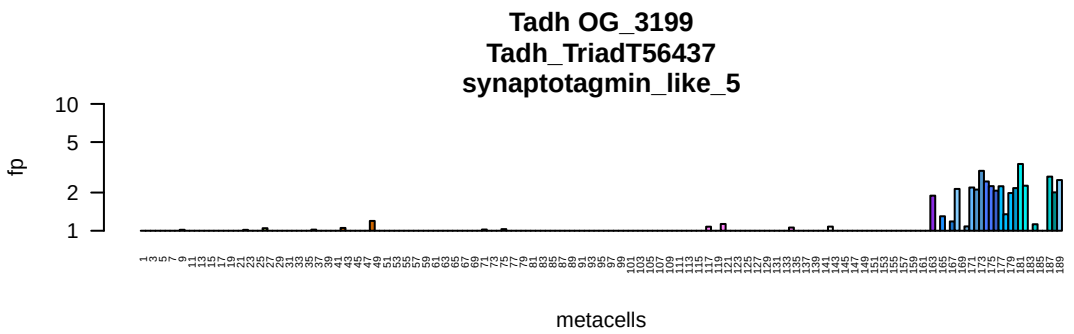
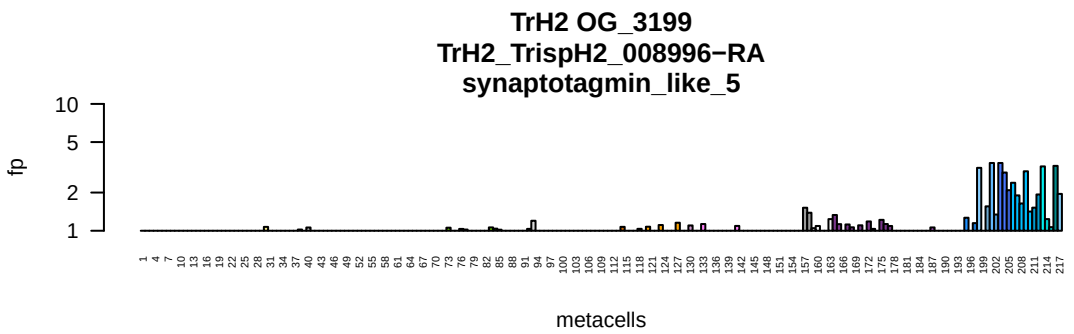
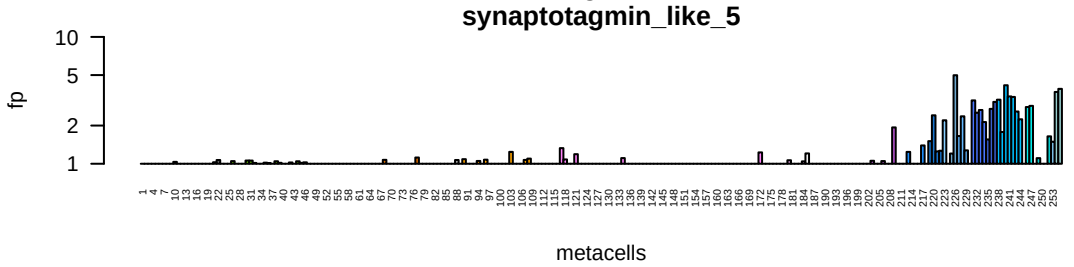

Tadh OG_3199
Tadh_TriadT56437
synaptotagmin_like_5
fp
metacells


TrH2 OG_3199
TrH2_TrispH2_008996−RA
synaptotagmin_like_5
fp
metacells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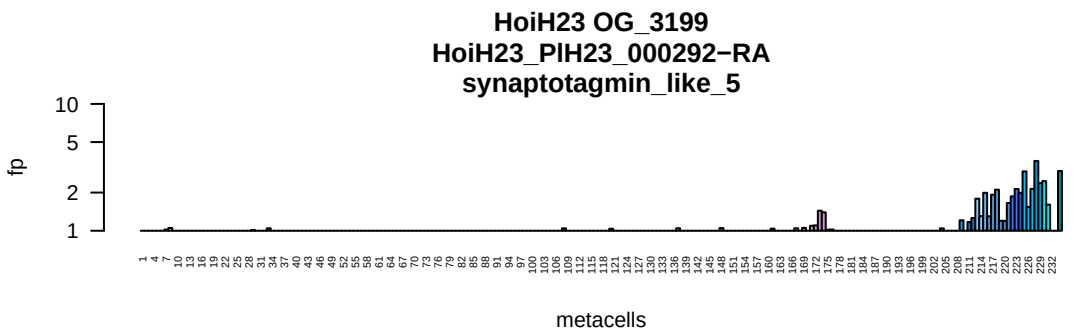
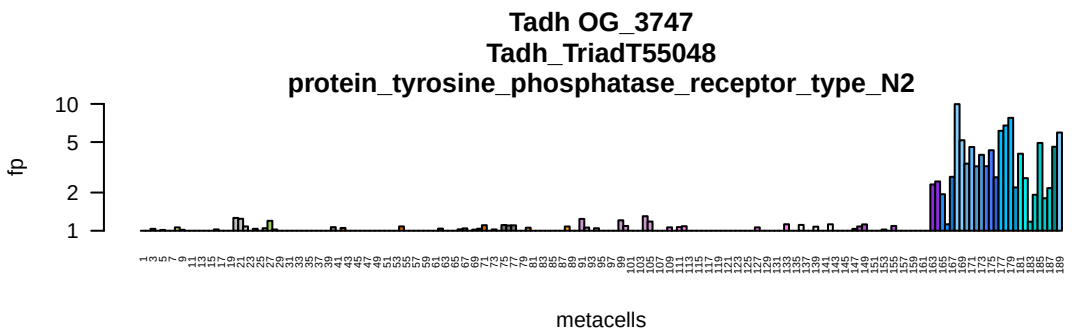
synaptotagmin_like_5
fp
metacells


HoiH23 OG_3199
HoiH23_PlH23_000292−RA
synaptotagmin_like_5
fp
metacells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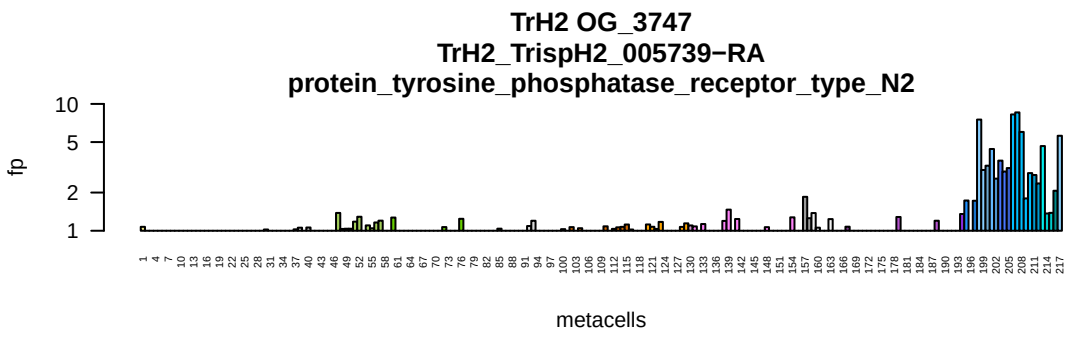
Tadh_TriadT55048
protein_tyrosine_phosphatase_receptor_type_N2
fp
metacells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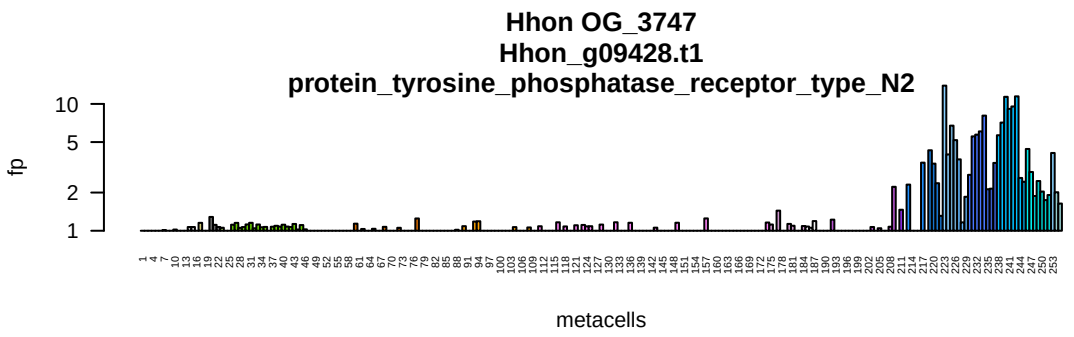
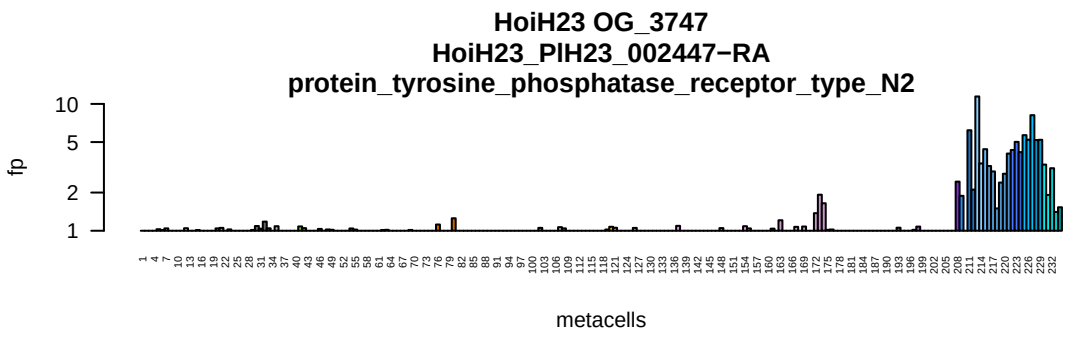
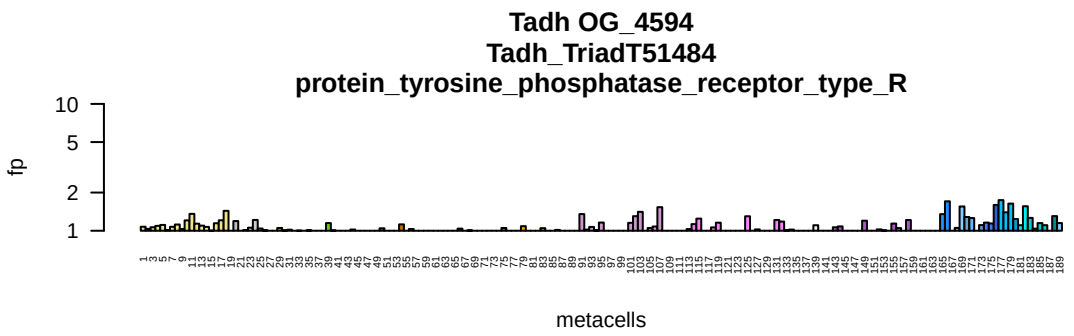
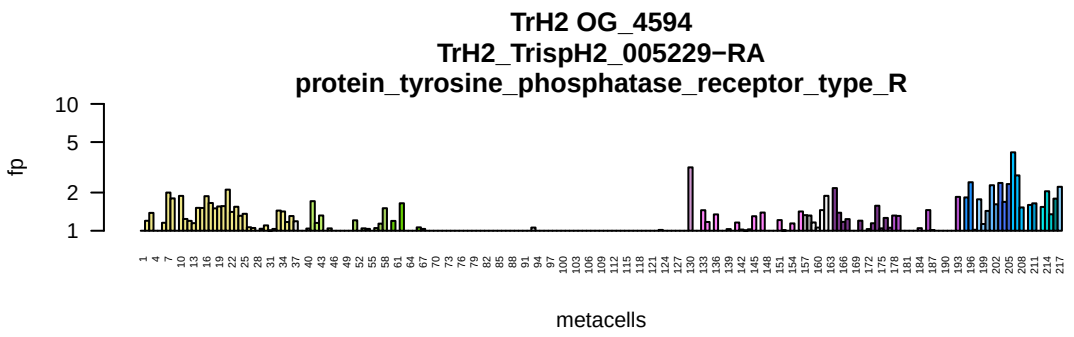
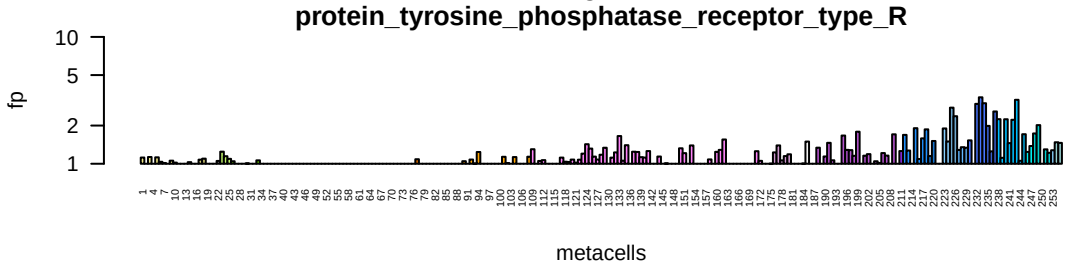
TrH2_TrispH2_005739−RA
protein_tyrosine_phosphatase_receptor_type_N2
fp
metacells

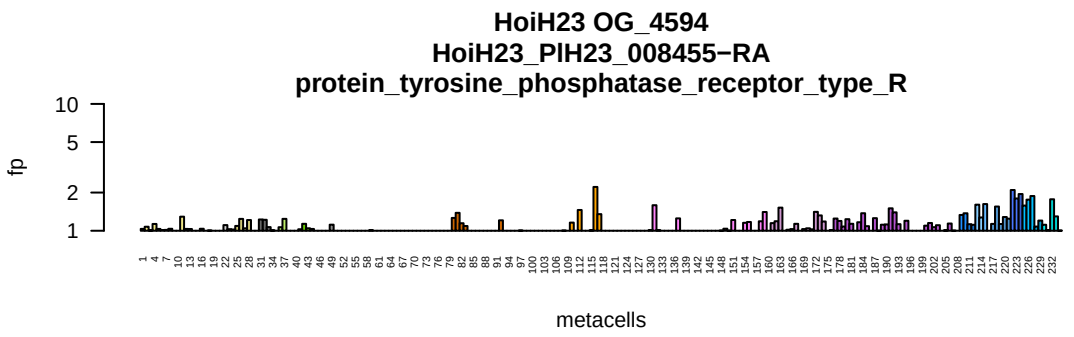

Hhon OG_3747
Hhon_g09428.t1
protein_tyrosine_phosphatase_receptor_type_N2
fp
metacells

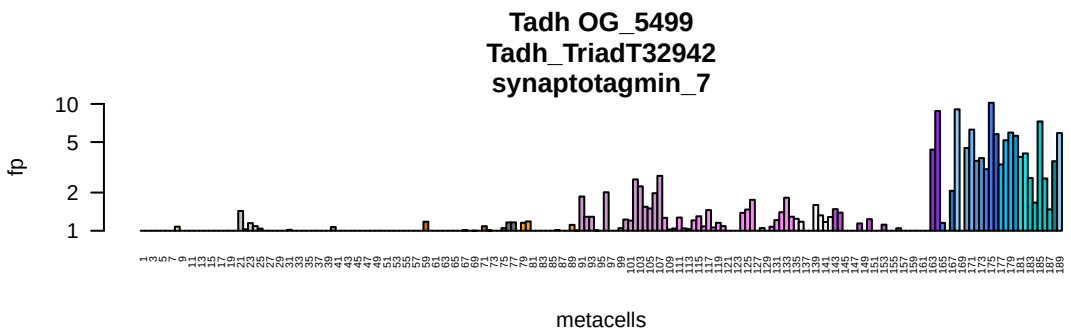

HoiH23 OG_3747
HoiH23_PlH23_002447−RA
protein_tyrosine_phosphatase_receptor_type_N2
fp
metacells

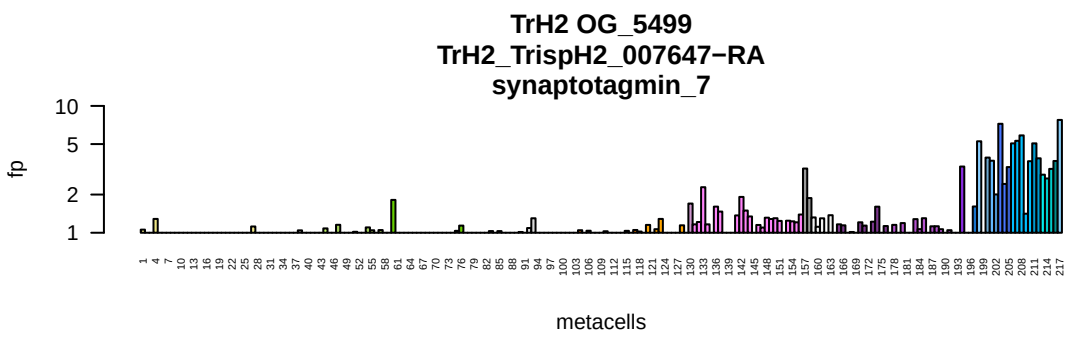

Tadh OG_4594
Tadh_TriadT51484
protein_tyrosine_phosphatase_receptor_type_R
fp
metacells

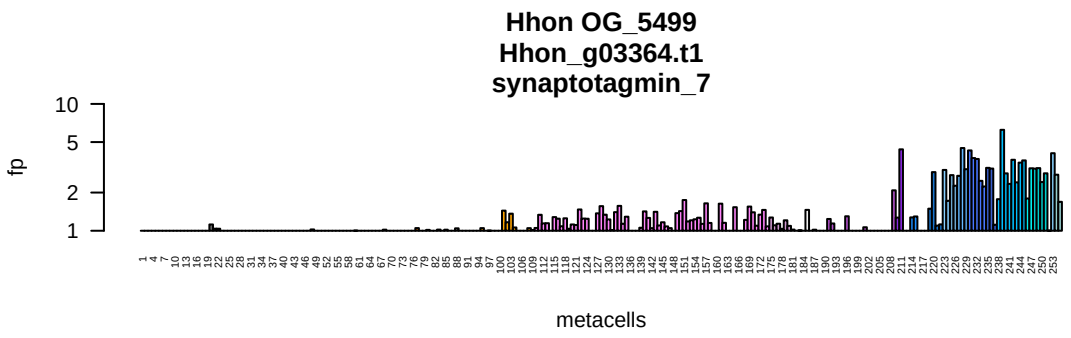

TrH2 OG_4594
TrH2_TrispH2_005229−RA
protein_tyrosine_phosphatase_receptor_type_R
fp
metacells

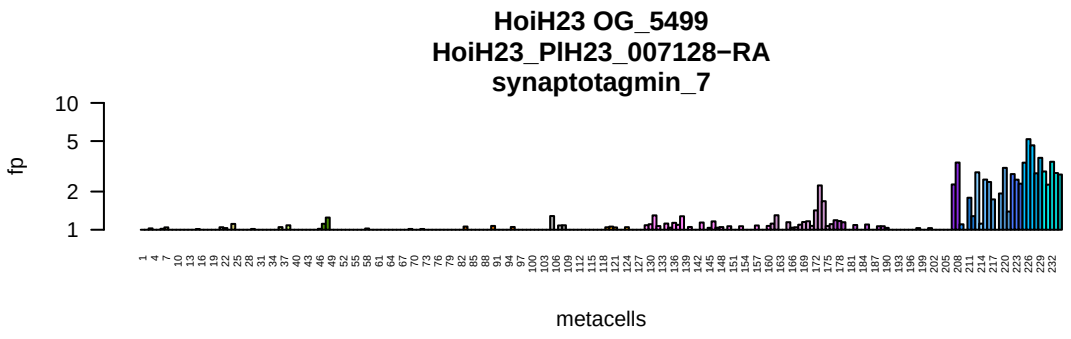

Hhon OG_4594
Hhon_g08135.t1
protein_tyrosine_phosphatase_receptor_type_R
fp
metacells


HoiH23 OG_4594
HoiH23_PlH23_008455−RA
protein_tyrosine_phosphatase_receptor_type_R
fp
metacells


Tadh OG_5499
Tadh_TriadT32942
synaptotagmin_7
fp
metacells


TrH2 OG_5499
TrH2_TrispH2_007647−RA
synaptotagmin_7
fp
metacells


Hhon OG_5499
Hhon_g03364.t1
synaptotagmin_7
fp
metacells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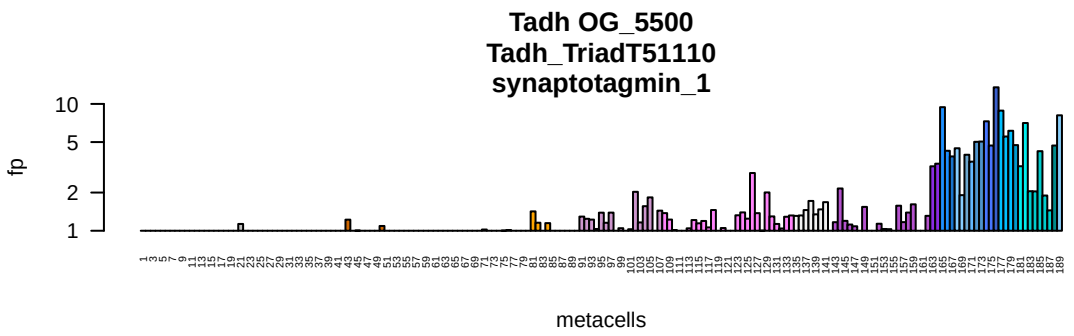
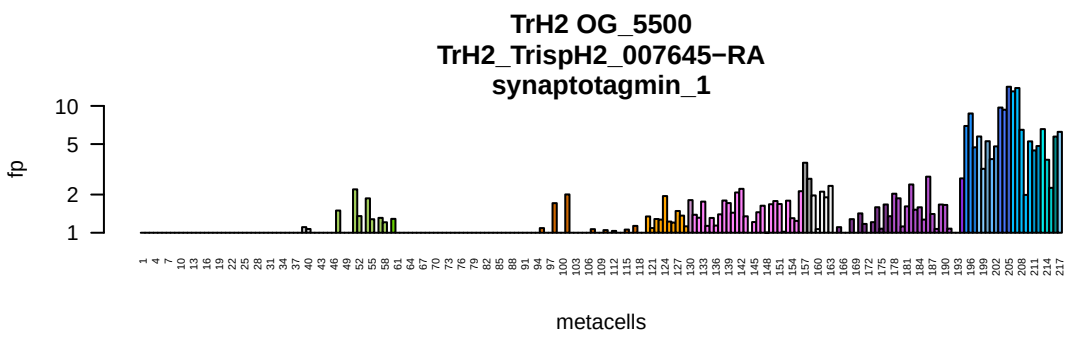
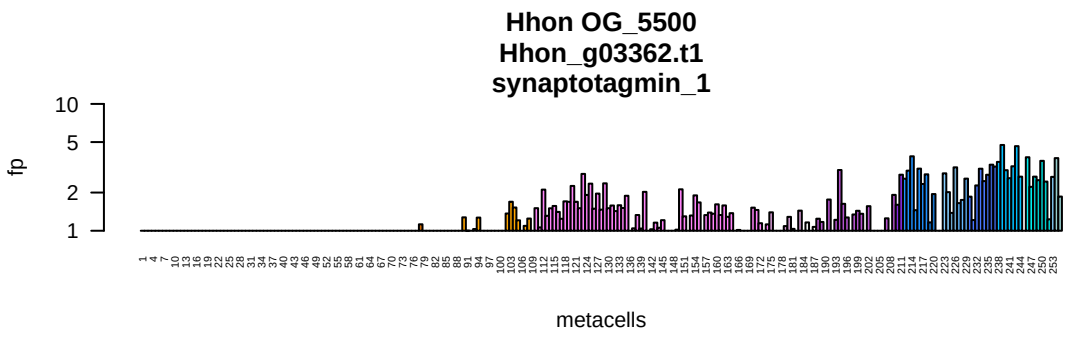
HoiH23_PlH23_007128−RA
synaptotagmin_7
fp
metacells


Tadh OG_5500
Tadh_TriadT51110
synaptotagmin_1
fp
metacells


TrH2 OG_5500
TrH2_TrispH2_007645−RA
synaptotagmin_1
fp
metacells


Hhon OG_5500
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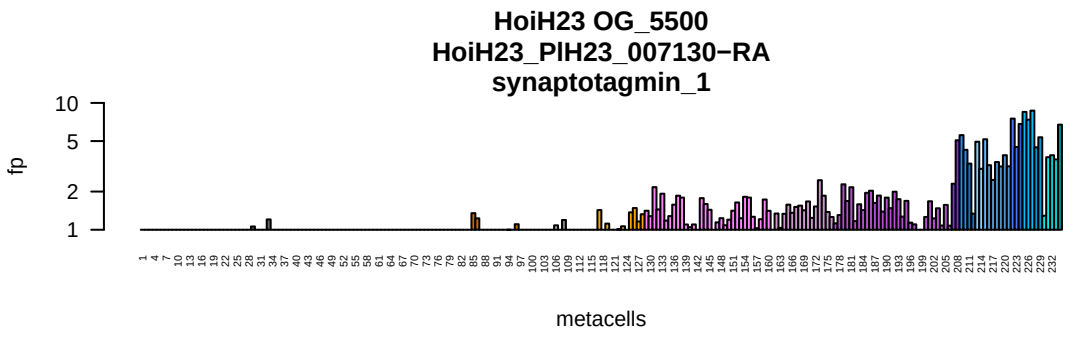
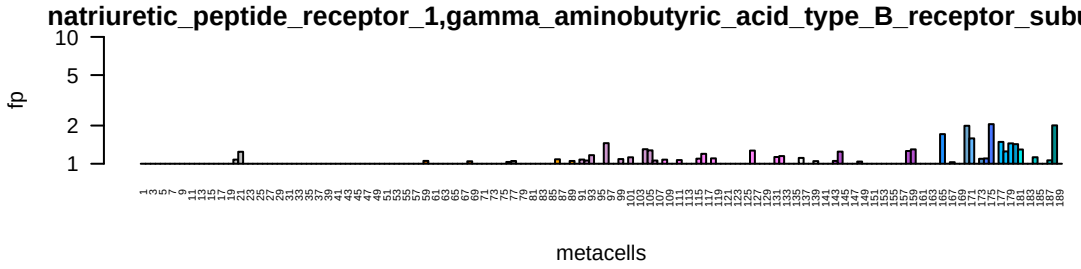
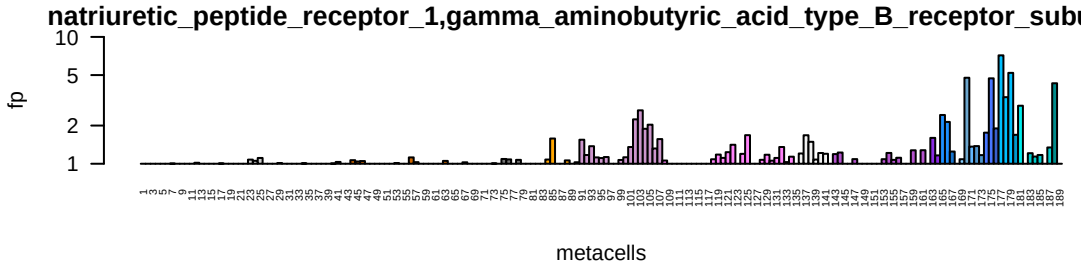
synaptotagmin_1
fp
metacells


HoiH23 OG_5500
HoiH23_PlH23_007130−RA
synaptotagmin_1
fp
metacells

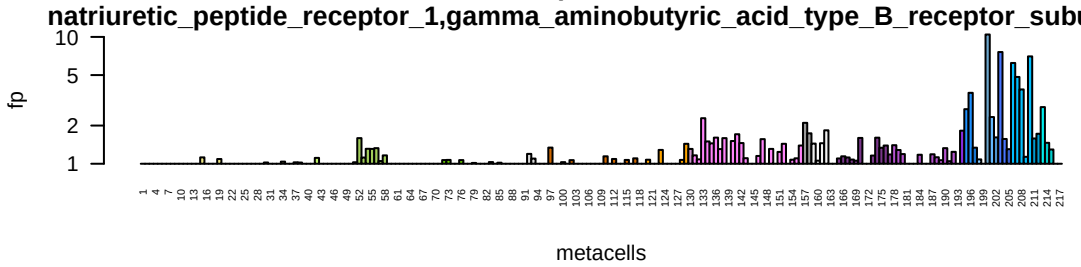
Tadh OG_5687
Tadh_TriadT59259
natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1



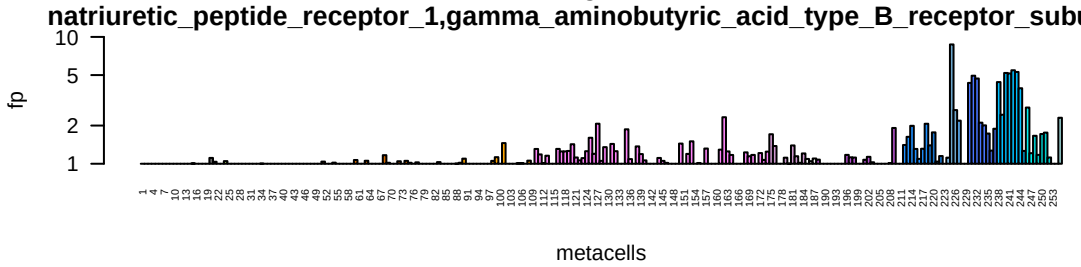
Tadh OG_5687
Tadh_TriadT29353
natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1



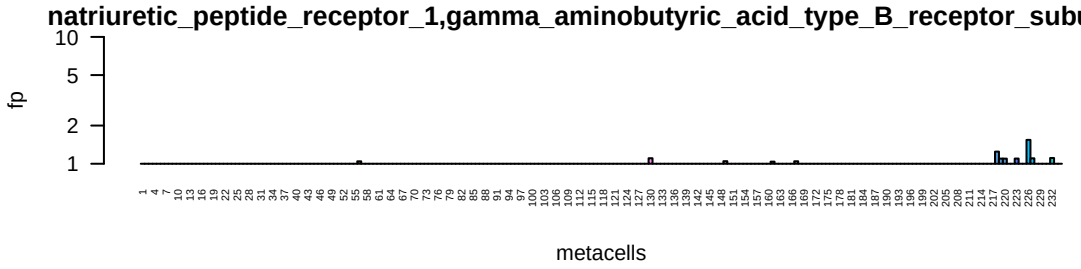
TrH2 OG_5687
TrH2_TrispH2_002051−RA
natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1



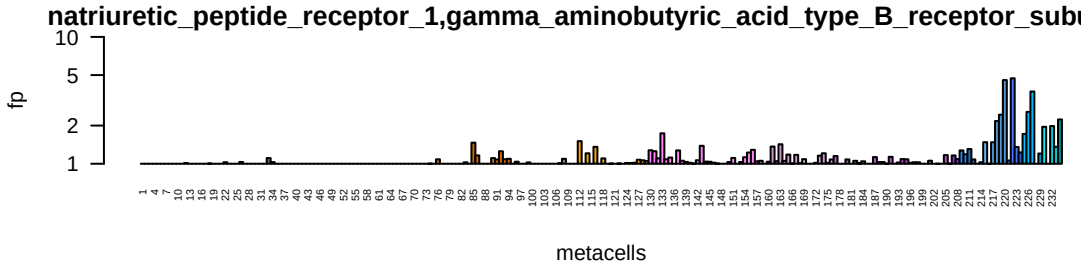
Hhon OG_5687
Hhon_g00715.t1
natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1

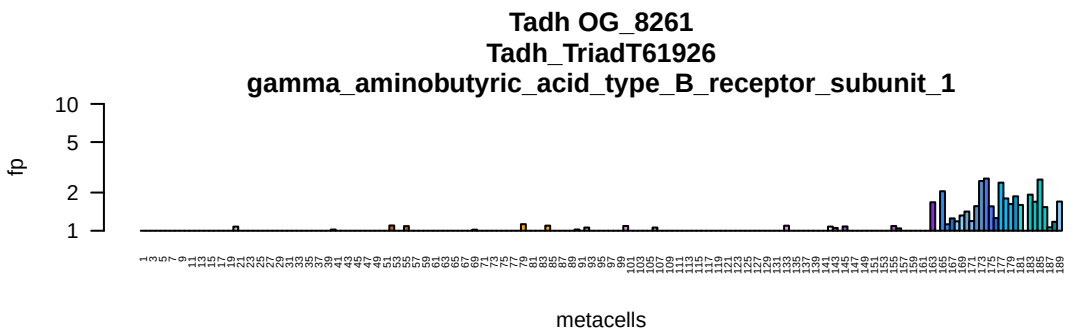


HoiH23 OG_5687
HoiH23_PlH23_009782−RA
natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1



HoiH23 OG_5687
HoiH23_PlH23_009783−RA
natriuretic_peptide_receptor_1,gamma_aminobutyric_acid_type_B_receptor_subunit_1

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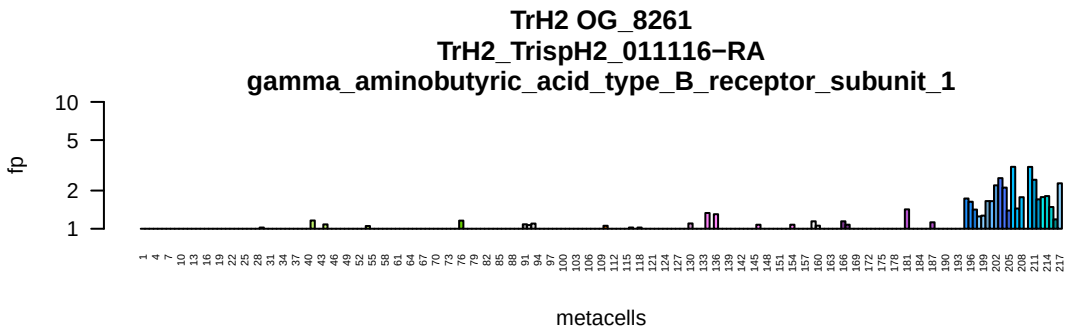
Tadh_TriadT61926
gamma_aminobutyric_acid_type_B_receptor_subunit_1
fp
10
5
2
1
metacells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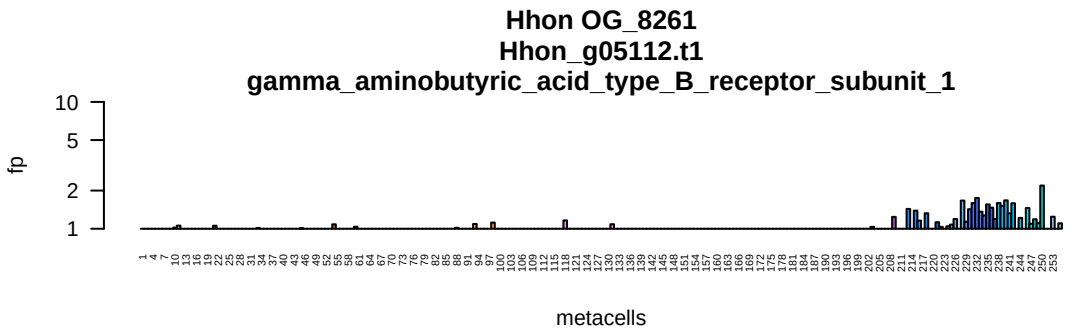
TrH2_TrispH2_011116−RA
gamma_aminobutyric_acid_type_B_receptor_subunit_1
fp
10
5
2
1
metacells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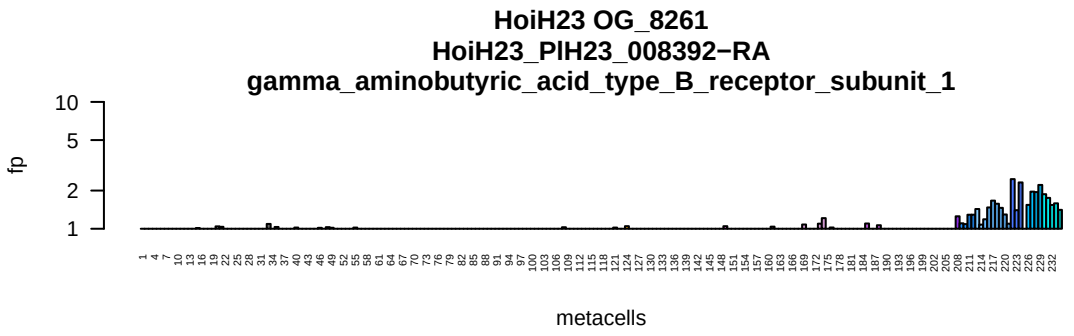
Hhon_g05112.t1
gamma_aminobutyric_acid_type_B_receptor_subunit_1
fp
10
5
2
1
metacells


HoiH23 OG_8261
HoiH23_PlH23_008392−RA
gamma_aminobutyric_acid_type_B_receptor_subunit_1
fp
10
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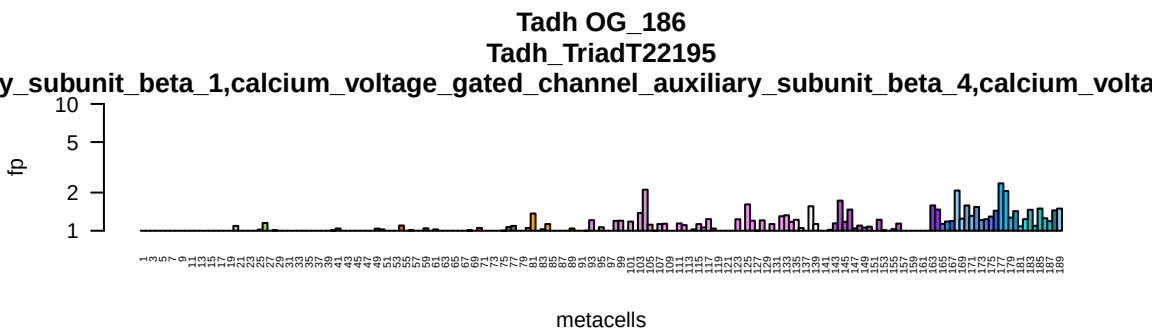
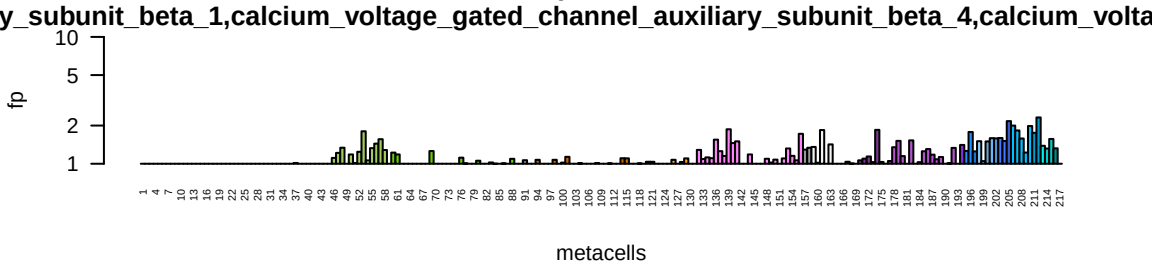
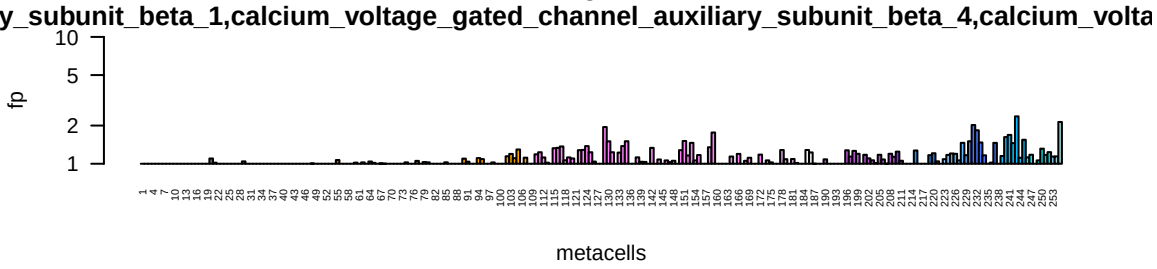
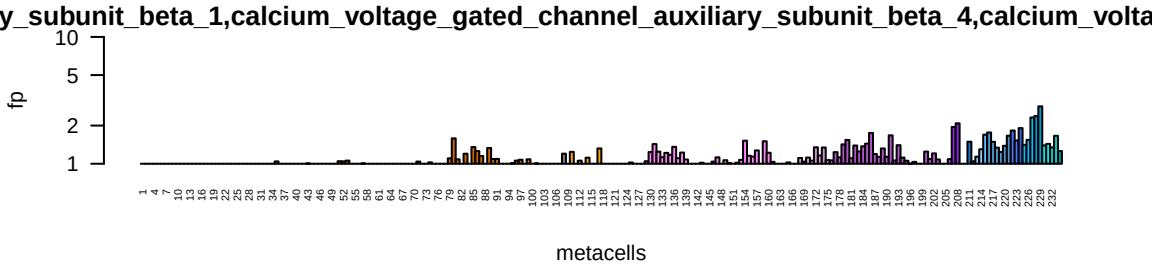
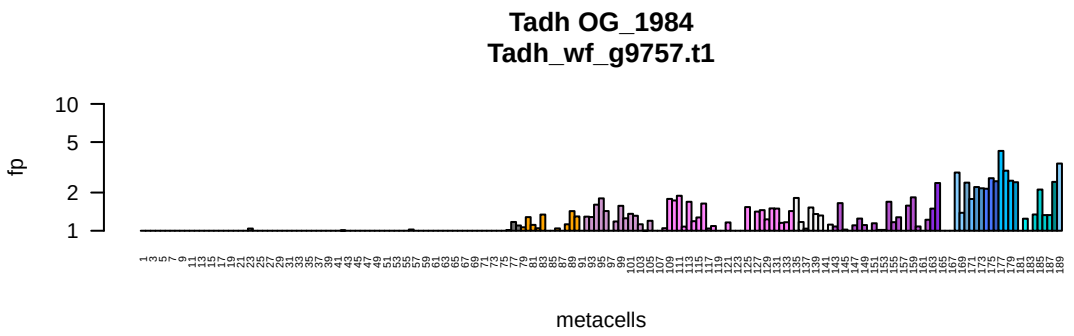
2
1
metacells

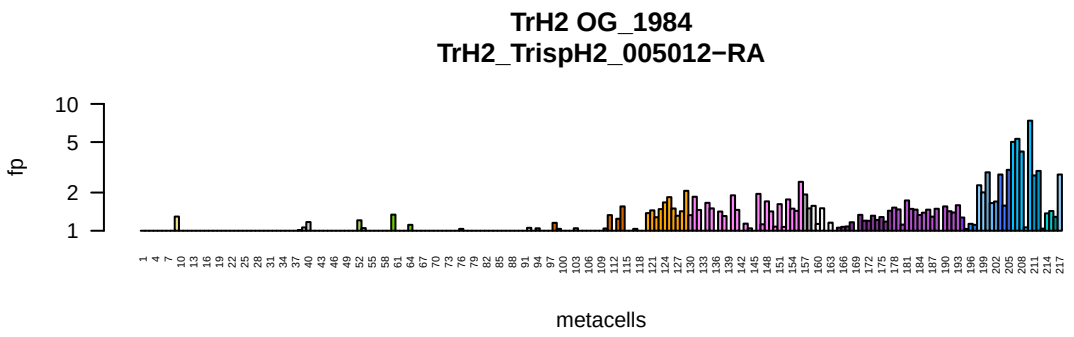

Tadh OG_186
Tadh_TriadT22195
y_subunit_beta_1,calcium_voltage_gated_channel_auxiliary_subunit_beta_4,calcium_volta
fp
metacells

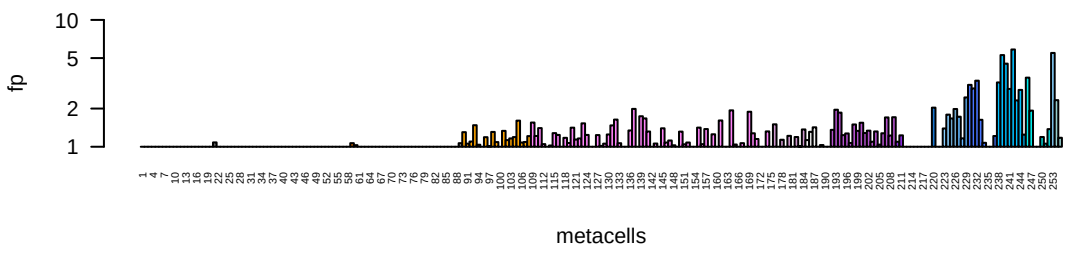

TrH2 OG_186
TrH2_TrispH2_007753−RA
y_subunit_beta_1,calcium_voltage_gated_channel_auxiliary_subunit_beta_4,calcium_volta
fp
metacells

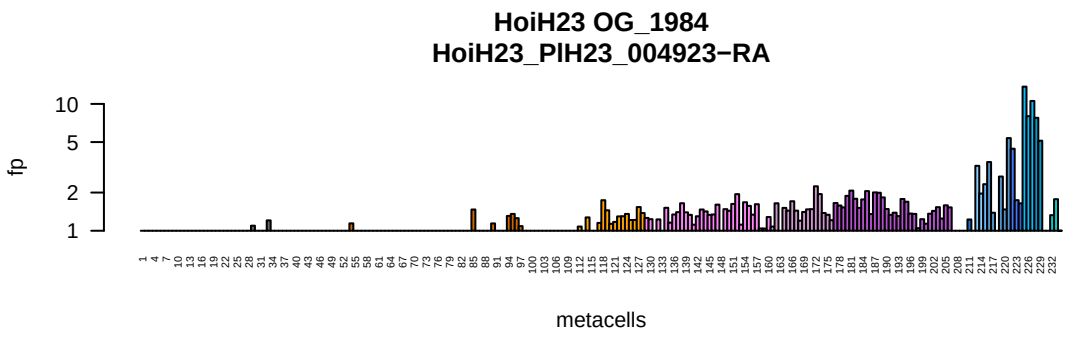

Hhon OG_186
Hhon_g00617.t1
y_subunit_beta_1,calcium_voltage_gated_channel_auxiliary_subunit_beta_4,calcium_volta
fp
metacells

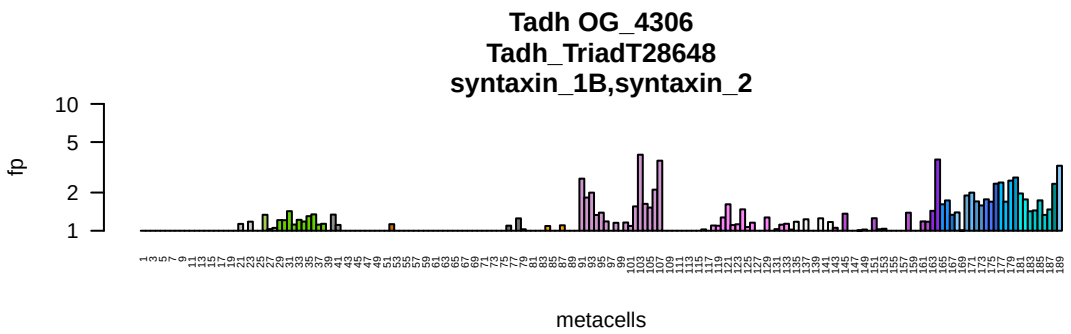

HoiH23 OG_186
HoiH23_PlH23_000883−RA
y_subunit_beta_1,calcium_voltage_gated_channel_auxiliary_subunit_beta_4,calcium_volta
fp
metacells

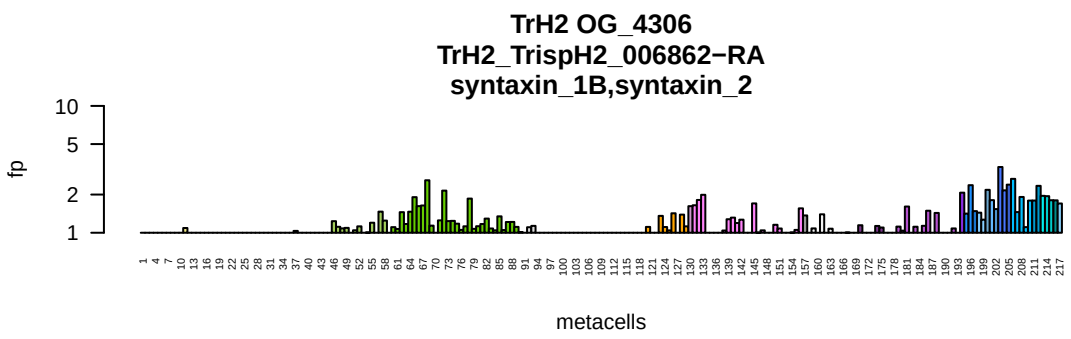

Tadh OG_1984
Tadh_wf_g9757.t1
fp
metacells

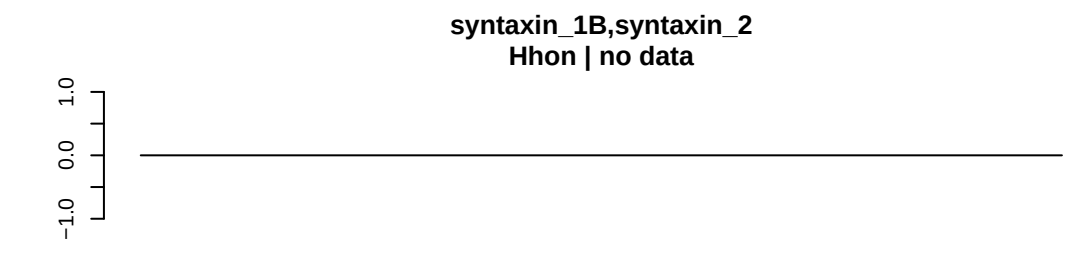

TrH2 OG_1984
TrH2_TrispH2_005012−RA
fp
metacells

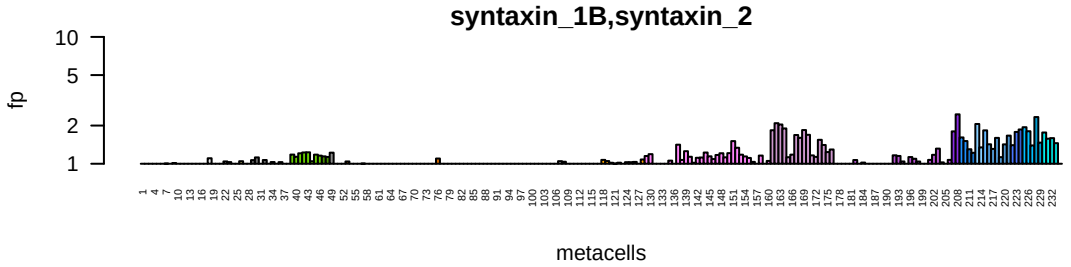

Hhon OG_1984
Hhon_g07161.t1
fp
metacells

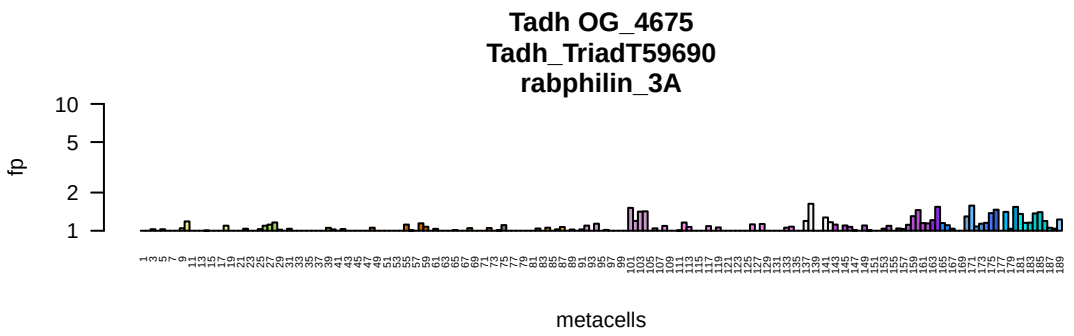

HoiH23 OG_1984
HoiH23_PlH23_004923−RA
fp
metacells

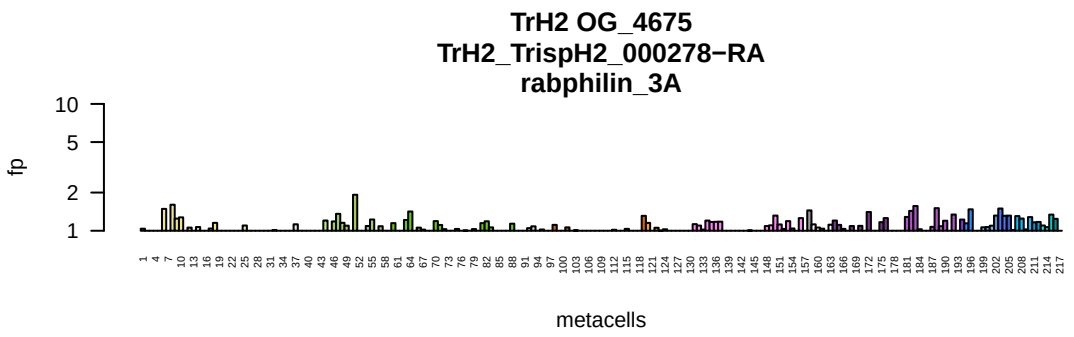

Tadh OG_4306
Tadh_TriadT28648
syntaxin_1B,syntaxin_2
fp
10
5
2
1
metacells

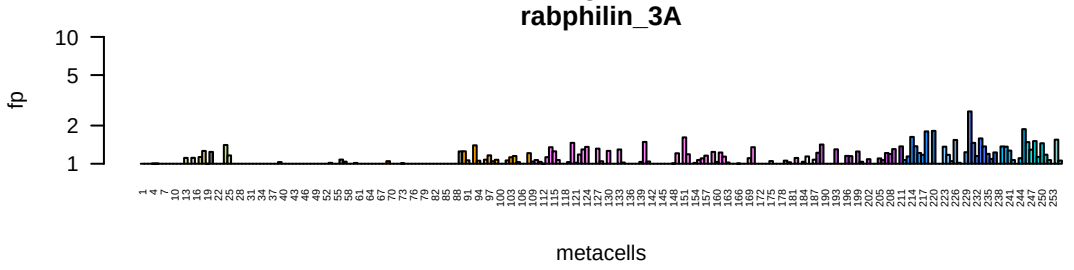

TrH2 OG_4306
TrH2_TrispH2_006862−RA
syntaxin_1B,syntaxin_2
fp
10
5
2
1
metacells

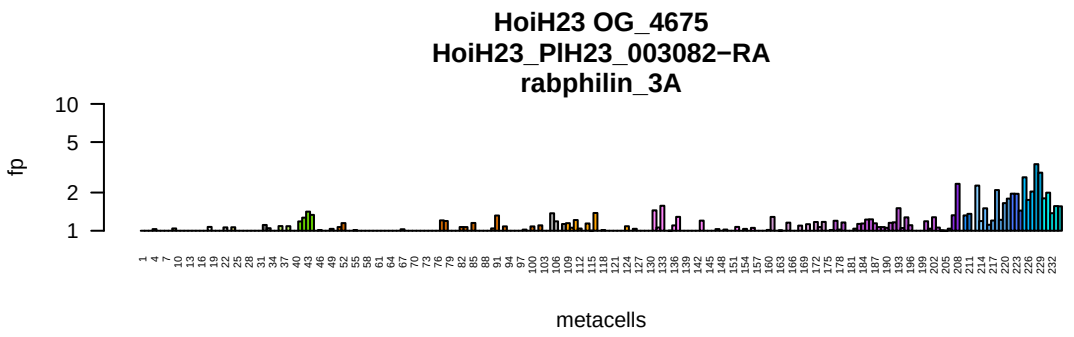

syntaxin_1B,syntaxin_2
Hhon | no data
1.0
0.0
−1.0

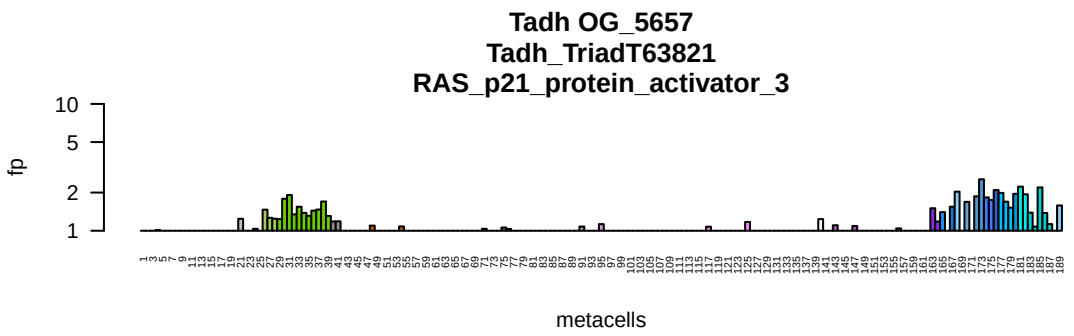

HoiH23 OG_4306
HoiH23_PlH23_000611−RA
syntaxin_1B,syntaxin_2
fp
10
5
2
1
metacells

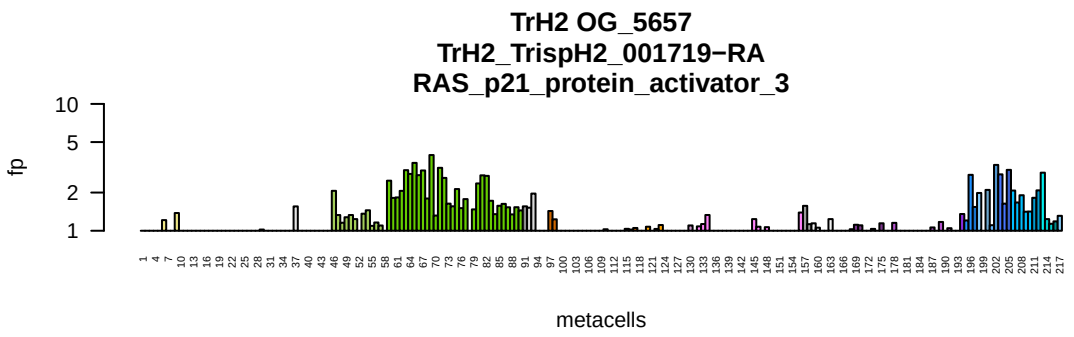

Tadh OG_4675
Tadh_TriadT59690
rabphilin_3A
fp
metacells

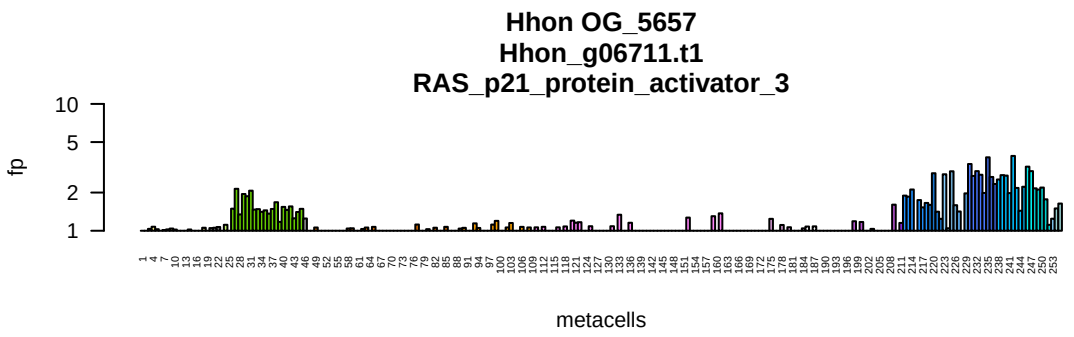

TrH2 OG_4675
TrH2_TrispH2_000278−RA
rabphilin_3A
fp
metacells

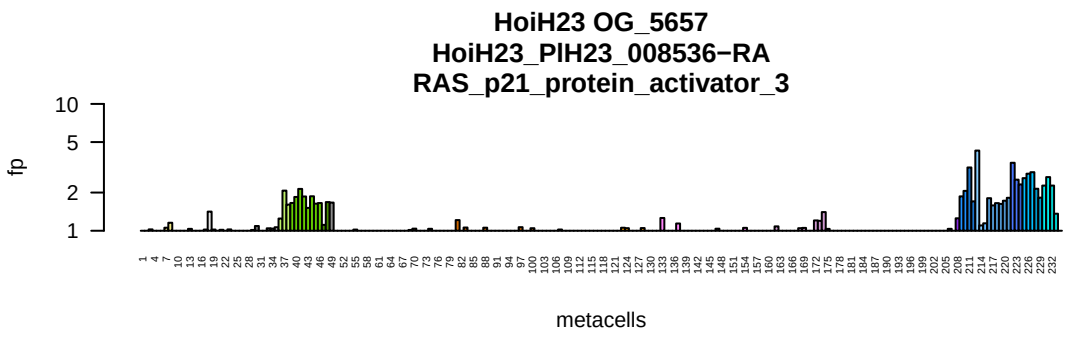

Hhon OG_4675
Hhon_g05981.t1
rabphilin_3A
fp
metacells

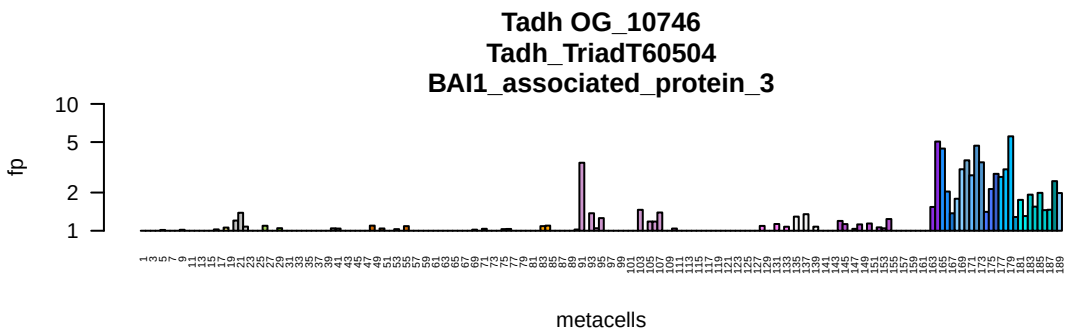

HoiH23 OG_4675
HoiH23_PlH23_003082−RA
rabphilin_3A
fp
metacells


Tadh OG_5657
Tadh_TriadT63821
RAS_p21_protein_activator_3
fp
metacells


TrH2 OG_5657
TrH2_TrispH2_001719−RA
RAS_p21_protein_activator_3
fp
metacells


Hhon OG_5657
Hhon_g06711.t1
RAS_p21_protein_activator_3
fp
metacells


HoiH23 OG_5657
HoiH23_PlH23_008536−RA
RAS_p21_protein_activator_3
fp
metacells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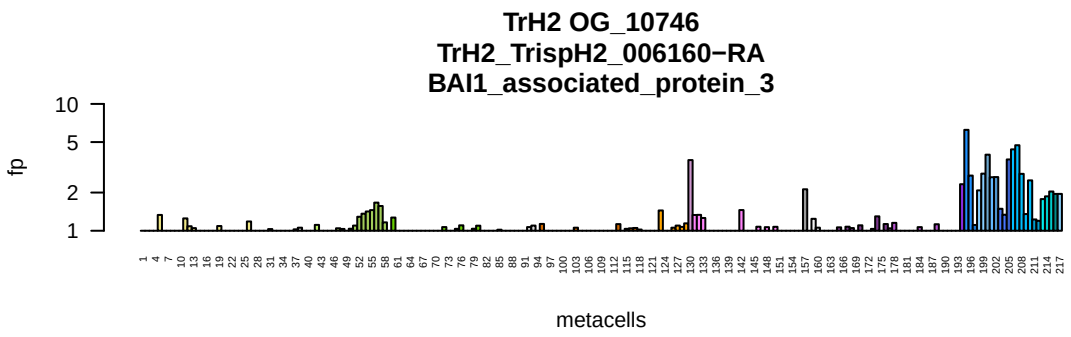
Tadh_TriadT60504
BAI1_associated_protein_3
fp
metacells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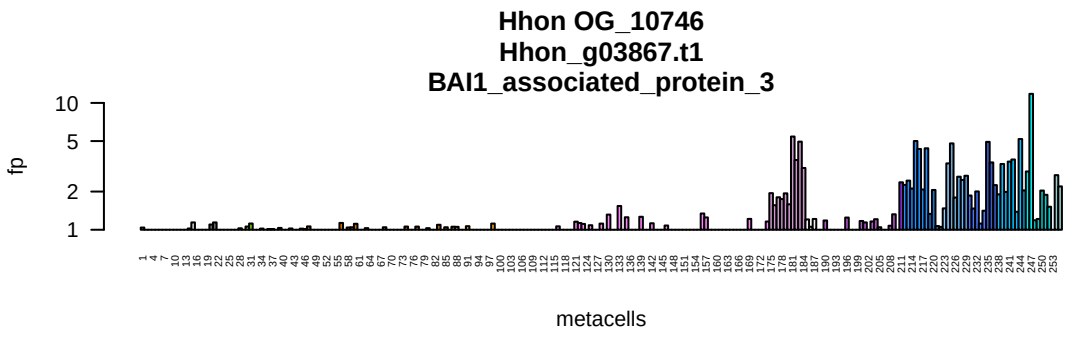
TrH2_TrispH2_006160−RA
BAI1_associated_protein_3
fp
metacells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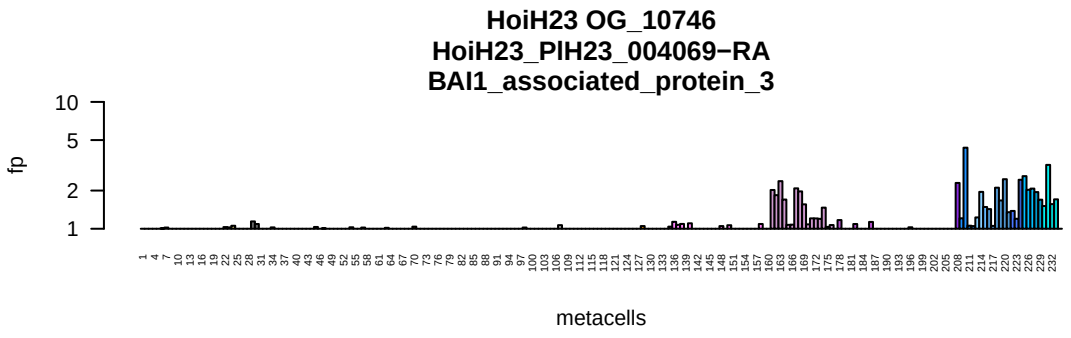
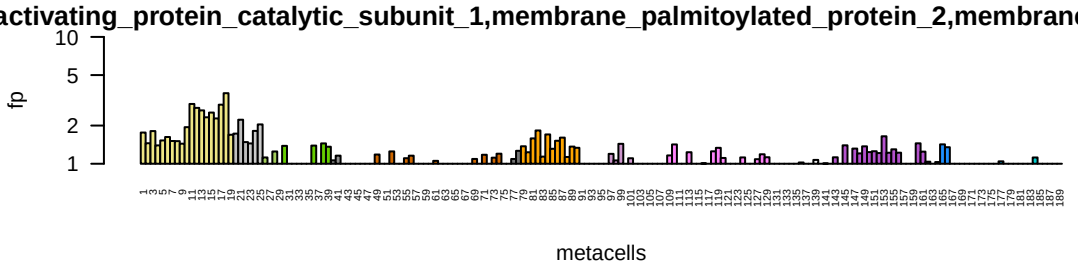
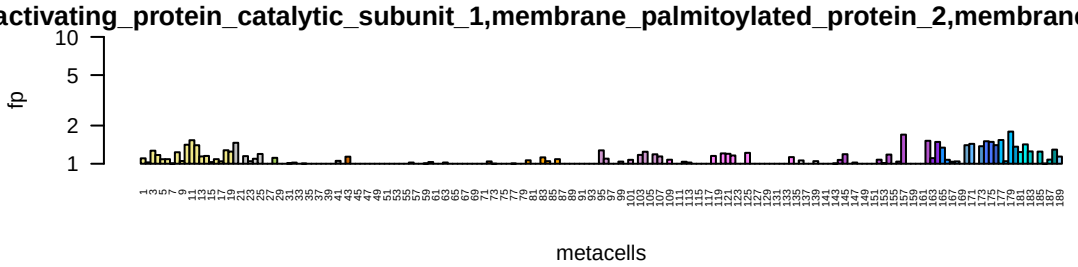
Hhon_g03867.t1
BAI1_associated_protein_3
fp
metacells


HoiH23 OG_10746
HoiH23_PlH23_004069−RA
BAI1_associated_protein_3
fp
metacells

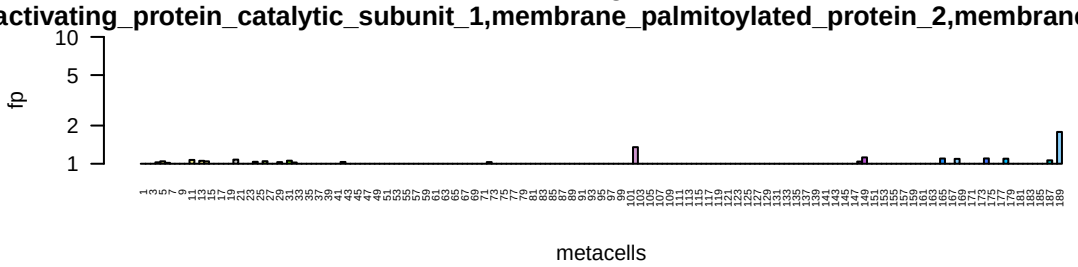
Tadh OG_3144
Tadh_TriadT63748
activating_protein_catalytic_subunit_1,membrane_palmitoylated_protein_2,membrane_palr



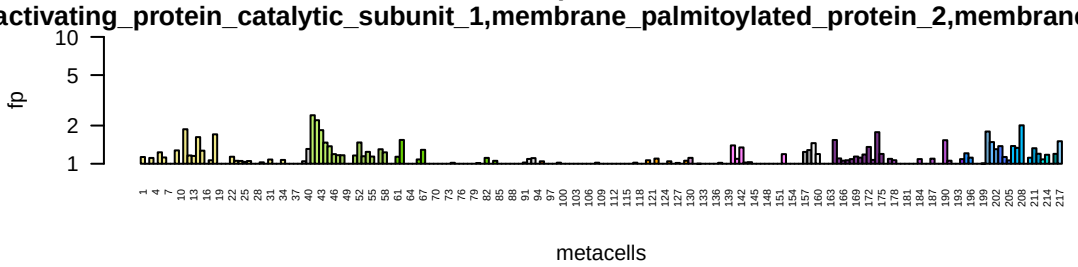
Tadh OG_3144
Tadh_TriadT23084
activating_protein_catalytic_subunit_1,membrane_palmitoylated_protein_2,membrane_palr



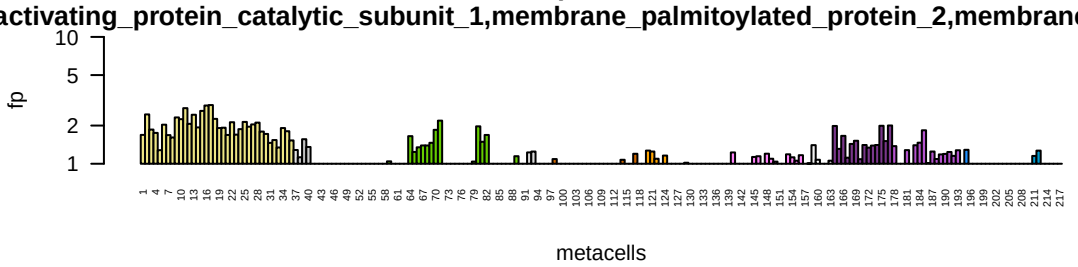
Tadh OG_3144
Tadh_wf_g3193.t1
activating_protein_catalytic_subunit_1,membrane_palmitoylated_protein_2,membrane_palr



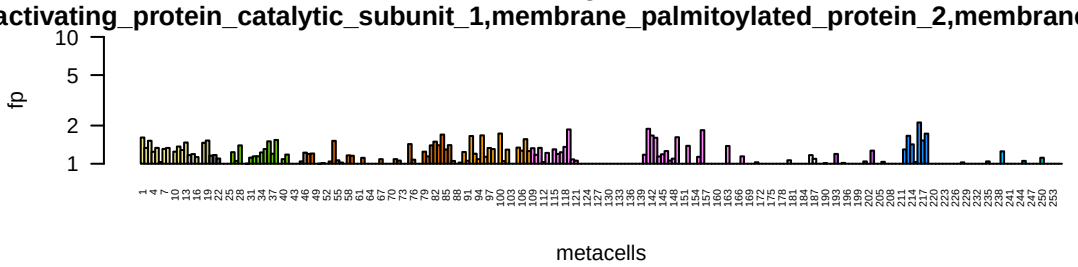
TrH2 OG_3144
TrH2_TrispH2_006042−RA
activating_protein_catalytic_subunit_1,membrane_palmitoylated_protein_2,membrane_palr



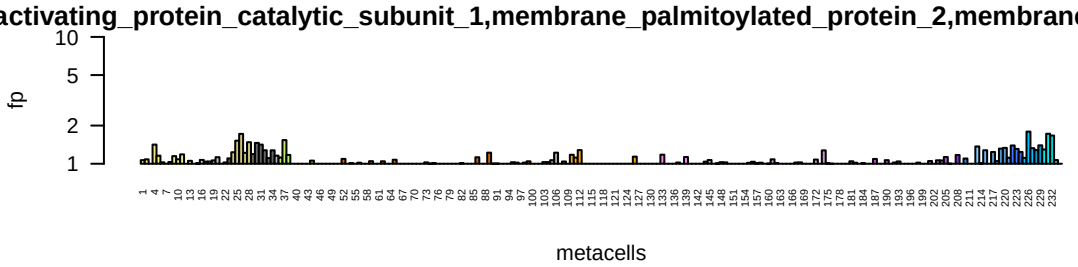
TrH2 OG_3144
TrH2_TrispH2_006041−RA
activating_protein_catalytic_subunit_1,membrane_palmitoylated_protein_2,membrane_palr



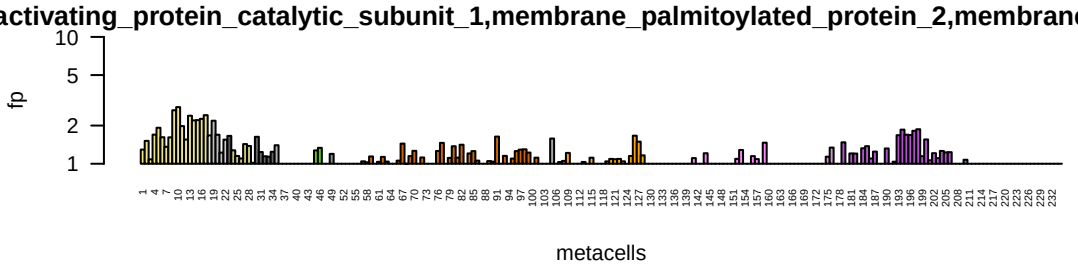
Hhon OG_3144
Hhon_g02440.t1
activating_protein_catalytic_subunit_1,membrane_palmitoylated_protein_2,membrane_palr

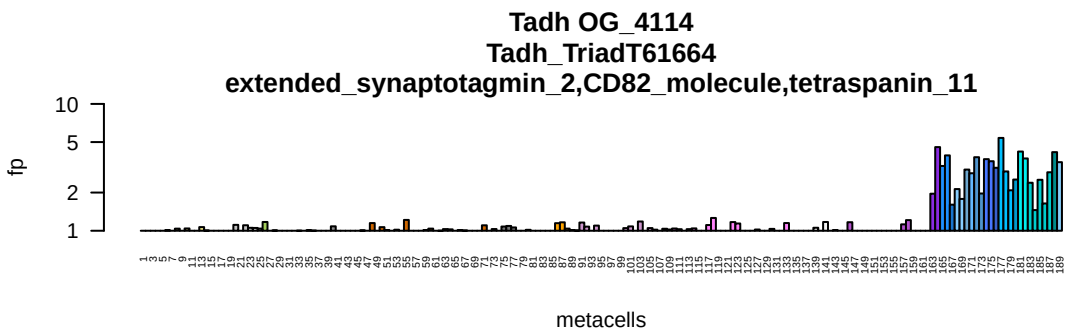


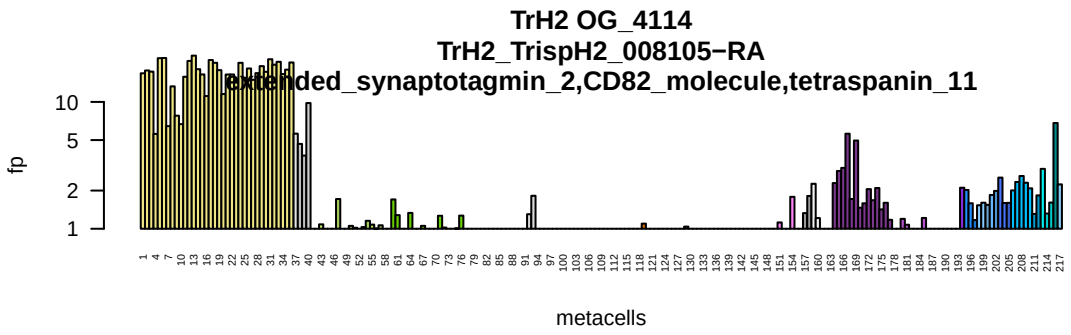
HoiH23 OG_3144
HoiH23_PlH23_002190−RA
activating_protein_catalytic_subunit_1,membrane_palmitoylated_protein_2,membrane_palr

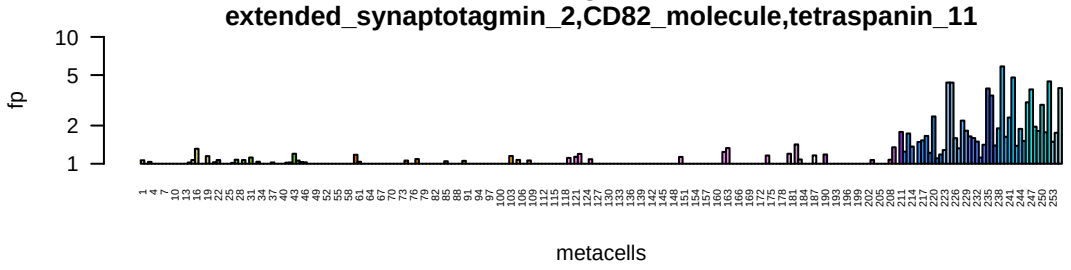


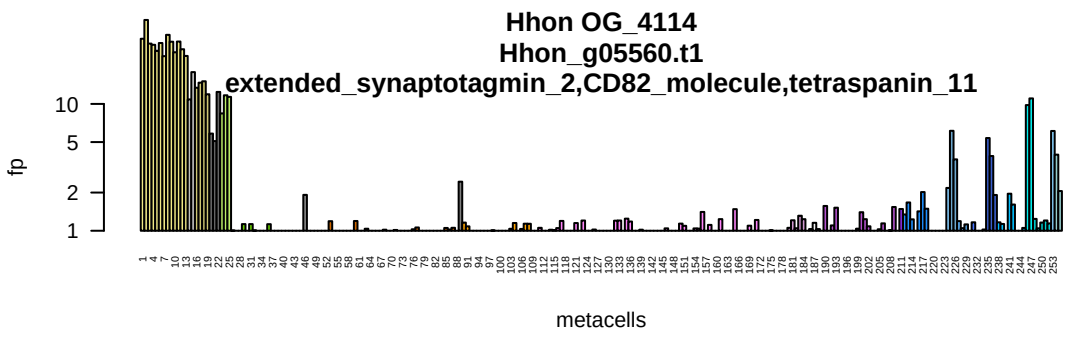
HoiH23 OG_3144
HoiH23_PlH23_002191−RA
activating_protein_catalytic_subunit_1,membrane_palmitoylated_protein_2,membrane_palr

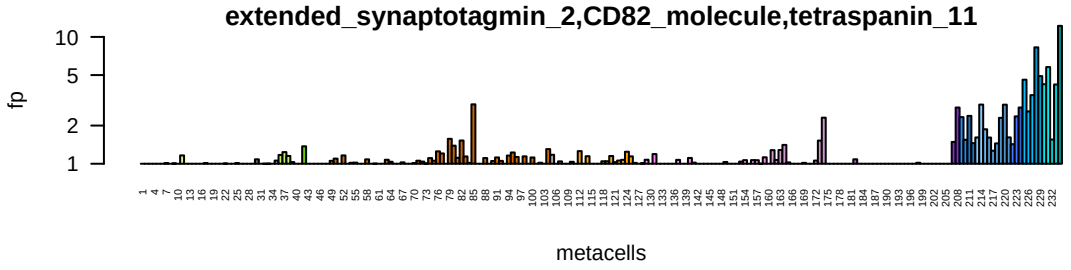


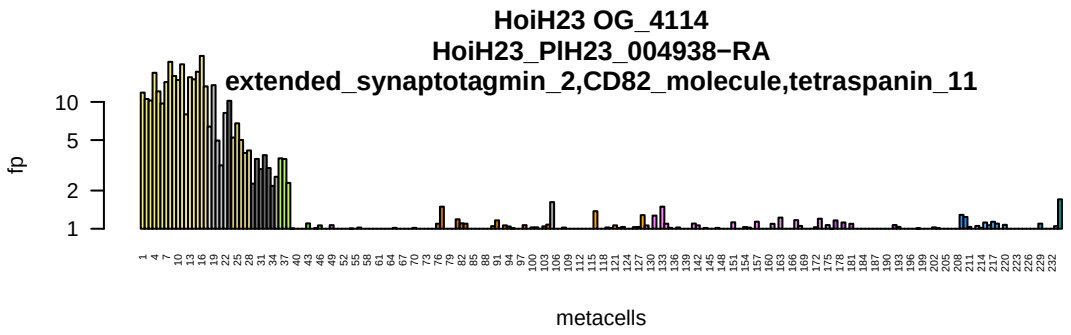

Tadh OG_4114
Tadh_TriadT61664
extended_synaptotagmin_2,CD82_molecule,tetraspanin_11
fp
10
5
2
1
metacells

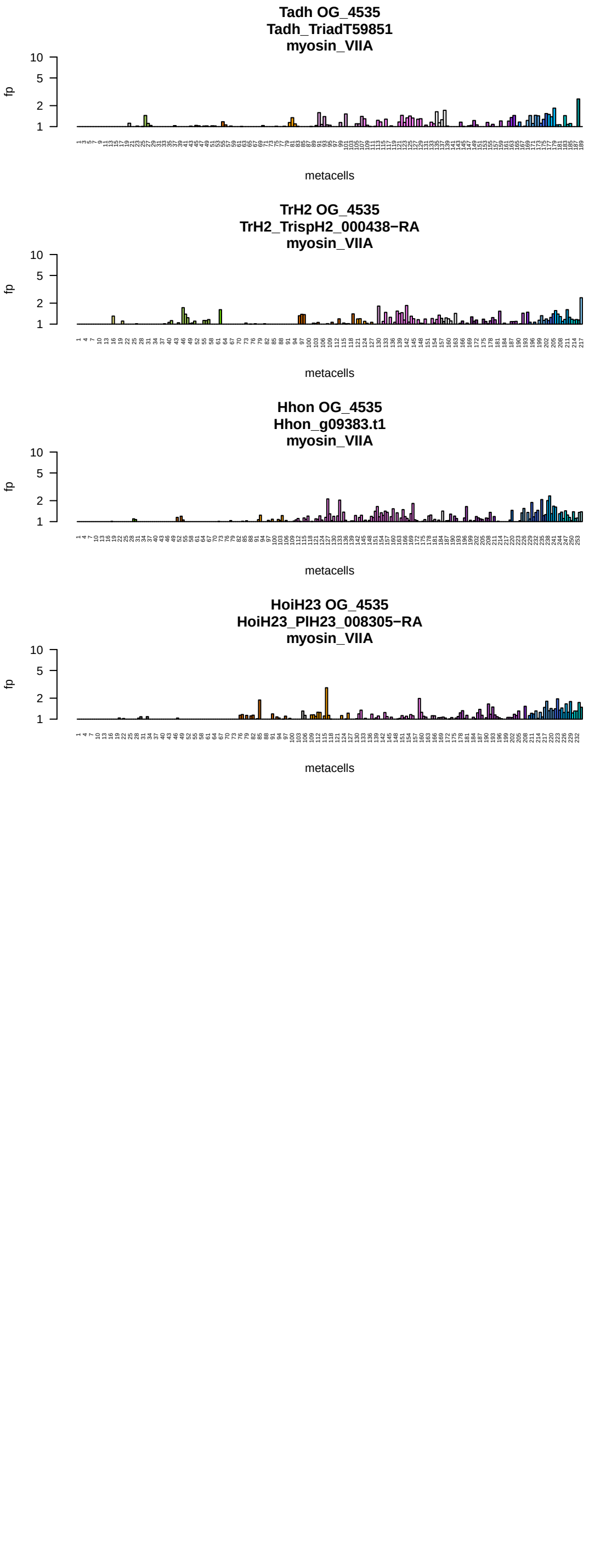

TrH2 OG_4114
TrH2_TrispH2_008105−RA
extended_synaptotagmin_2,CD82_molecule,tetraspanin_11
fp
10
5
2
1
metacells

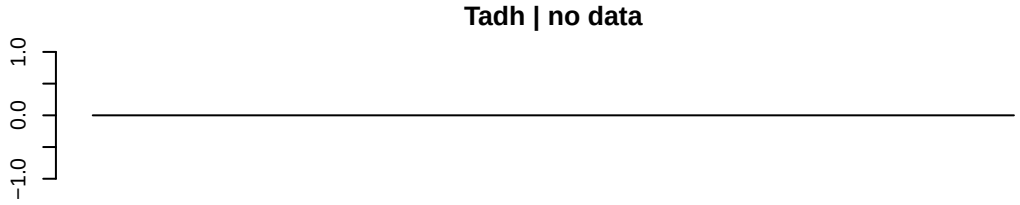

Hhon OG_4114
Hhon_g05559.t1
extended_synaptotagmin_2,CD82_molecule,tetraspanin_11
fp
10
5
2
1
metacells

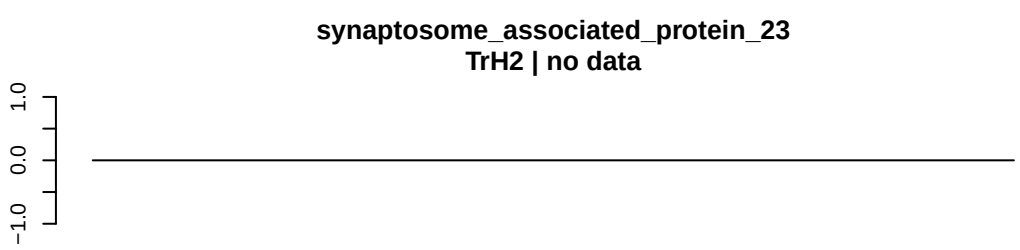

Hhon OG_4114
Hhon_g05560.t1
extended_synaptotagmin_2,CD82_molecule,tetraspanin_11
fp
10
5
2
1
metacells

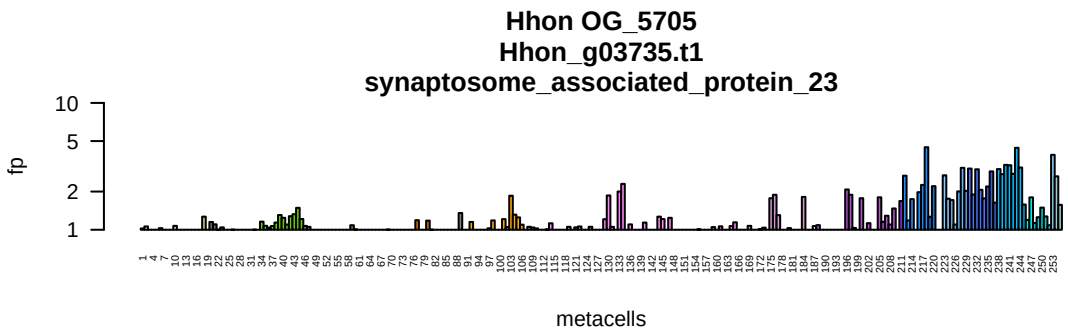

HoiH23 OG_4114
HoiH23_PlH23_004937−RA
extended_synaptotagmin_2,CD82_molecule,tetraspanin_11
fp
10
5
2
1
metacells

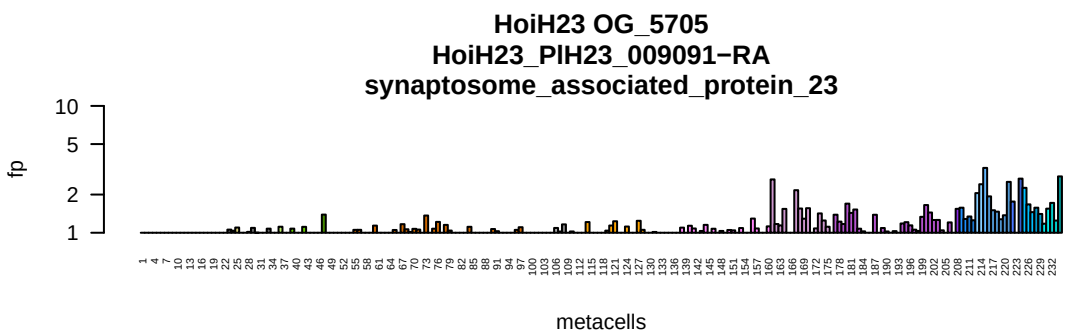

HoiH23 OG_4114
HoiH23_PlH23_004938−RA
extended_synaptotagmin_2,CD82_molecule,tetraspanin_11
fp
10
5
2
1
metacells


Tadh OG_4535
Tadh_TriadT59851
myosin_VIIA
fp
metacells
TrH2 OG_4535
TrH2_TrispH2_000438−RA
myosin_VIIA
fp
metacells
Hhon OG_4535
Hhon_g09383.t1
myosin_VIIA
fp
metacells
HoiH23 OG_4535
HoiH23_PlH23_008305−RA
myosin_VIIA
fp
metacells

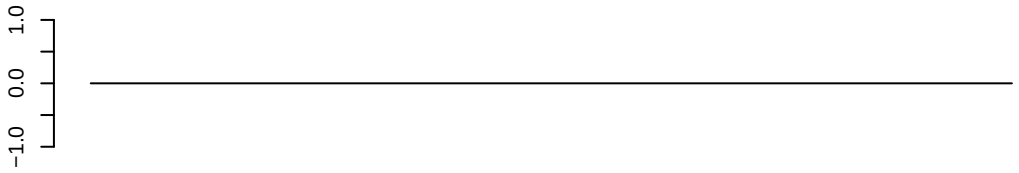

synaptosome_associated_protein_23
Tadh | no data
1.0
0.0
−1.0


synaptosome_associated_protein_23
TrH2 | no data
1.0
0.0
−1.0

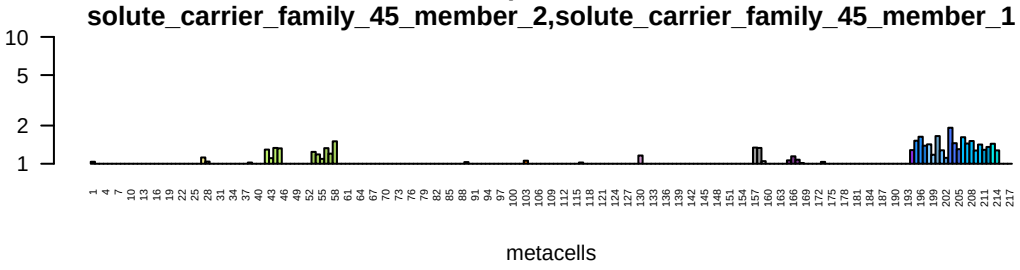

Hhon OG_5705
Hhon_g03735.t1
synaptosome_associated_protein_23
fp
10
5
2
1
metacells


HoiH23 OG_5705
HoiH23_PlH23_009091−RA
synaptosome_associated_protein_23
fp
10
5
2
1
metacells

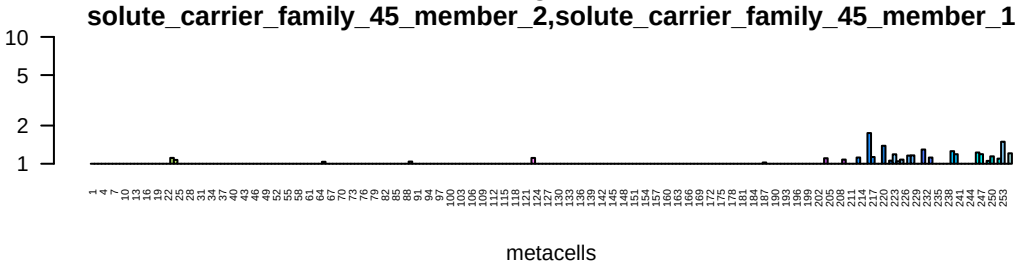
solute_carrier_family_45_member_2,solute_carrier_family_45_member_1
Tadh | no data



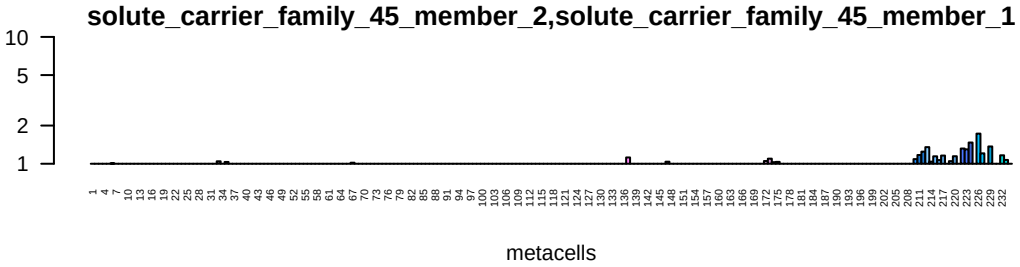
TrH2 OG_5858
TrH2_TrispH2_002756−RA
solute_carrier_family_45_member_2,solute_carrier_family_45_member_1



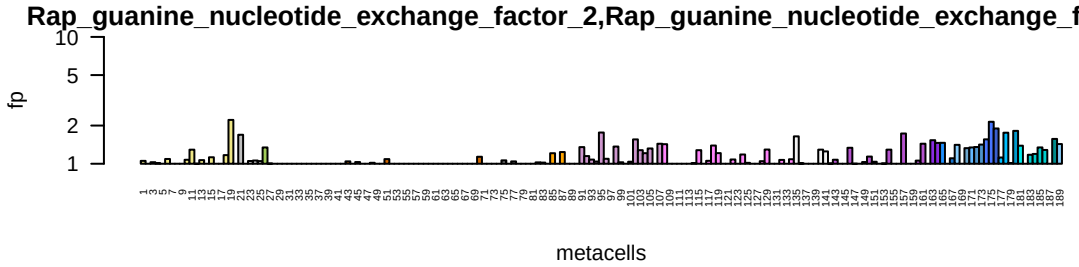
Hhon OG_5858
Hhon_g00430.t1
solute_carrier_family_45_member_2,solute_carrier_family_45_member_1



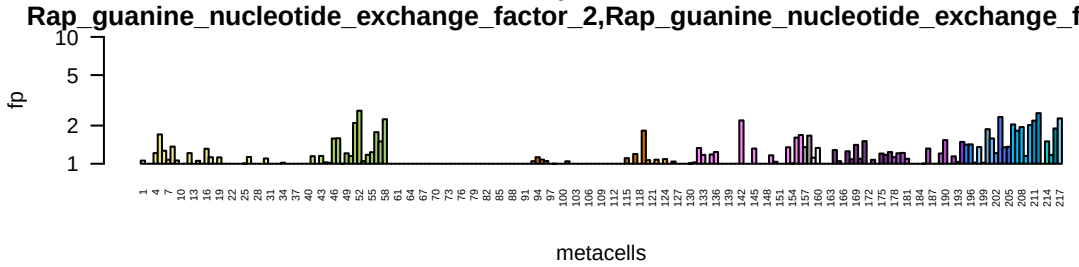
HoiH23 OG_5858
HoiH23_PlH23_001806−RA
solute_carrier_family_45_member_2,solute_carrier_family_45_member_1



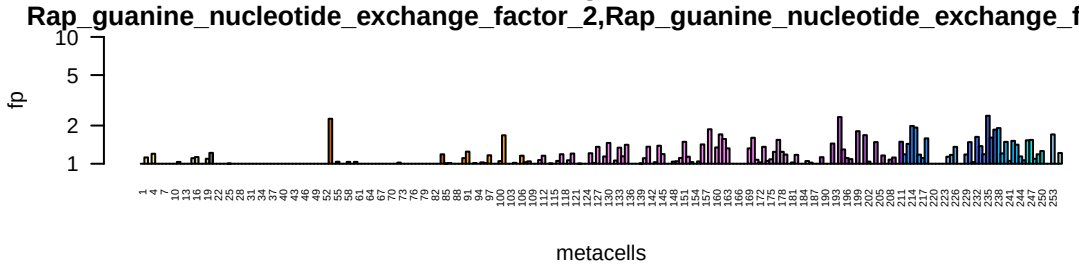
Tadh OG_6886
Tadh_TriadT59148
Rap_guanine_nucleotide_exchange_factor_2,Rap_guanine_nucleotide_exchange_factor_



TrH2 OG_6886
TrH2_TrispH2_009461−RA
Rap_guanine_nucleotide_exchange_factor_2,Rap_guanine_nucleotide_exchange_factor_

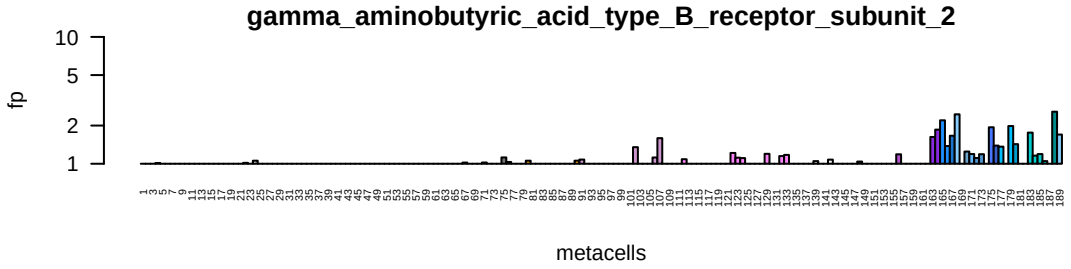


Hhon OG_6886
Hhon_g09898.t1
Rap_guanine_nucleotide_exchange_factor_2,Rap_guanine_nucleotide_exchange_factor_

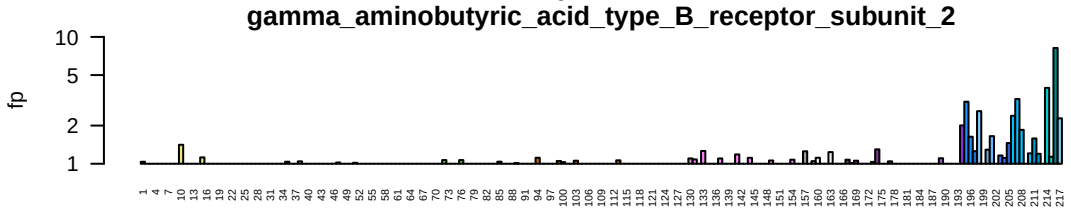


HoiH23 OG_6886
HoiH23_PlH23_000593−RA
Rap_guanine_nucleotide_exchange_factor_2,Rap_guanine_nucleotide_exchange_factor_

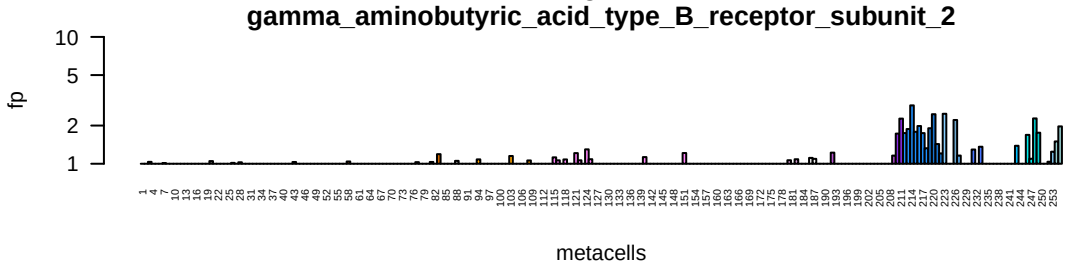
Tadh OG_8182
Tadh_TriadT57193
gamma_aminobutyric_acid_type_B_receptor_subunit_2



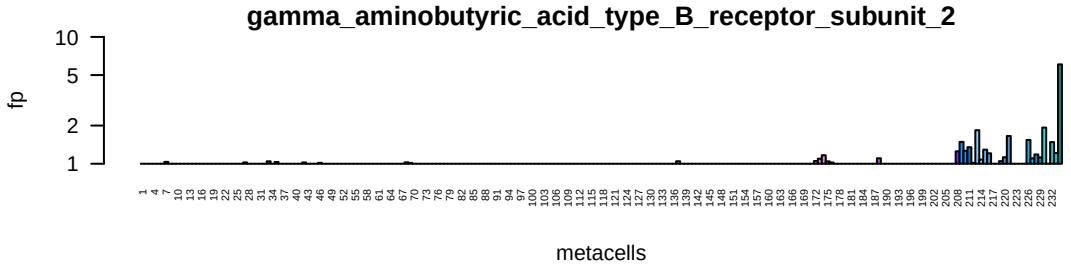
TrH2 OG_8182
TrH2_TrispH2_003275−RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2



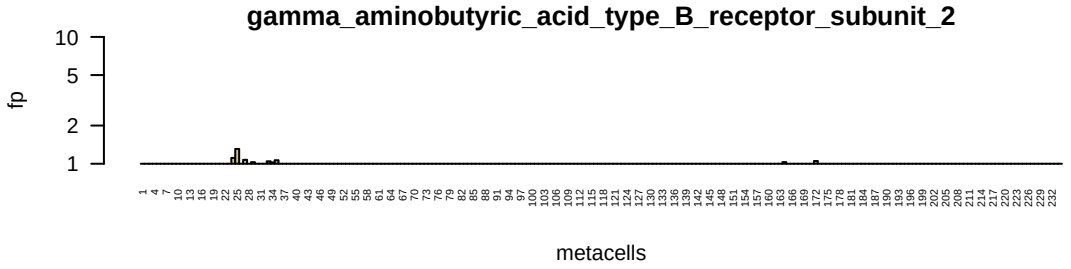
Hhon OG_8182
Hhon_g04629.t1
gamma_aminobutyric_acid_type_B_receptor_subunit_2



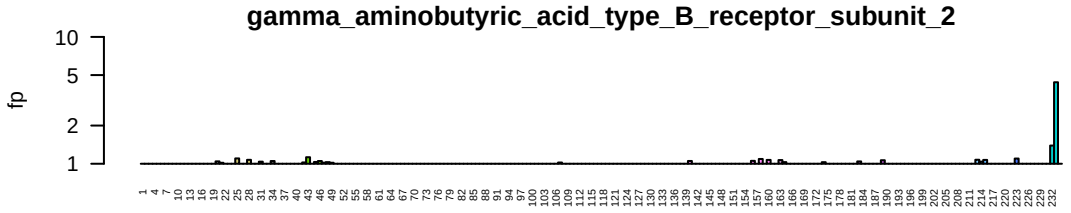
HoiH23 OG_8182
HoiH23_PlH23_010698−RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2



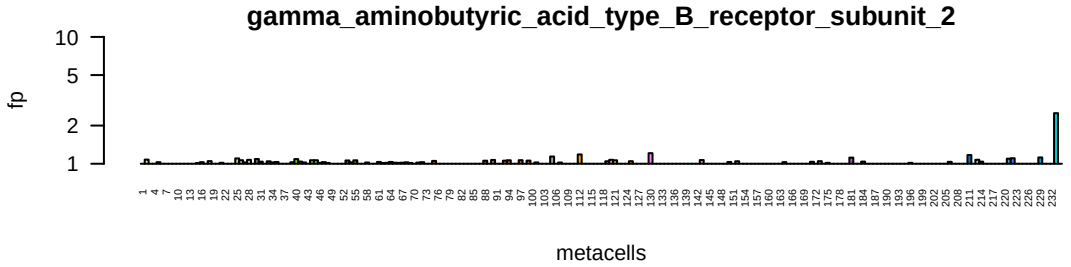
HoiH23 OG_8182
HoiH23_PlH23_005534−RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2

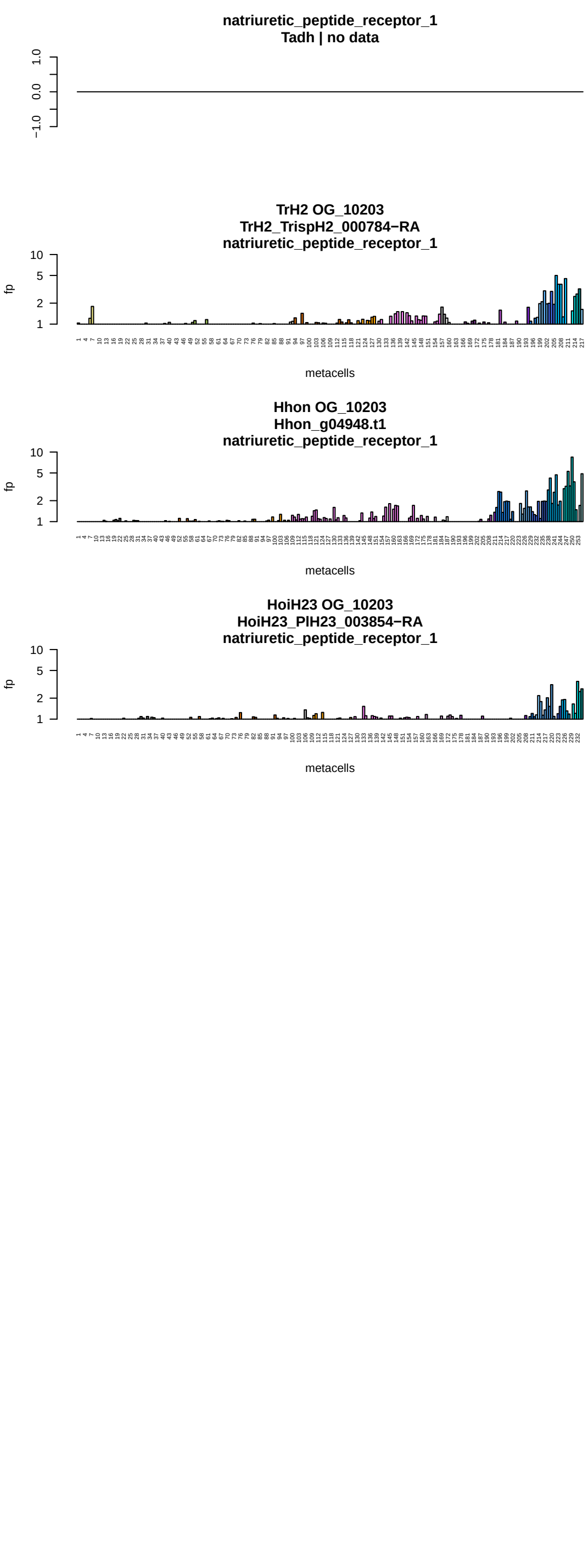


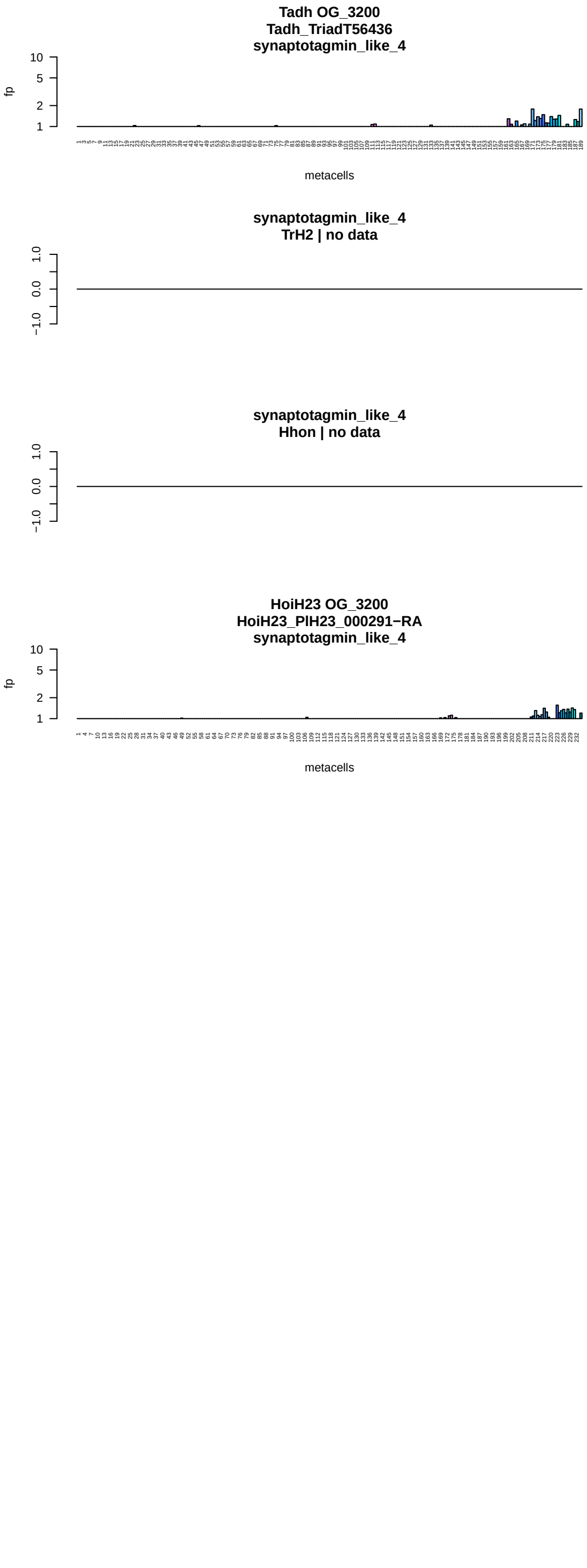
HoiH23 OG_8182
HoiH23_PlH23_005535−RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2



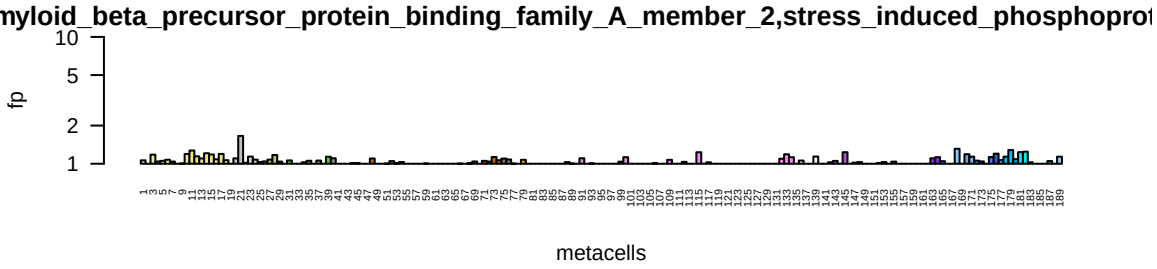
HoiH23 OG_8182
HoiH23_PlH23_005536−RA
gamma_aminobutyric_acid_type_B_receptor_subunit_2



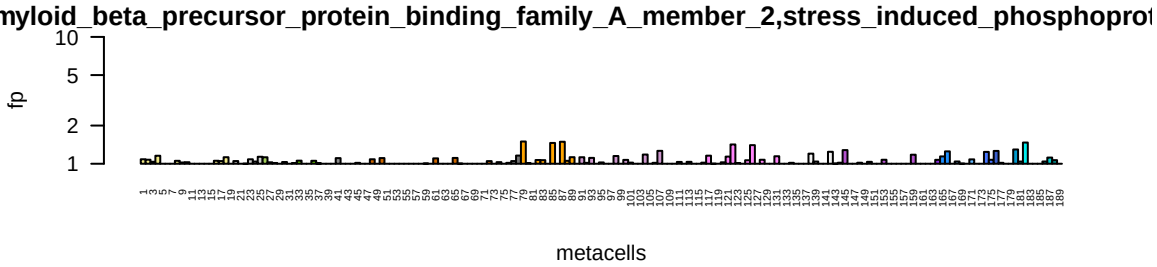

natriuretic_peptide_receptor_1
Tadh | no data
1.0
0.0
−1.0
TrH2 OG_10203
TrH2_TrispH2_000784−RA
natriuretic_peptide_receptor_1
fp
10
5
2
1
metacells
Hhon OG_10203
Hhon_g04948.t1
natriuretic_peptide_receptor_1
fp
10
5
2
1
metacells
HoiH23 OG_10203
HoiH23_PlH23_003854−RA
natriuretic_peptide_receptor_1
fp
10
5
2
1
metacells


Tadh OG_3200
Tadh_TriadT56436
synaptotagmin_like_4
fp
10
5
2
1
metacells
synaptotagmin_like_4
TrH2 | no data
1.0
0.0
−1.0
synaptotagmin_like_4
Hhon | no data
1.0
0.0
−1.0
HoiH23 OG_3200
HoiH23_PlH23_000291−RA
synaptotagmin_like_4
fp
10
5
2
1
metacells

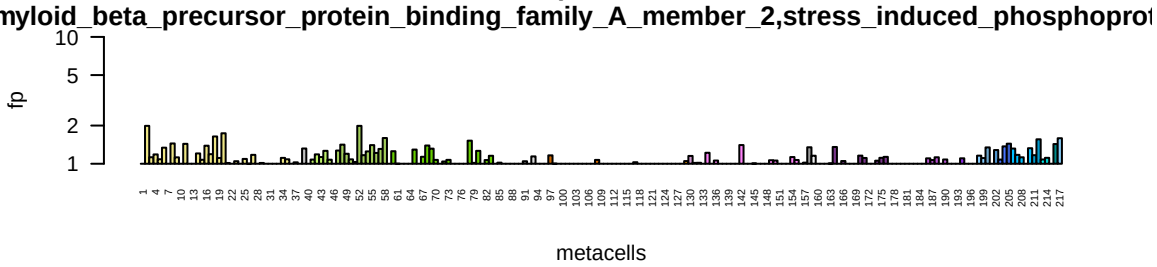
Tadh OG_3728
Tadh_TriadT23350
myloid_beta_precursor_protein_binding_family_A_member_2,stress_induced_phosphoprot



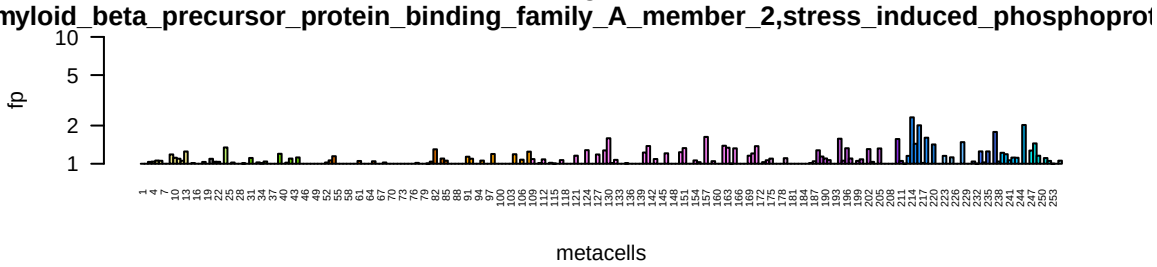
Tadh OG_3728
Tadh_TriadT23199
myloid_beta_precursor_protein_binding_family_A_member_2,stress_induced_phosphoprot



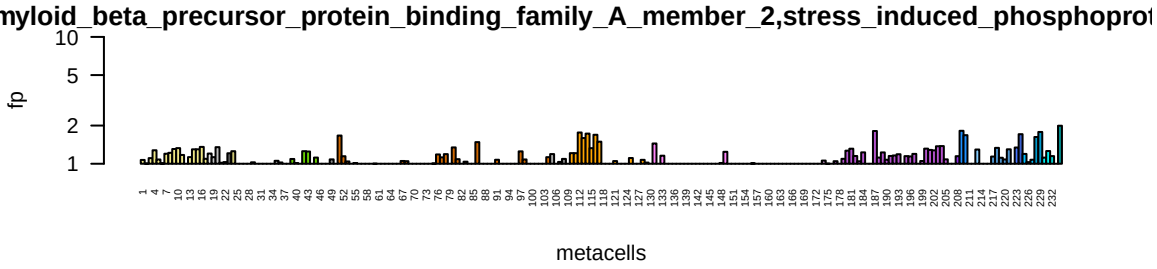
TrH2 OG_3728
TrH2_TrispH2_006762−RA
myloid_beta_precursor_protein_binding_family_A_member_2,stress_induced_phosphoprot



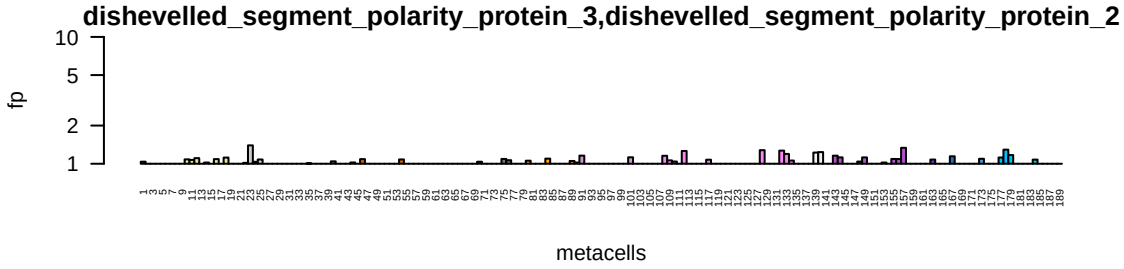
Hhon OG_3728
Hhon_g08341.t1
myloid_beta_precursor_protein_binding_family_A_member_2,stress_induced_phosphoprot



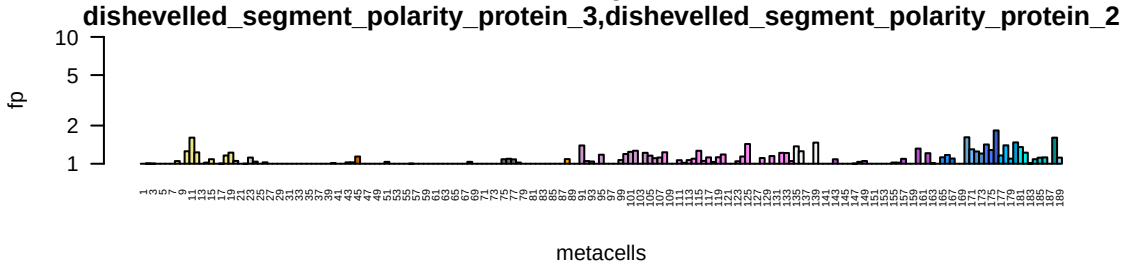
HoiH23 OG_3728
HoiH23_PlH23_004813−RA
myloid_beta_precursor_protein_binding_family_A_member_2,stress_induced_phosphoprot



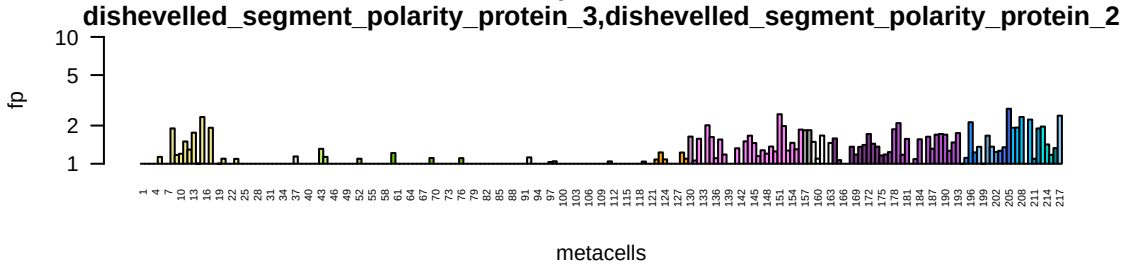
Tadh OG_3858
Tadh_TriadT26894
dishevelled_segment_polarity_protein_3,dishevelled_segment_polarity_protein_2



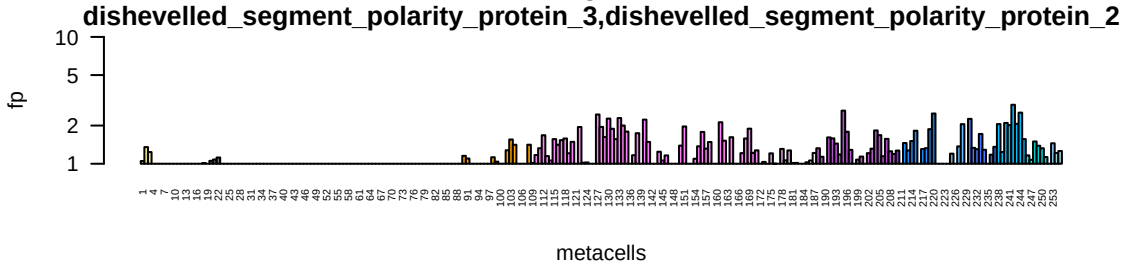
Tadh OG_3858
Tadh_wf_g6315.t1
dishevelled_segment_polarity_protein_3,dishevelled_segment_polarity_protein_2



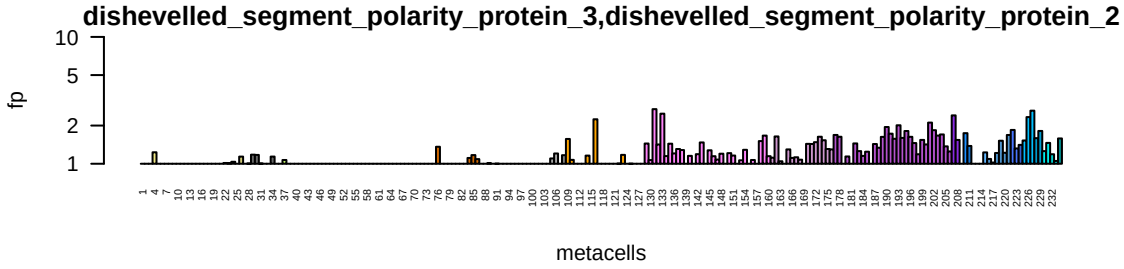
TrH2 OG_3858
TrH2_TrispH2_009138−RA
dishevelled_segment_polarity_protein_3,dishevelled_segment_polarity_protein_2



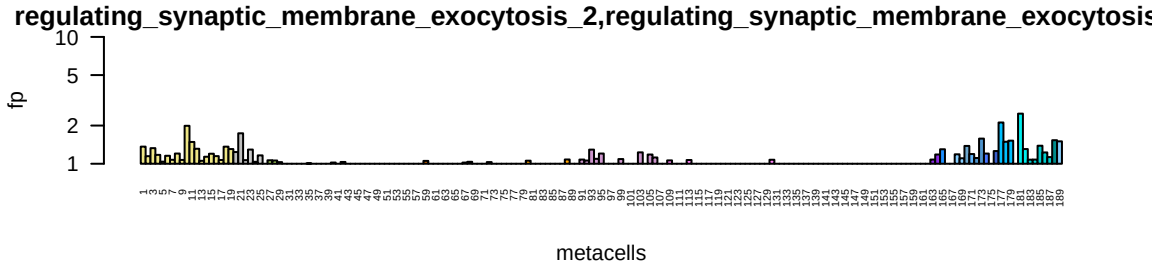
Hhon OG_3858
Hhon_g01922.t1
dishevelled_segment_polarity_protein_3,dishevelled_segment_polarity_protein_2



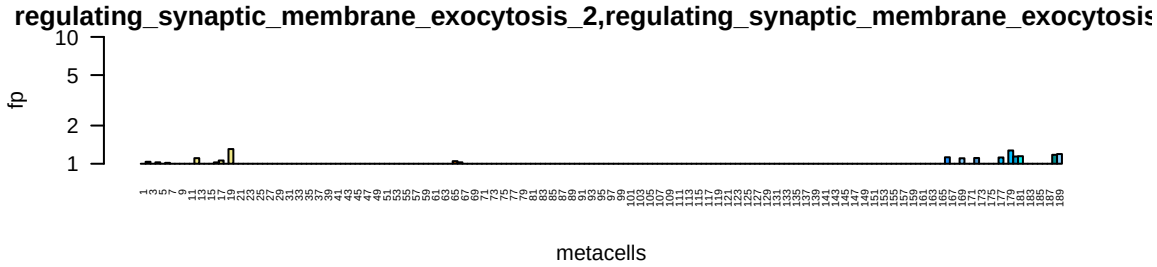
HoiH23 OG_3858
HoiH23_PlH23_001486−RA
dishevelled_segment_polarity_protein_3,dishevelled_segment_polarity_protein_2



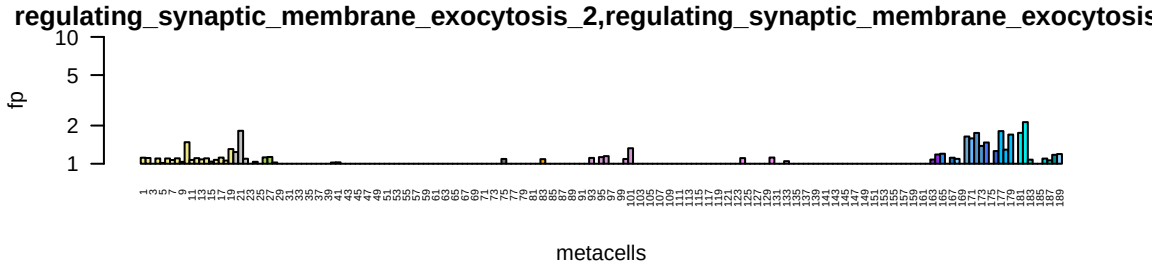
Tadh OG_3902
Tadh_TriadT58203
regulating_synaptic_membrane_exocytosis_2,regulating_synaptic_membrane_exocytosis



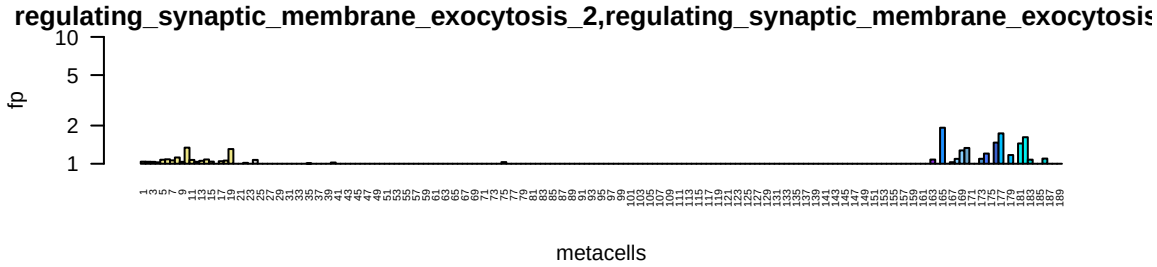
Tadh OG_3902
Tadh_TriadT58201
regulating_synaptic_membrane_exocytosis_2,regulating_synaptic_membrane_exocytosis



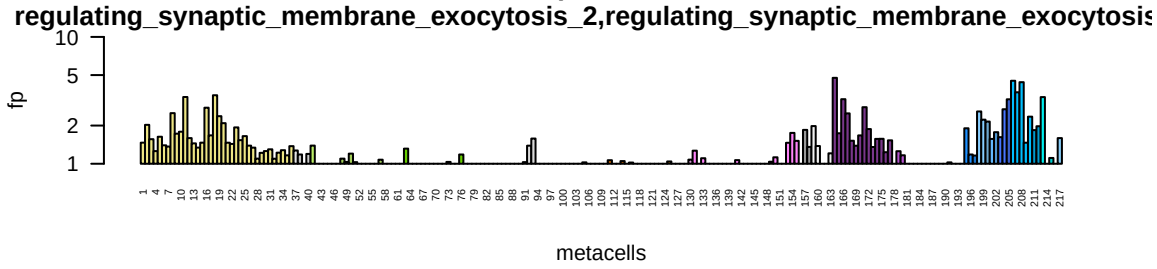
Tadh OG_3902
Tadh_TriadT58202
regulating_synaptic_membrane_exocytosis_2,regulating_synaptic_membrane_exocytosis



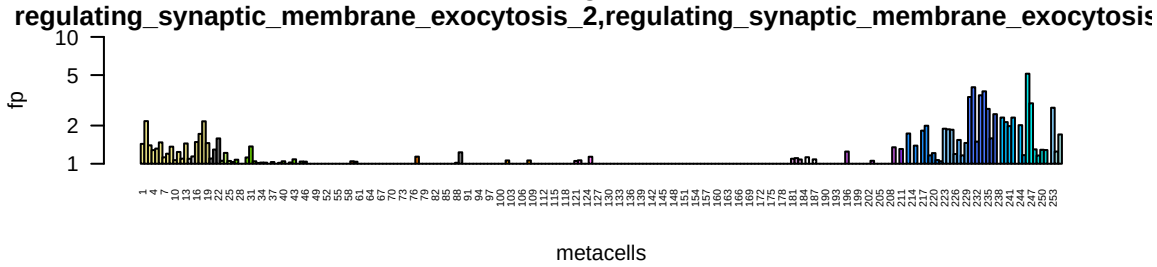
Tadh OG_3902
Tadh_TriadT58200
regulating_synaptic_membrane_exocytosis_2,regulating_synaptic_membrane_exocytosis



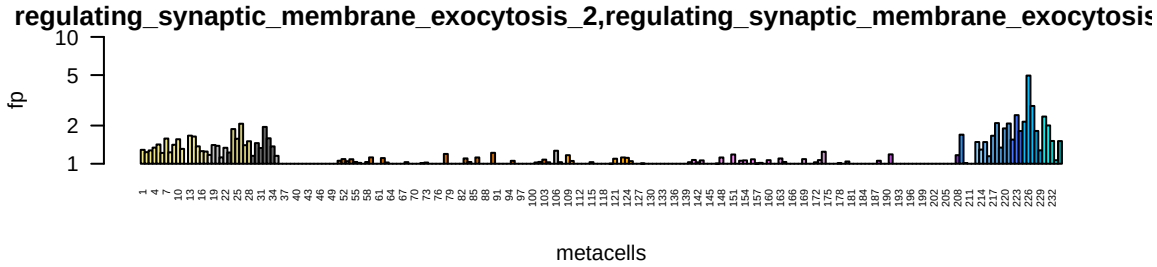
TrH2 OG_3902
TrH2_TrispH2_000666−RA
regulating_synaptic_membrane_exocytosis_2,regulating_synaptic_membrane_exocytosis

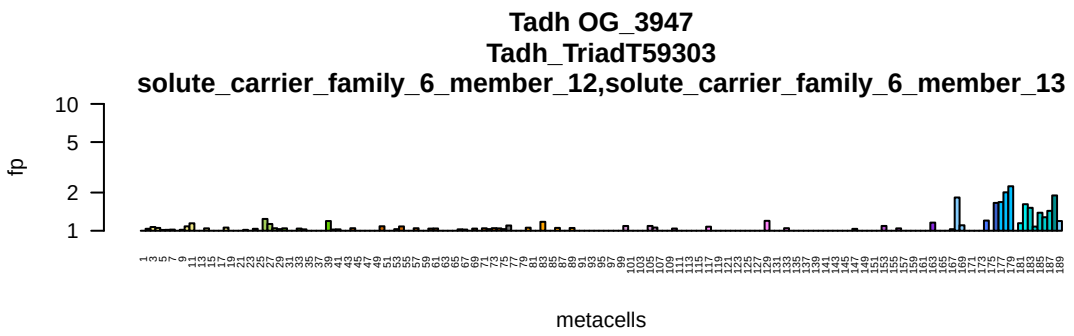


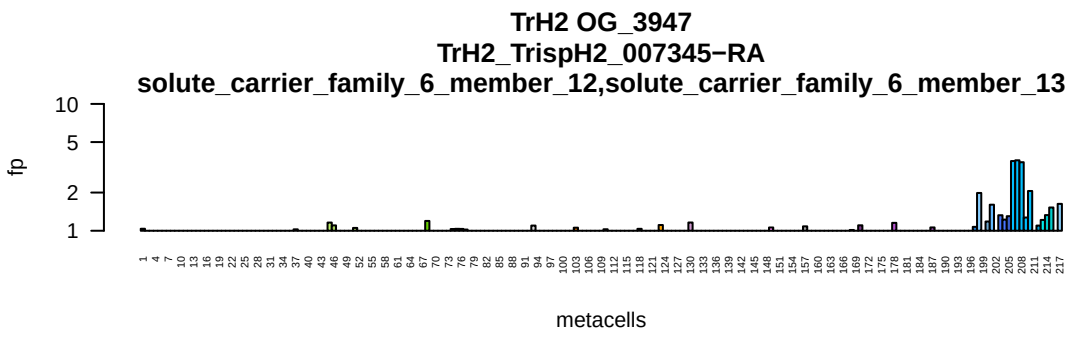
Hhon OG_3902
Hhon_g00025.t1
regulating_synaptic_membrane_exocytosis_2,regulating_synaptic_membrane_exocytosis

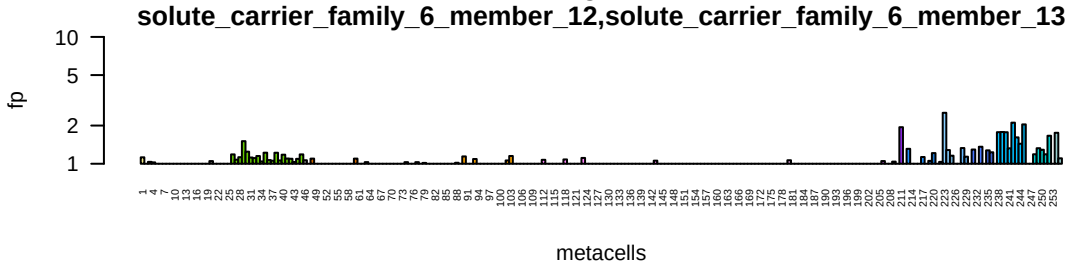


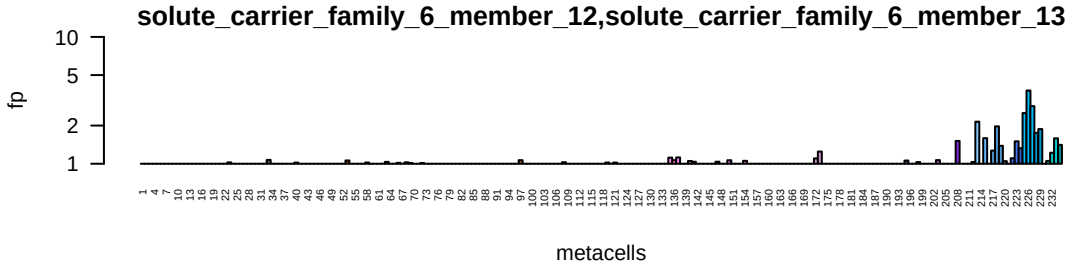
HoiH23 OG_3902
HoiH23_PlH23_008516−RA
regulating_synaptic_membrane_exocytosis_2,regulating_synaptic_membrane_exocytosis

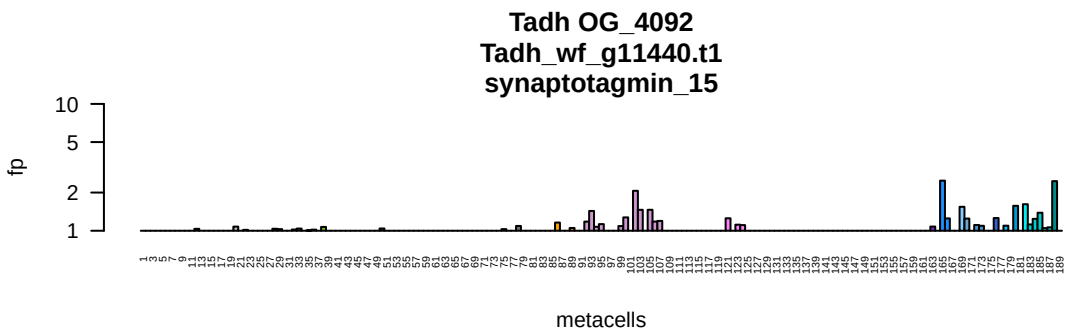


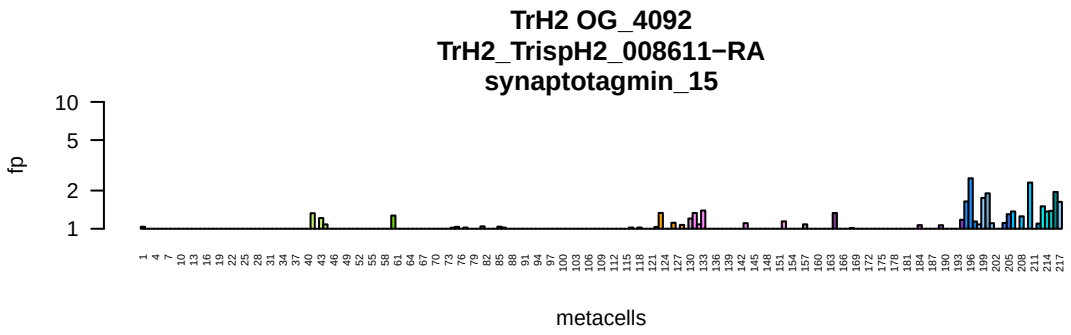

Tadh OG_3947
Tadh_TriadT59303
solute_carrier_family_6_member_12,solute_carrier_family_6_member_13
fp
metacells

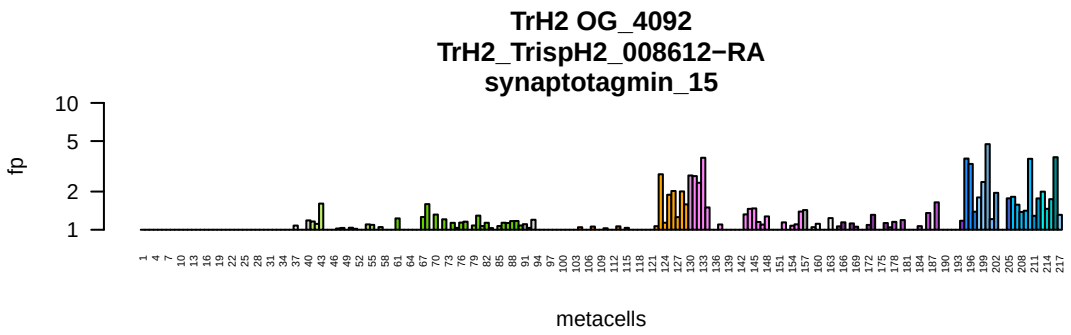

TrH2 OG_3947
TrH2_TrispH2_007345−RA
solute_carrier_family_6_member_12,solute_carrier_family_6_member_13
fp
metacells

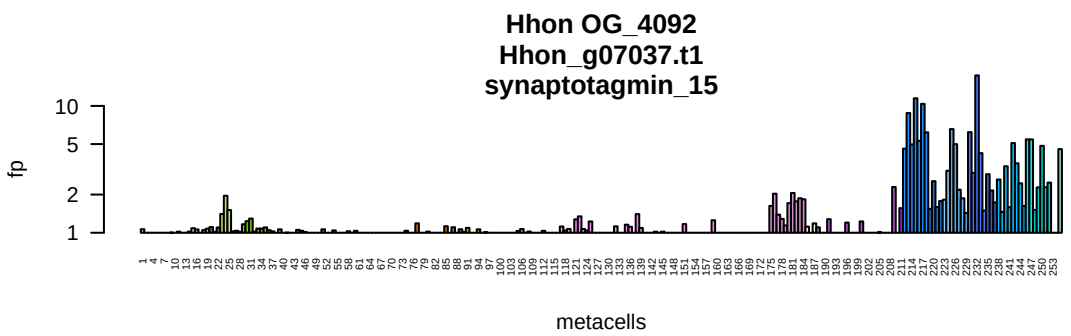

Hhon OG_3947
Hhon_g09021.t1
solute_carrier_family_6_member_12,solute_carrier_family_6_member_13
fp
metacells

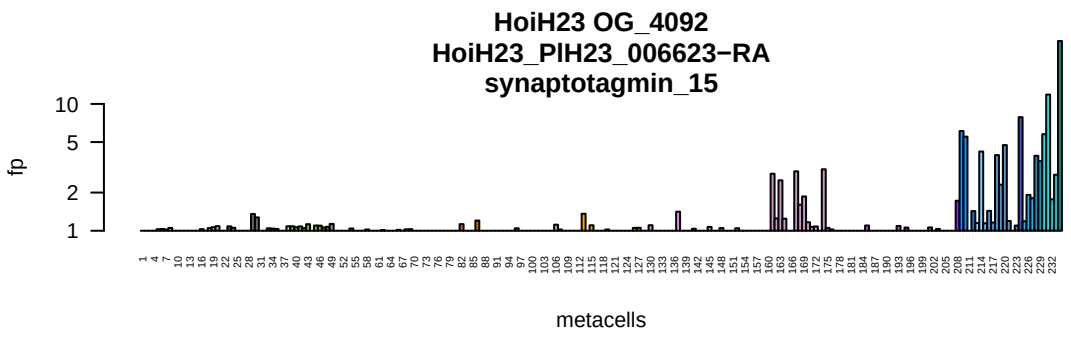

HoiH23 OG_3947
HoiH23_PlH23_007652−RA
solute_carrier_family_6_member_12,solute_carrier_family_6_member_13
fp
metacells


Tadh OG_4092
Tadh_wf_g11440.t1
synaptotagmin_15
fp
10
5
2
1
metacells

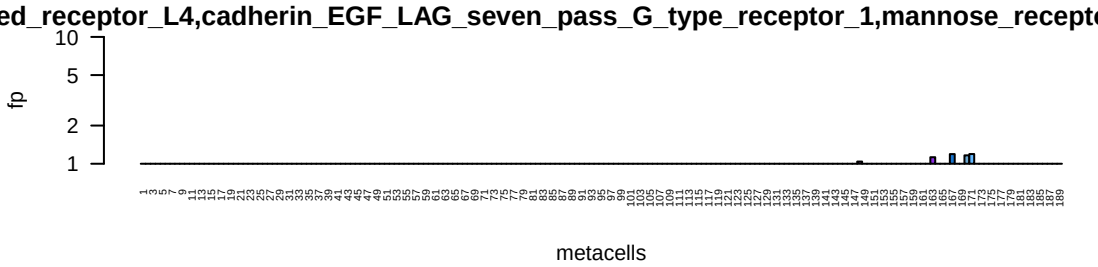

TrH2 OG_4092
TrH2_TrispH2_008611−RA
synaptotagmin_15
fp
10
5
2
1
metacells


TrH2 OG_4092
TrH2_TrispH2_008612−RA
synaptotagmin_15
fp
10
5
2
1
metacells

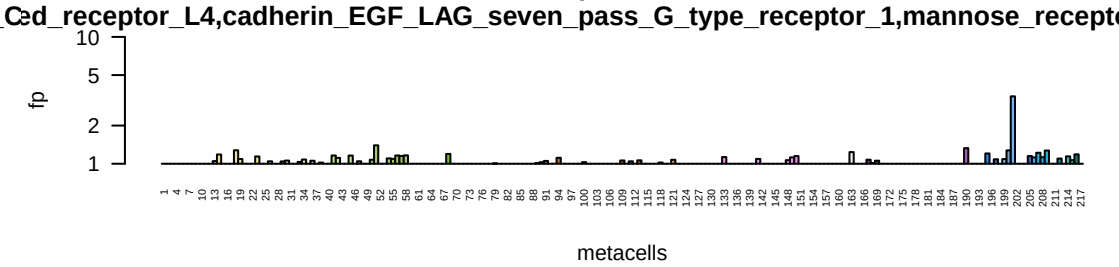

Hhon OG_4092
Hhon_g07037.t1
synaptotagmin_15
fp
10
5
2
1
metacells


HoiH23 OG_4092
HoiH23_PlH23_006623−RA
synaptotagmin_15
fp
10
5
2
1
metacells

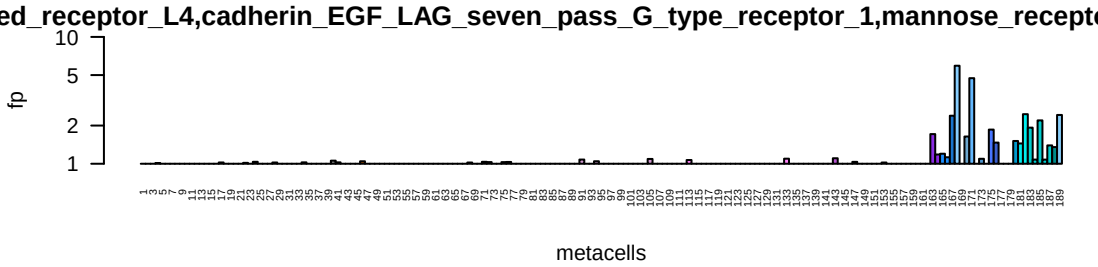
Tadh OG_4305
Tadh_TriadT60119
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



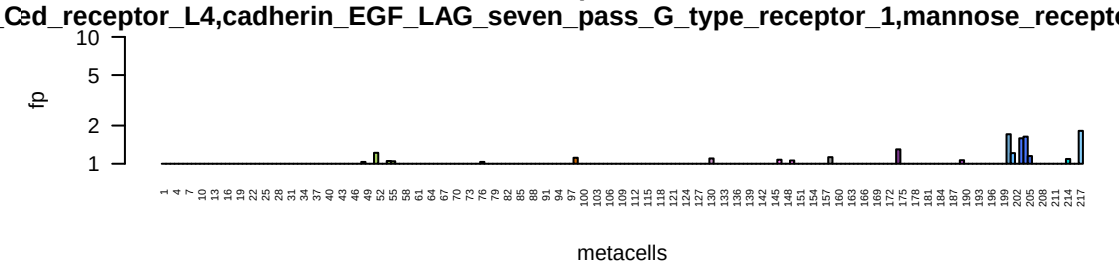
TrH2 OG_4305
TrH2_TrispH2_011070−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



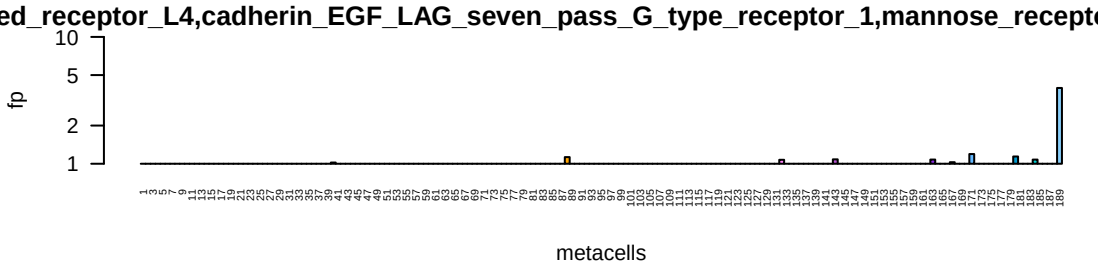
Tadh OG_4305
Tadh_TriadT60118
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



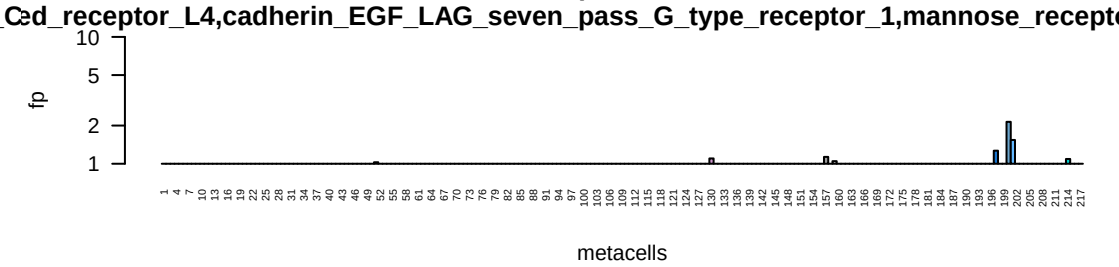
TrH2 OG_4305
TrH2_TrispH2_000685−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



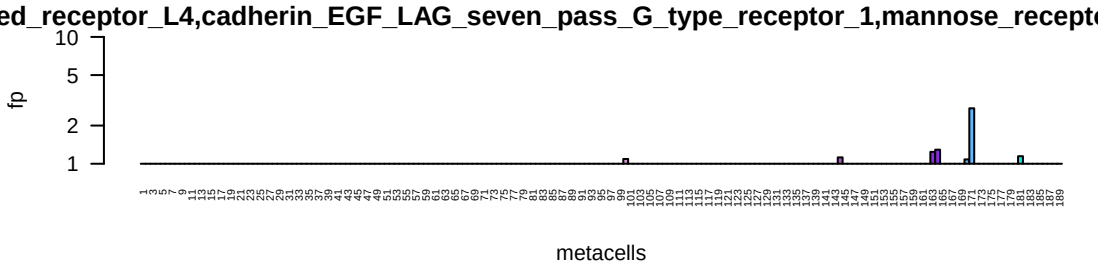
Tadh OG_4305
Tadh_TriadT60121
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



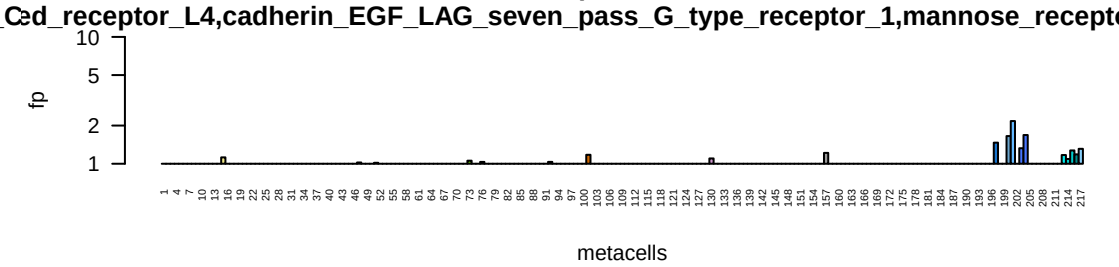
TrH2 OG_4305
TrH2_TrispH2_011793−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



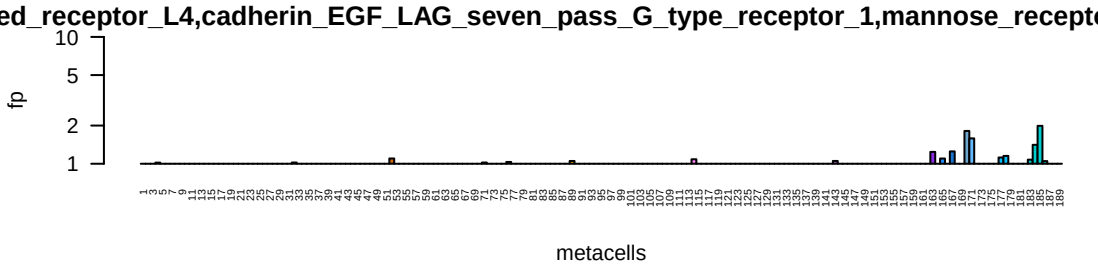
Tadh OG_4305
Tadh_TriadT60117
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



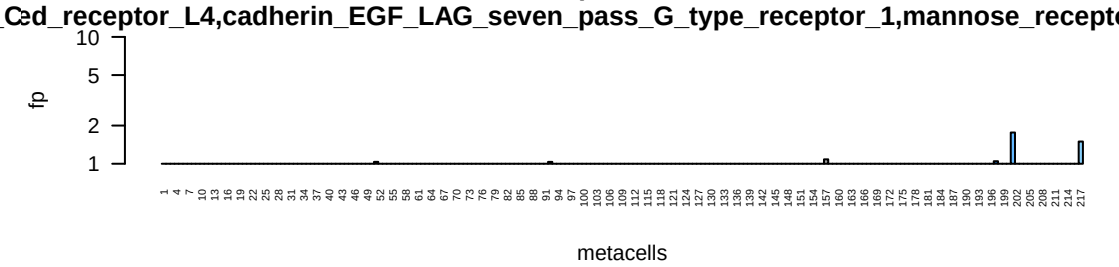
TrH2 OG_4305
TrH2_TrispH2_000683−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



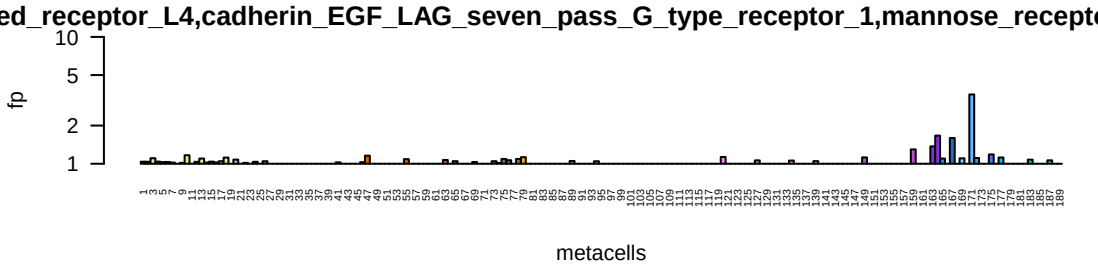
Tadh OG_4305
Tadh_TriadT60120
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



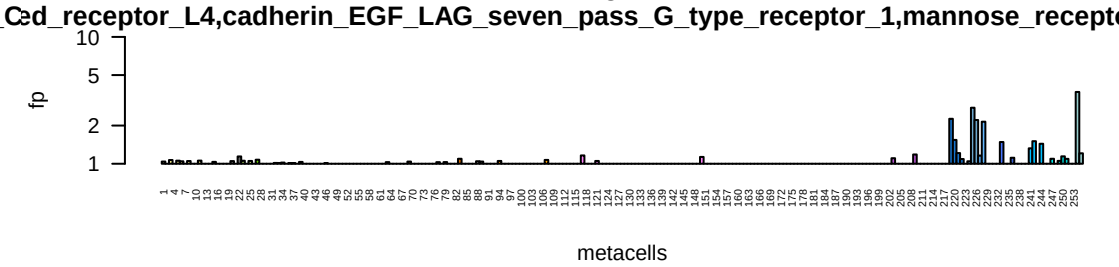
TrH2 OG_4305
TrH2_TrispH2_011071−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



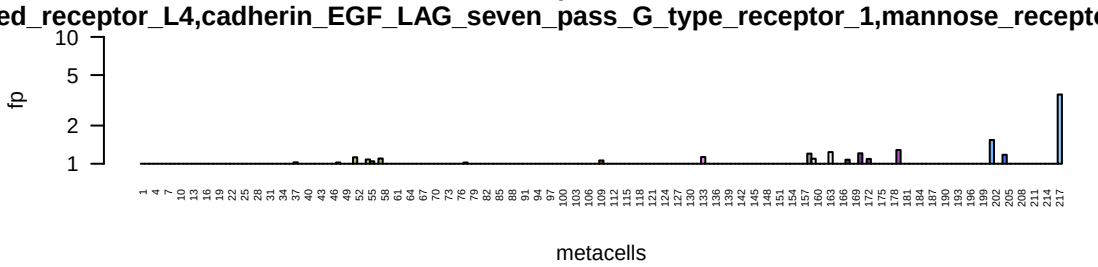
Tadh OG_4305
Tadh_TriadT59938
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



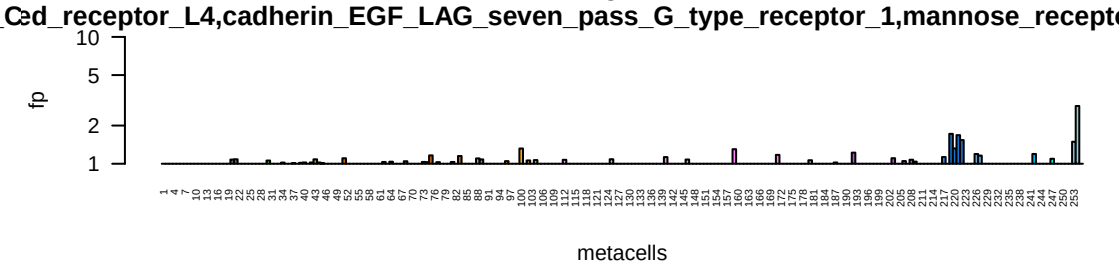
Hhon OG_4305
Hhon_g10982.t1
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



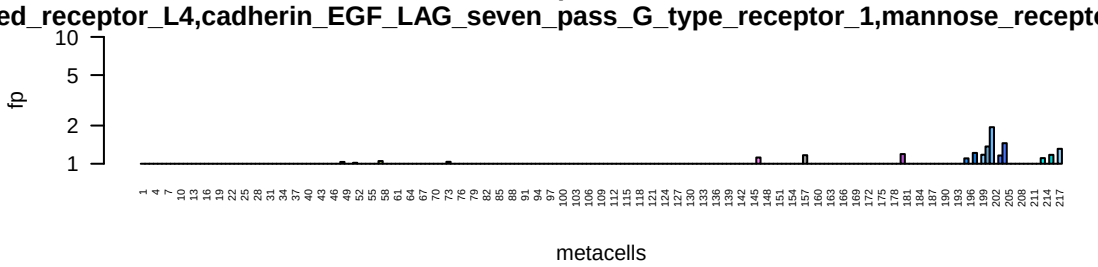
TrH2 OG_4305
TrH2_TrispH2_000684−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C



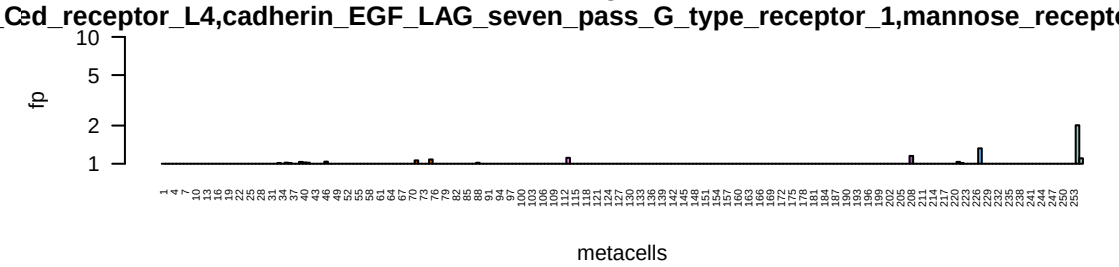
Hhon OG_4305
Hhon_g11834.t1
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

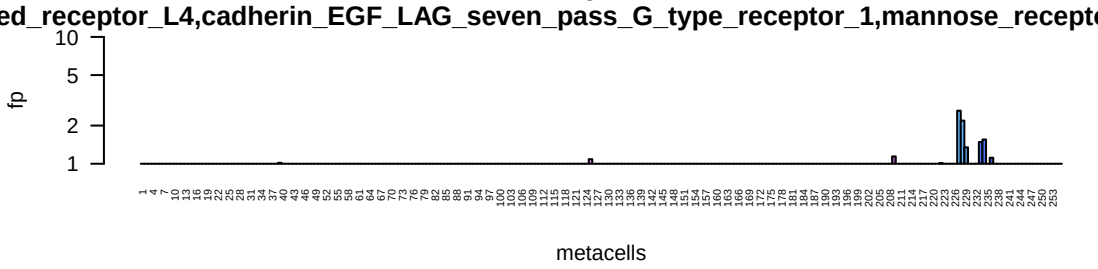


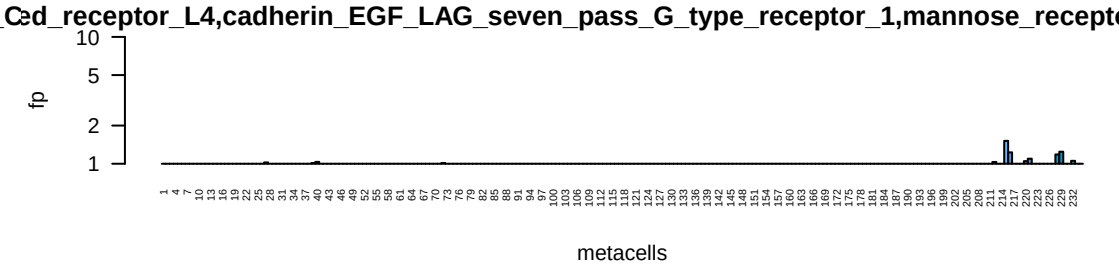
TrH2 OG_4305
TrH2_TrispH2_011671−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

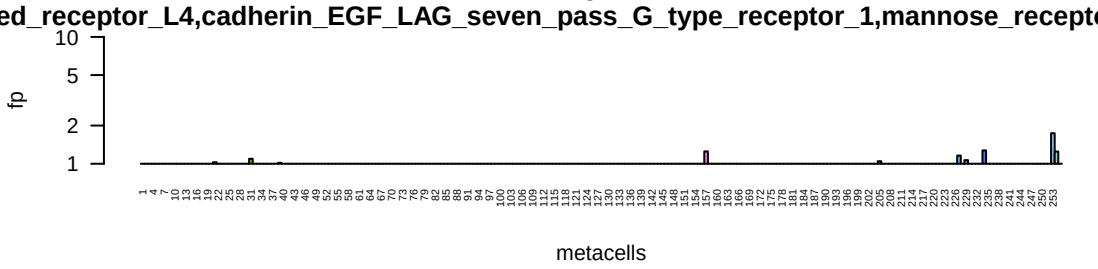


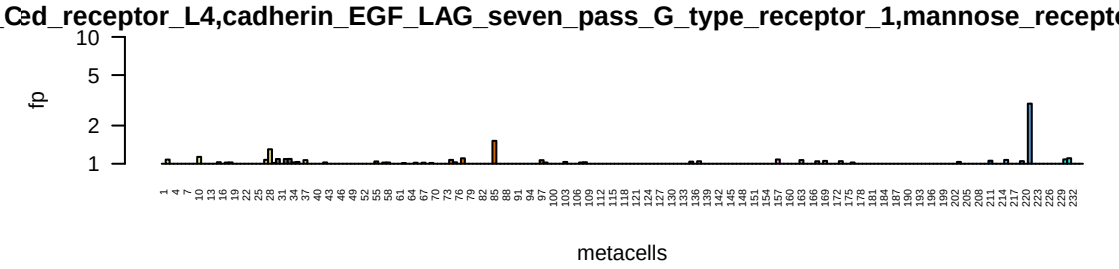
Hhon OG_4305
Hhon_g09074.t1
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C

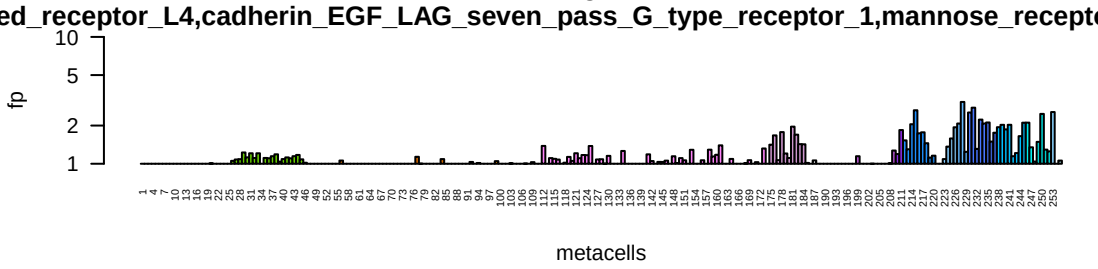



Hhon OG_4305
Hhon_g11448.t1
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C
fp
metacells


HoiH23 OG_4305
HoiH23_PlH23_010761−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C
fp
metacells


Hhon OG_4305
Hhon_g11449.t1
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C
fp
metacells


HoiH23 OG_4305
HoiH23_PlH23_011325−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C
fp
metacells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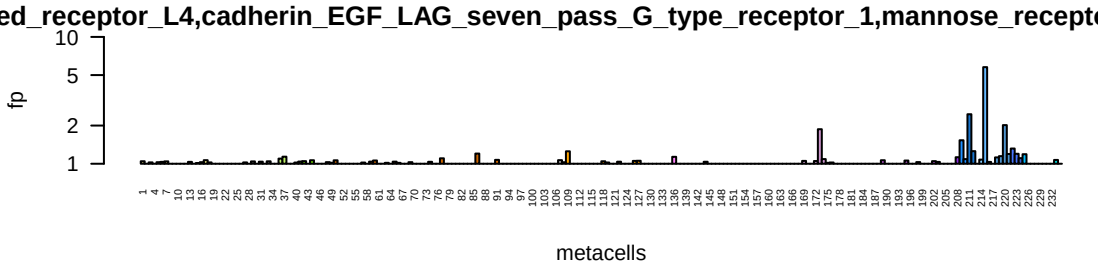
Hhon_g07820.t1
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C
fp
metacells


HoiH23 OG_4305
HoiH23_PlH23_005510−RA
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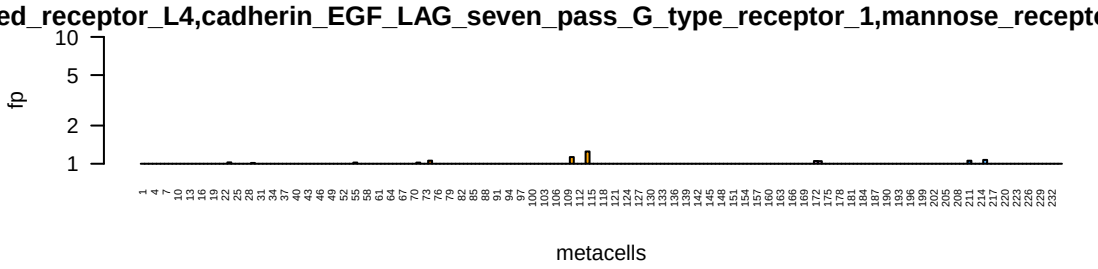
fp
metacells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HoiH23_PlH23_010760−RA
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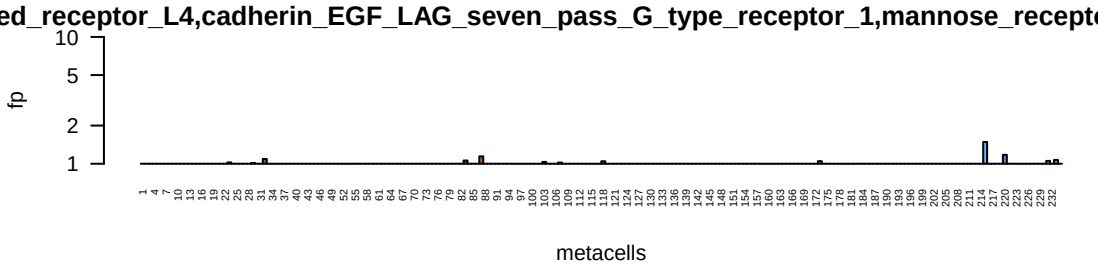
fp
metacells


HoiH23 OG_4305
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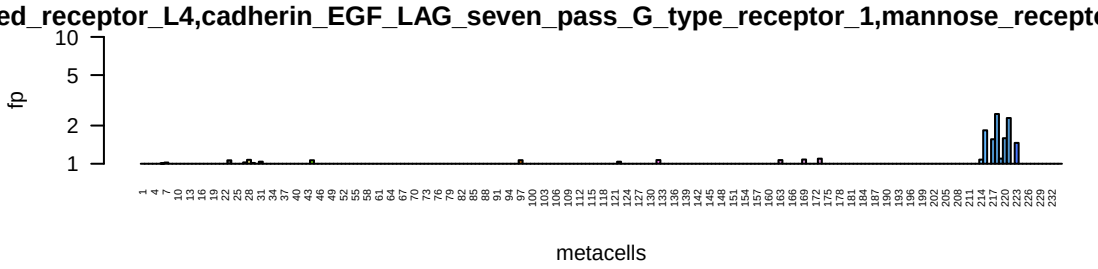
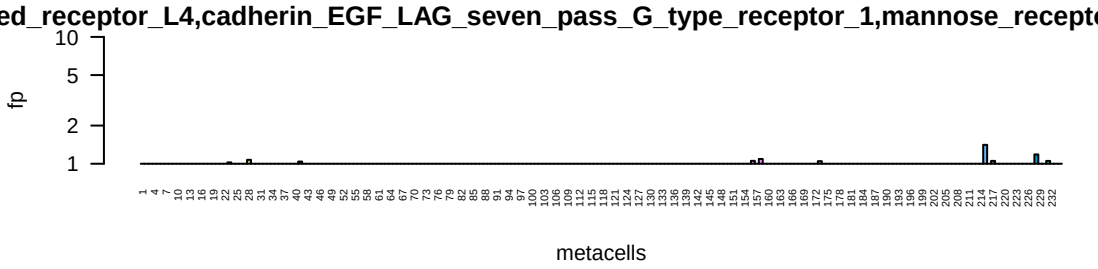
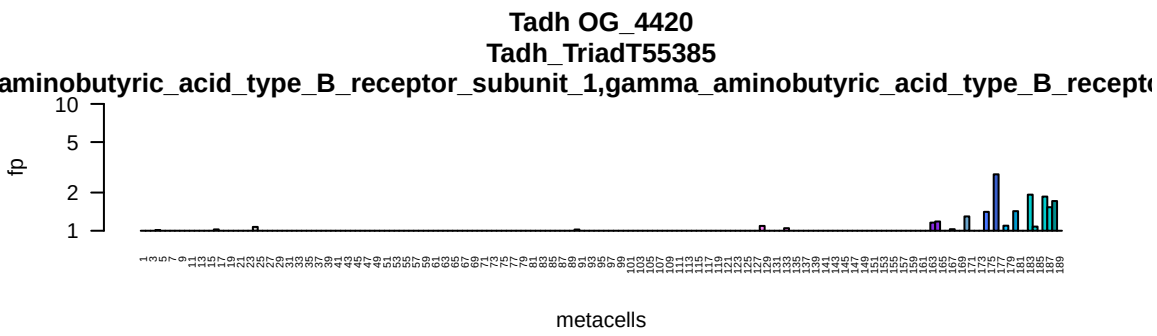
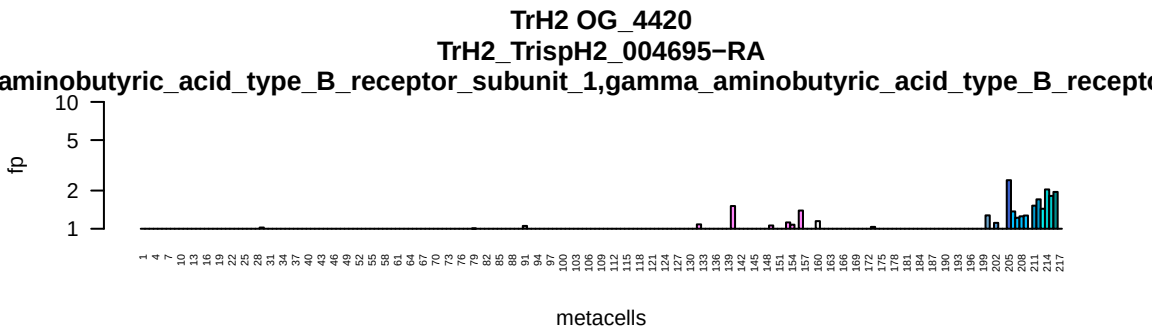
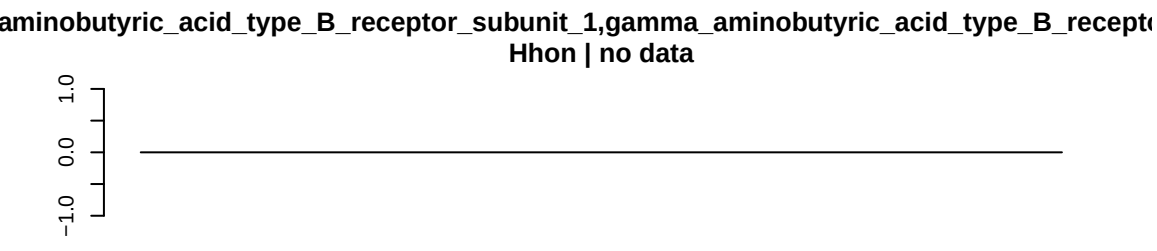
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C
fp
metacells

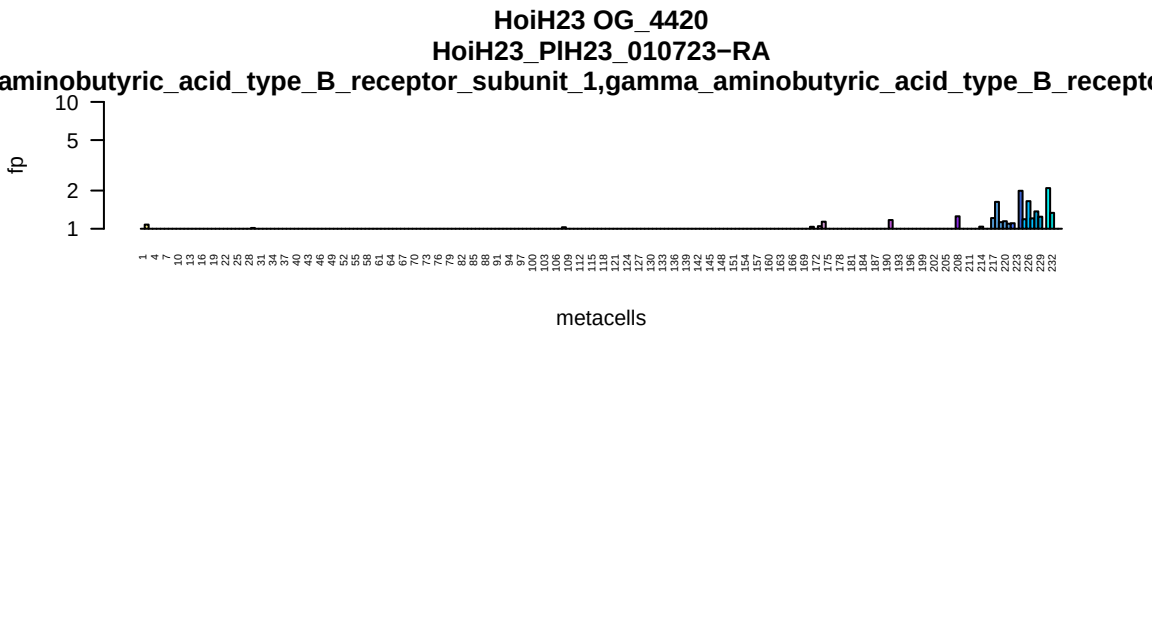

HoiH23 OG_4305
HoiH23_PlH23_009282−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C
fp
metacells

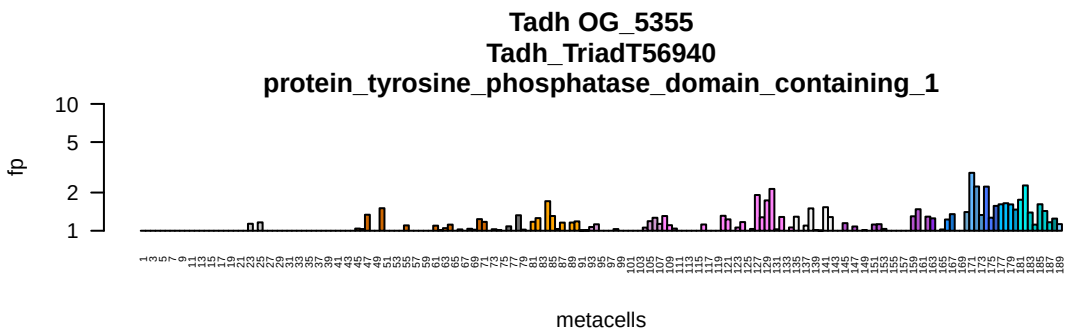

HoiH23 OG_4305
HoiH23_PlH23_011178−RA
ed_receptor_L4,cadherin_EGF_LAG_seven_pass_G_type_receptor_1,mannose_receptor_C
fp
metacells

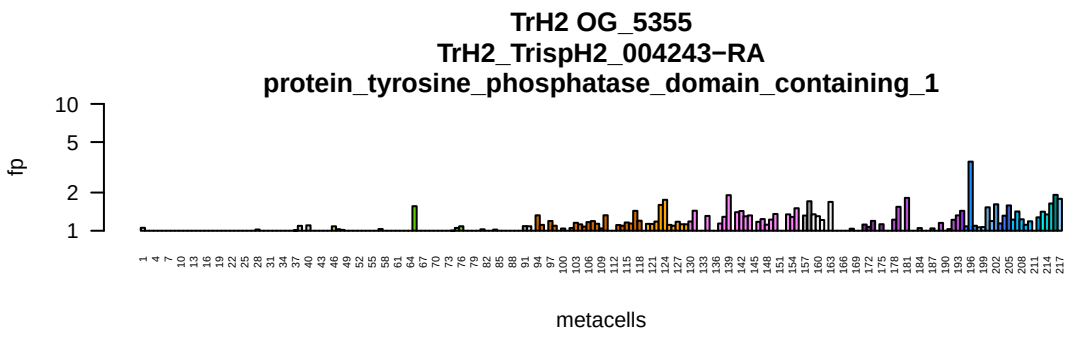

Tadh OG_4420
Tadh_TriadT55385
aminobutyric_acid_type_B_receptor_subunit_1,gamma_aminobutyric_acid_type_B_recepto
fp
metacells

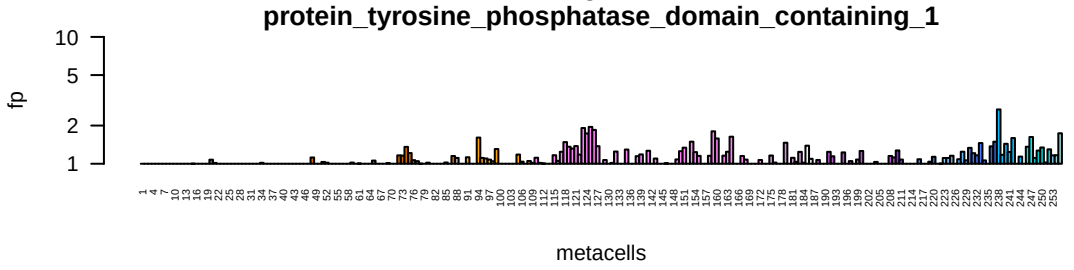

TrH2 OG_4420
TrH2_TrispH2_004695−RA
aminobutyric_acid_type_B_receptor_subunit_1,gamma_aminobutyric_acid_type_B_recepto
fp
metacells

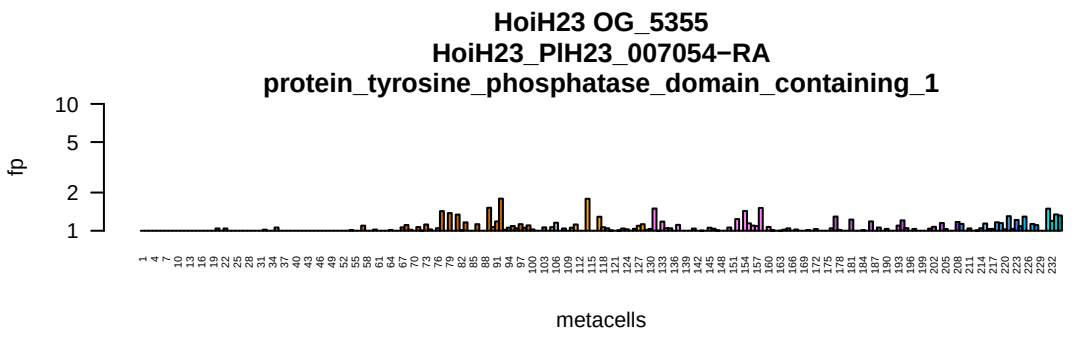

aminobutyric_acid_type_B_receptor_subunit_1,gamma_aminobutyric_acid_type_B_recepto
Hhon | no data

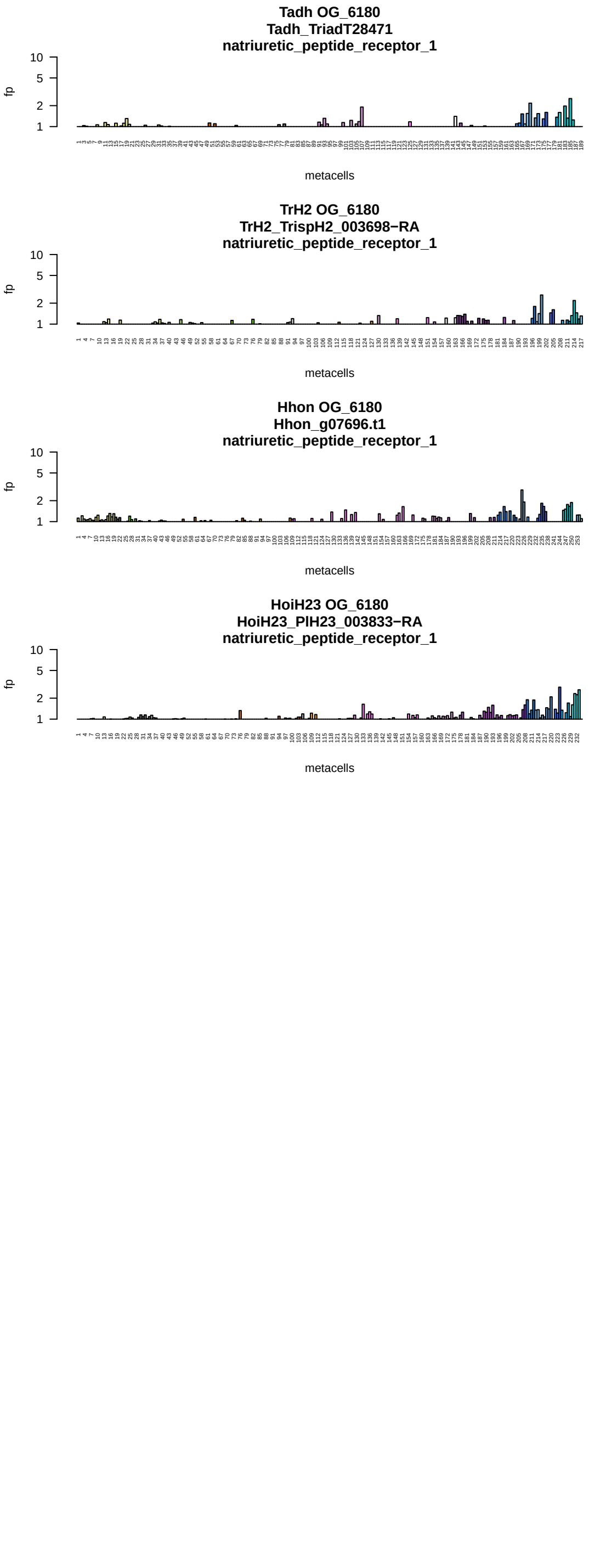

HoiH23 OG_4420
HoiH23_PlH23_010723−RA
aminobutyric_acid_type_B_receptor_subunit_1,gamma_aminobutyric_acid_type_B_recepto
fp
metacells

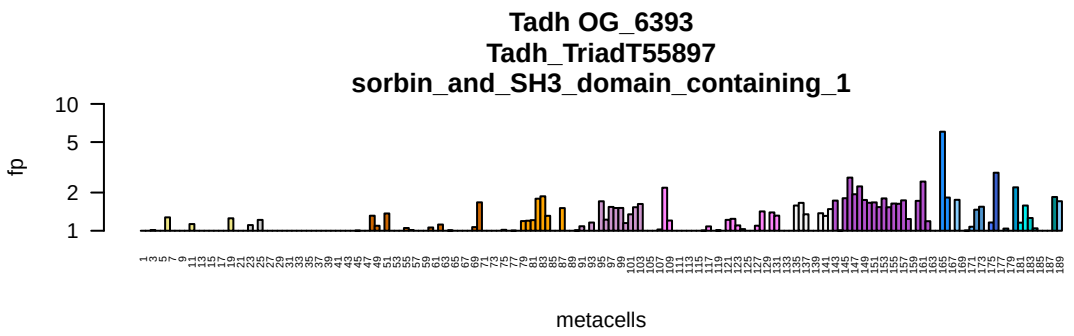

Tadh OG_5355
Tadh_TriadT56940
protein_tyrosine_phosphatase_domain_containing_1
fp
metacells

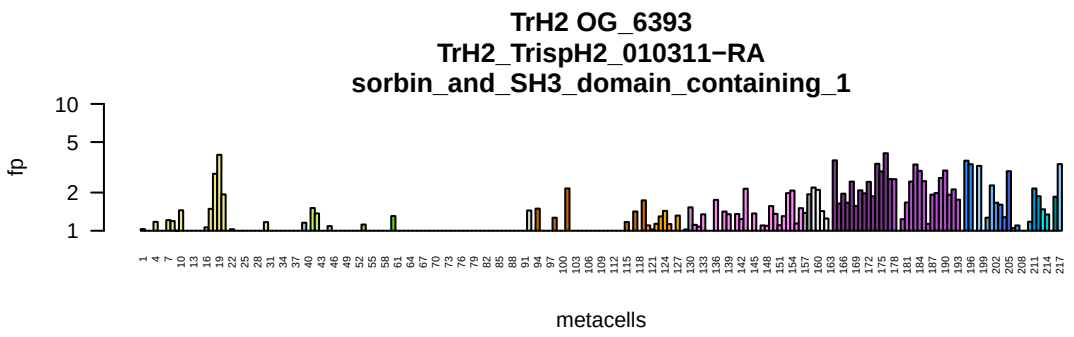

TrH2 OG_5355
TrH2_TrispH2_004243−RA
protein_tyrosine_phosphatase_domain_containing_1
fp
metacells

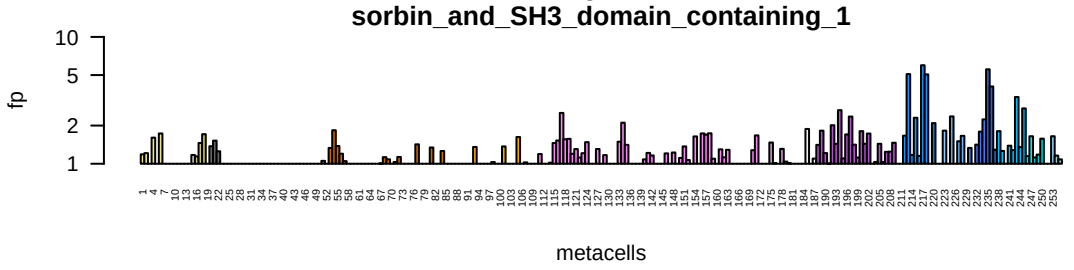

Hhon OG_5355
Hhon_g05675.t1
protein_tyrosine_phosphatase_domain_containing_1
fp
metacells

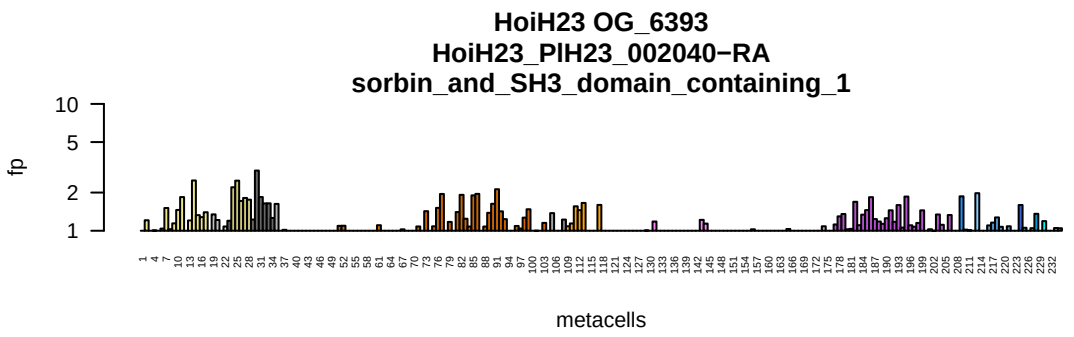

HoiH23 OG_5355
HoiH23_PlH23_007054−RA
protein_tyrosine_phosphatase_domain_containing_1
fp
metacells

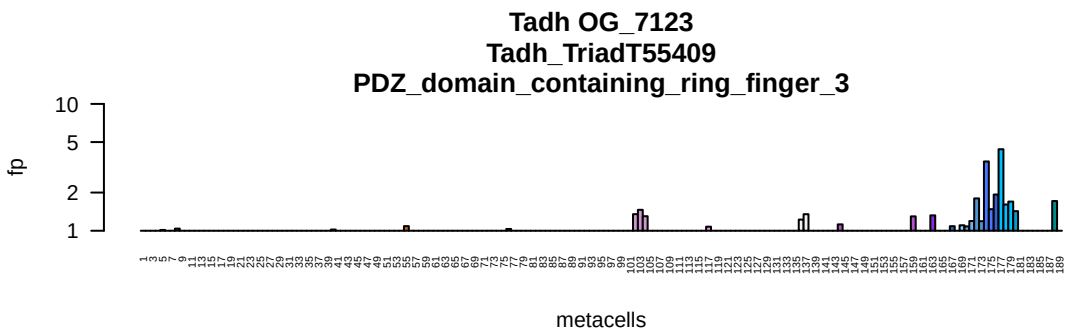

Tadh OG_6180
Tadh_TriadT28471
natriuretic_peptide_receptor_1
fp
metacells
TrH2 OG_6180
TrH2_TrispH2_003698−RA
natriuretic_peptide_receptor_1
fp
metacells
Hhon OG_6180
Hhon_g07696.t1
natriuretic_peptide_receptor_1
fp
metacells
HoiH23 OG_6180
HoiH23_PlH23_003833−RA
natriuretic_peptide_receptor_1
fp
metacells

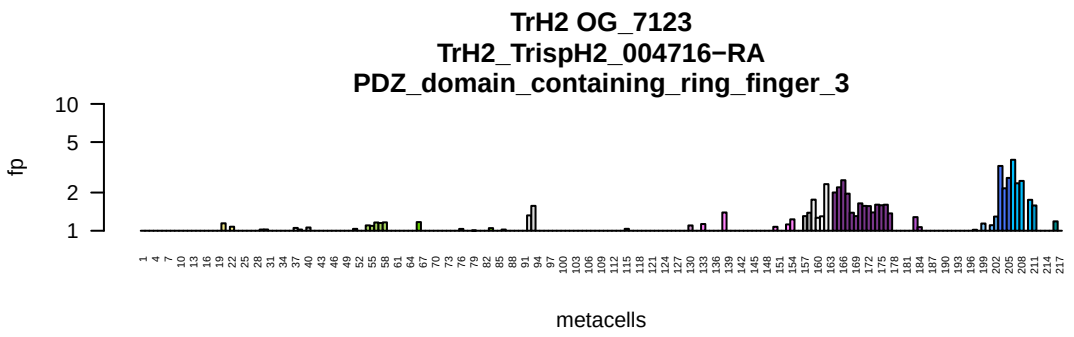

Tadh OG_6393
Tadh_TriadT55897
sorbin_and_SH3_domain_containing_1
fp
metacells

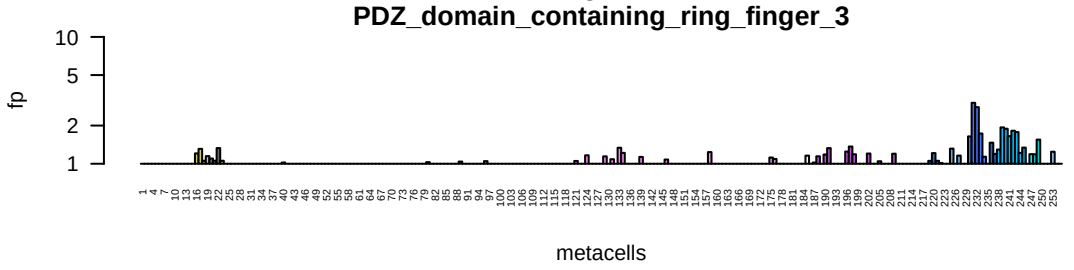

TrH2 OG_6393
TrH2_TrispH2_010311−RA
sorbin_and_SH3_domain_containing_1
fp
metacells

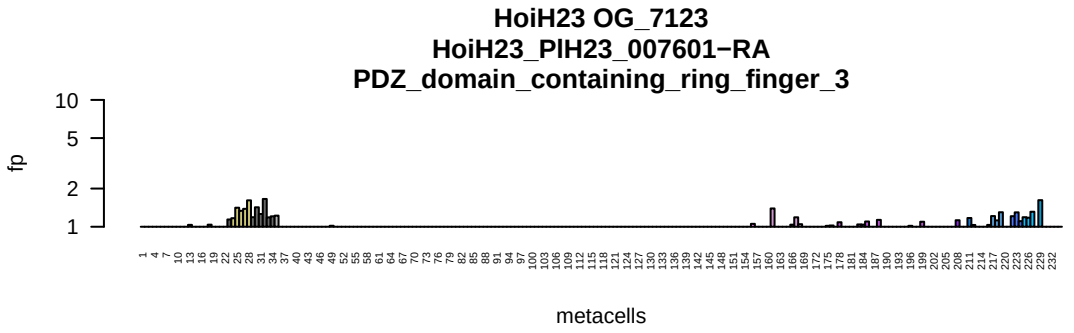

Hhon OG_6393
Hhon_g03284.t1
sorbin_and_SH3_domain_containing_1
fp
metacells

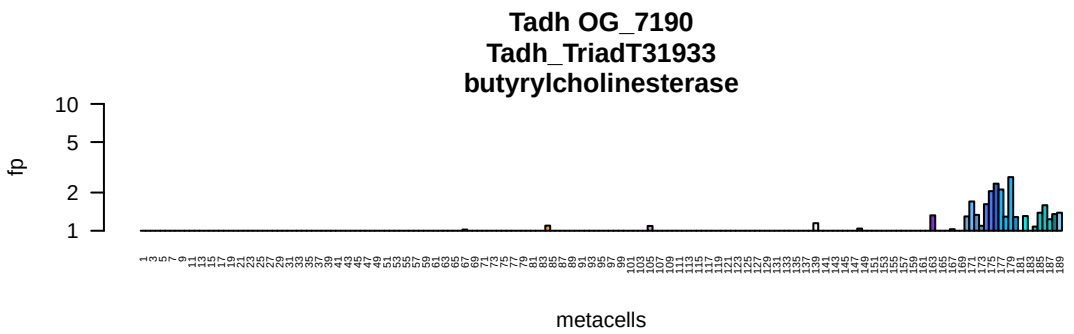

HoiH23 OG_6393
HoiH23_PlH23_002040−RA
sorbin_and_SH3_domain_containing_1
fp
metacells

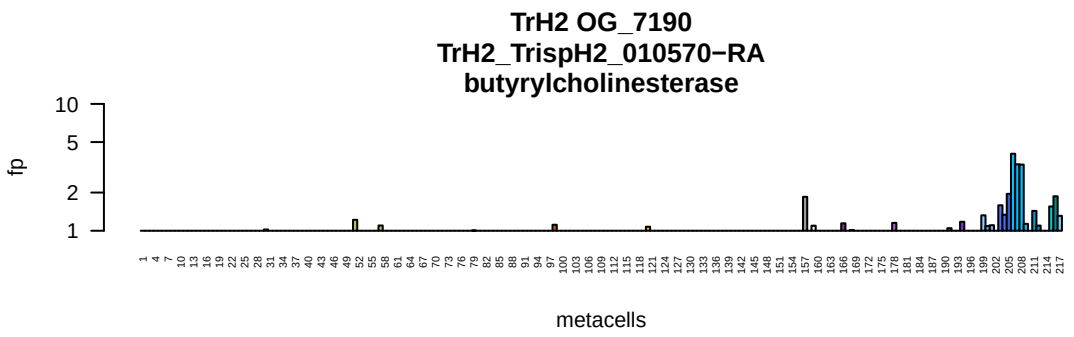

Tadh OG_7123
Tadh_TriadT55409
PDZ_domain_containing_ring_finger_3
fp
metacells

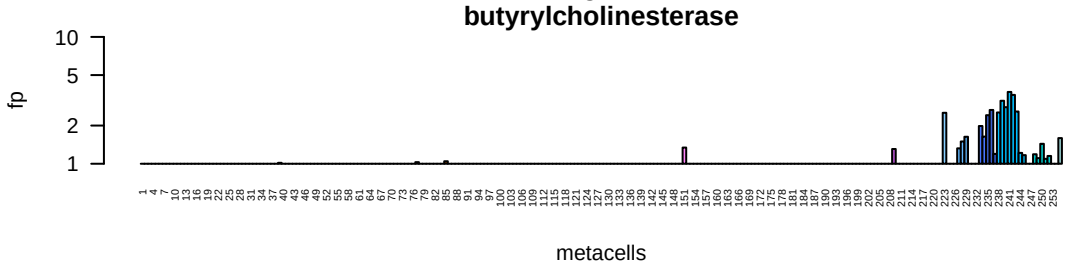

TrH2 OG_7123
TrH2_TrispH2_004716−RA
PDZ_domain_containing_ring_finger_3
fp
metacells

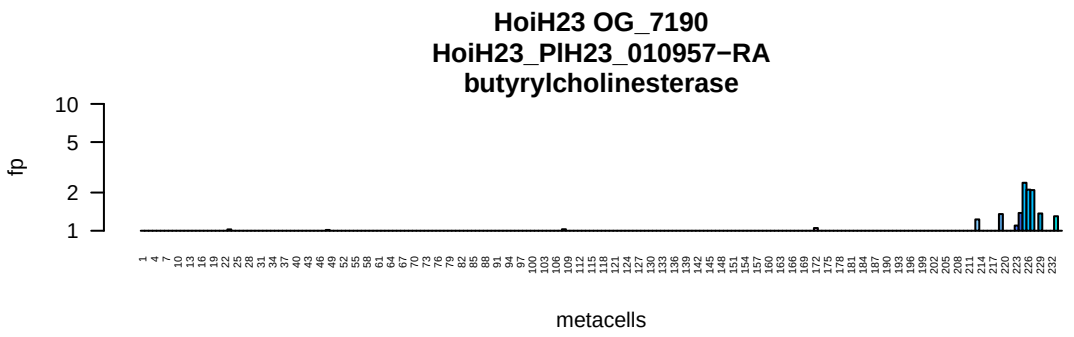

Hhon OG_7123
Hhon_g09054.t1
PDZ_domain_containing_ring_finger_3
fp
metacells


HoiH23 OG_7123
HoiH23_PlH23_007601−RA
PDZ_domain_containing_ring_finger_3
fp
metacells


Tadh OG_7190
Tadh_TriadT31933
butyrylcholinesterase
fp
metacells


TrH2 OG_7190
TrH2_TrispH2_010570−RA
butyrylcholinesterase
fp
metacells


Hhon OG_7190
Hhon_g01153.t1
butyrylcholinesterase
fp
metacells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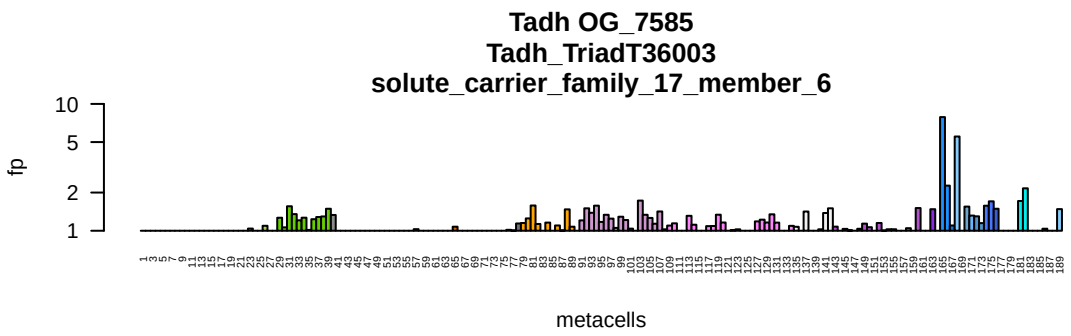
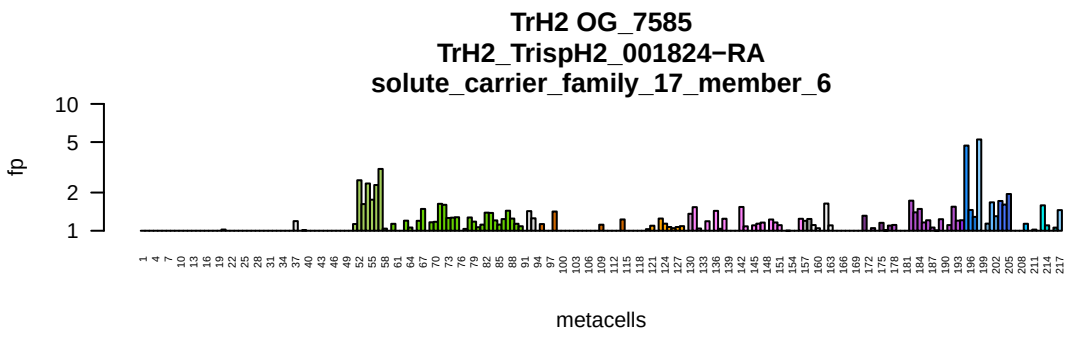
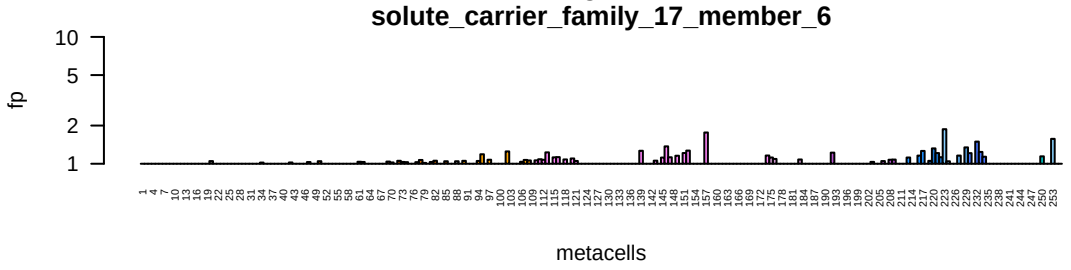
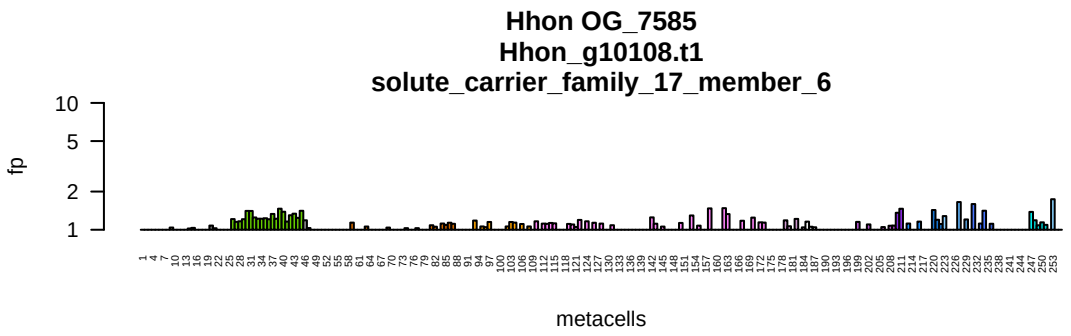
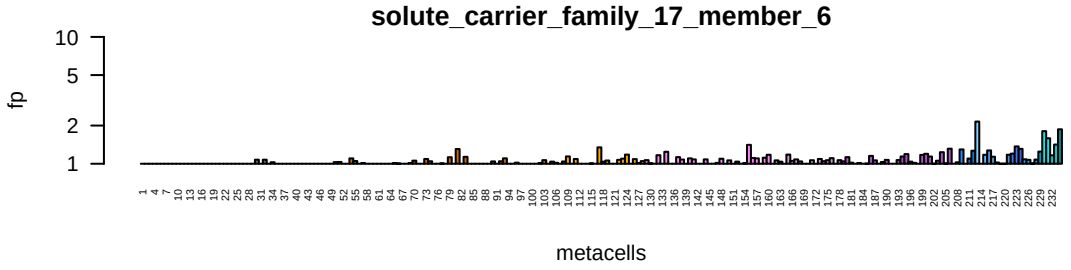
HoiH23_PlH23_010957−RA
butyrylcholinesterase
fp
metacells

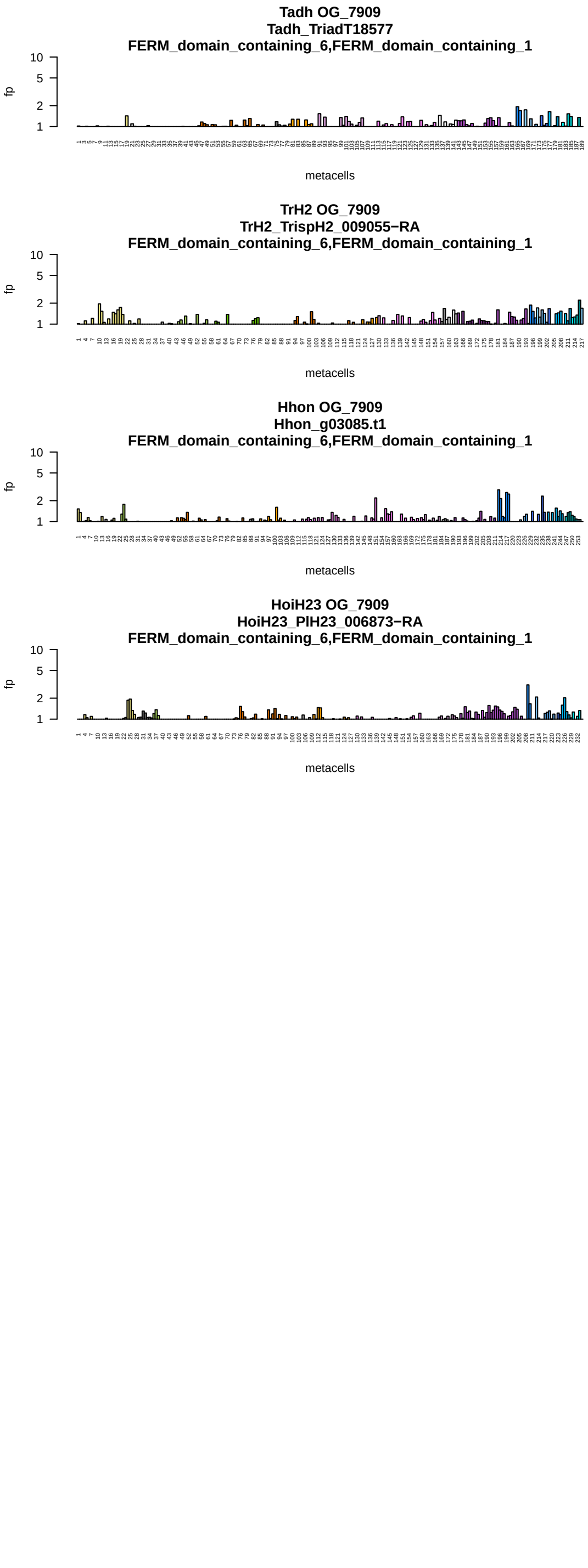

Tadh OG_7585
Tadh_TriadT36003
solute_carrier_family_17_member_6
fp
10
5
2
1
metacells

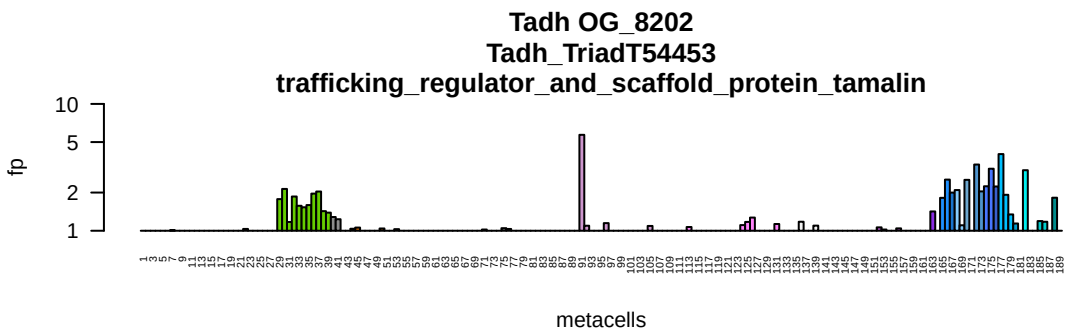

TrH2 OG_7585
TrH2_TrispH2_001824−RA
solute_carrier_family_17_member_6
fp
10
5
2
1
metacells

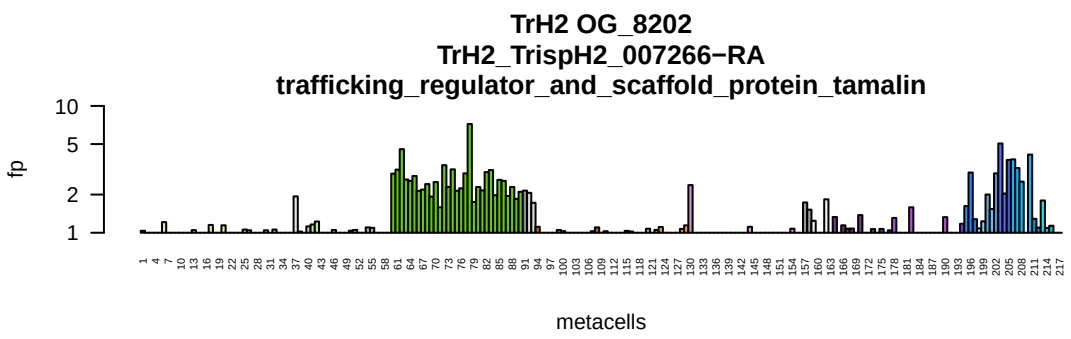

Hhon OG_7585
Hhon_g11731.t1
solute_carrier_family_17_member_6
fp
10
5
2
1
metacells

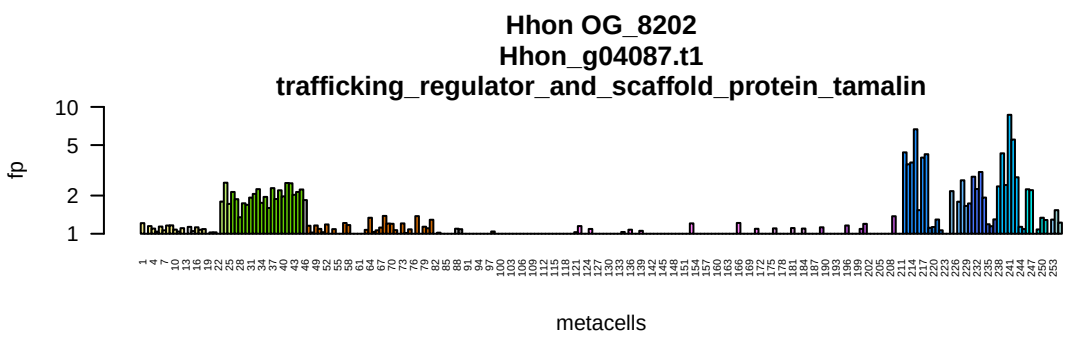

Hhon OG_7585
Hhon_g10108.t1
solute_carrier_family_17_member_6
fp
10
5
2
1
metacells

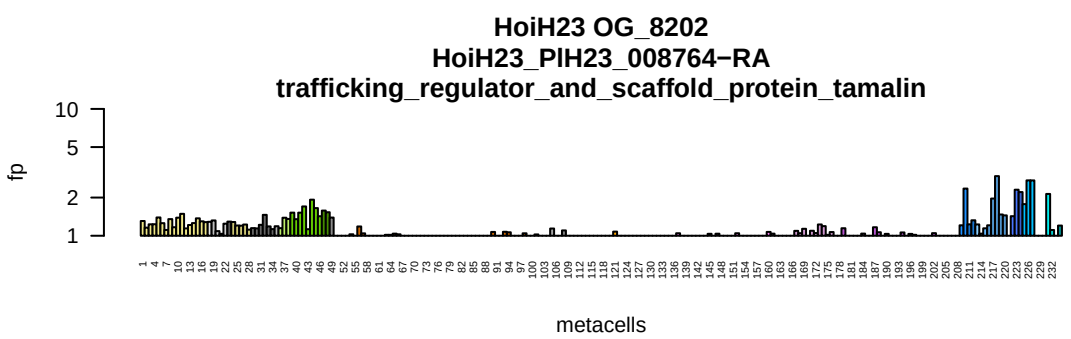

HoiH23 OG_7585
HoiH23_PlH23_000319−RA
solute_carrier_family_17_member_6
fp
10
5
2
1
metacells


Tadh OG_7909
Tadh_TriadT18577
FERM_domain_containing_6,FERM_domain_containing_1
fp
metacells
TrH2 OG_7909
TrH2_TrispH2_009055−RA
FERM_domain_containing_6,FERM_domain_containing_1
fp
metacells
Hhon OG_7909
Hhon_g03085.t1
FERM_domain_containing_6,FERM_domain_containing_1
fp
metacells
HoiH23 OG_7909
HoiH23_PlH23_006873−RA
FERM_domain_containing_6,FERM_domain_containing_1
fp
metacells

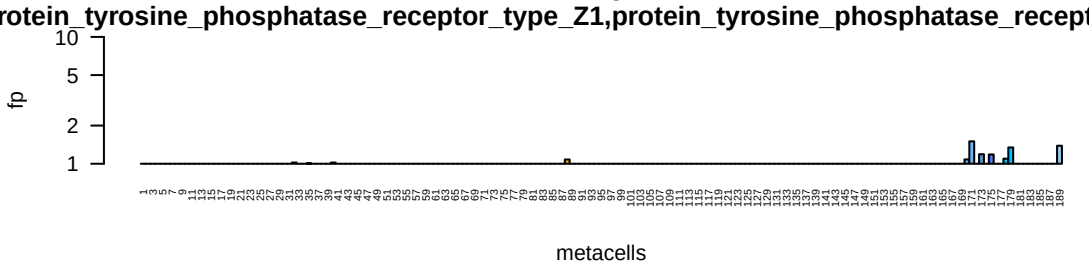

Tadh OG_8202
Tadh_TriadT54453
trafficking_regulator_and_scaffold_protein_tamalin
fp
metacells


TrH2 OG_8202
TrH2_TrispH2_007266−RA
trafficking_regulator_and_scaffold_protein_tamalin
fp
metacells

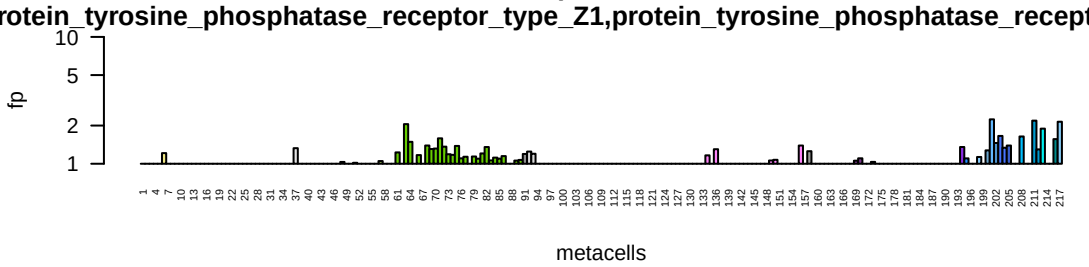

Hhon OG_8202
Hhon_g04087.t1
trafficking_regulator_and_scaffold_protein_tamalin
fp
metacells


HoiH23 OG_8202
HoiH23_PlH23_008764−RA
trafficking_regulator_and_scaffold_protein_tamalin
fp
metacells

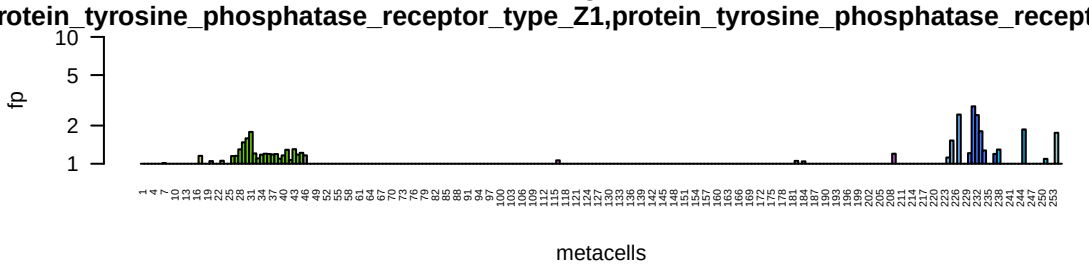
Tadh OG_8999
Tadh_wf_g1014.t1
rotein_tyrosine_phosphatase_receptor_type_Z1,protein_tyrosine_phosphatase_receptor_ty



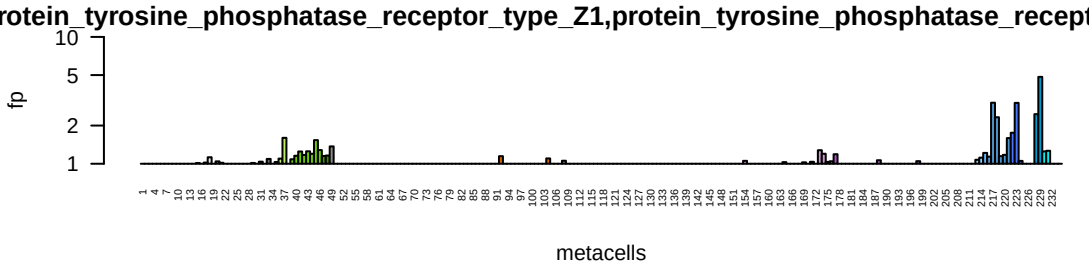
TrH2 OG_8999
TrH2_TrispH2_001036−RA
rotein_tyrosine_phosphatase_receptor_type_Z1,protein_tyrosine_phosphatase_receptor_ty

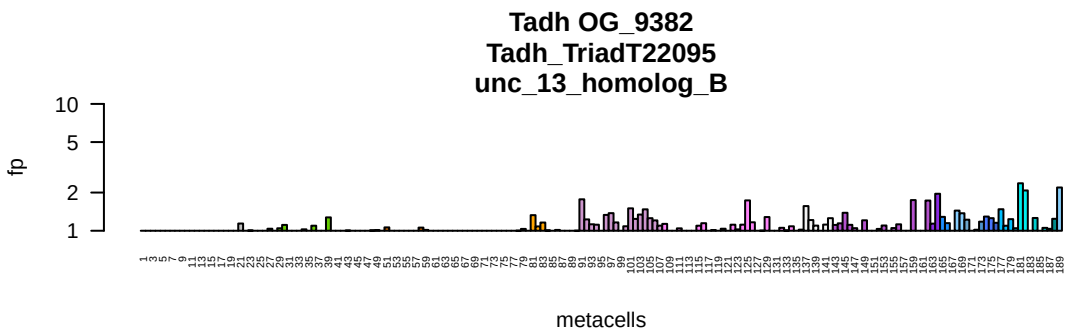


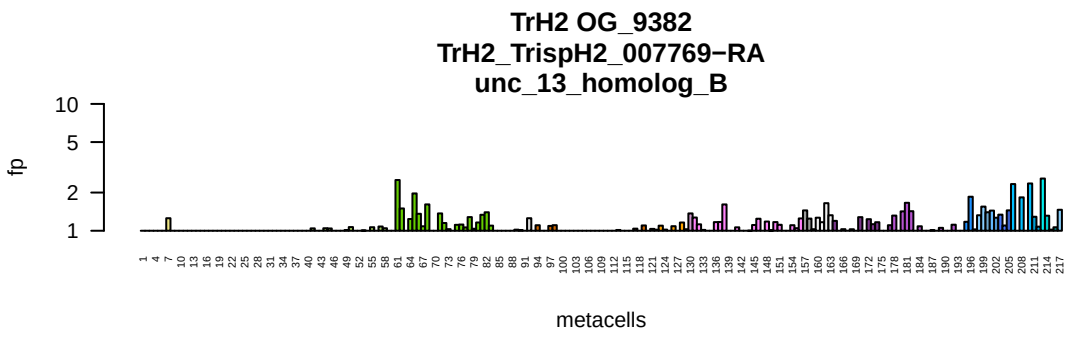
Hhon OG_8999
Hhon_g02885.t1
rotein_tyrosine_phosphatase_receptor_type_Z1,protein_tyrosine_phosphatase_receptor_ty



HoiH23 OG_8999
HoiH23_PlH23_008288−RA
rotein_tyrosine_phosphatase_receptor_type_Z1,protein_tyrosine_phosphatase_receptor_ty




Tadh OG_9382
Tadh_TriadT22095
unc_13_homolog_B
fp
metacells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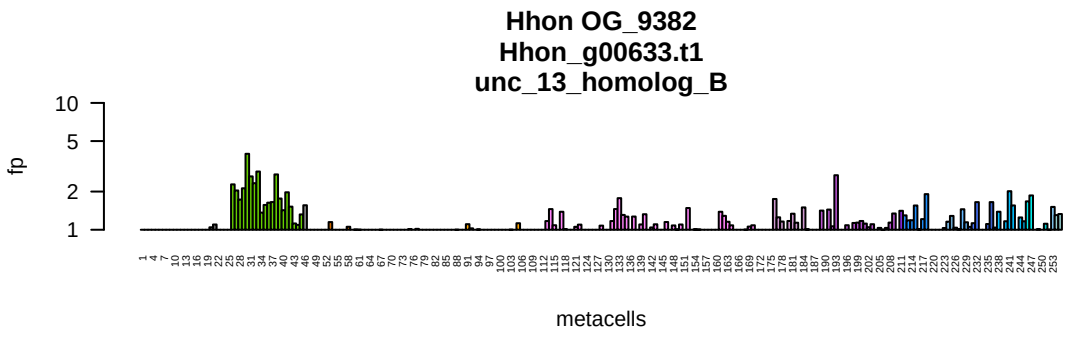
metacells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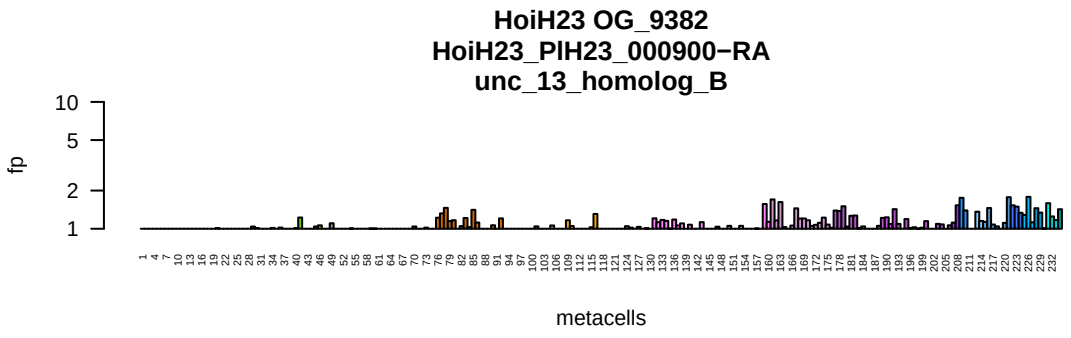
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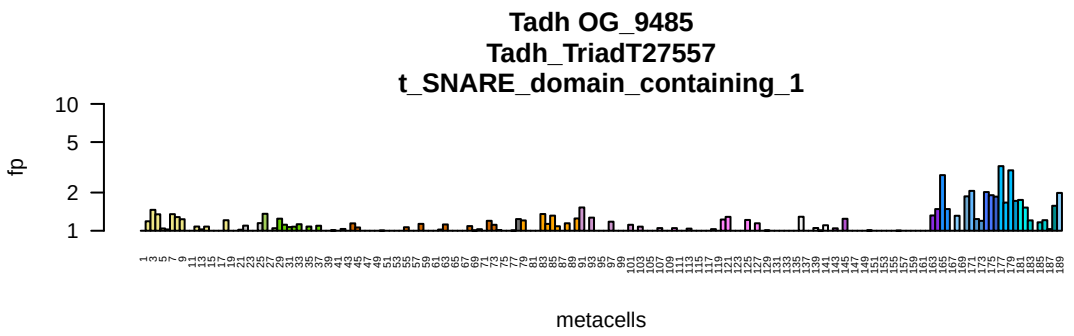
metacells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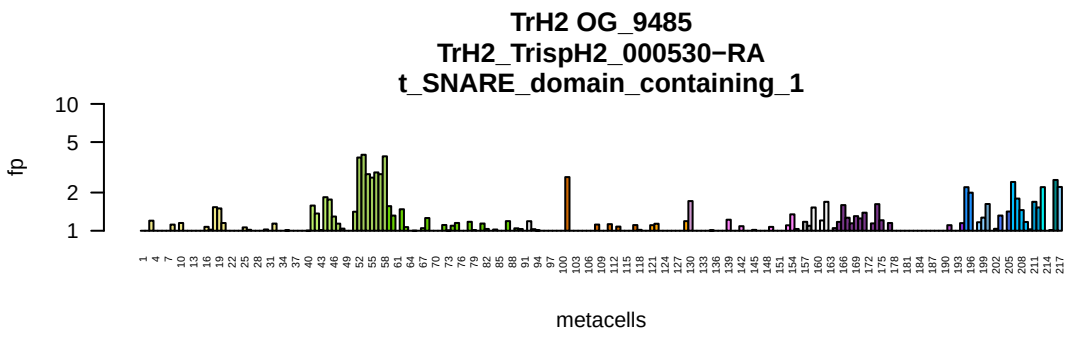
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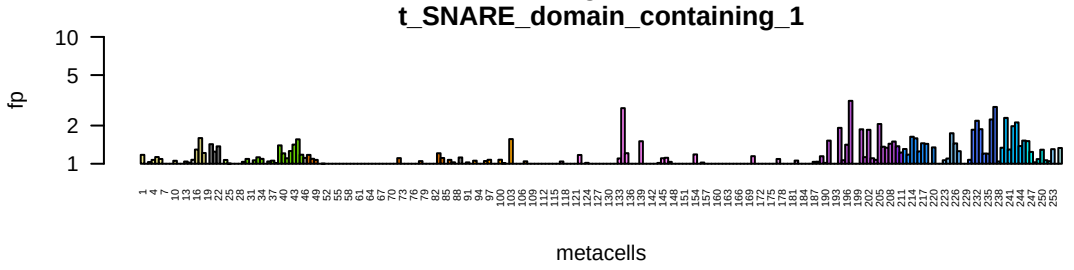
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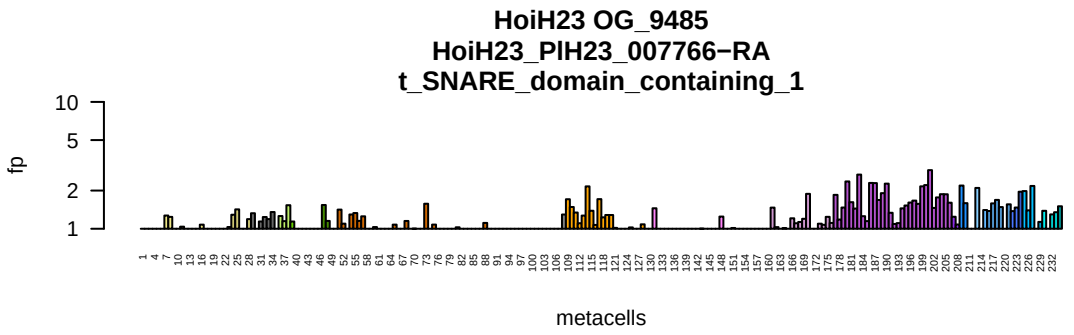
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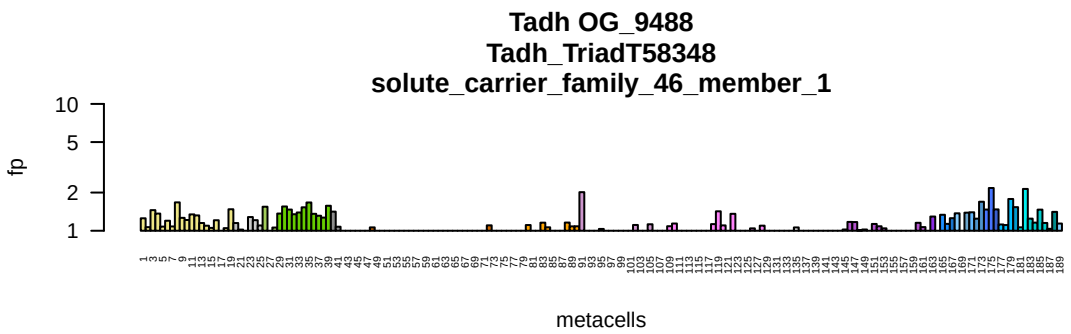
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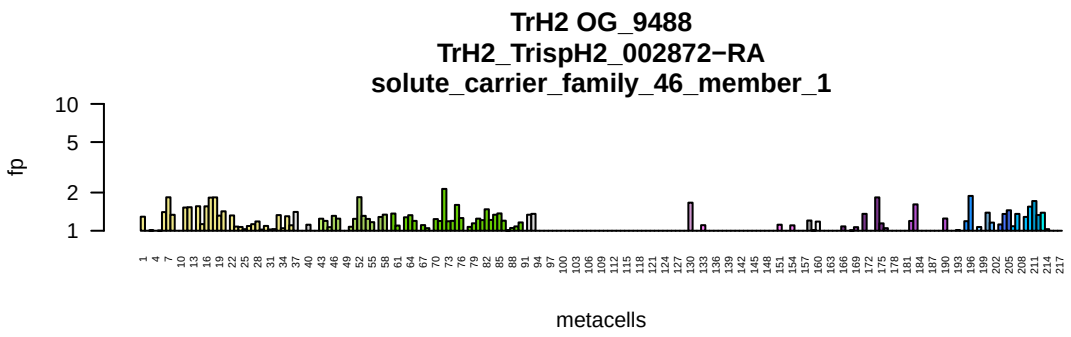
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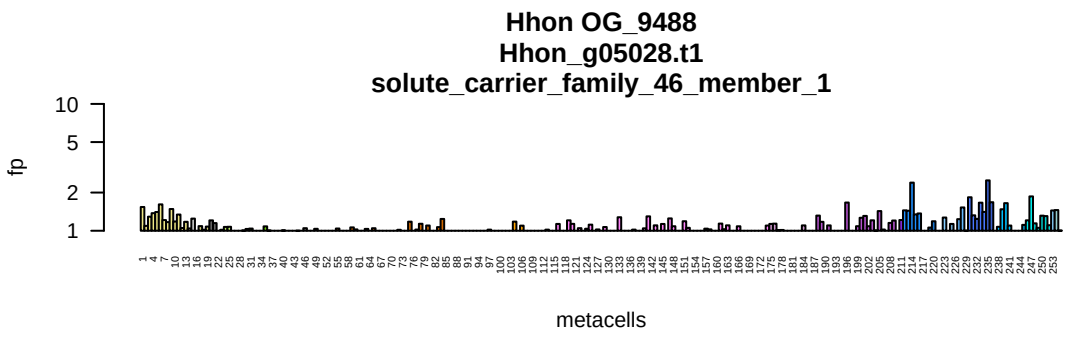
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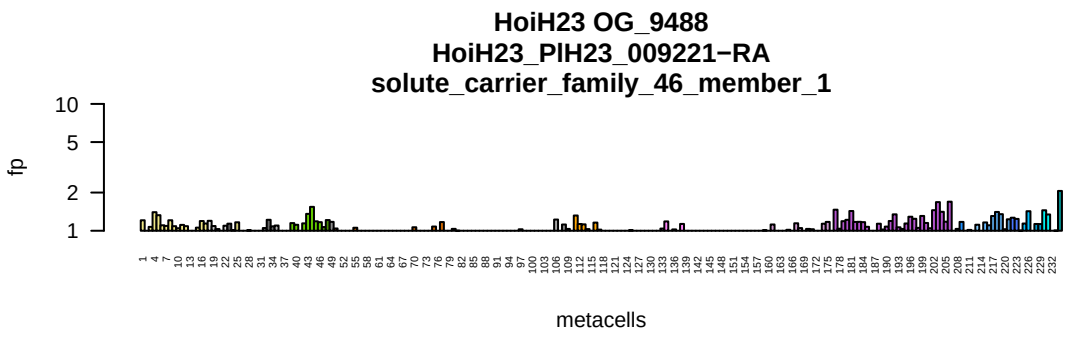
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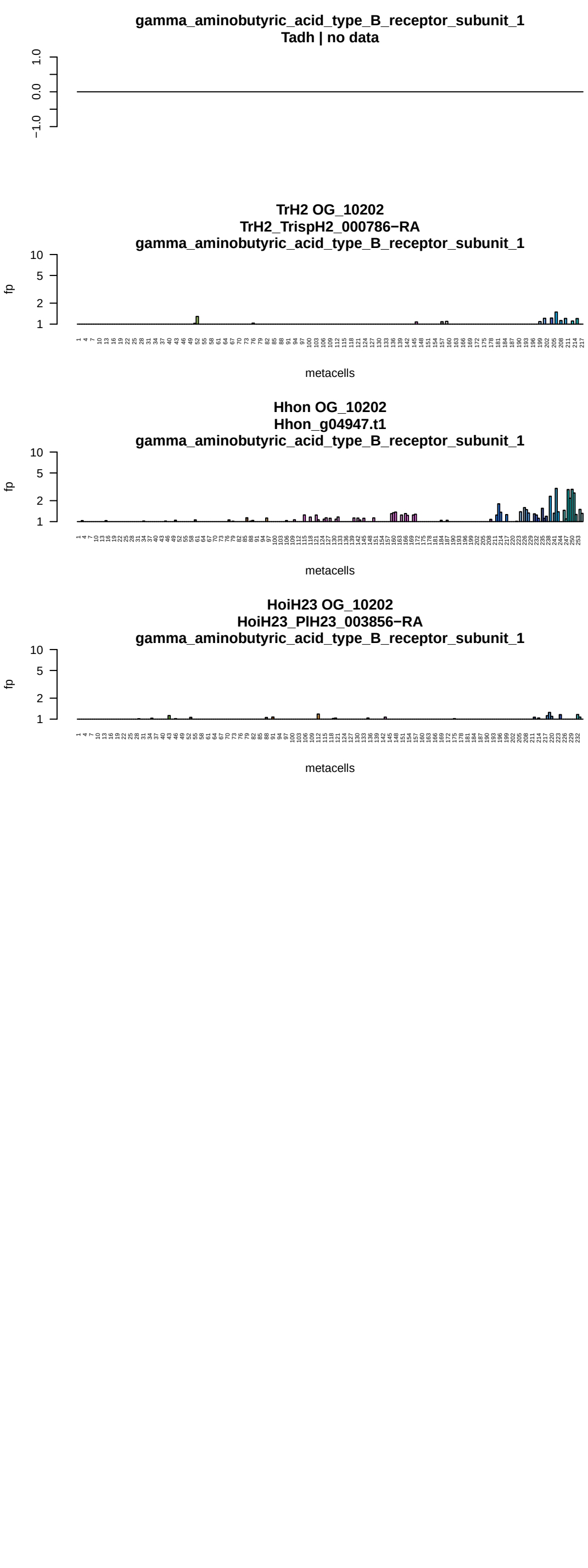
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0.0
−1.0
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fp
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2
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metacells
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fp
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2
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metacells