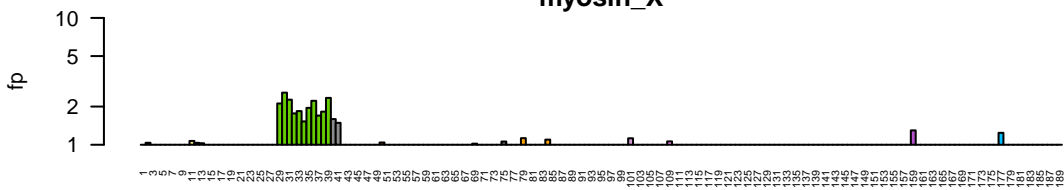
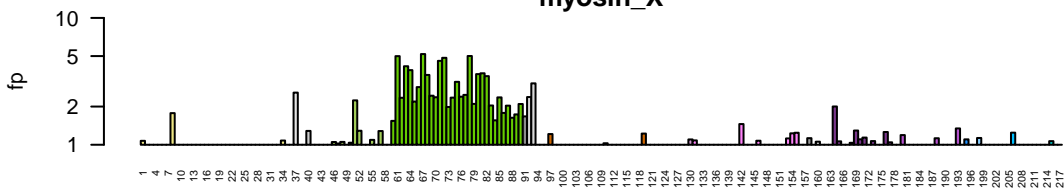


Tadh OG_7184
Tadh_TriadT54062
myosin_X



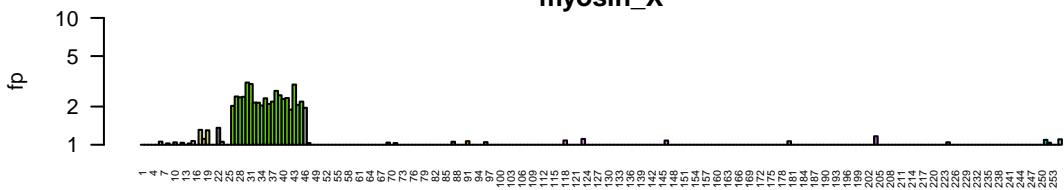
metacells

TrH2 OG_7184
TrH2_TrispH2_004795-RA
myosin_X



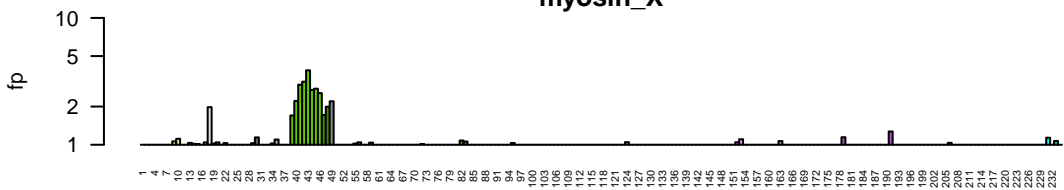
metacells

Hhon OG_7184
Hhon_g02820.t1
myosin_X

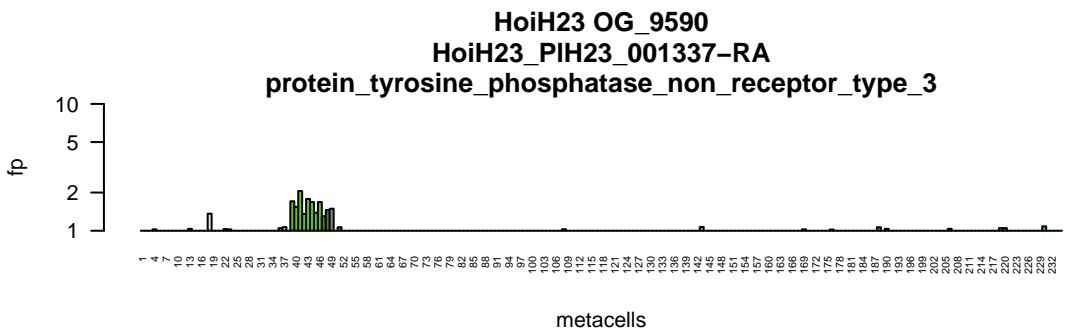
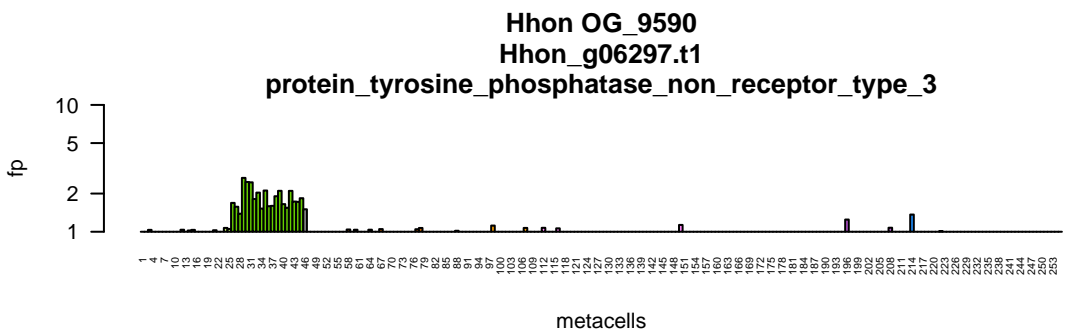
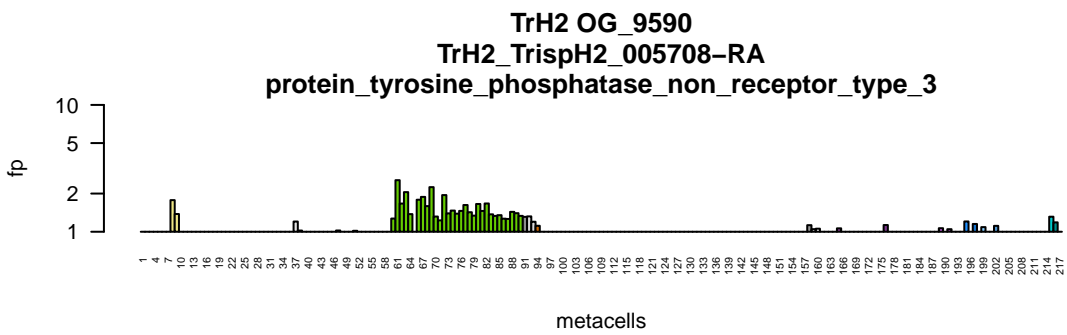
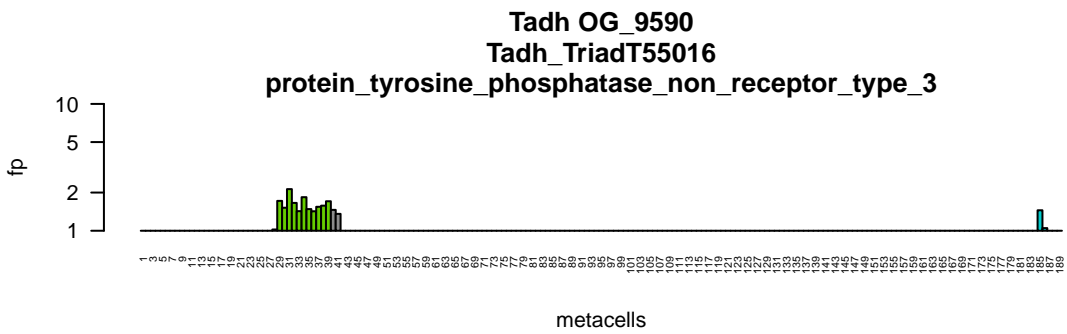


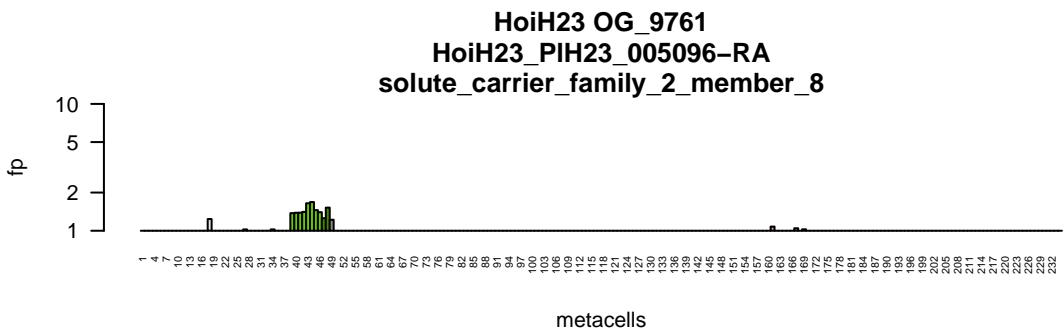
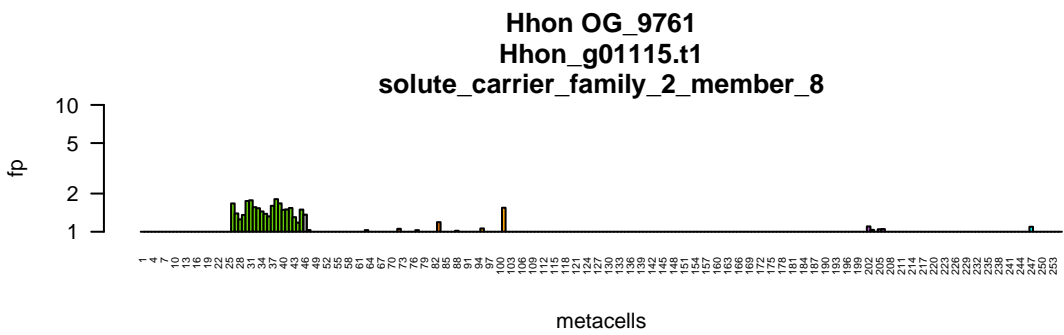
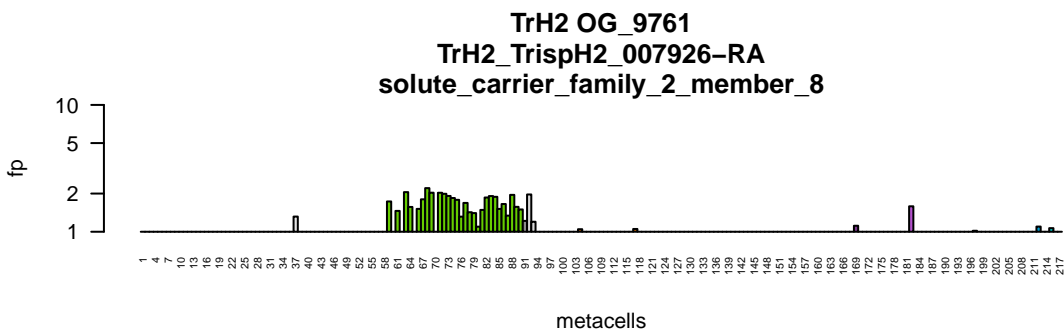
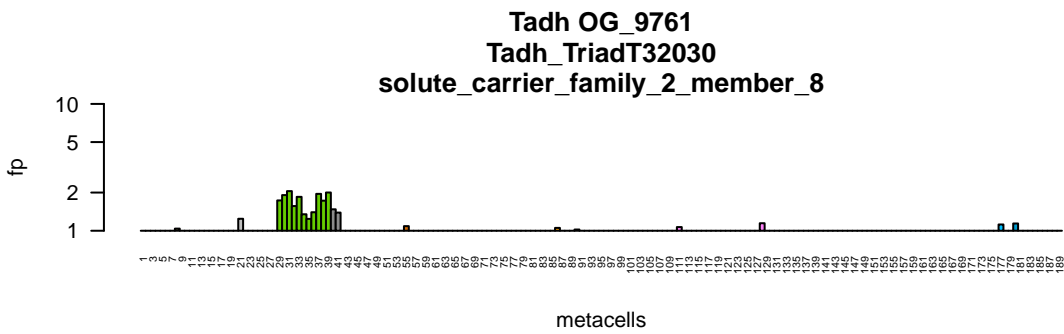
metacells

HoiH23 OG_7184
HoiH23_PIH23_009900-RA
myosin_X

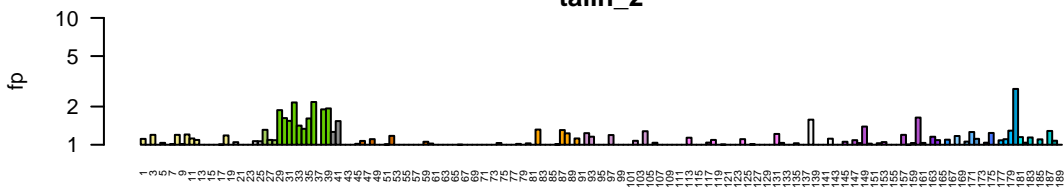


metacells



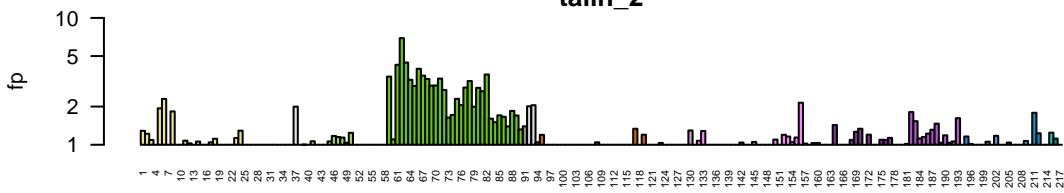


Tadh OG_10376
Tadh_TriadT19795
talin_2



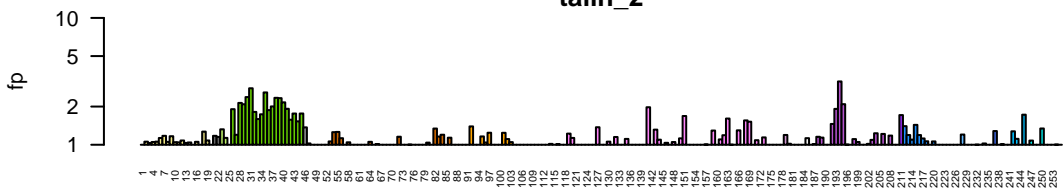
metacells

TrH2 OG_10376
TrH2_TrispH2_007006-RA
talin_2



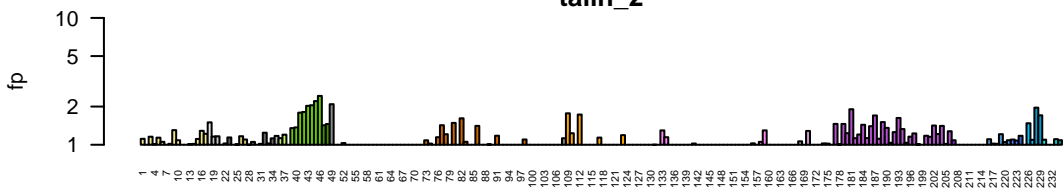
metacells

Hhon OG_10376
Hhon_g04057.t1
talin_2



metacells

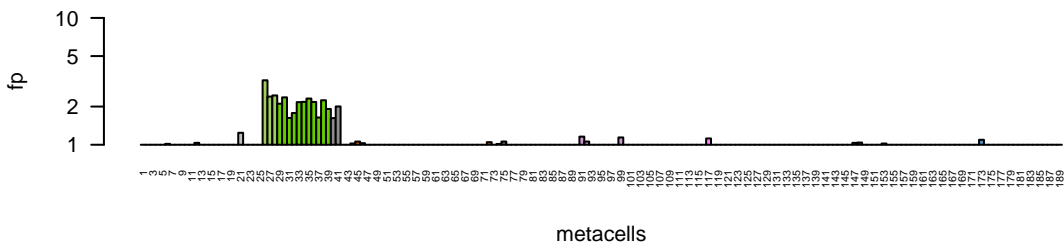
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HoiH23_PIH23_000825-RA
talin_2



metacells

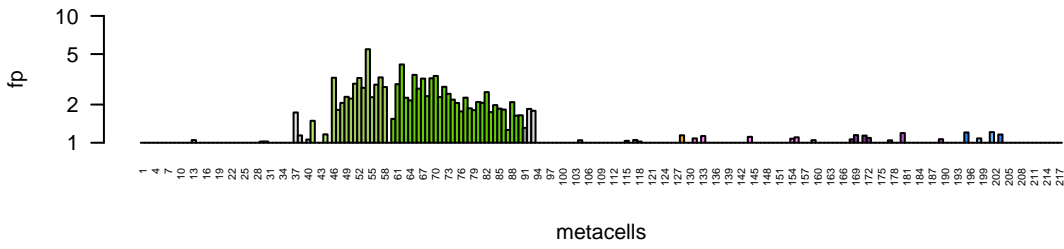
Tadh OG_1141

Tadh_TriadT54495



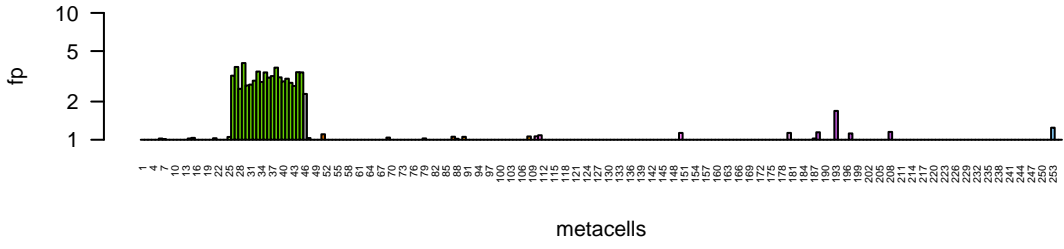
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TrH2_TrispH2_001223-RA



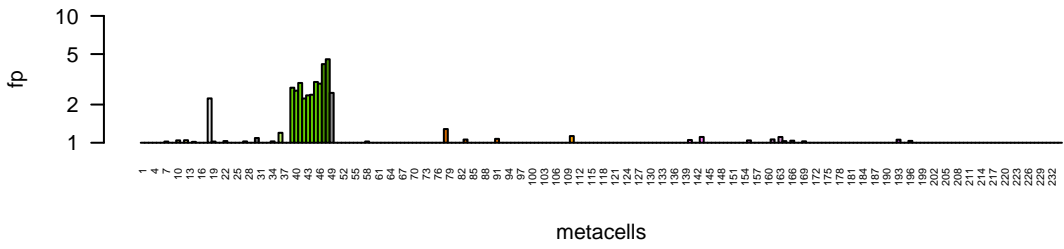
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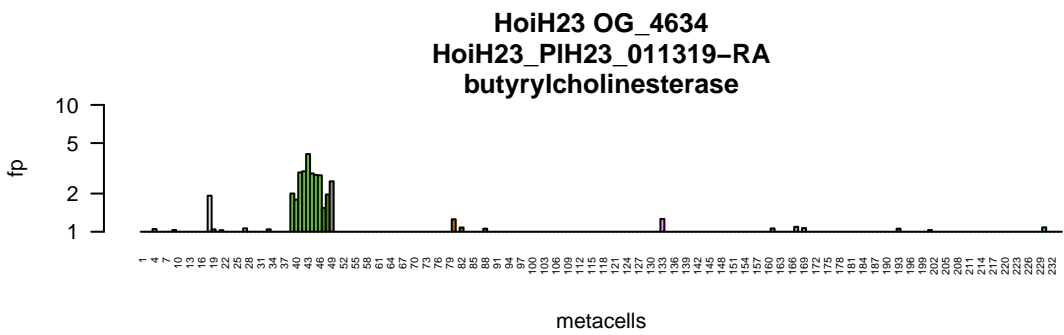
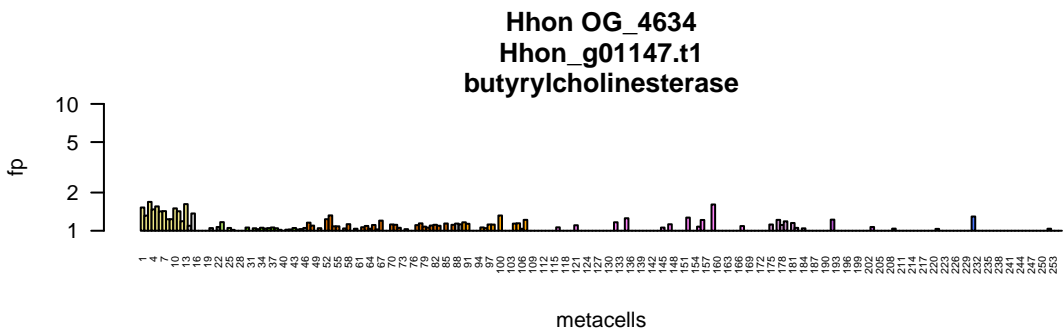
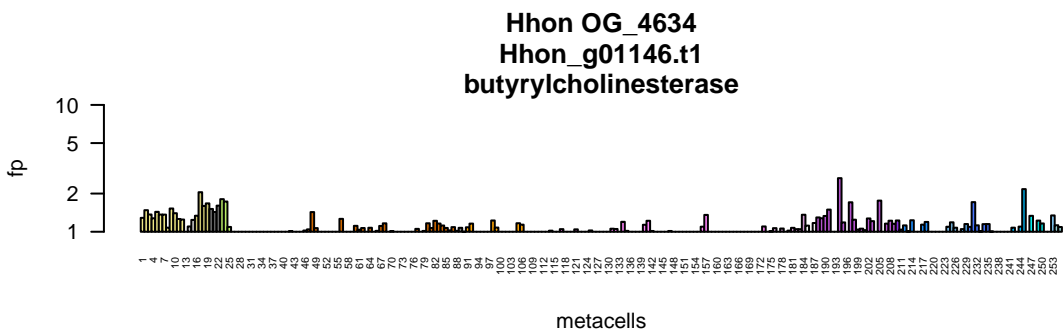
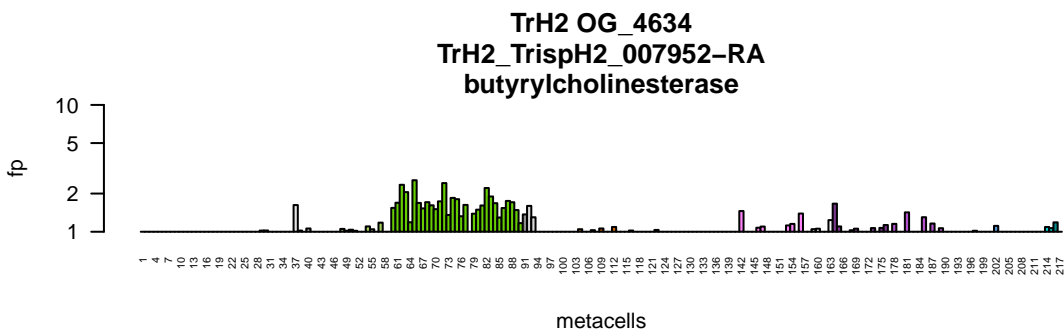
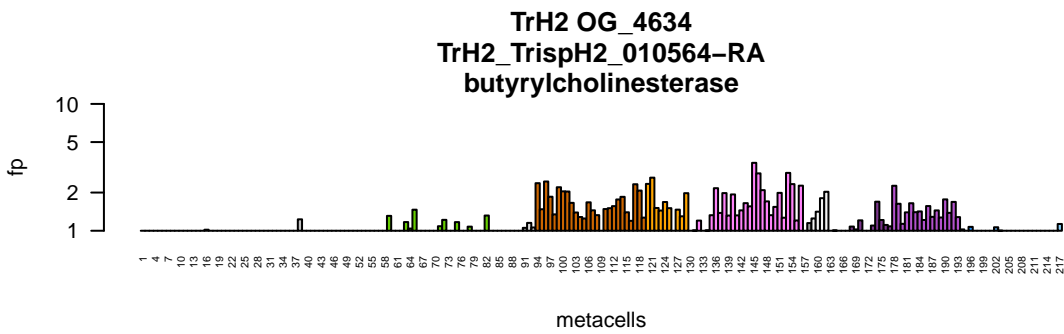
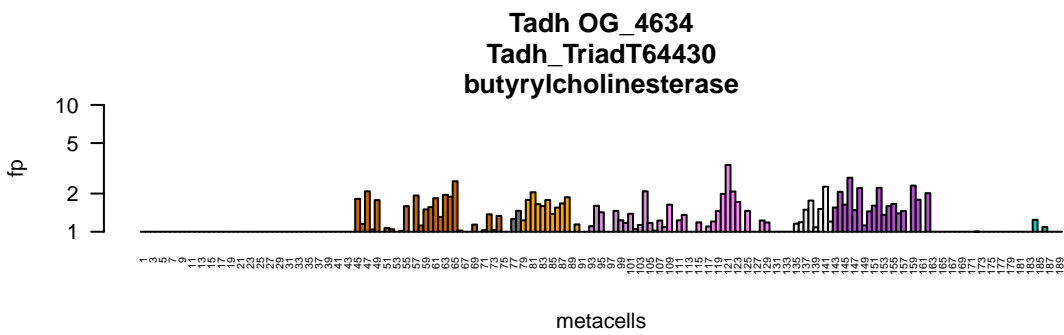
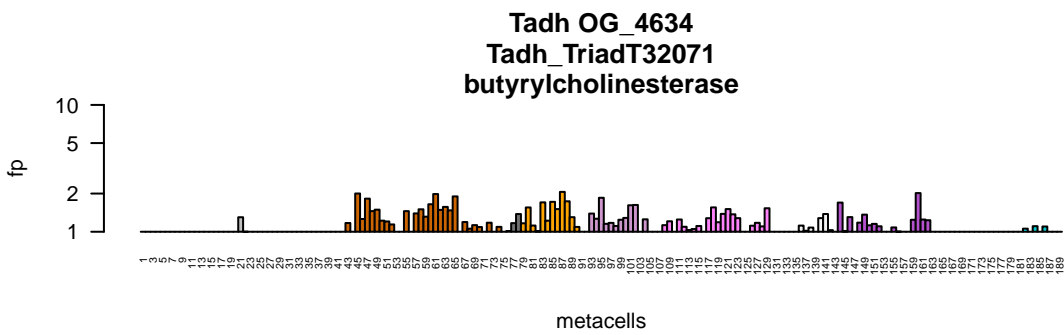
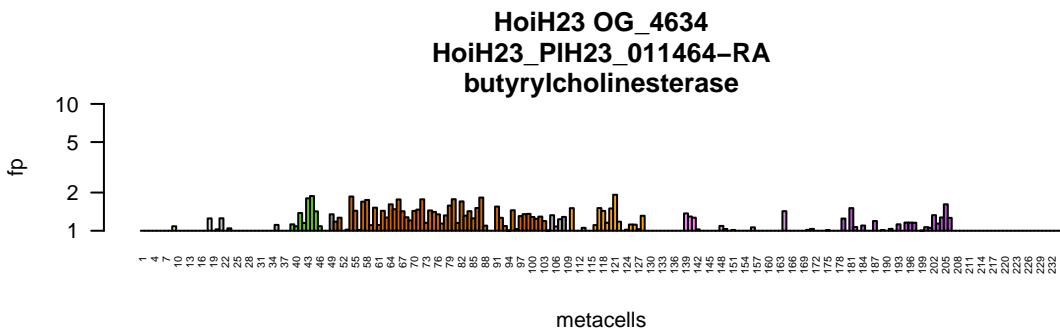
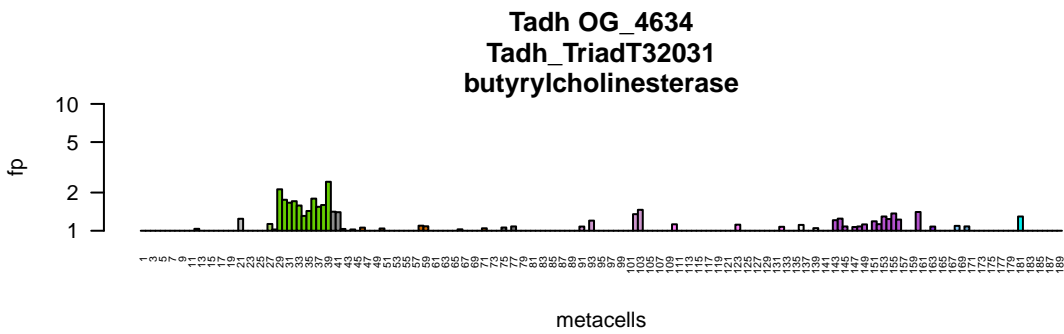
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HoiH23 OG_1141

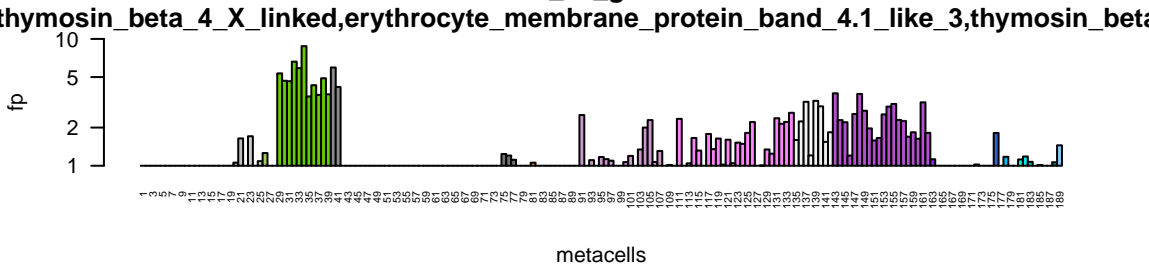
HoiH23_PIH23_003059-RA





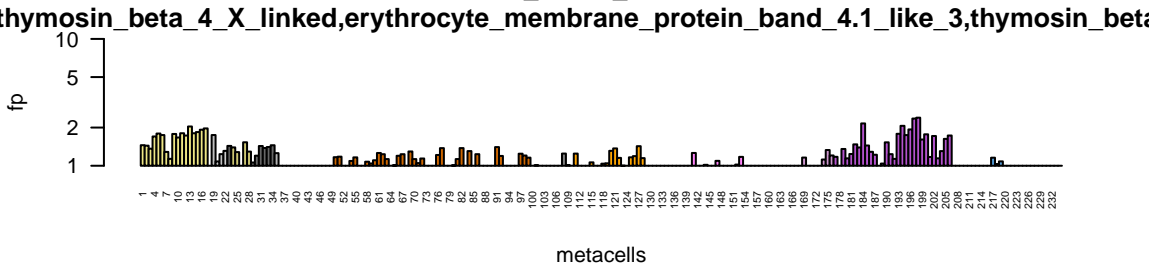
Tadh OG 5067

Tadh_wf_g5526.t1



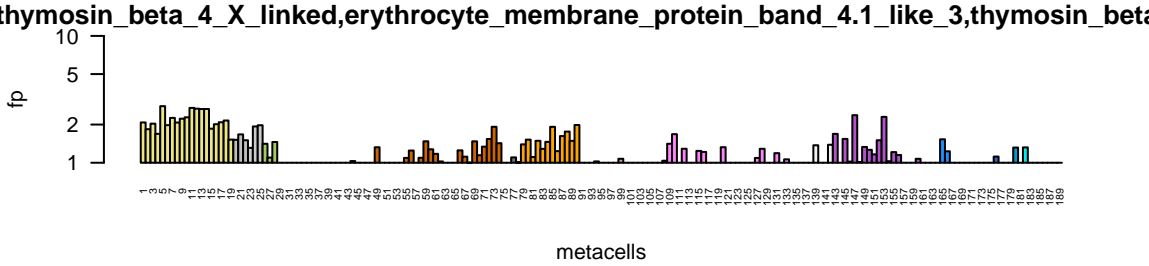
HoiH23 OG 5067

HoiH23_PIH23_001116-RA



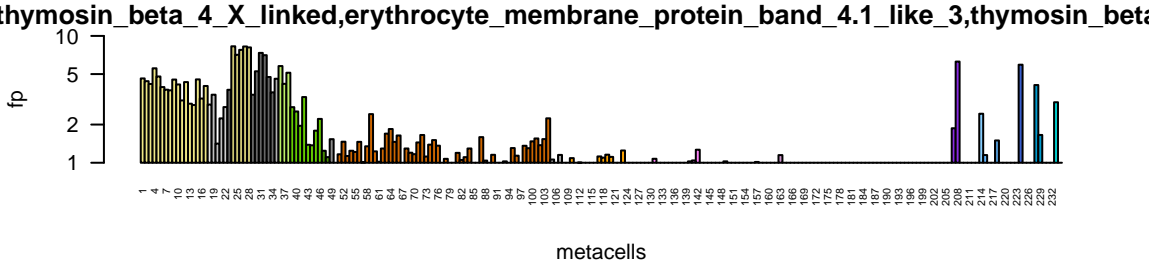
Tadh OG_5067

Tadh_TriadT64069



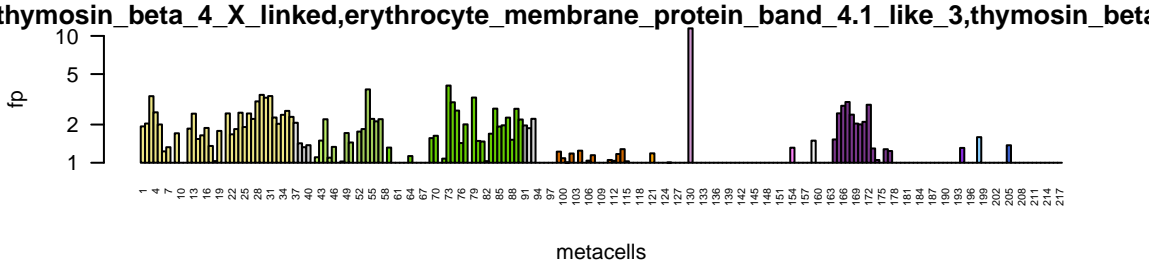
HoiH23 OG_5067

HoiH23_PIH23_006991-RA



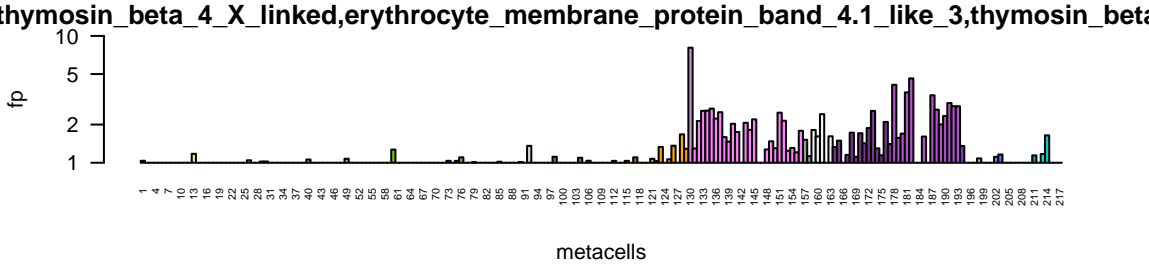
TrH2 OG_5067

TrH2_TrispH2_011135-RA



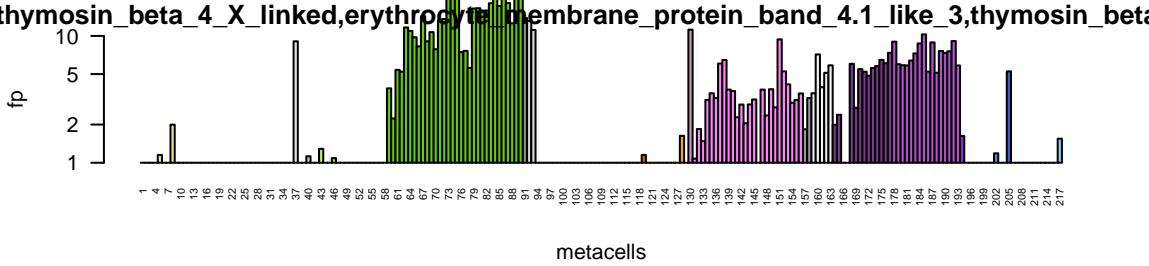
TrH2 OG_5067

TrH2_TrispH2_007984-RA



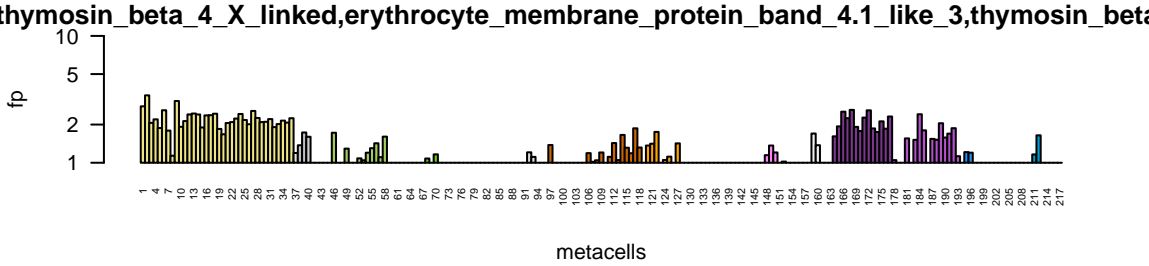
TrH2 OG_5067

TrH2_TrispH2_011134-RA



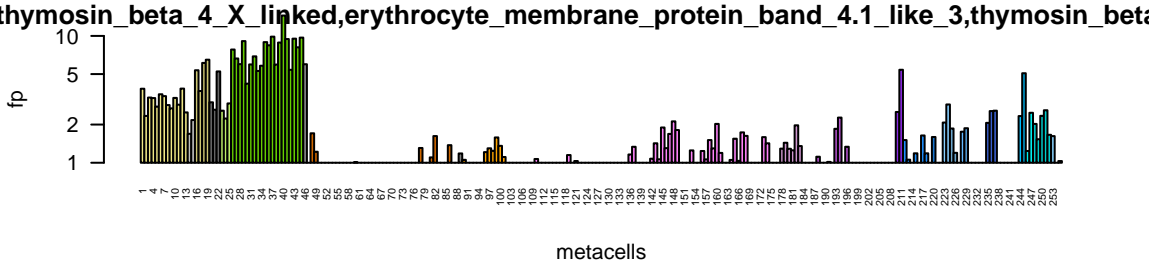
TrH2 OG_5067

TrH2_TrispH2_001493-RA



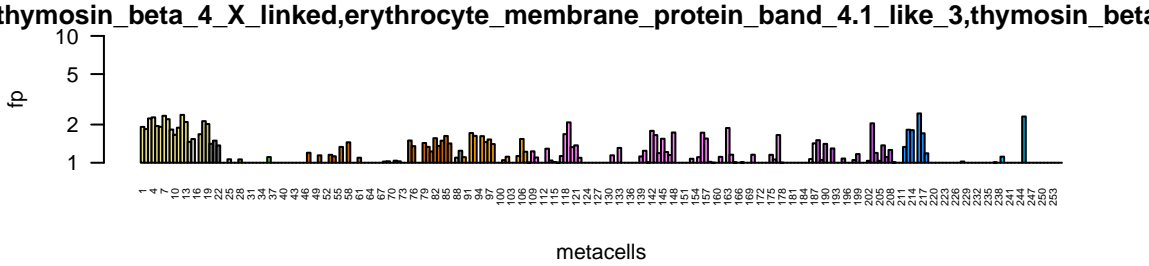
Hhon OG_5067

Hhon_g05621.t1

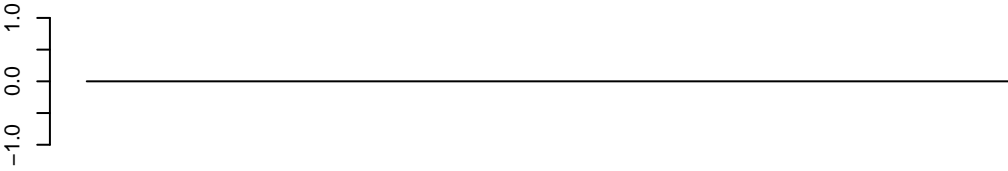


Hhon OG_5067

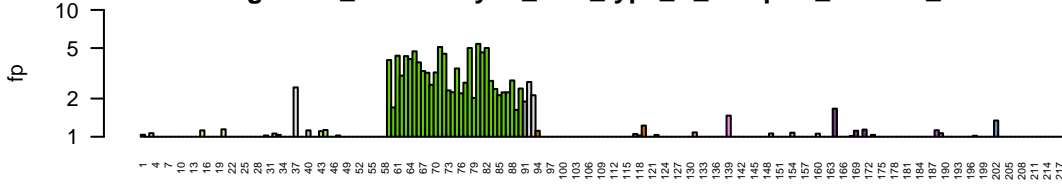
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gamma_aminobutyric_acid_type_B_receptor_subunit_2
Tadh | no data

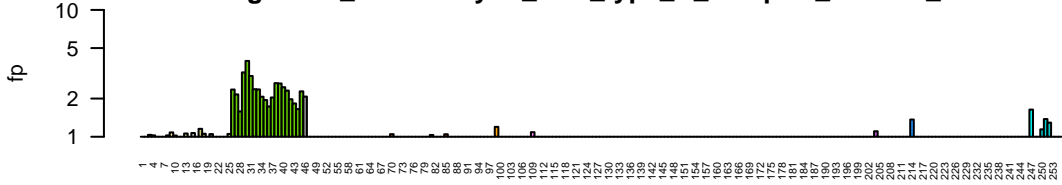


TrH2 OG_7215
TrH2_TrispH2_007186-RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2



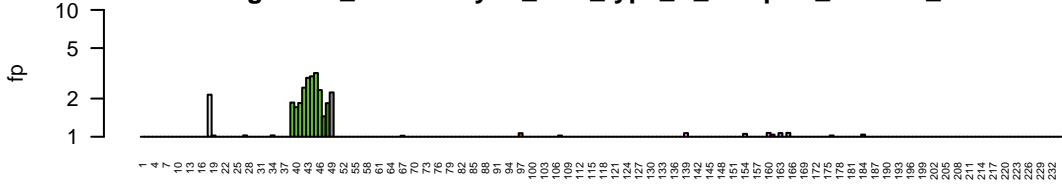
metacells

Hhon OG_7215
Hhon_g06834.t1
gamma_aminobutyric_acid_type_B_receptor_subunit_2

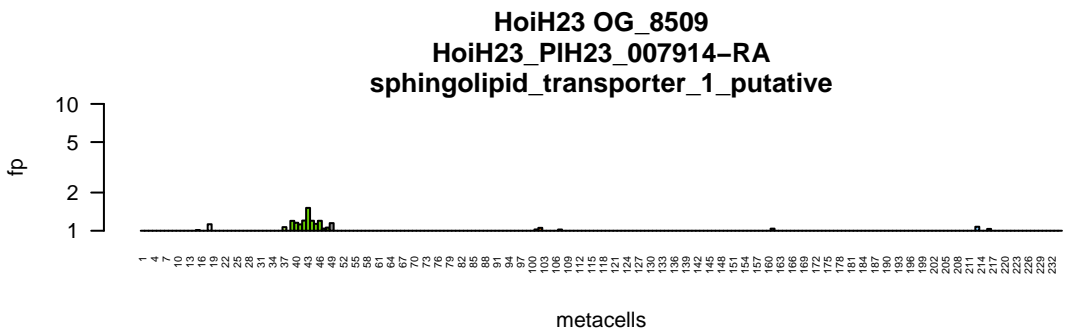
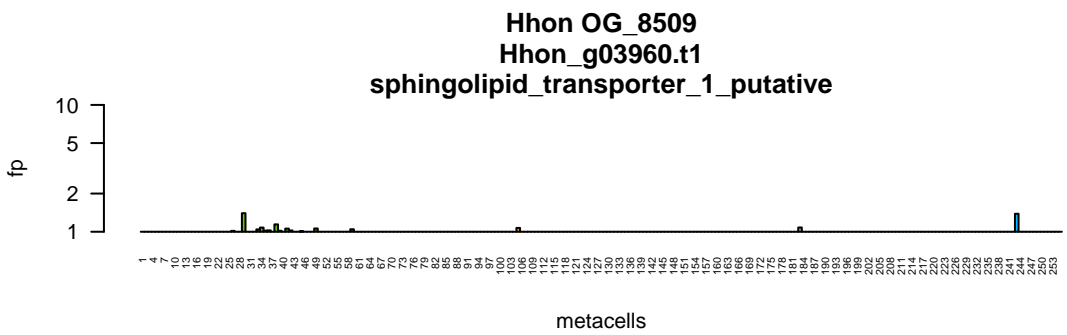
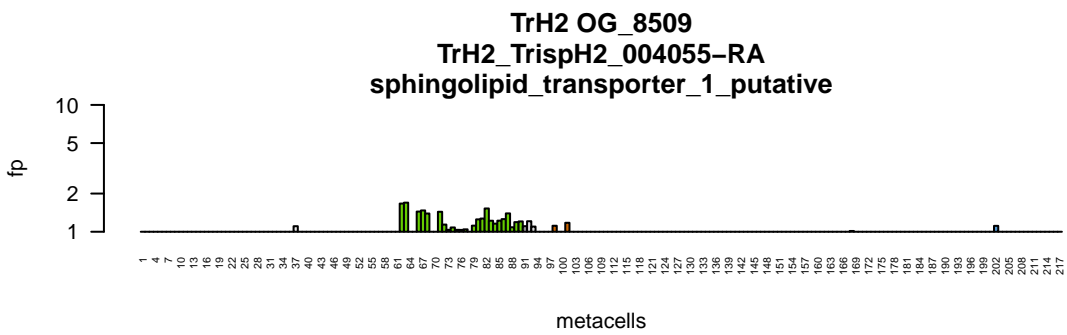
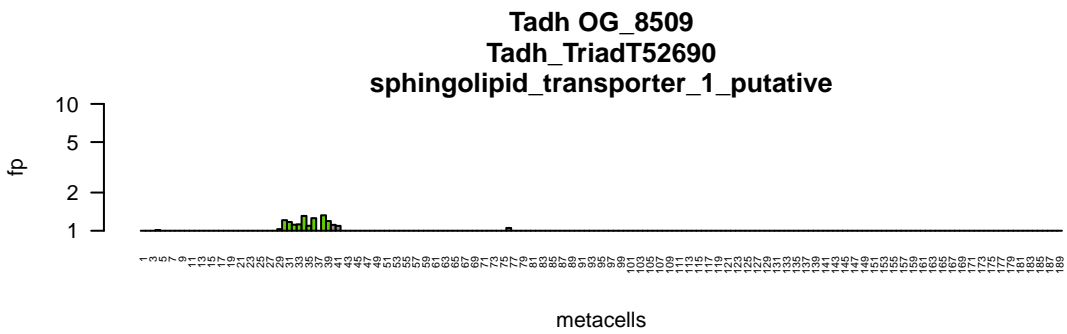


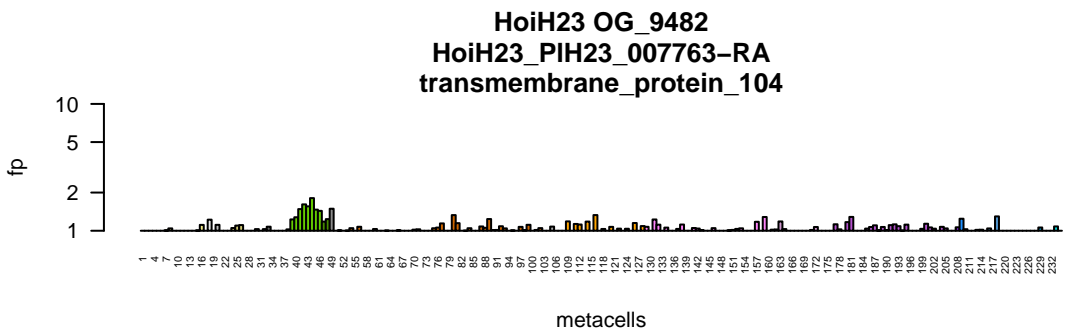
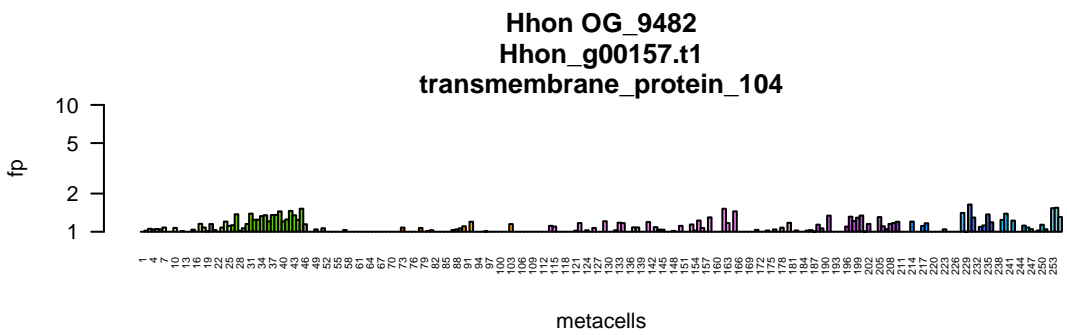
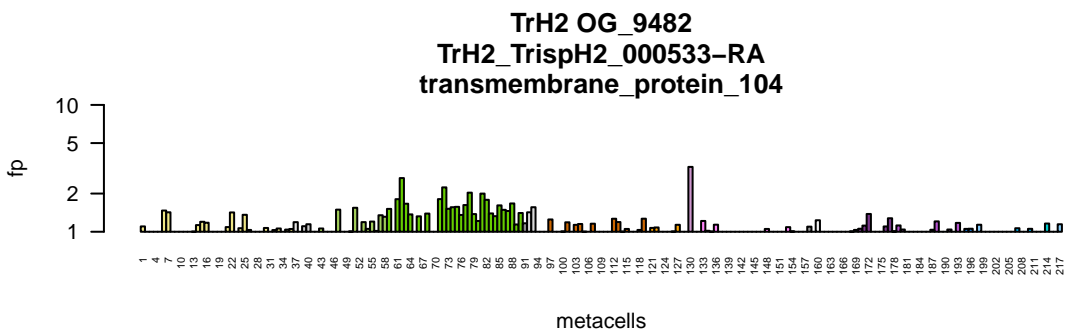
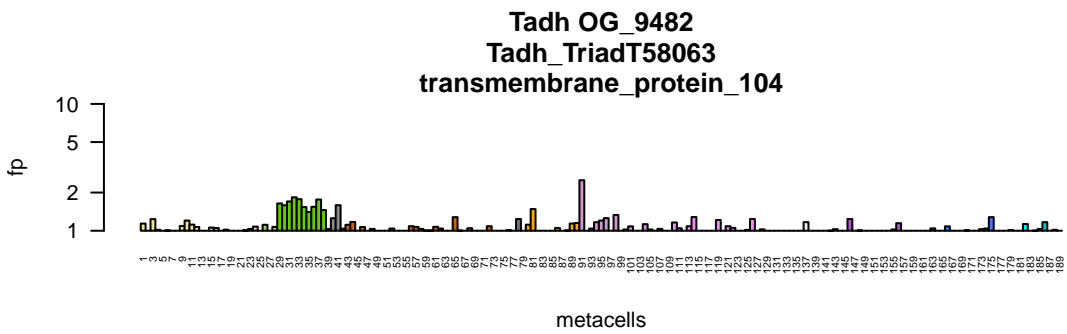
metacells

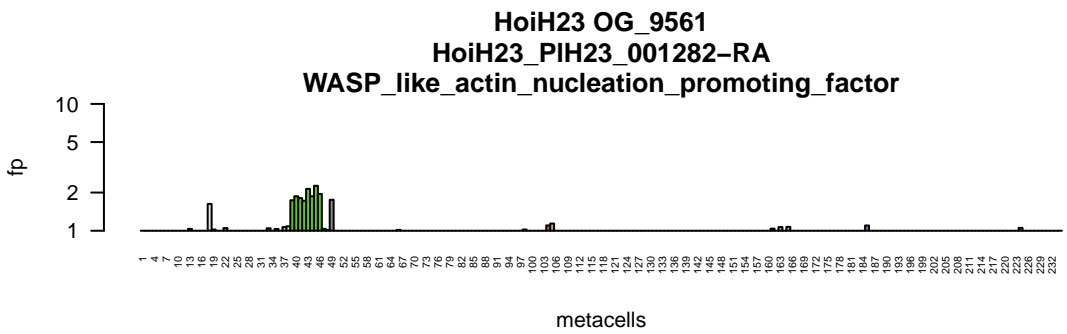
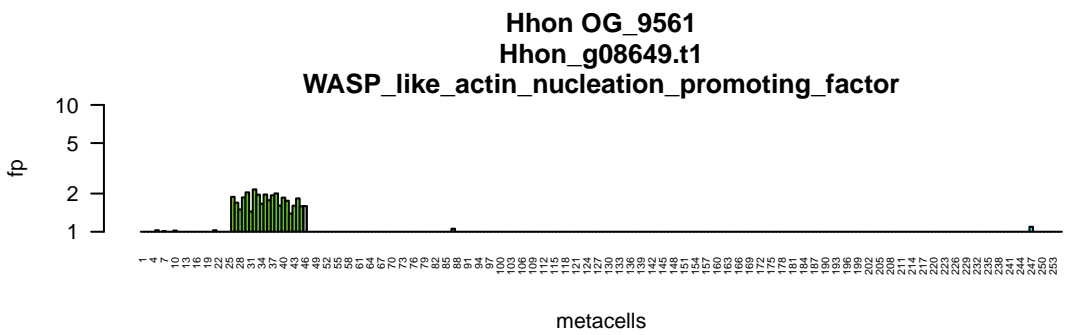
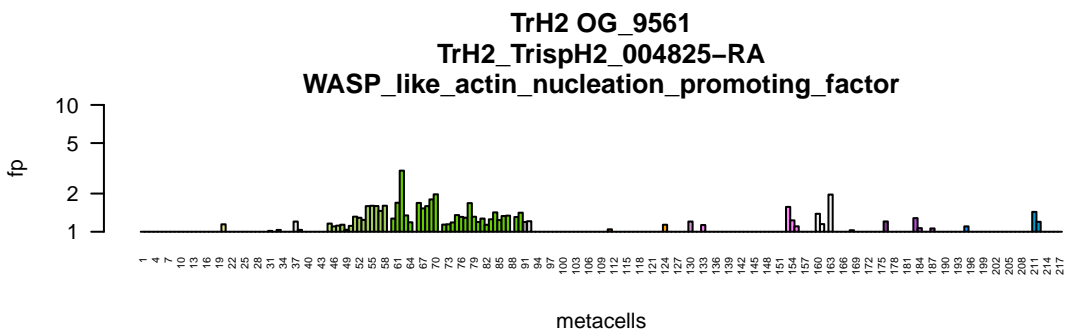
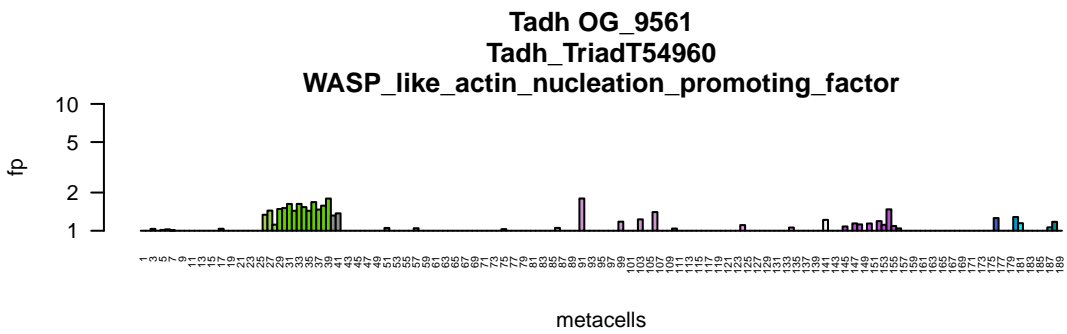
HoiH23 OG_7215
HoiH23_PIH23_003953-RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2

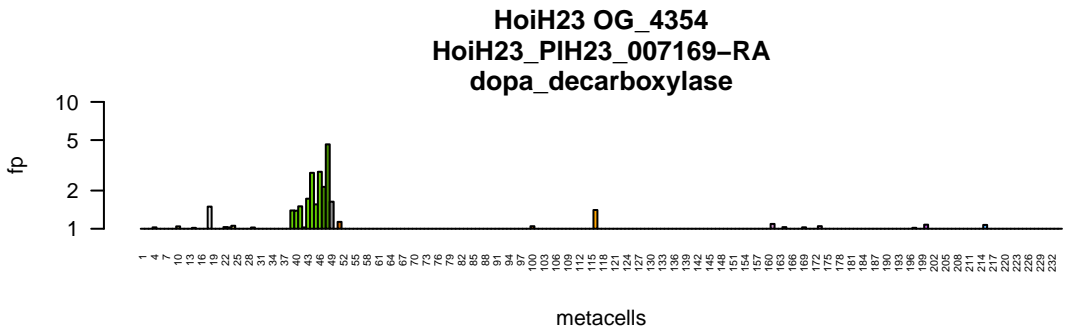
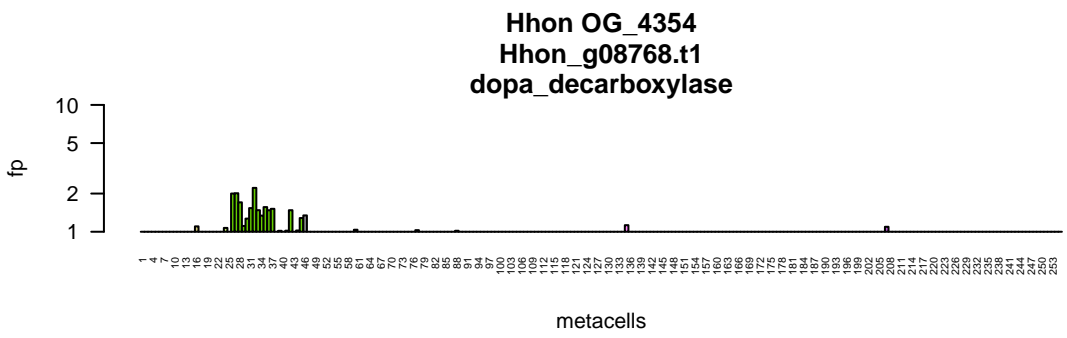
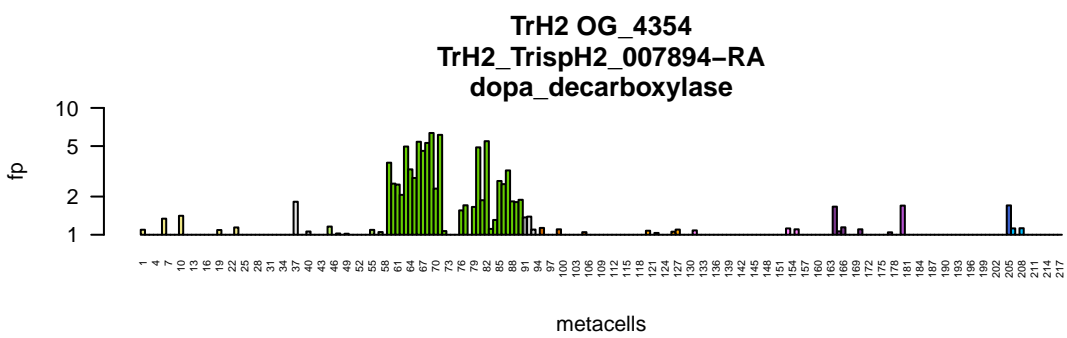
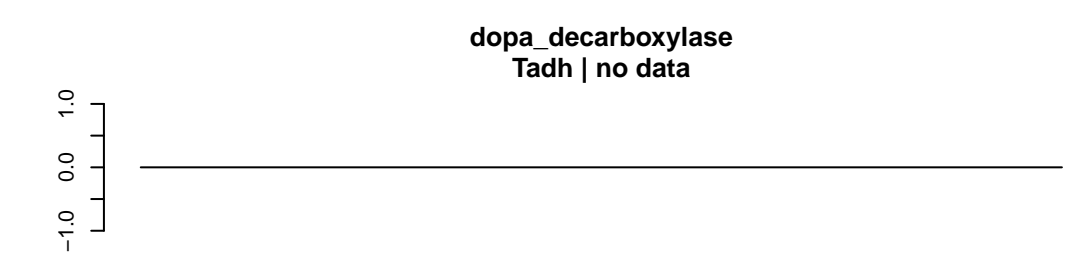


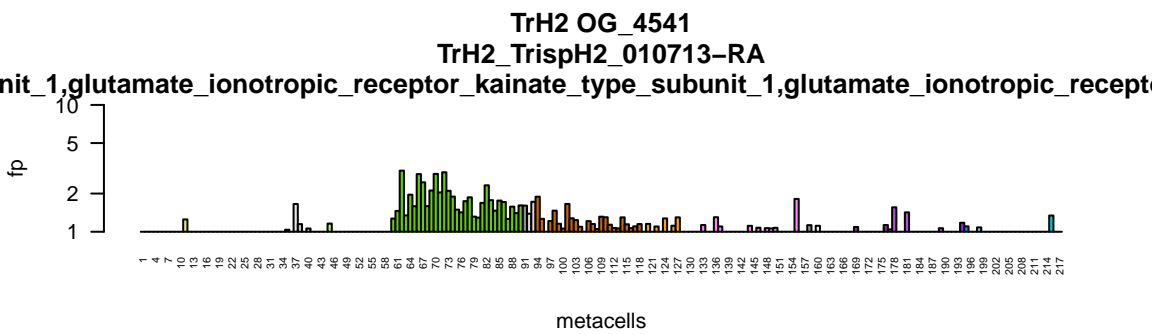
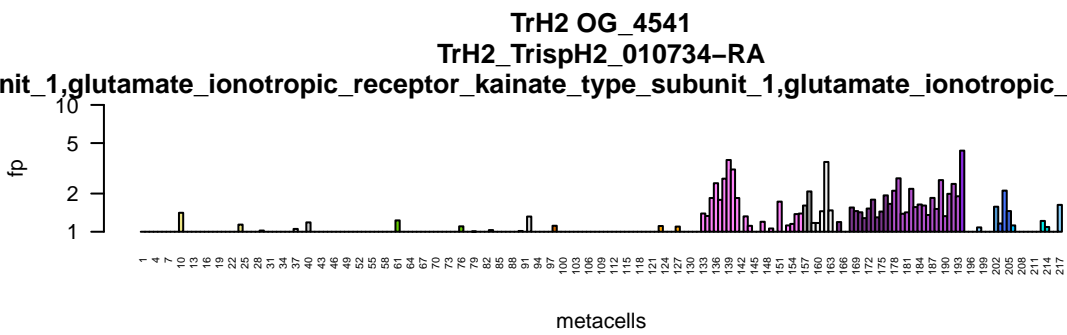
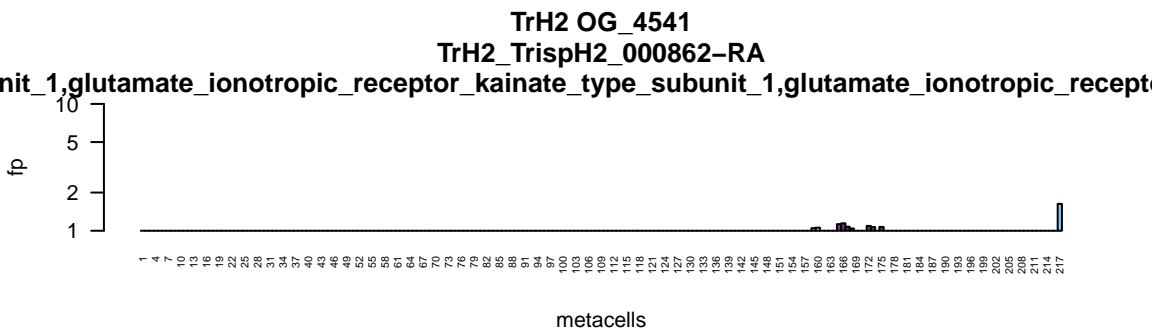
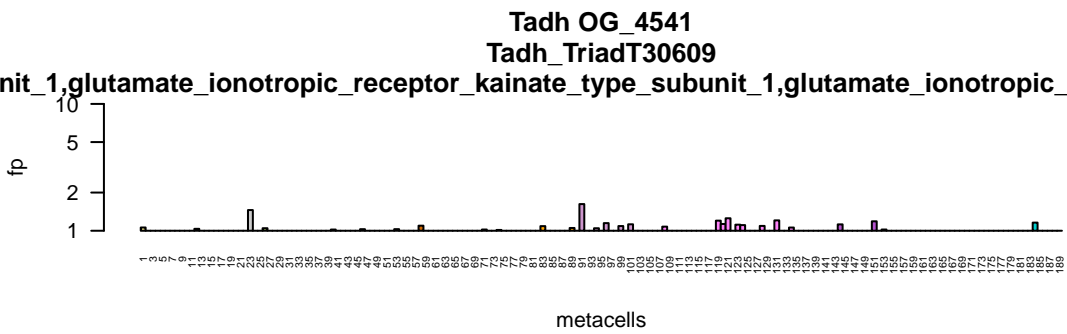
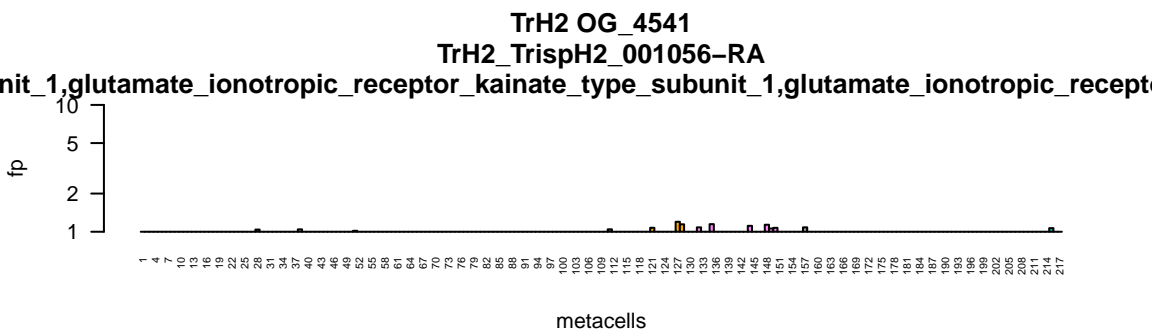
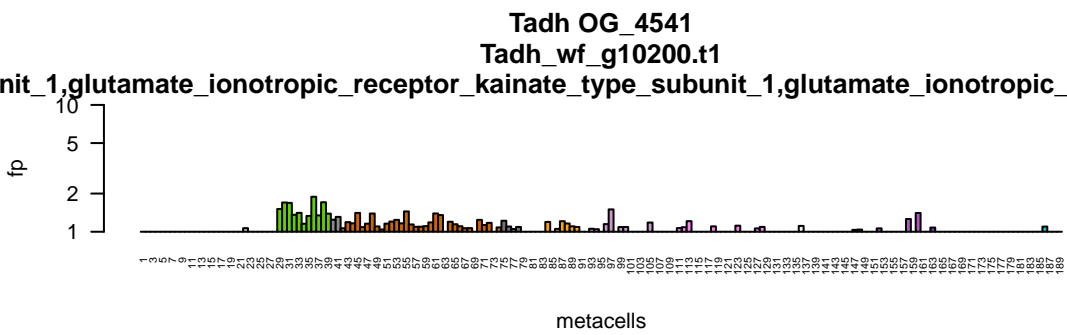
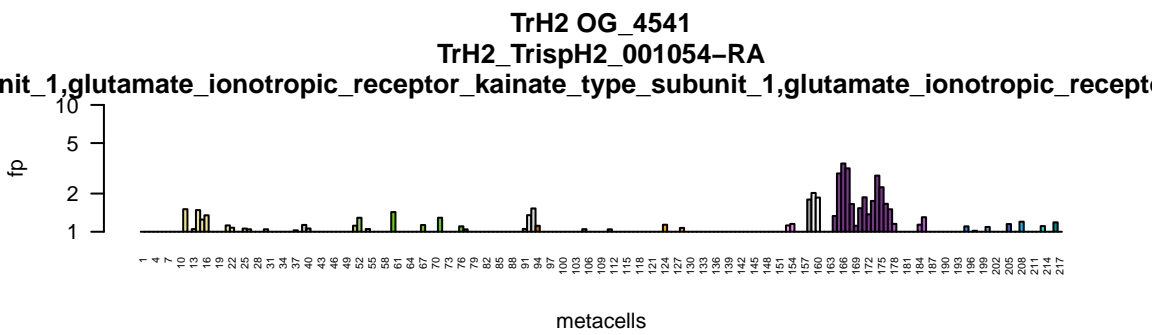
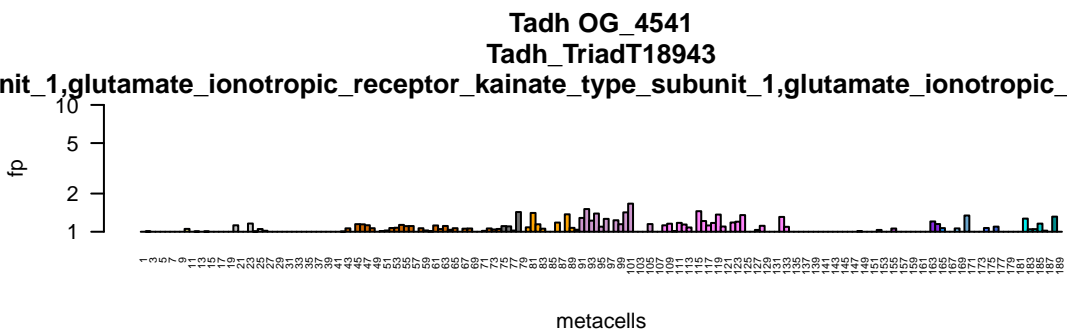
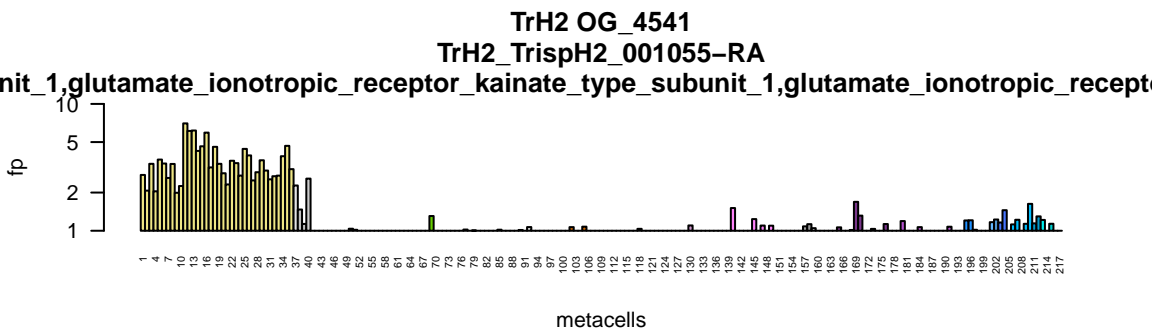
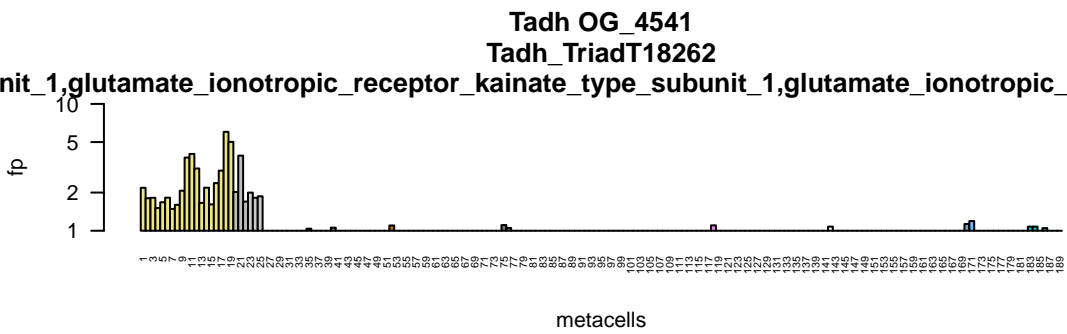
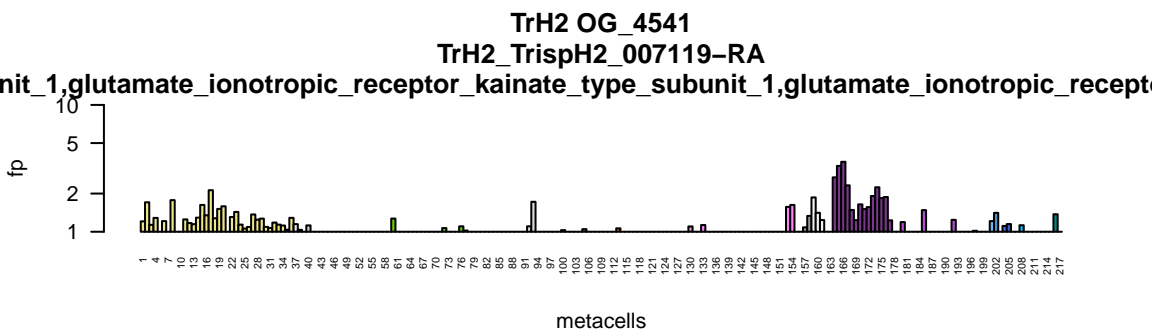
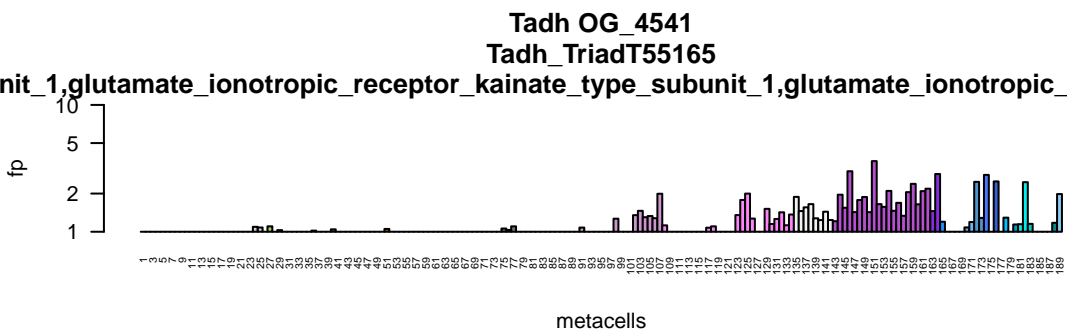
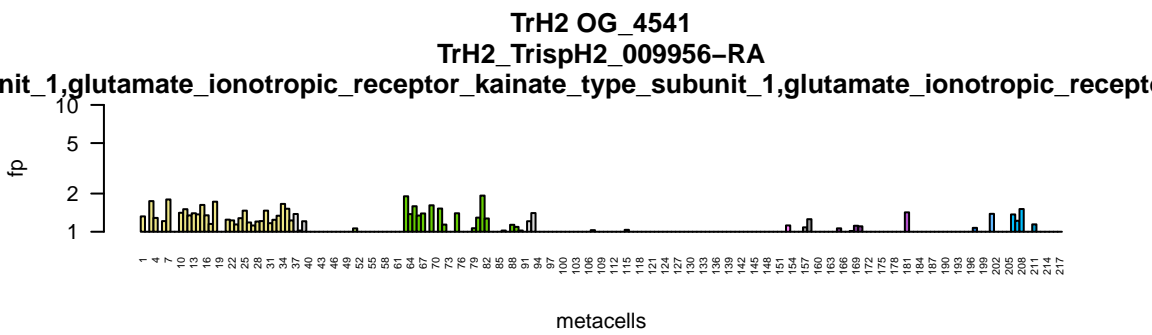
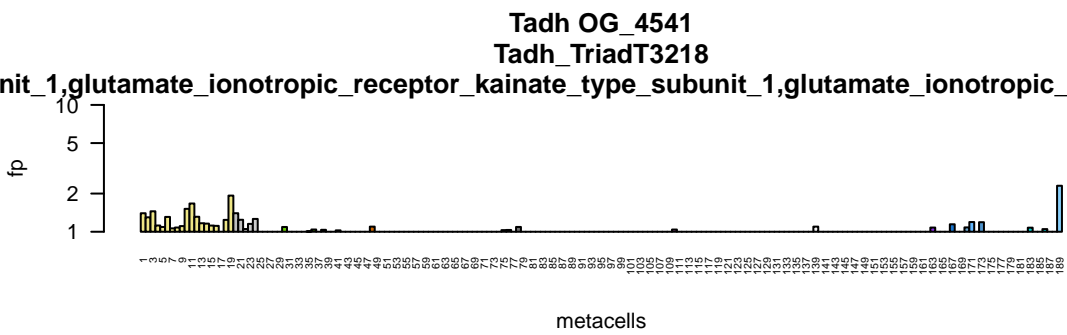
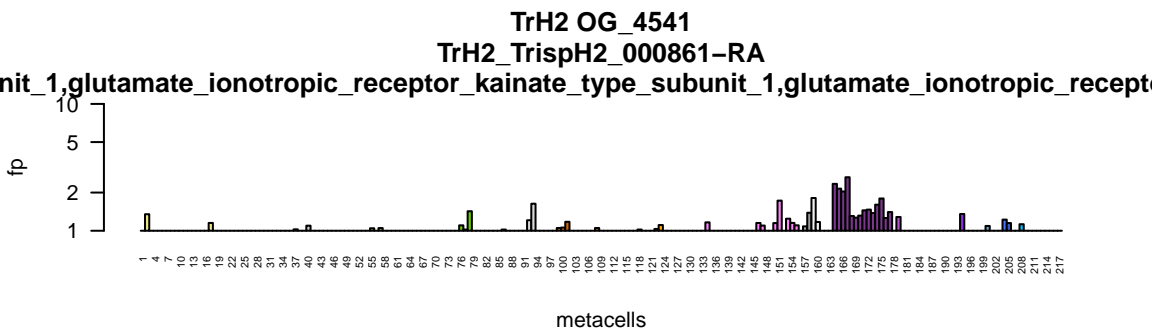
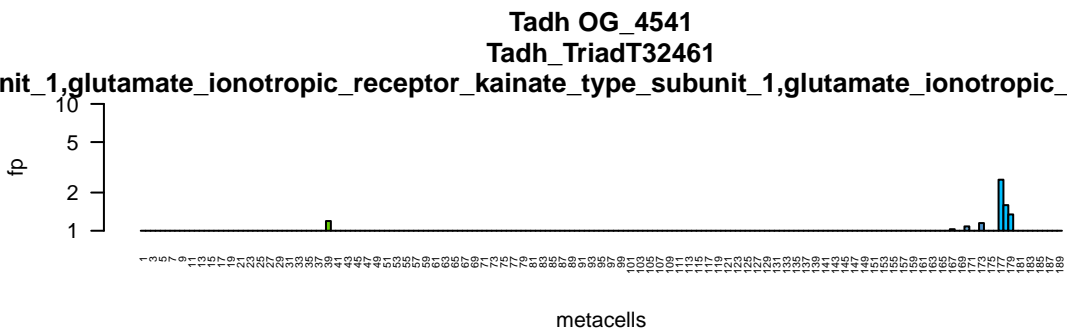
metacells

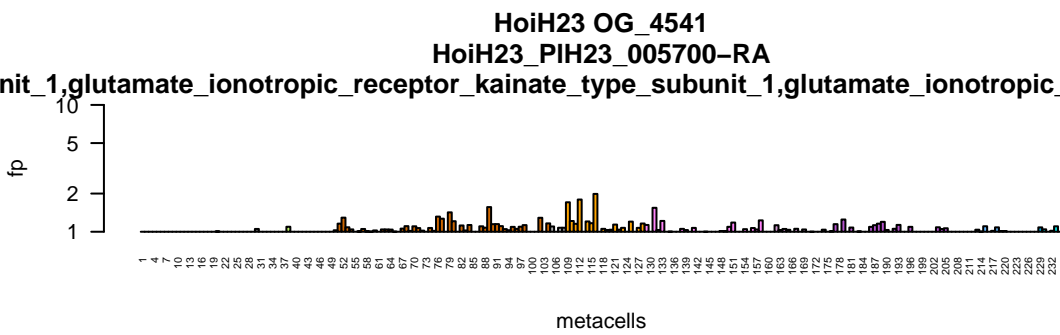
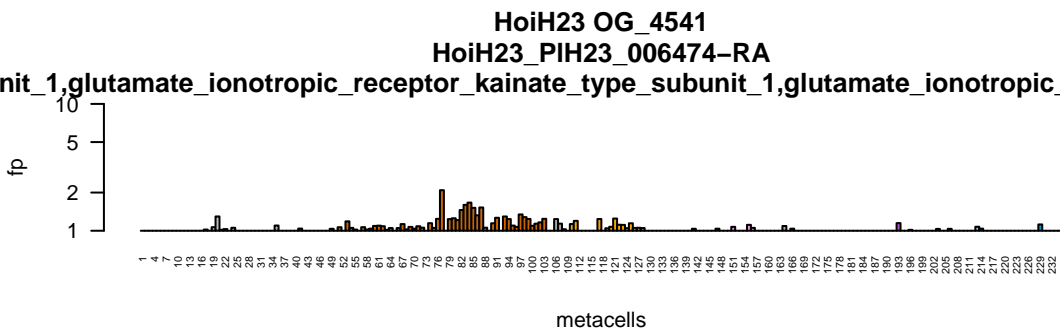
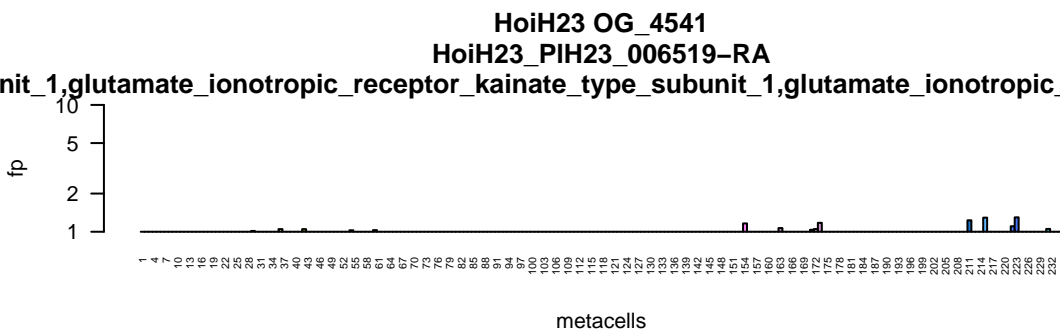
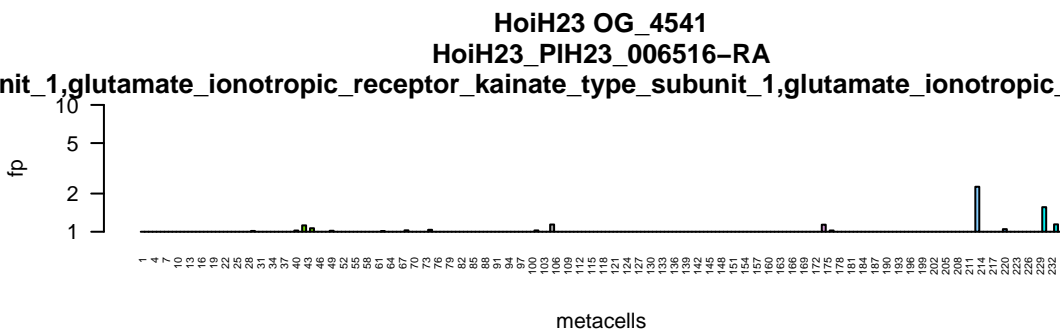
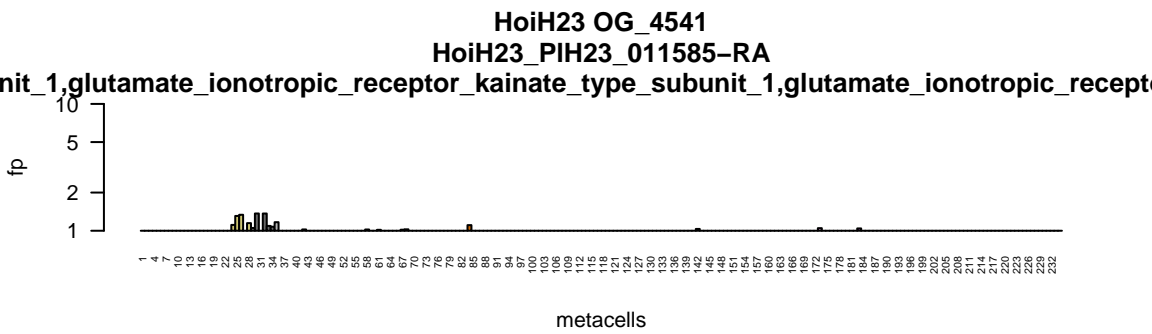
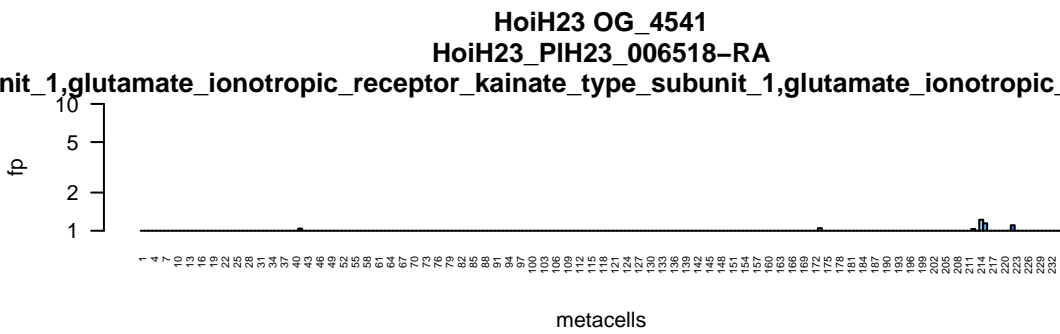
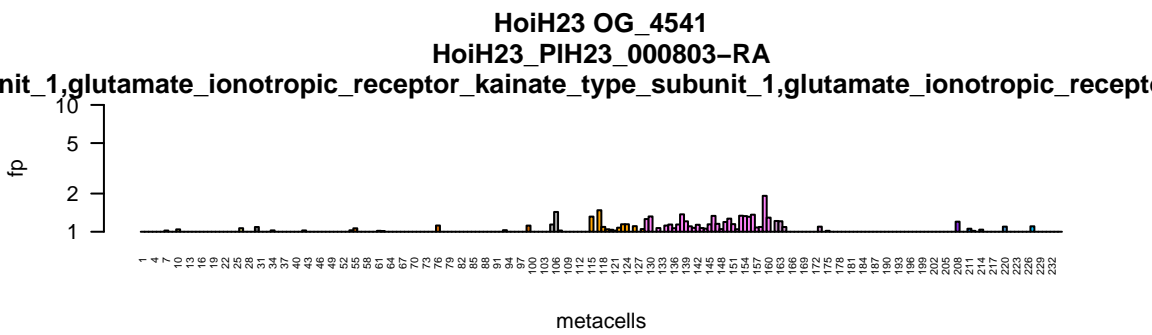
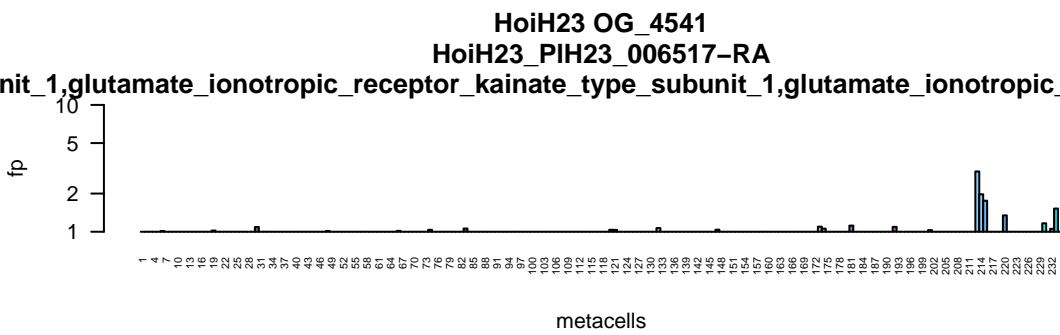
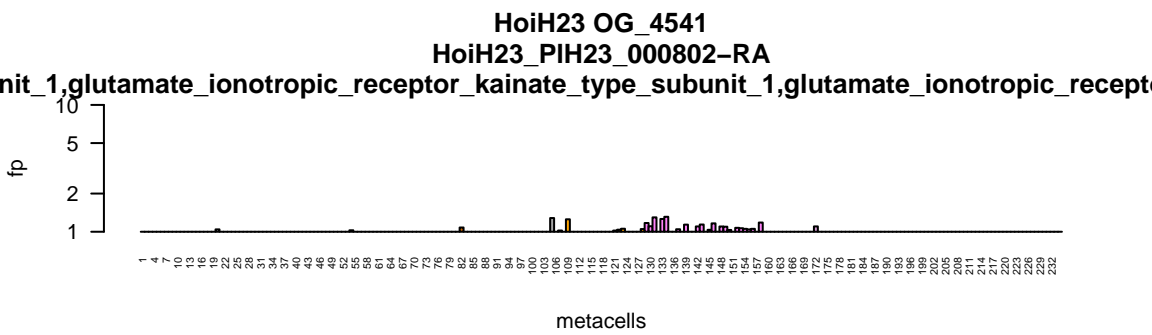
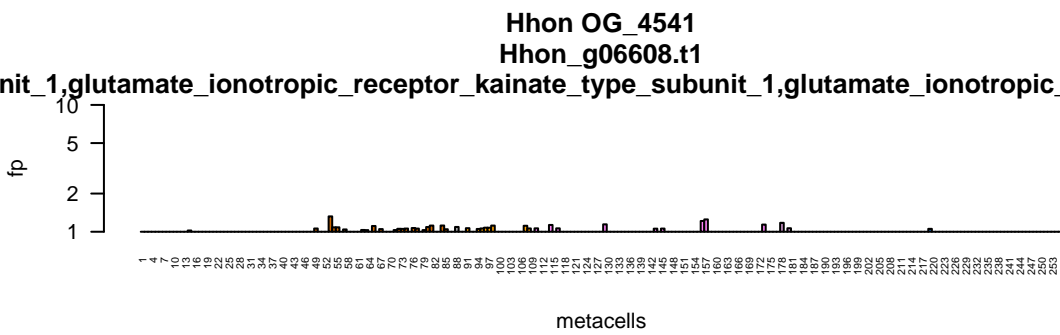
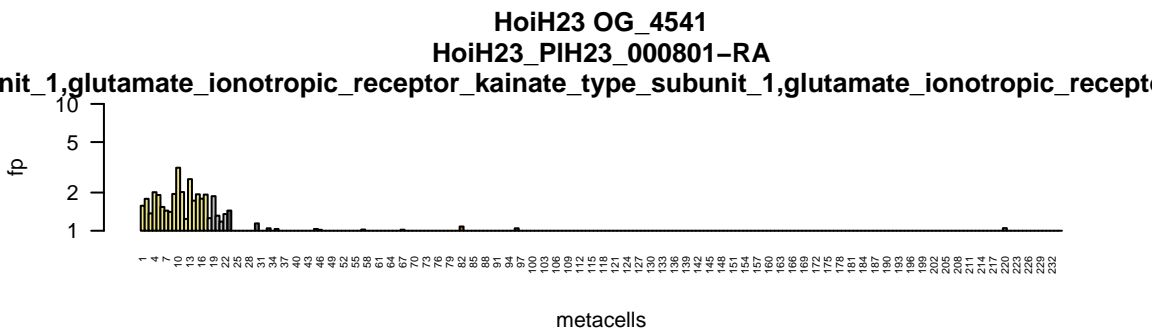
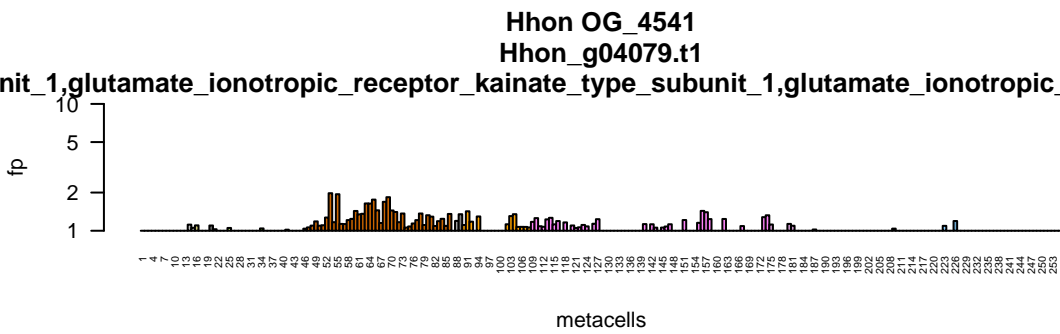


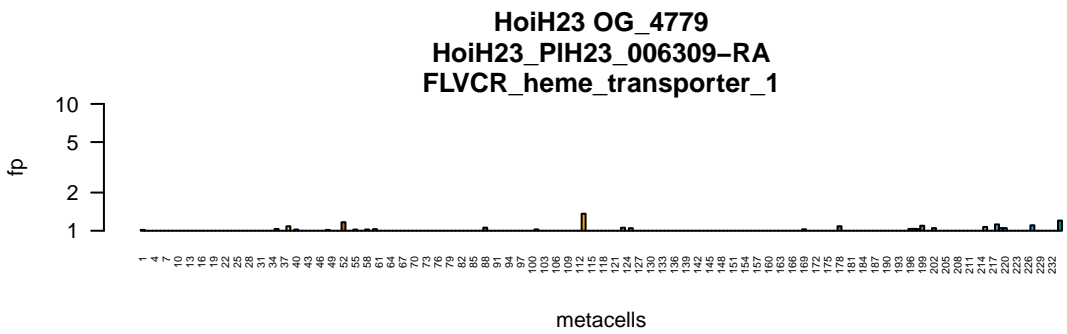
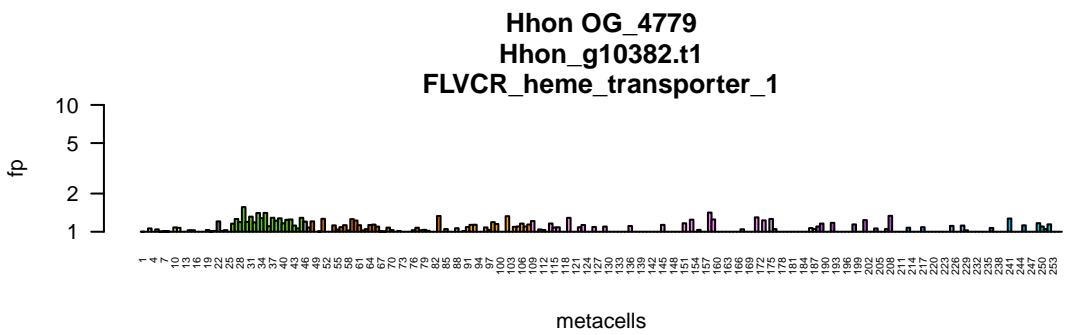
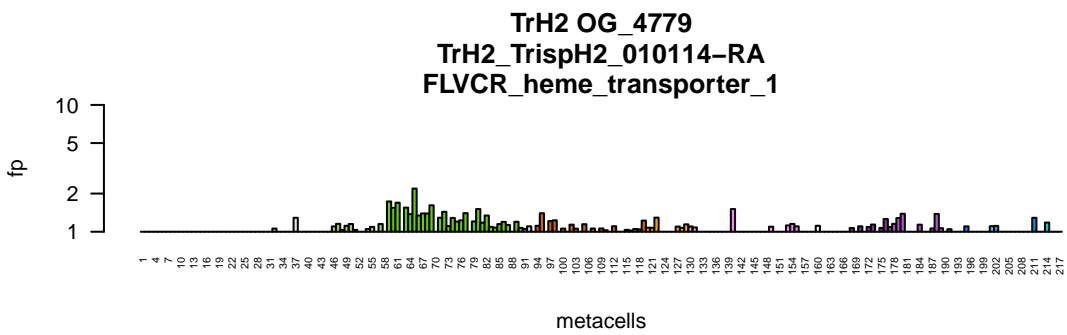
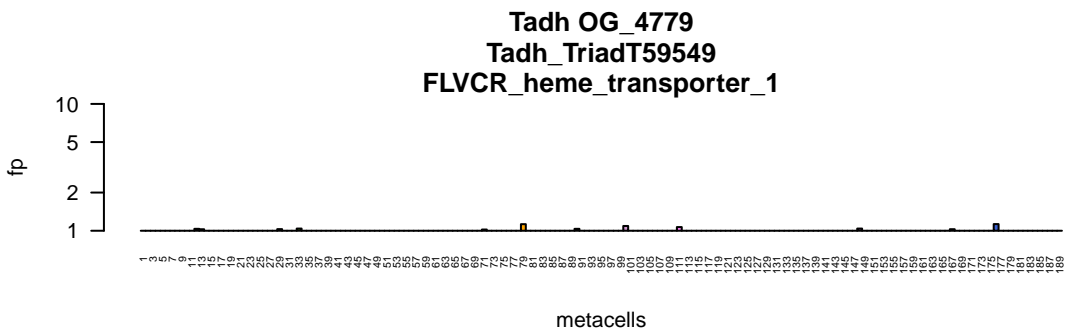


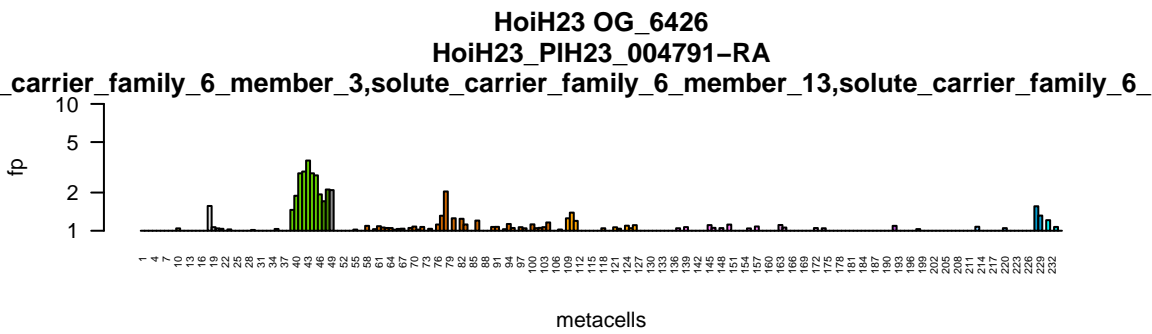
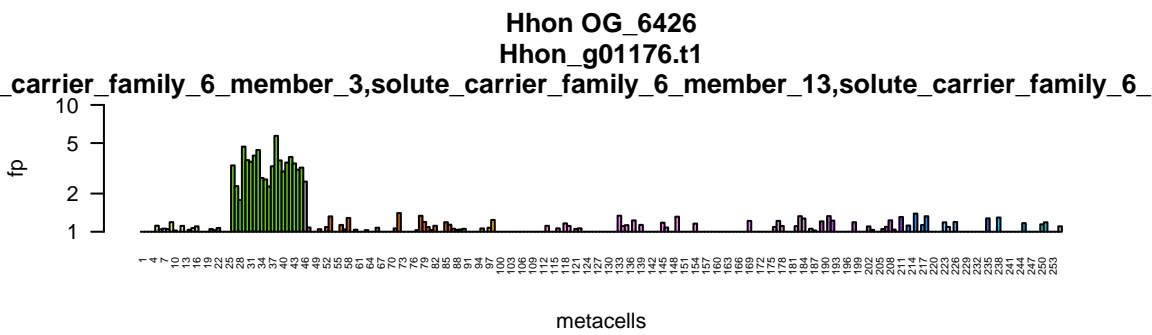
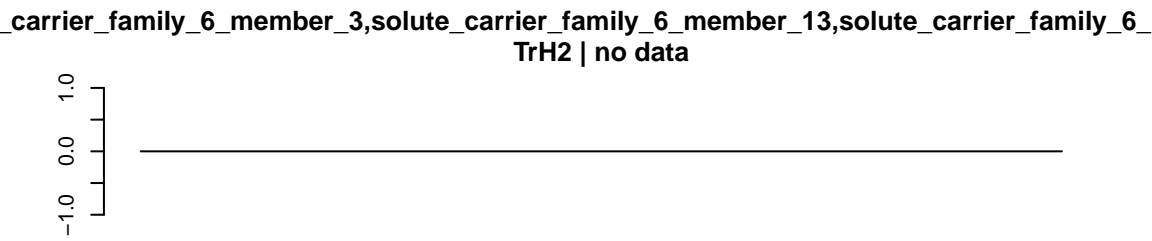
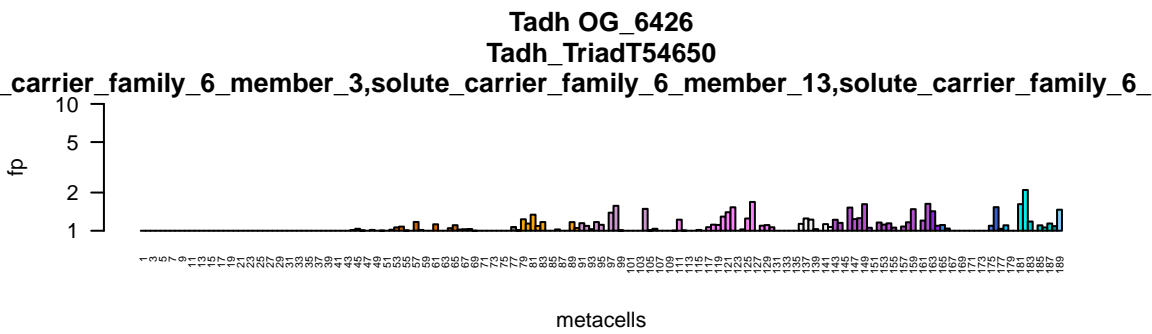


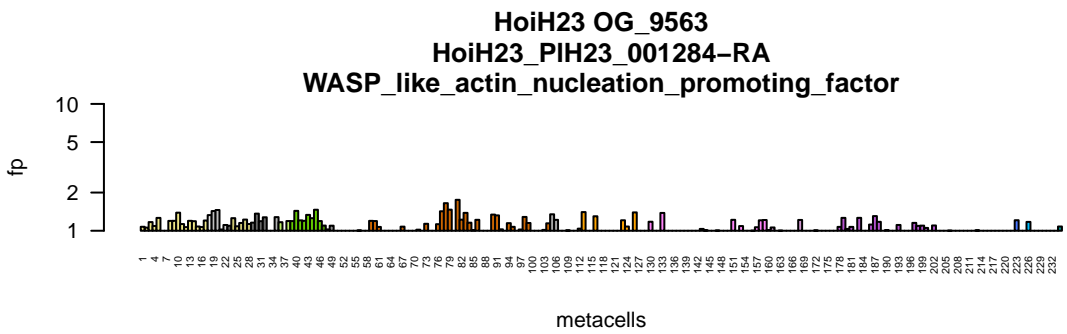
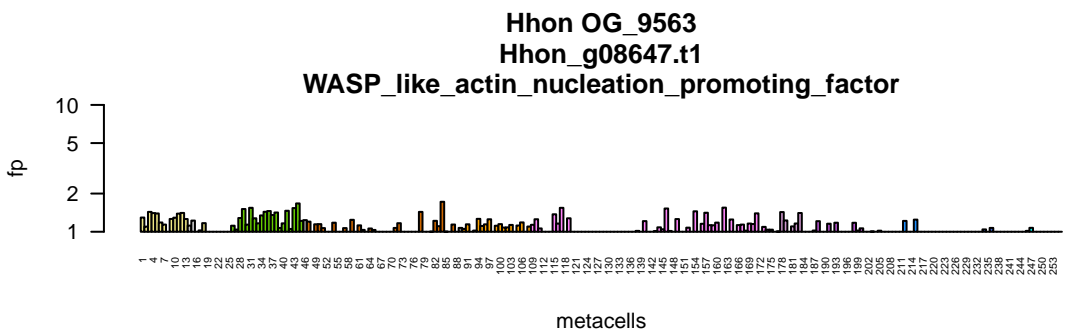
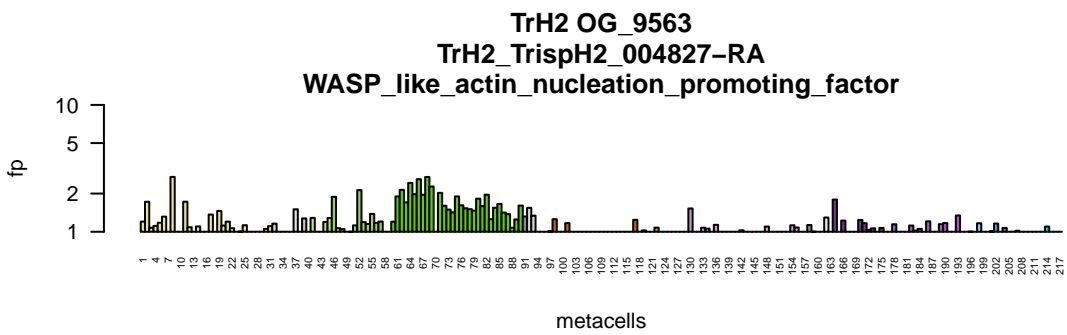
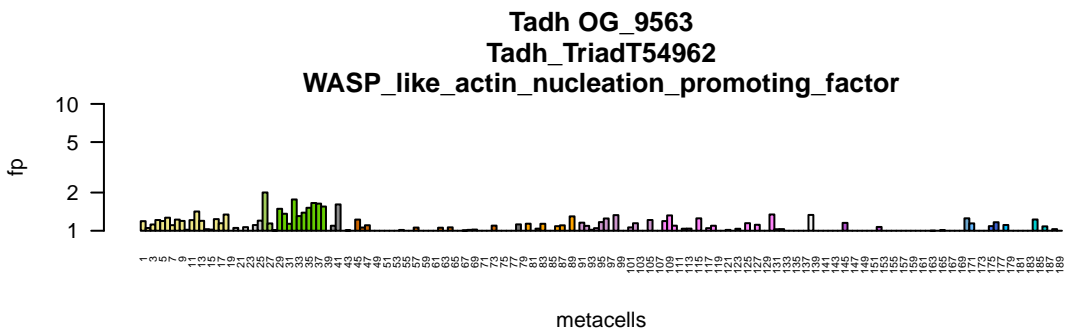


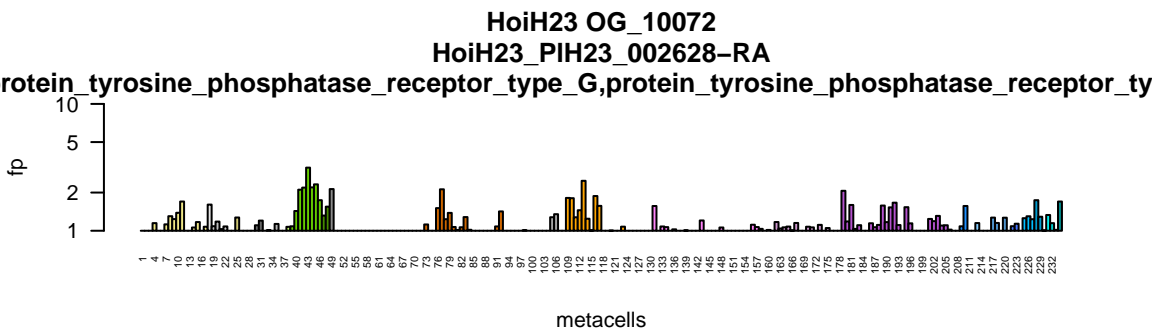
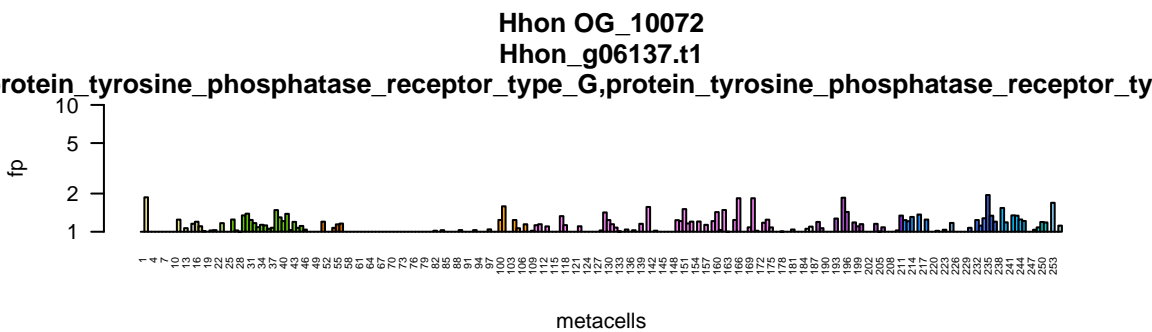
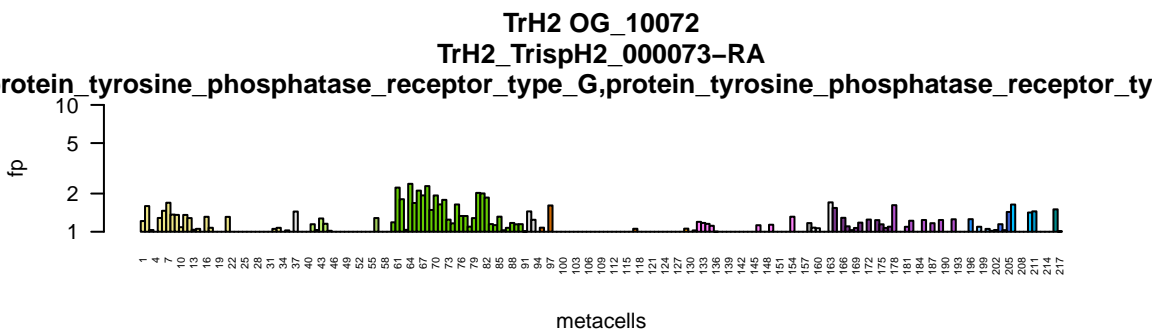
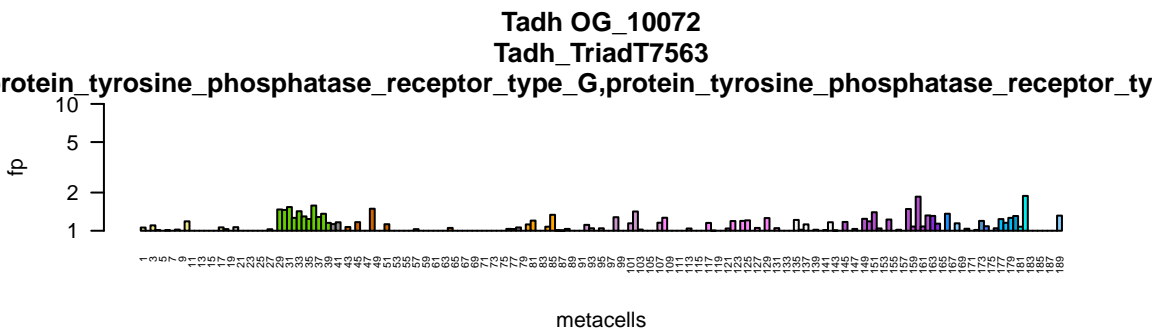


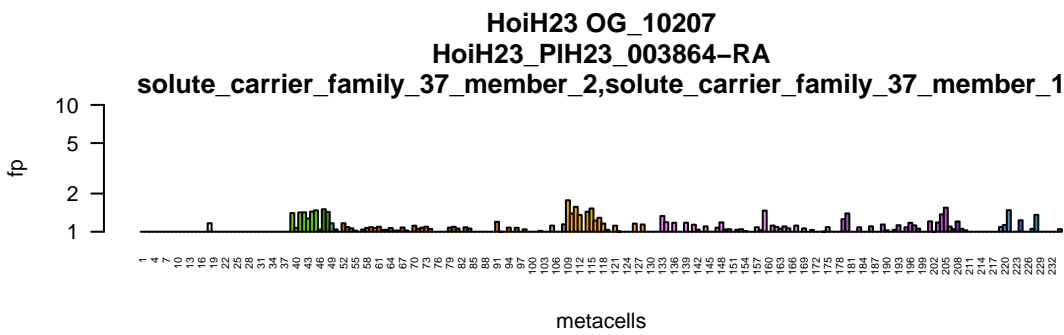
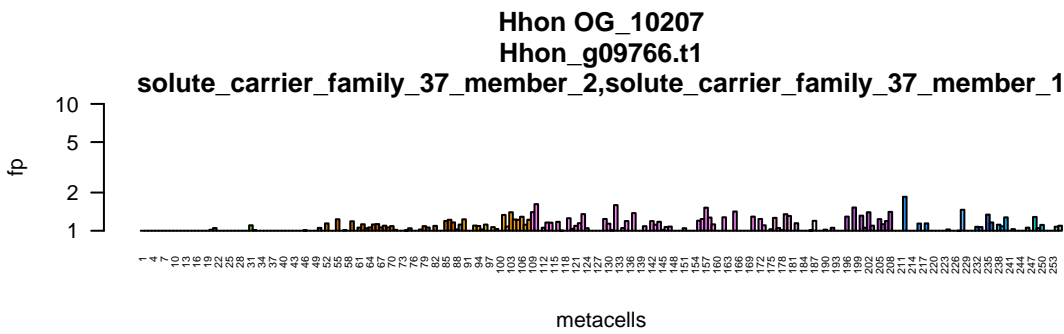
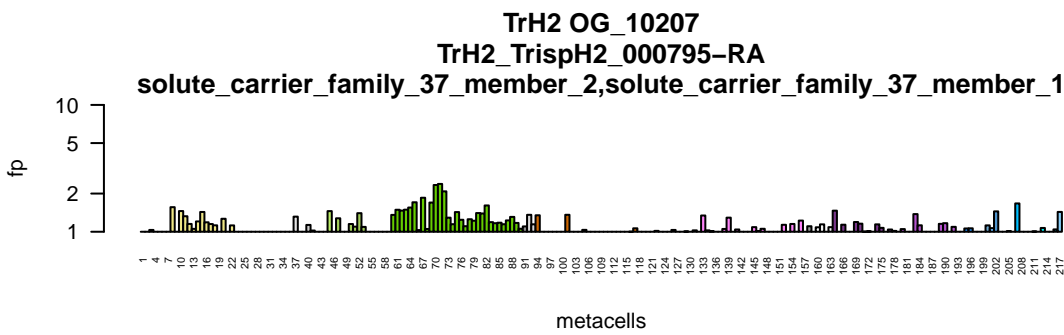
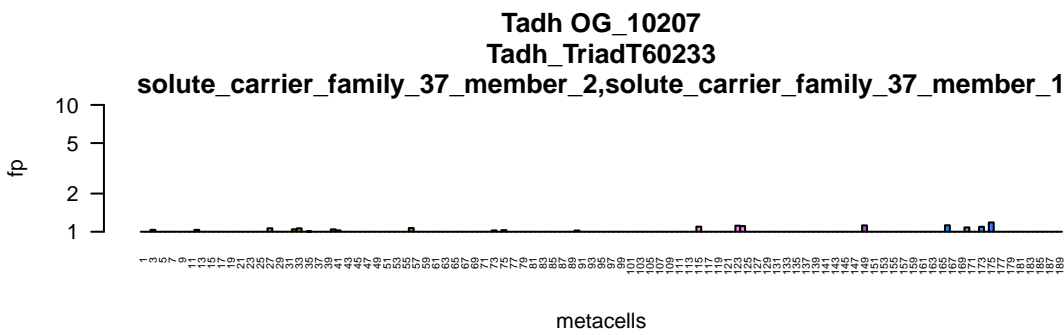
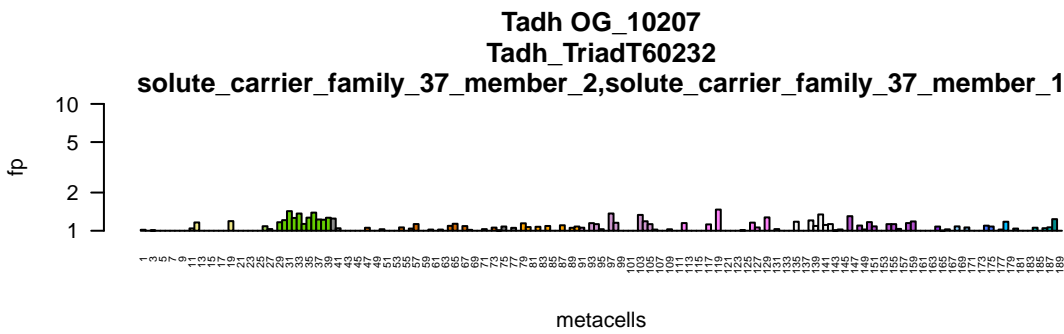


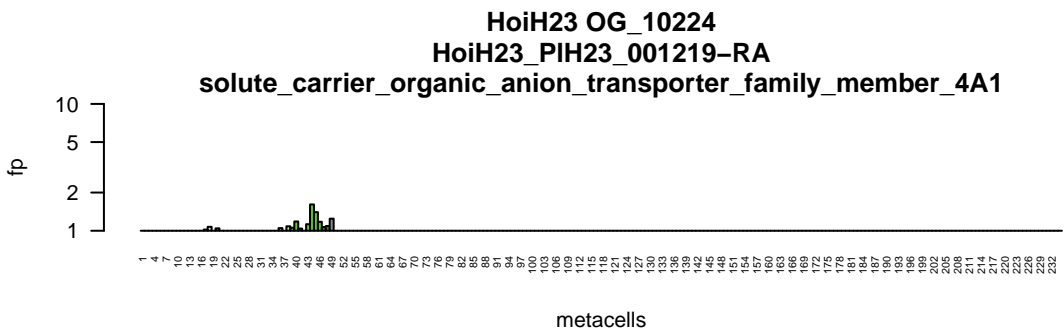
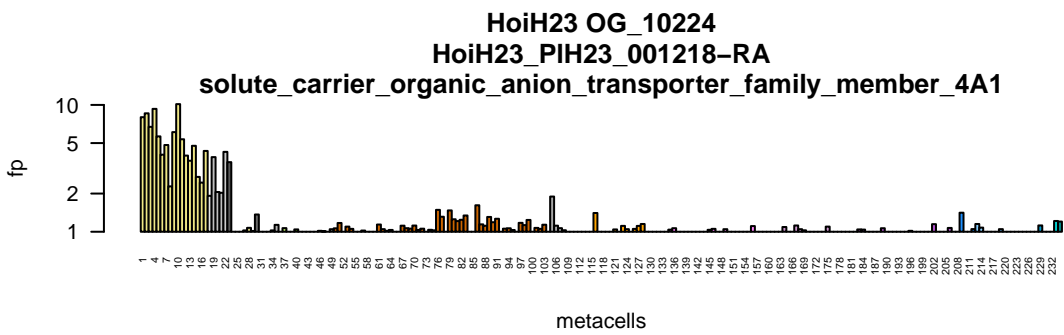
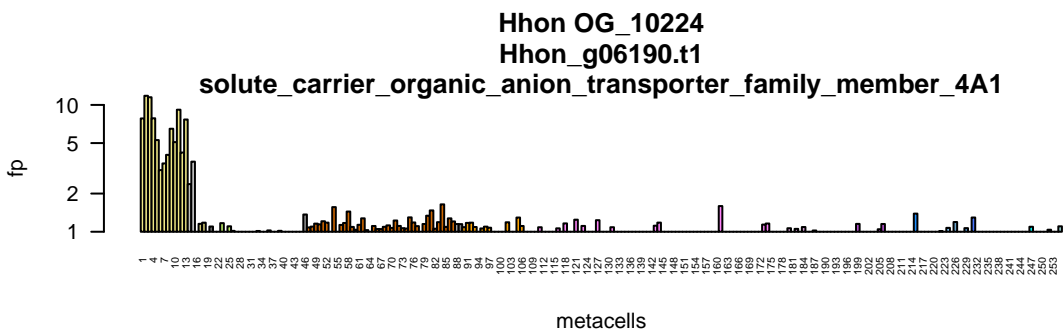
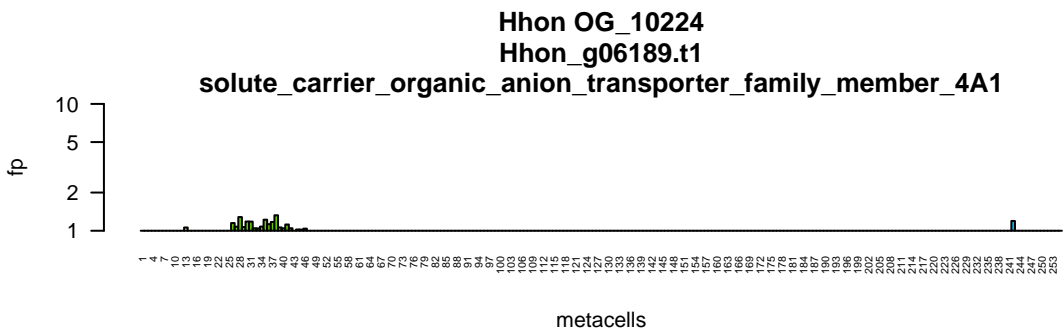
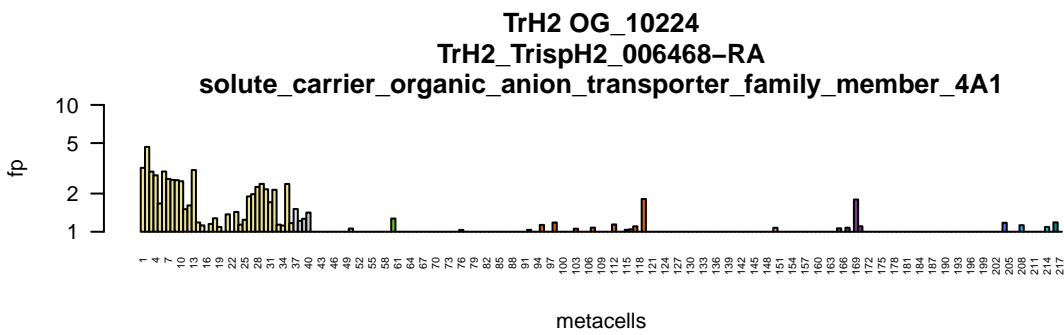
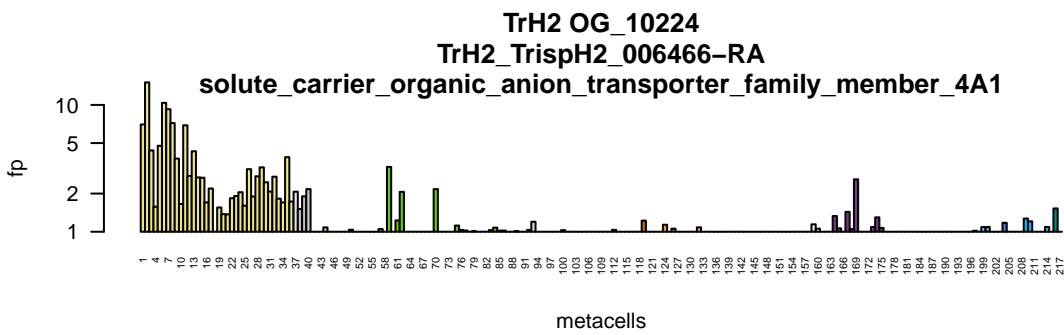
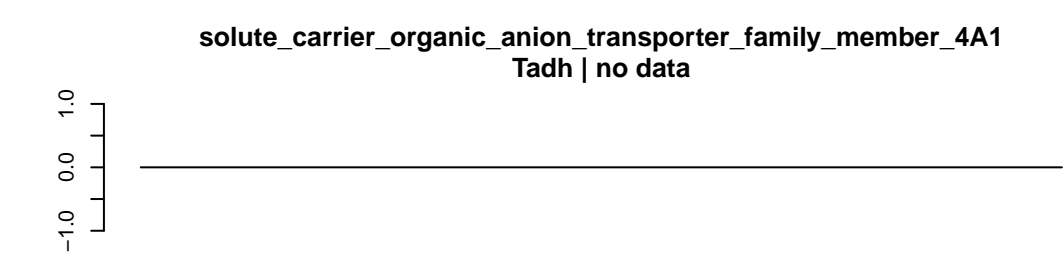


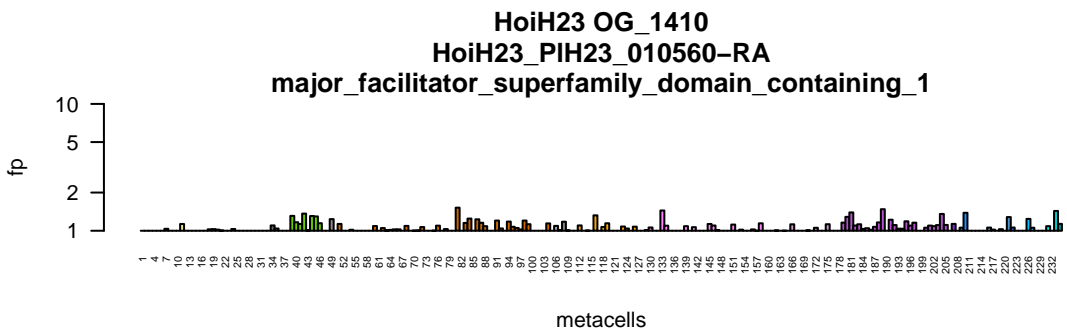
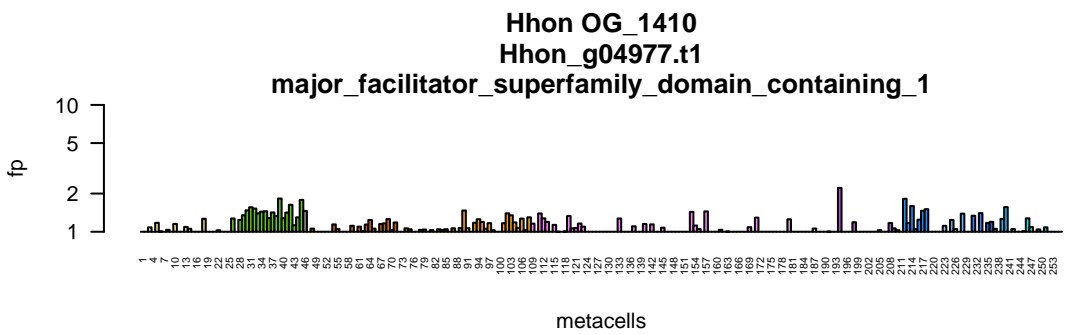
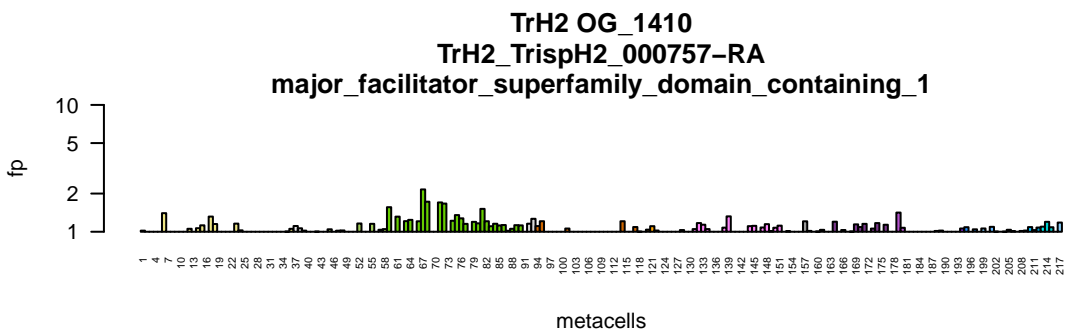
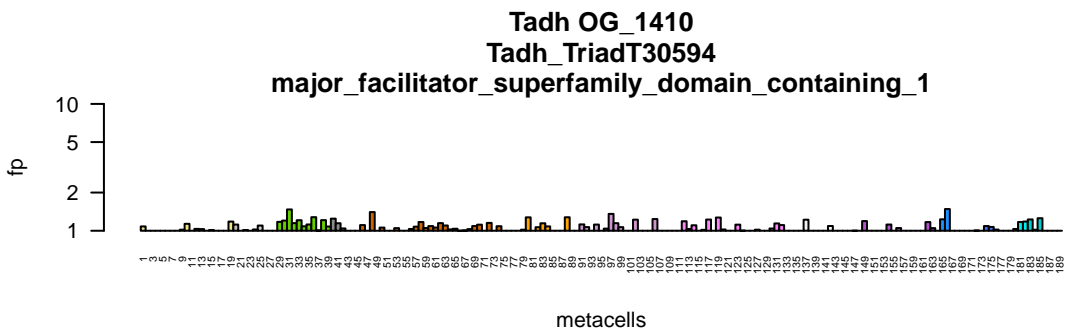


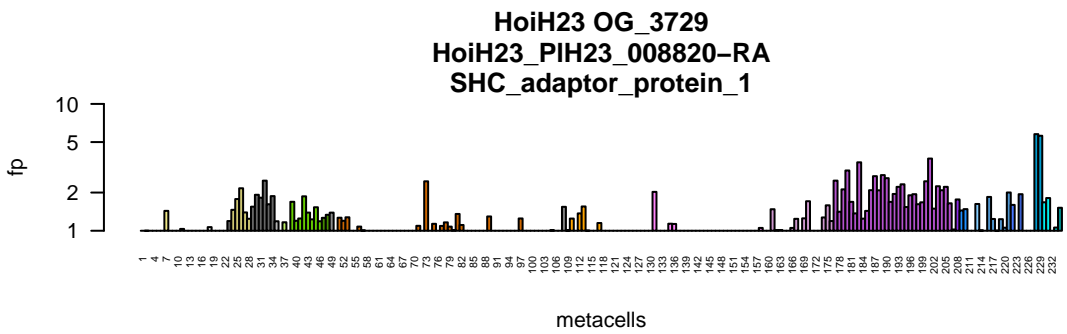
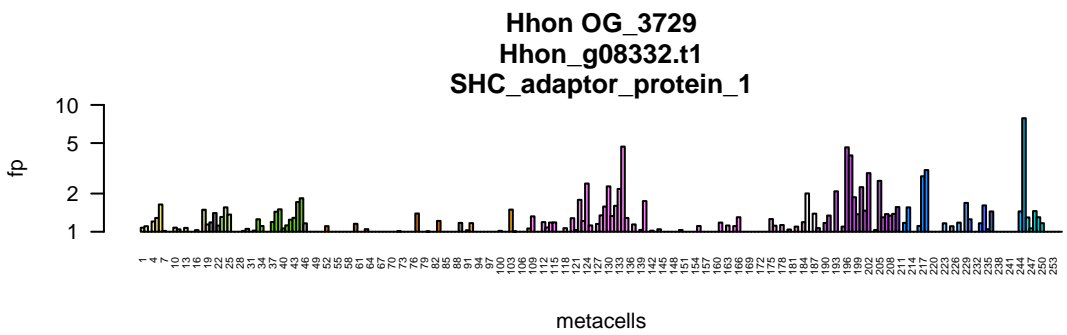
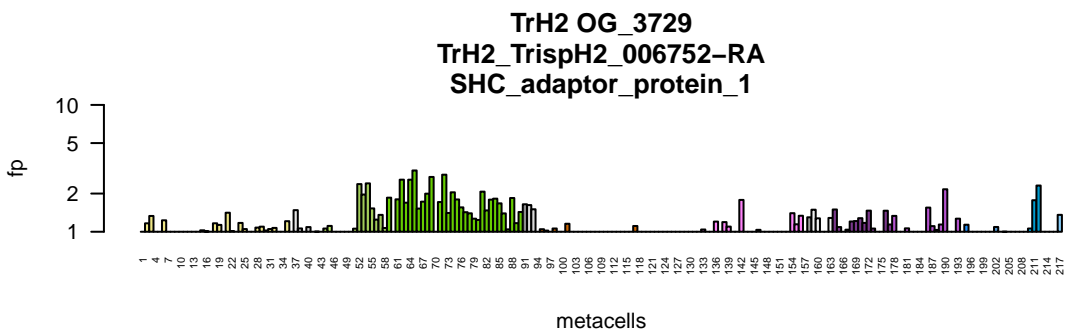
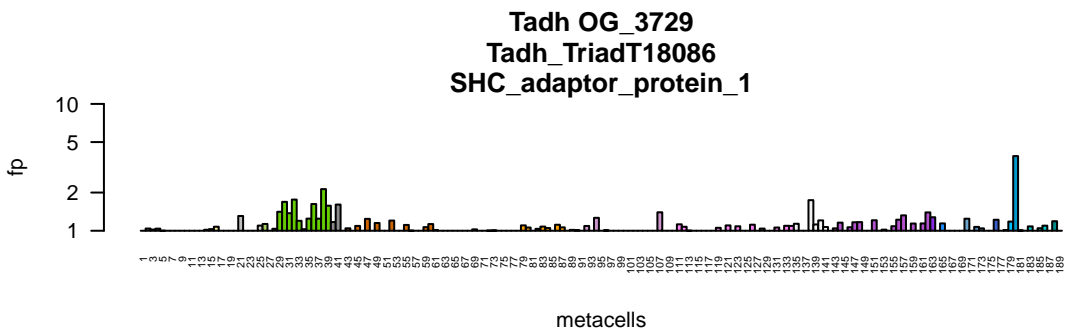








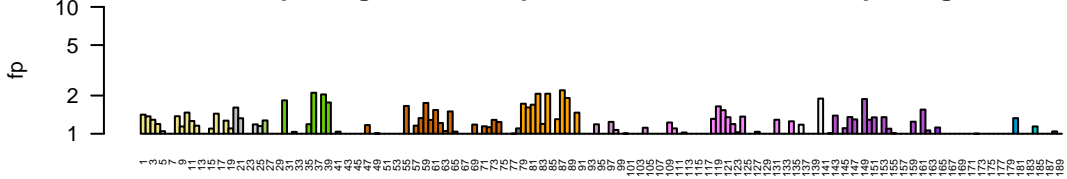




Tadh OG_4109

Tadh_TriadT63163

ATPase_Na_K_transporting_subunit_alpha_1,ATPase_Na_K_transporting_subunit_alpha

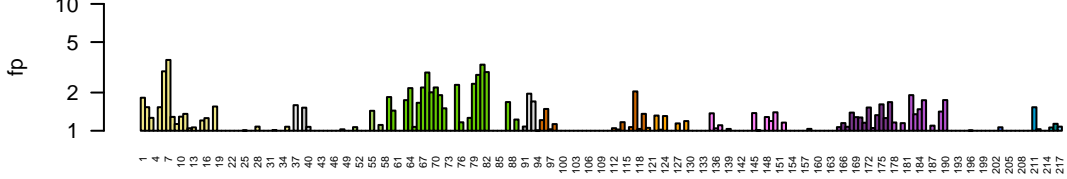


metacells

TrH2 OG_4109

TrH2_TrispH2_011484-RA

ATPase_Na_K_transporting_subunit_alpha_1,ATPase_Na_K_transporting_subunit_alpha

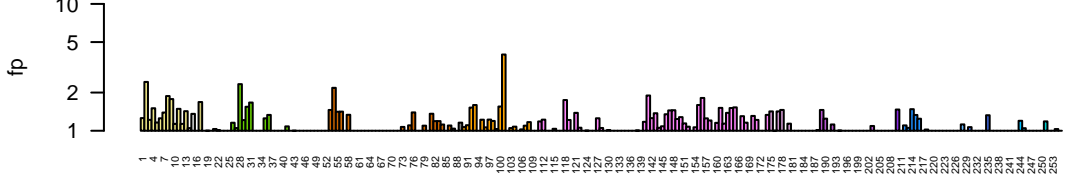


metacells

Hhon OG_4109

Hhon_g07340.t1

ATPase_Na_K_transporting_subunit_alpha_1,ATPase_Na_K_transporting_subunit_alpha

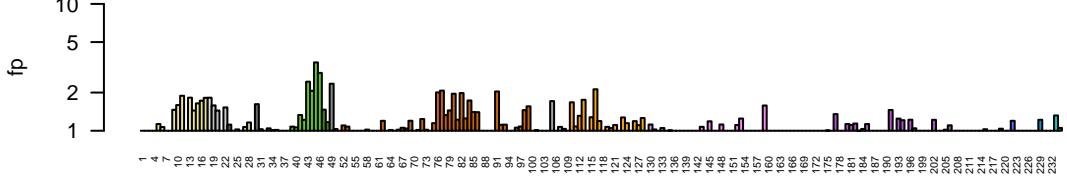


metacells

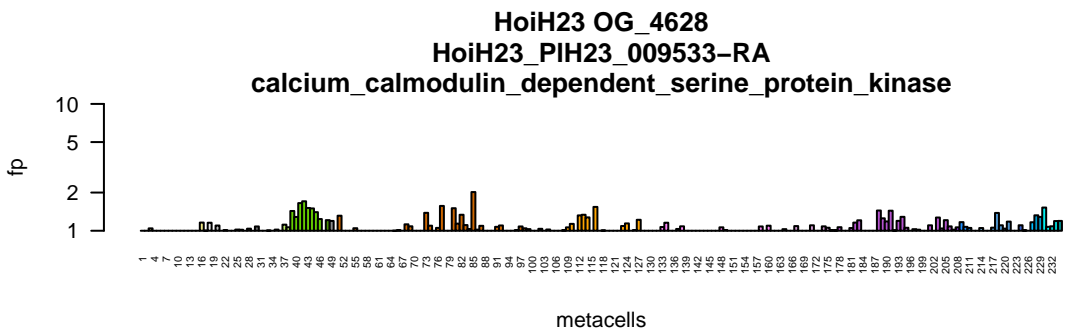
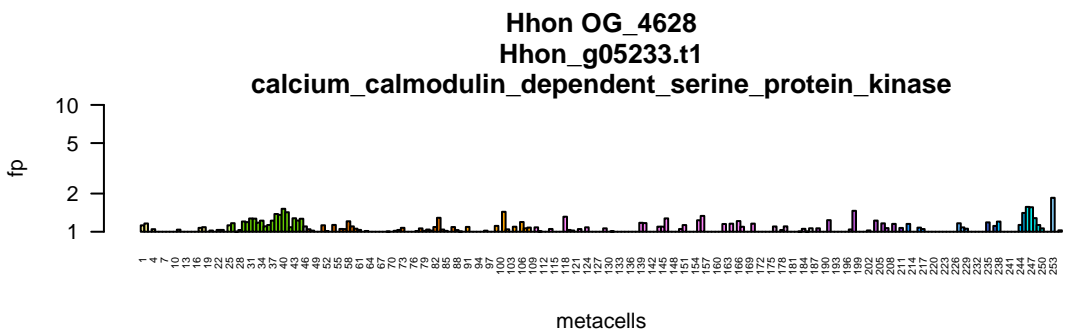
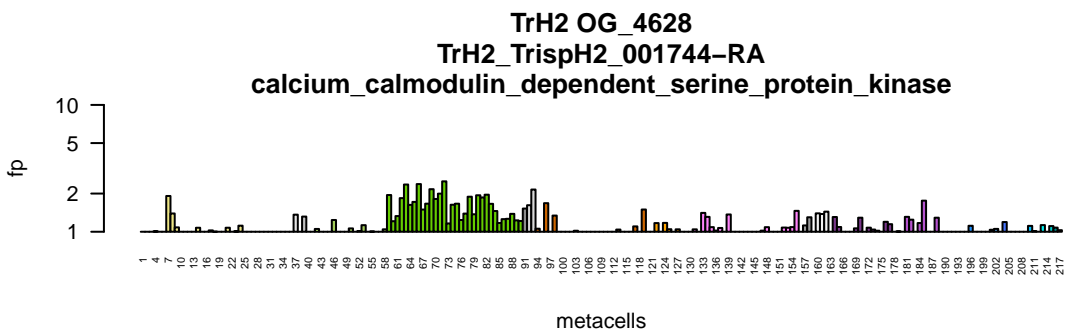
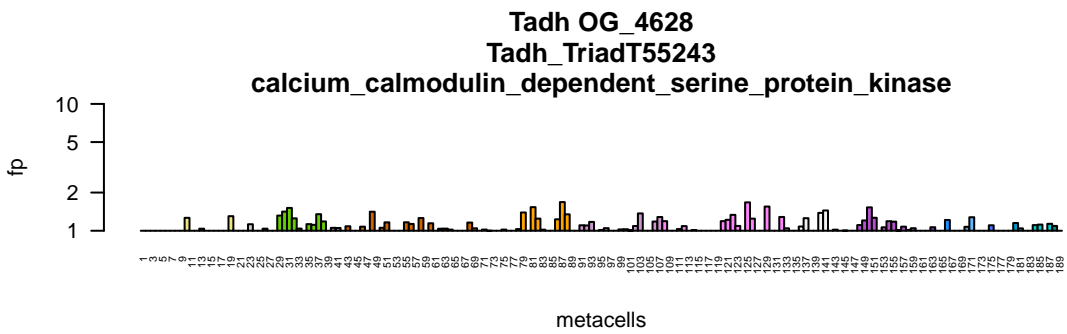
HoiH23 OG_4109

HoiH23_PIH23_011185-RA

ATPase_Na_K_transporting_subunit_alpha_1,ATPase_Na_K_transporting_subunit_alpha



metacells



A bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and ranges from 1 to 217. The y-axis is labeled 'fp' and ranges from 1 to 10. The chart shows that most metacells have a frequency of 1, with a small cluster of higher frequencies around metacell 154.

metacells	fp
1	1
4	1
10	1
13	1
16	1
22	1
23	1
25	1
28	1
31	1
32	1
37	1
40	1
43	1
44	1
49	1
55	1
56	1
59	1
64	1
67	1
73	1
76	1
79	1
82	1
88	1
91	1
94	1
100	1
103	1
108	1
110	1
112	1
115	1
118	1
124	1
127	1
130	1
133	1
136	1
139	1
142	1
145	1
148	1
151	1
154	2
155	1
157	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
202	1
205	1
208	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 indices from 1 to 217. Most metacells have 1 false positive, but metacells 55 and 56 have 2, and metacells 190 and 191 have 1.5.

metacell	fp
1	1
4	1
10	1
13	1
16	1
22	1
23	1
25	1
28	1
31	1
32	1
37	1
40	1
43	1
49	1
52	1
55	2
56	2
61	1
64	1
67	1
73	1
76	1
79	1
82	1
88	1
91	1
94	1
100	1
103	1
108	1
112	1
115	1
118	1
121	1
124	1
127	1
130	1
133	1
139	1
142	1
145	1
148	1
151	1.5
154	1
157	1
163	1
166	1
169	1
172	1
175	1
178	1
181	1
184	1
187	1
190	1.5
193	1
196	1
200	1
202	1
205	1
208	1
211	1
214	1
217	2

A bar chart showing the frequency of metacells. The x-axis is labeled 'metacells' and the y-axis is labeled 'fp'. The y-axis has a non-linear scale with values 1, 2, 5, and 10. The x-axis lists metacell IDs from 1 to 189. Most metacells have a frequency of 1, represented by a single purple bar. A few metacells have higher frequencies: metacell 103 has a frequency of 2, metacell 111 has a frequency of 2, metacell 113 has a frequency of 2, metacell 117 has a frequency of 2, metacell 123 has a frequency of 2, metacell 133 has a frequency of 2, metacell 135 has a frequency of 2, metacell 137 has a frequency of 2, metacell 141 has a frequency of 2, metacell 144 has a frequency of 2, metacell 145 has a frequency of 2, metacell 146 has a frequency of 2, metacell 151 has a frequency of 2, metacell 152 has a frequency of 2, metacell 153 has a frequency of 2, metacell 154 has a frequency of 2, metacell 155 has a frequency of 2, metacell 156 has a frequency of 2, metacell 157 has a frequency of 2, metacell 158 has a frequency of 2, metacell 159 has a frequency of 2, metacell 161 has a frequency of 2, metacell 162 has a frequency of 2, metacell 163 has a frequency of 2, metacell 164 has a frequency of 2, metacell 165 has a frequency of 2, metacell 166 has a frequency of 2, metacell 167 has a frequency of 2, metacell 168 has a frequency of 2, metacell 169 has a frequency of 2, metacell 171 has a frequency of 2, metacell 172 has a frequency of 2, metacell 173 has a frequency of 2, metacell 174 has a frequency of 2, metacell 175 has a frequency of 2, metacell 176 has a frequency of 2, metacell 177 has a frequency of 2, metacell 178 has a frequency of 2, metacell 179 has a frequency of 2, metacell 180 has a frequency of 2, metacell 181 has a frequency of 2, metacell 182 has a frequency of 2, metacell 183 has a frequency of 2, metacell 184 has a frequency of 2, metacell 185 has a frequency of 2, metacell 186 has a frequency of 2, metacell 187 has a frequency of 2, metacell 188 has a frequency of 2, metacell 189 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 1 to 10. The x-axis is labeled 'metacells' and lists indices from 1 to 217. Most metacells have a false positive count of 1, with a few having counts of 2 or 4.

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

Bar chart showing the number of false positives (fp) for various metacells. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists 40 metacells. Most metacells have a false positive count of 1, while a few have counts of 2 or 3.

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

Bar chart showing the number of false positives (fp) for each metacell. The y-axis is labeled 'fp' and ranges from 1 to 10. The x-axis is labeled 'metacells' and lists 100 metacells. Most metacells have a false positive count of 1, with a few having counts of 2 or 3.

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

[illegible]

Figure 10: A bar chart showing the frequency of the top 253 metacells. The x-axis is labeled 'metacells' and lists metacell IDs from 1 to 253. The y-axis is labeled 'fp' (frequency) and ranges from 0 to 10. The chart shows a distribution of frequencies across the metacells, with most having a frequency of 1, and a few having higher frequencies (up to 2).

glycoprotein_2A,synaptic_vesicle_glycoprotein_2B,synaptic_vesicle_glycoprotein_2C

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	1
205	1
208	1
211	1
214	1
217	1

[illegible]

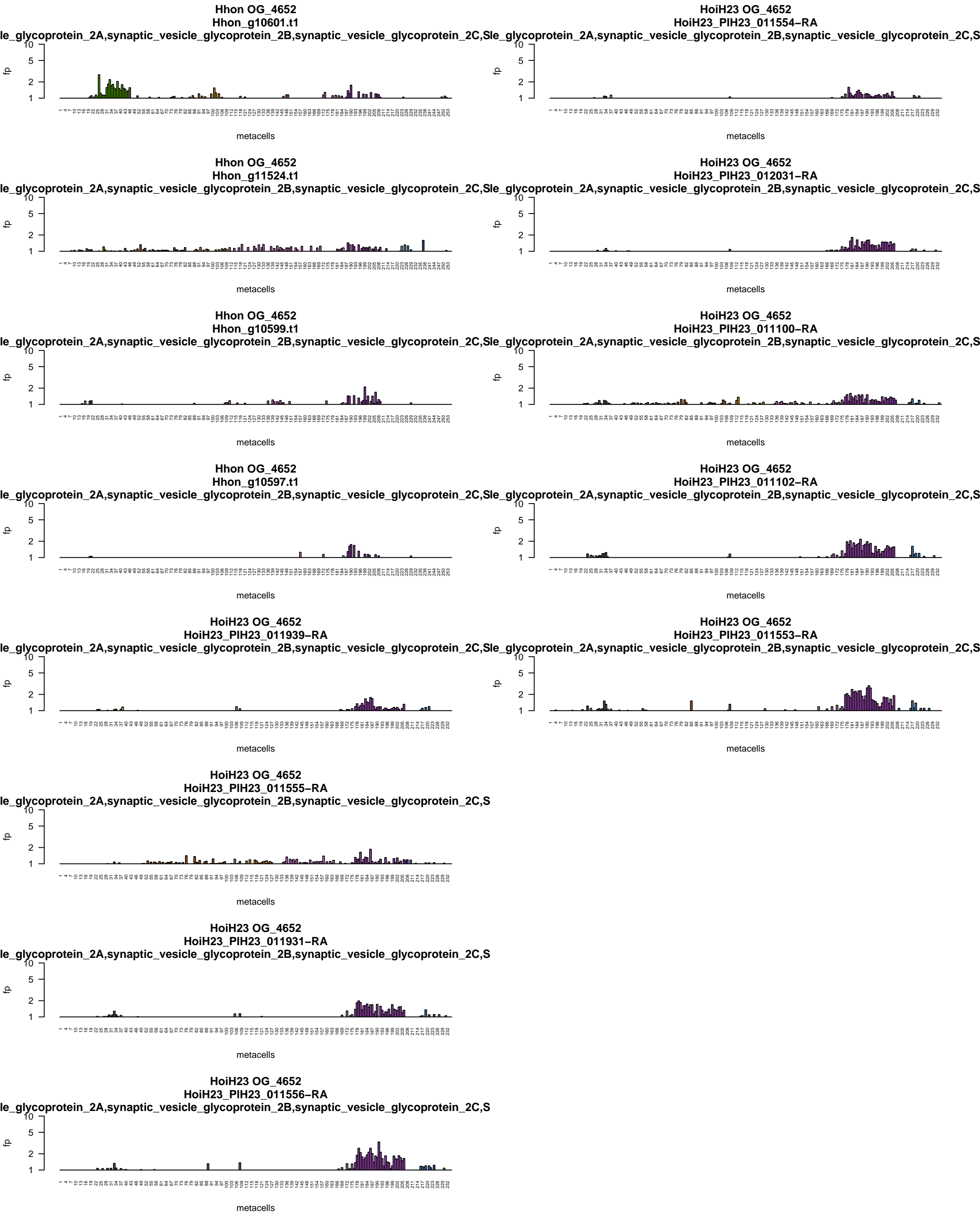
glycoprotein_2A,synaptic_vesicle_glycoprotein_2B,synaptic_vesicle_glycoprotein_2C

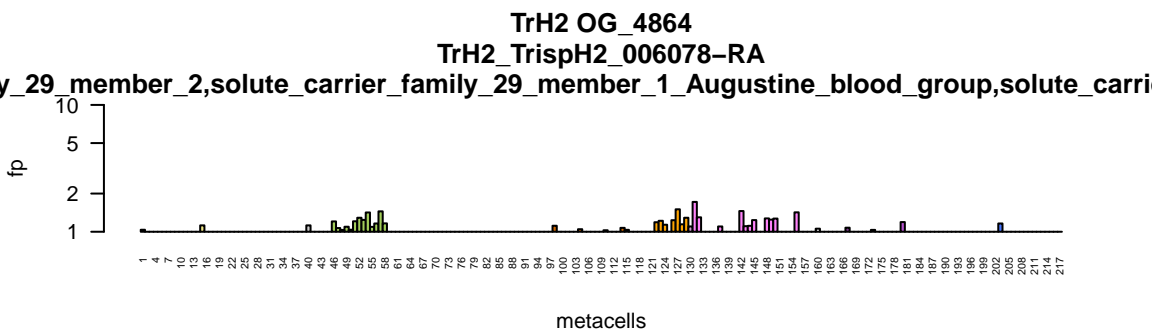
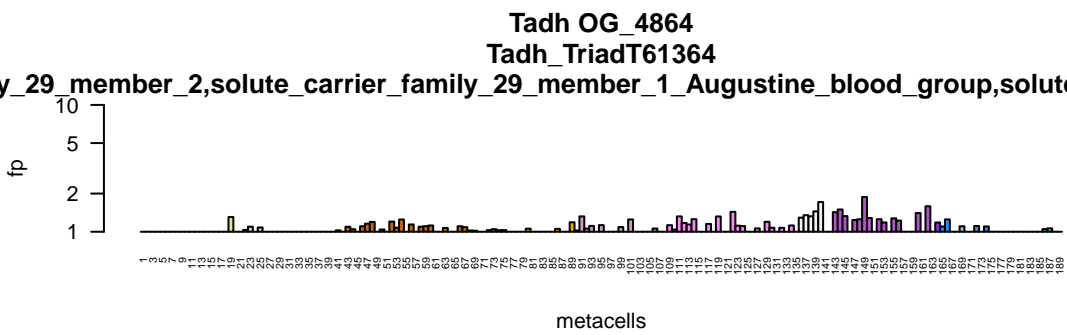
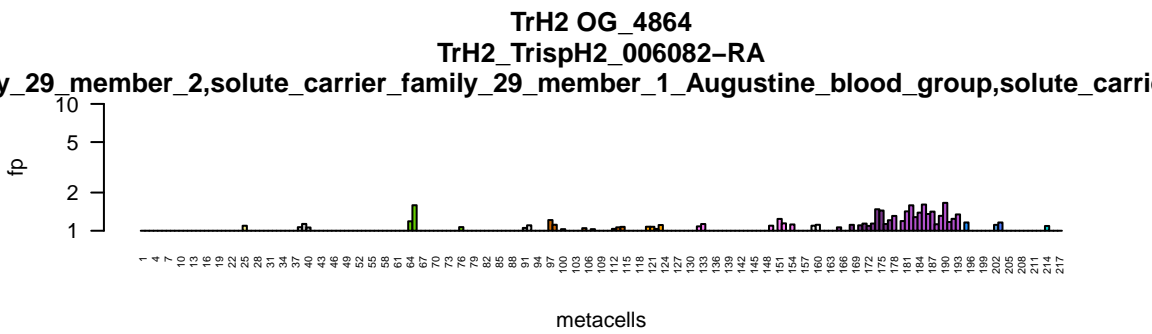
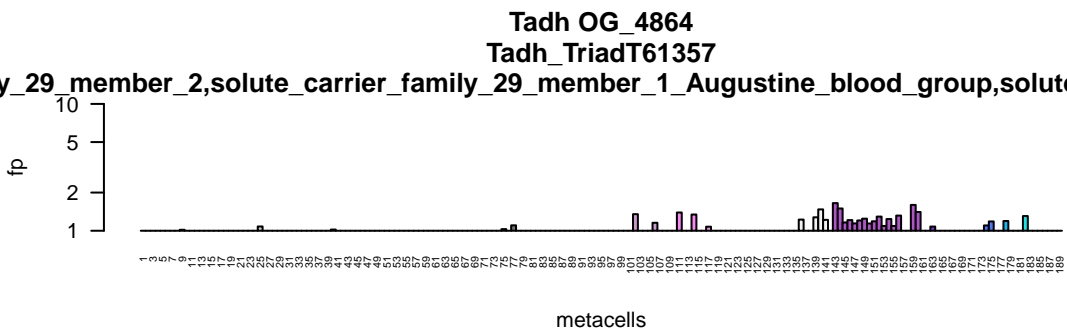
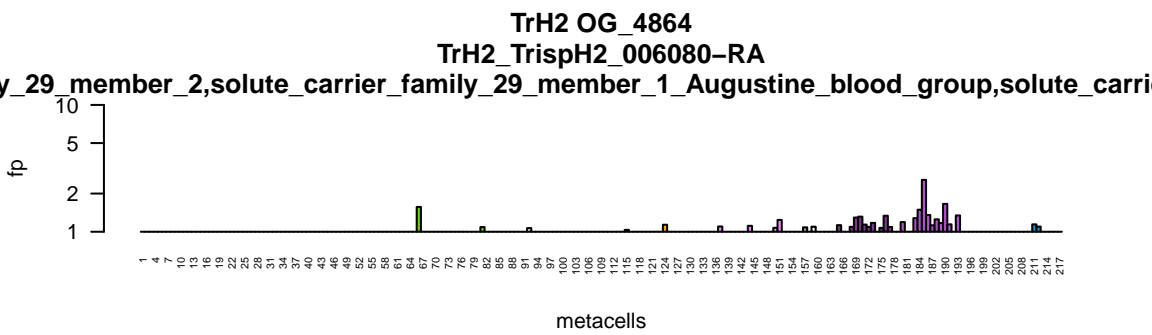
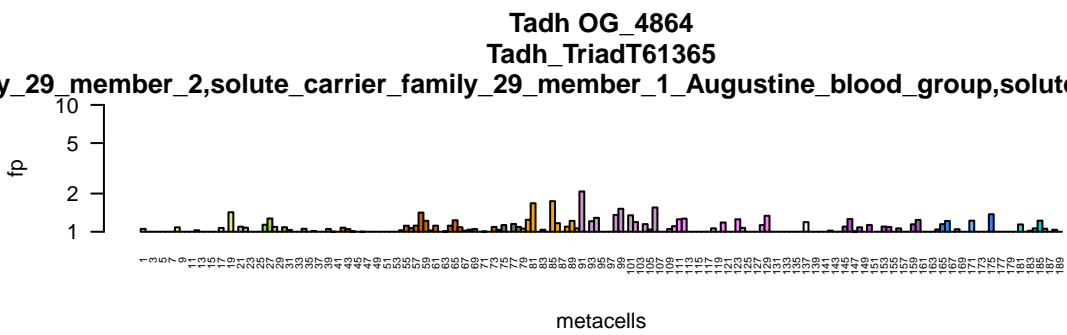
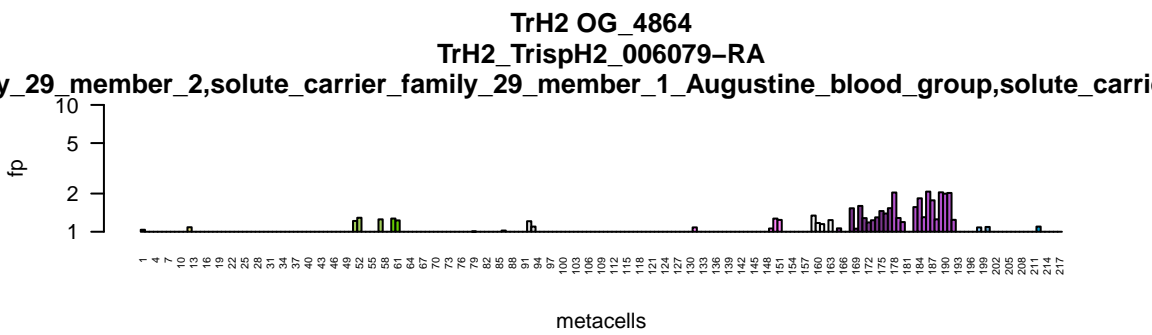
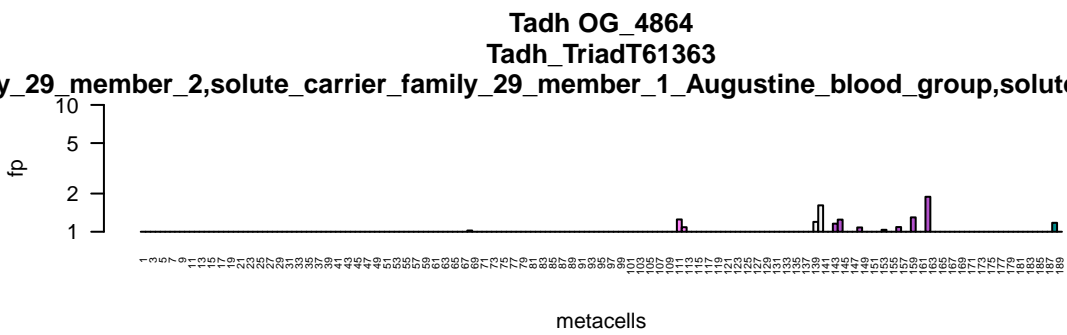
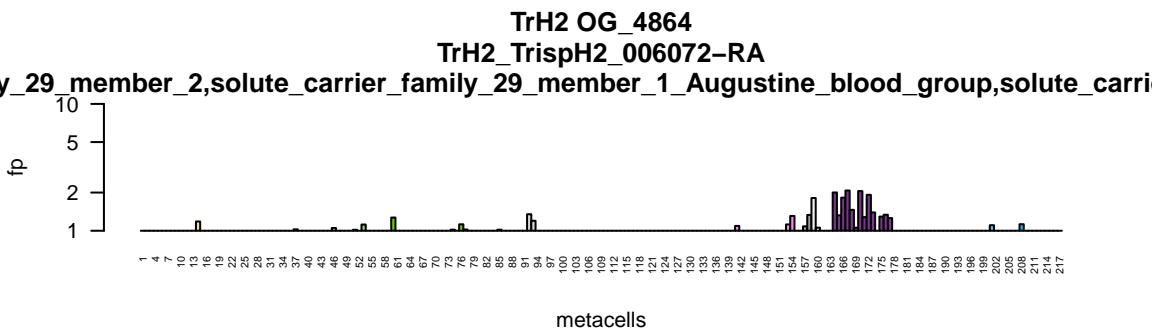
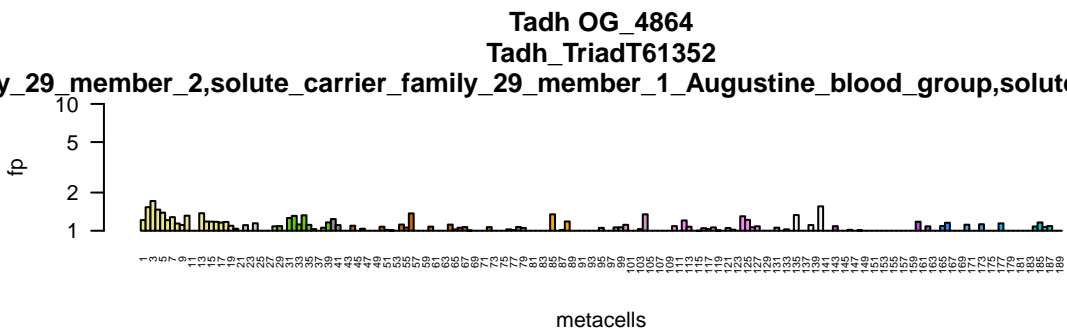
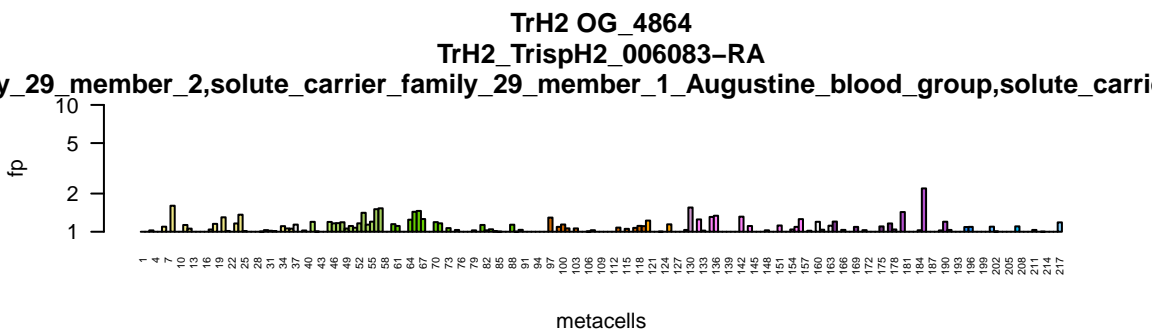
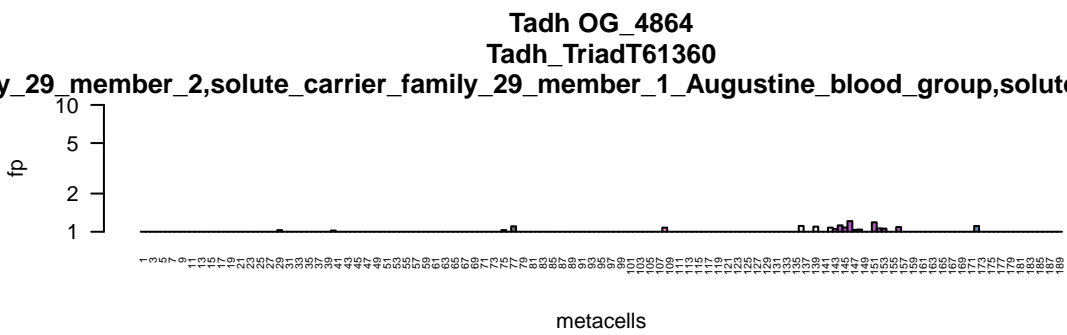
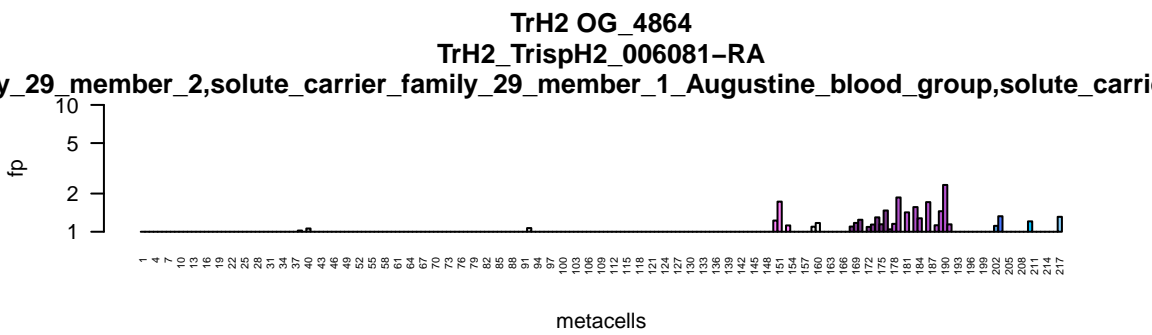
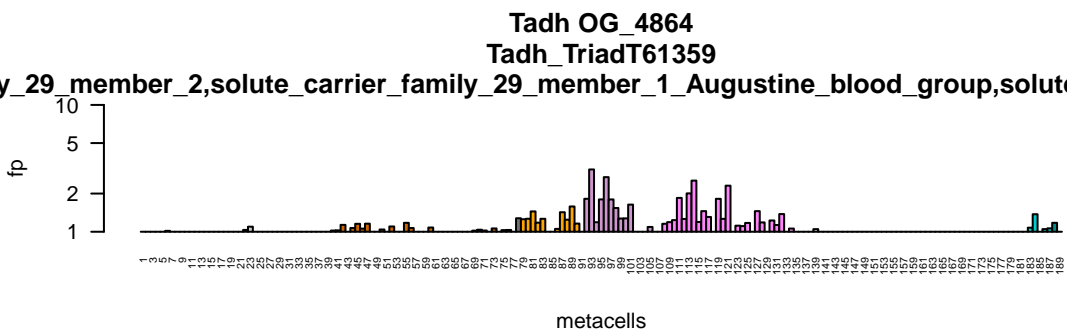
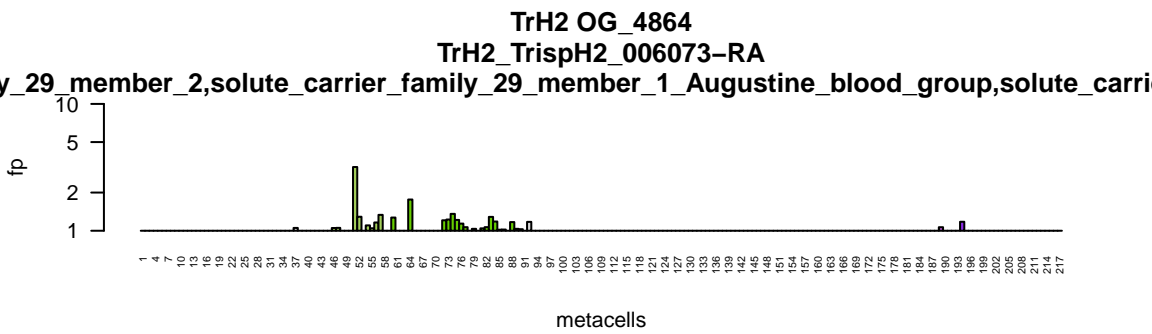
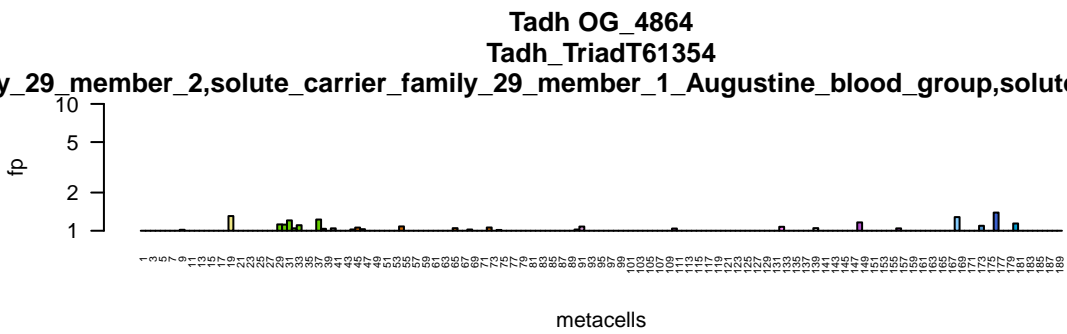
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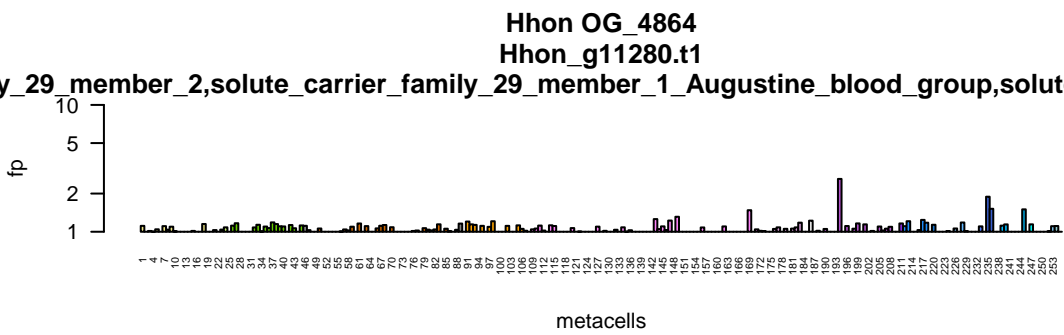
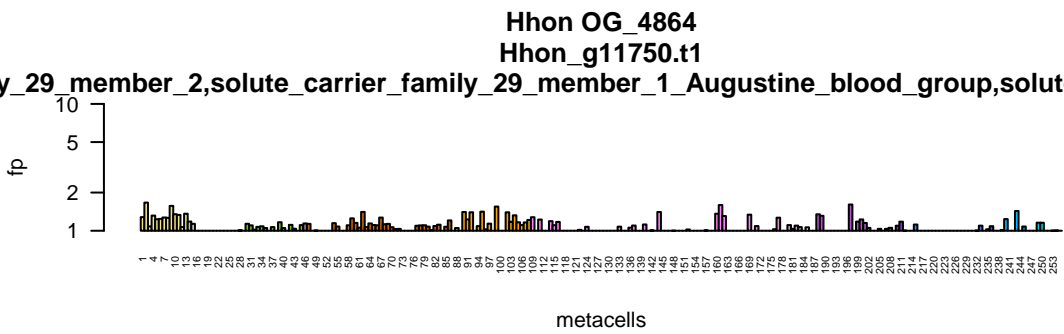
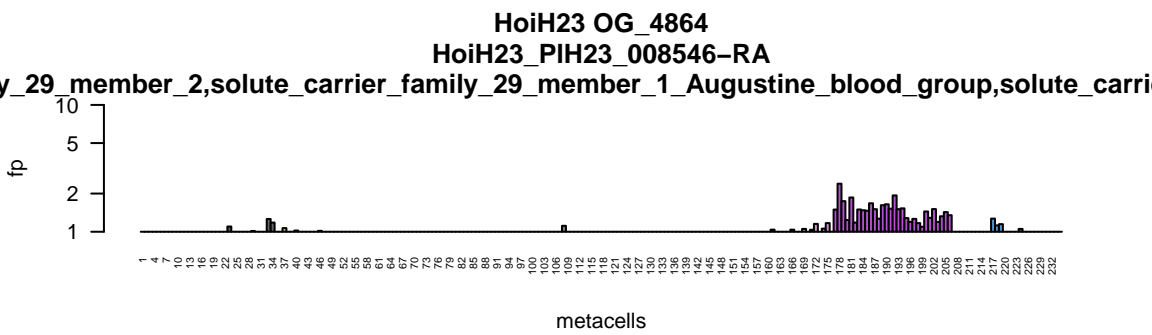
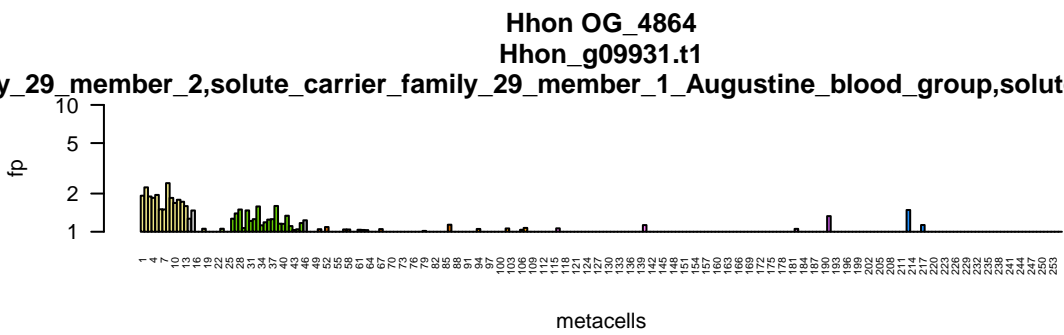
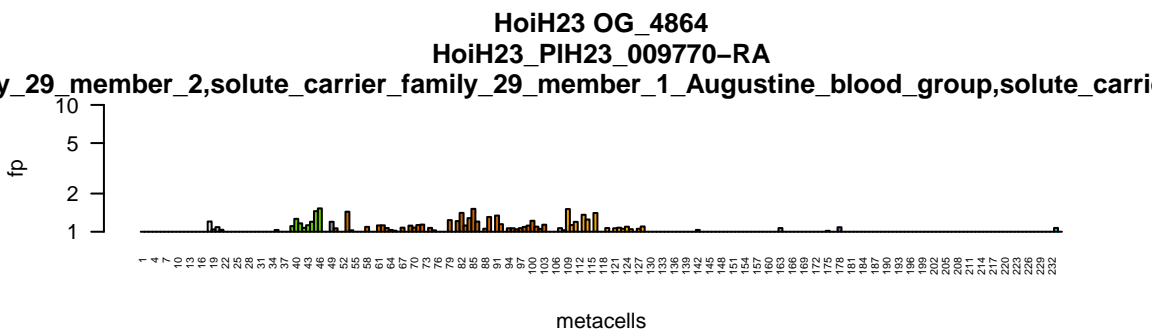
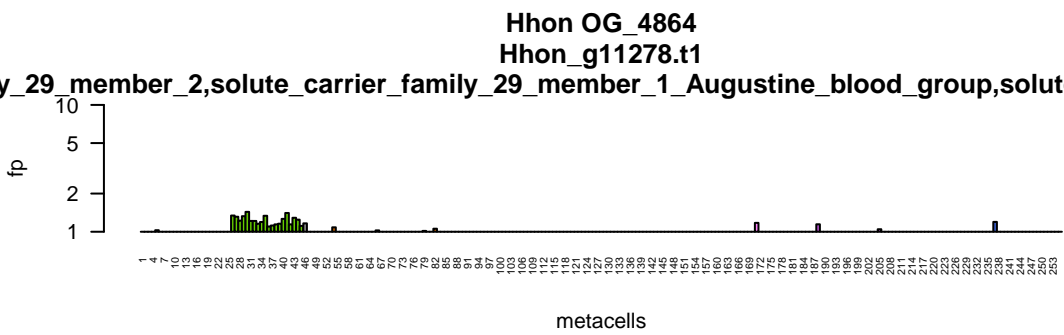
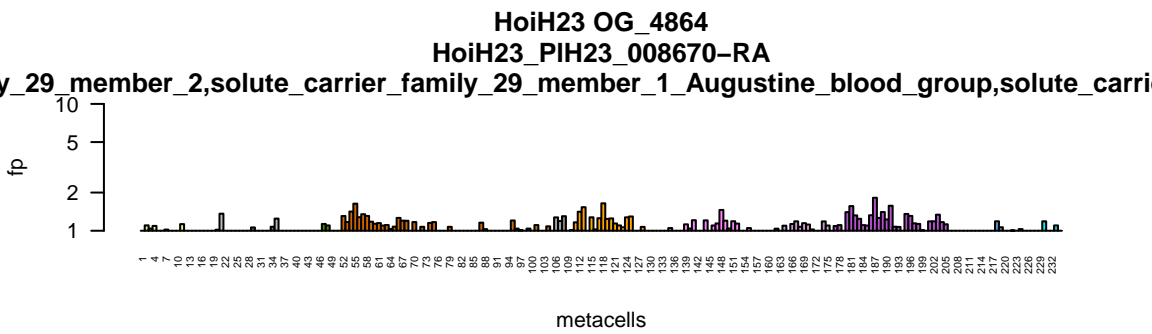
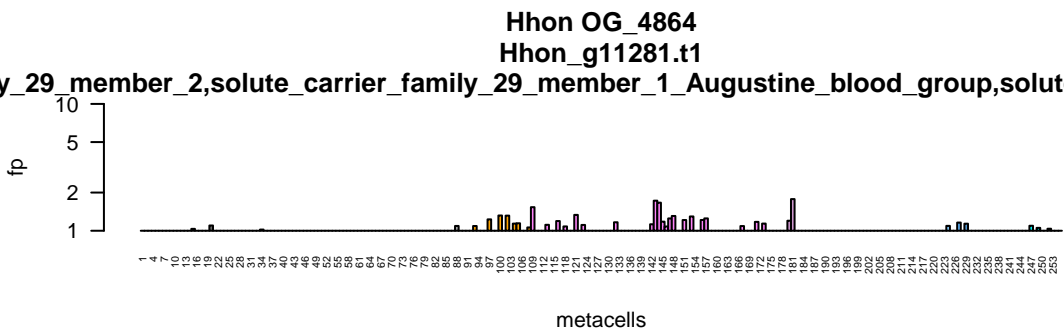
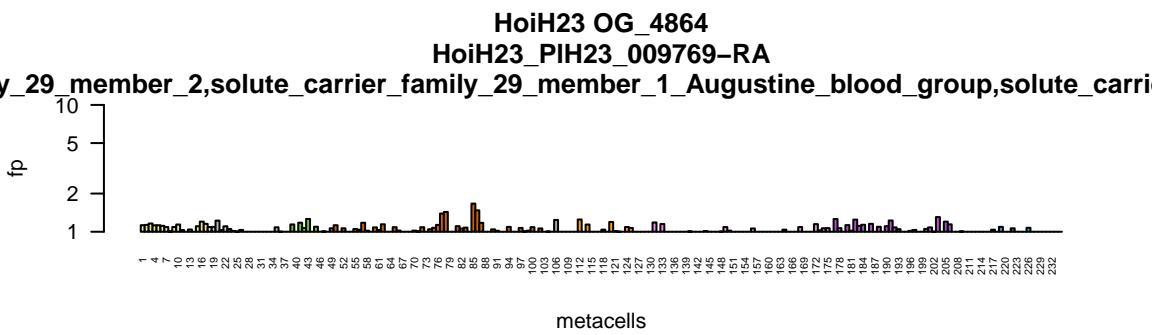
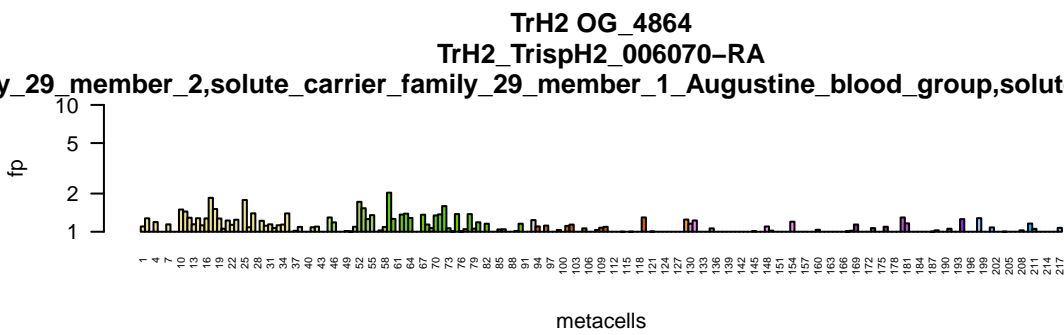
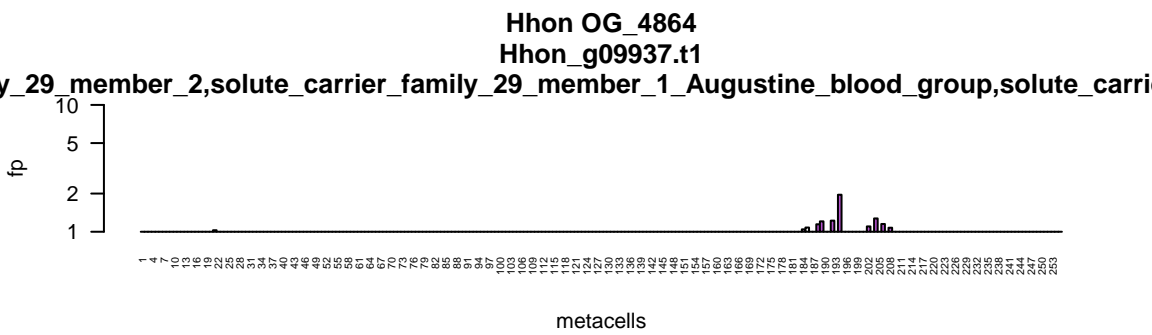
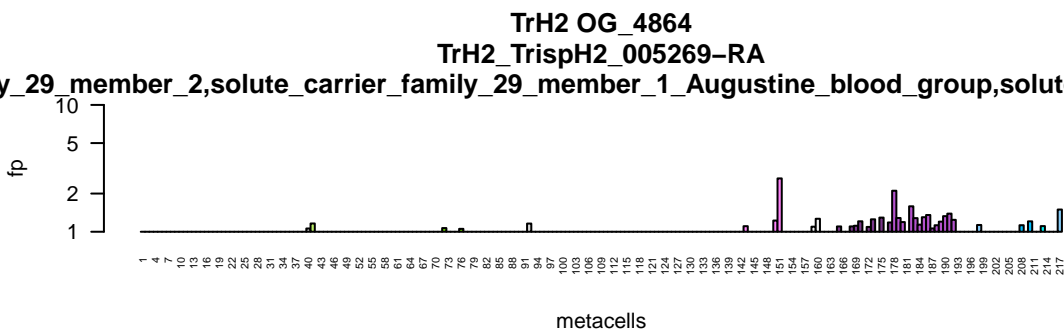
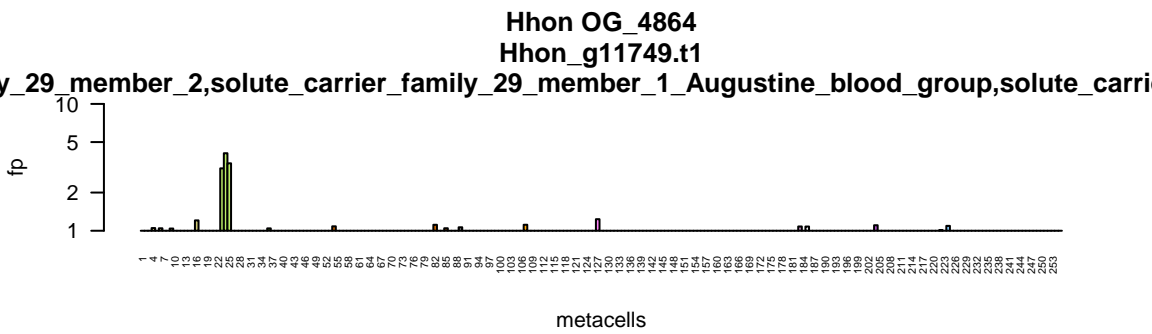
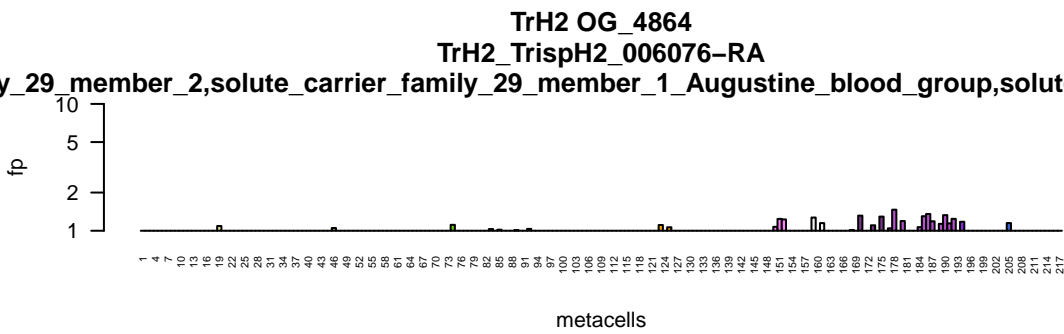
metacells

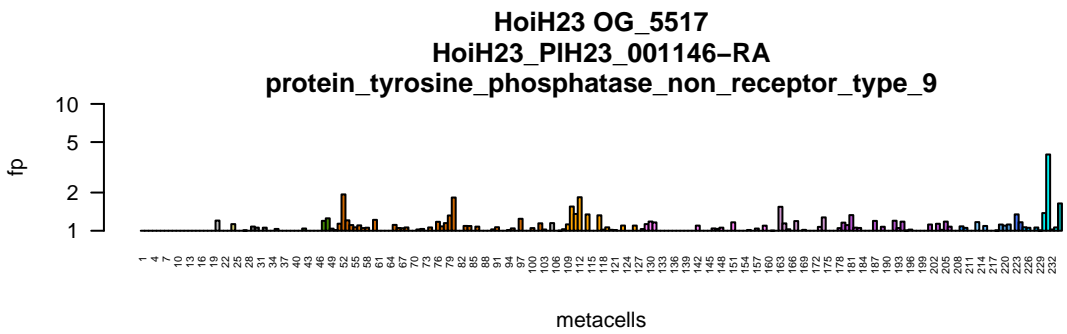
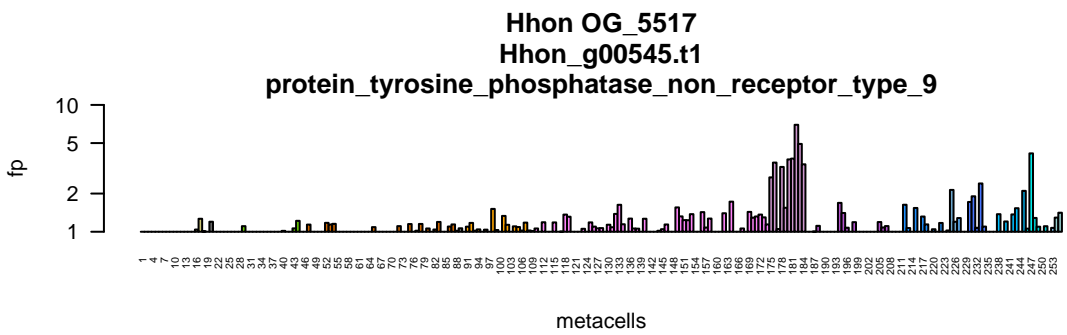
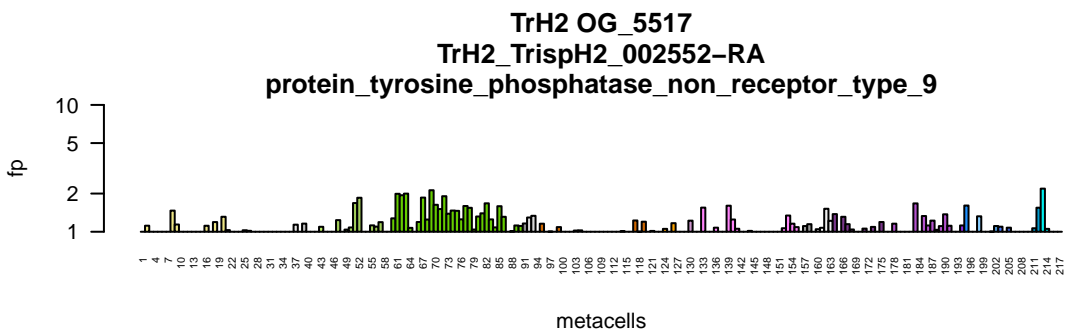
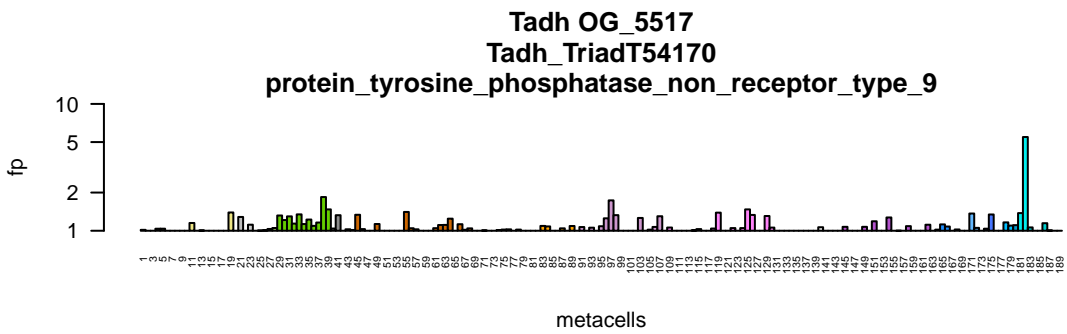
glycoprotein_2A,synaptic_vesicle_glycoprotein_2B,synaptic_vesicle_glycoprotein

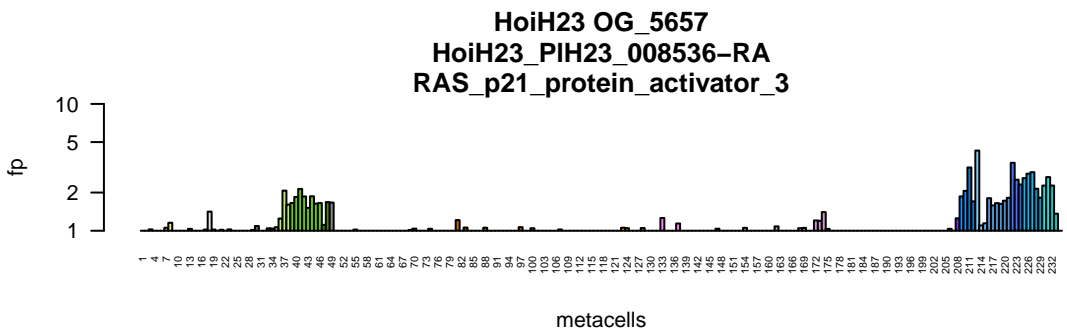
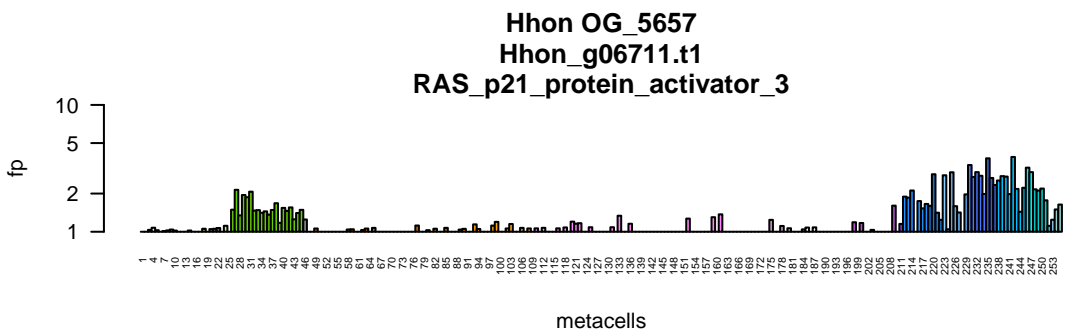
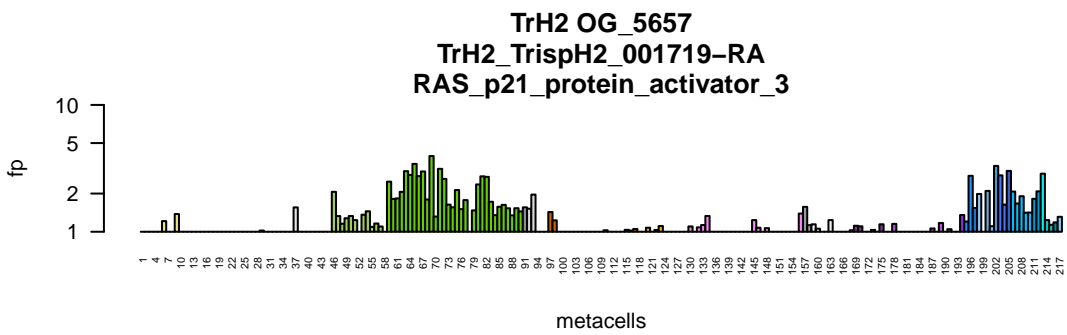
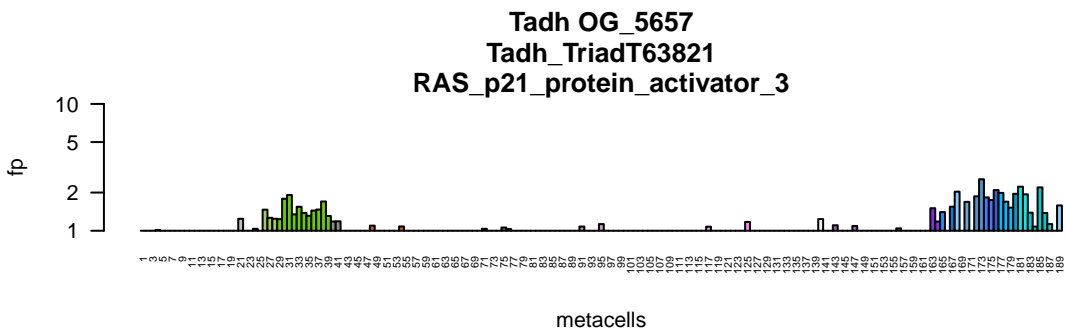
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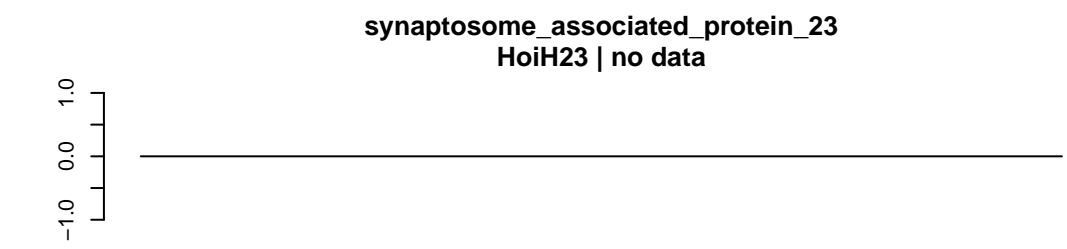
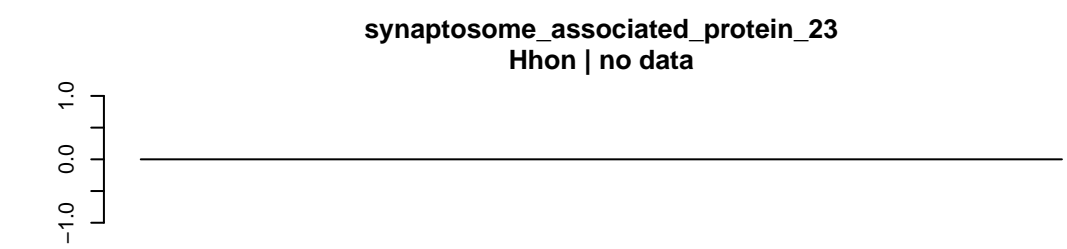
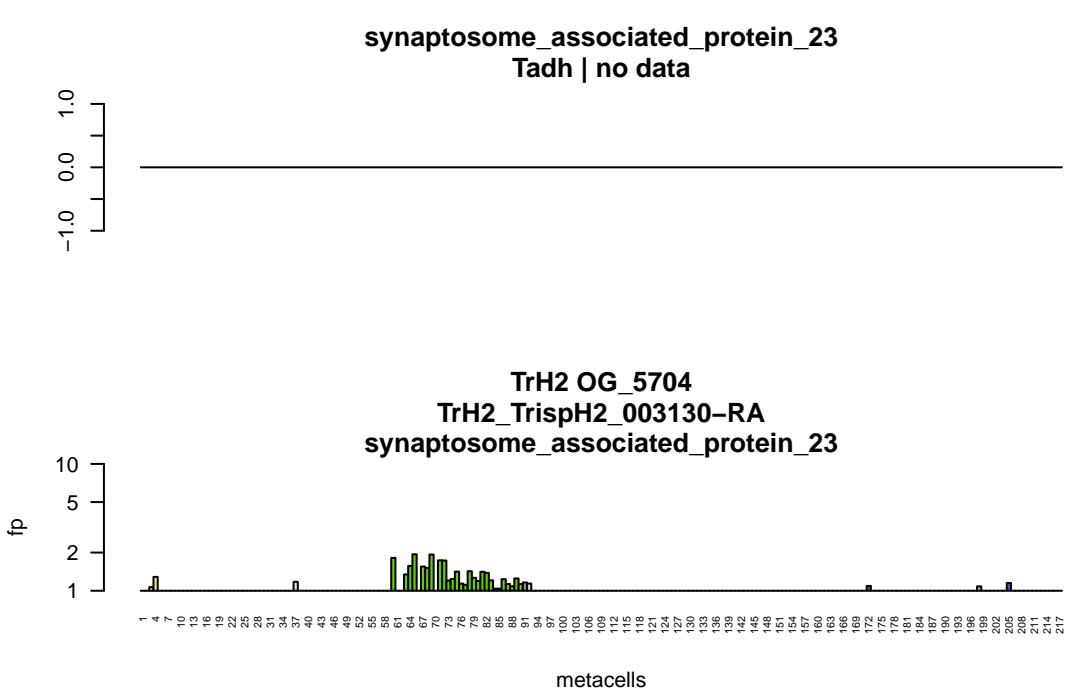


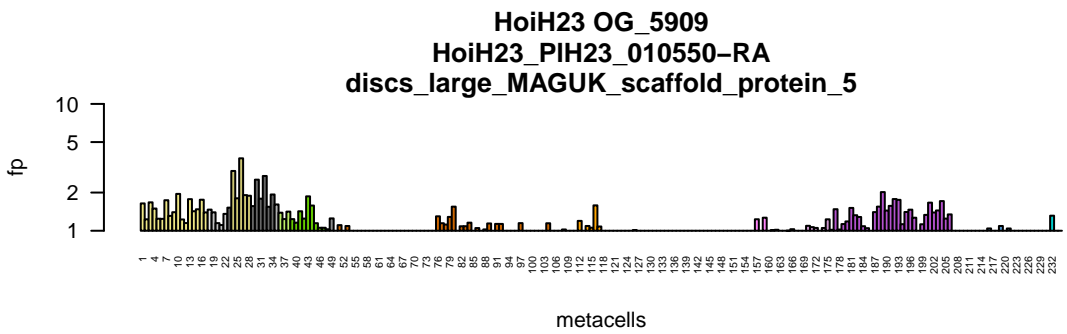
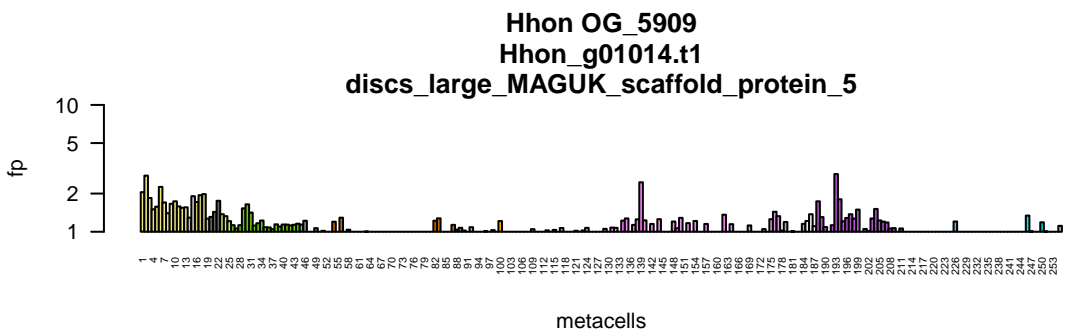
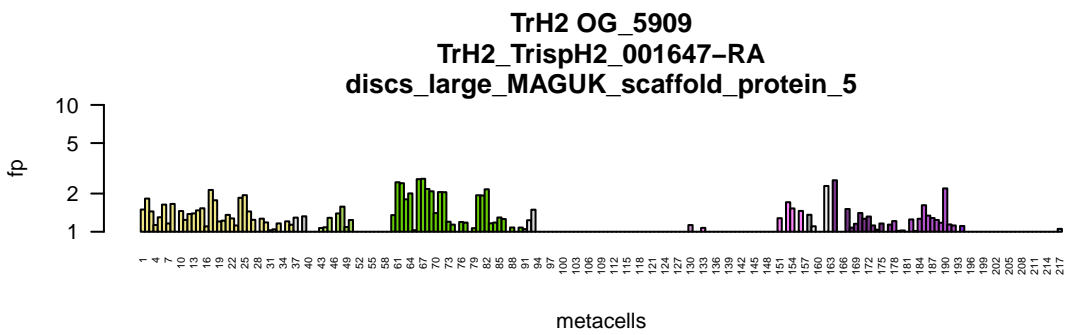
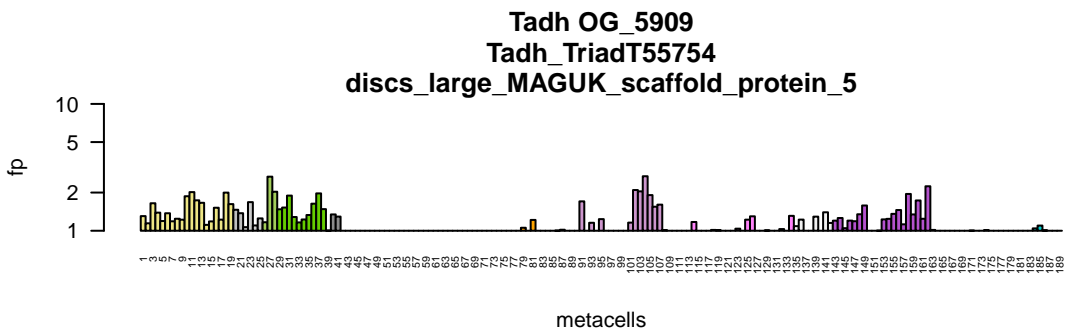




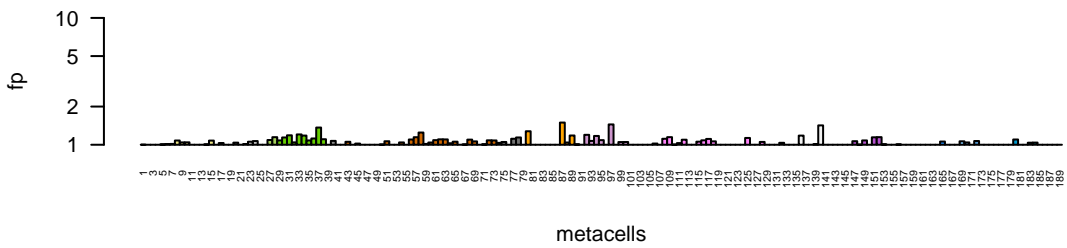




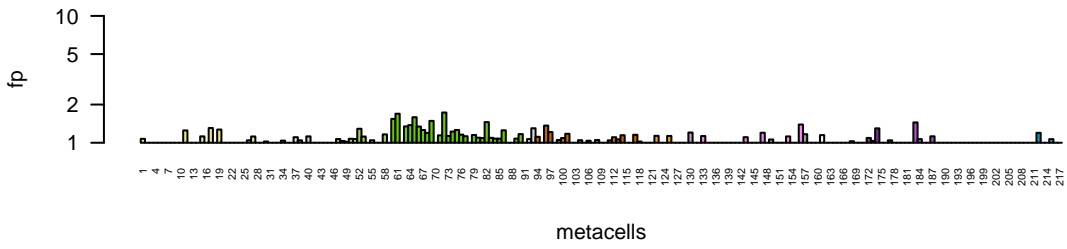




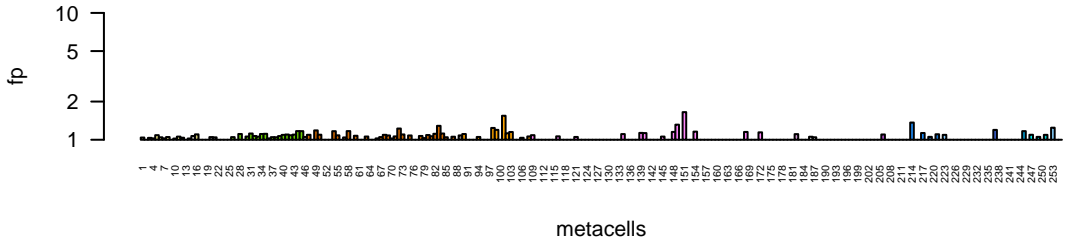
Tadh OG_6248
Tadh_TriadT59935



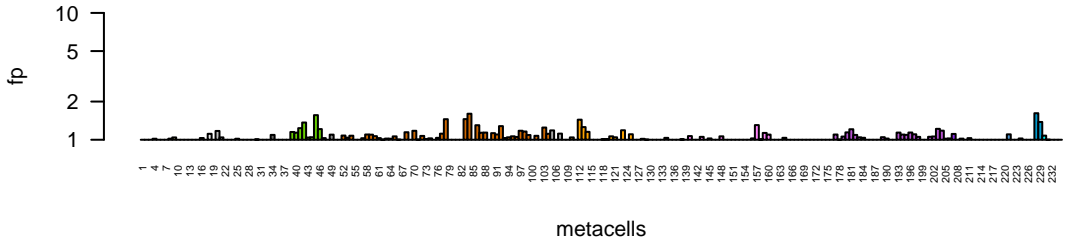
TrH2 OG_6248
TrH2_TrispH2_007622-RA



Hhon OG_6248
Hhon_g09079.t1



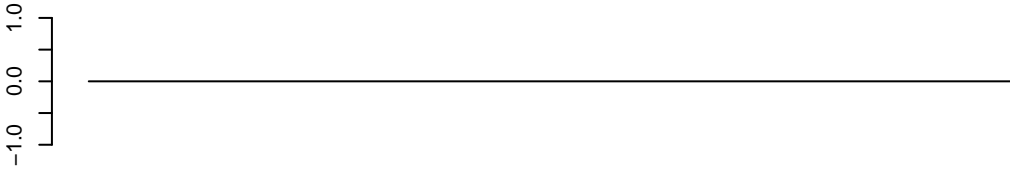
HoiH23 OG_6248
HoiH23_PIH23_005506-RA



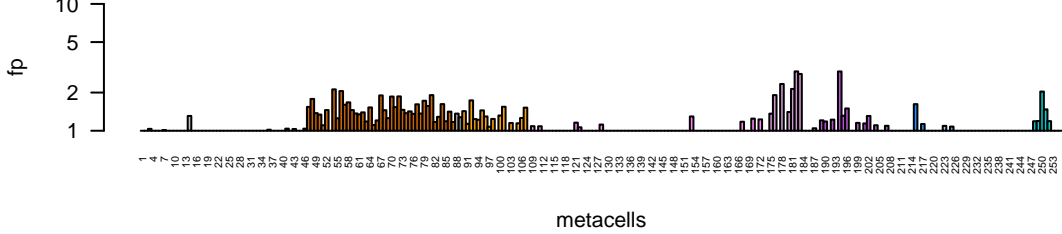
solute_carrier_family_6_member_12
Tadh | no data



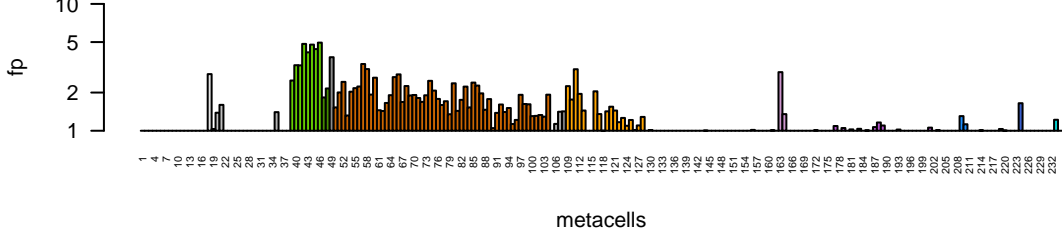
solute_carrier_family_6_member_12
TrH2 | no data

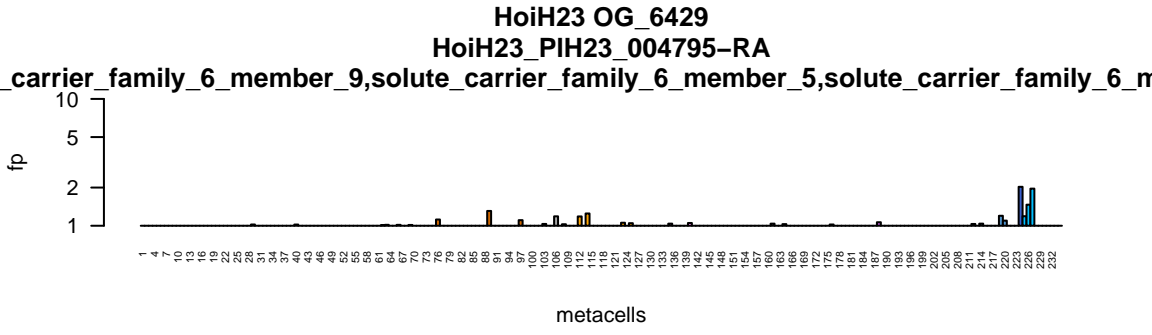
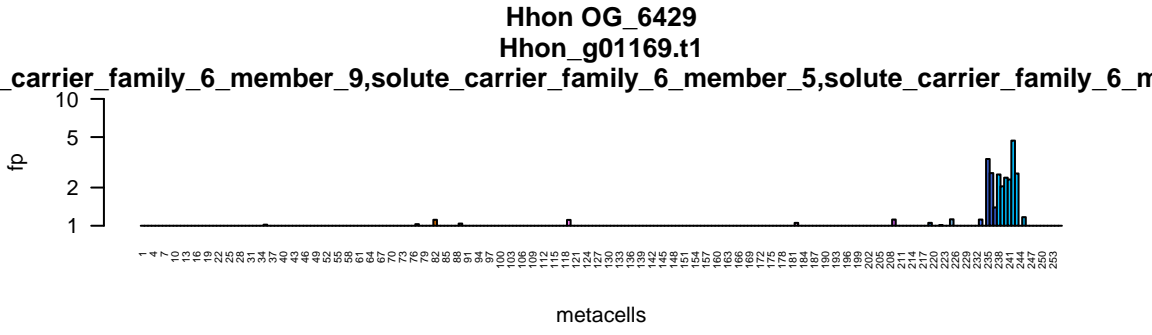
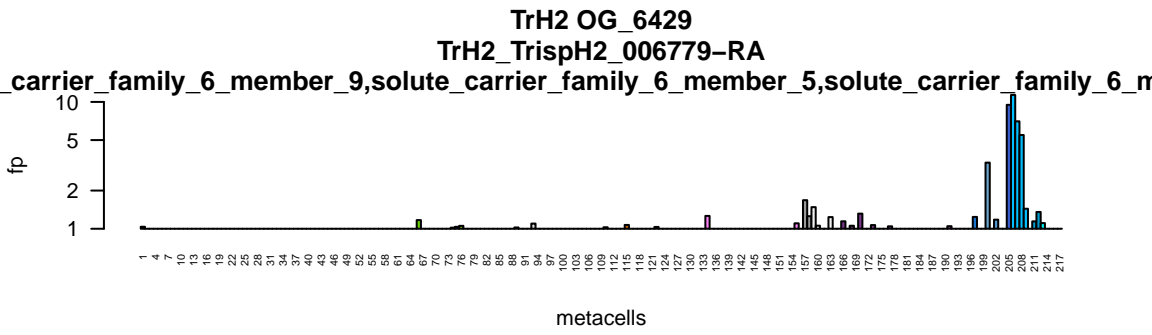
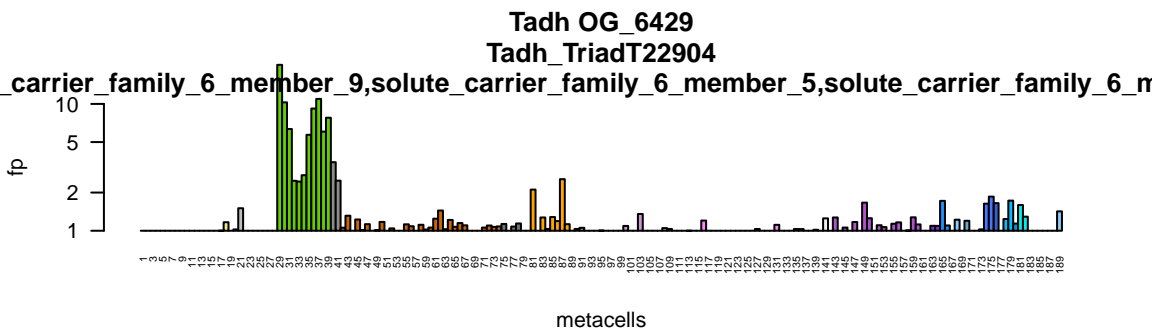


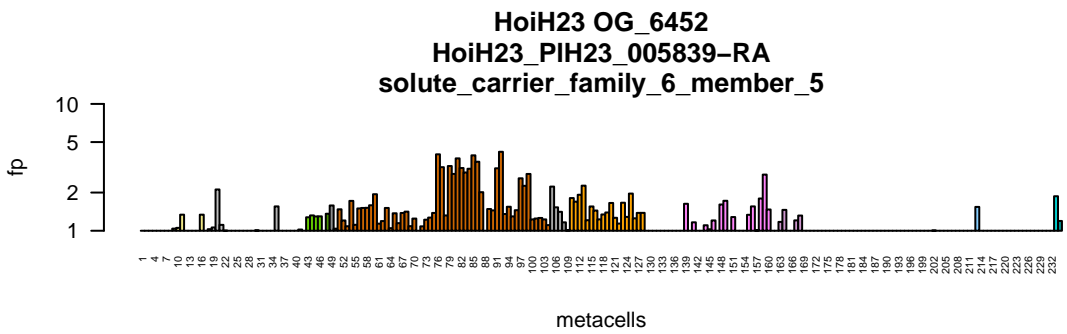
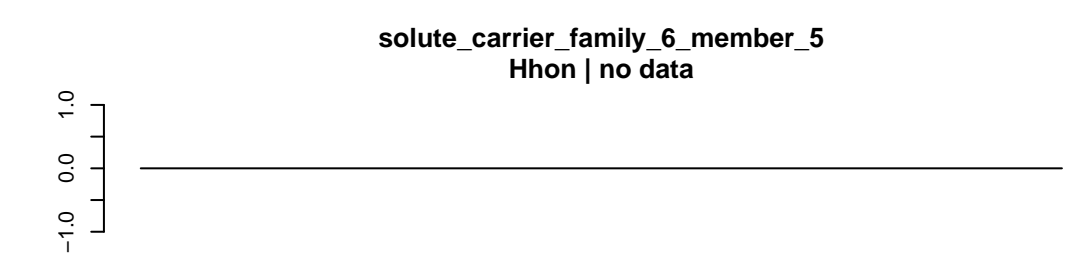
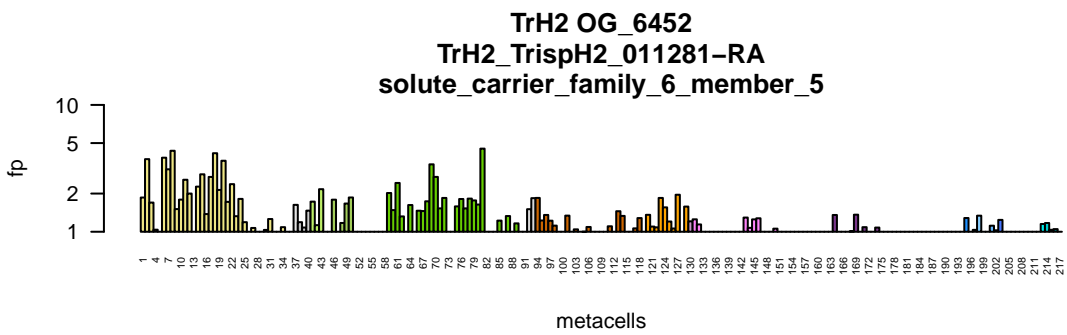
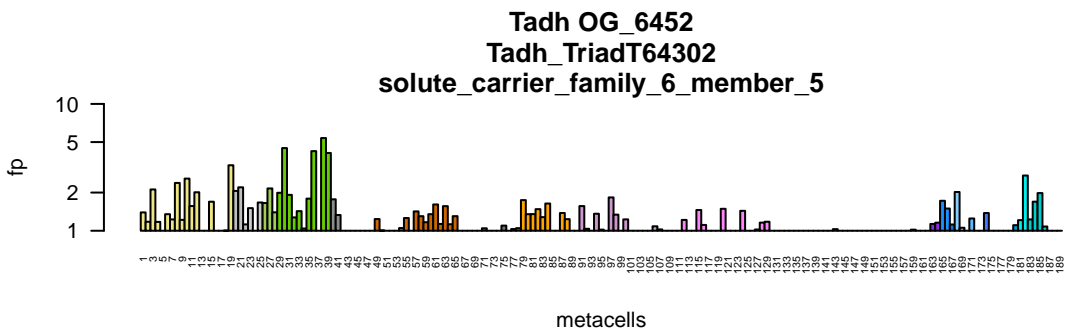
Hhon OG_6428
Hhon_g01172.t1
solute_carrier_family_6_member_12



HoiH23 OG_6428
HoiH23_PIH23_004794-RA
solute_carrier_family_6_member_12







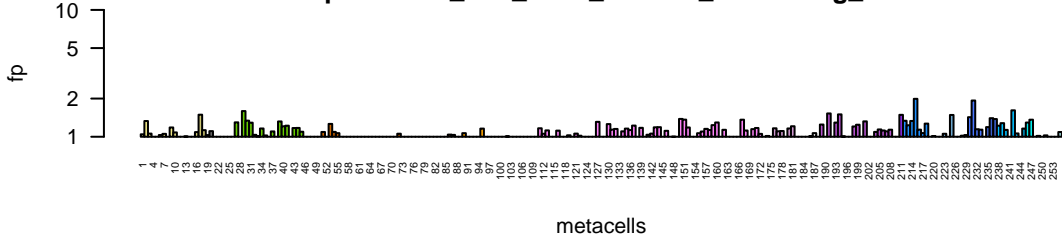
pleckstrin_and_Sec7_domain_containing_2
Tadh | no data



pleckstrin_and_Sec7_domain_containing_2
TrH2 | no data

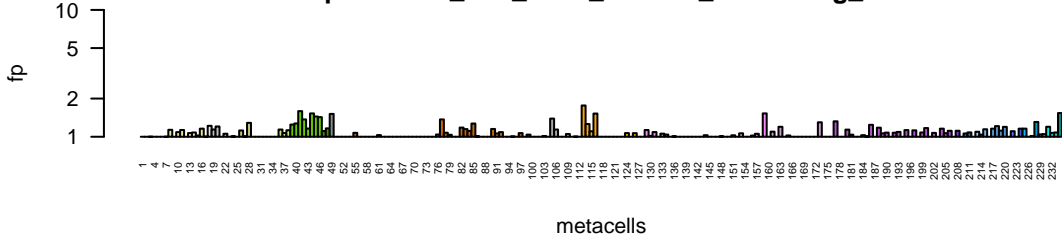


Hhon OG_6544
Hhon_g09899.t1
pleckstrin_and_Sec7_domain_containing_2

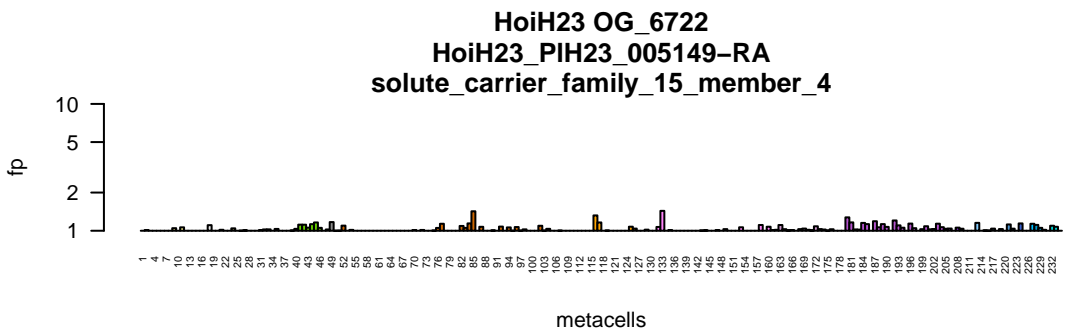
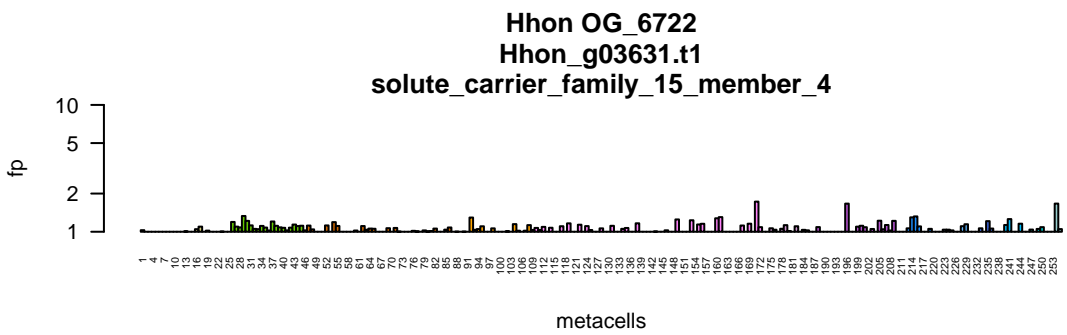
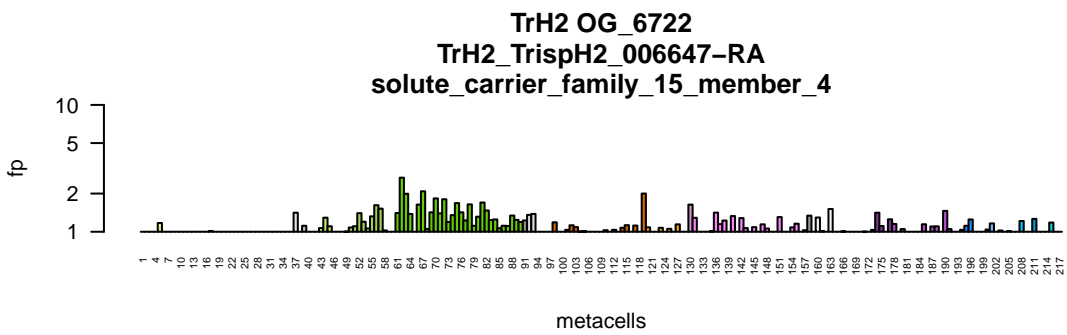
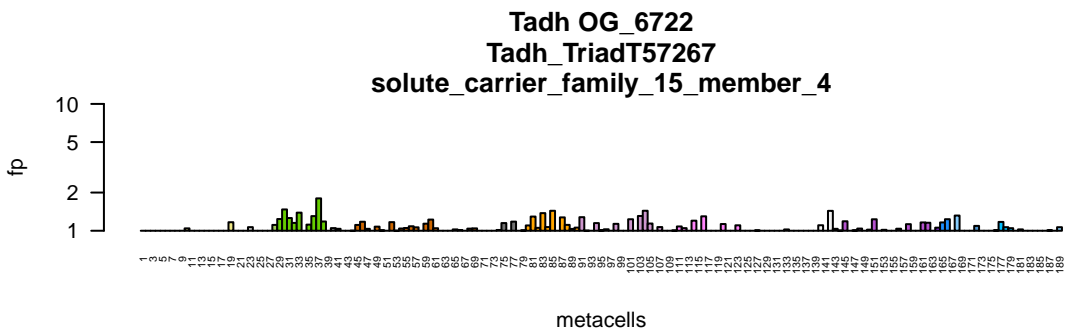


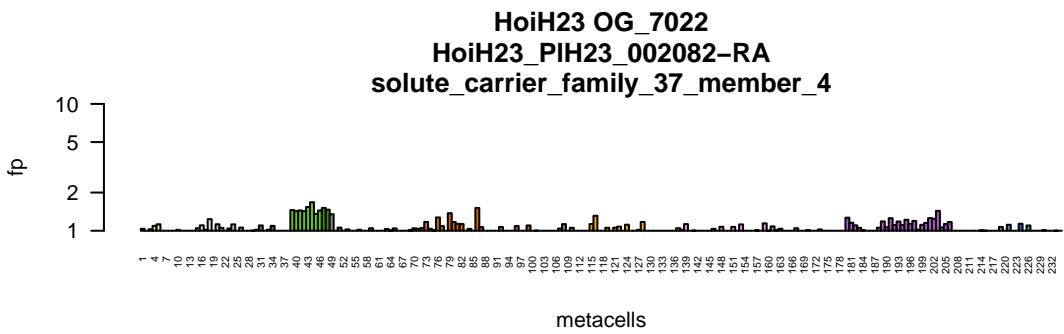
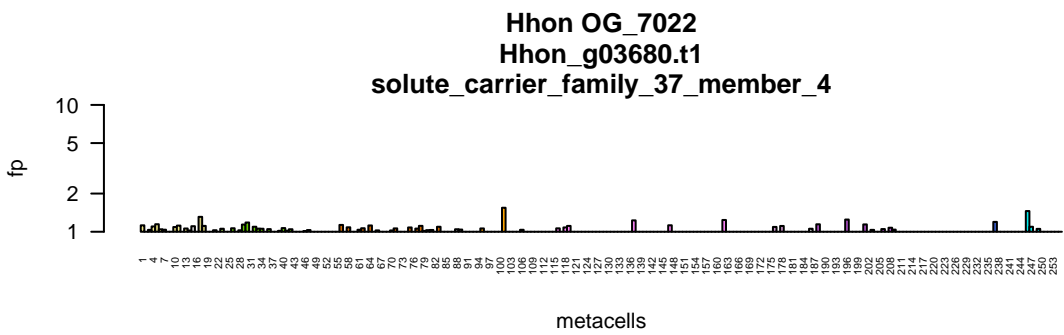
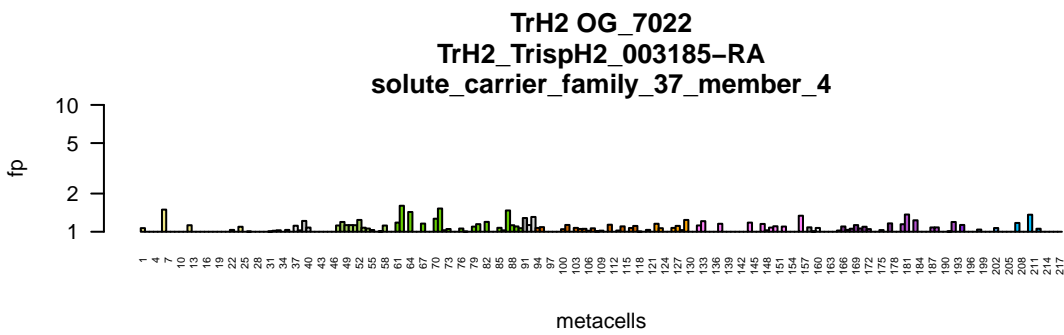
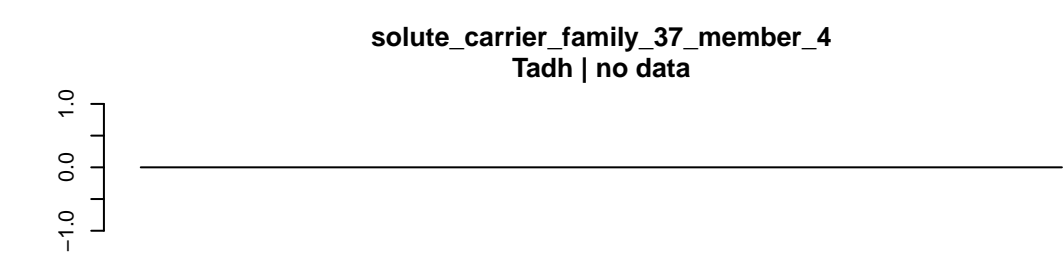
metacells

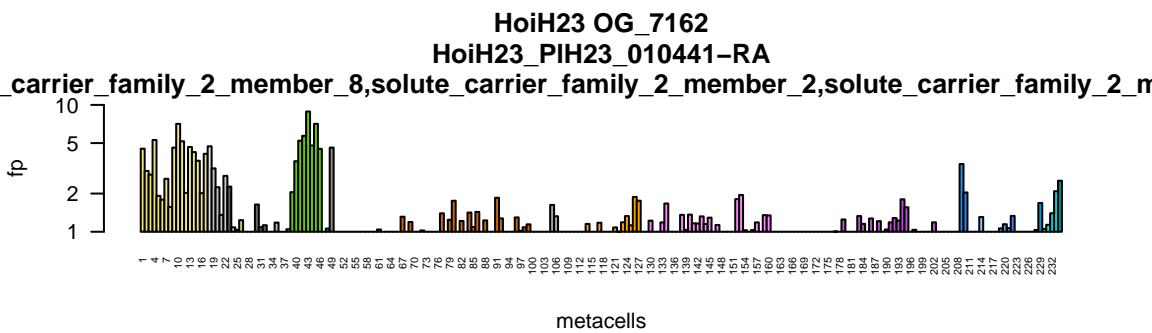
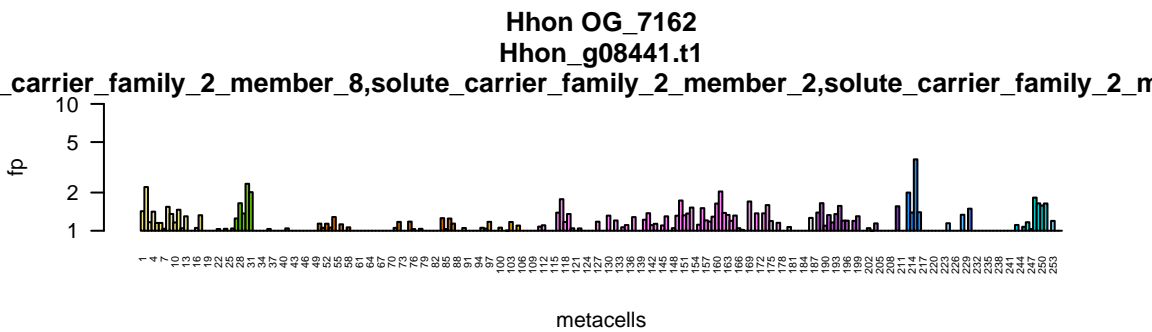
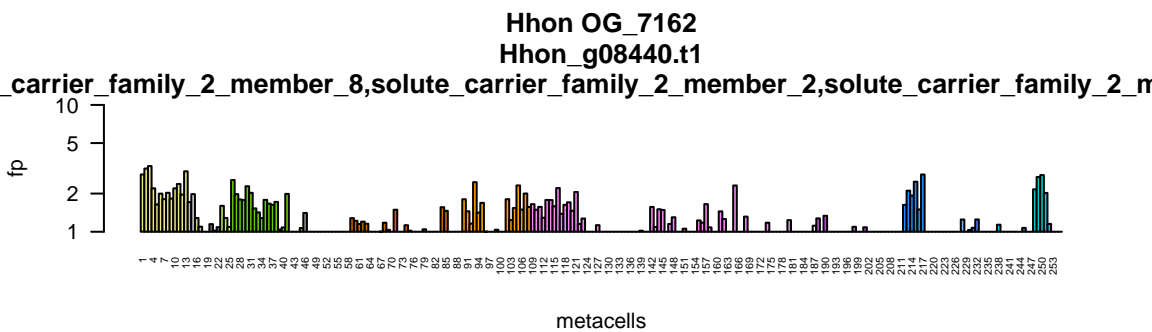
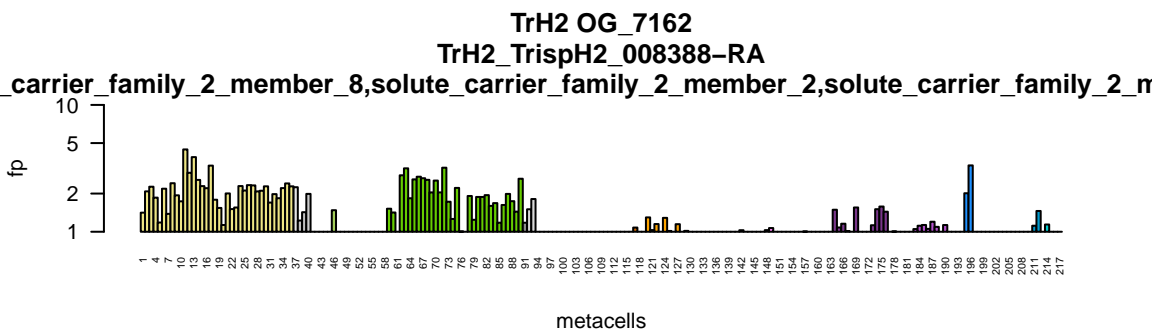
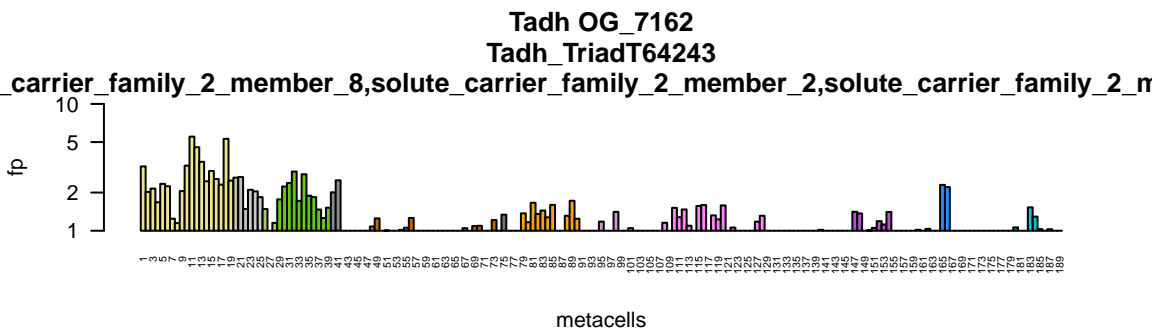
HoiH23 OG_6544
HoiH23_PIH23_000592-RA
pleckstrin_and_Sec7_domain_containing_2



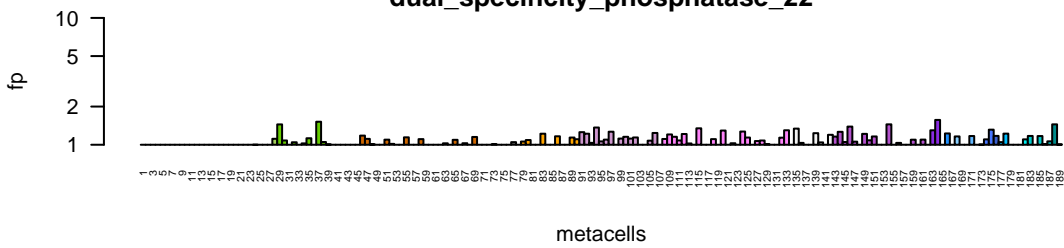
metacells



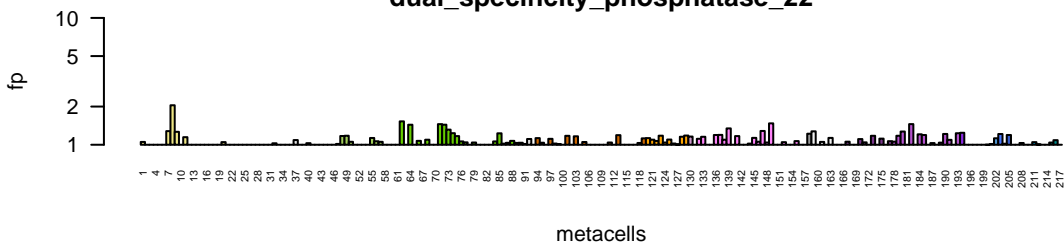




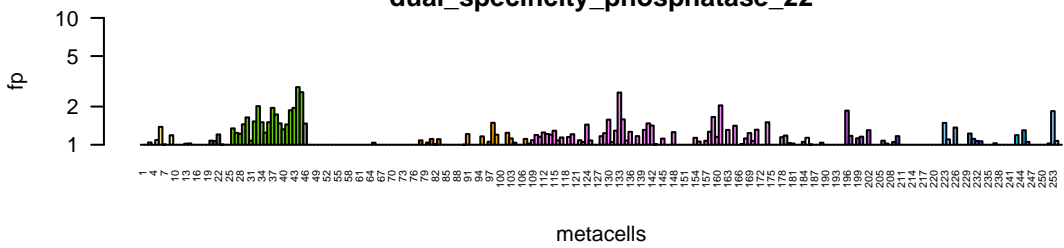
Tadh OG_7170
Tadh_TriadT60150
dual_specificity_phosphatase_22



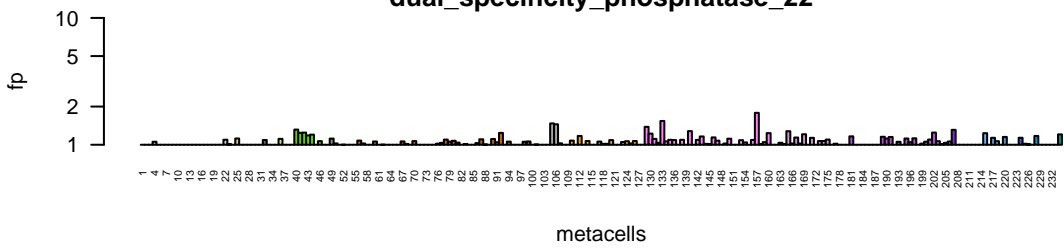
TrH2 OG_7170
TrH2_TrispH2_000712-RA
dual_specificity_phosphatase_22

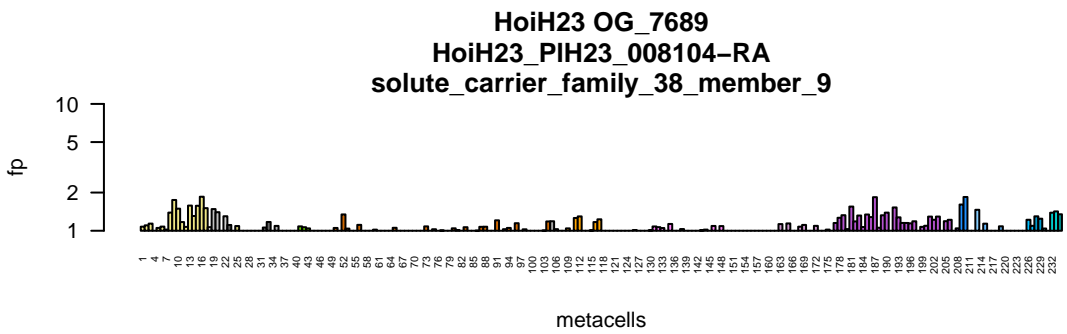
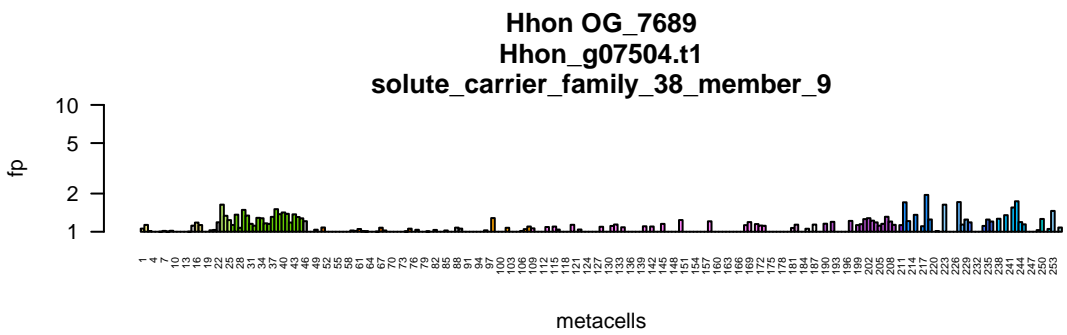
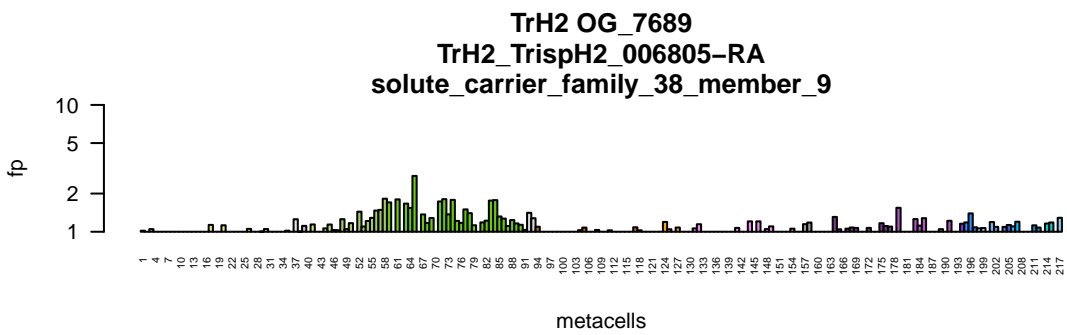
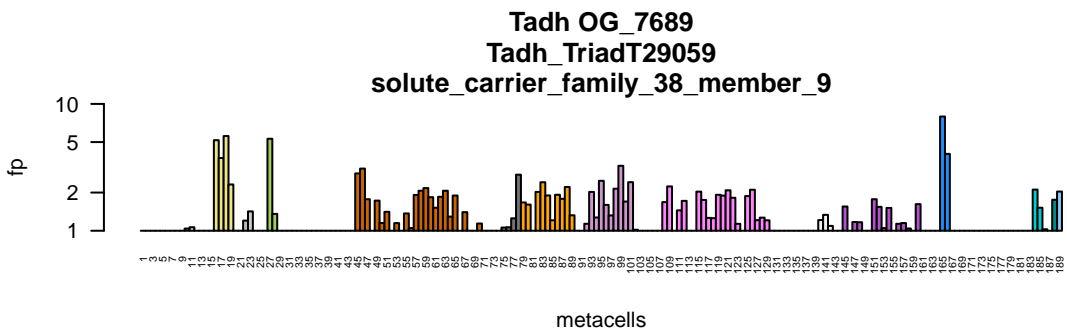


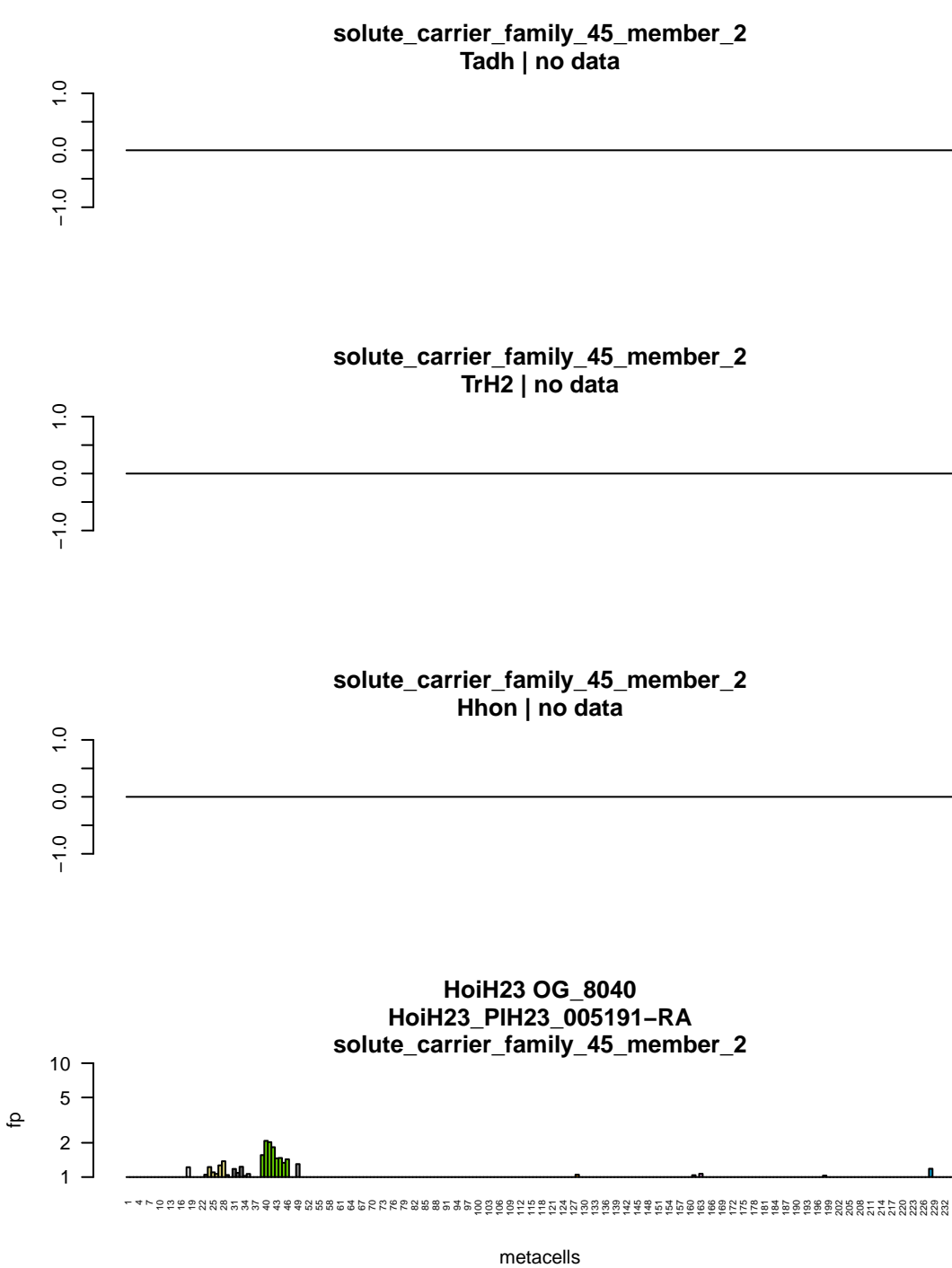
Hhon OG_7170
Hhon_g05020.t1
dual_specificity_phosphatase_22

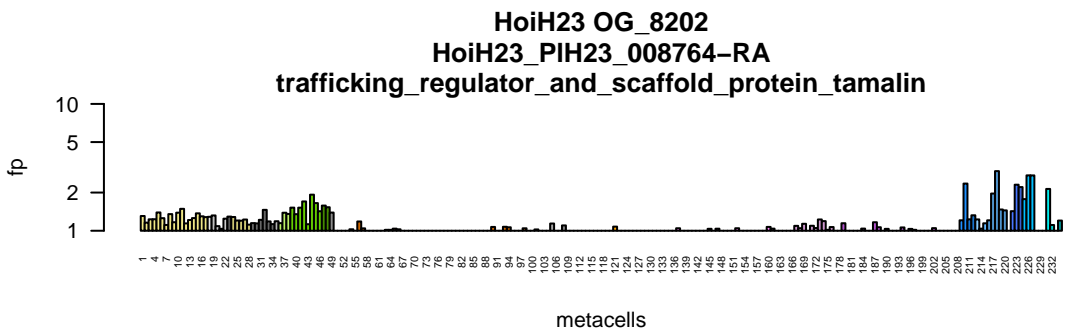
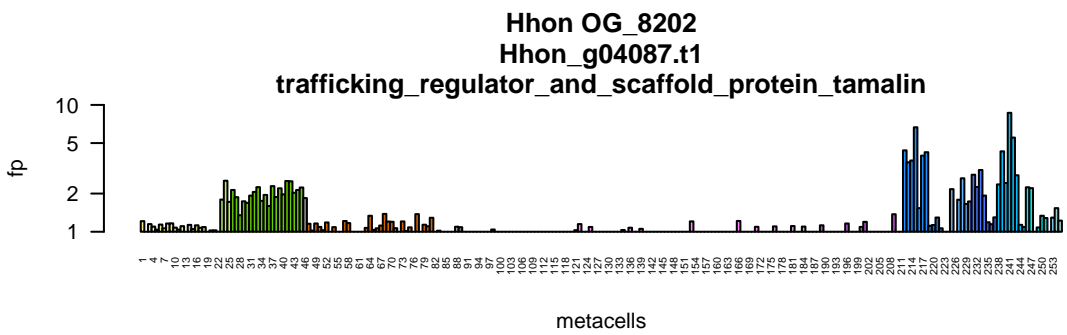
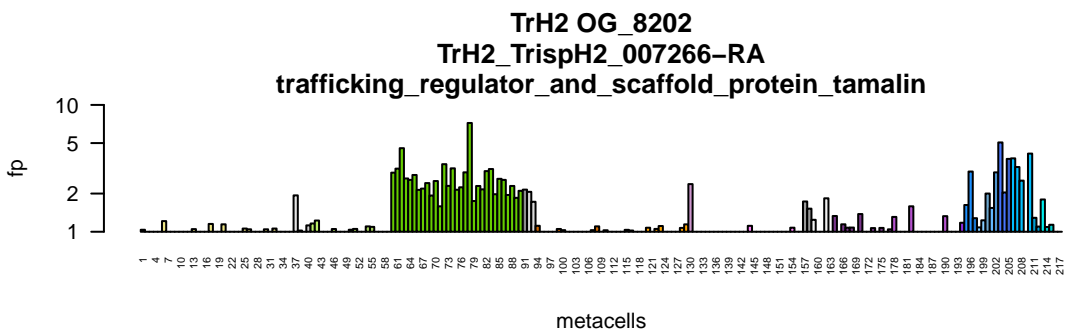
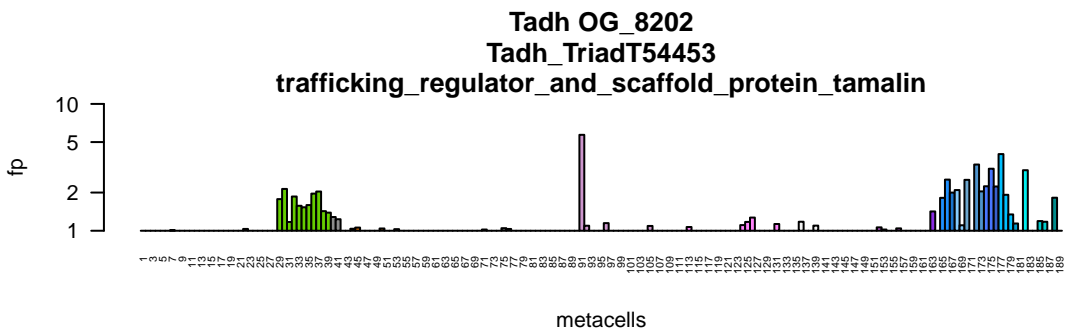


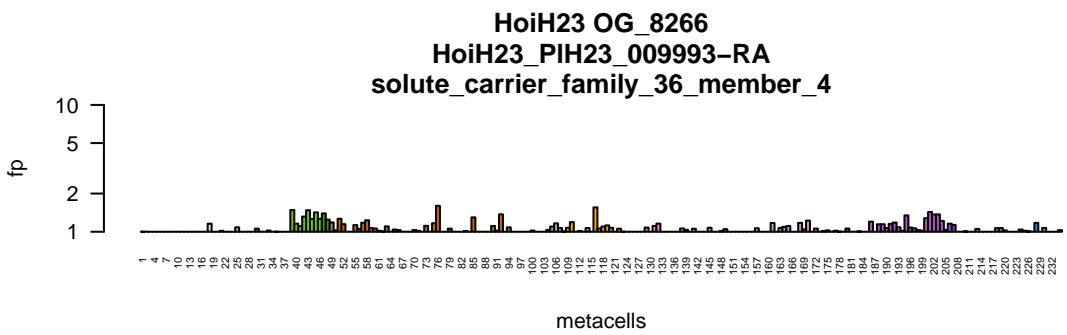
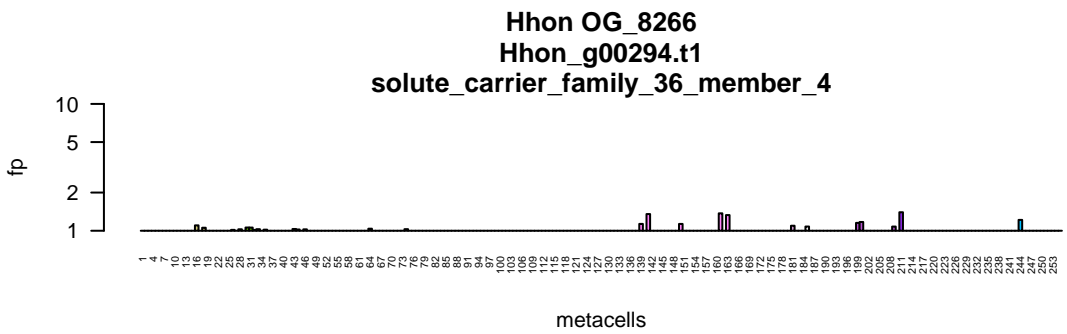
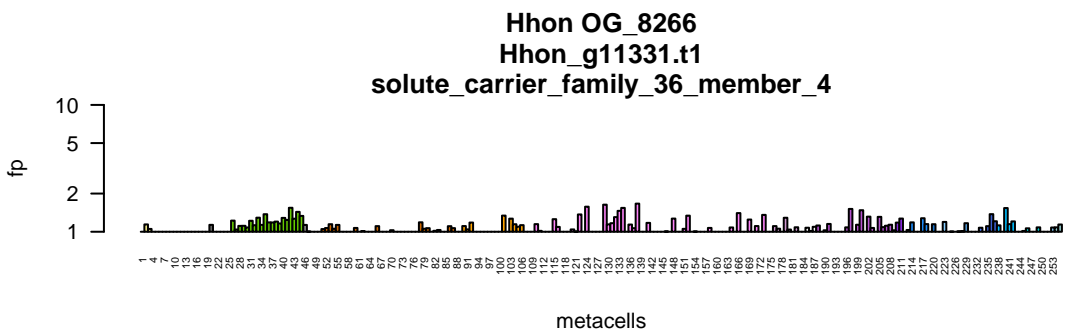
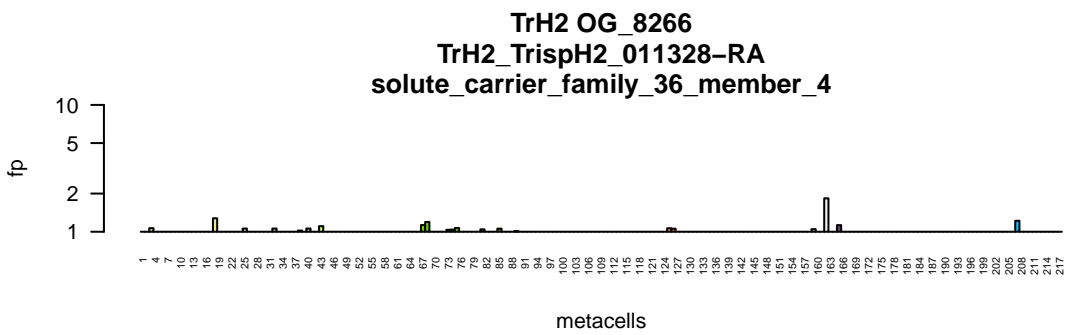
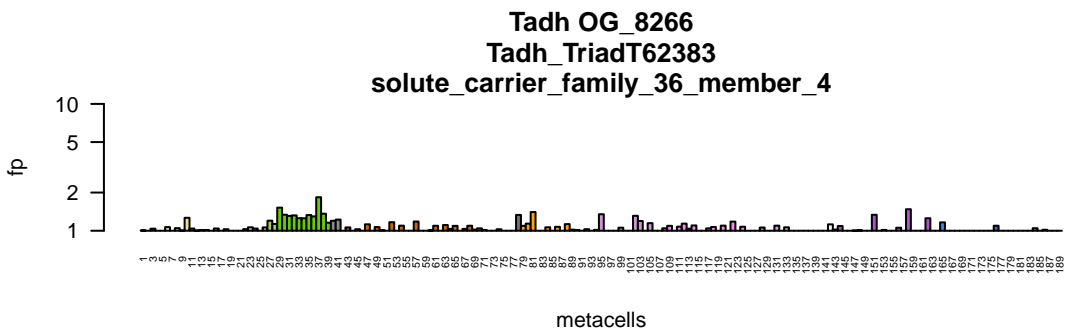
HoiH23 OG_7170
HoiH23_PIH23_008031-RA
dual_specificity_phosphatase_22



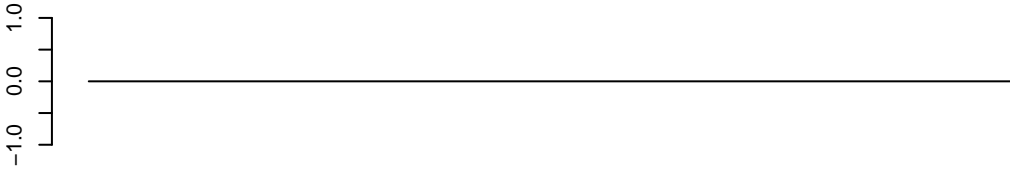




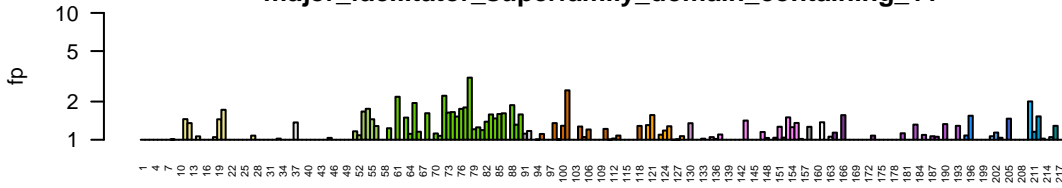




major_facilitator_superfamily_domain_containing_11
Tadh | no data

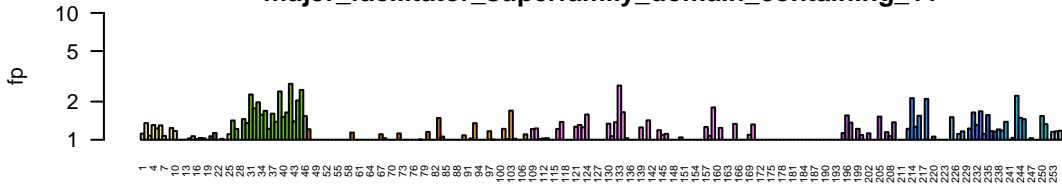


TrH2 OG_8331
TrH2_TrispH2_009301-RA
major_facilitator_superfamily_domain_containing_11



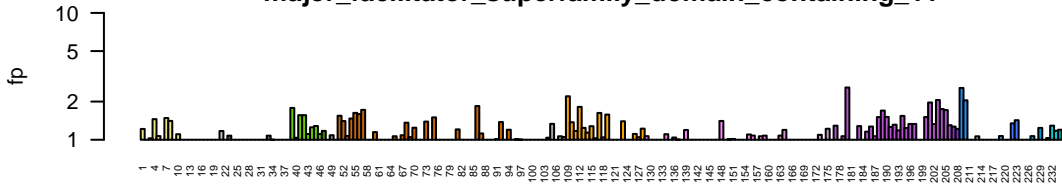
metacells

Hhon OG_8331
Hhon_g01417.t1
major_facilitator_superfamily_domain_containing_11

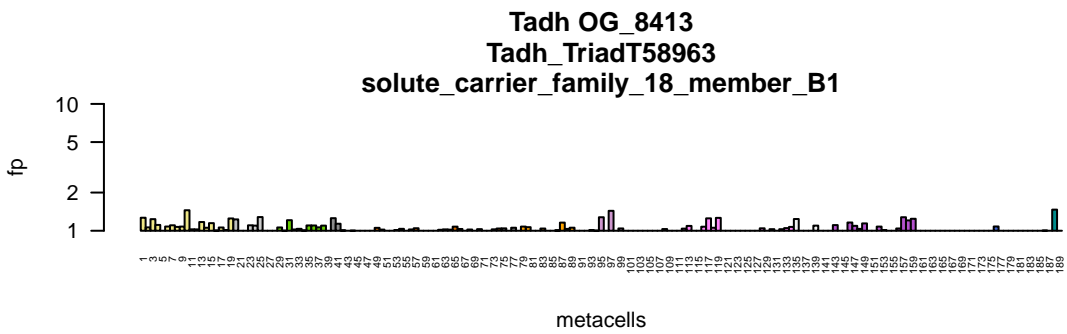


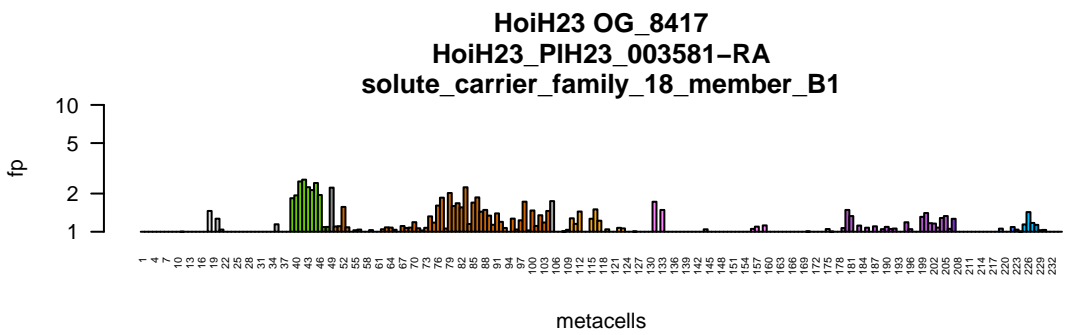
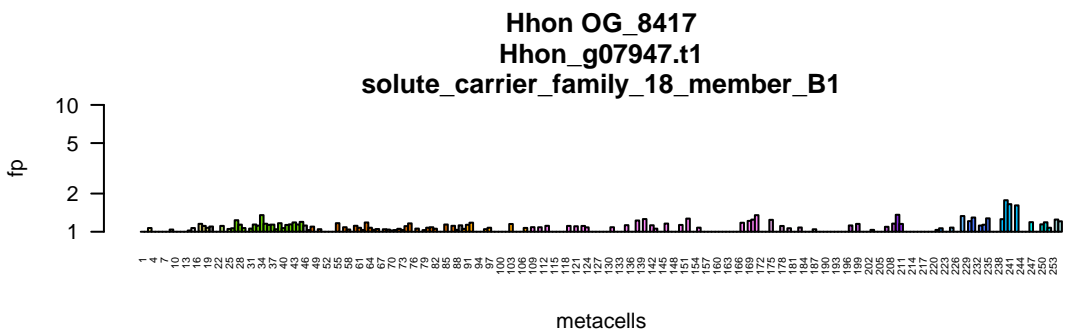
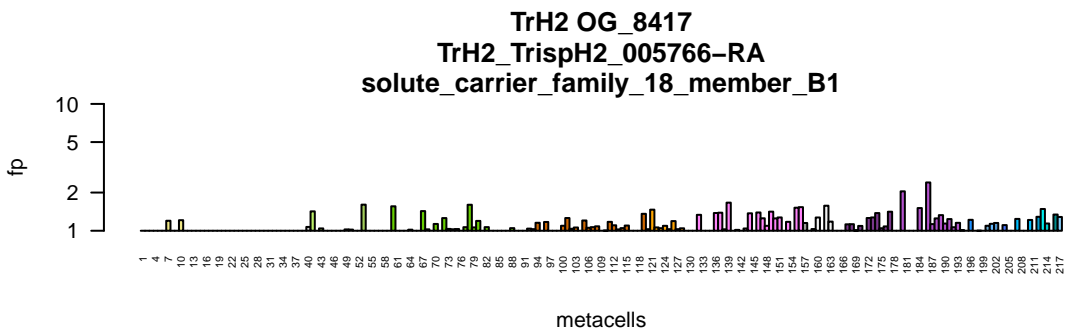
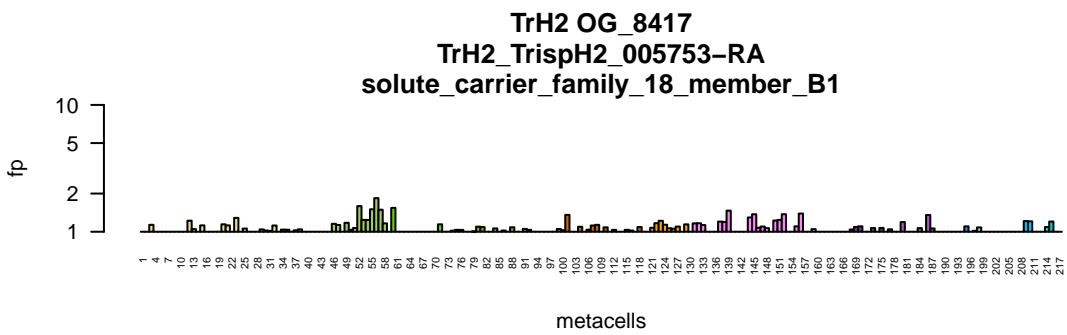
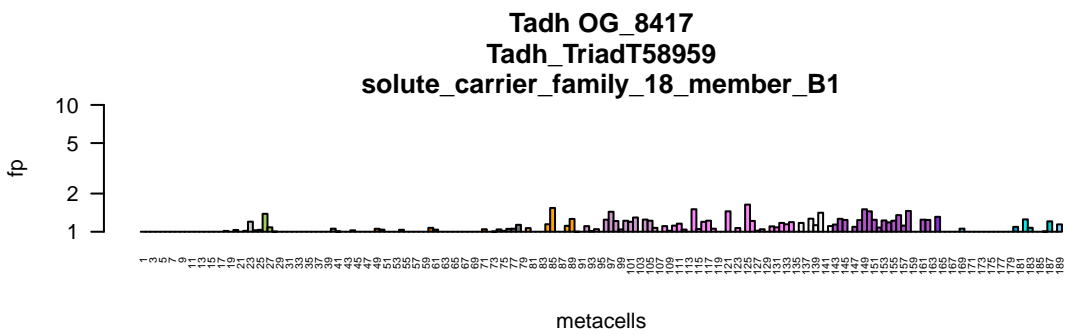
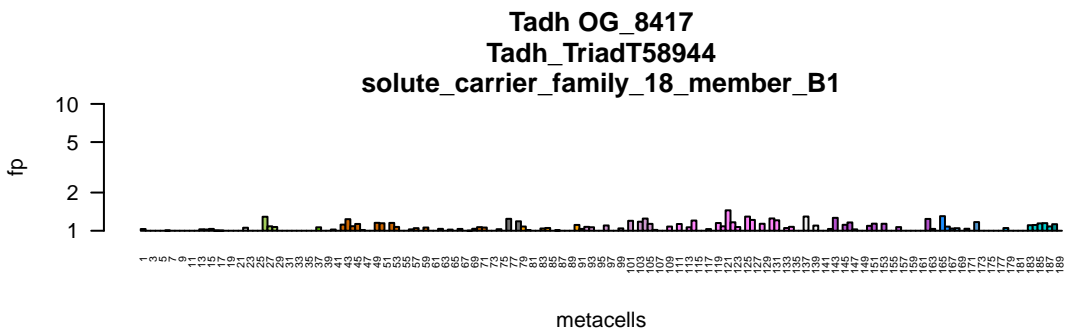
metacells

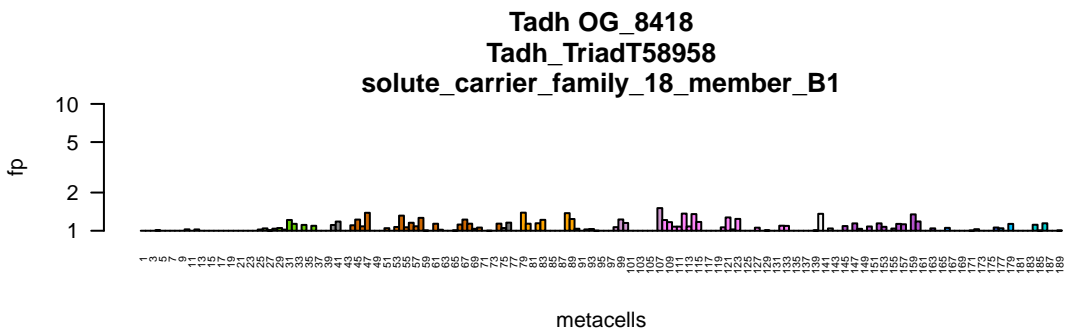
HoiH23 OG_8331
HoiH23_PIH23_002902-RA
major_facilitator_superfamily_domain_containing_11

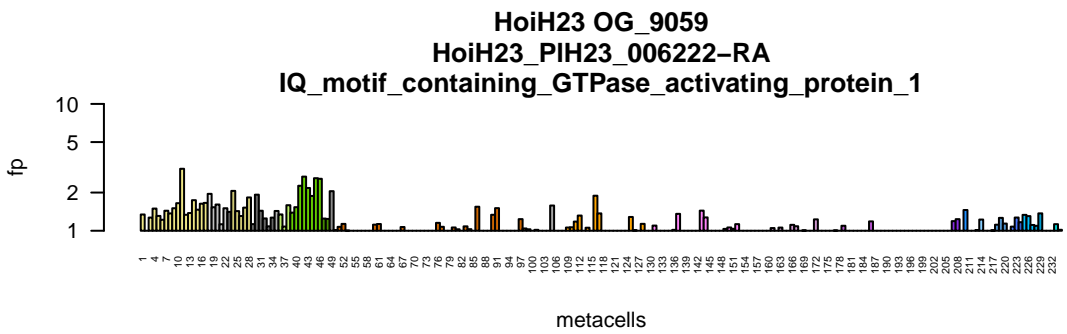
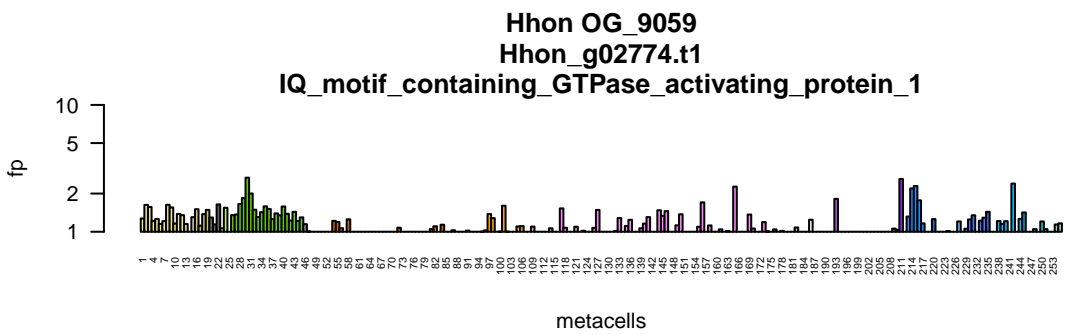
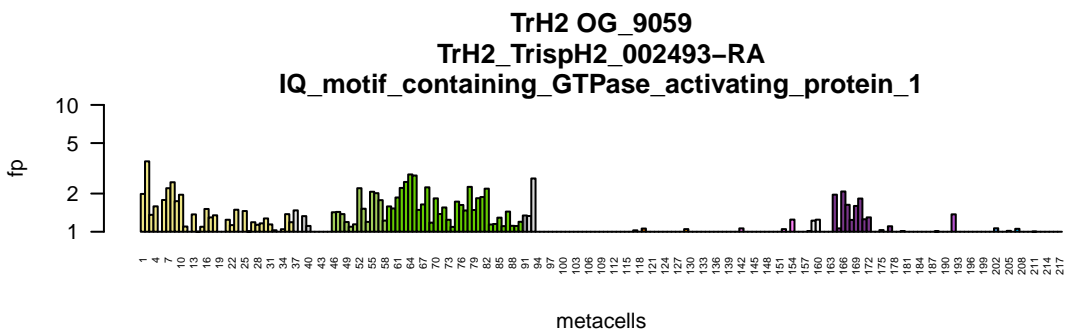
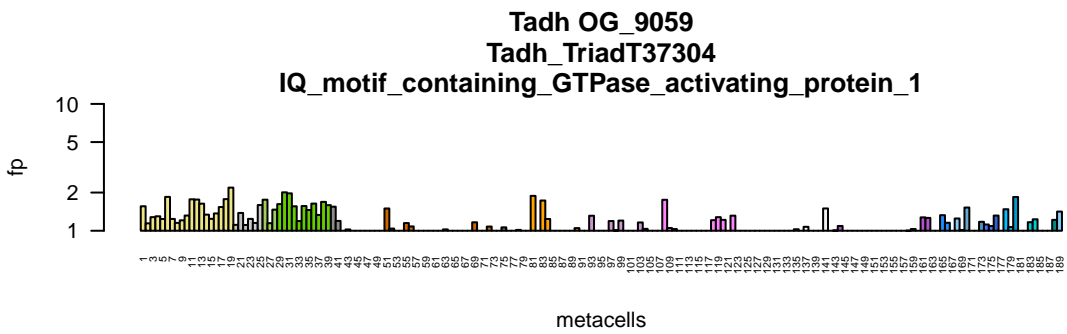


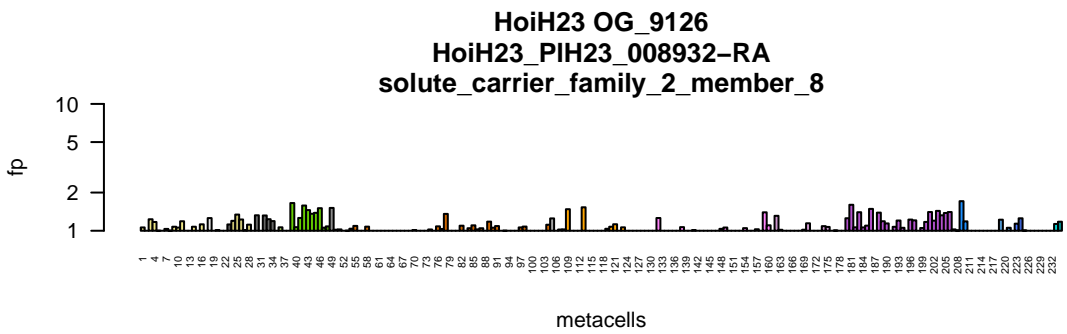
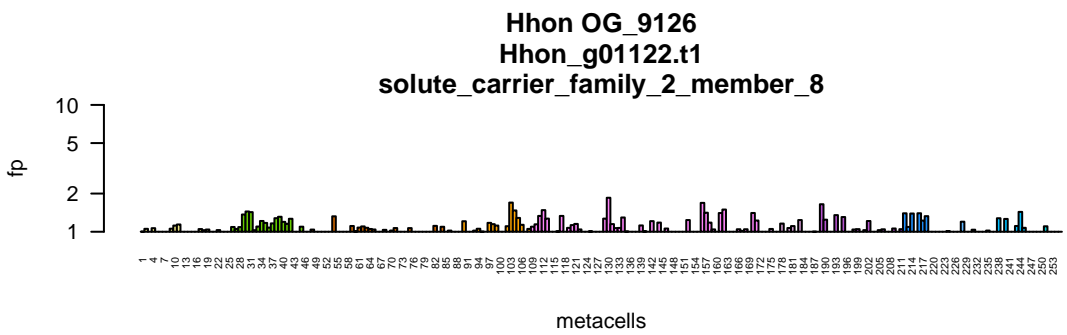
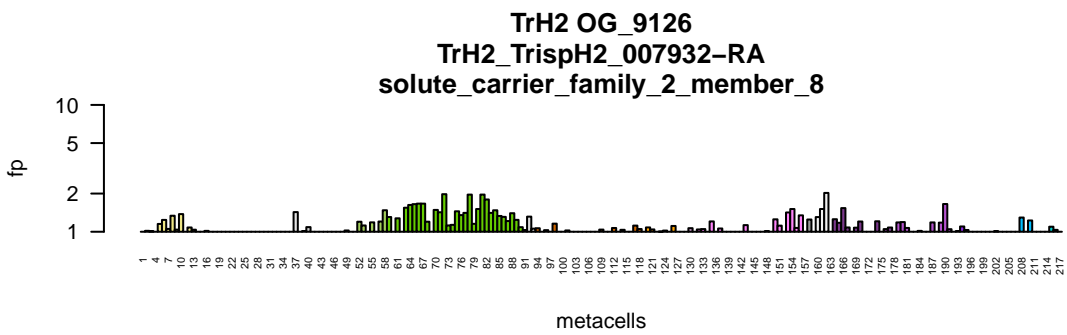
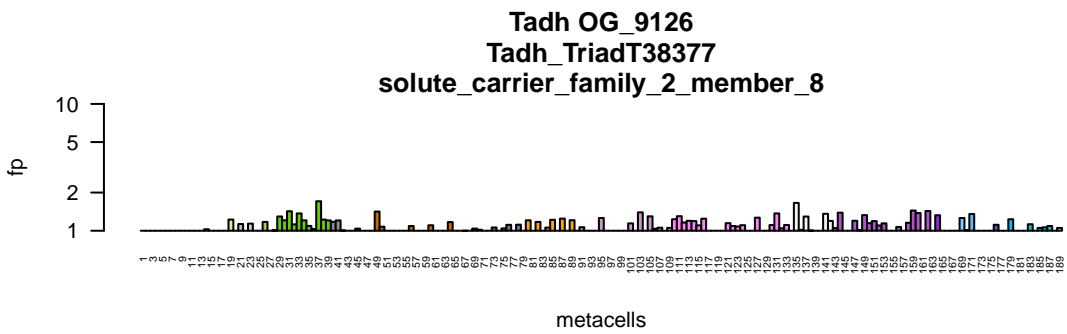
metacells

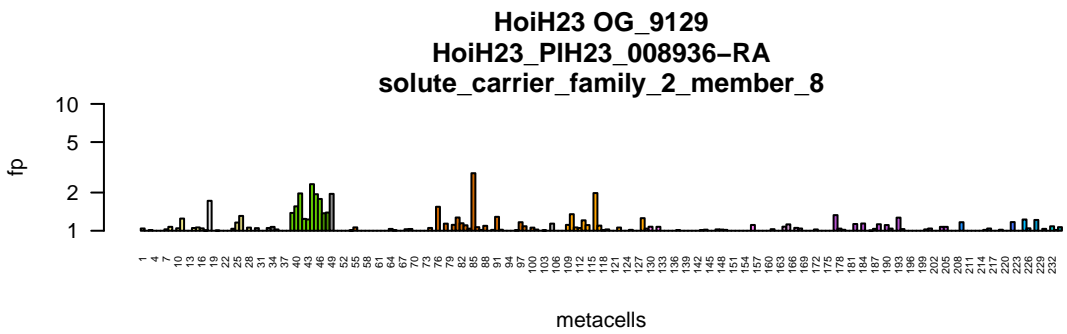
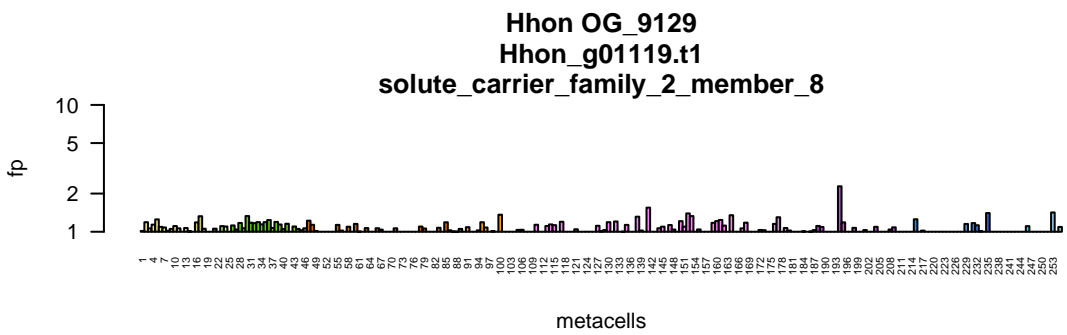
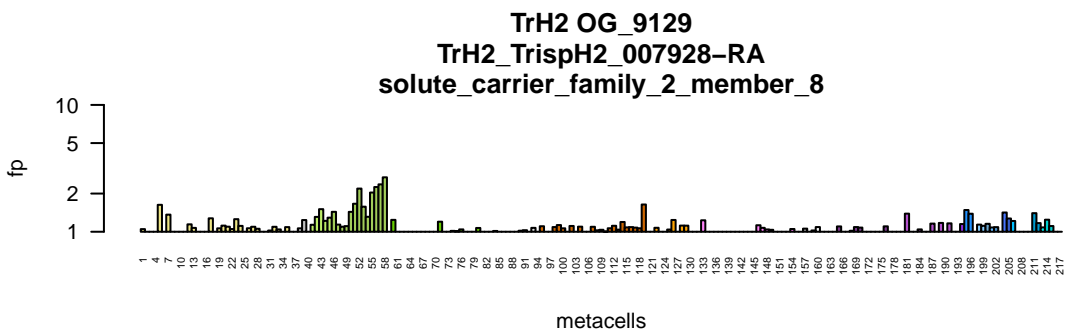
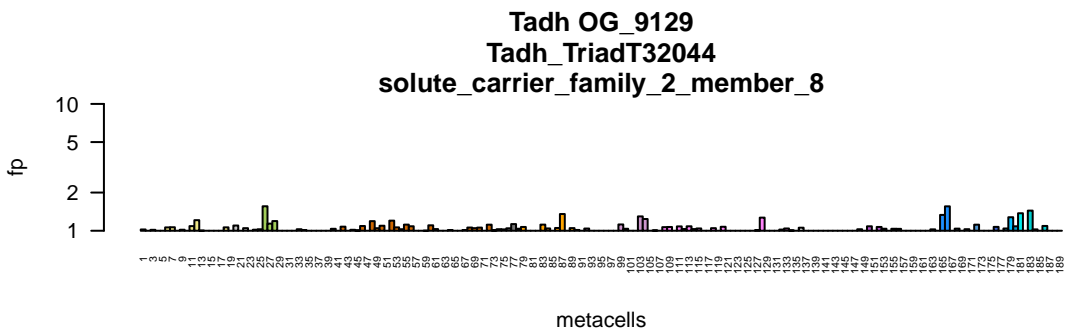


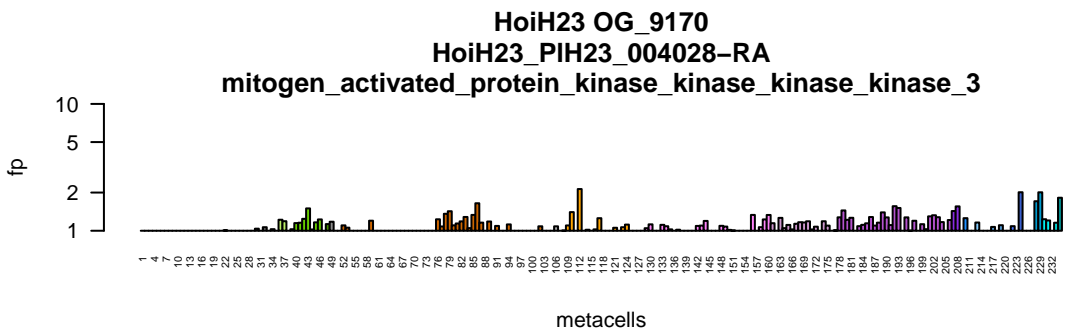
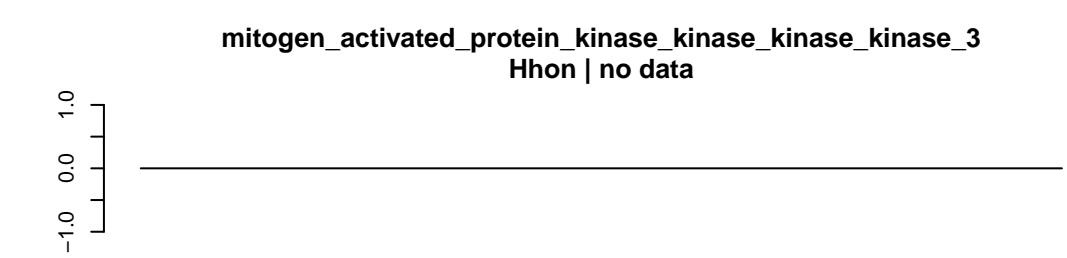
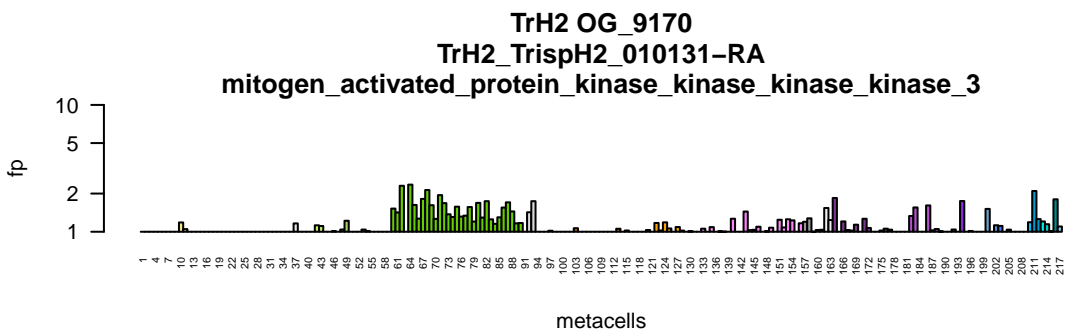
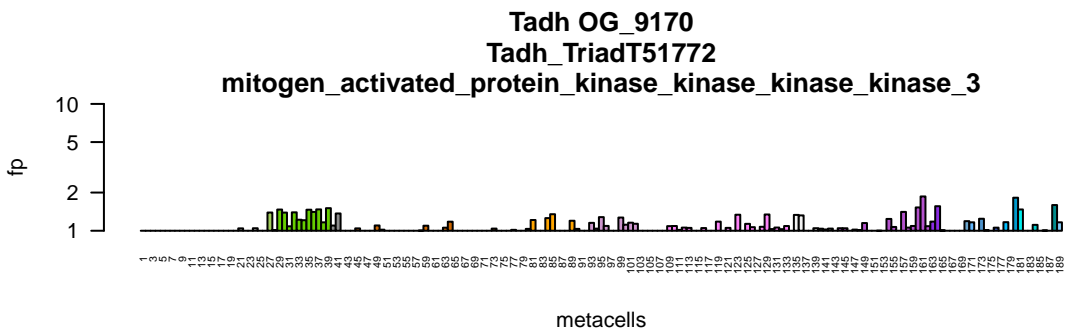


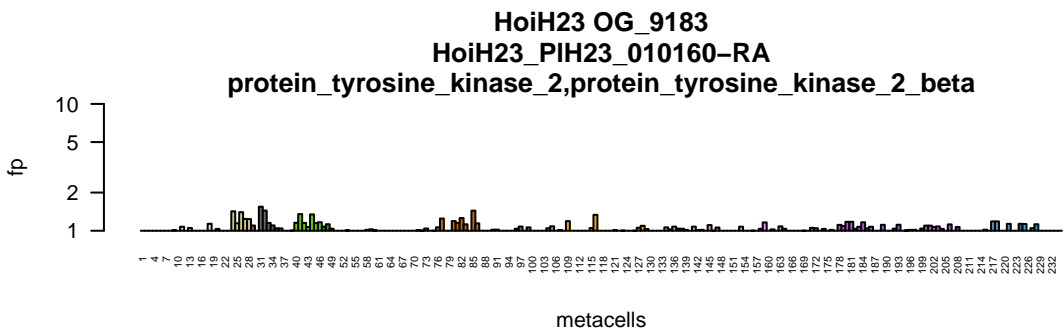
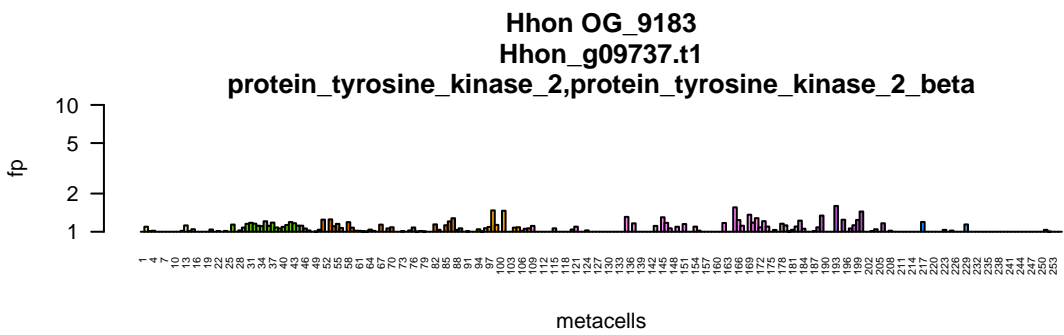
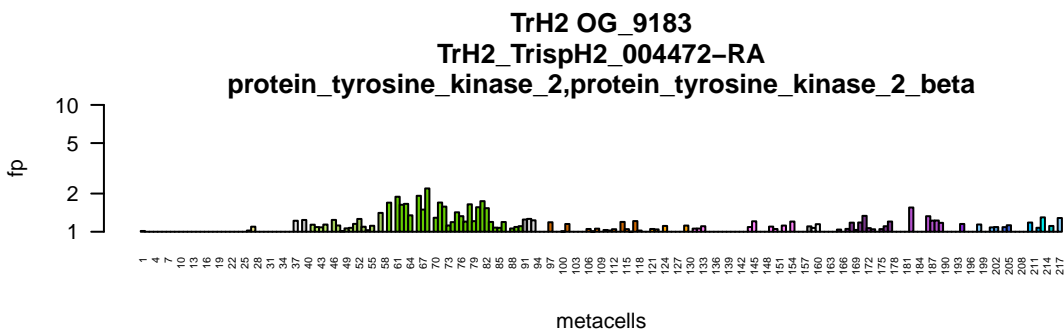
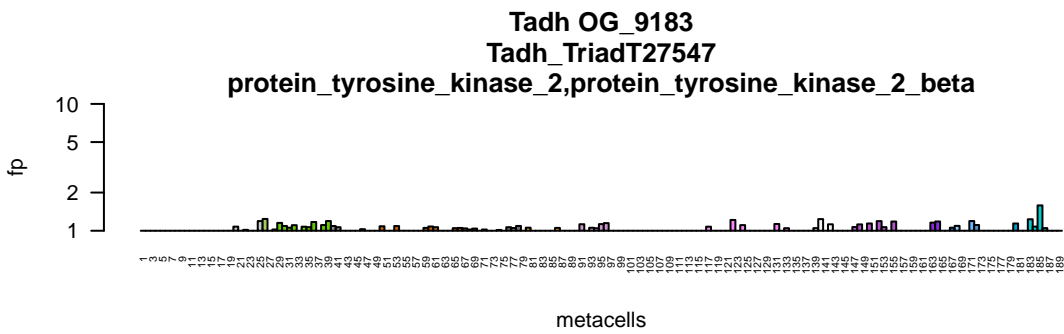


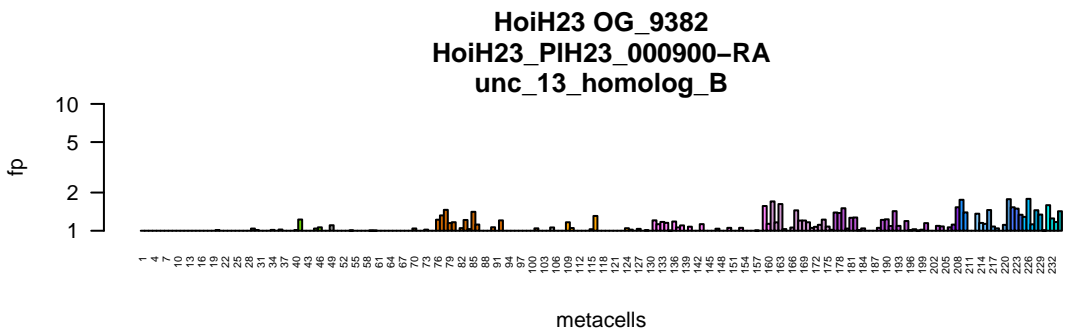
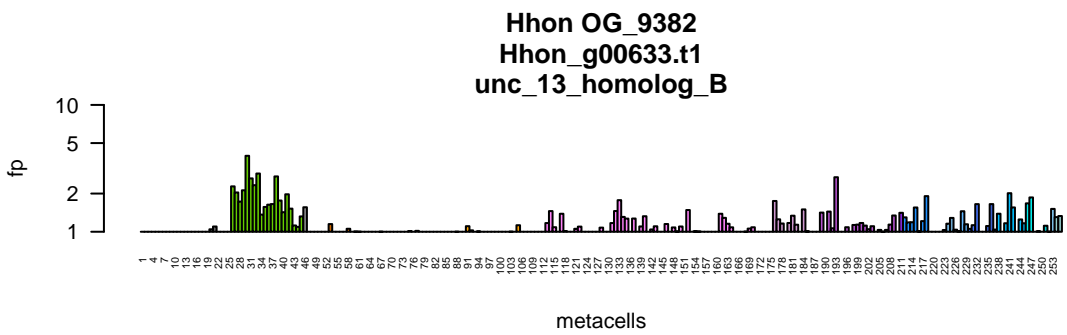
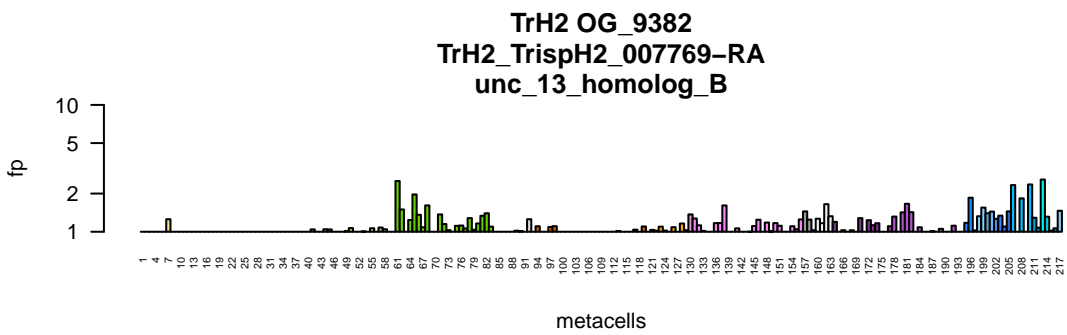
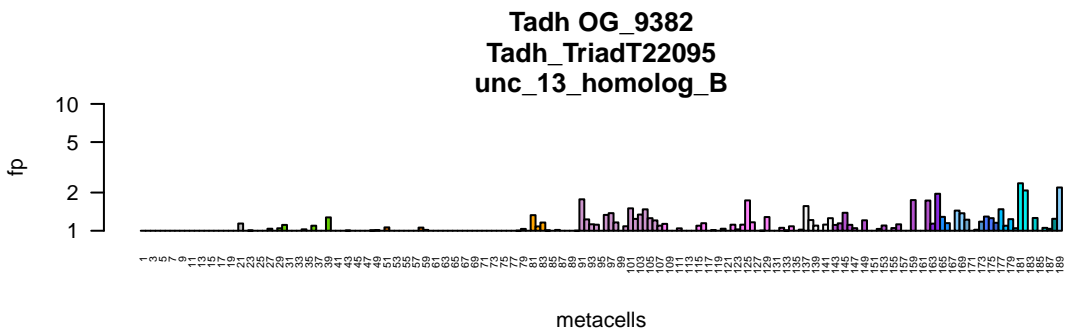


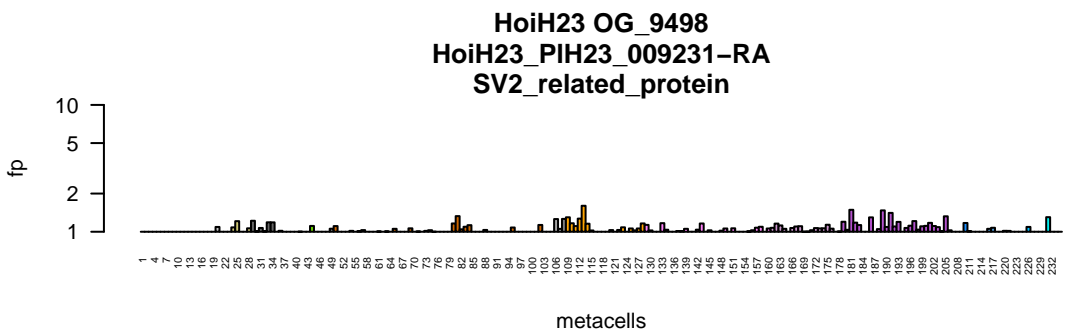
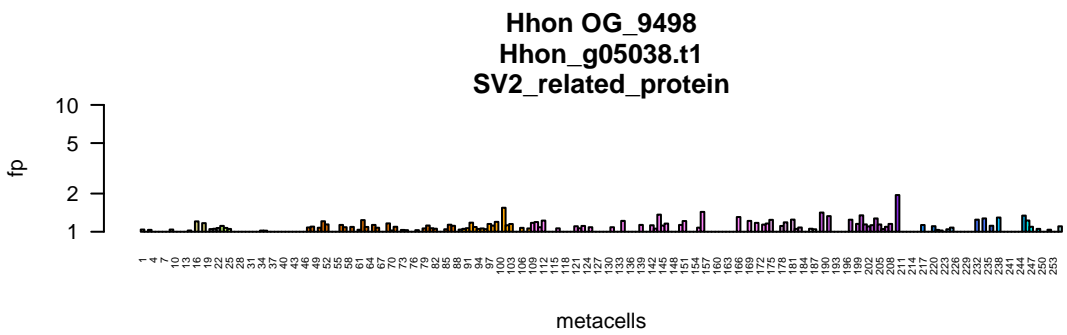
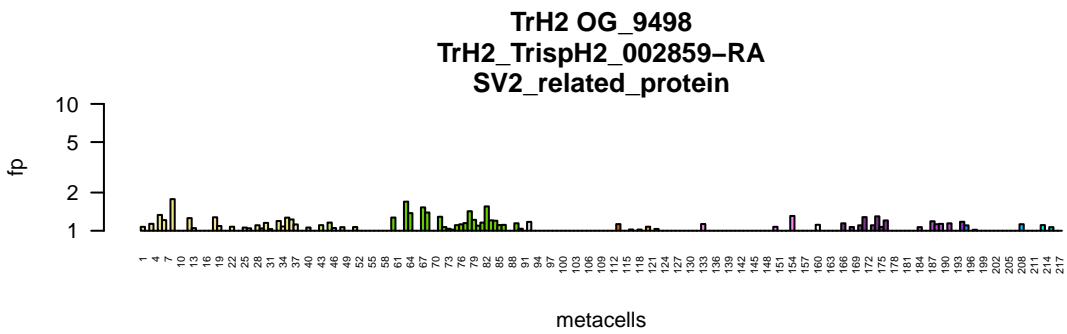
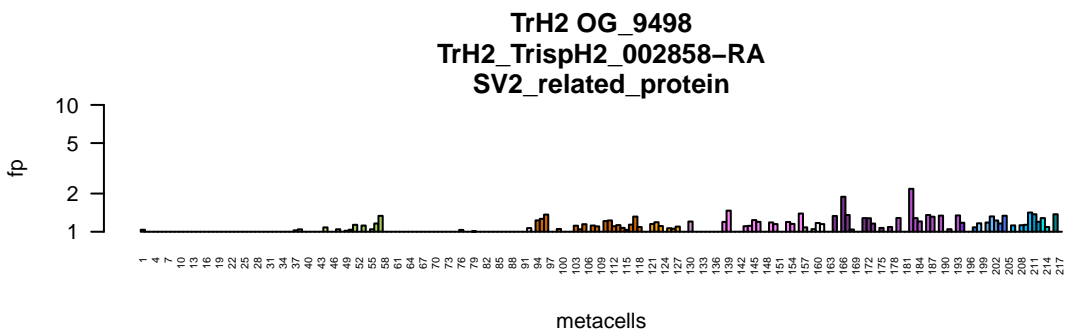
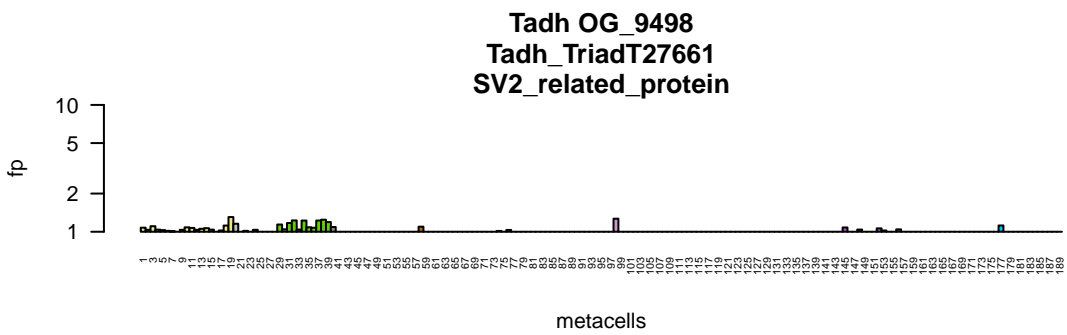
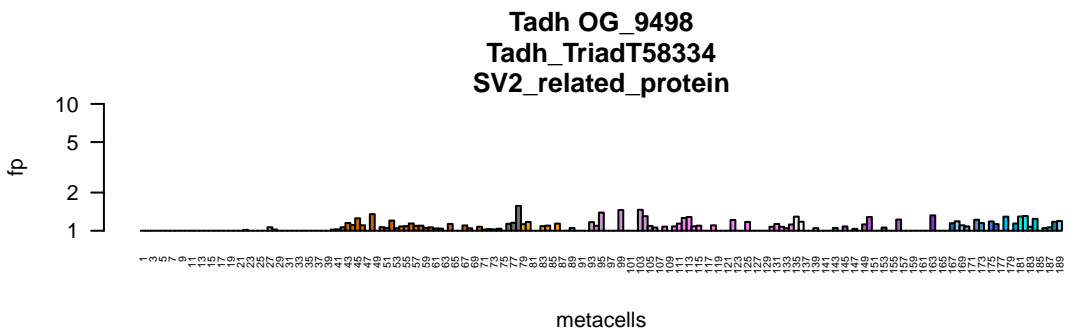


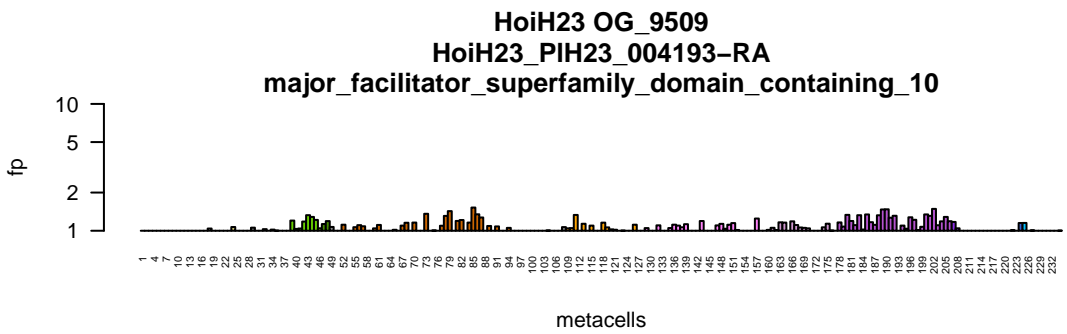
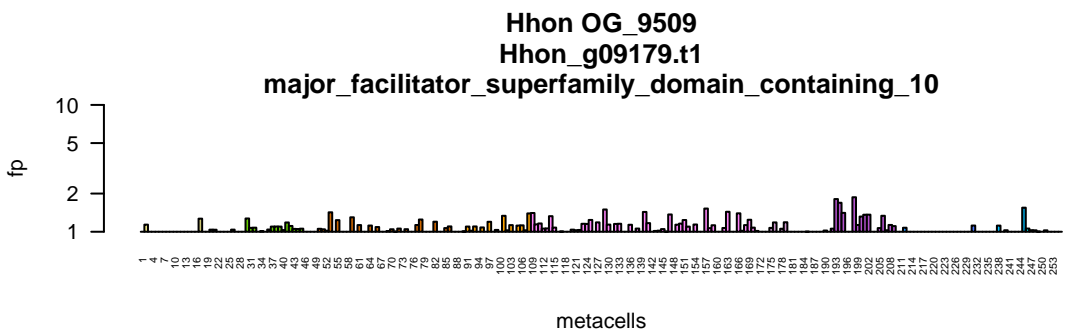
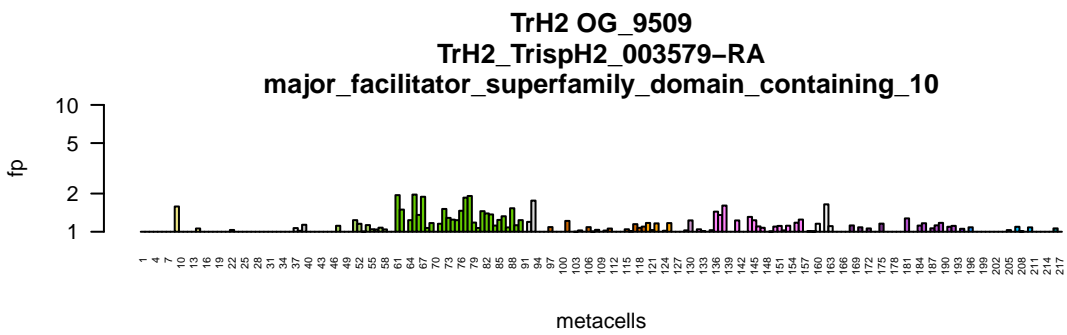
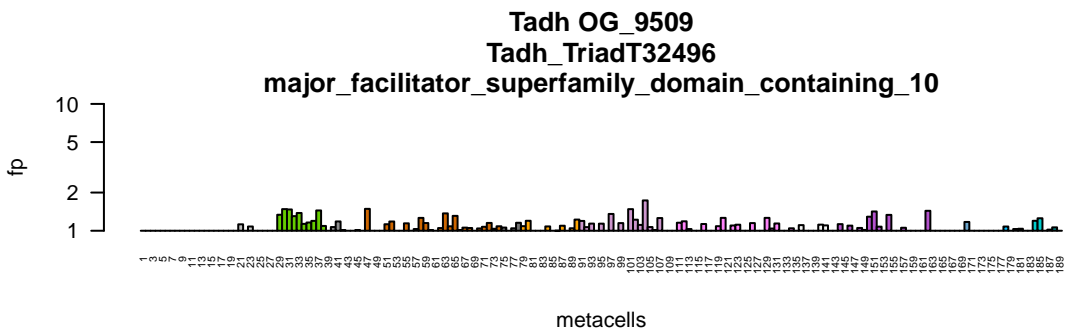


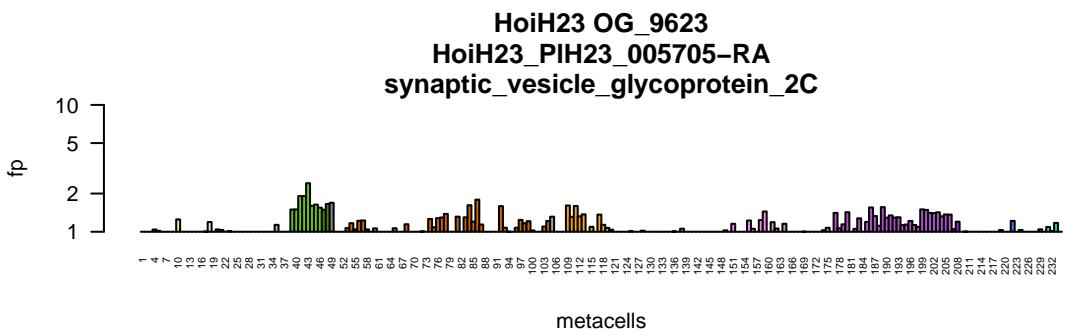
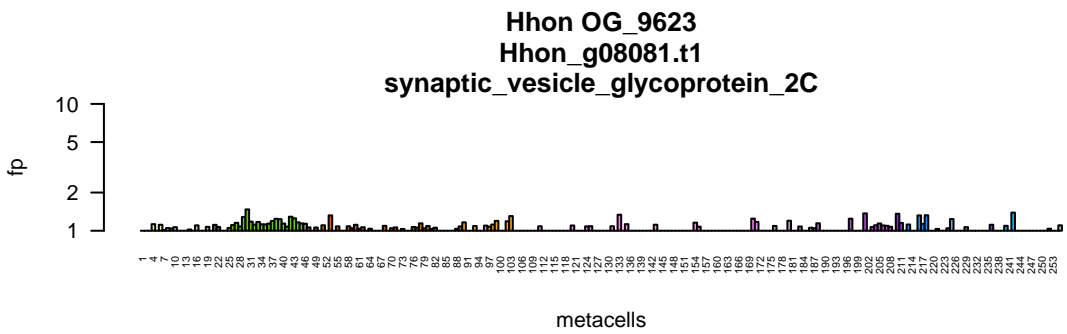
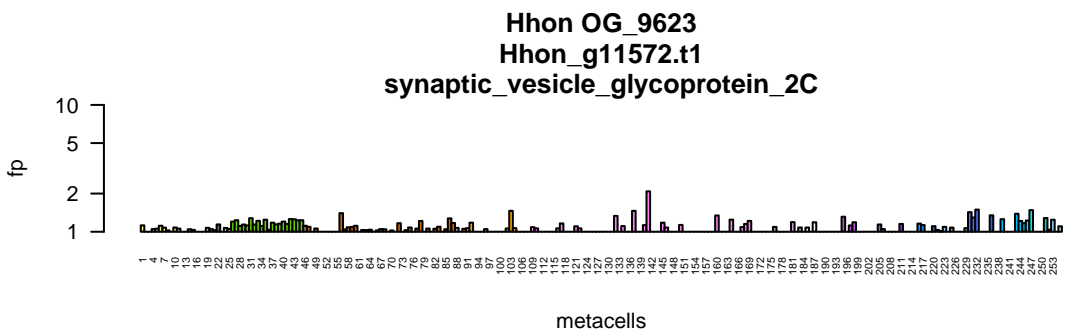
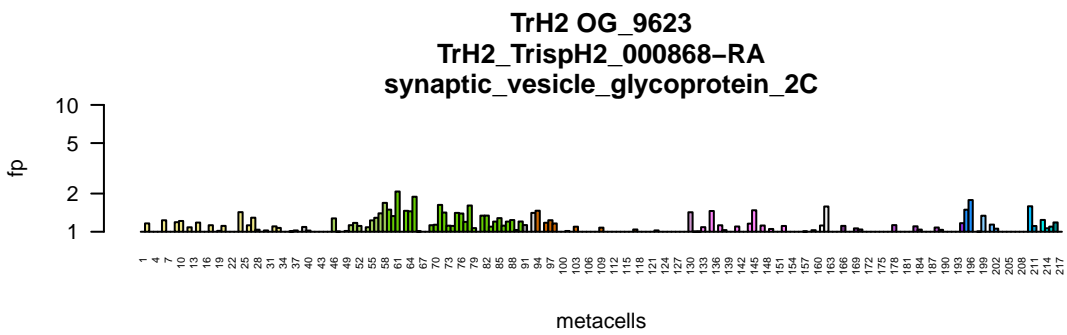
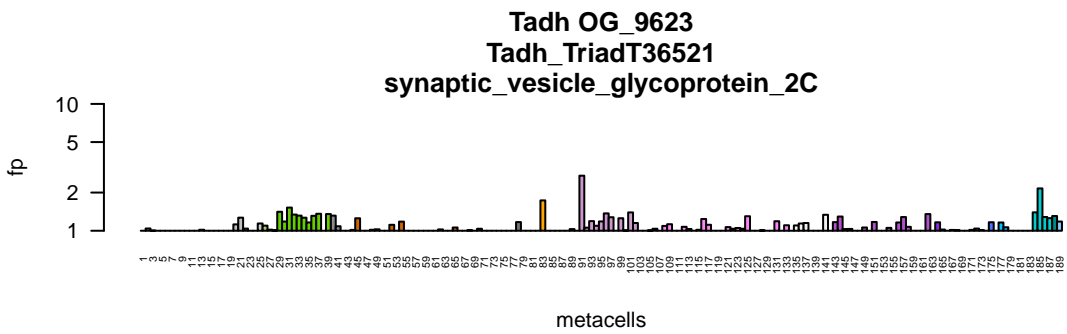








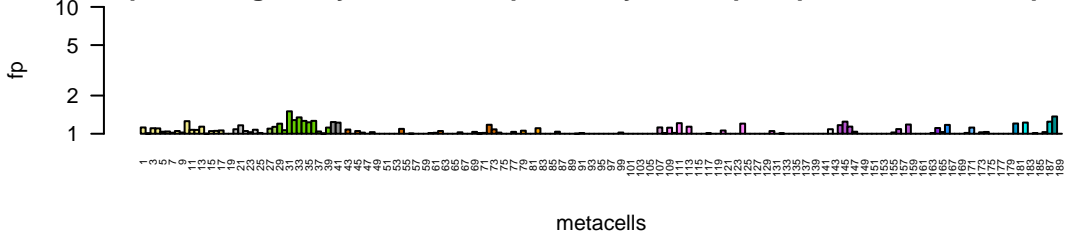




Tadh OG_9681

Tadh_TriadT18456

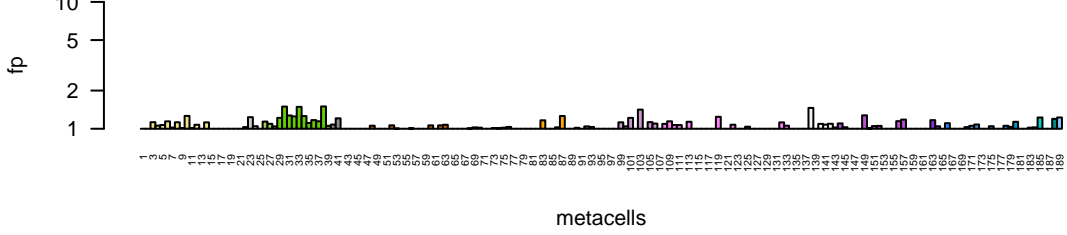
ALG8_alpha_1_3_glucosyltransferase,protein_tyrosine_phosphatase_non_receptor_type



Tadh OG_9681

Tadh_TriadT52495

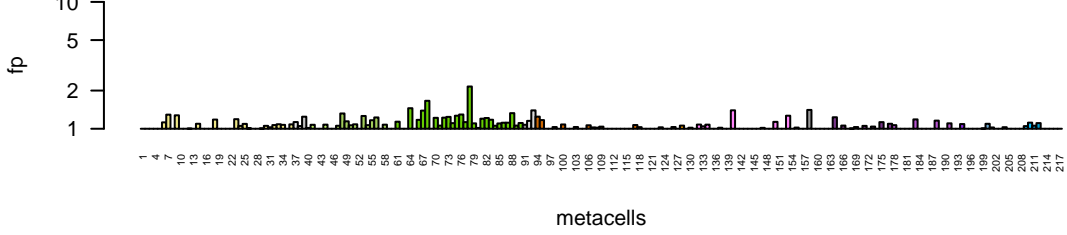
ALG8_alpha_1_3_glucosyltransferase,protein_tyrosine_phosphatase_non_receptor_type



TrH2 OG_9681

TrH2_TrispH2_000149-RA

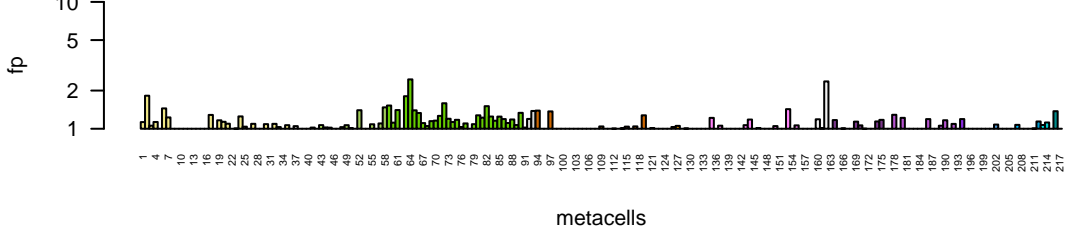
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TrH2 OG_9681

TrH2_TrispH2_000150-RA

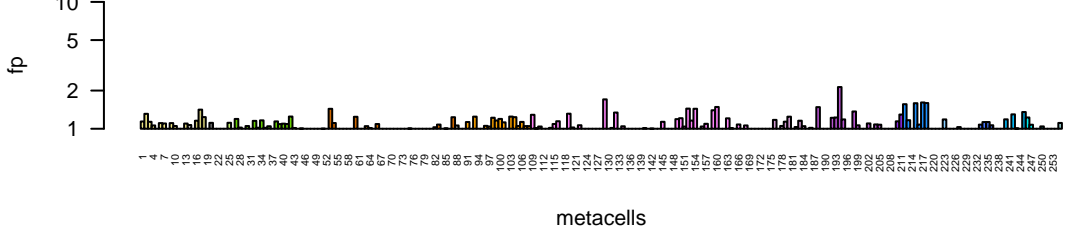
ALG8_alpha_1_3_glucosyltransferase,protein_tyrosine_phosphatase_non_receptor_type



Hhon OG_9681

Hhon_g06178.t1

ALG8_alpha_1_3_glucosyltransferase,protein_tyrosine_phosphatase_non_receptor_type



HoiH23 OG_9681

HoiH23_PIH23_001877-RA

ALG8_alpha_1_3_glucosyltransferase,protein_tyrosine_phosphatase_non_receptor_type

