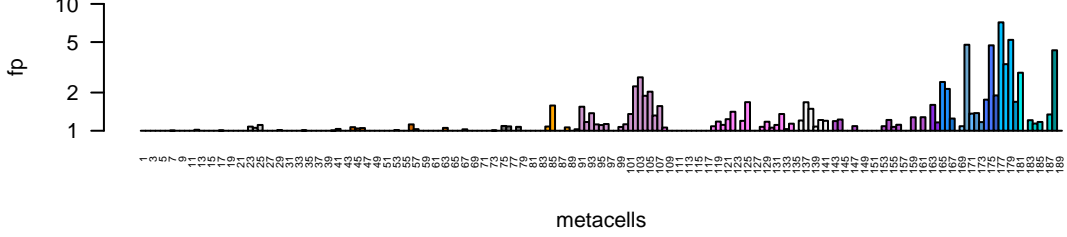
Tadh OG_5687
Tadh_TriadT59259

natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1



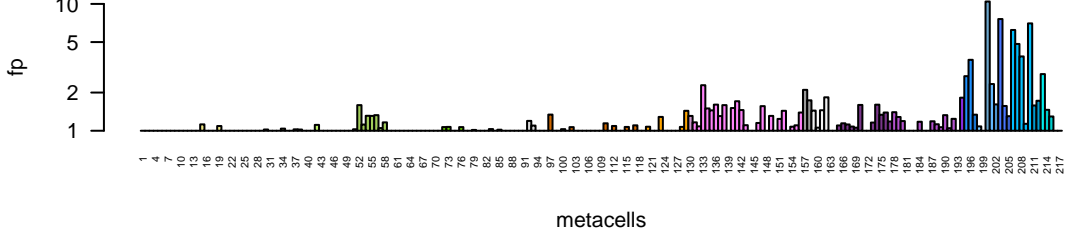
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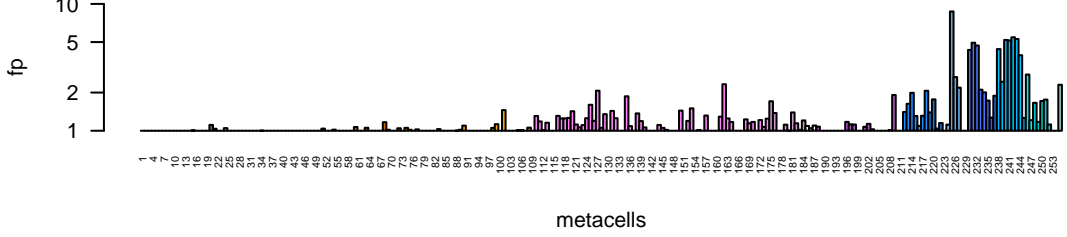
TrH2 OG_5687
TrH2_TrispH2_002051-RA

natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1



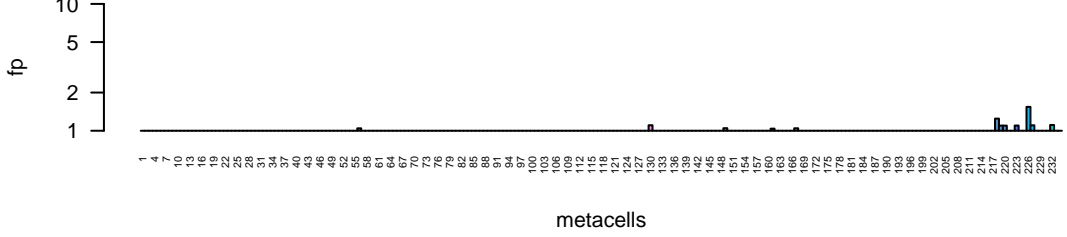
Hhon OG_5687
Hhon_g00715.t1

natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1



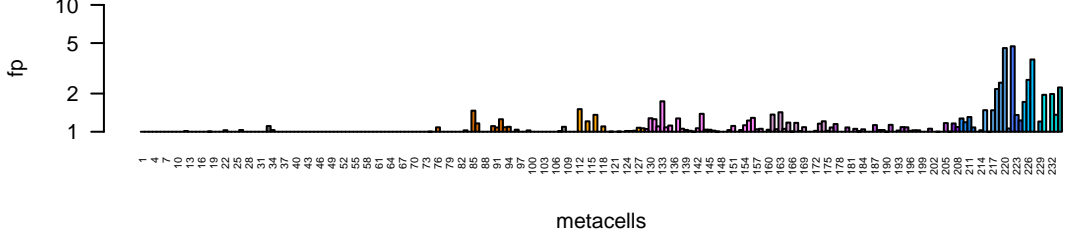
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HoiH23_PIH23_009782-RA

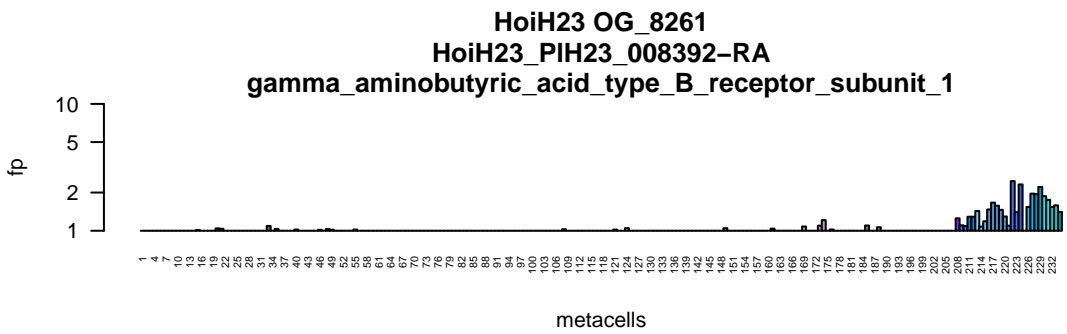
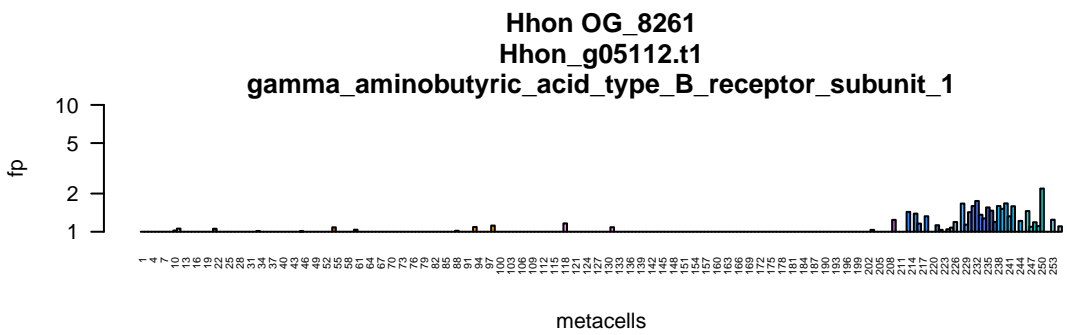
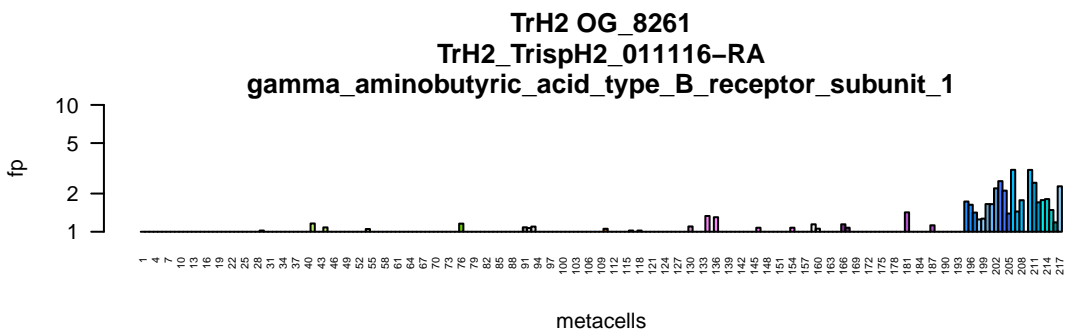
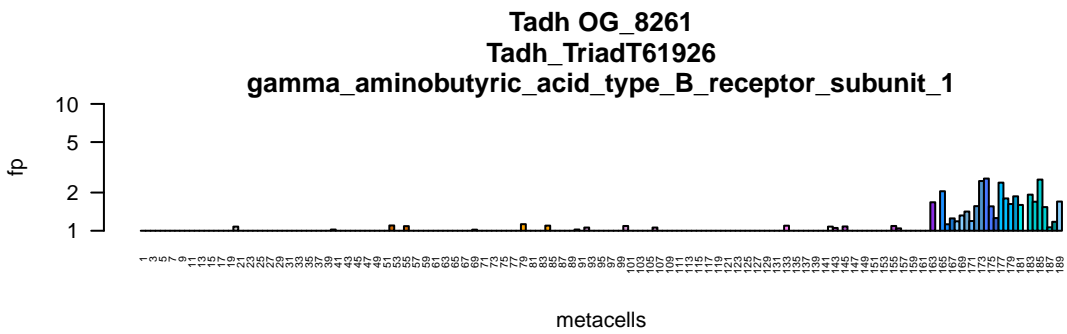
natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1

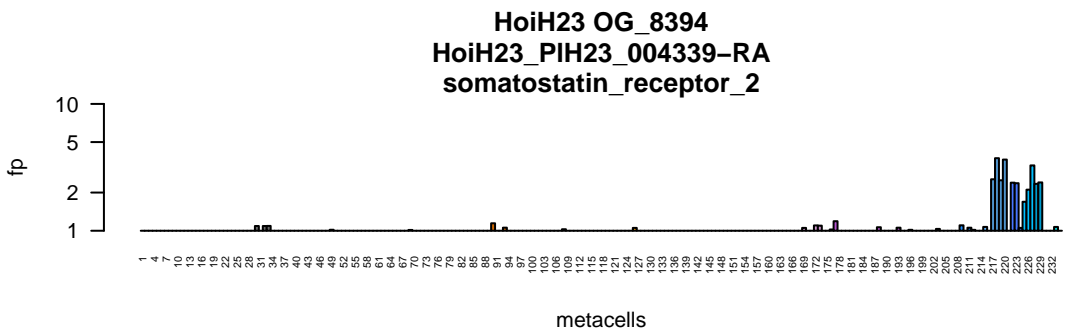
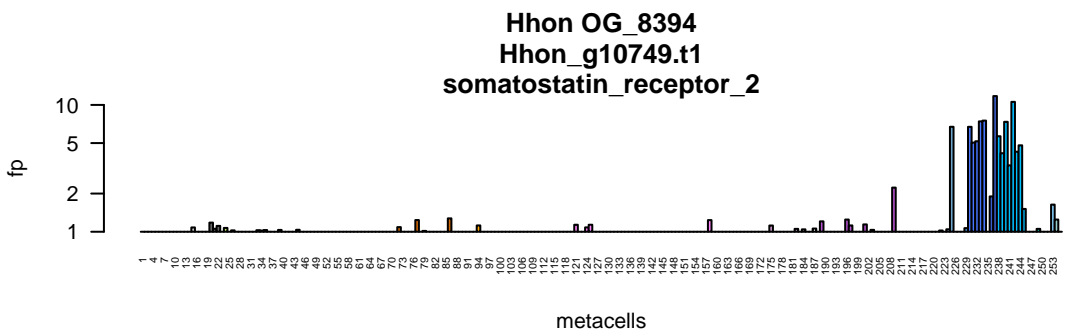
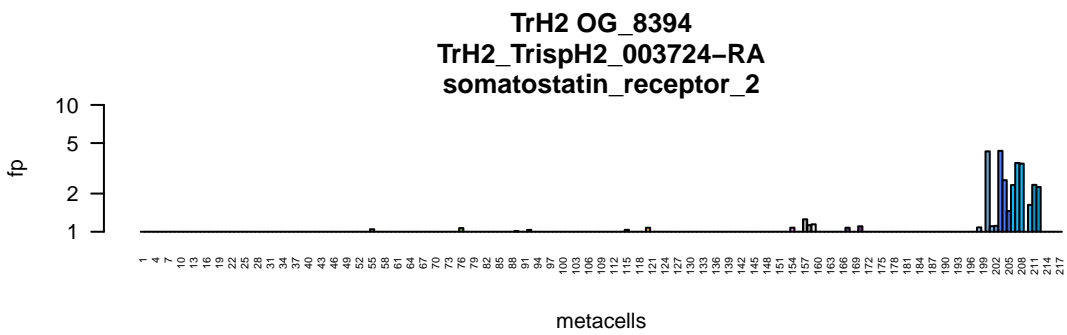
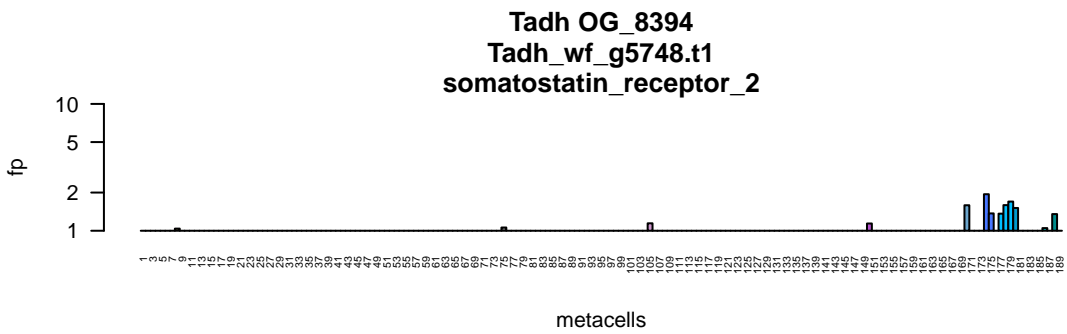


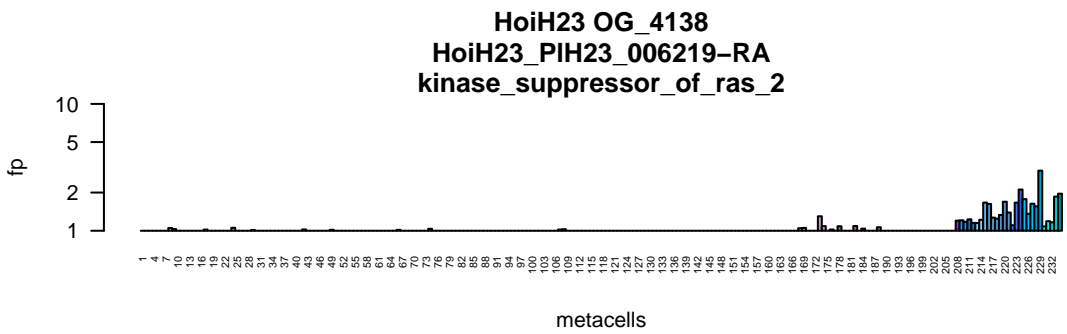
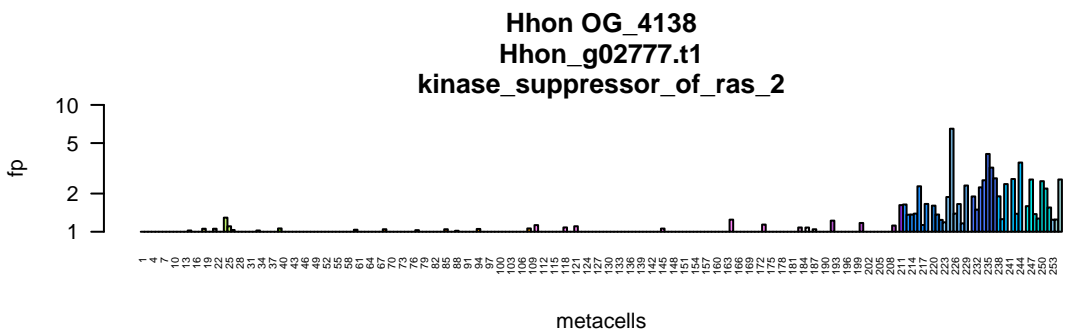
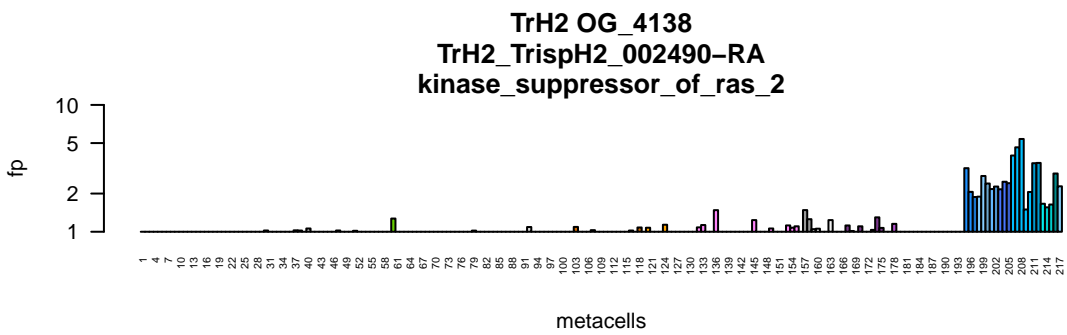
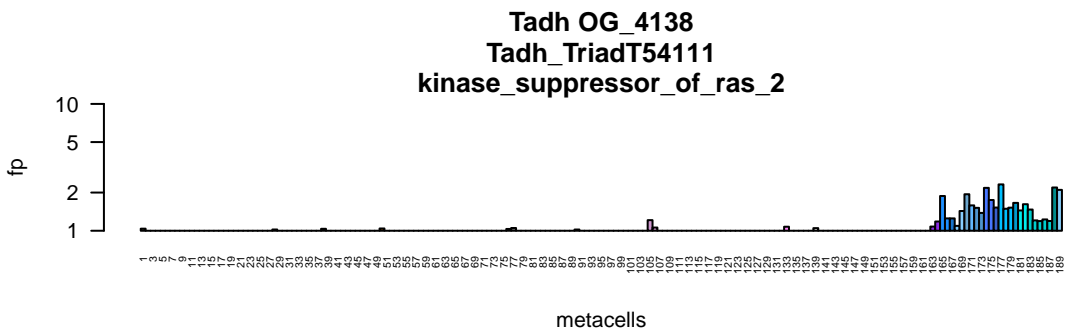
HoiH23 OG_5687
HoiH23_PIH23_009783-RA

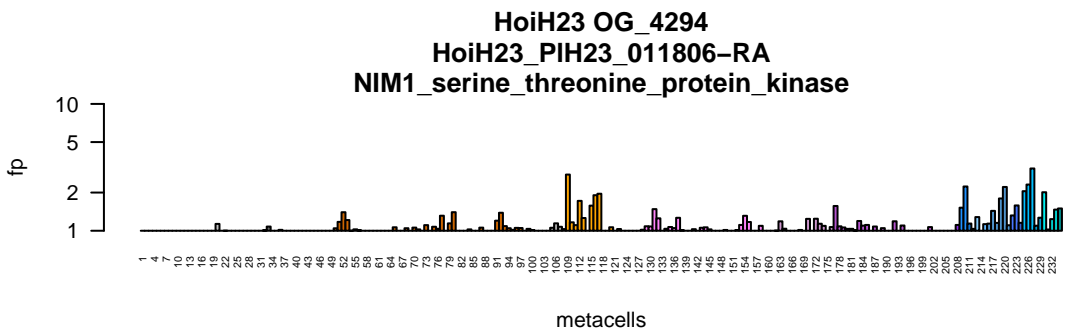
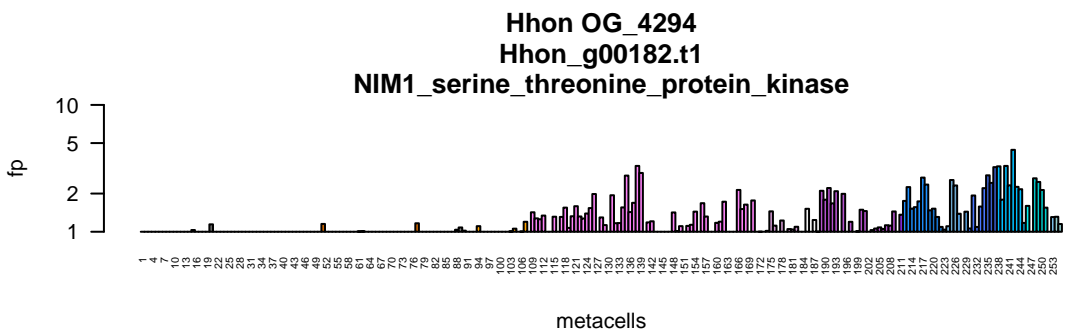
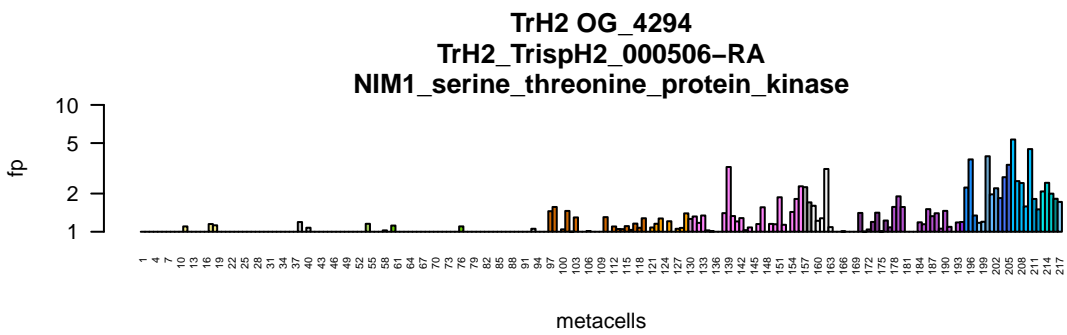
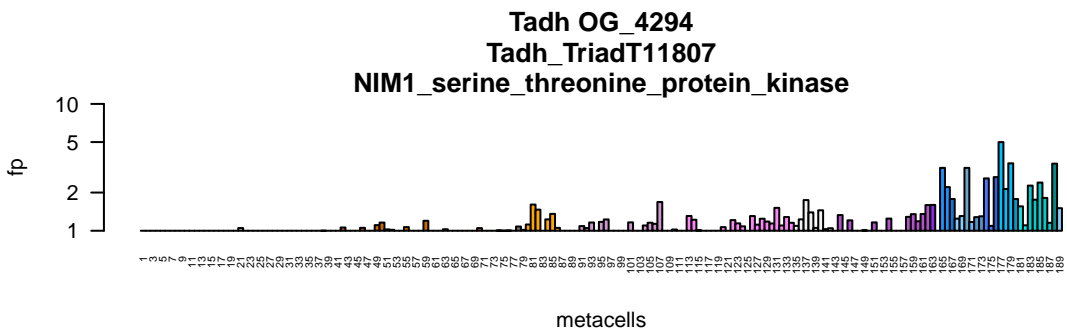
natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1



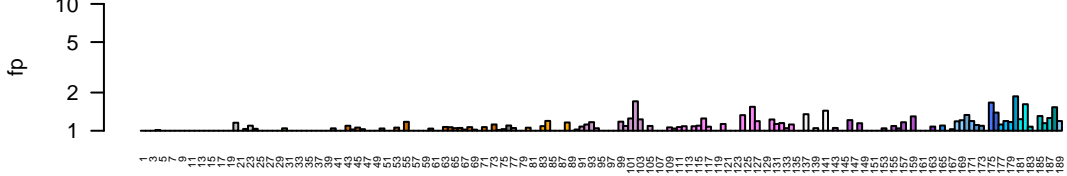




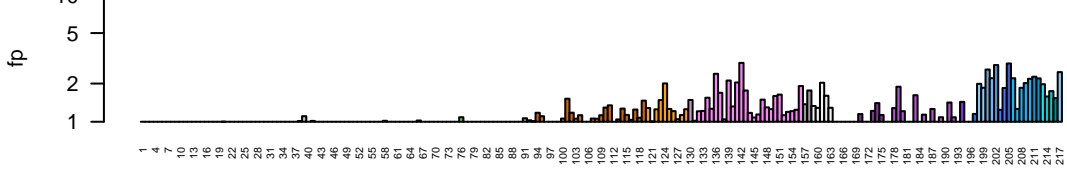




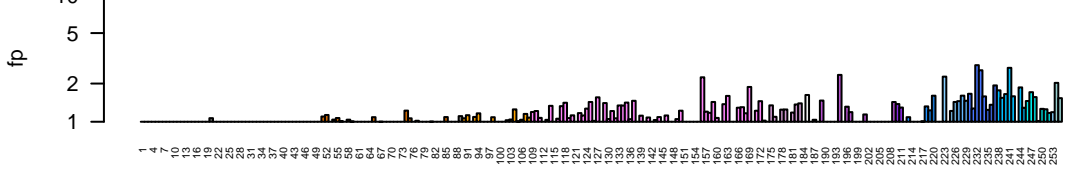
Tadh OG_5702
Tadh_TriadT19575
cyclin_dependent_kinase_like_1



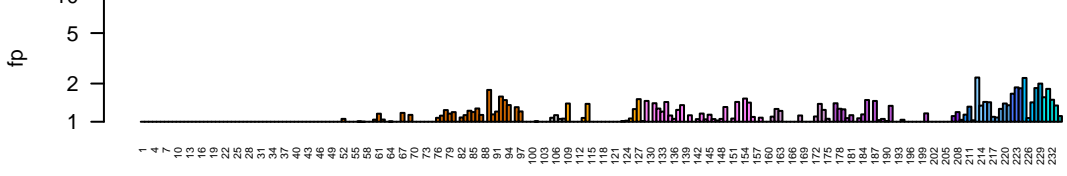
TrH2 OG_5702
TrH2_TrispH2_003126-RA
cyclin_dependent_kinase_like_1

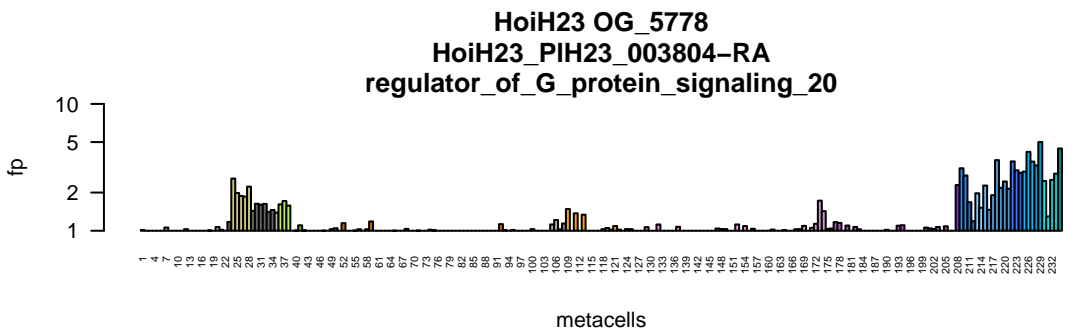
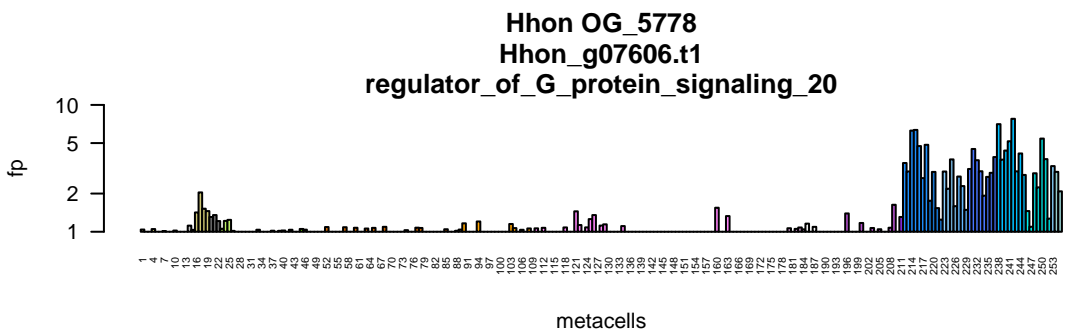
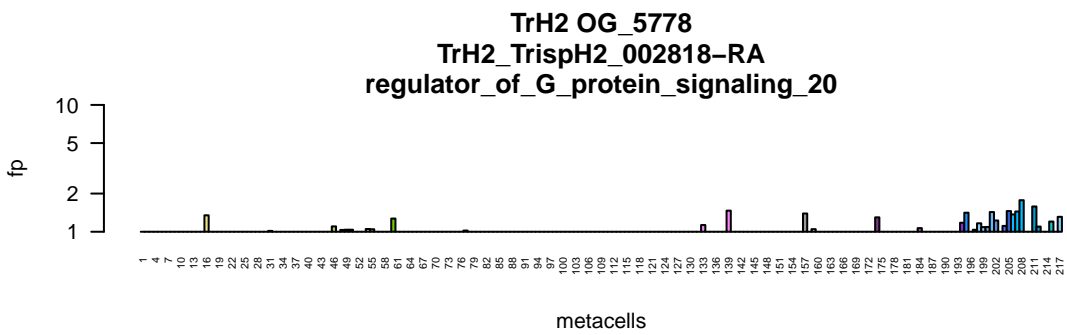
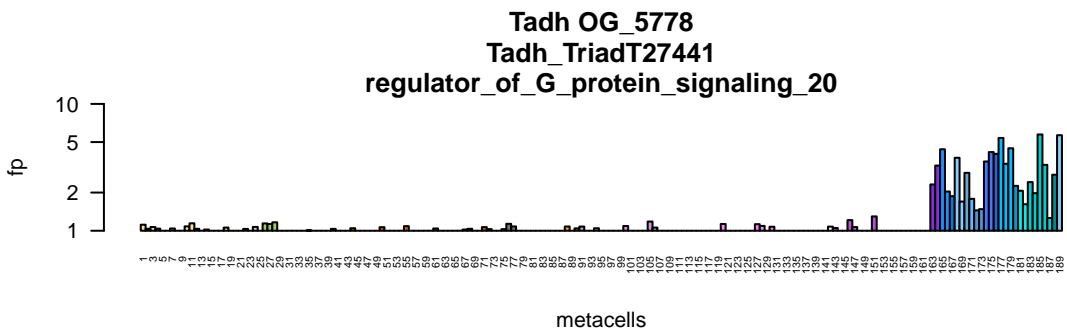


Hhon OG_5702
Hhon_g03738.t1
cyclin_dependent_kinase_like_1

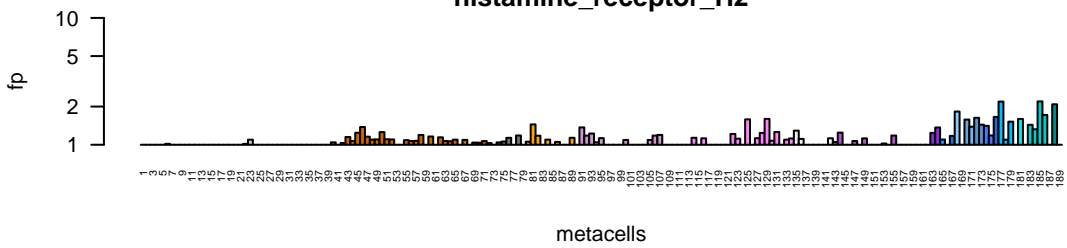


HoiH23 OG_5702
HoiH23_PIH23_009088-RA
cyclin_dependent_kinase_like_1

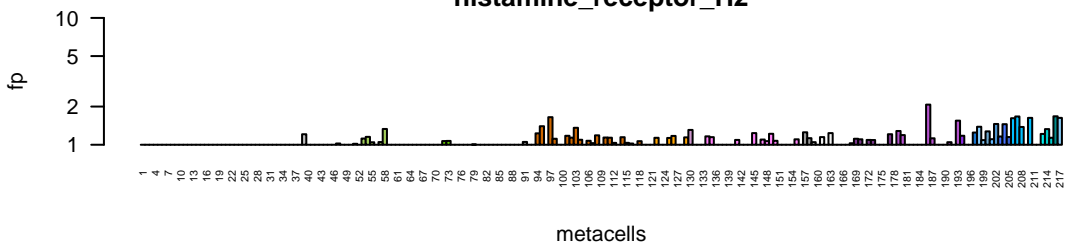




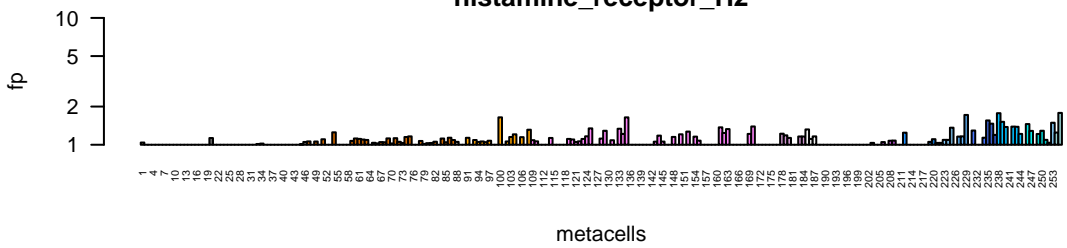
Tadh OG_6463
Tadh_TriadT60646
histamine_receptor_H2



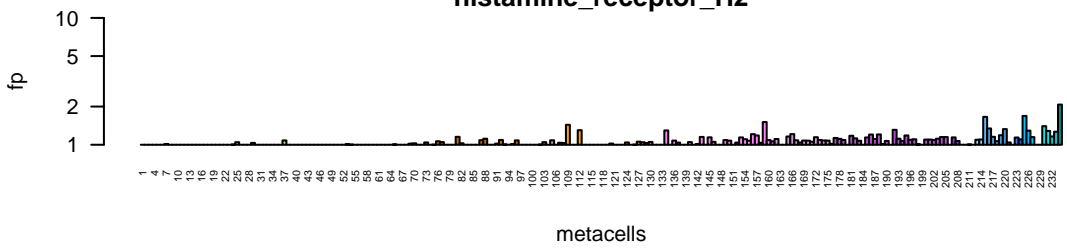
TrH2 OG_6463
TrH2_TrispH2_009668-RA
histamine_receptor_H2

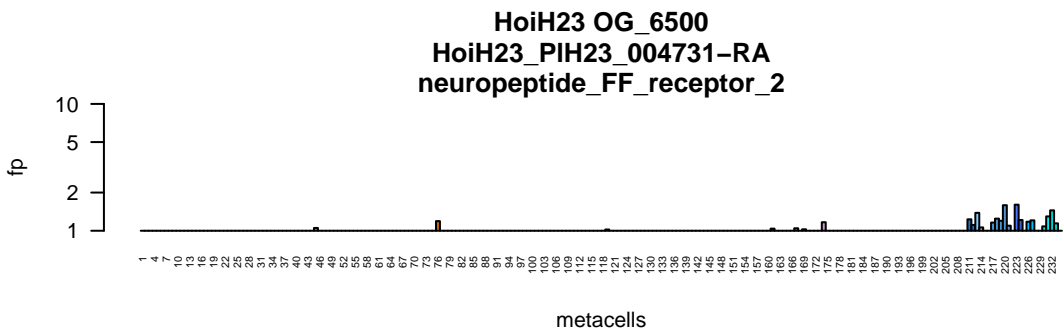
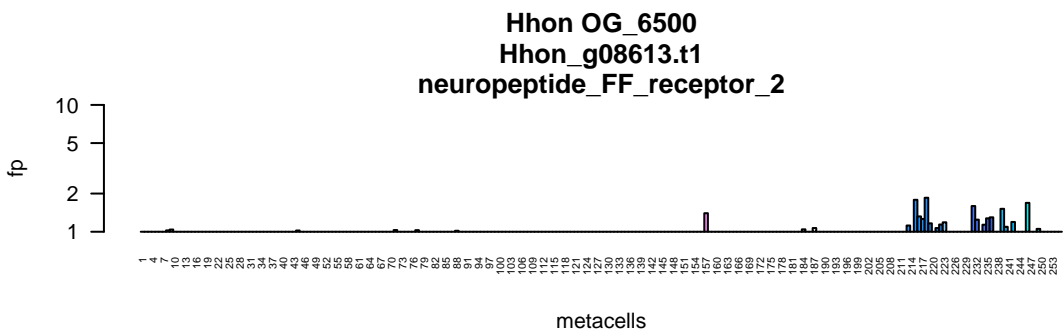
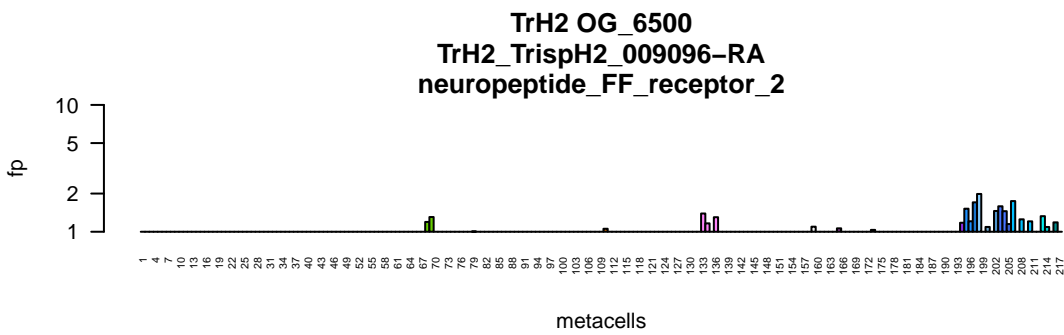
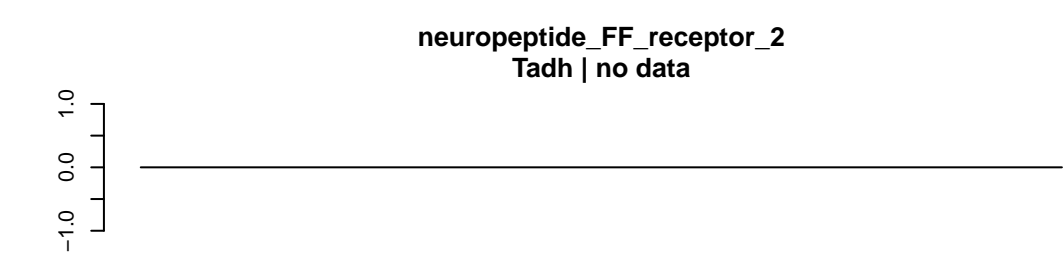


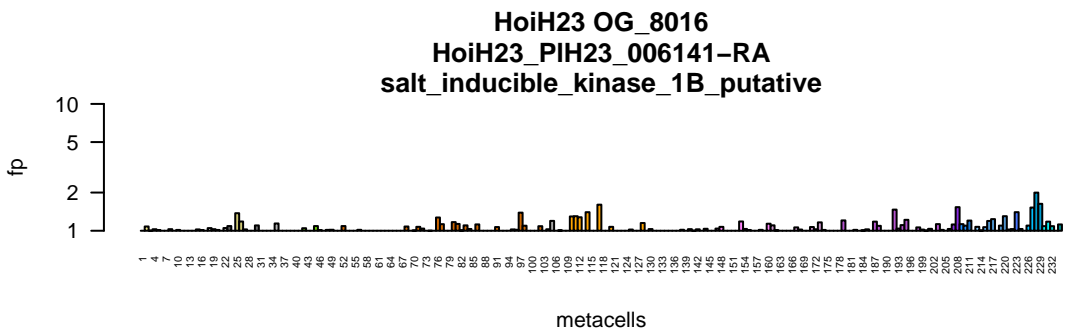
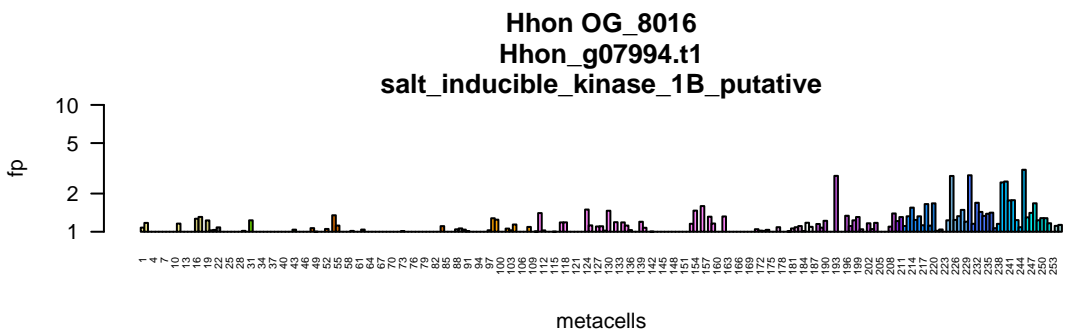
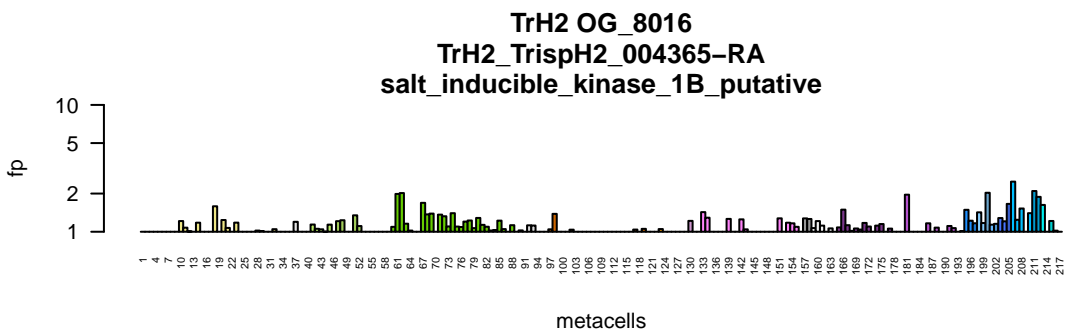
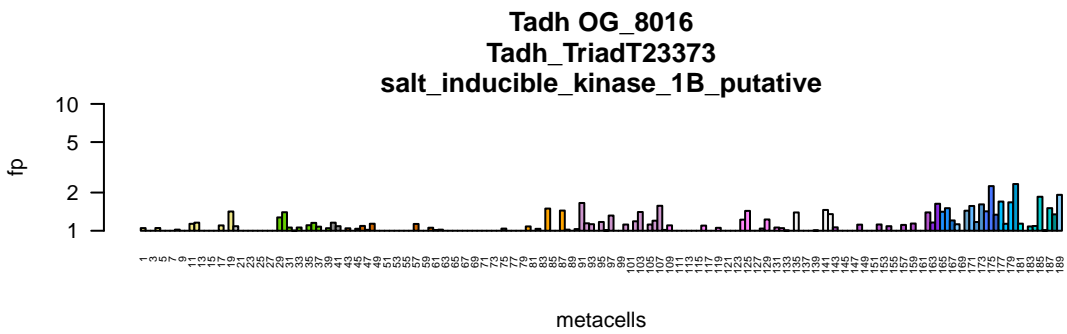
Hhon OG_6463
Hhon_g08668.t1
histamine_receptor_H2

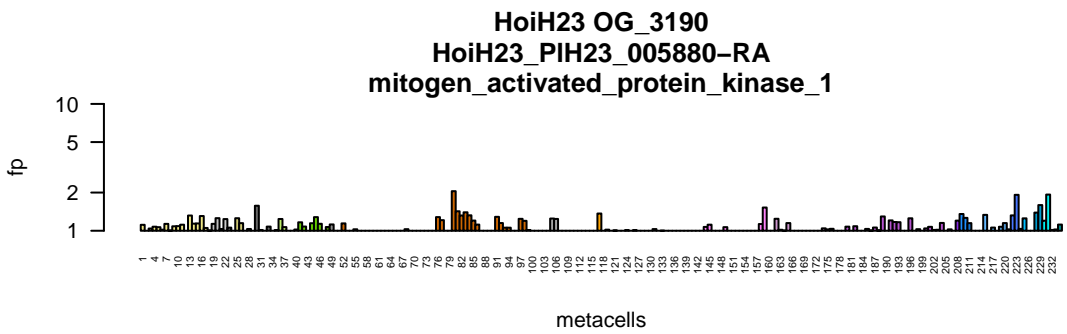
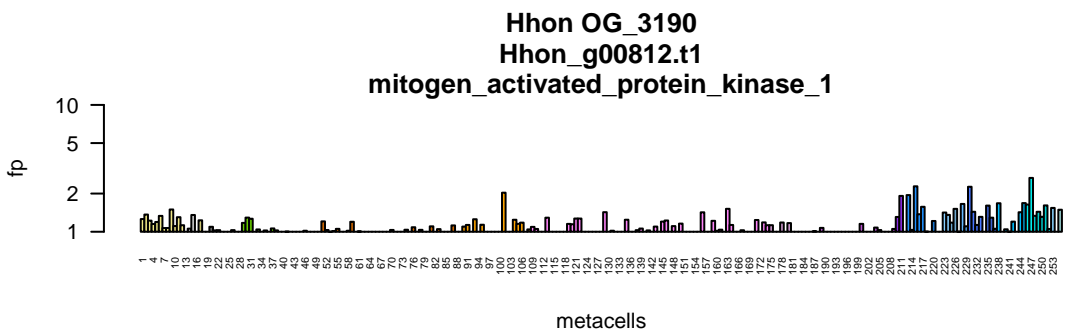
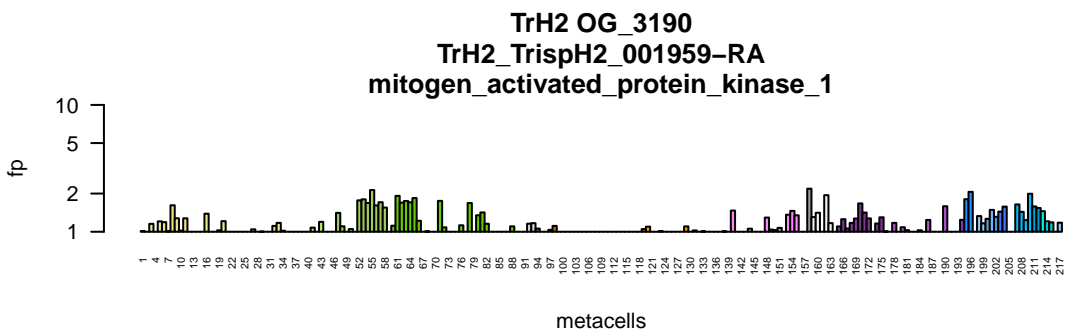
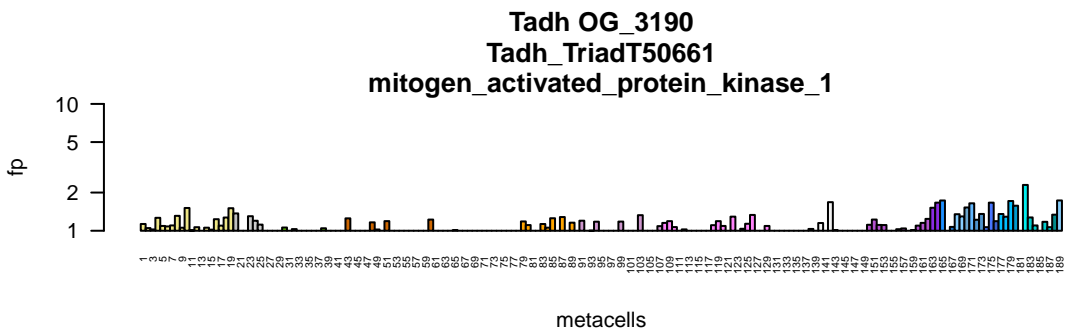


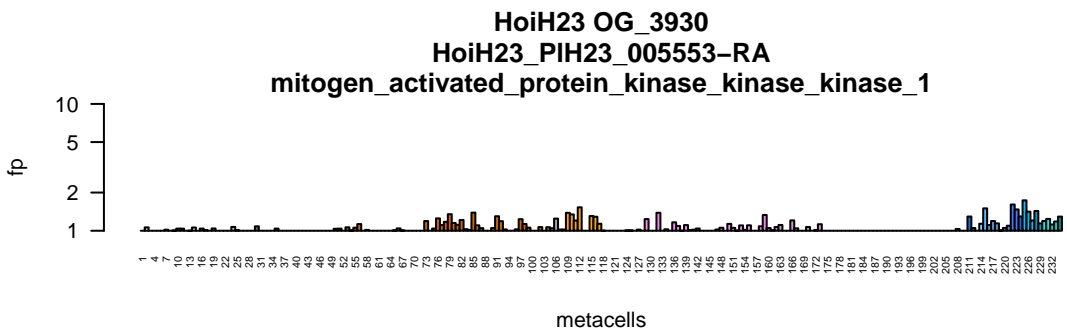
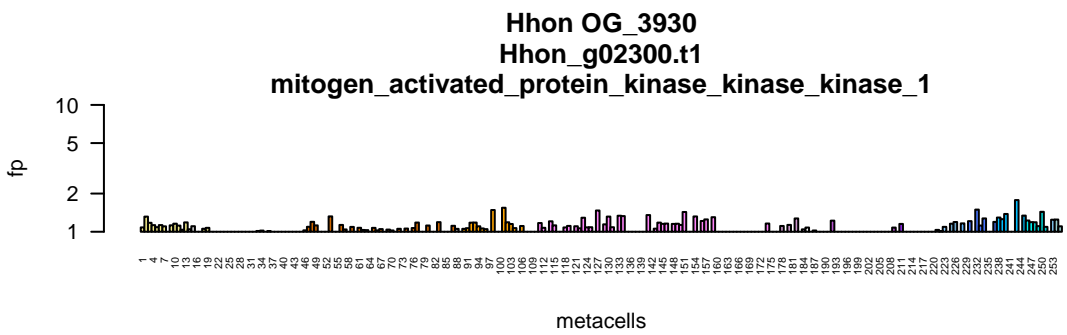
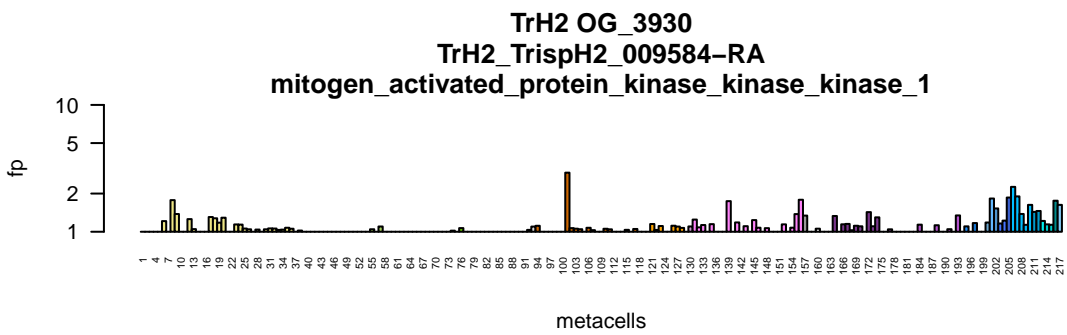
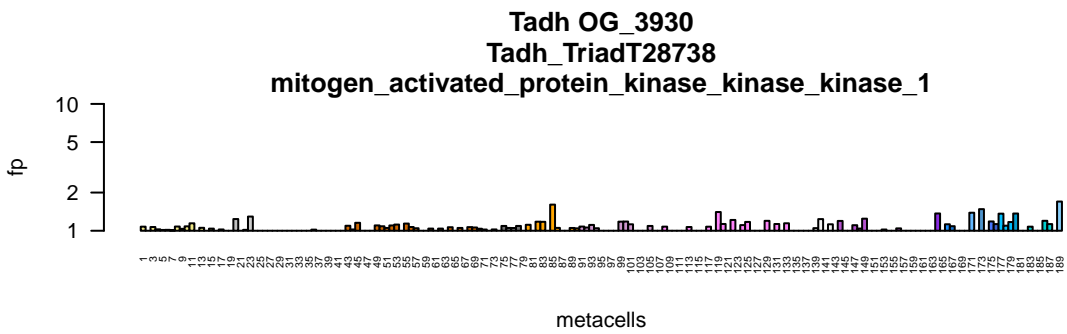
HoiH23 OG_6463
HoiH23_PIH23_005854-RA
histamine_receptor_H2

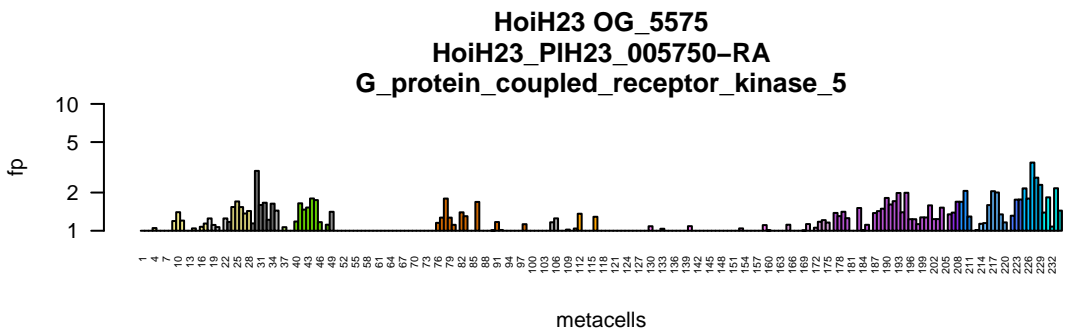
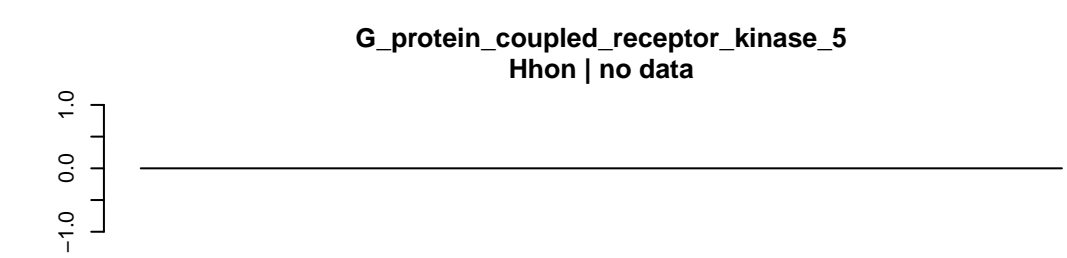
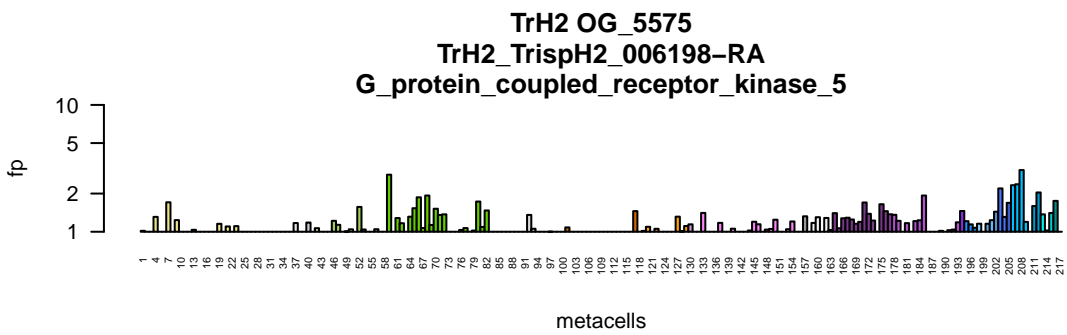
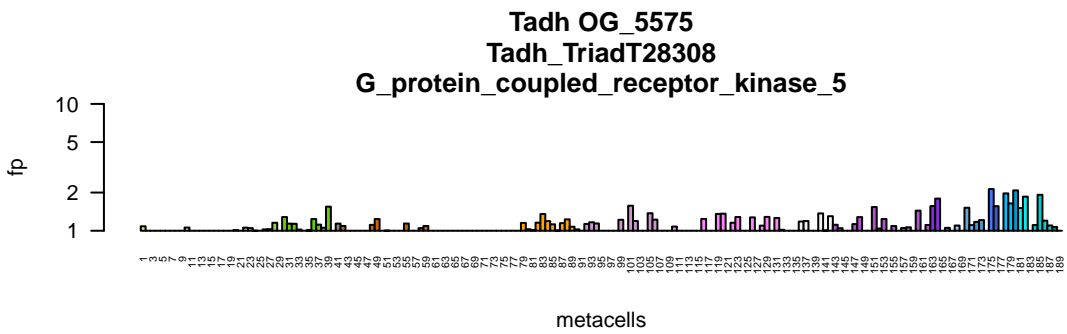


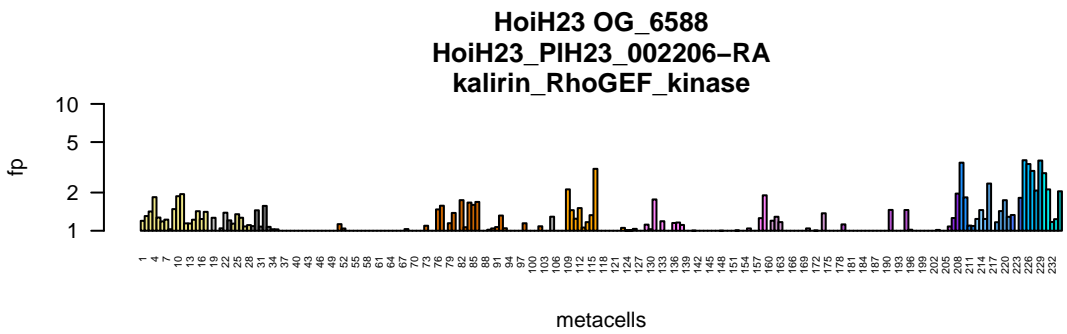
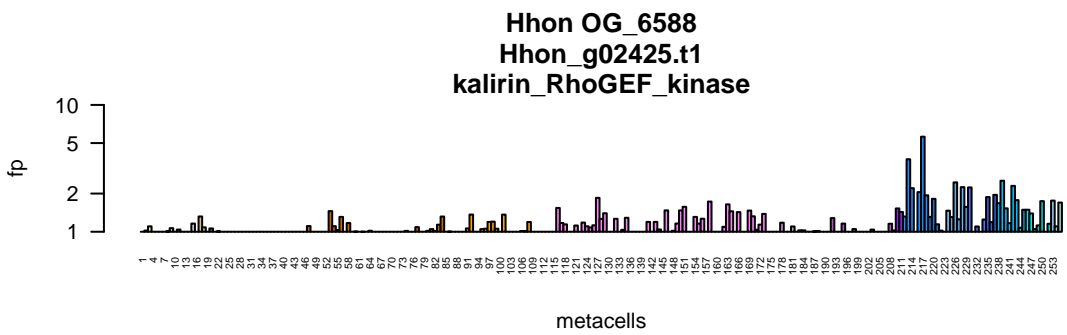
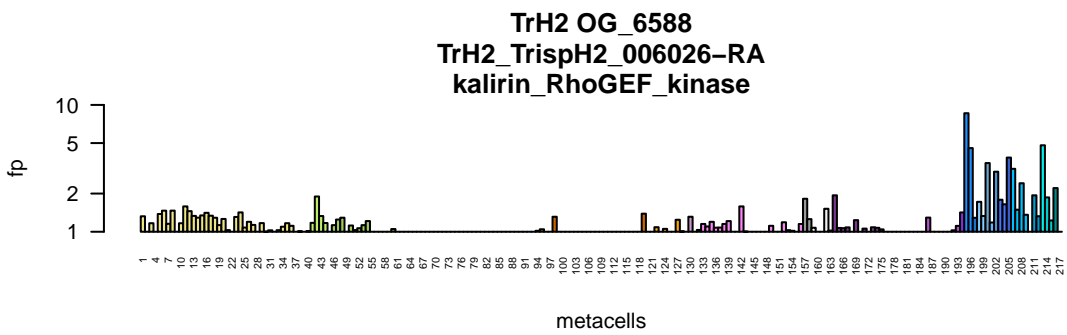
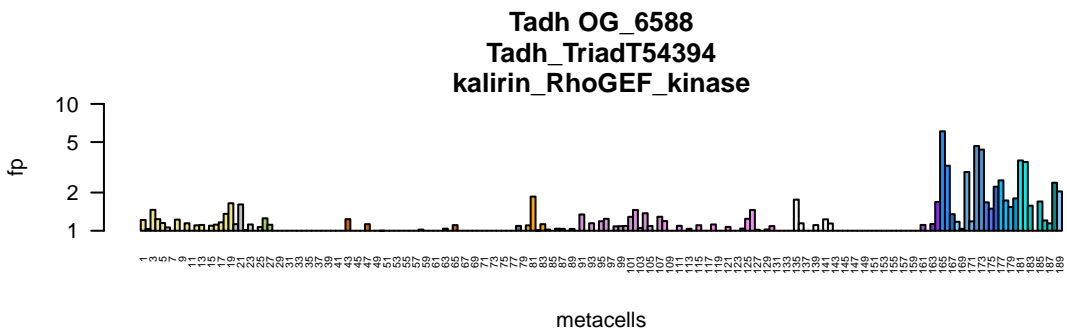


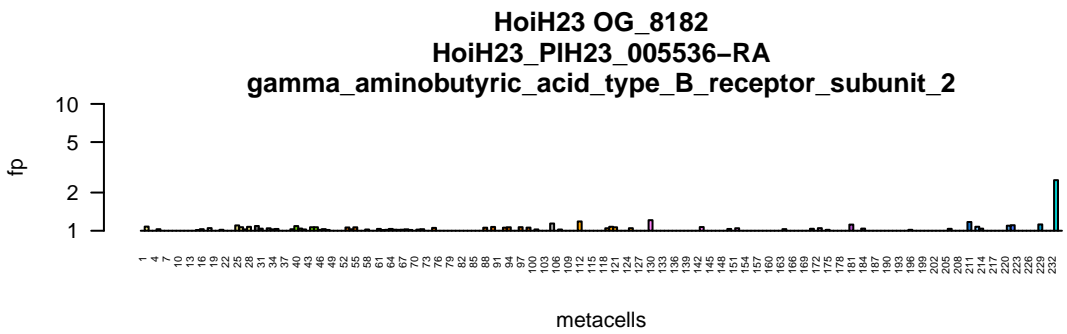
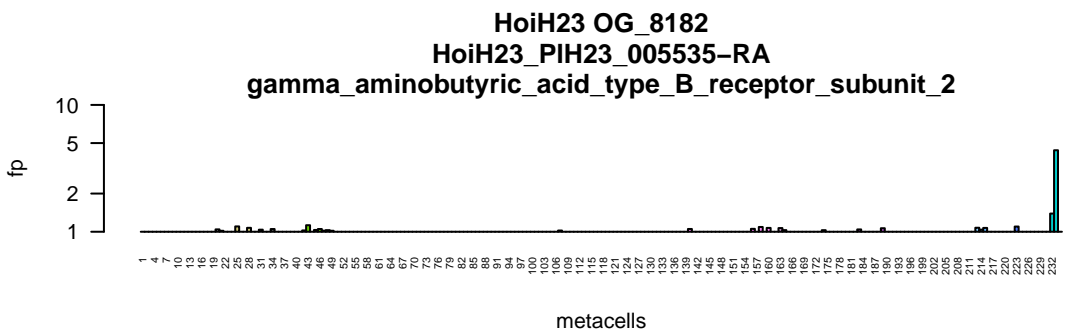
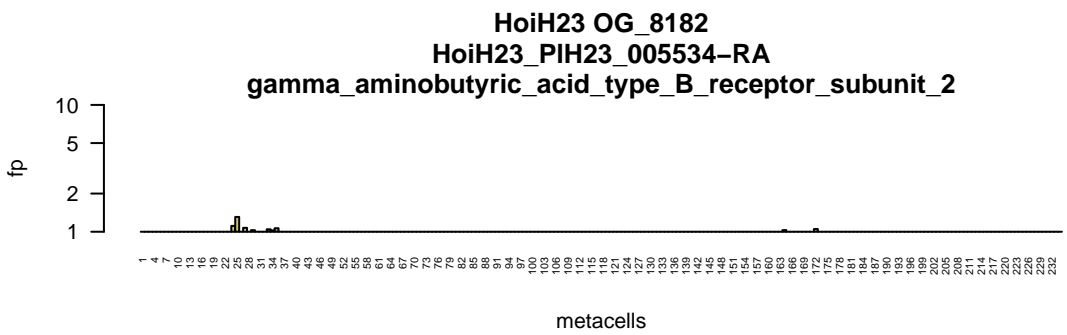
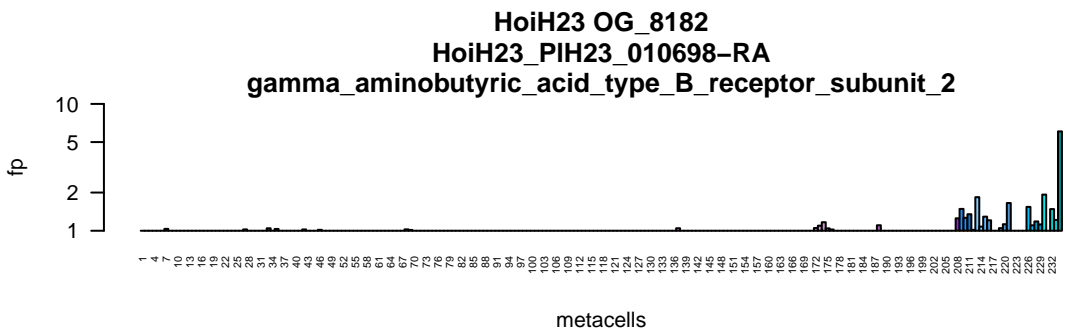
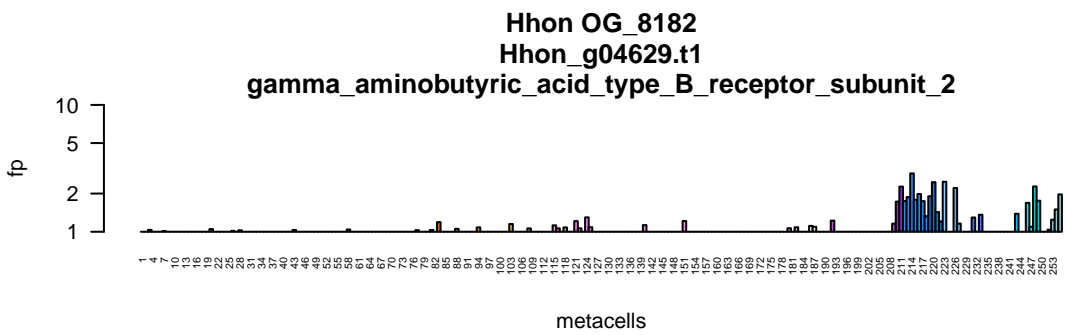
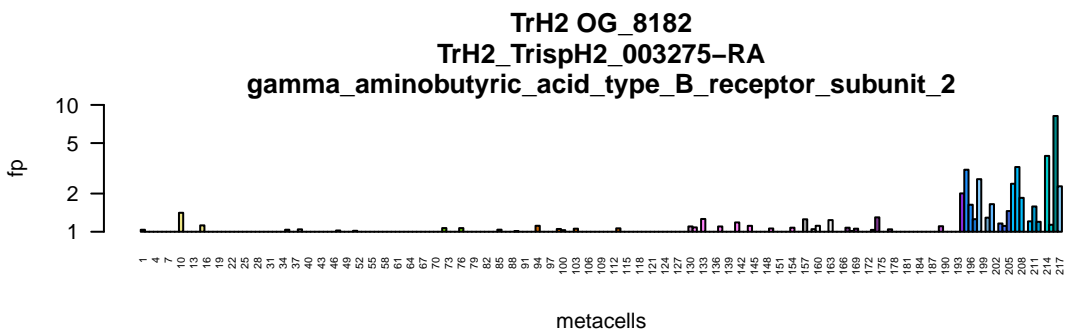
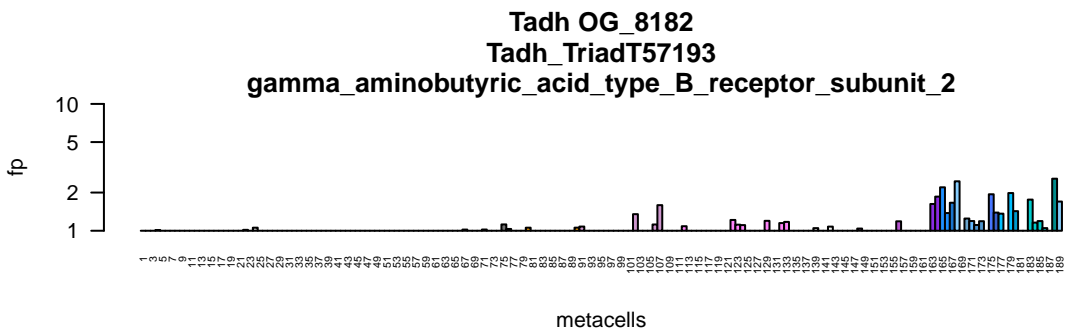






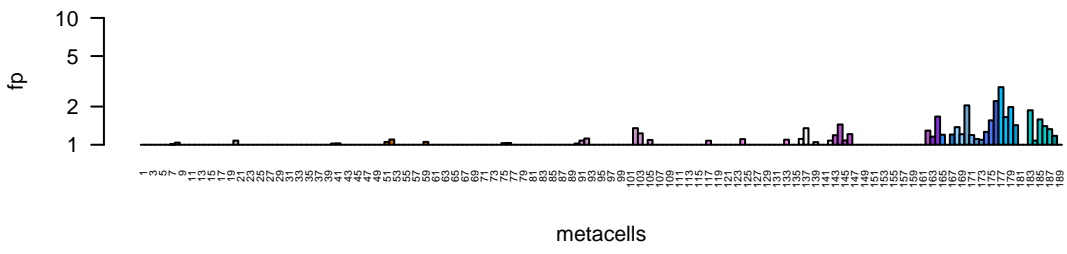






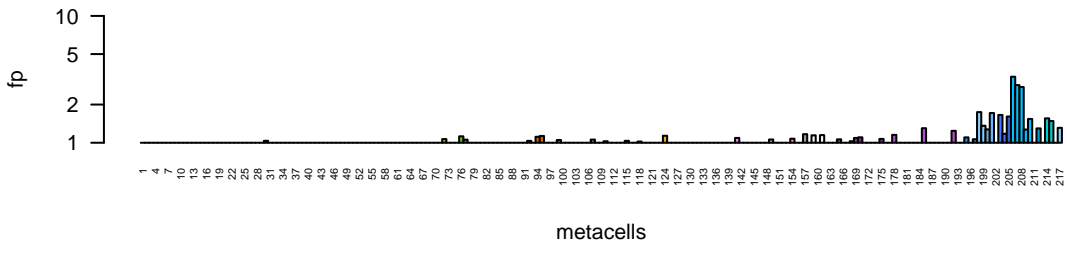
Tadh OG_8295

Tadh_TriadT54897



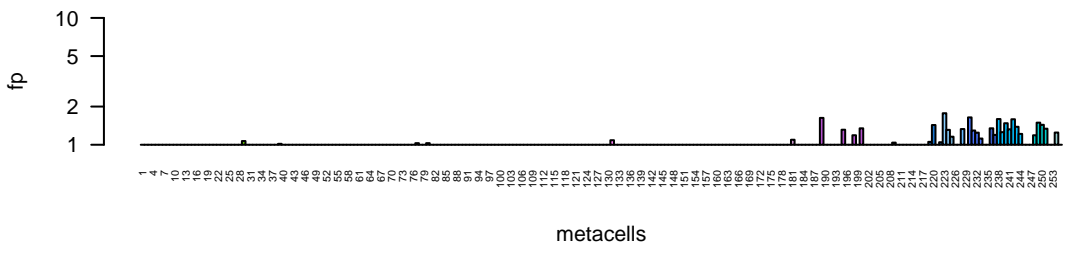
TrH2 OG_8295

TrH2_TrispH2_007497-RA



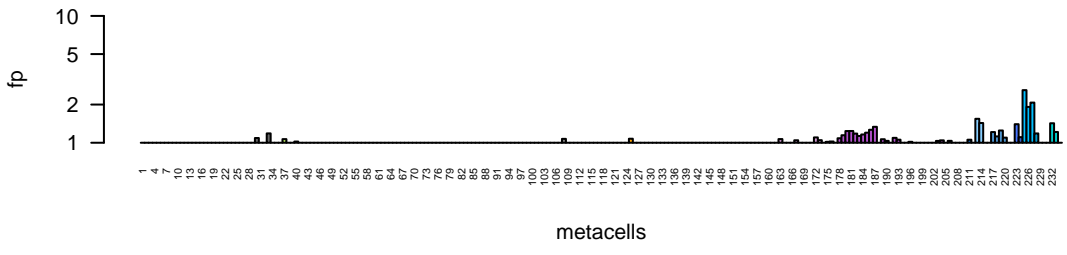
Hhon OG_8295

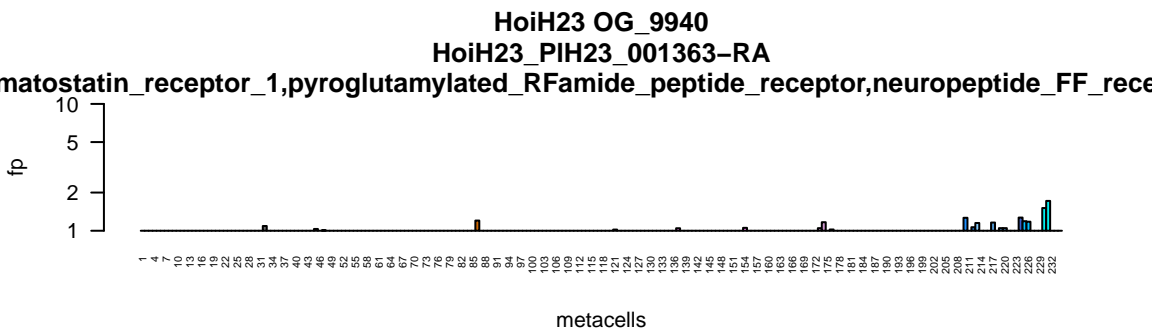
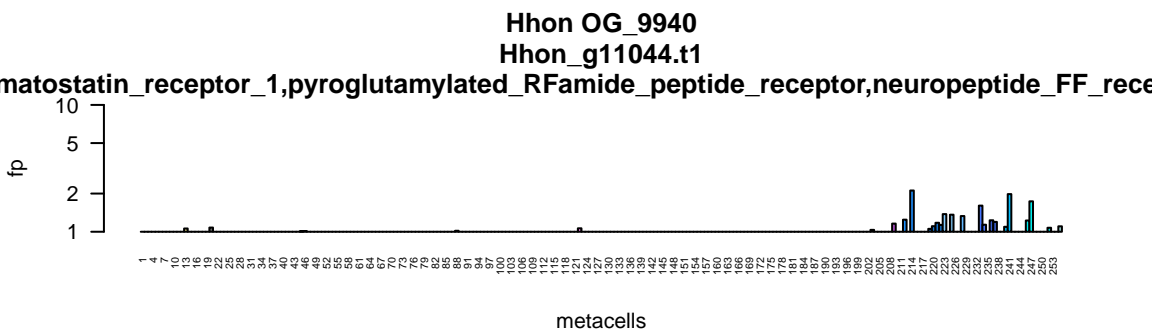
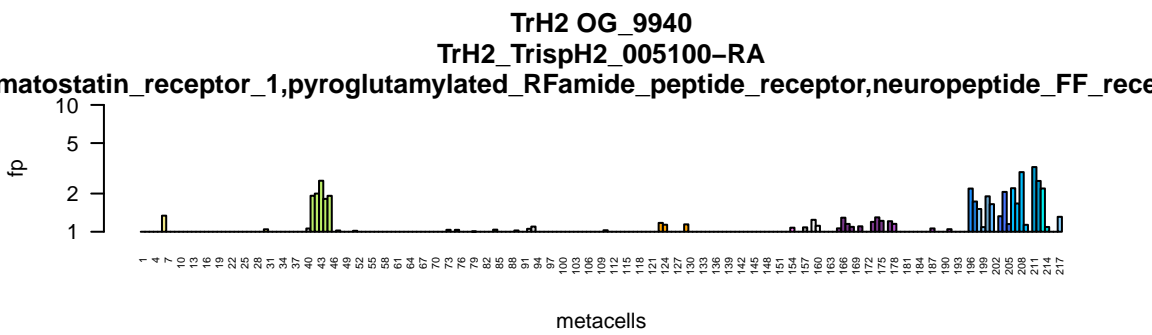
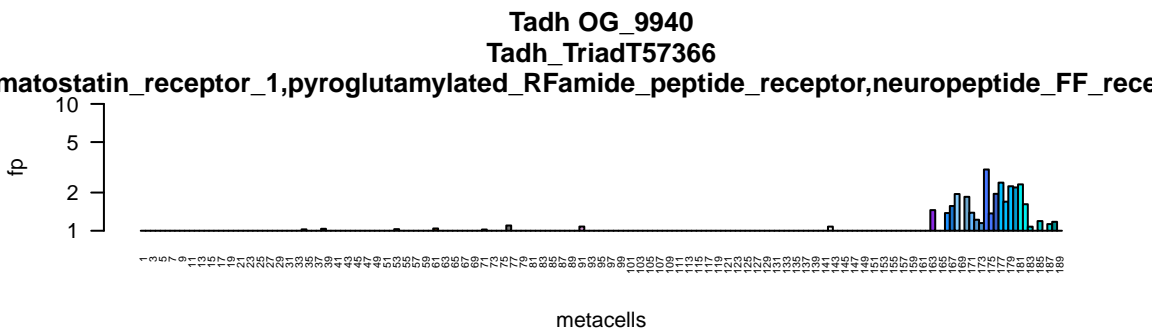
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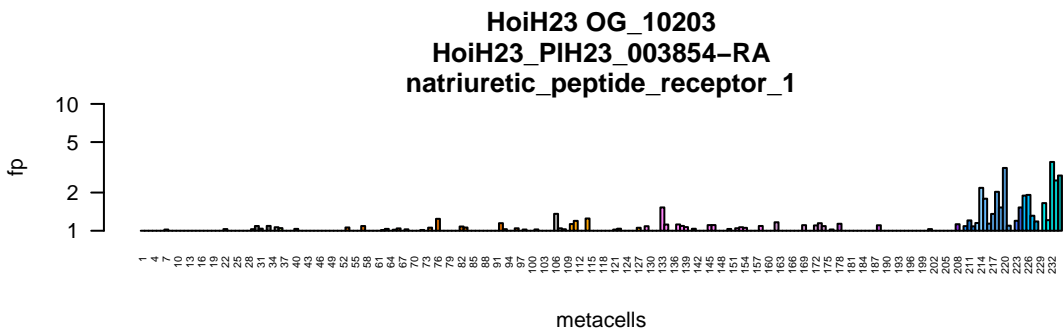
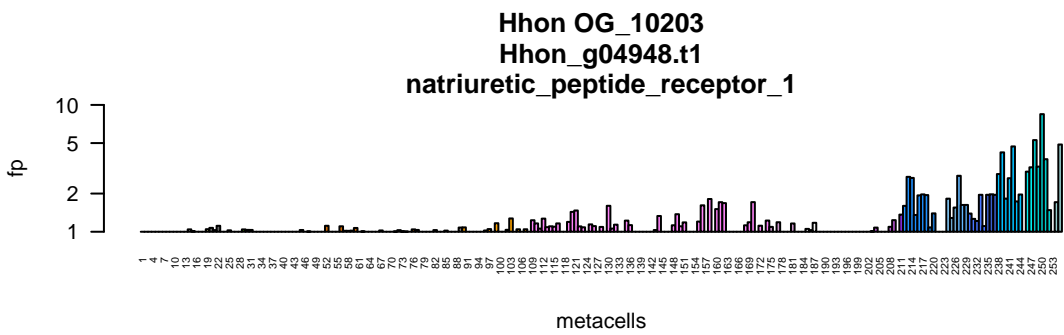
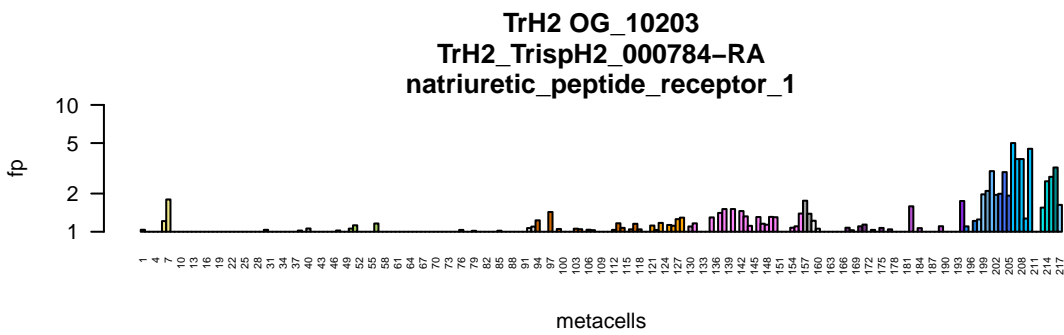
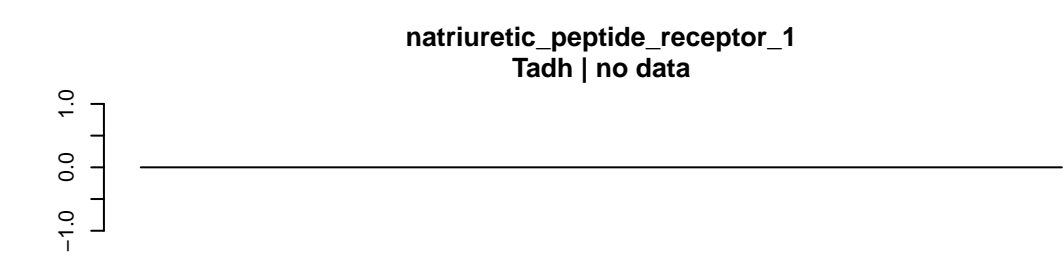


HoiH23 OG_8295

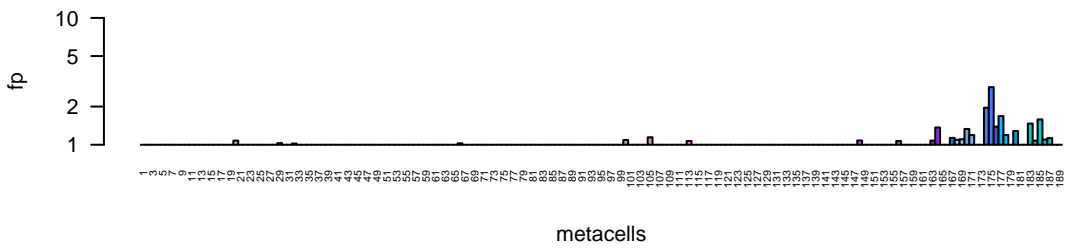
HoiH23_PIH23_002701-RA



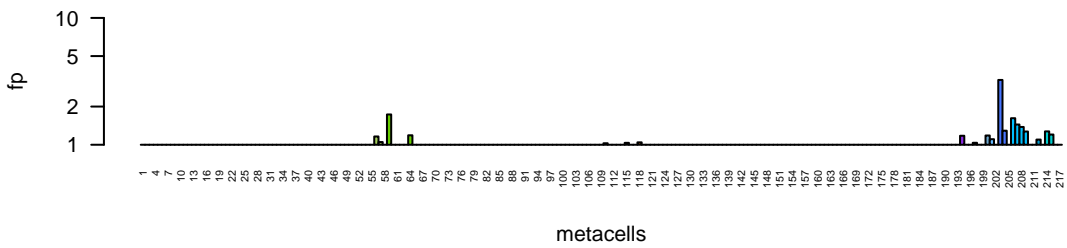




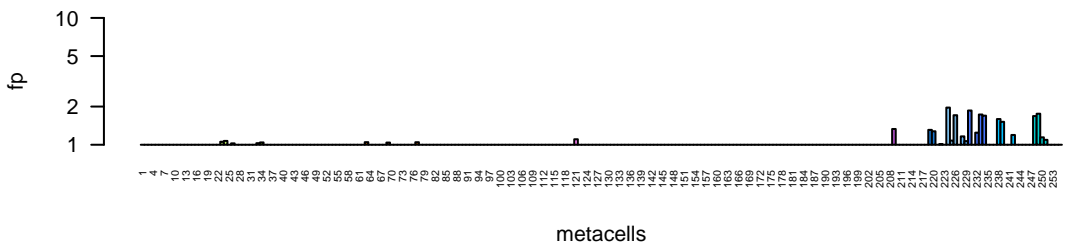
Tadh OG_515
Tadh_TriadT57467



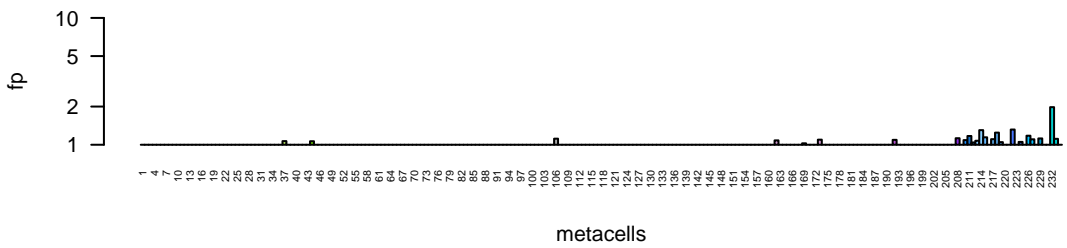
TrH2 OG_515
TrH2_TrispH2_007315-RA

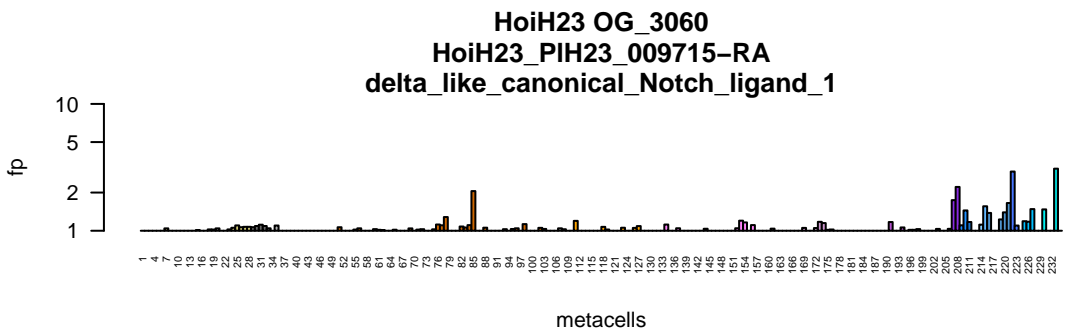
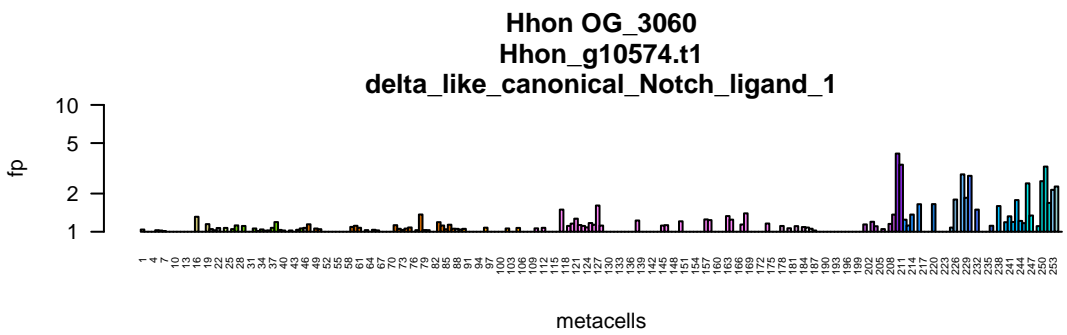
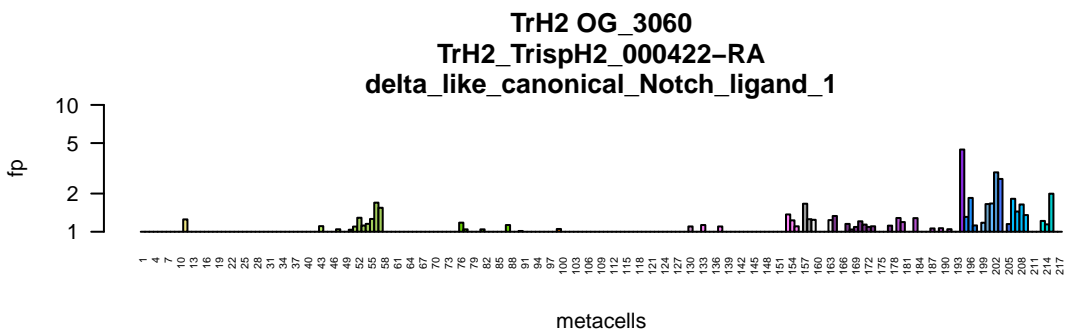
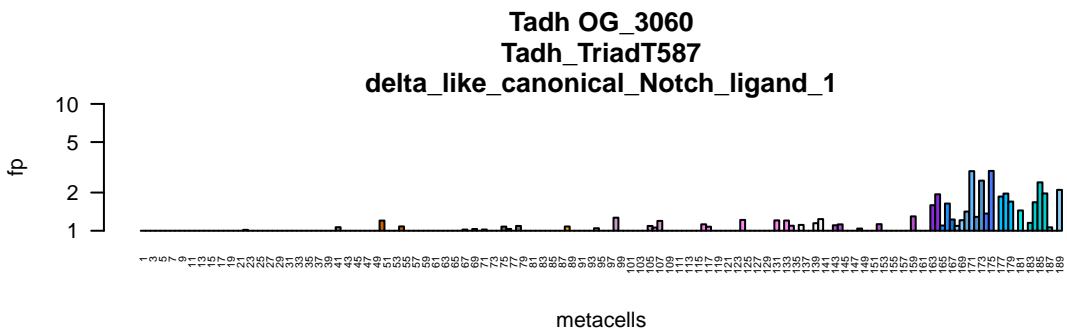


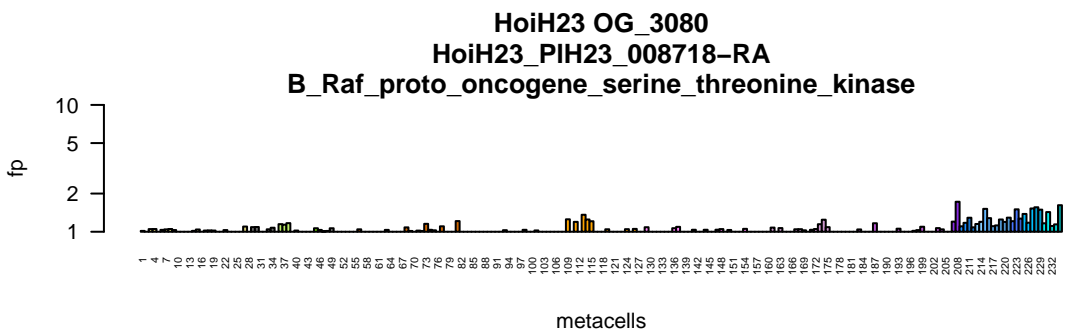
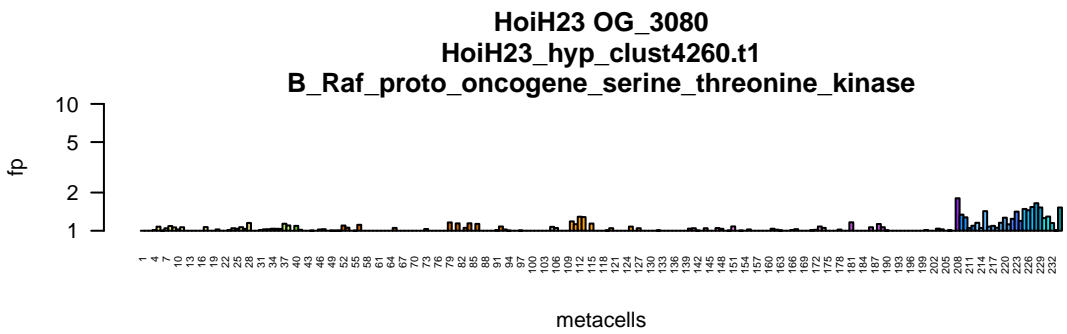
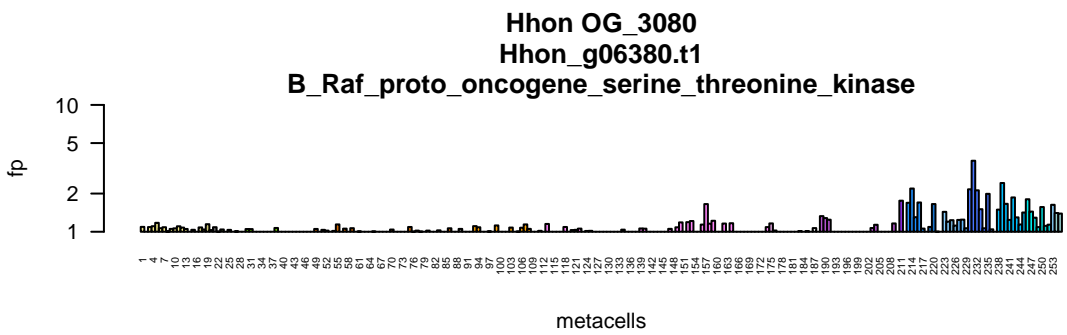
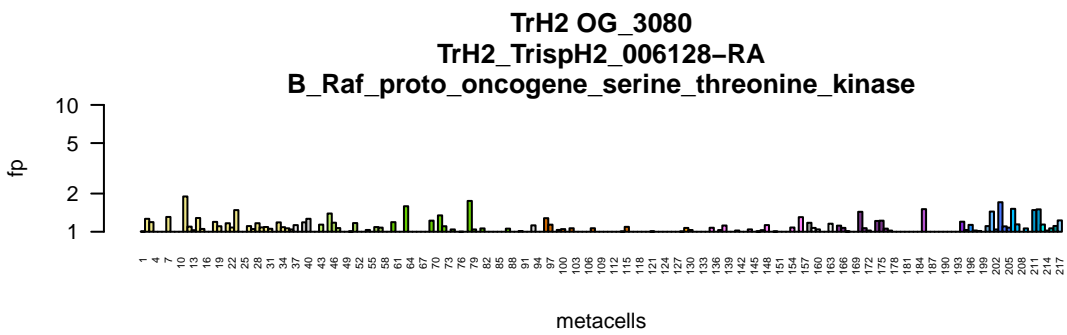
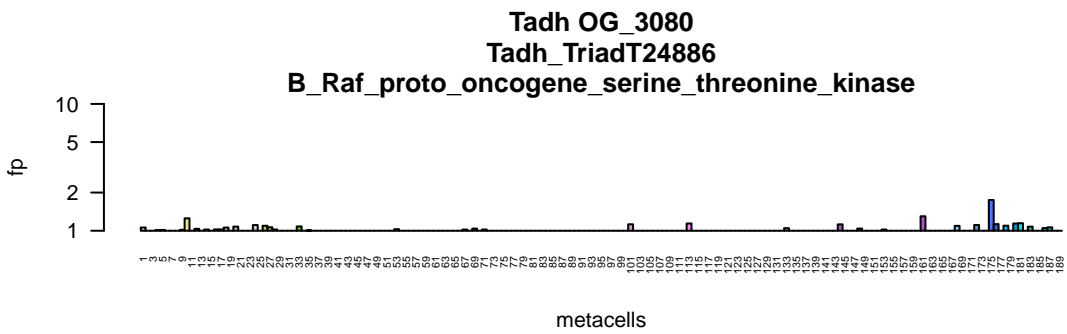
Hhon OG_515
Hhon_g01896.t1

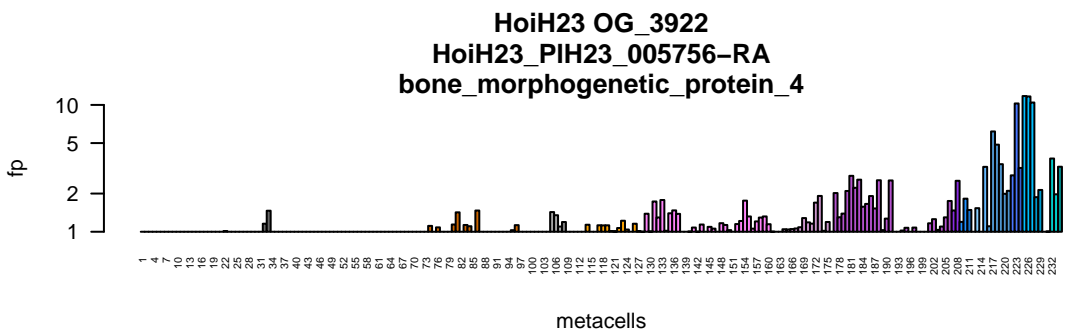
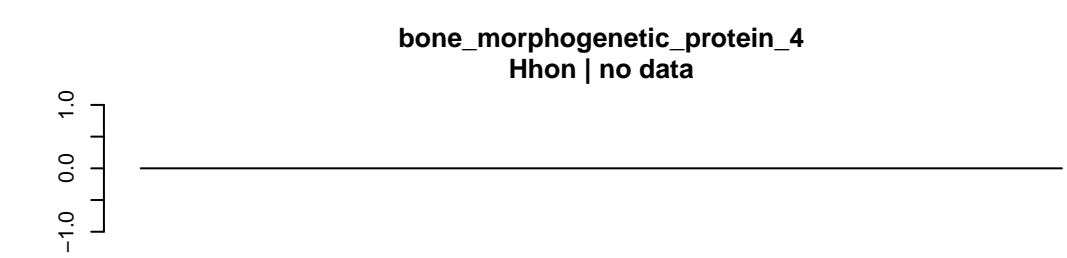
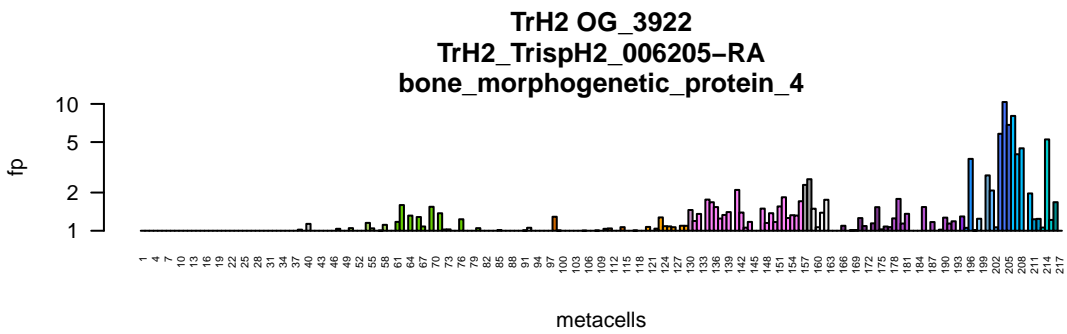
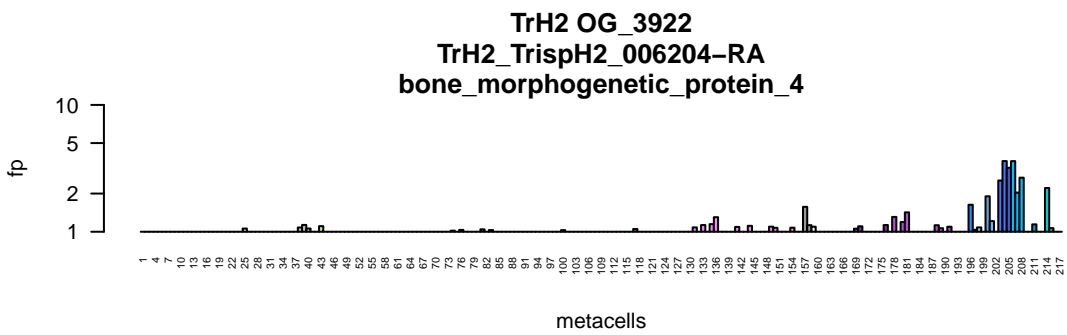
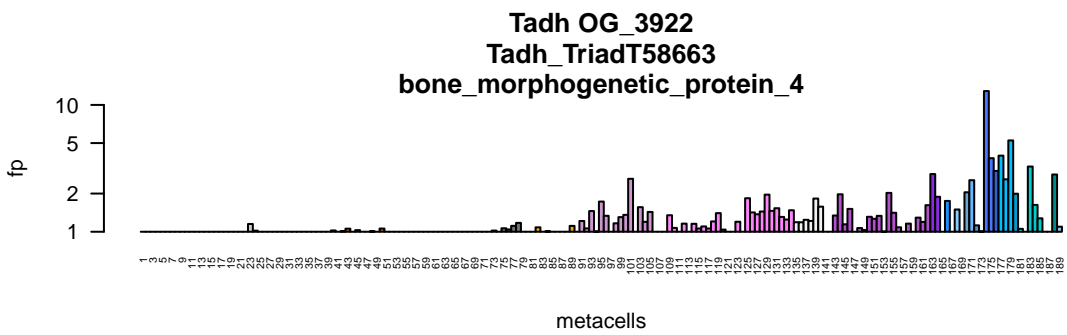
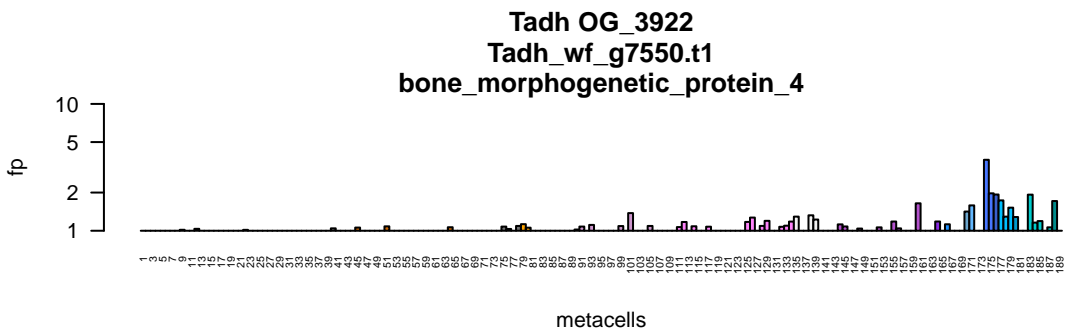


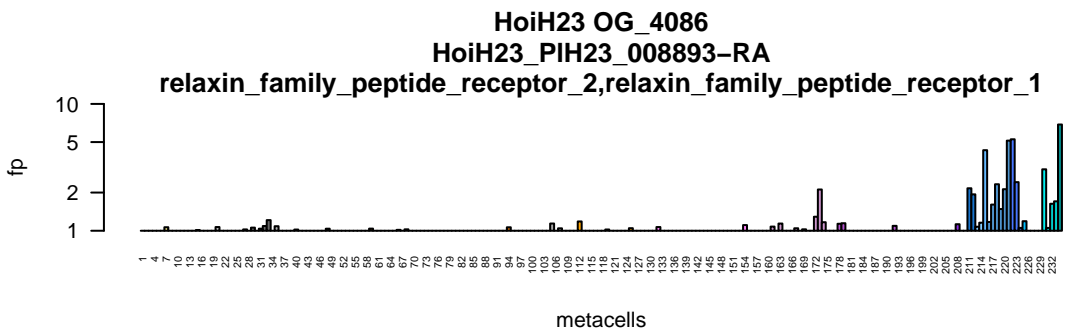
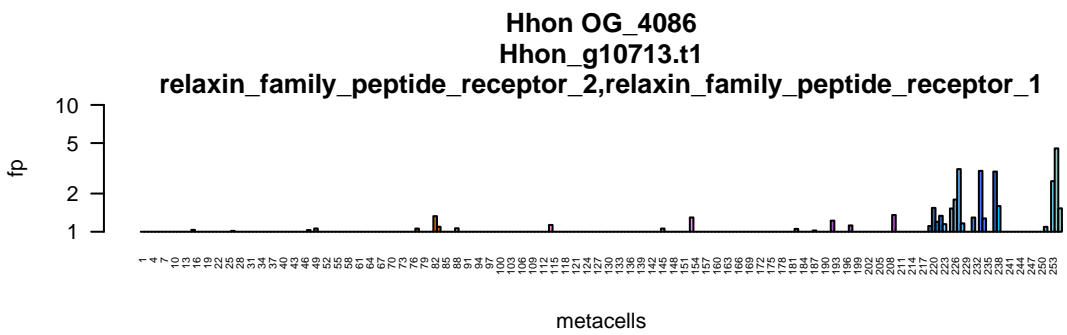
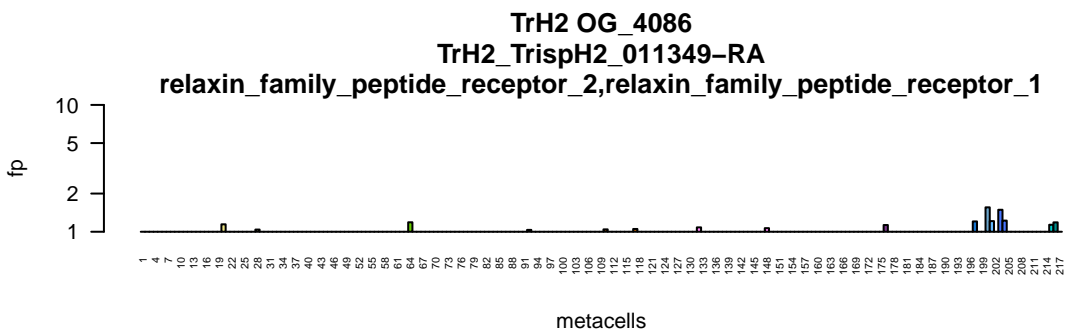
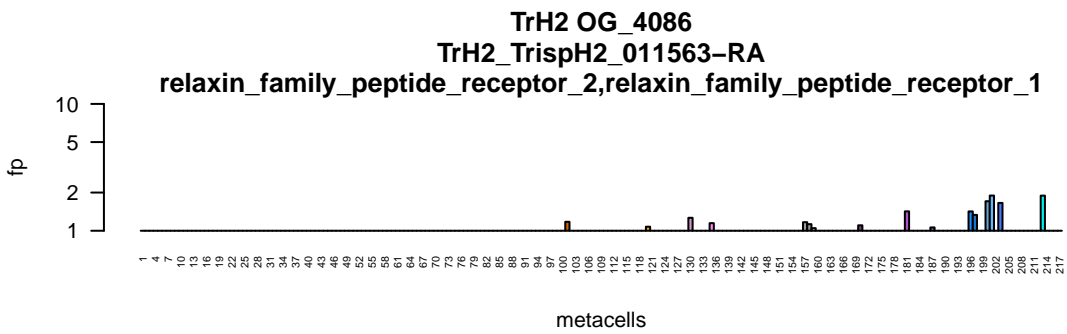
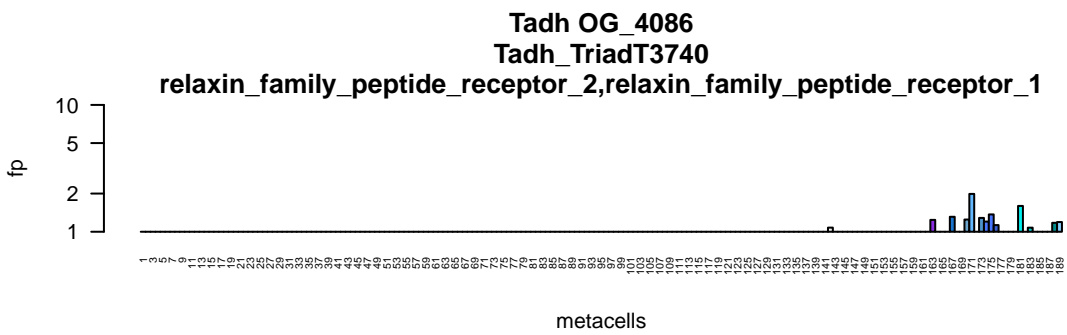
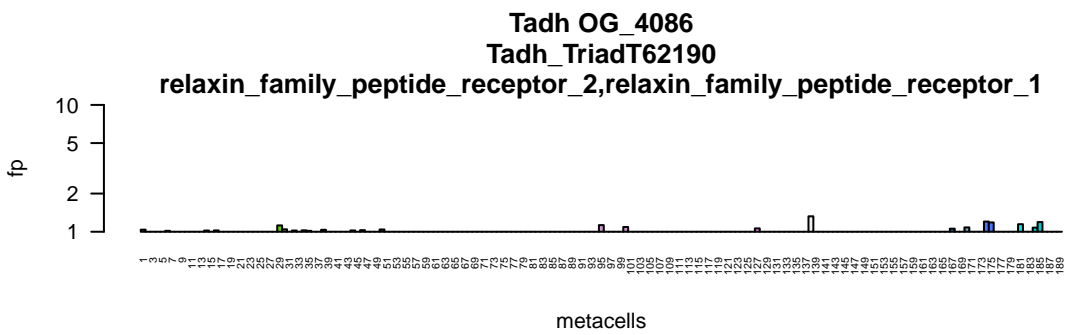
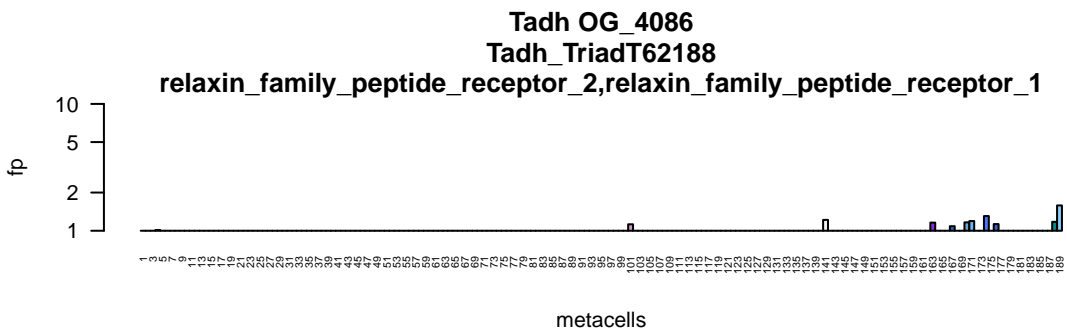
HoiH23 OG_515
HoiH23_PIH23_001462-RA

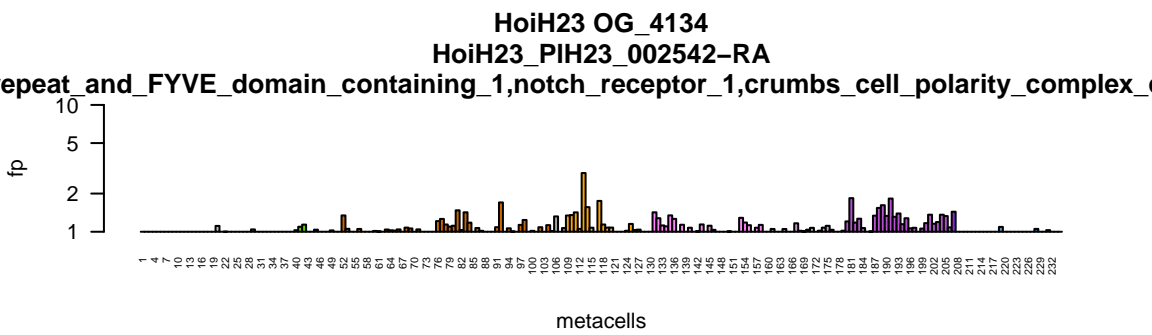
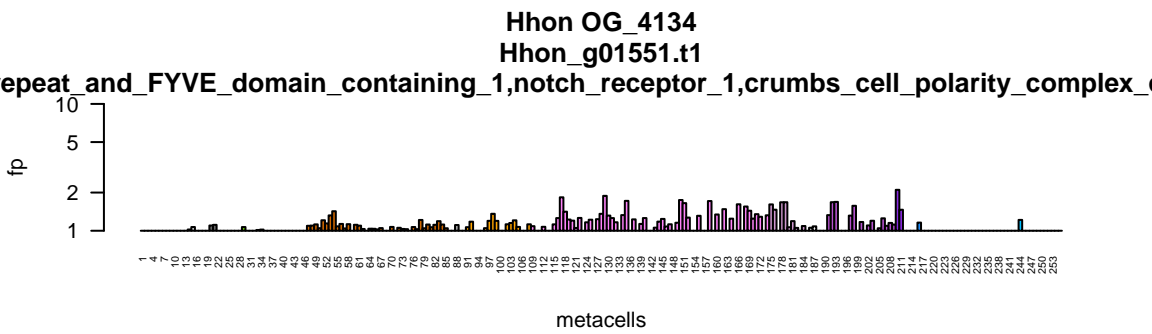
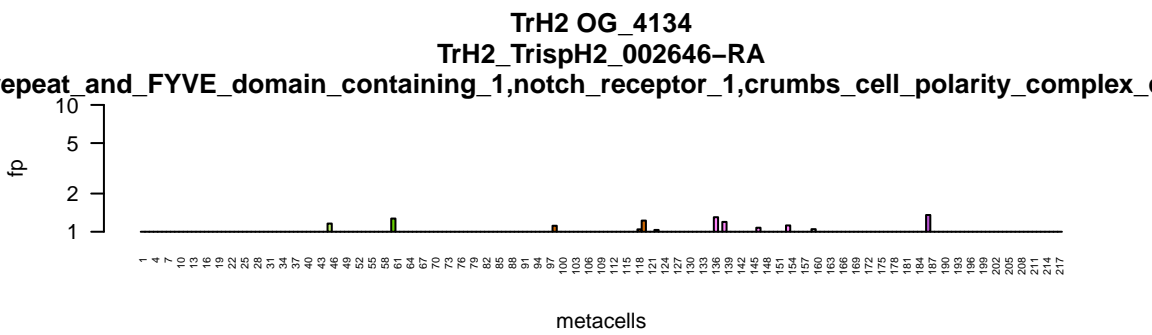
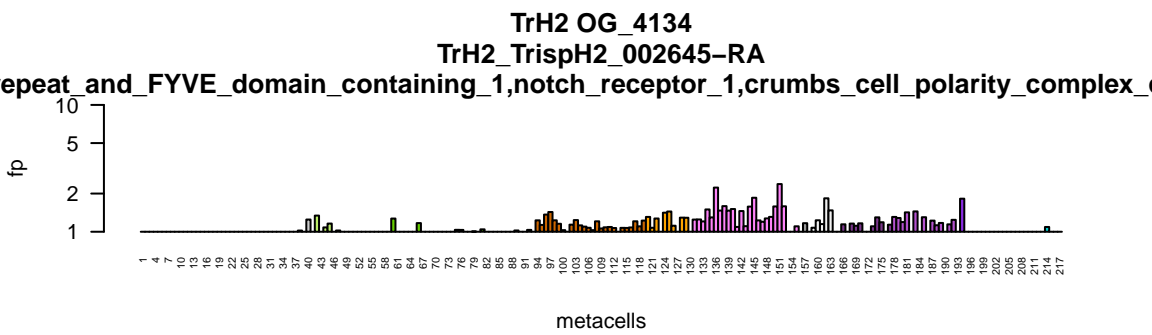
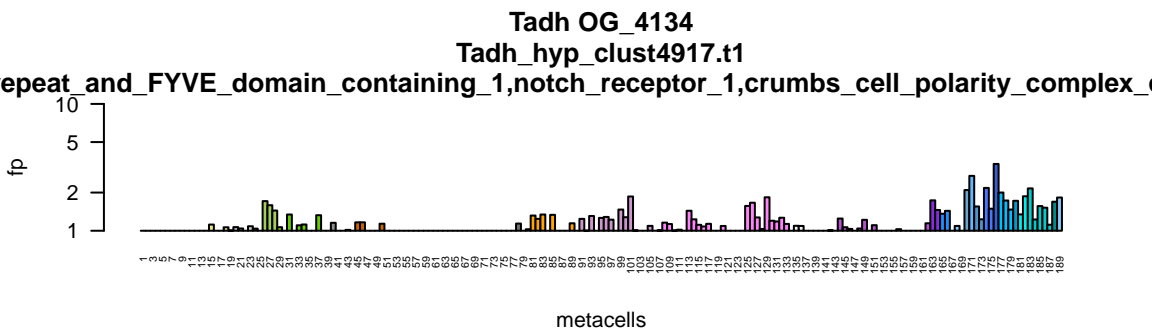




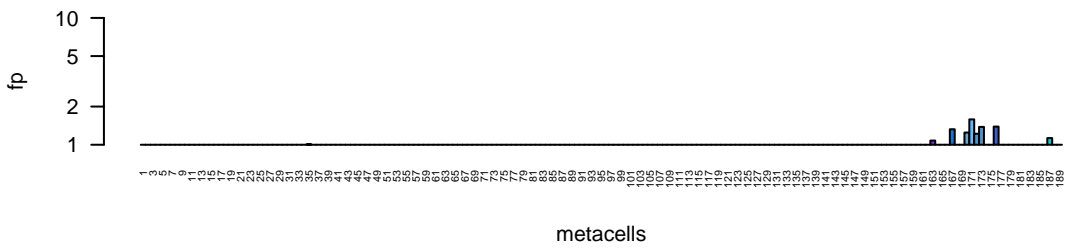




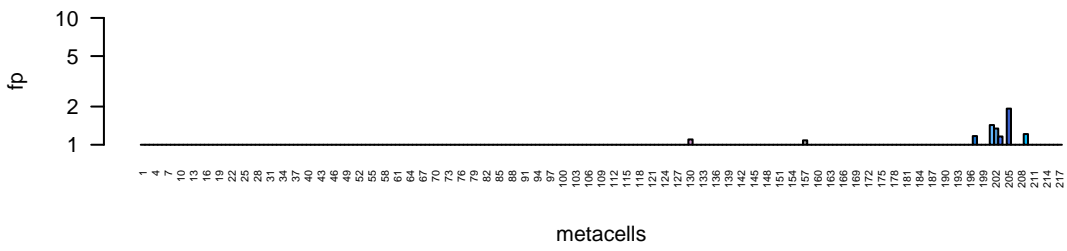




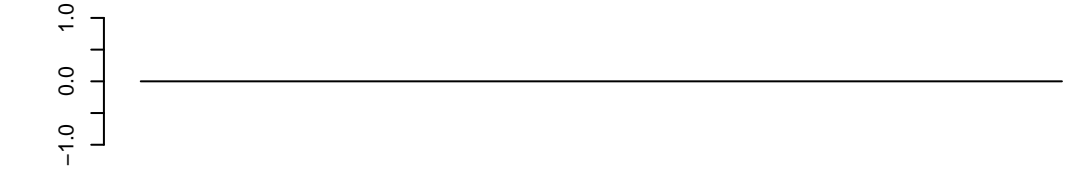
Tadh OG_4270
Tadh_wf_g4875.t1



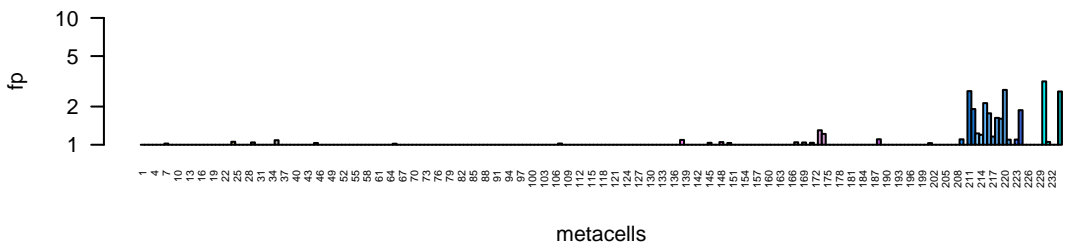
TrH2 OG_4270
TrH2_TrispH2_008489-RA



Hhon | no data



HoiH23 OG_4270
HoiH23_PIH23_009325-RA



metacells

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 0 false positives, indicated by a horizontal line at fp=0. A few metacells have 1 false positive (e.g., 46, 52, 97, 127, 145, 154, 160, 175, 190, 214), and a small group at the end (metacells 199-217) has 2 false positives.

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
157	0
160	1
163	0
166	0
169	0
172	0
175	0
178	0
181	0
184	0
187	0
190	0
193	0
196	1
199	2
200	2
201	1
202	1
203	0
204	0
207	0
210	0
211	0
214	1
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

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 40 metacells. Most metacells have 1 false positive, but metacells 202 and 214 have 2 false positives.

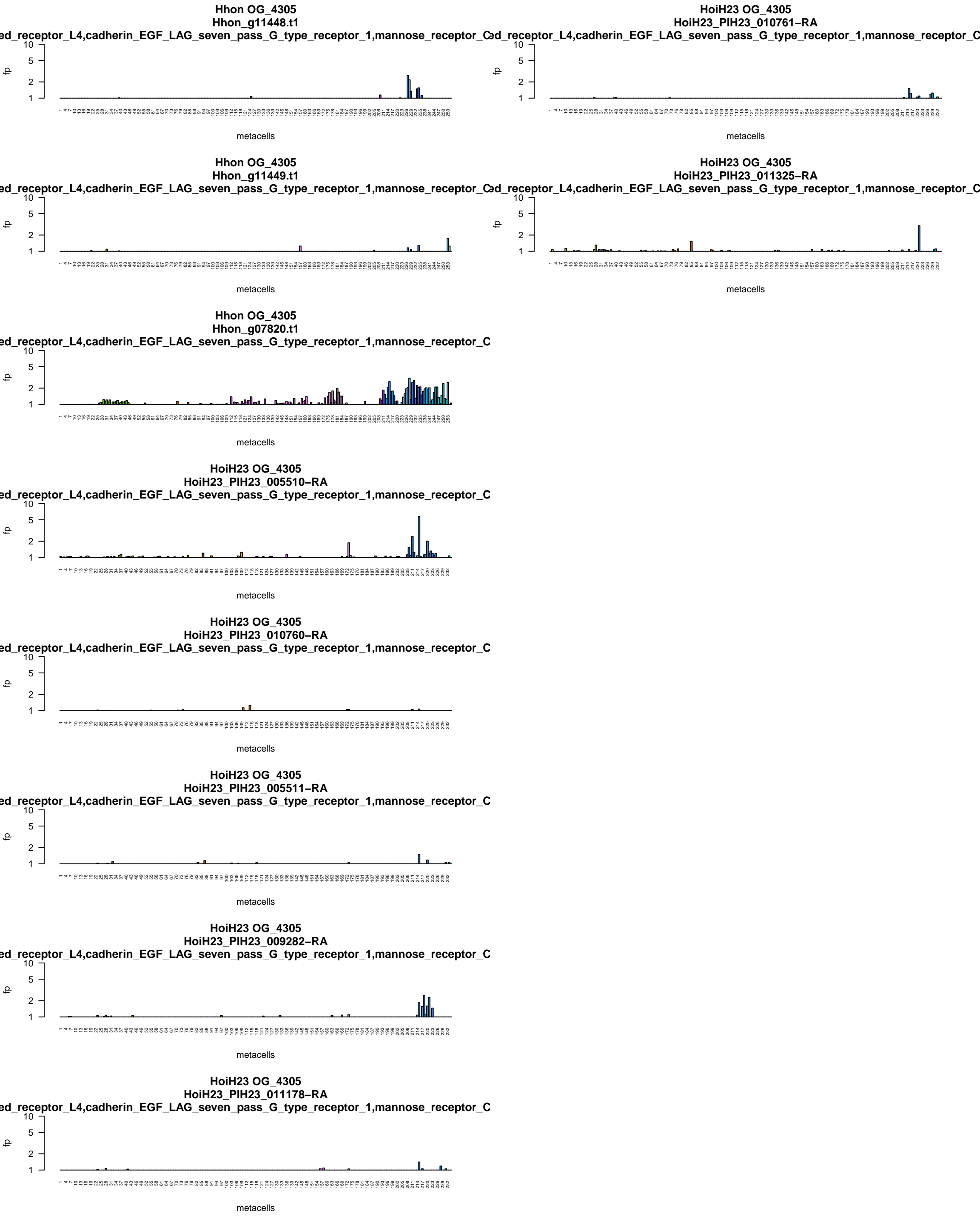
metacells	fp
1	1
4	1
7	1
10	1
13	1
16	1
19	1
22	1
25	1
28	1
31	1
34	1
37	1
40	1
43	1
46	1
49	1
52	1
55	1
58	1
61	1
64	1
67	1
70	1
73	1
76	1
79	1
82	1
85	1
88	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
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

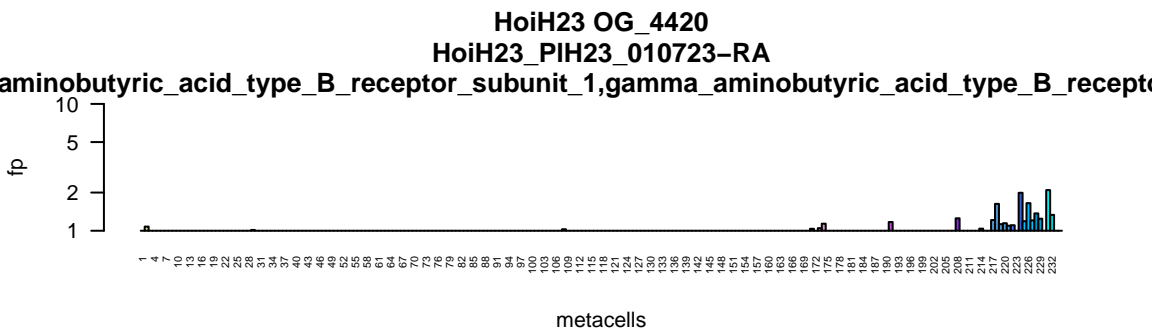
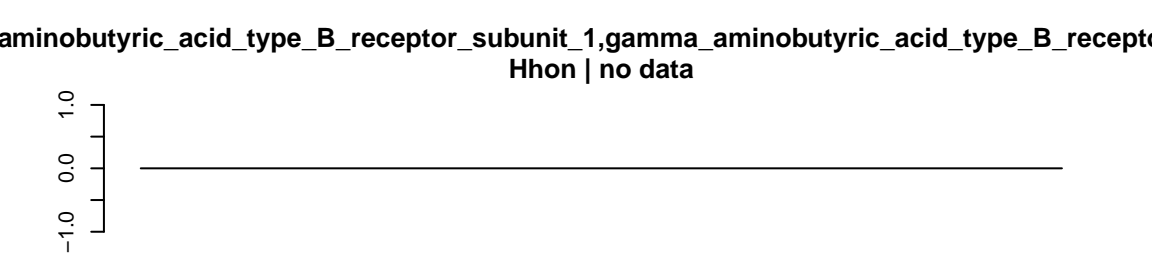
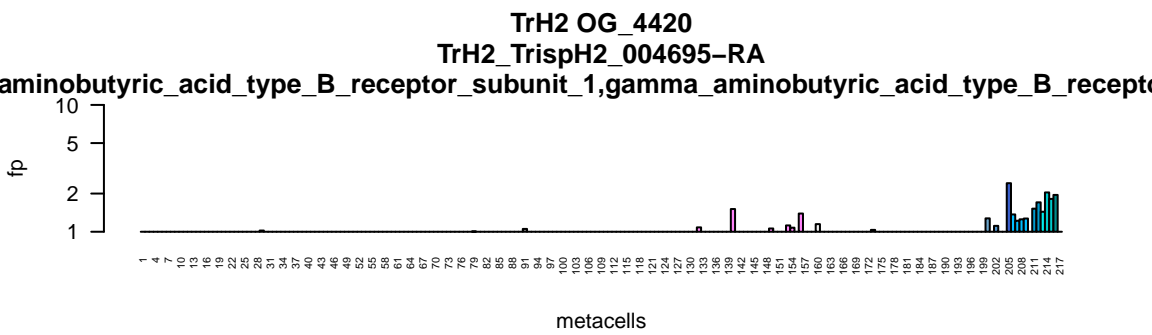
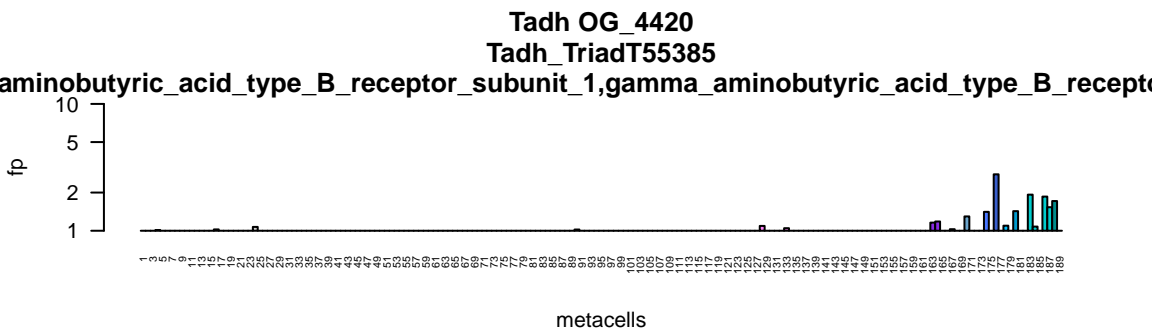
Bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis ranges from 1 to 253, and the y-axis ranges from 1 to 10. The chart shows a distribution of feature counts across metacells, with most metacells having 1 or 2 features. A notable peak is observed at metacell 253, which has 4 features.

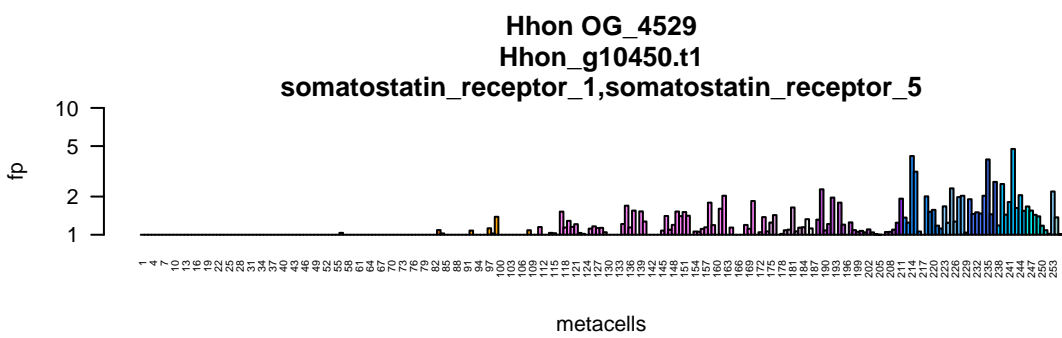
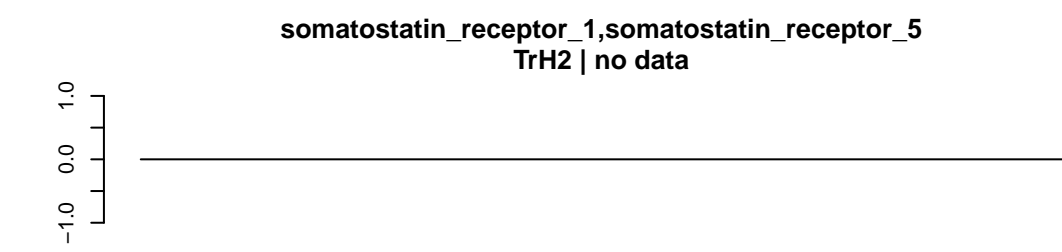
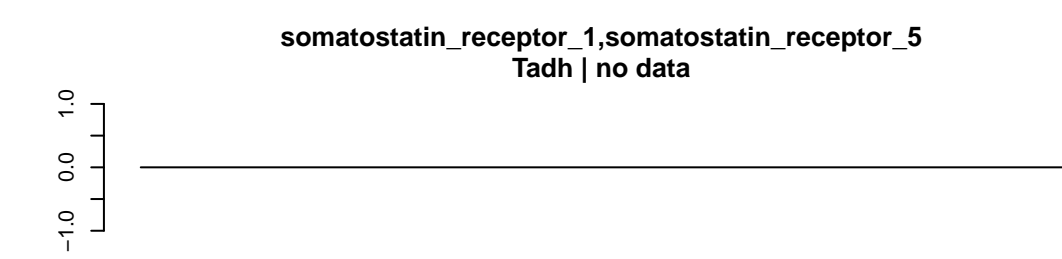
Bar chart showing the frequency of metacells (x-axis) versus the number of features (fp, y-axis). The x-axis lists metacells from 1 to 253. The y-axis shows fp values from 1 to 10. Most metacells have a frequency of 1, with a few having higher frequencies (e.g., metacell 220 has a frequency of 2).

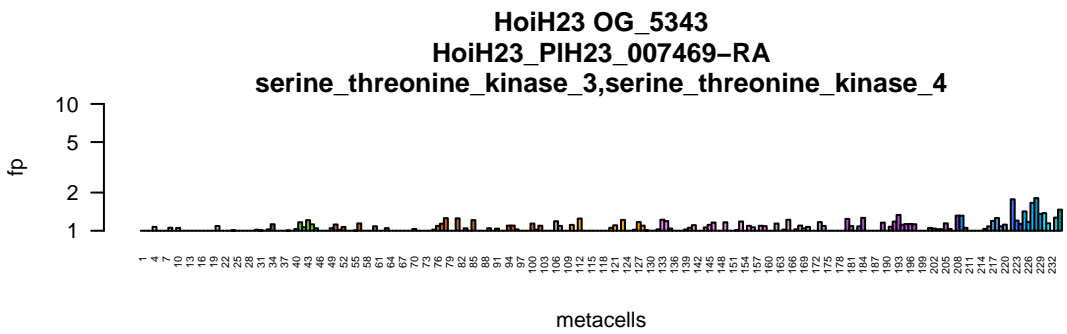
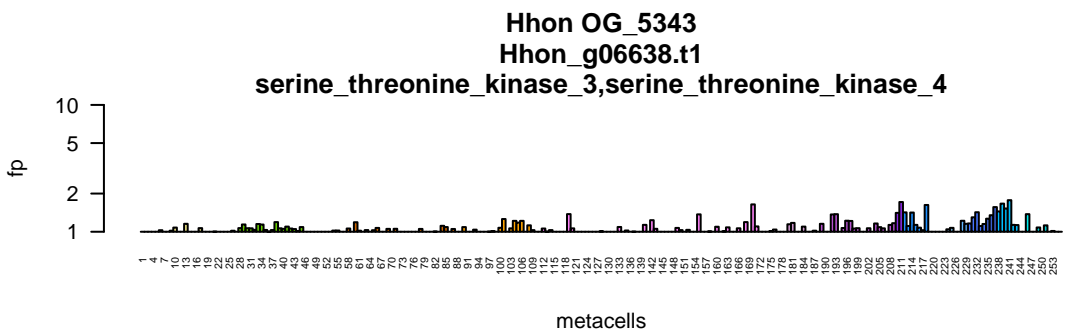
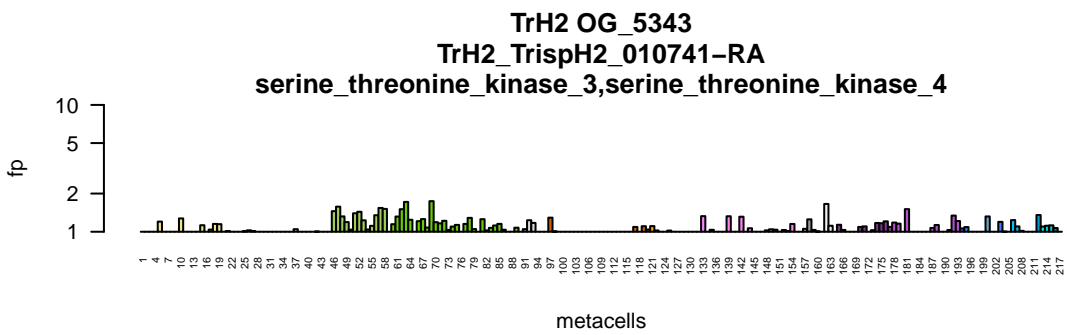
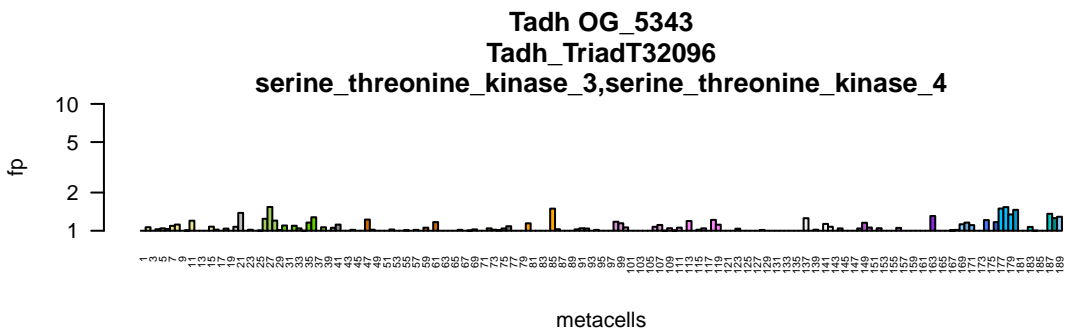
Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 0 to 10. The x-axis is labeled 'metacells' and lists metacell IDs from 1 to 253. Most metacells have 0 false positives, but some have 1 or 2. Metacell 253 has the highest number of false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

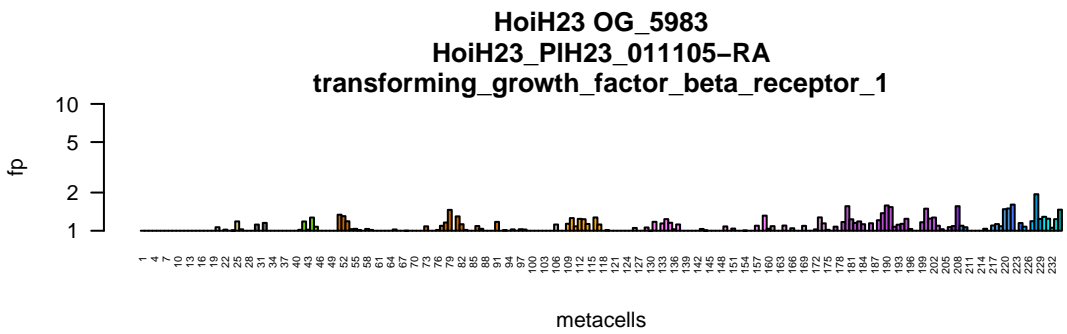
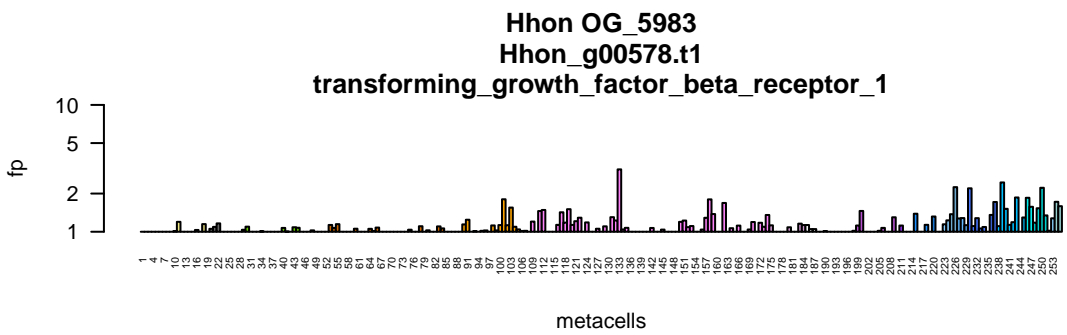
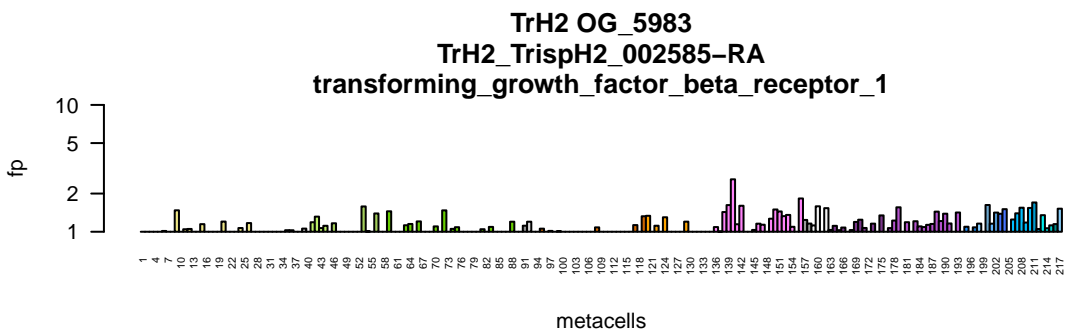
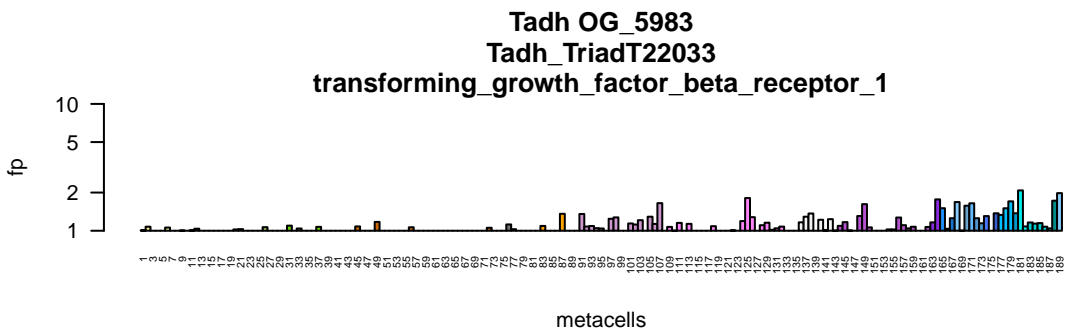
metacell	fp
1	0
4	0
10	0
13	0
19	0
22	0
28	0
31	0
37	0
40	0
46	0
49	0
55	0
64	0
70	0
73	0
76	0
82	0
85	0
88	0
91	0
97	0
100	0
106	0
109	0
115	0
118	0
127	0
129	0
134	0
136	0
142	0
145	0
154	0
157	0
160	0
163	0
169	0
172	0
178	0
184	0
187	0
191	0
193	0
195	0
200	0
202	0
205	0
214	0
219	0
223	0
229	0
232	0
233	0
238	0
241	0
247	0
250	0
253	2



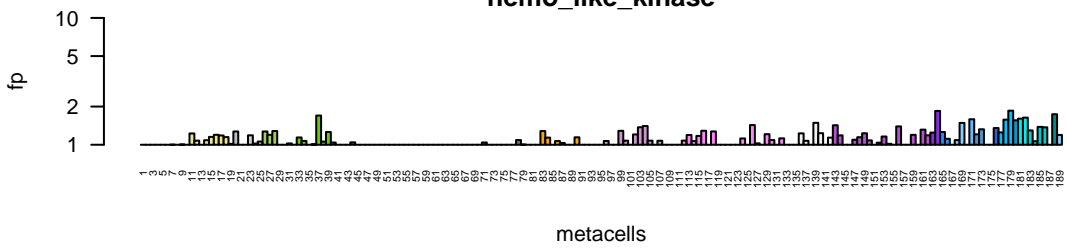






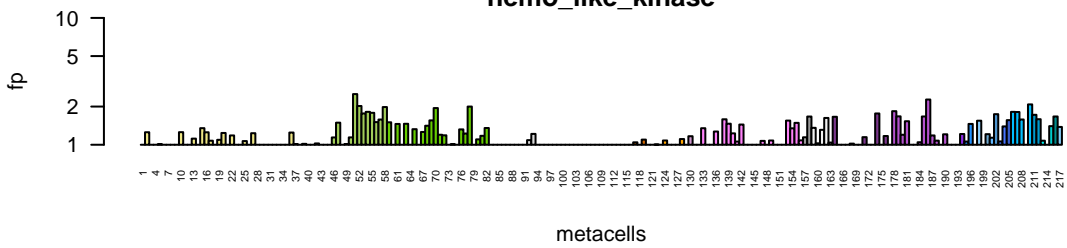


Tadh OG_6114
Tadh_TriadT38054
nemo_like_kinase



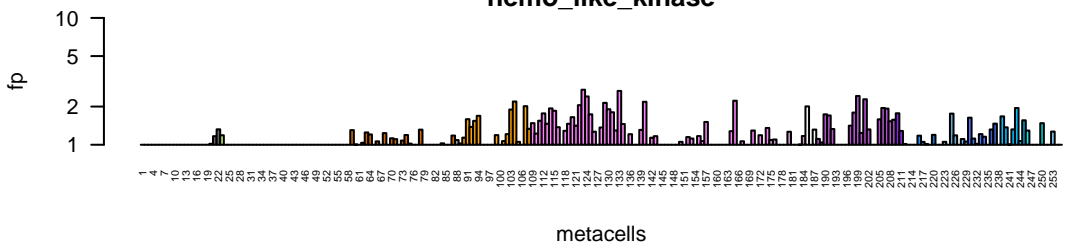
metacells

TrH2 OG_6114
TrH2_TrispH2_006894-RA
nemo_like_kinase



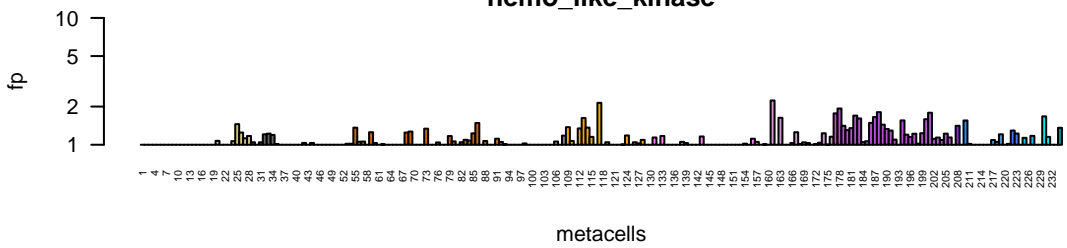
metacells

Hhon OG_6114
Hhon_g02719.t1
nemo_like_kinase

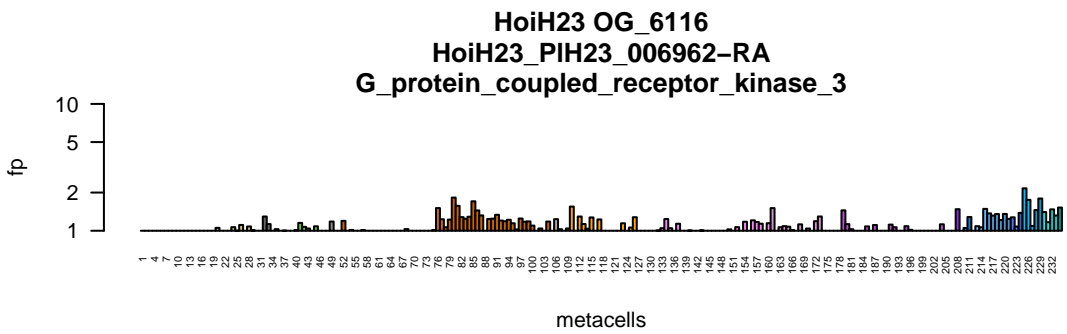
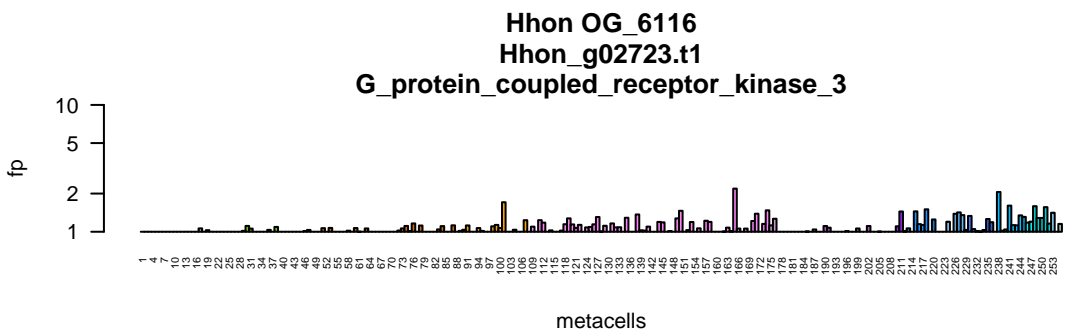
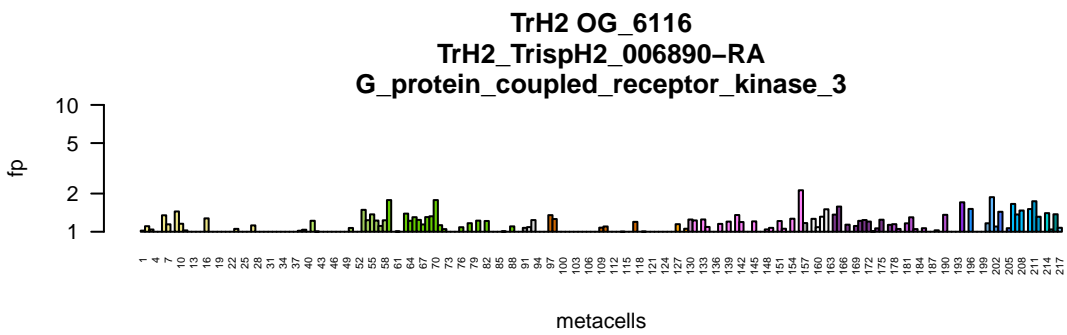
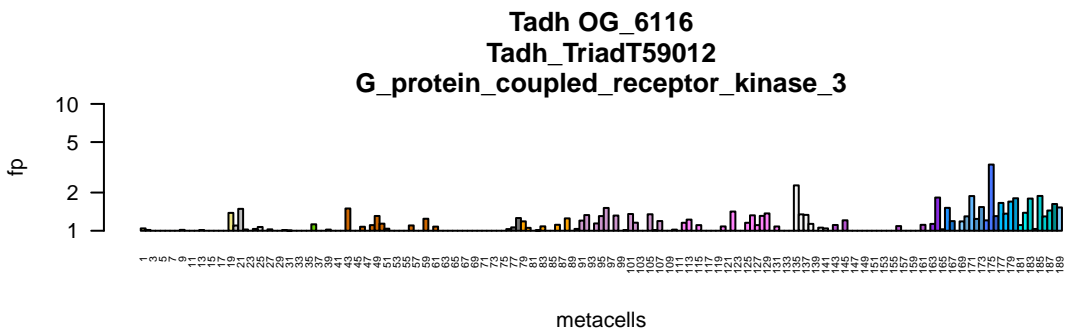


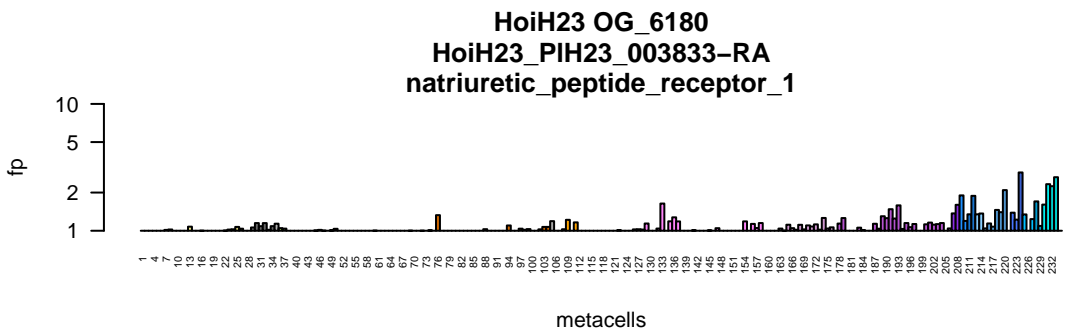
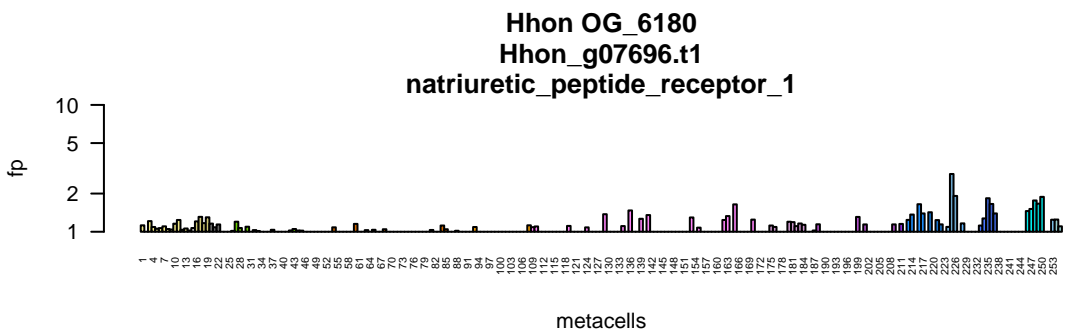
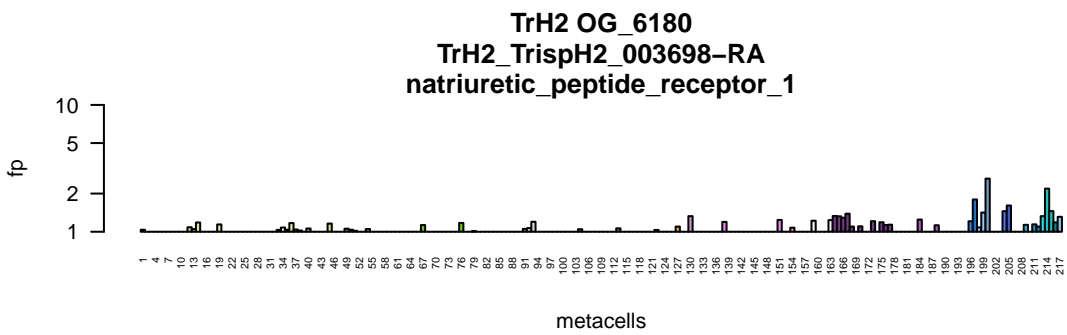
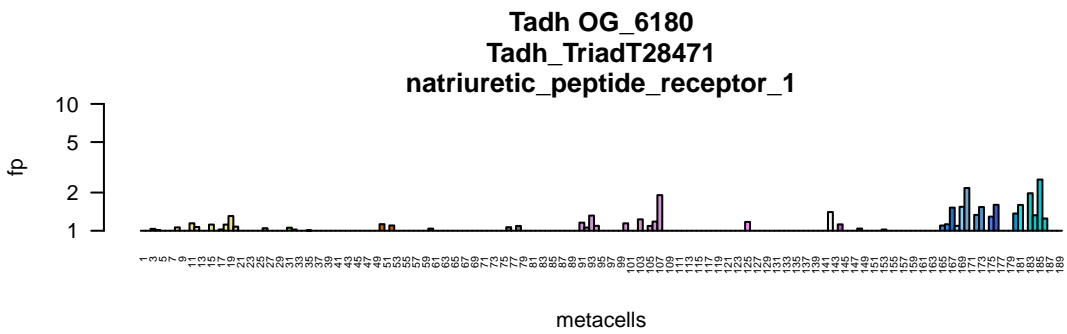
metacells

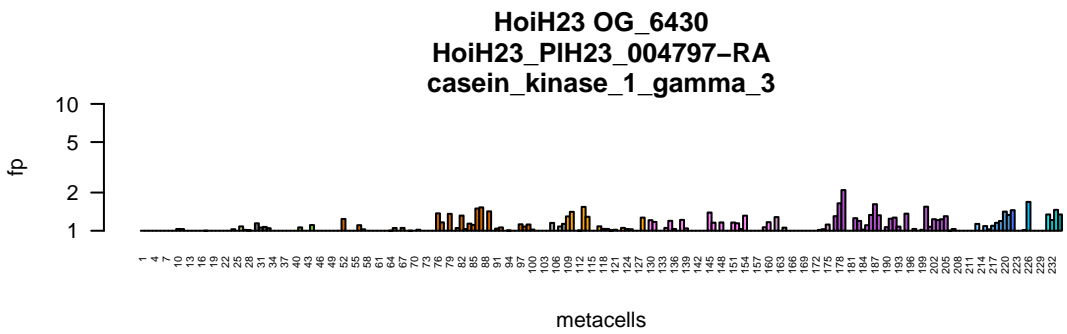
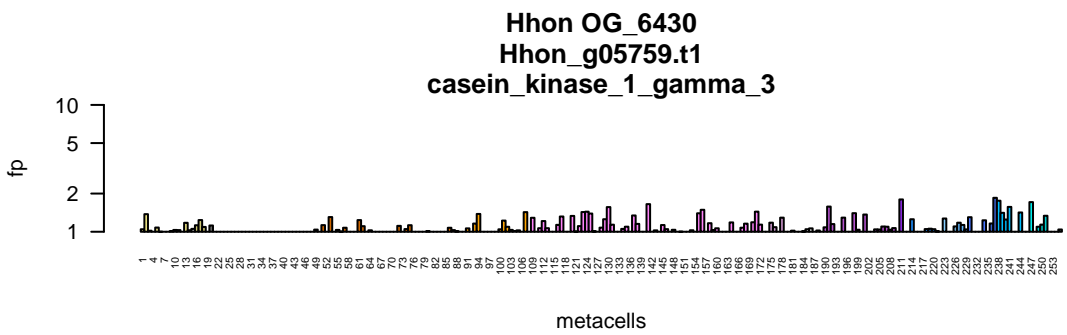
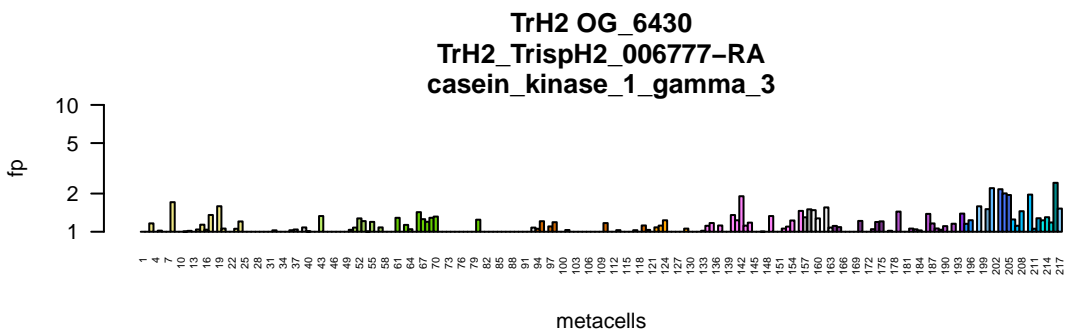
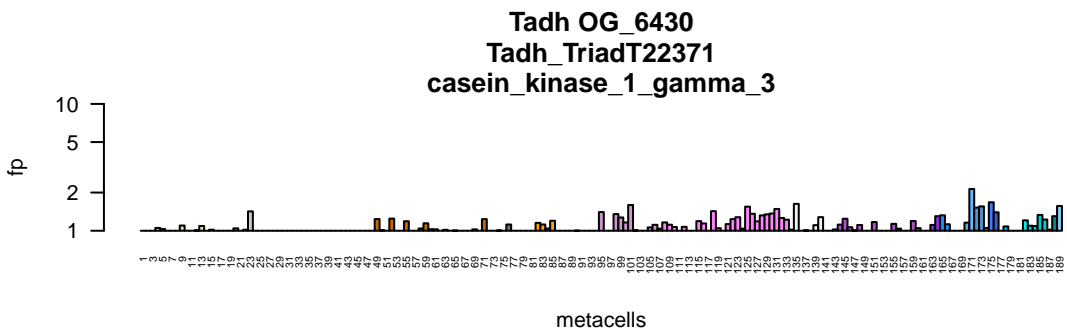
HoiH23 OG_6114
HoiH23_PIH23_006958-RA
nemo_like_kinase

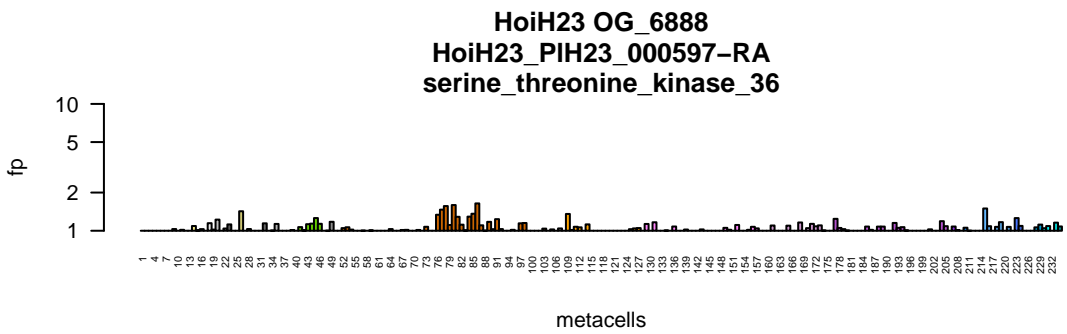
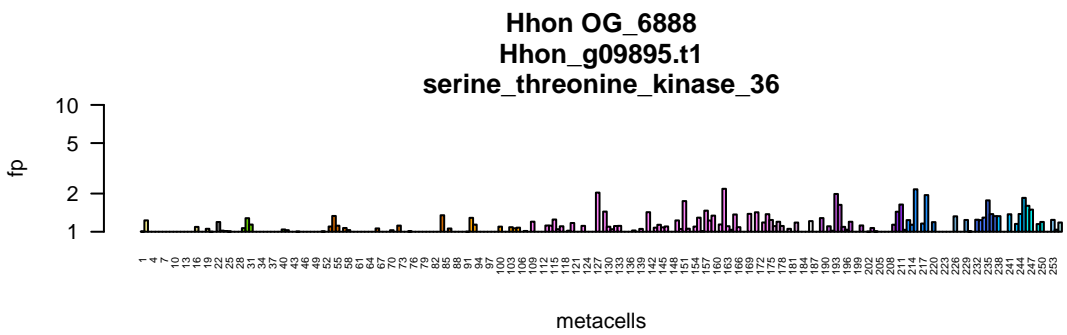
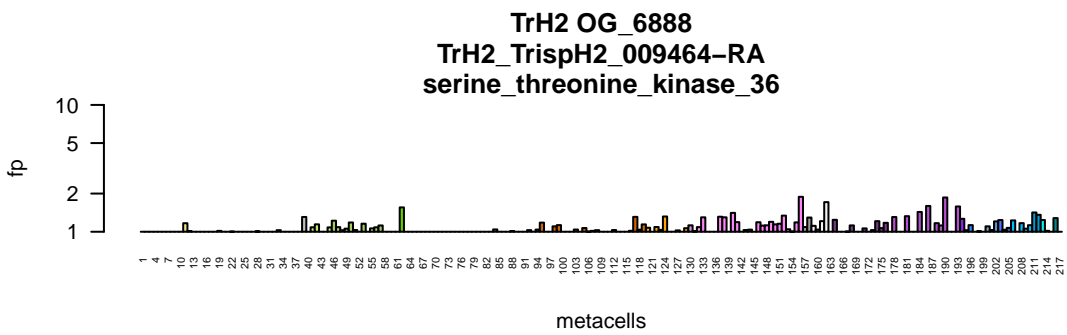
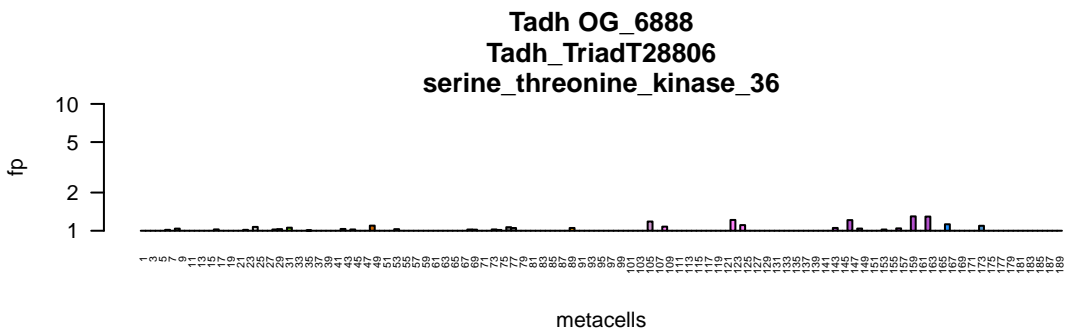


metacells





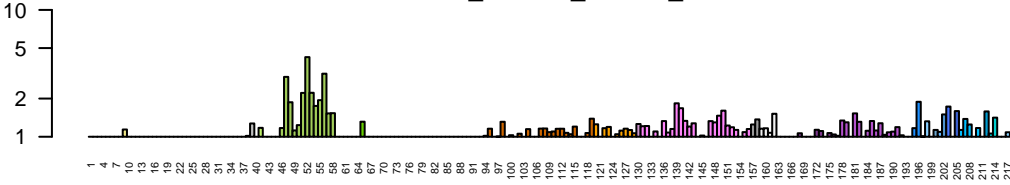




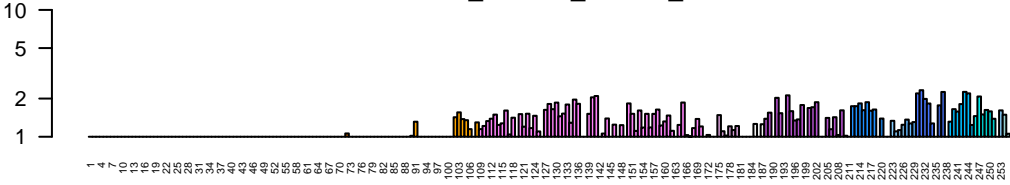
NIMA_related_kinase_7
Tadh | no data



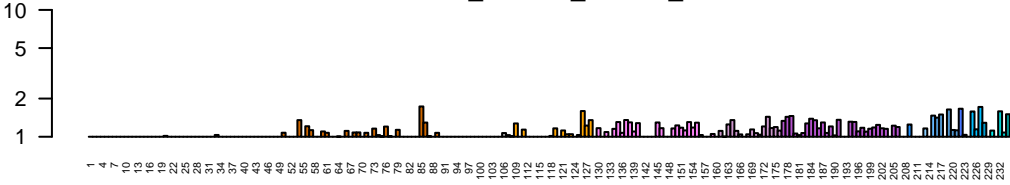
TrH2 OG_6968
TrH2_TrispH2_004112-RA
NIMA_related_kinase_7

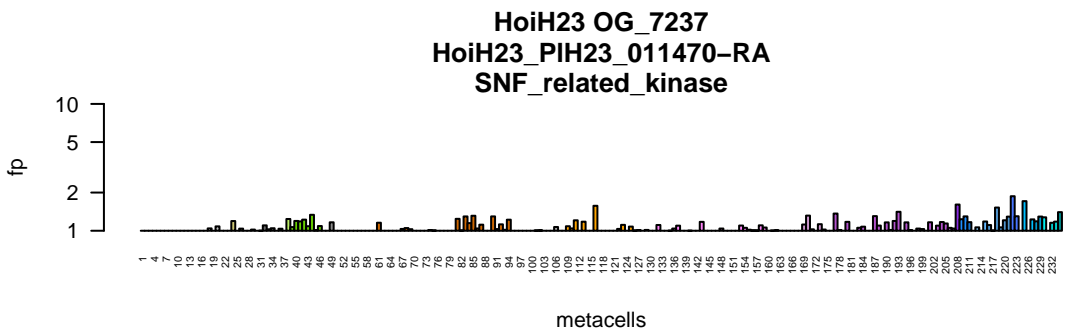
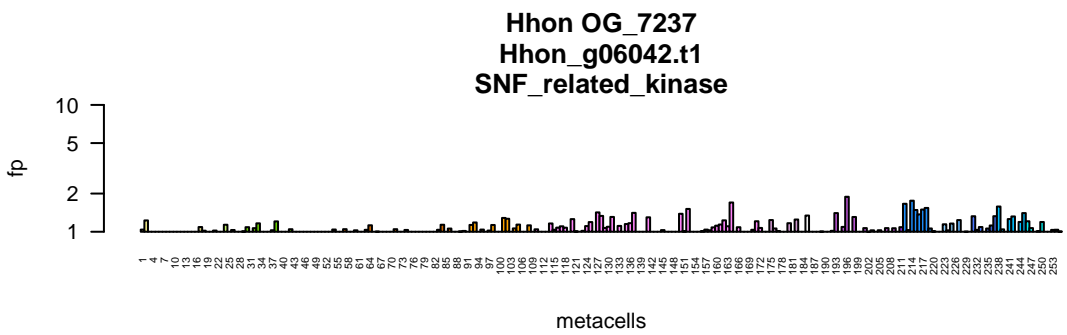
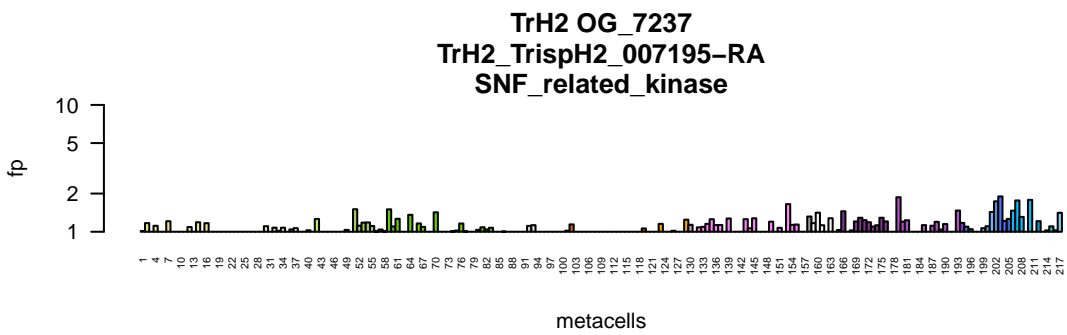
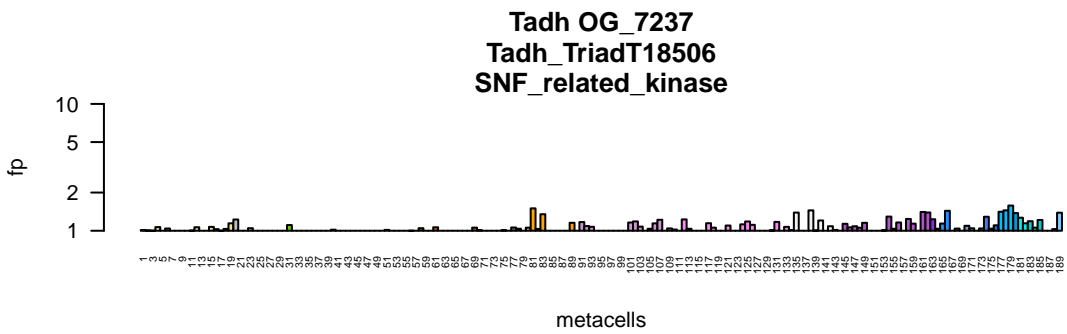


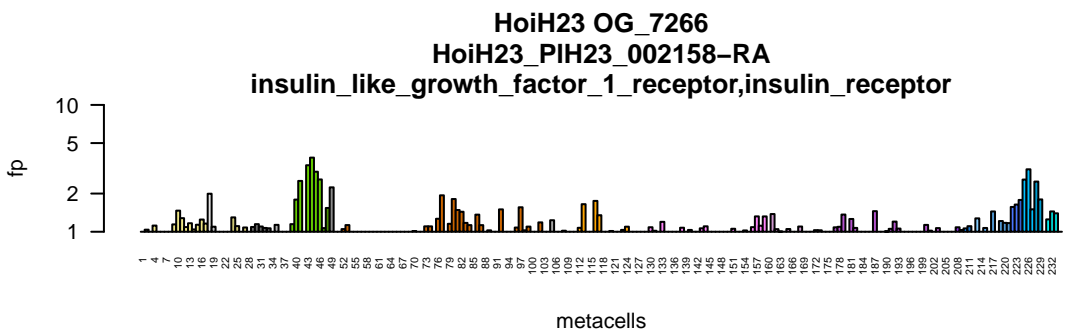
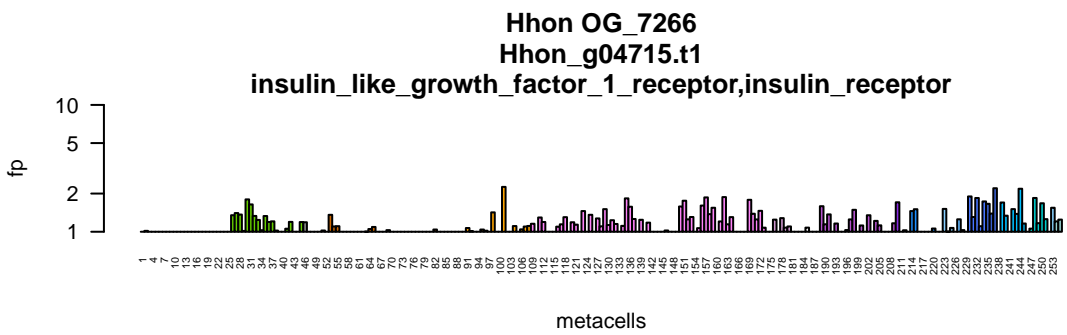
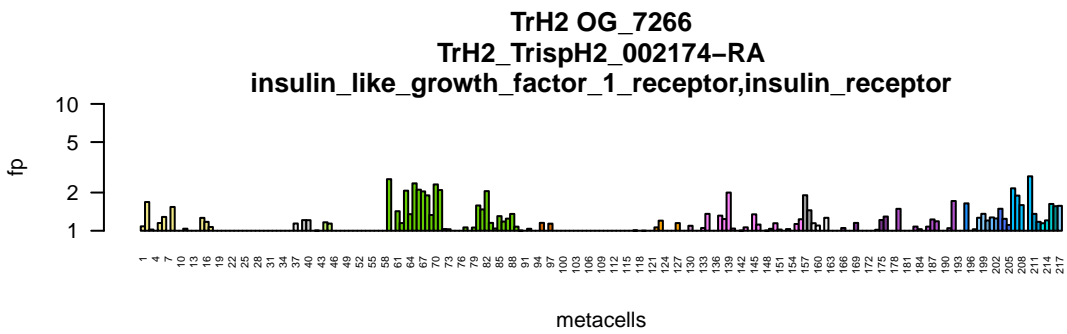
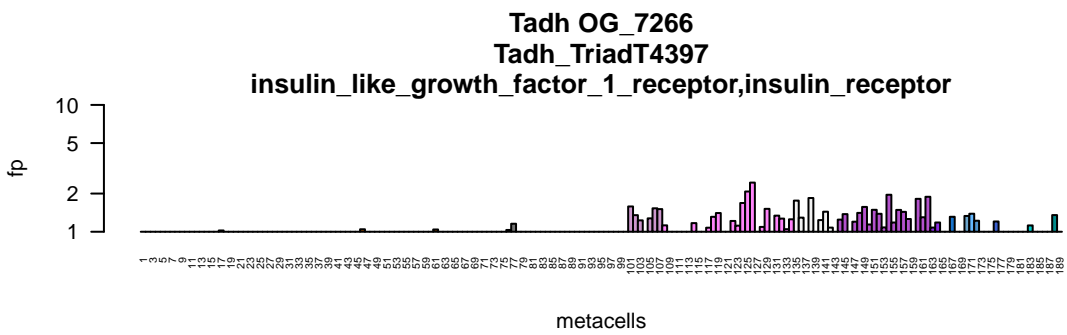
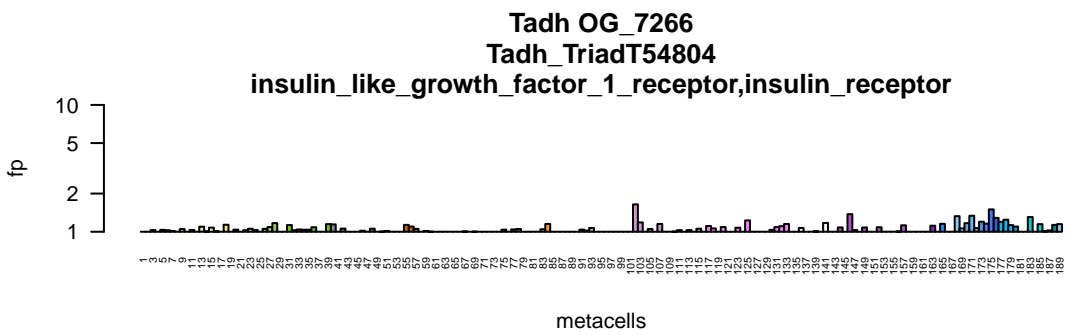
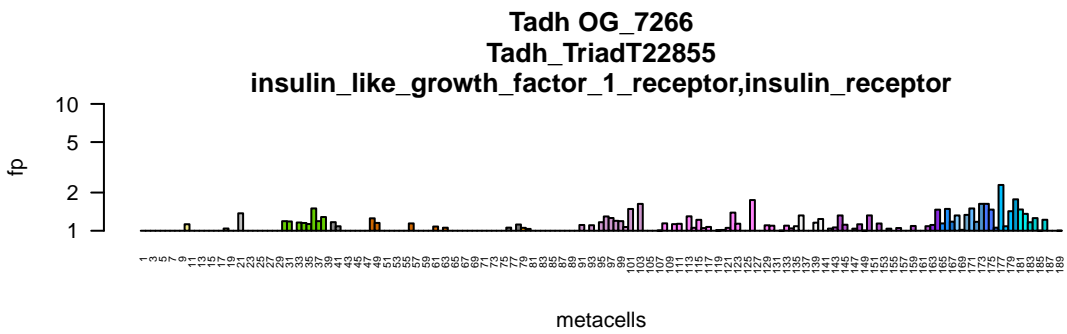
Hhon OG_6968
Hhon_g01675.t1
NIMA_related_kinase_7

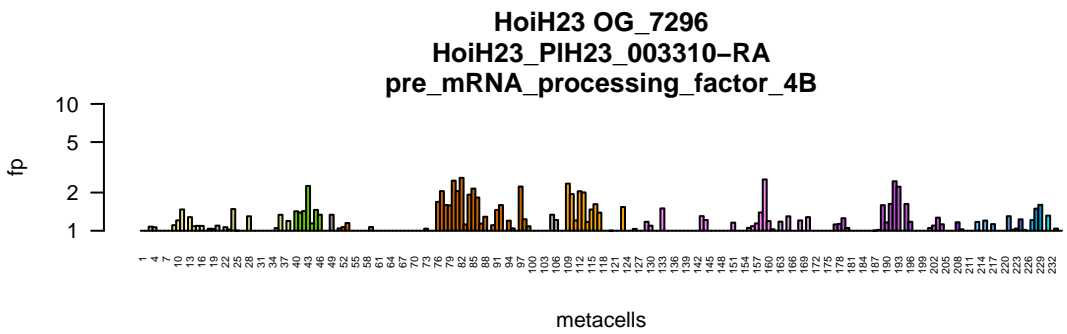
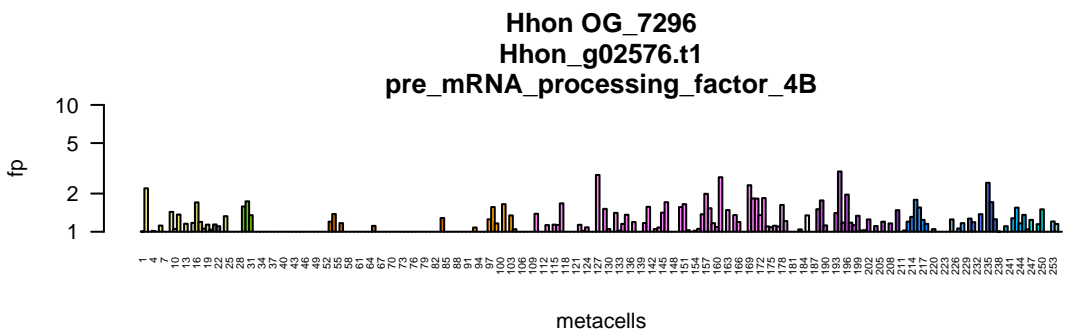
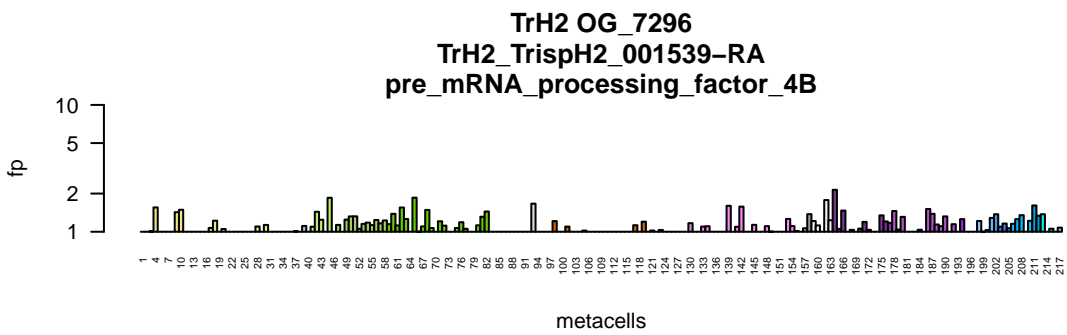
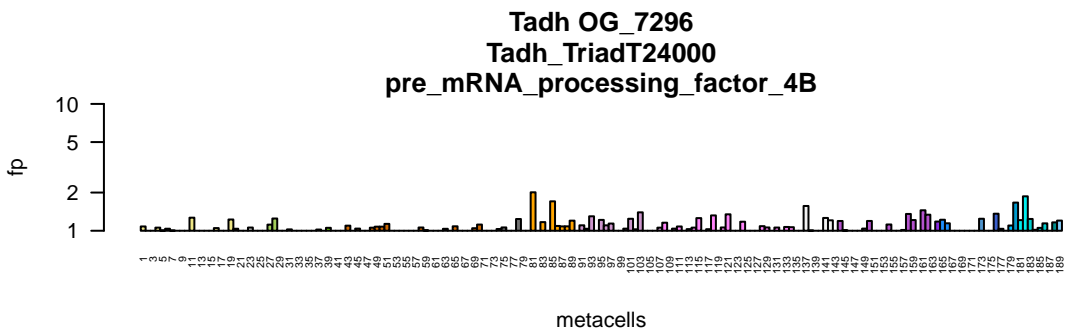


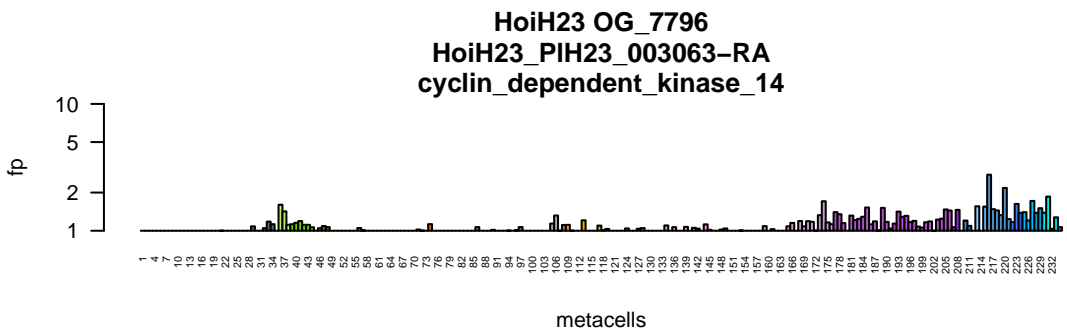
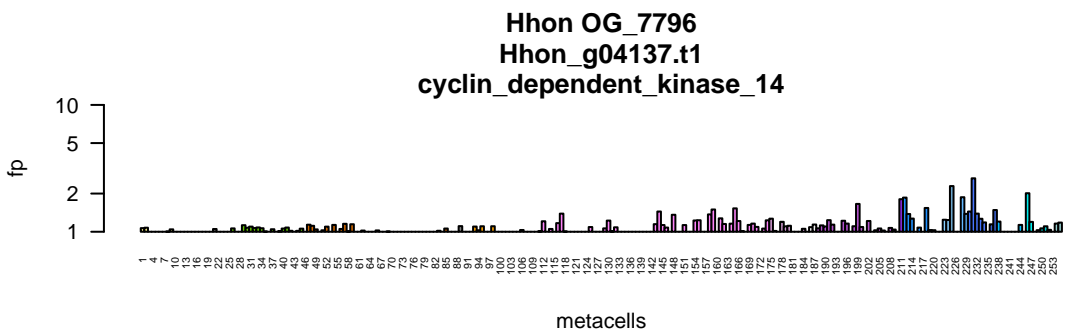
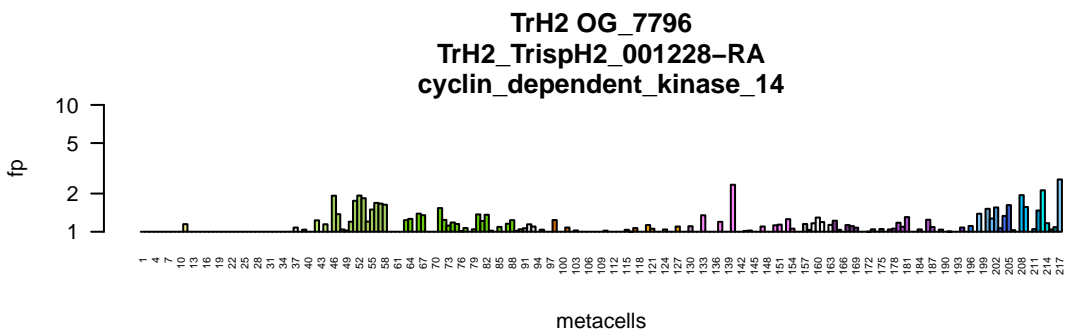
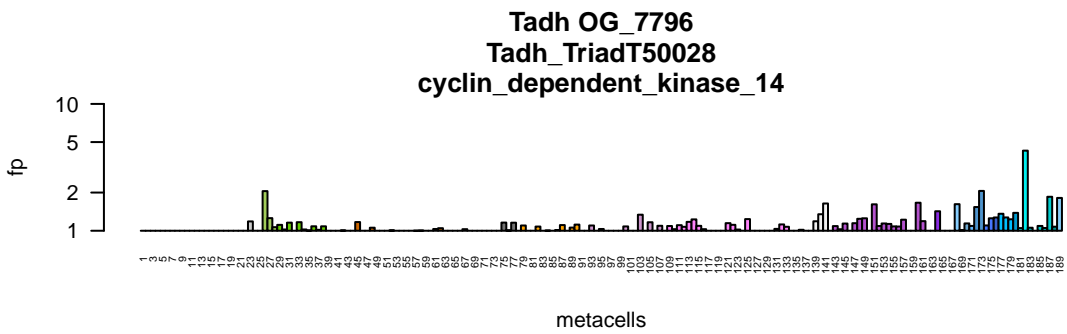
HoiH23 OG_6968
HoiH23_PIH23_005337-RA
NIMA_related_kinase_7

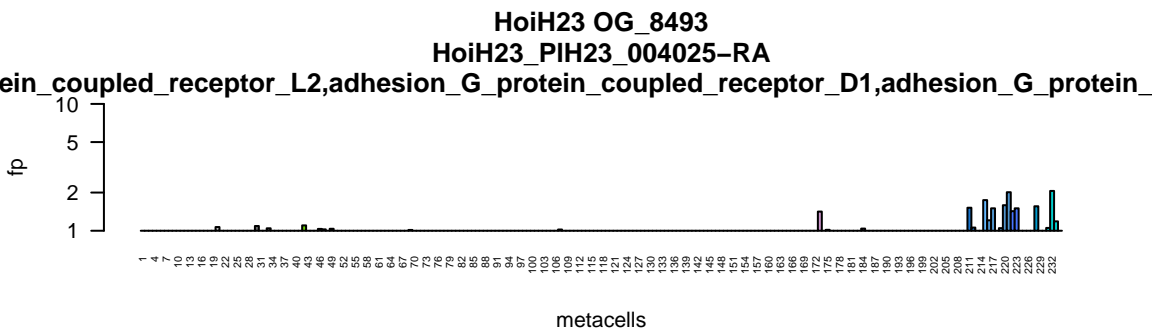
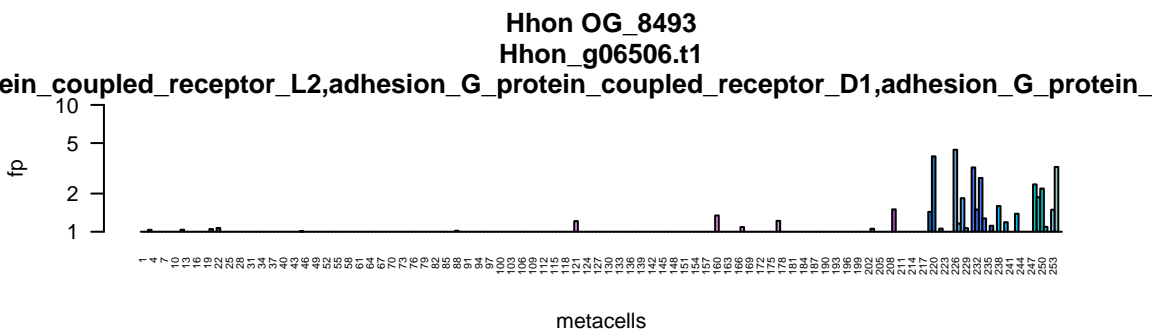
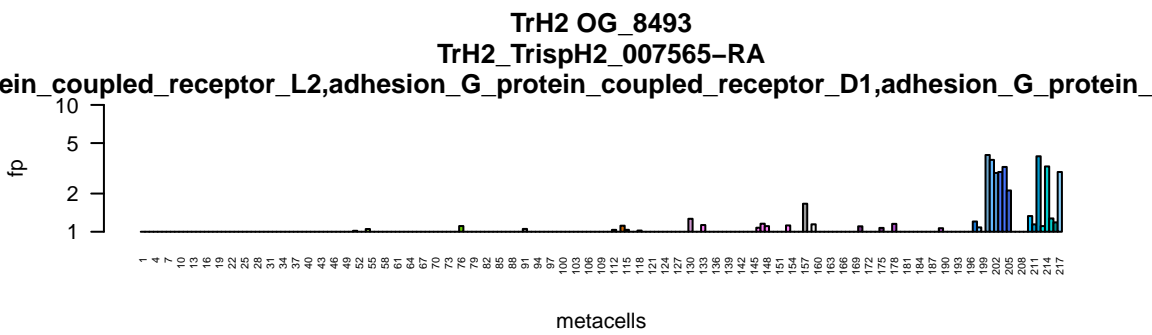
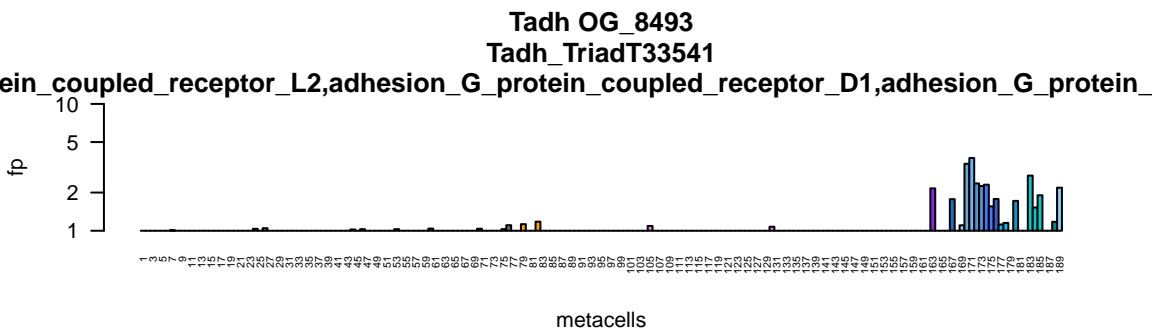


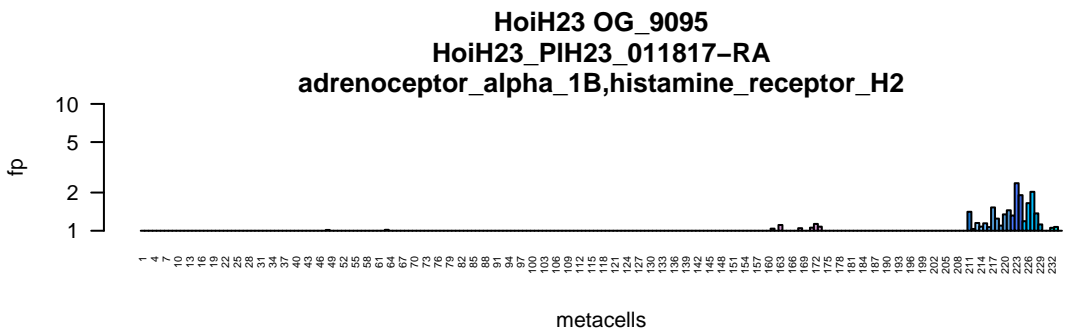
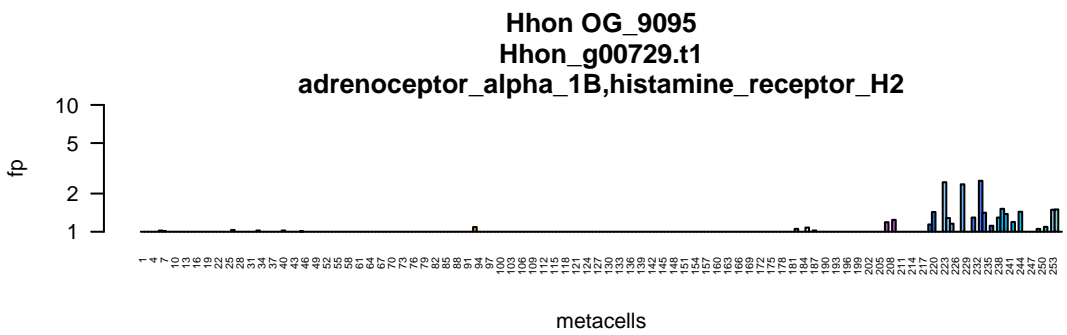
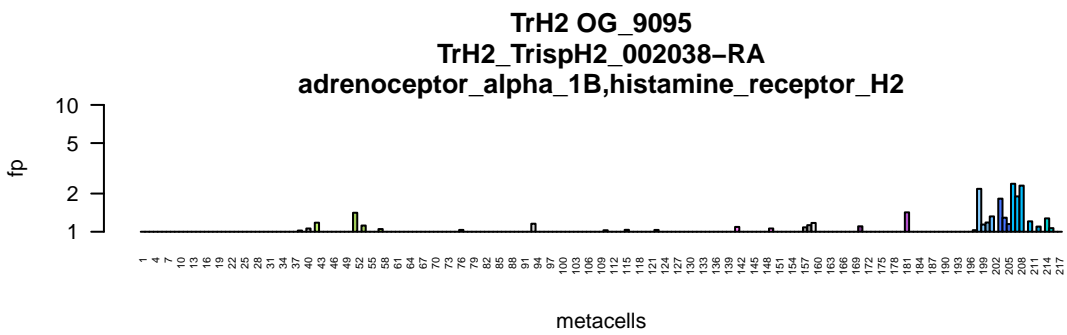
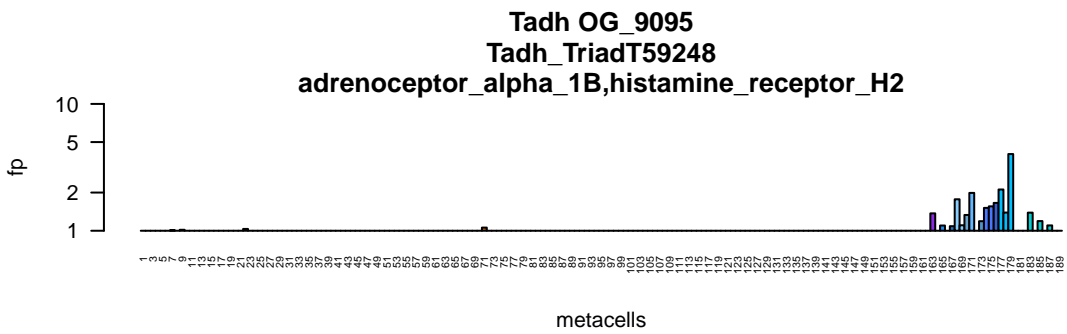








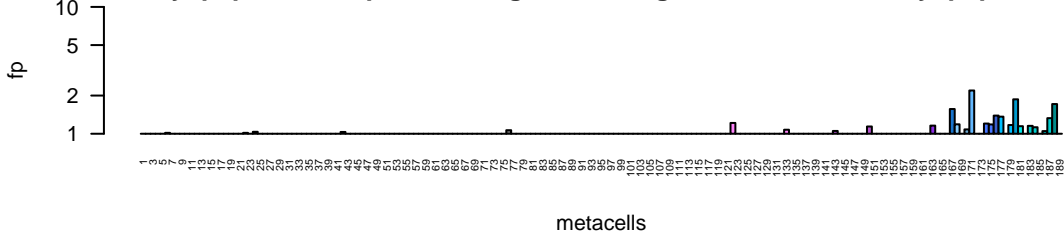




Tadh OG_9223

Tadh_TriadT51796

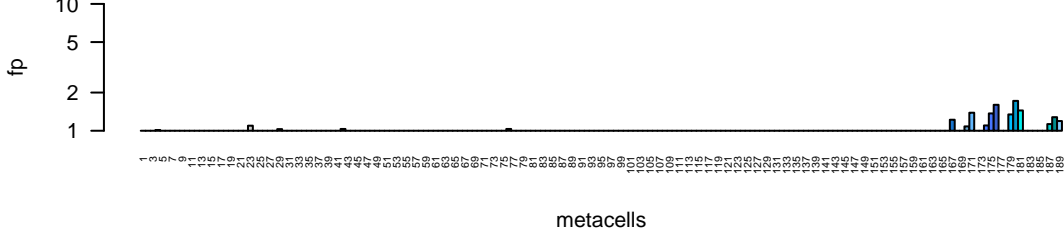
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Tadh OG_9223

Tadh_TriadT51797

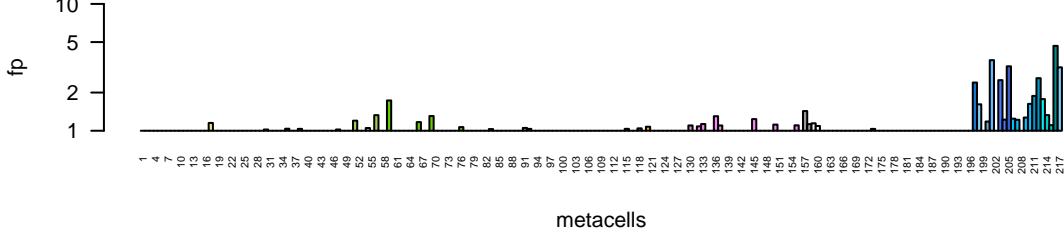
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



TrH2 OG_9223

TrH2_TrispH2_008387-RA

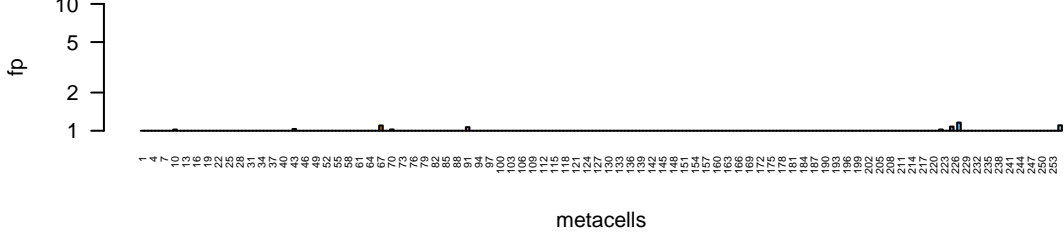
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Hhon OG_9223

Hhon_g11380.t1

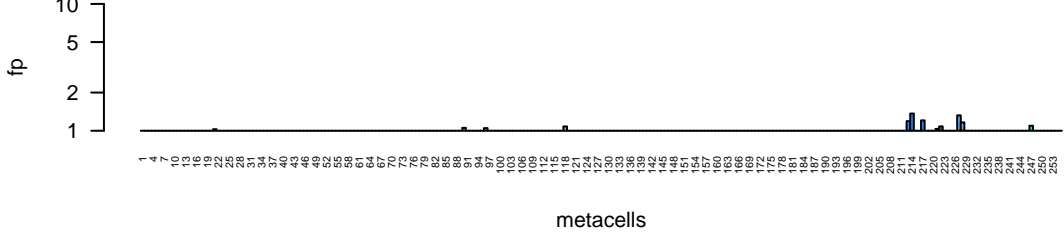
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Hhon OG_9223

Hhon_g01959.t1

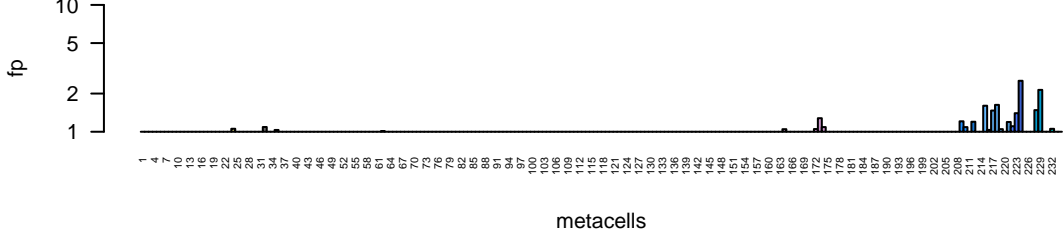
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto

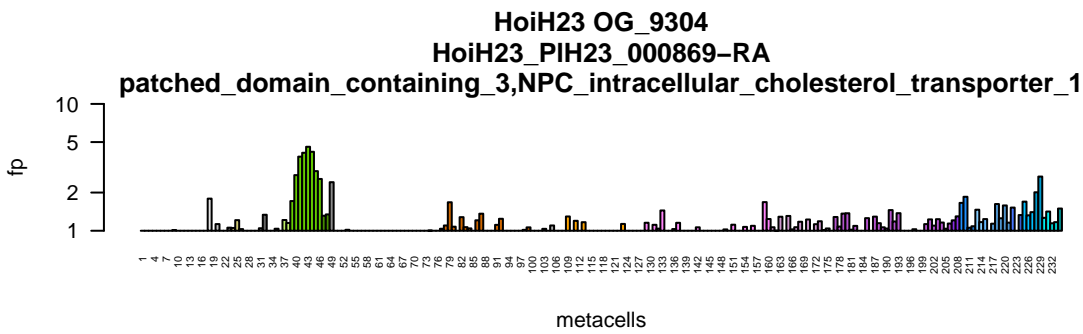
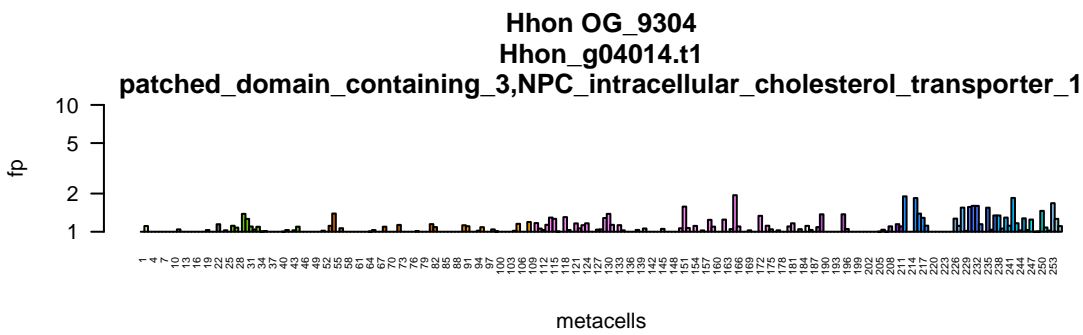
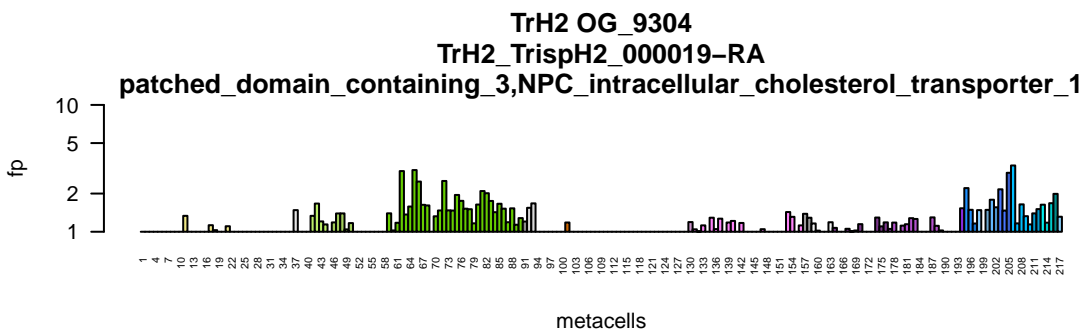
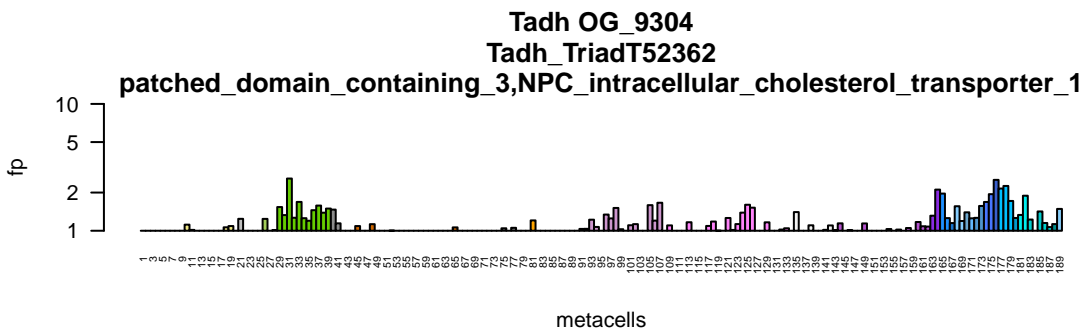


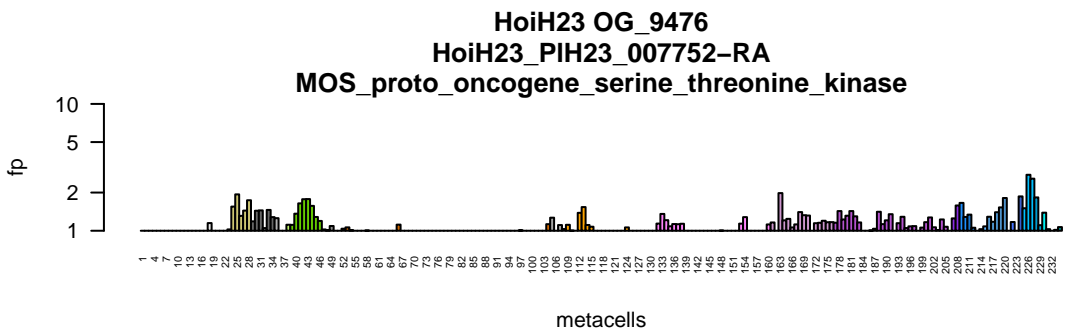
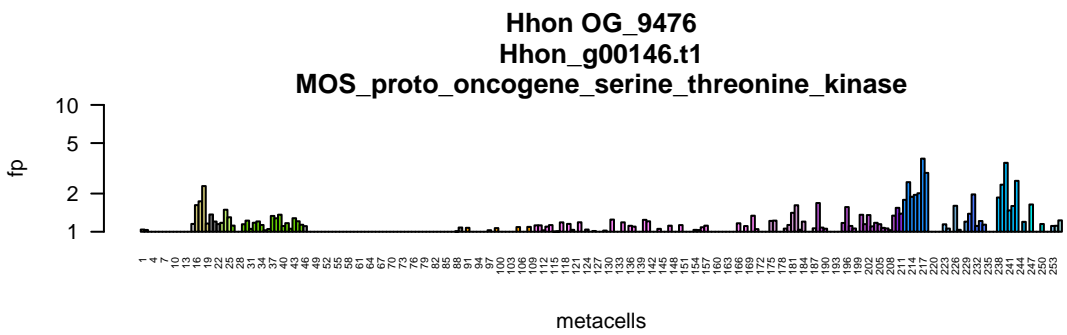
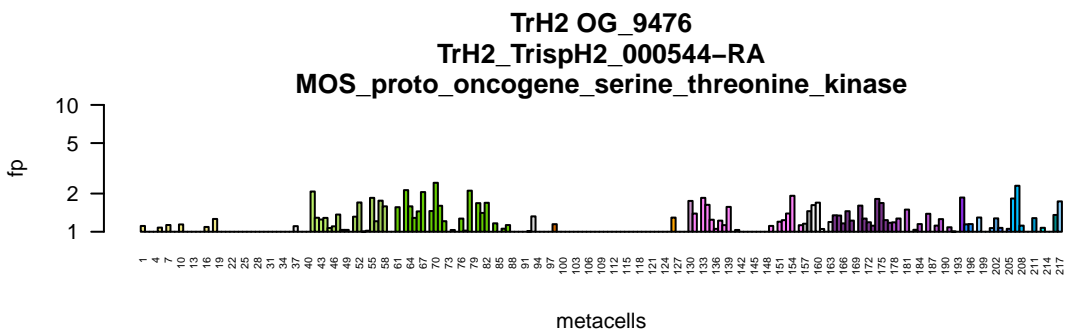
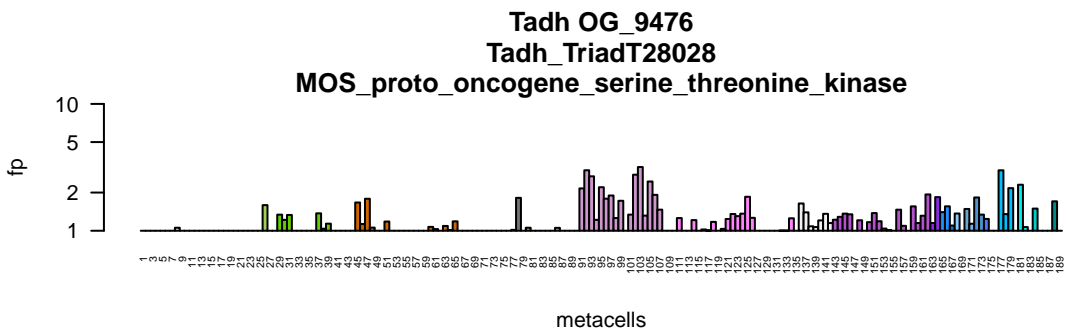
HoiH23 OG_9223

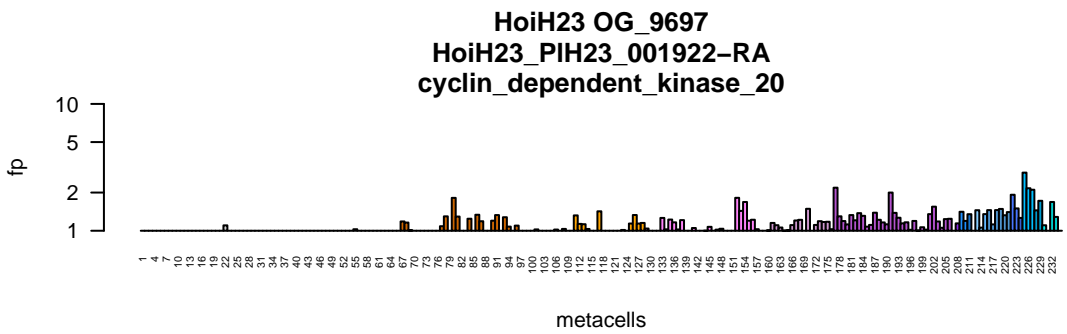
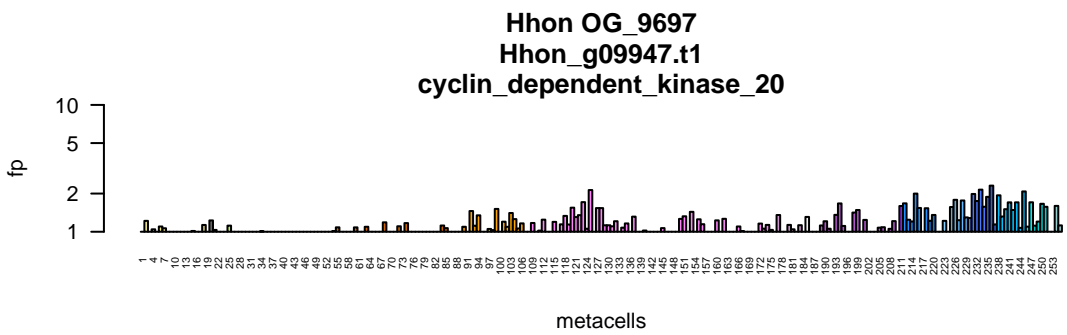
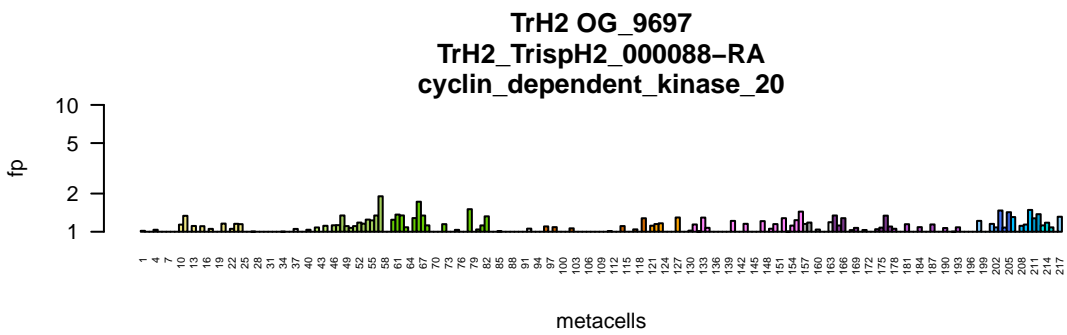
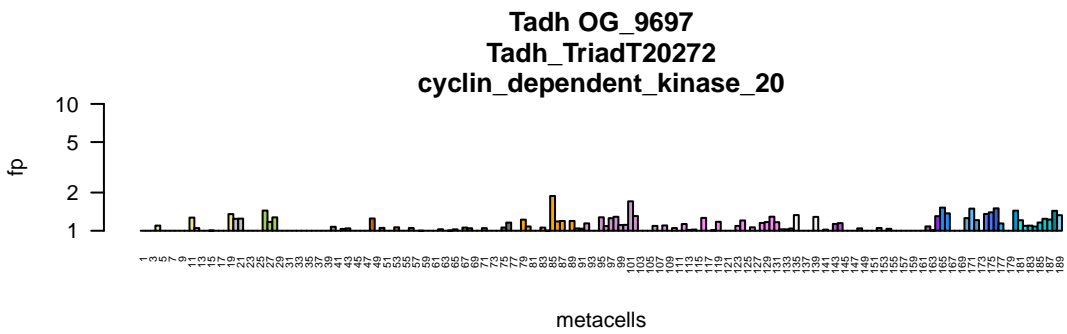
HoiH23_PIH23_004873-RA

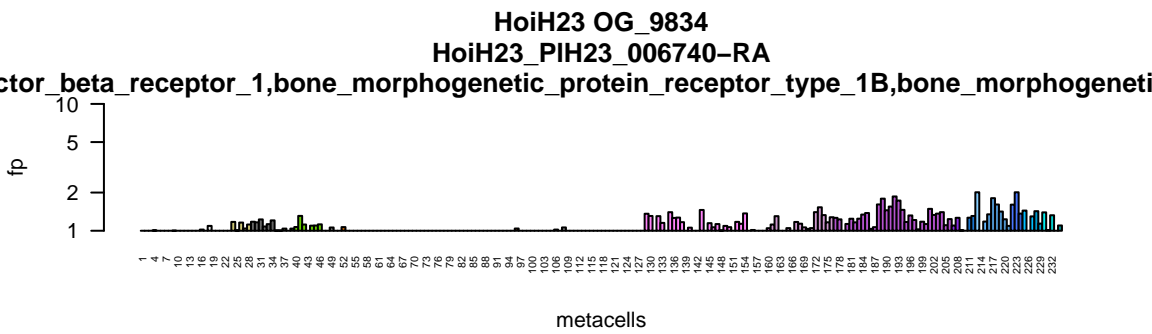
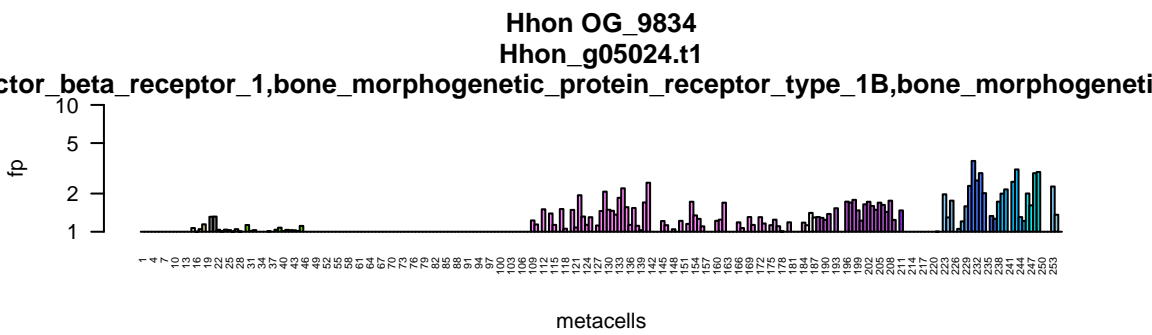
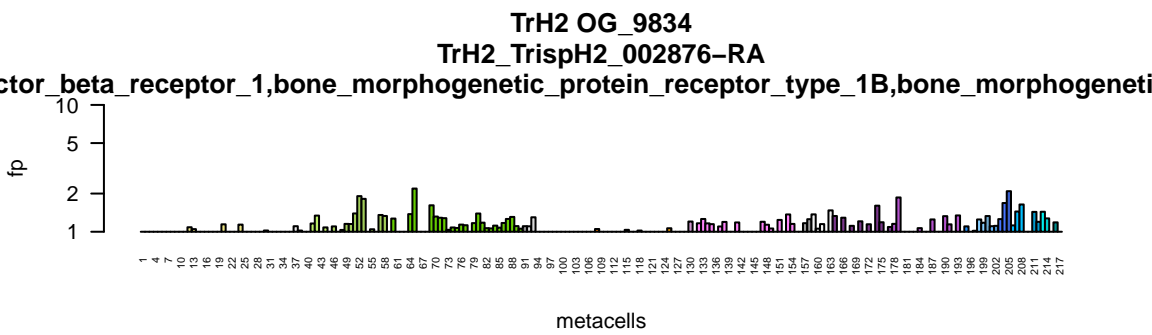
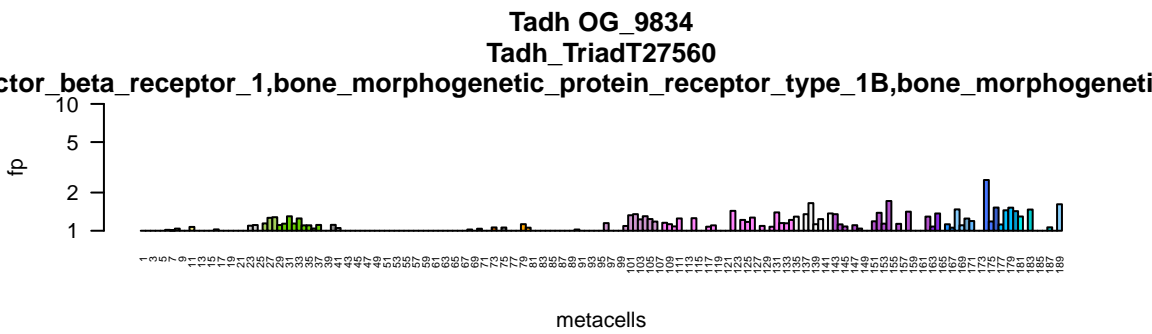
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto

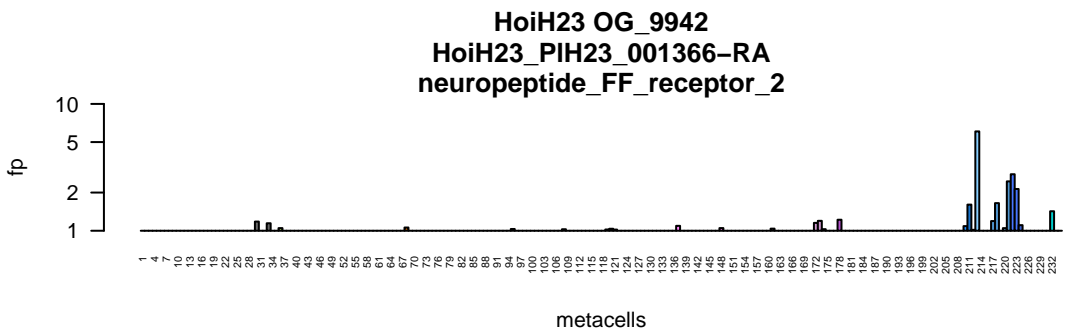
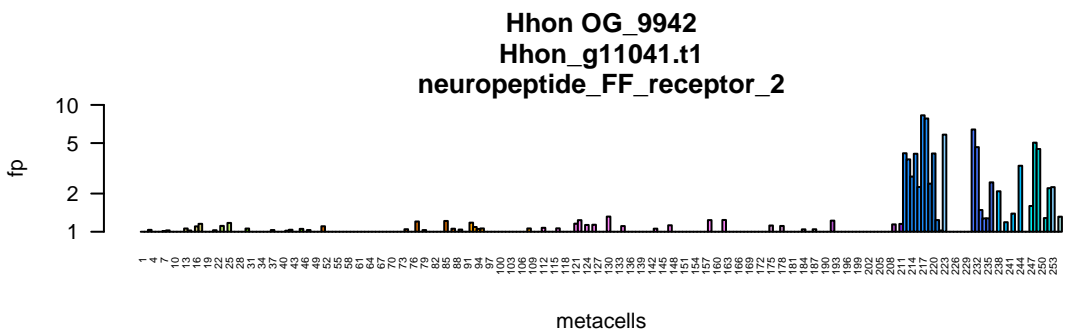
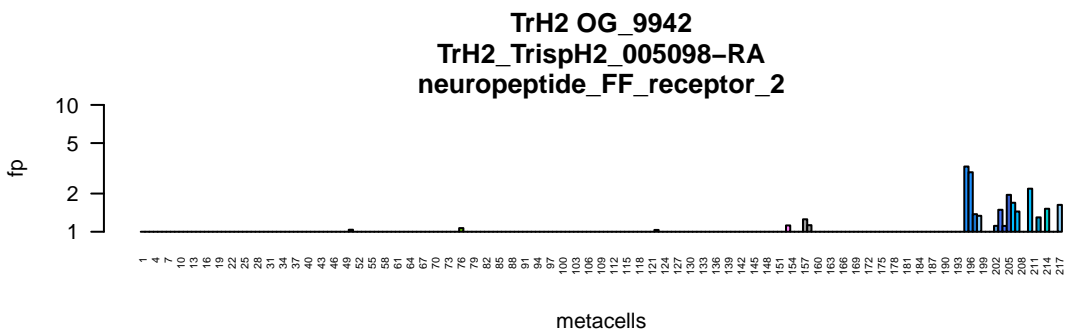
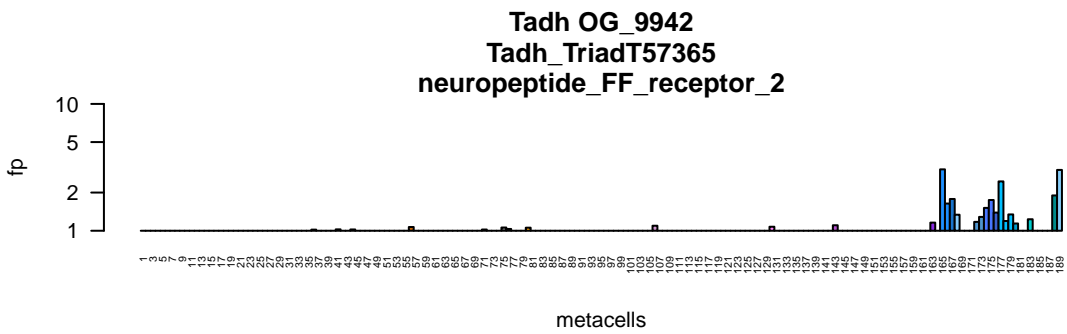


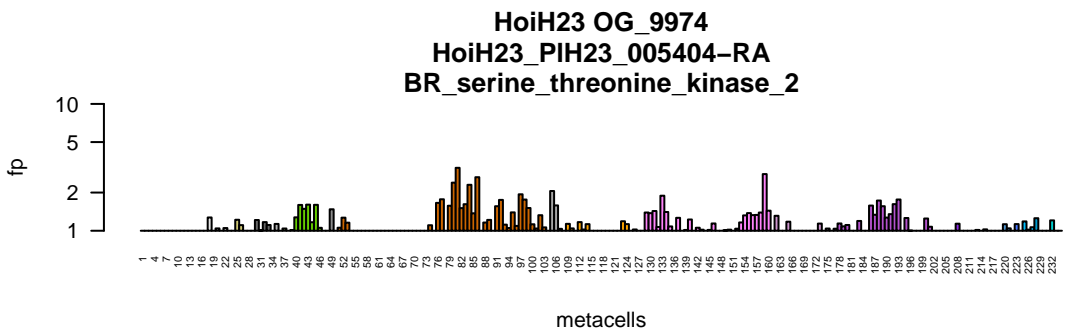
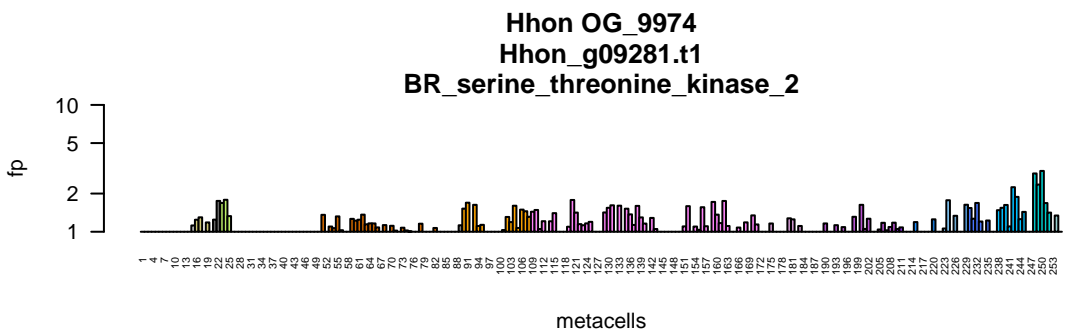
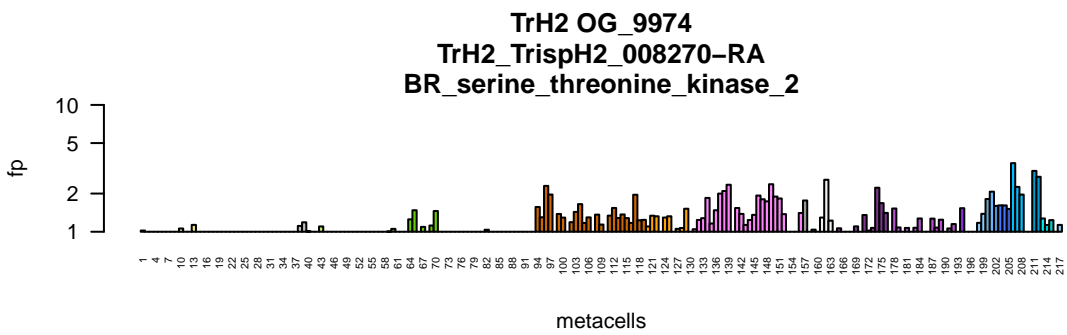
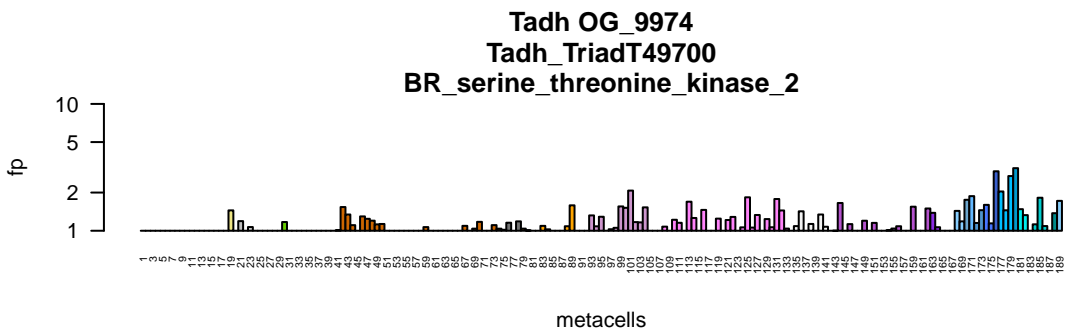








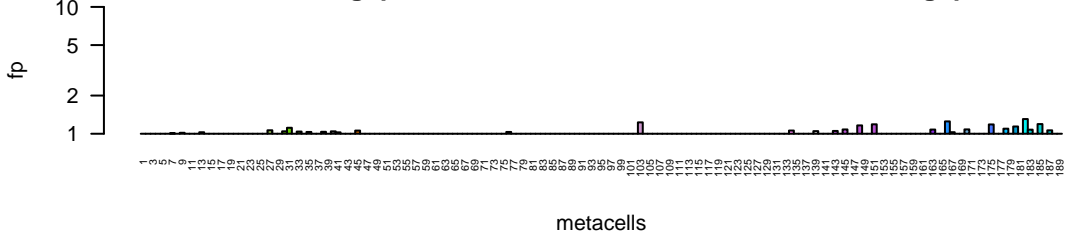




Tadh OG_9990

Tadh_TriadT32832

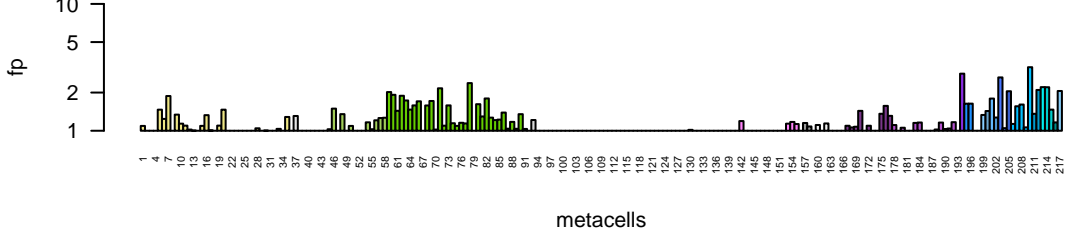
homeodomain_interacting_protein_kinase_2,homeodomain_interacting_protein_kinase_2



TrH2 OG_9990

TrH2_TrispH2_010386-RA

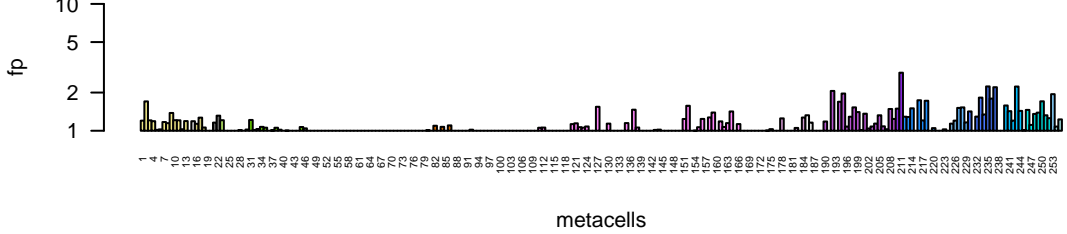
homeodomain_interacting_protein_kinase_2,homeodomain_interacting_protein_kinase_2



Hhon OG_9990

Hhon_g05557.t1

homeodomain_interacting_protein_kinase_2,homeodomain_interacting_protein_kinase_2



HoiH23 OG_9990

HoiH23_PIH23_004935-RA

homeodomain_interacting_protein_kinase_2,homeodomain_interacting_protein_kinase_2

