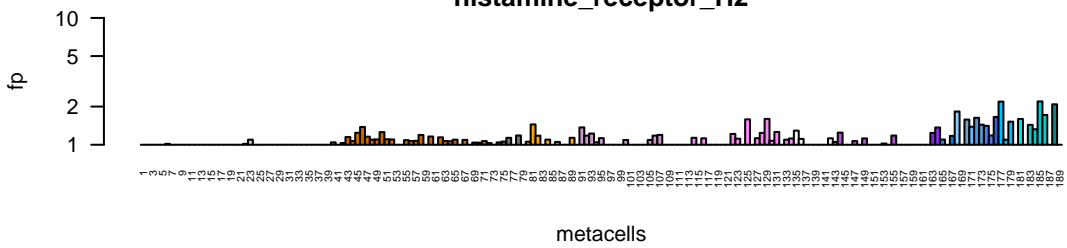
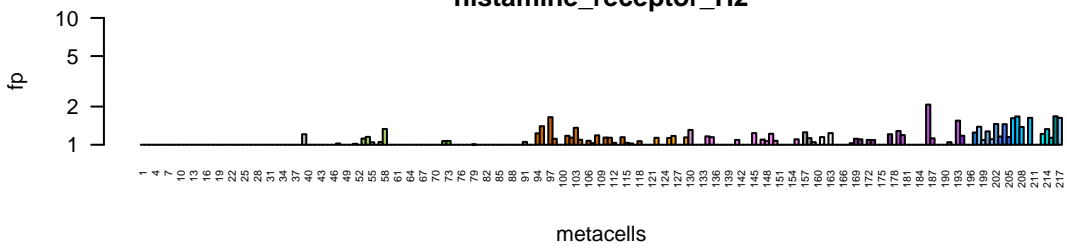


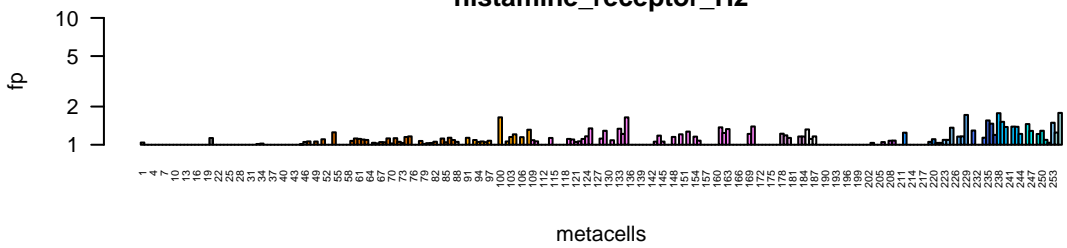
Tadh OG_6463
Tadh_TriadT60646
histamine_receptor_H2



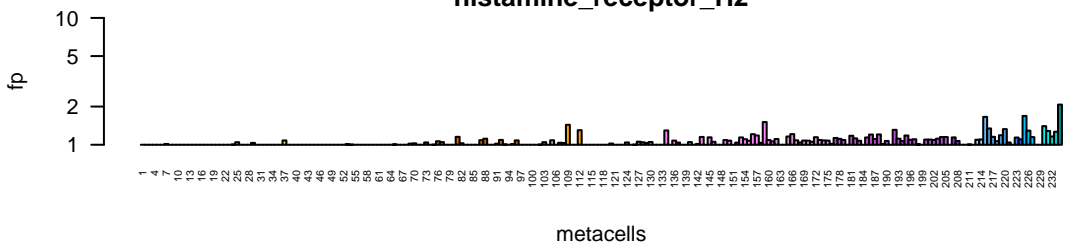
TrH2 OG_6463
TrH2_TrispH2_009668-RA
histamine_receptor_H2

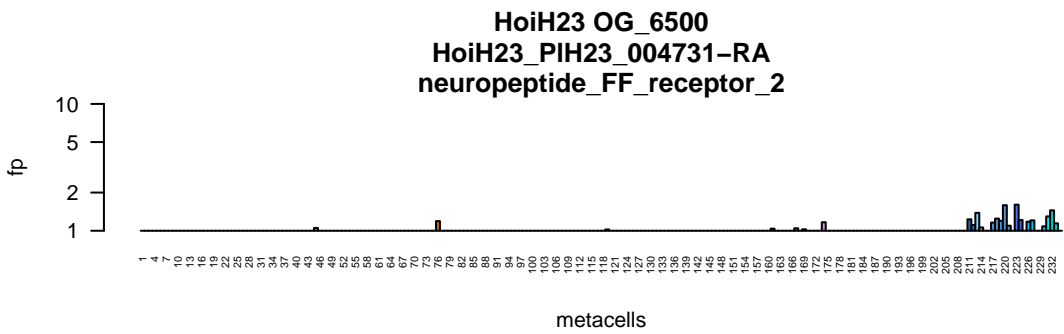
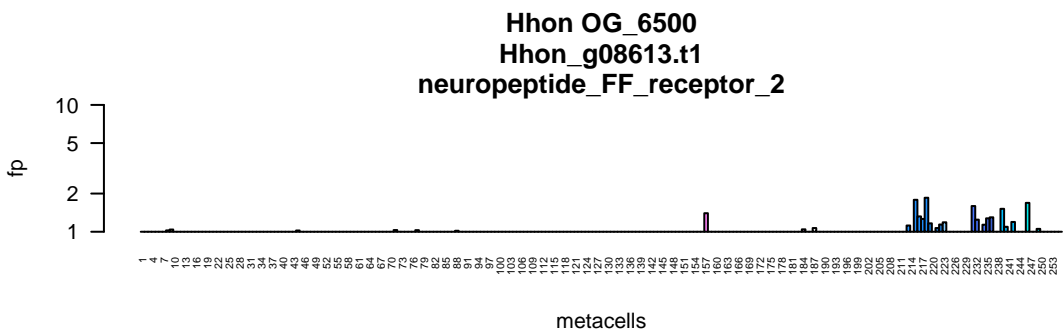
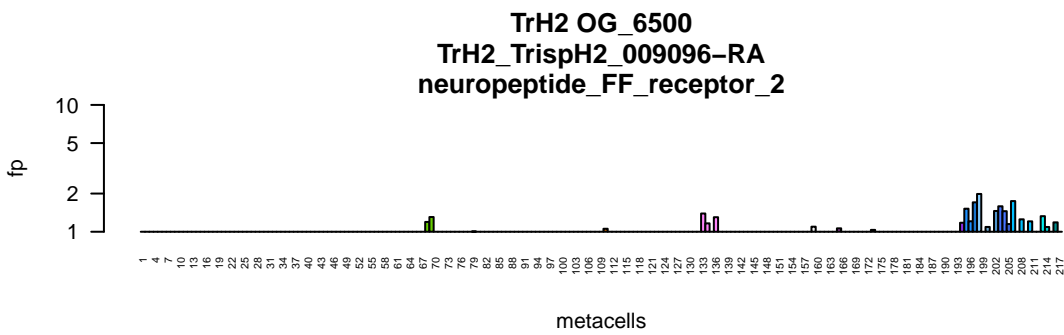
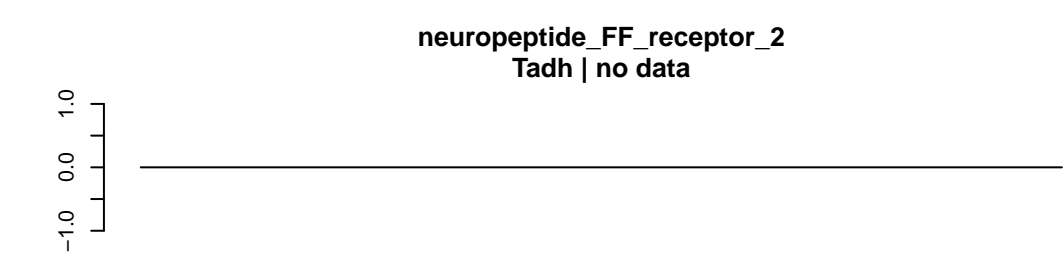


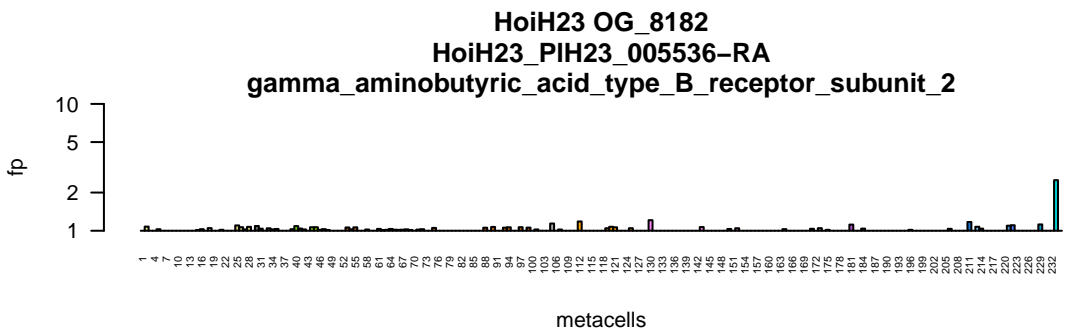
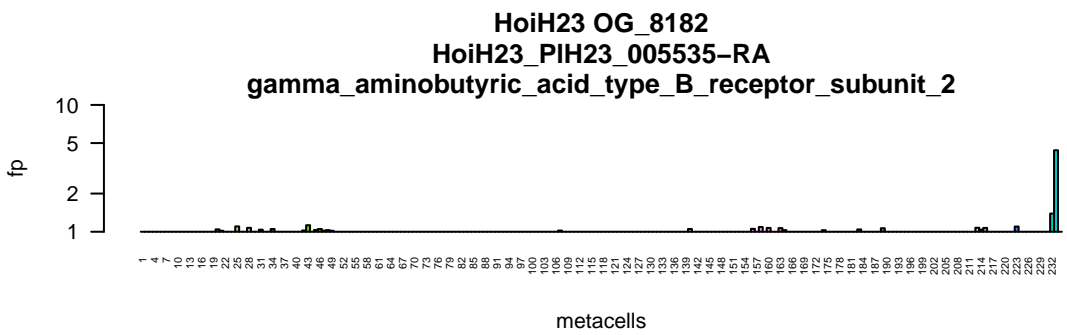
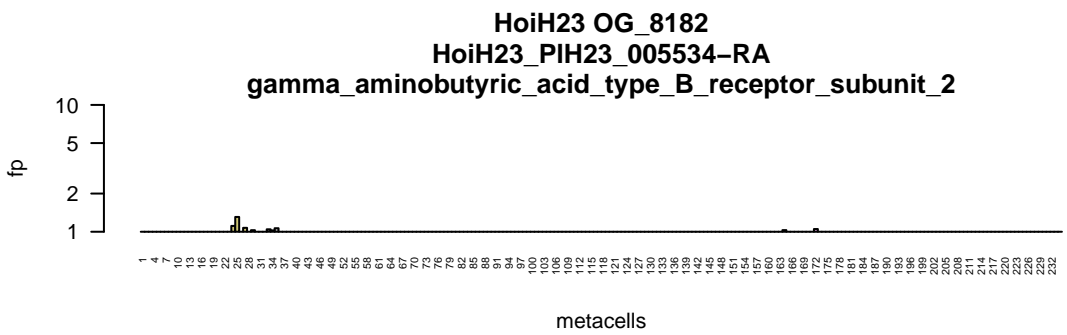
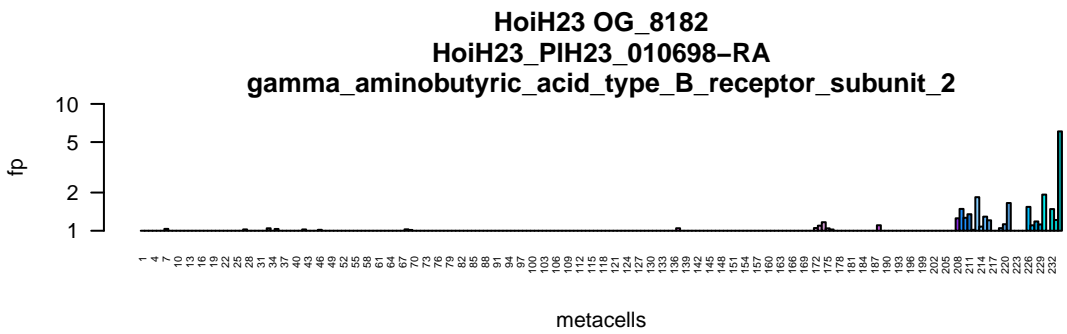
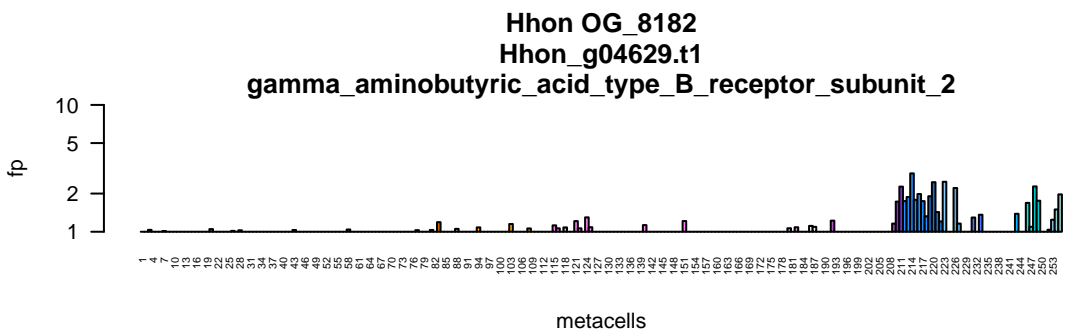
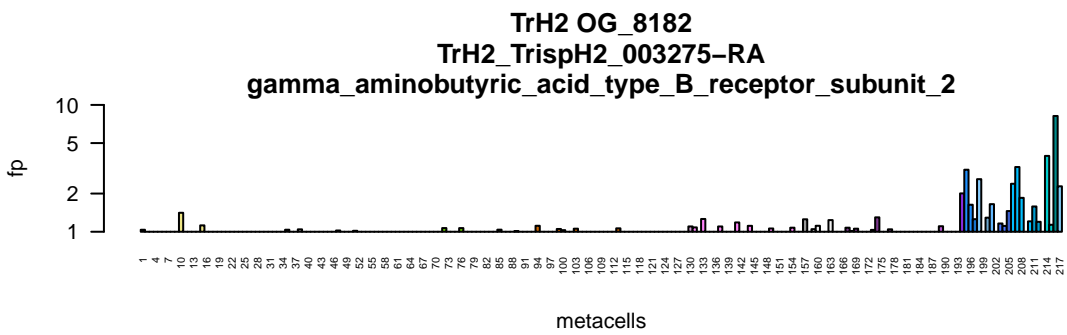
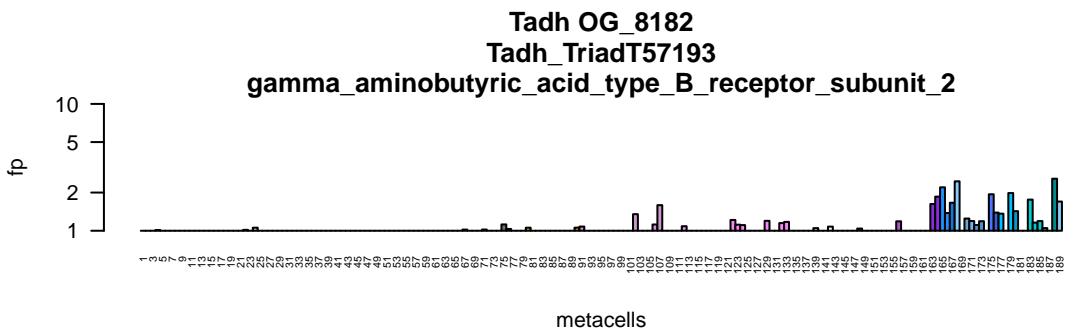
Hhon OG_6463
Hhon_g08668.t1
histamine_receptor_H2



HoiH23 OG_6463
HoiH23_PIH23_005854-RA
histamine_receptor_H2

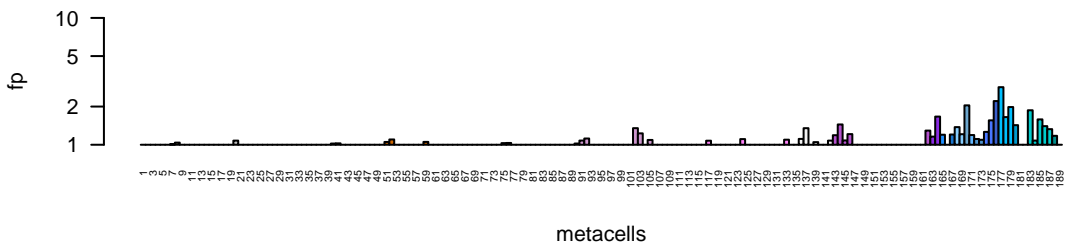






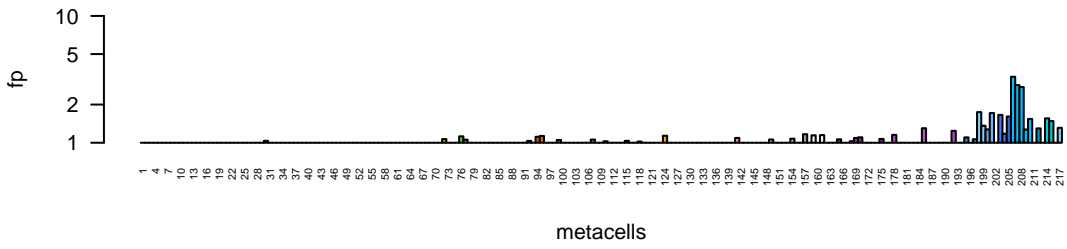
Tadh OG_8295

Tadh_TriadT54897



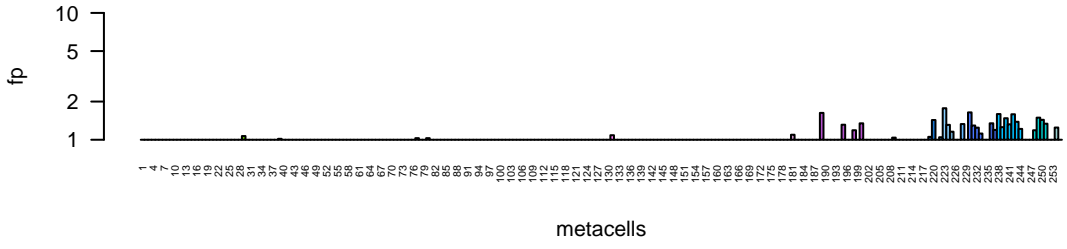
TrH2 OG_8295

TrH2_TrispH2_007497-RA



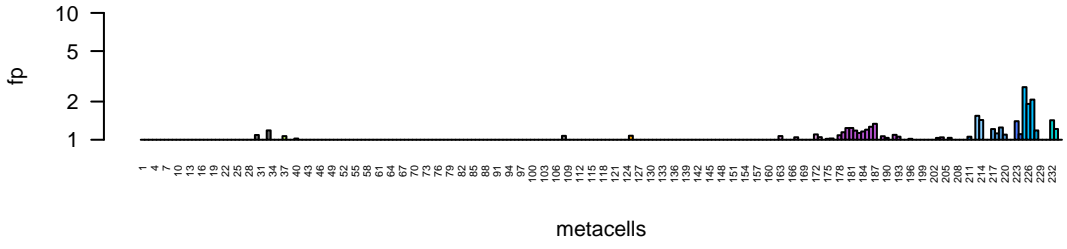
Hhon OG_8295

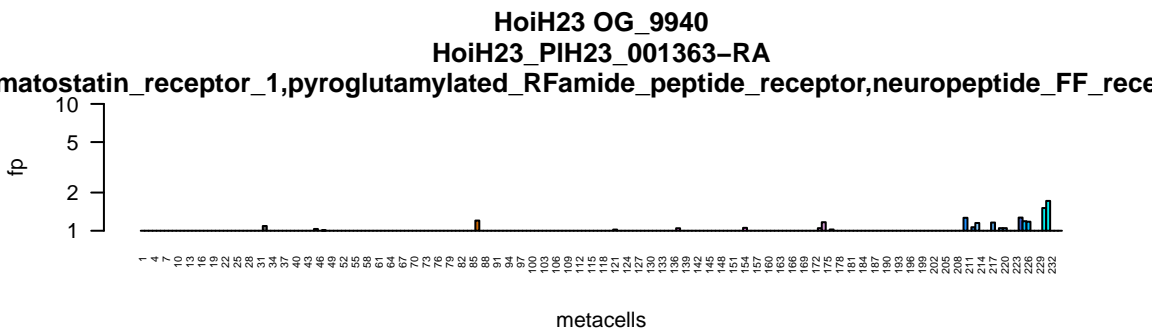
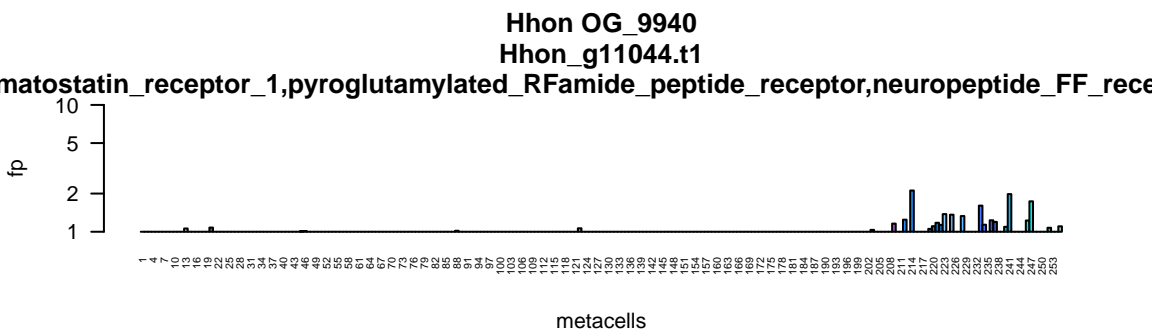
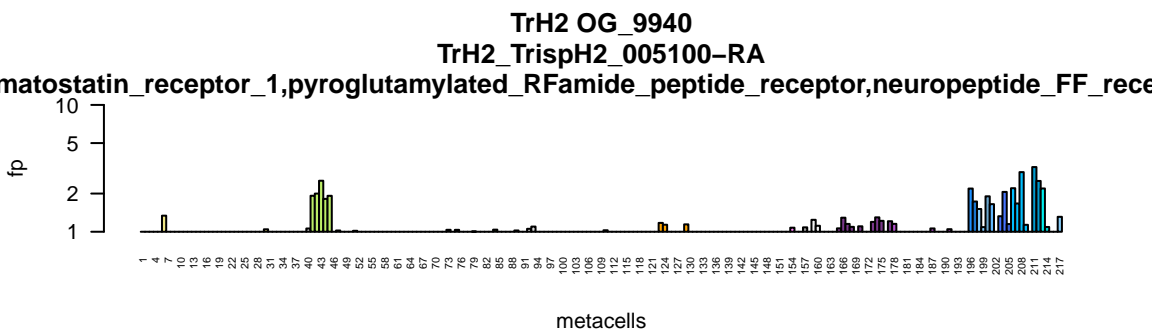
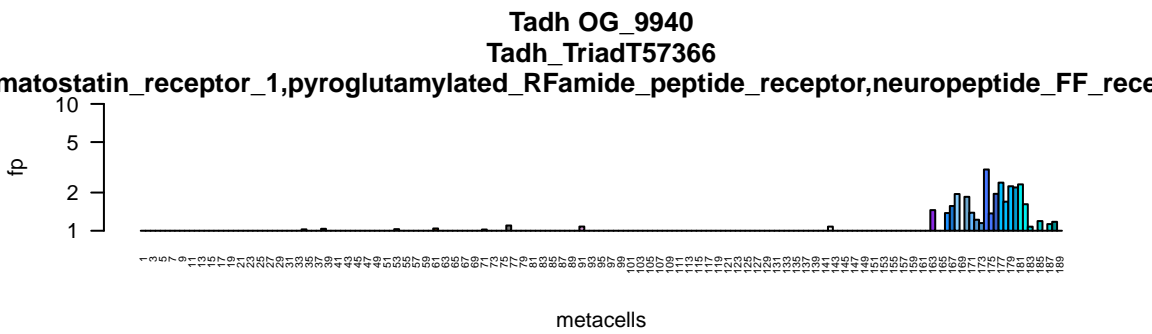
Hhon_g07101.t1



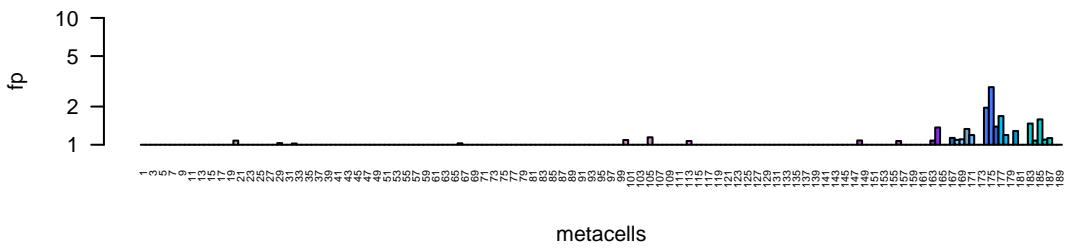
HoiH23 OG_8295

HoiH23_PIH23_002701-RA

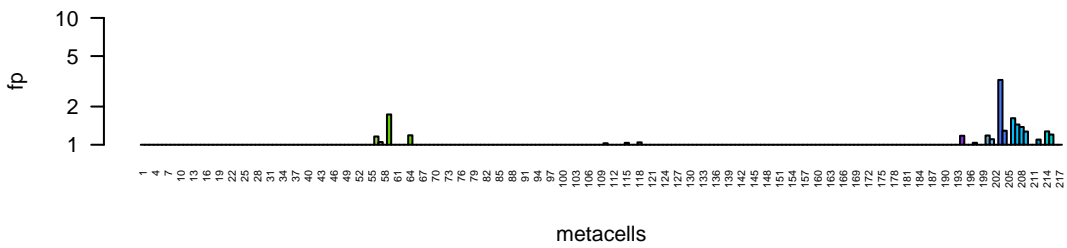




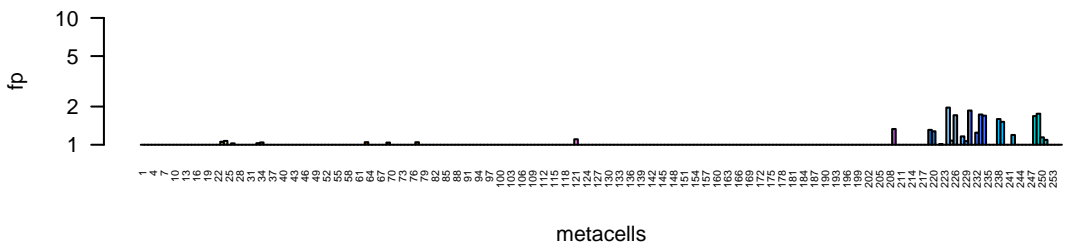
Tadh OG_515
Tadh_TriadT57467



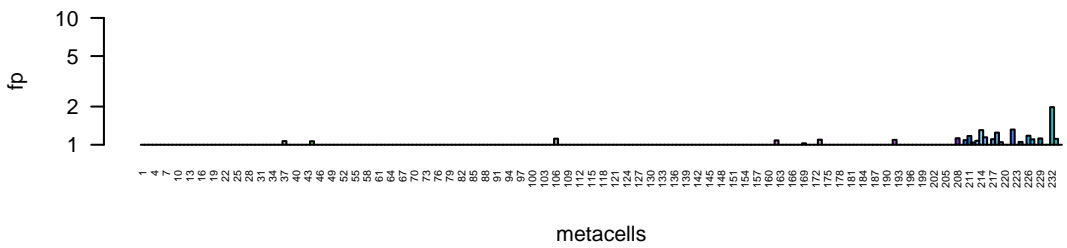
TrH2 OG_515
TrH2_TrispH2_007315-RA

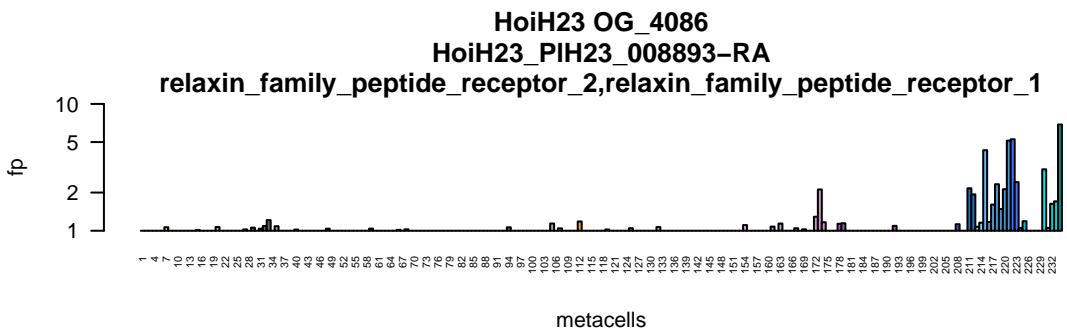
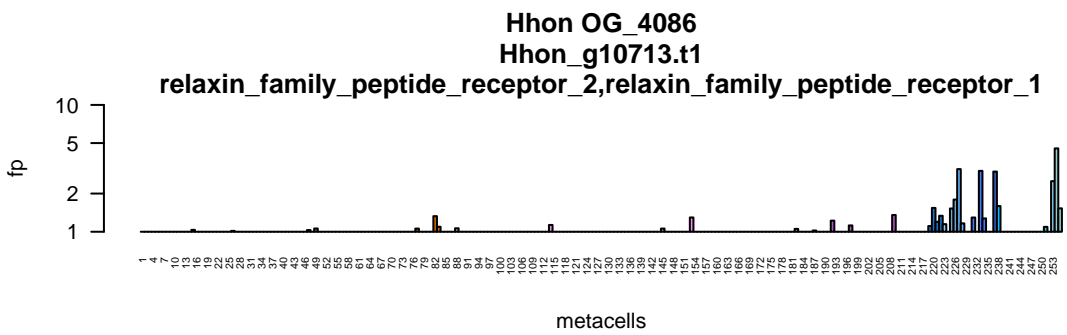
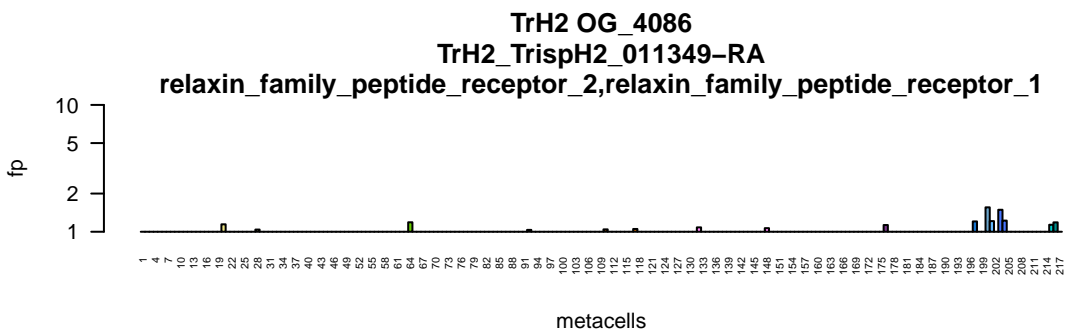
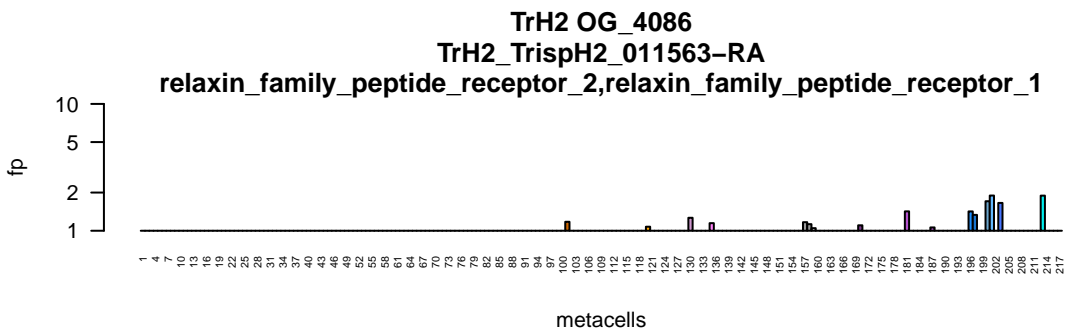
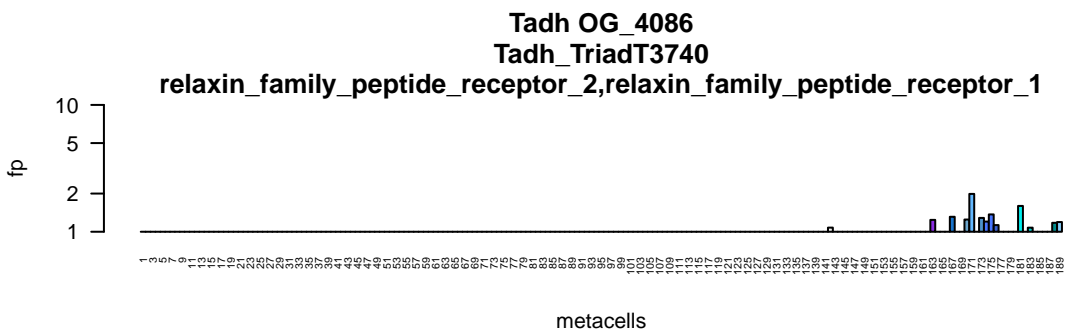
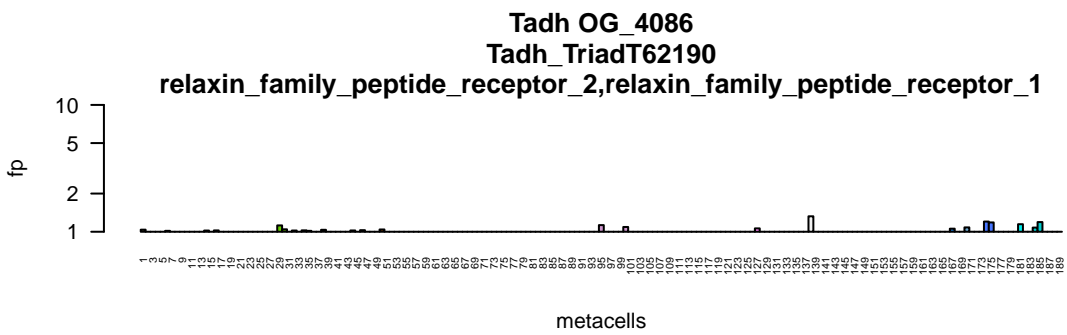
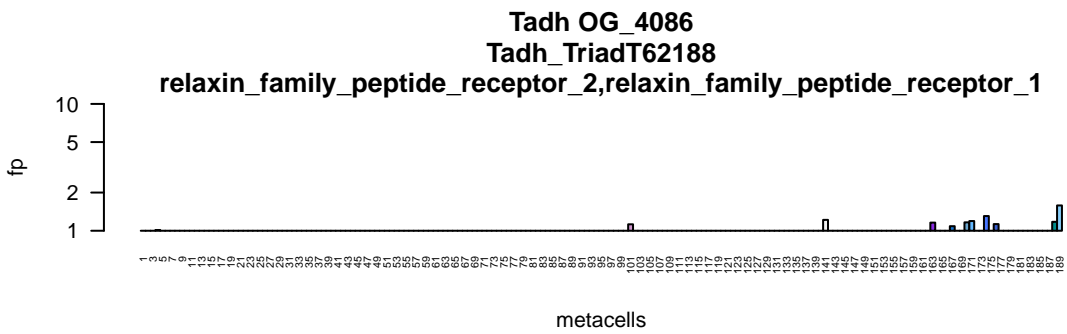


Hhon OG_515
Hhon_g01896.t1

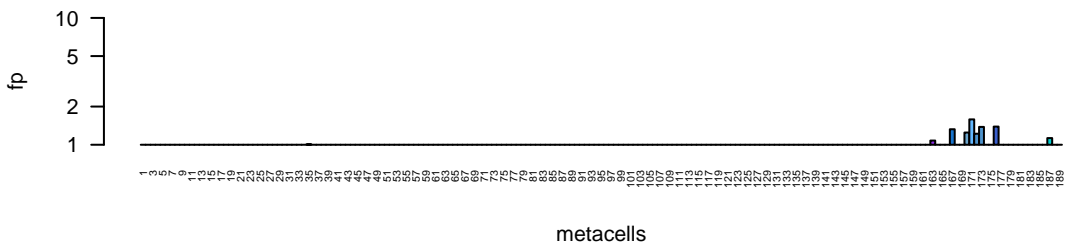


HoiH23 OG_515
HoiH23_PIH23_001462-RA

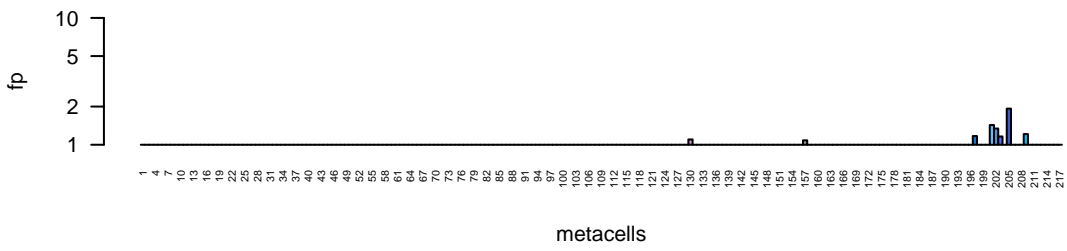




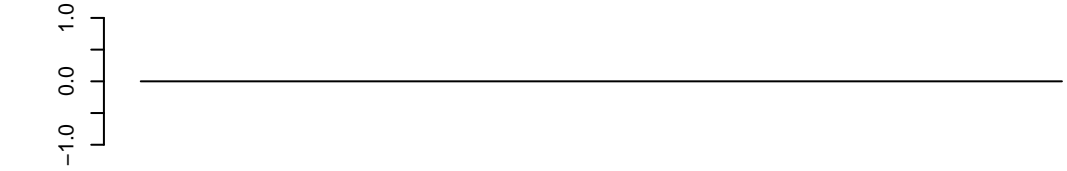
Tadh OG_4270
Tadh_wf_g4875.t1



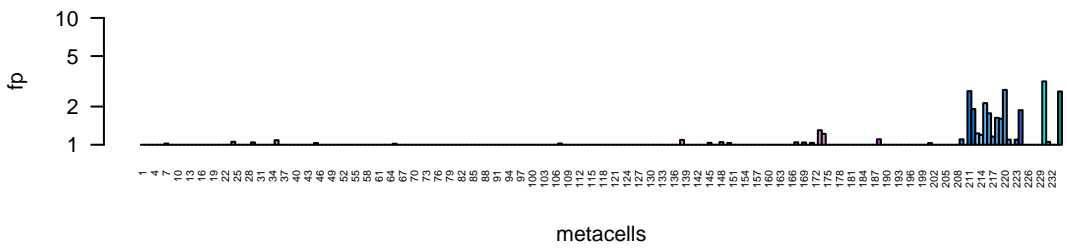
TrH2 OG_4270
TrH2_TrispH2_008489-RA



Hhon | no data



HoiH23 OG_4270
HoiH23_PIH23_009325-RA



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 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a notable peak at metacell 202 reaching a frequency of 4. Other metacells with frequency 1 include 13, 16, 19, 25, 28, 31, 34, 37, 40, 43, 46, 49, 52, 55, 58, 61, 64, 67, 70, 73, 76, 79, 82, 85, 88, 91, 94, 97, 100, 103, 106, 109, 112, 115, 118, 121, 124, 127, 130, 133, 136, 139, 142, 145, 148, 151, 154, 157, 160, 163, 166, 169, 172, 175, 178, 181, 184, 187, 190, 193, 196, 199, 202, 205, 208, 211, 214, and 217.

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacell	fp
1	1
4	1
7	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	1.2
58	1
61	1
67	1
70	1
73	1
76	1
79	1
85	1
86	1
89	1
94	1
97	1.2
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1.2
130	1
133	1
136	1
139	1
142	1
145	1.2
148	1
151	1.2
154	1
157	1.2
160	1.2
163	1
166	1
169	1
172	1
175	1.5
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	1.8
202	1.8
203	1.2
205	1.2
208	1
211	1
214	1.2
217	2

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacells	fp
1	1
4	1
7	1
13	1
16	1
19	1
25	1
28	1
31	1
34	1
40	1
43	1
46	1
52	1
55	1
58	1
61	1
67	1
70	1
73	1
76	1
79	1
85	1
86	1
91	1
94	1
97	1
100	1
103	1
106	1
109	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	1
157	1
160	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1
193	1
196	1
199	2
202	2.5
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

metacells	fp
1	1
4	1
13	1
16	1
17	1
25	1
26	1
28	1
31	1
34	1
40	1
43	1
46	1
52	1
55	1
58	1
61	1
67	1
70	1
73	1
75	1
79	1
85	1
86	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
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	1
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

metacells	fp
1	1
4	1
7	1
13	1
16	1
19	1
25	1
28	1
31	1
34	1
40	1
43	1
46	1
52	1
55	1
59	1
61	1
67	1
70	1
73	1
76	1
79	1
85	1
86	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	1
217	2

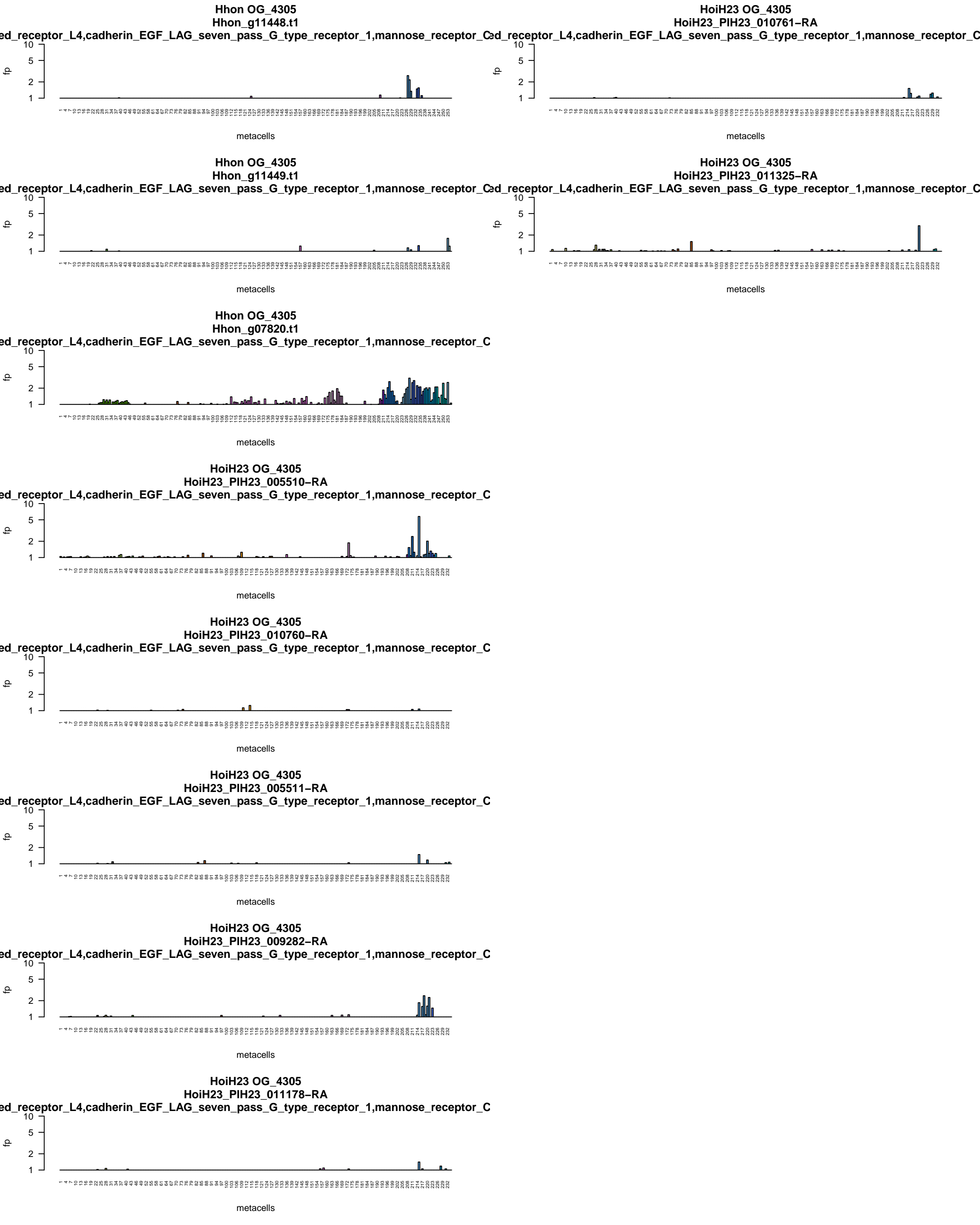
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

fp

metacells

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

metacells	fp
1	1
4	1
10	1
13	1
19	1
22	1
28	1
31	1
37	1
40	1
46	1
52	1
55	1
64	1
68	1
70	1
73	1
79	1
88	1
91	1
97	1
100	1
106	1
109	1
115	1
118	1
127	1
124	1
133	1
142	1
145	1
151	1
154	1
163	1
169	1
172	1
178	1
184	1
187	1
193	1
196	1
202	1
205	1
211	1
214	1
220	1
223	1
229	1
232	1
238	1
241	1
247	1
250	1
253	2



HoiH23 OG_4305

HoiH23_PIH23_010760-RA

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



metacells

HoiH23 OG_4305

HoiH23_PIH23_005511-RA

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



metacells

HoiH23 OG_4305

HoiH23_PIH23_009282-RA

ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



metacells

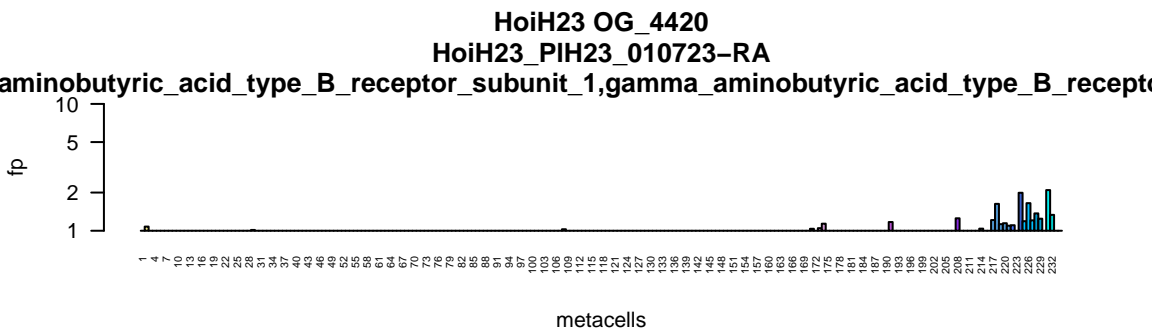
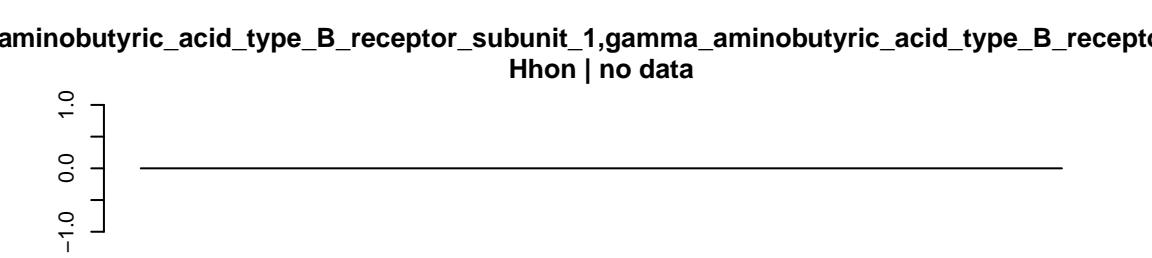
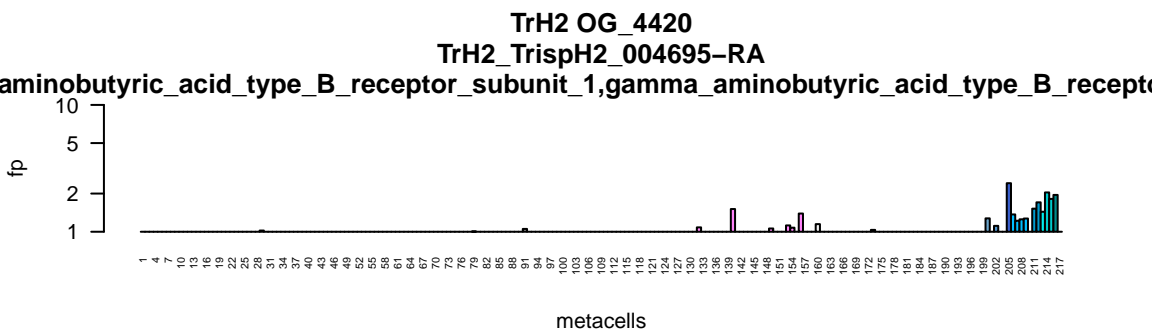
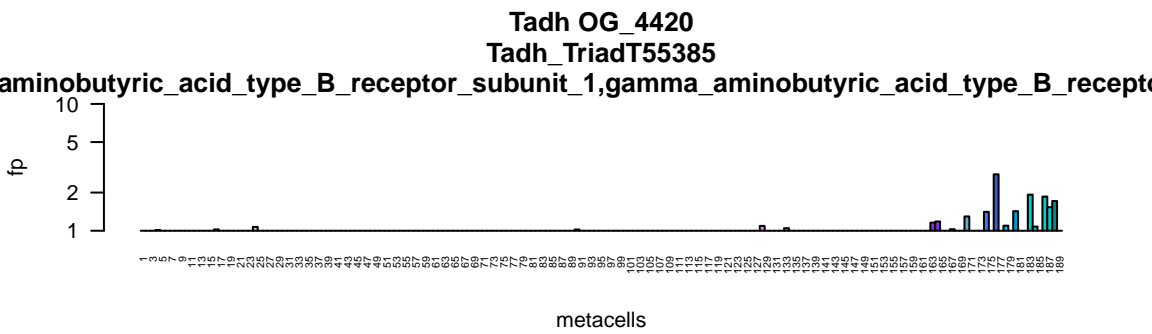
HoiH23 OG_4305

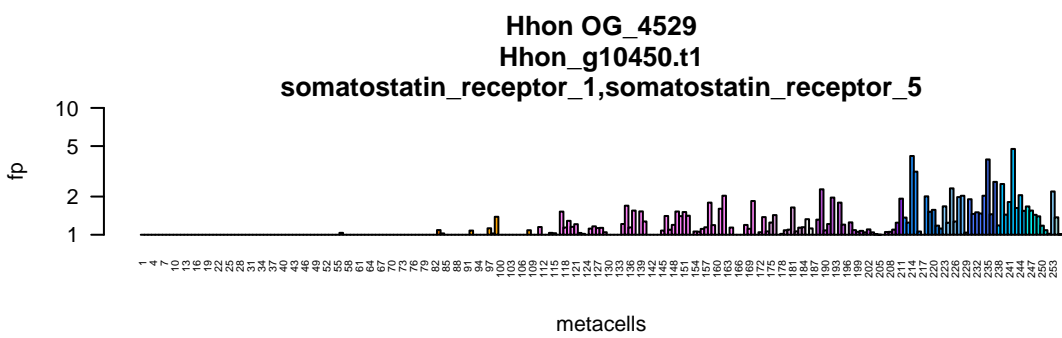
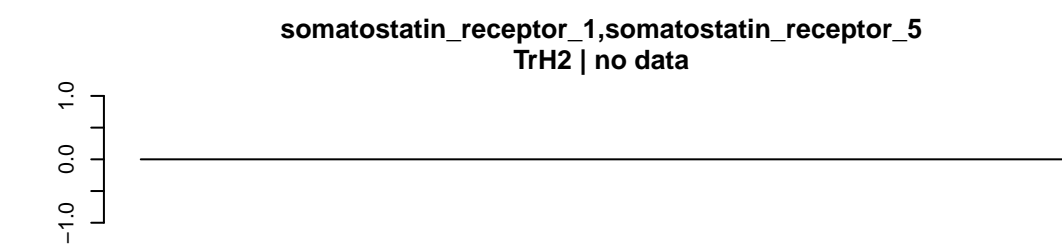
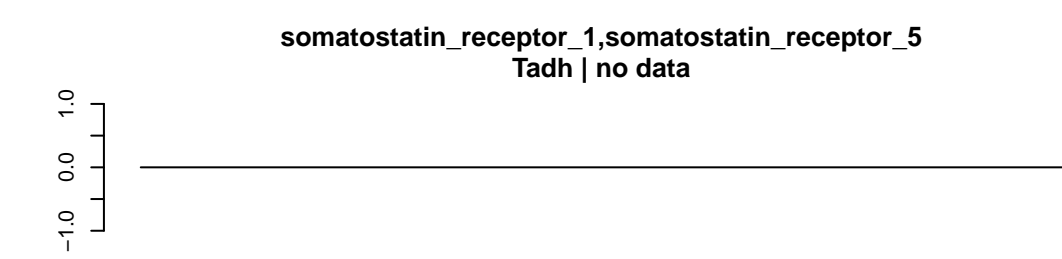
HoiH23_PIH23_011178-RA

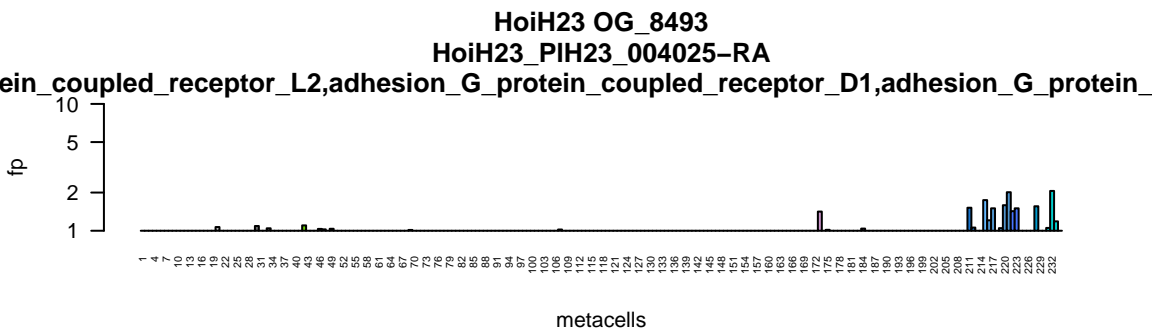
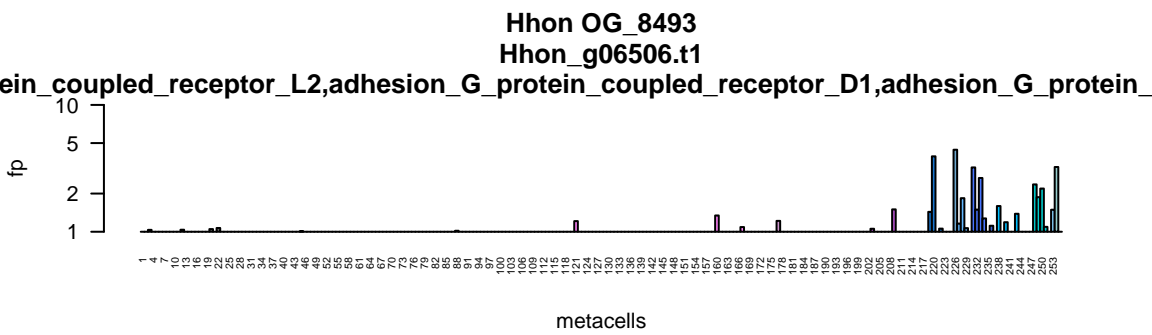
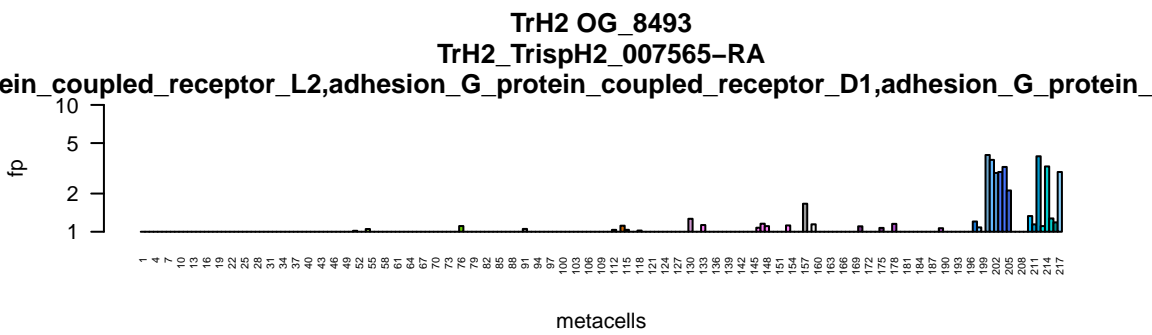
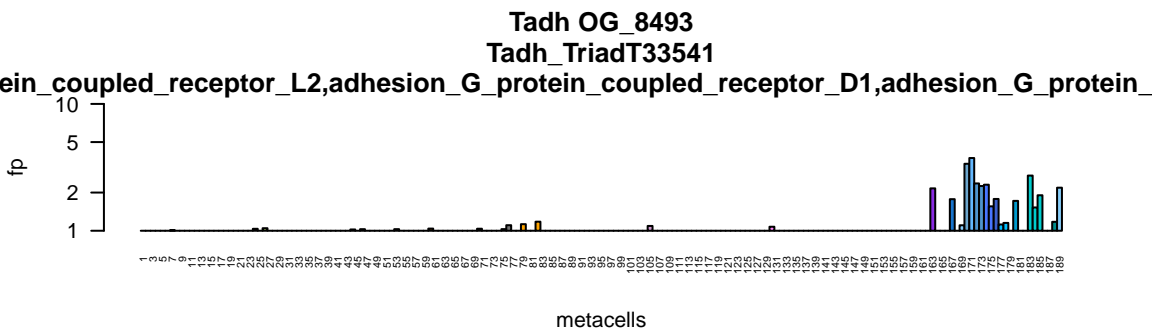
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

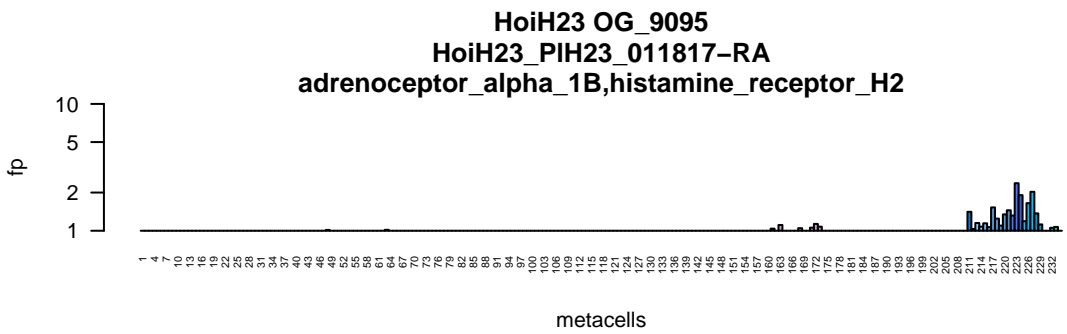
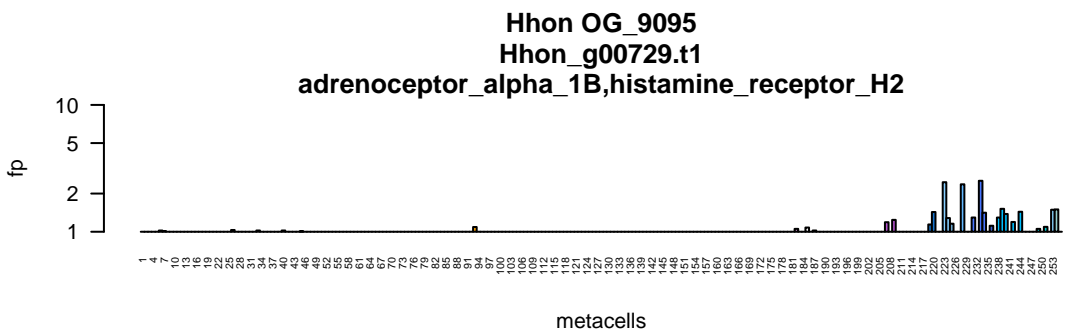
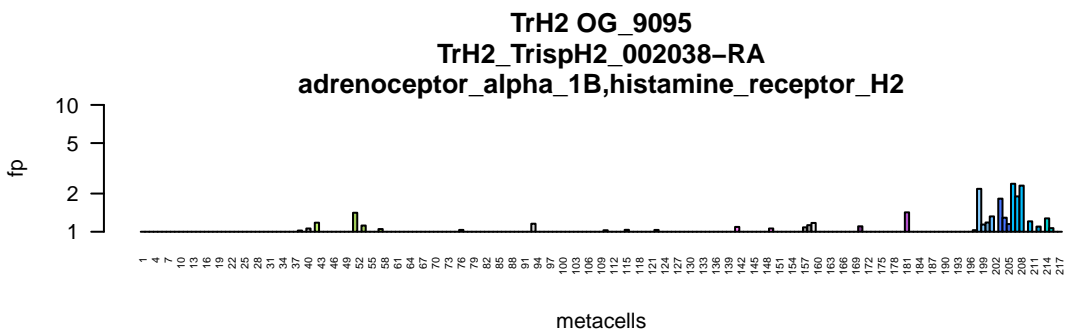
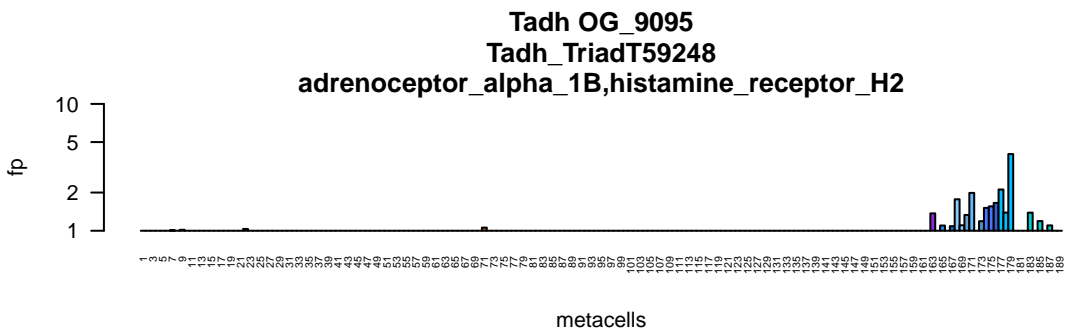


metacells





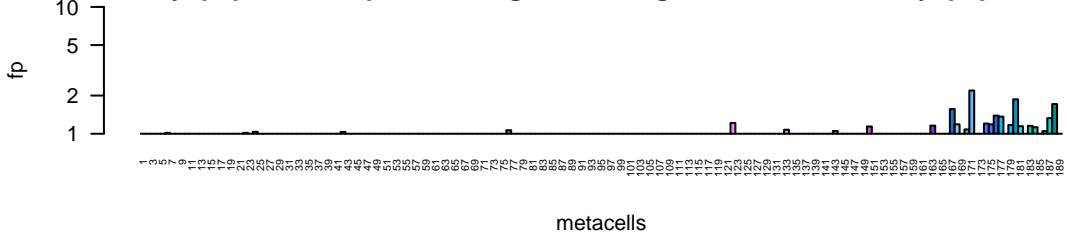




Tadh OG_9223

Tadh_TriadT51796

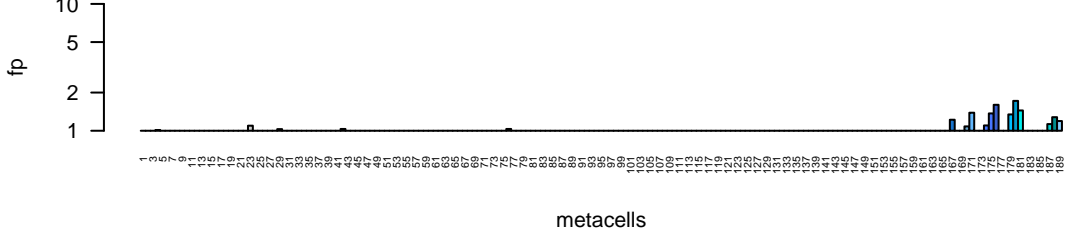
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Tadh OG_9223

Tadh_TriadT51797

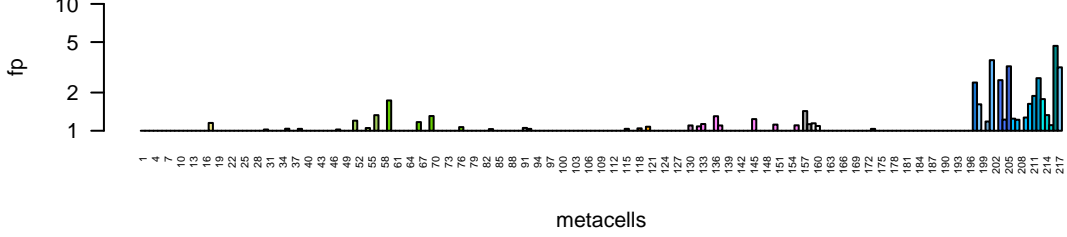
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



TrH2 OG_9223

TrH2_TrispH2_008387-RA

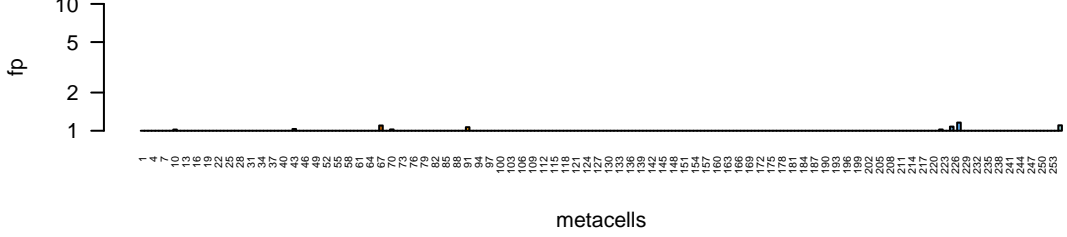
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



Hhon OG_9223

Hhon_g11380.t1

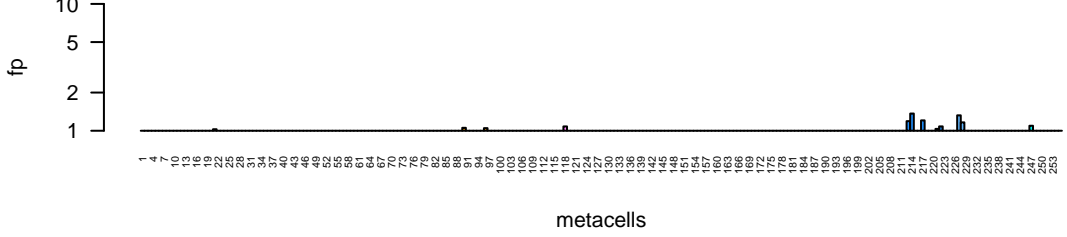
relaxin_family_peptide_receptor_1,slit_guidance_ligand_1,relaxin_family_peptide_recepto



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

