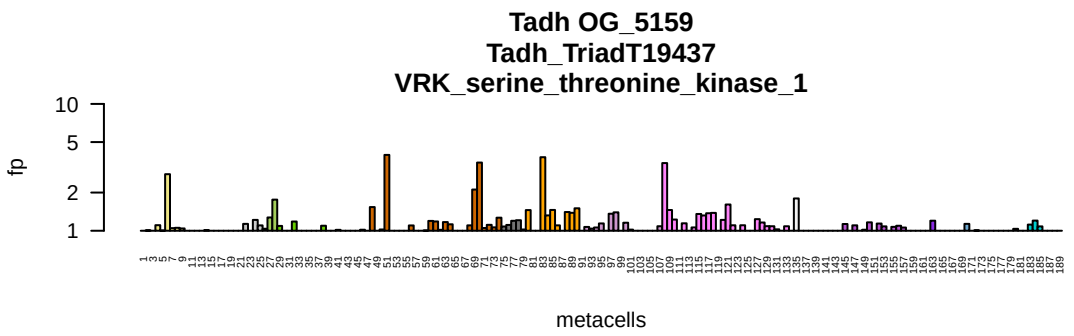
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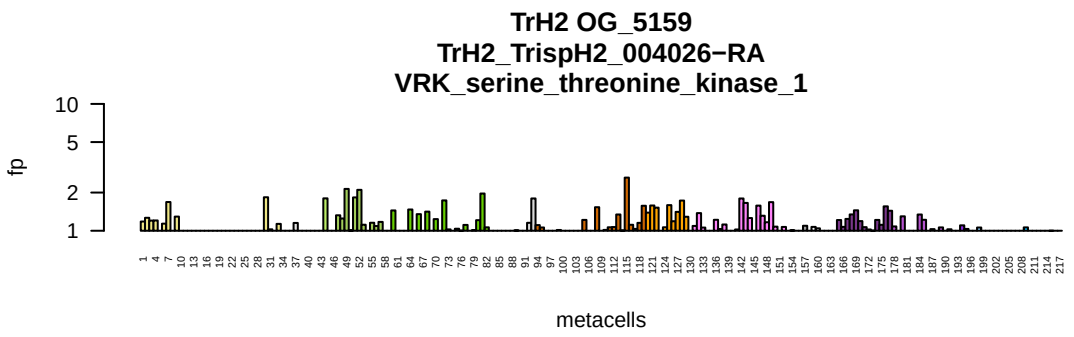
VRK_serine_threonine_kinase_1
fp
metacells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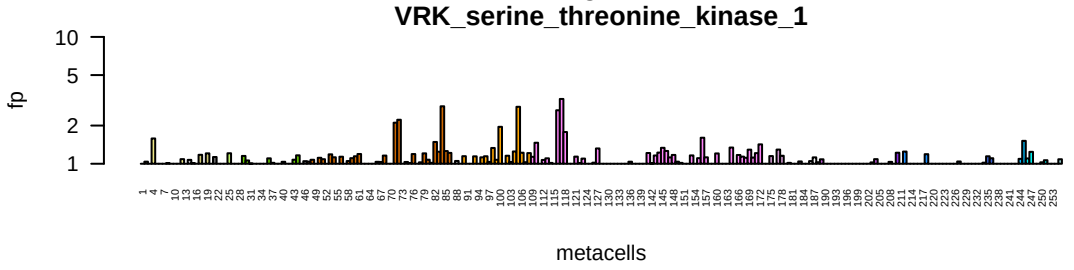
VRK_serine_threonine_kinase_1
fp
metacells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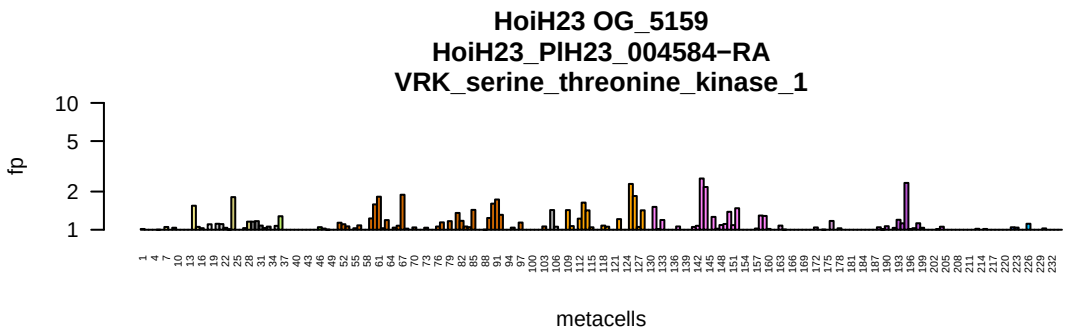
VRK_serine_threonine_kinase_1
fp
metacells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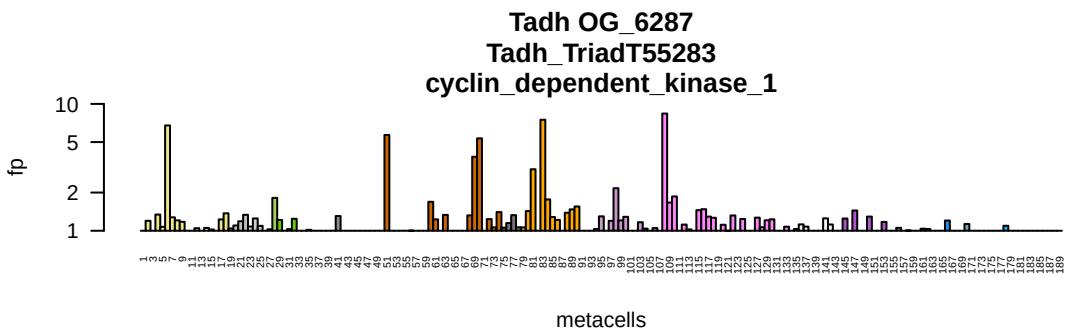
HoiH23_PlH23_004584−RA
VRK_serine_threonine_kinase_1
fp
metacells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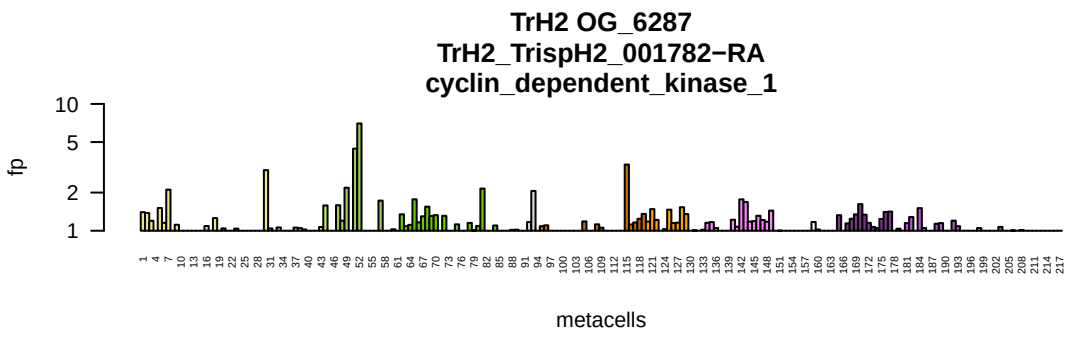
cyclin_dependent_kinase_1
fp
metacells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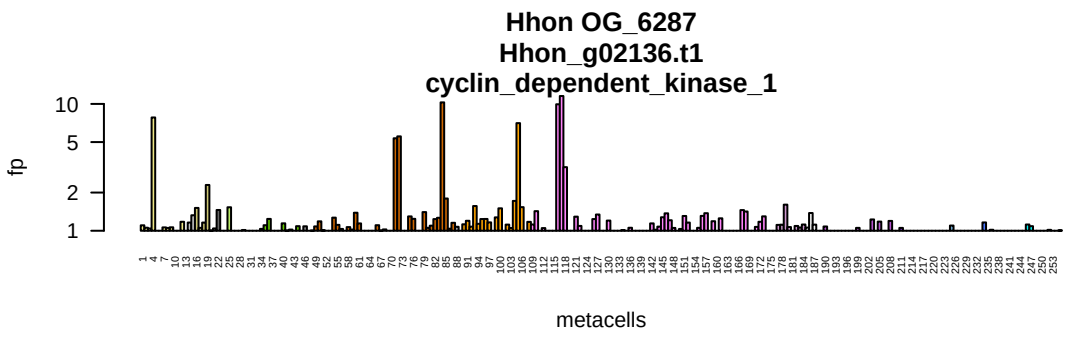
TrH2_TrispH2_001782−RA
cyclin_dependent_kinase_1
fp
metacells


Hhon OG_6287
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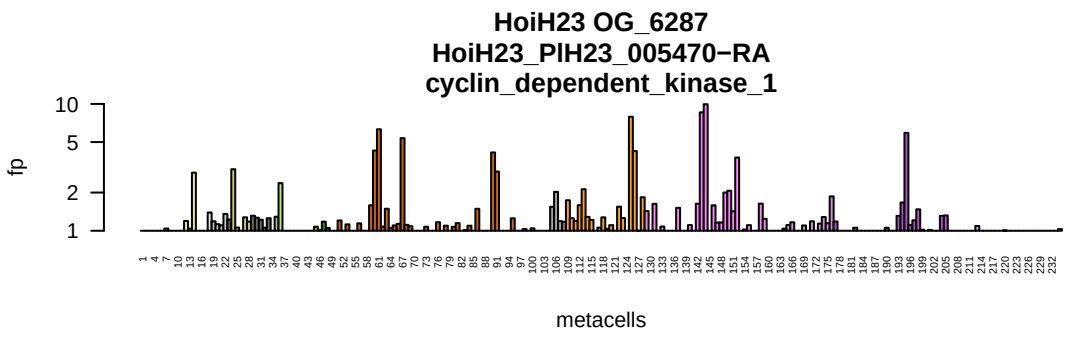
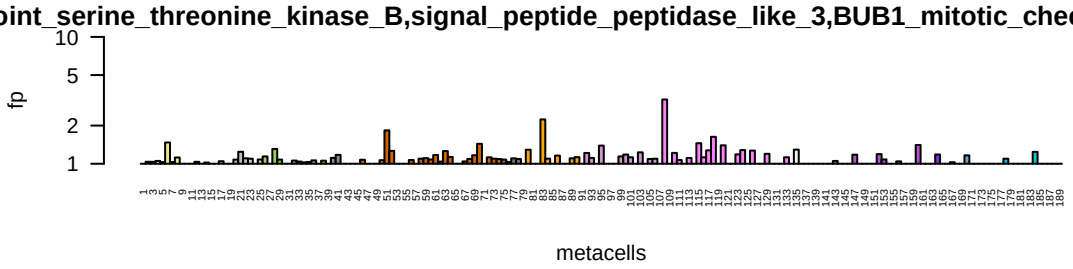
cyclin_dependent_kinase_1
fp
metacells


HoiH23 OG_6287
HoiH23_PlH23_005470−RA
cyclin_dependent_kinase_1
fp
metacells

Tadh OG_6929
Tadh_TriadT5779
oint_serine_threonine_kinase_B,signal_peptide_peptidase_like_3,BUB1_mitotic_checkpoin



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Tadh_TriadT59065
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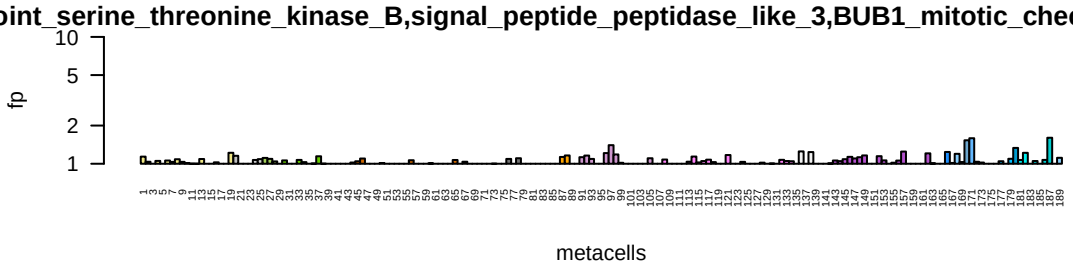
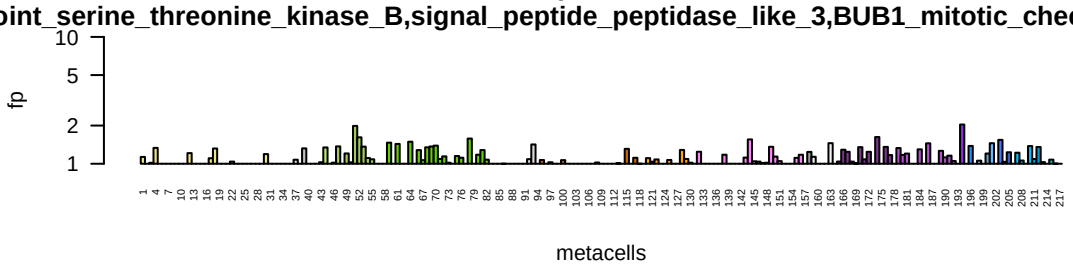


TrH2 OG_6929
TrH2_TrispH2_005950−RA
oint_serine_threonine_kinase_B,signal_peptide_peptidase_like_3,BUB1_mitotic_checkpoin



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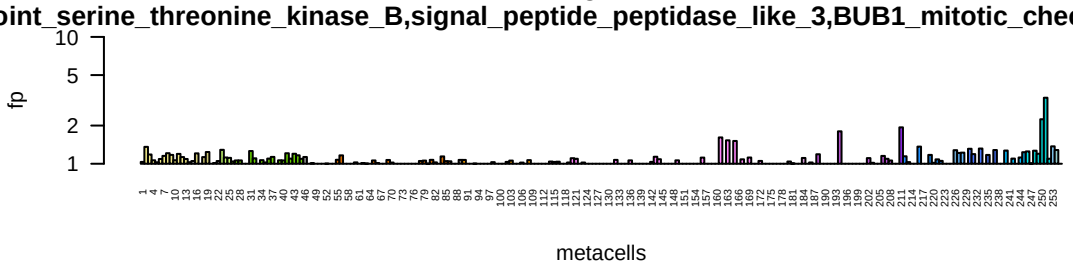
oint_serine_threonine_kinase_B,signal_peptide_peptidase_like_3,BUB1_mitotic_checkpoin



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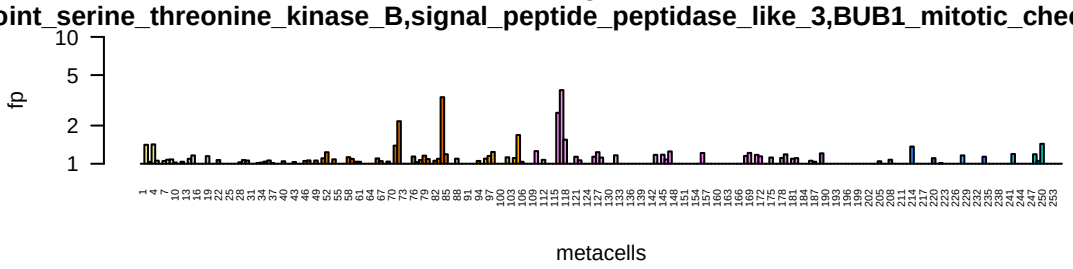
oint_serine_threonine_kinase_B,signal_peptide_peptidase_like_3,BUB1_mitotic_checkpoin



HoiH23 OG_6929
HoiH23_PlH23_000675−RA
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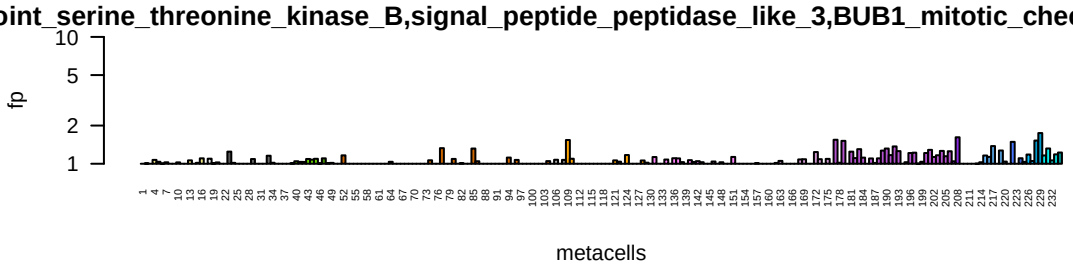

HoiH23 OG_6929
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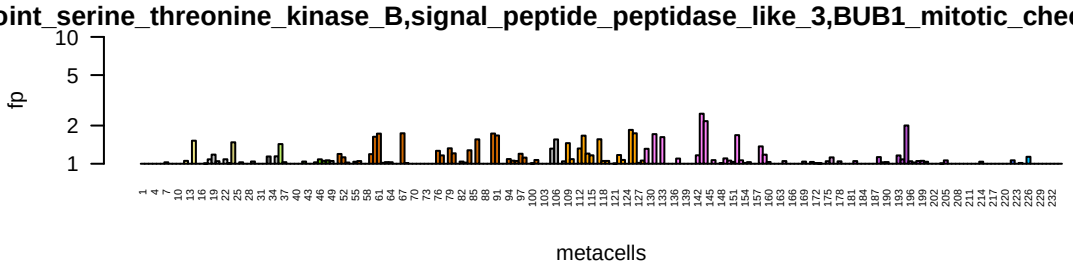

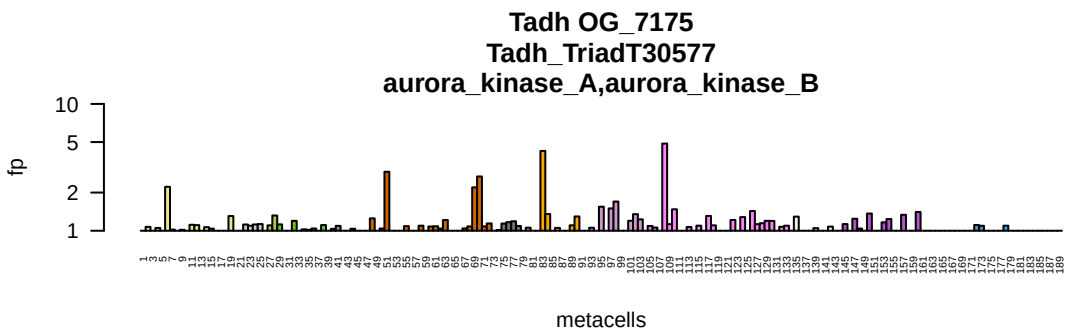
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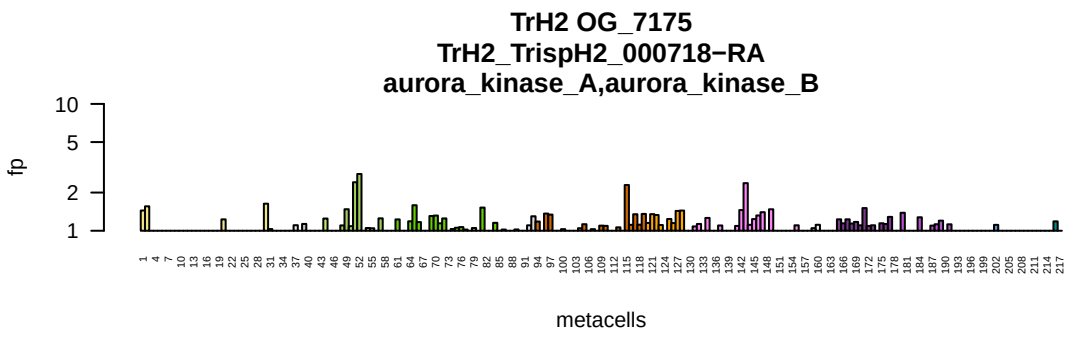
Tadh_TriadT30577
aurora_kinase_A,aurora_kinase_B
fp
metacells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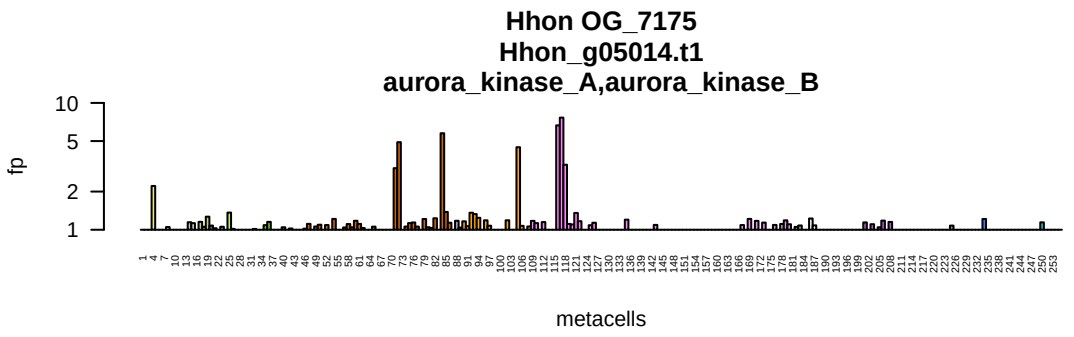
TrH2_TrispH2_000718−RA
aurora_kinase_A,aurora_kinase_B
fp
metacells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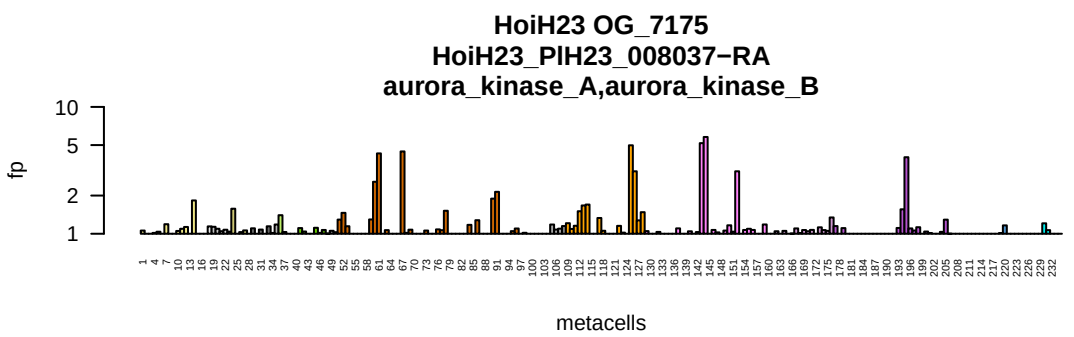
aurora_kinase_A,aurora_kinase_B
fp
metacells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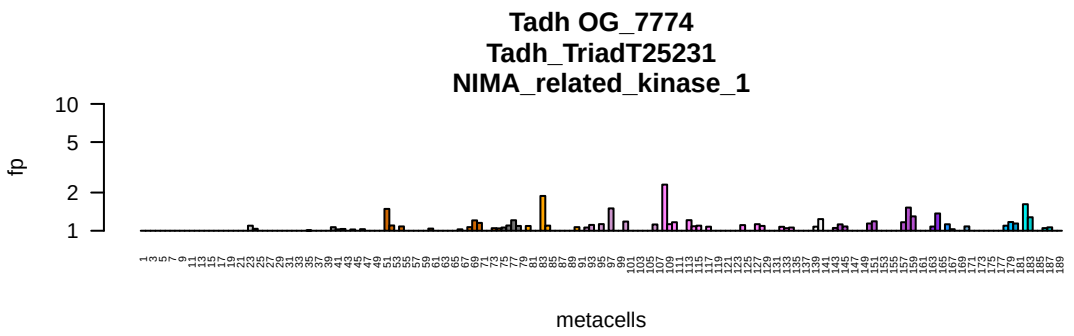
HoiH23_PlH23_008037−RA
aurora_kinase_A,aurora_kinase_B
fp
metacells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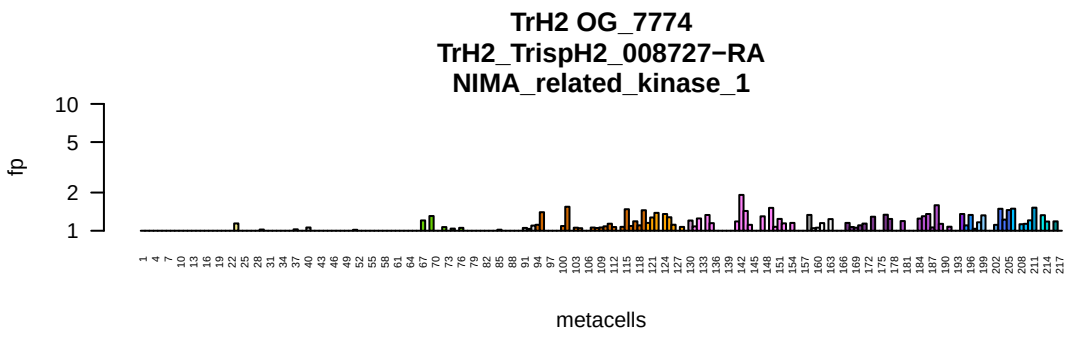
Tadh_TriadT25231
NIMA_related_kinase_1
fp
metacells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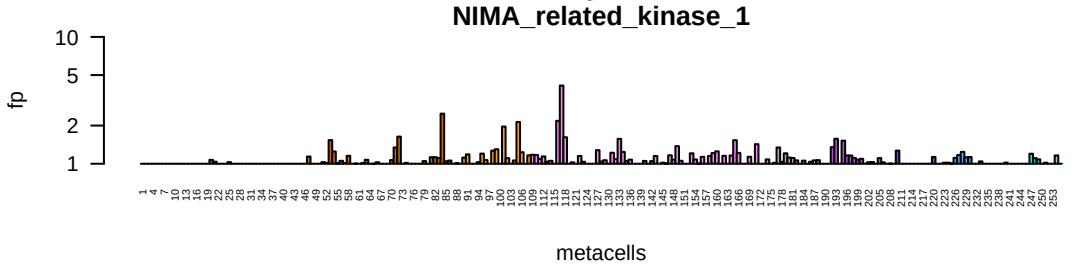
NIMA_related_kinase_1
fp
metacells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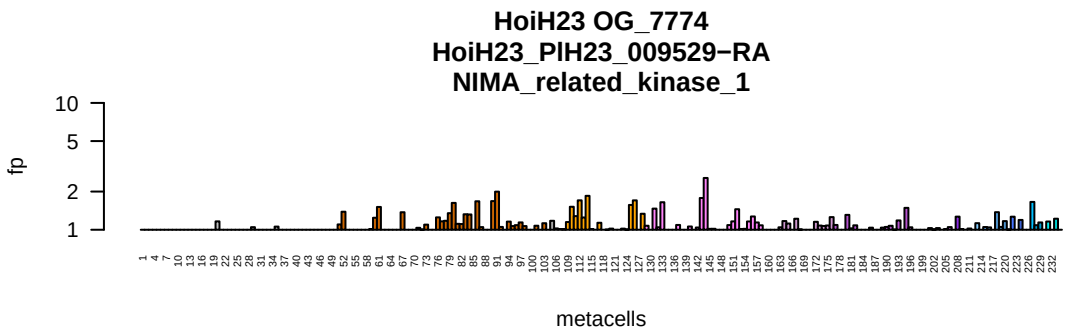
NIMA_related_kinase_1
fp
metacells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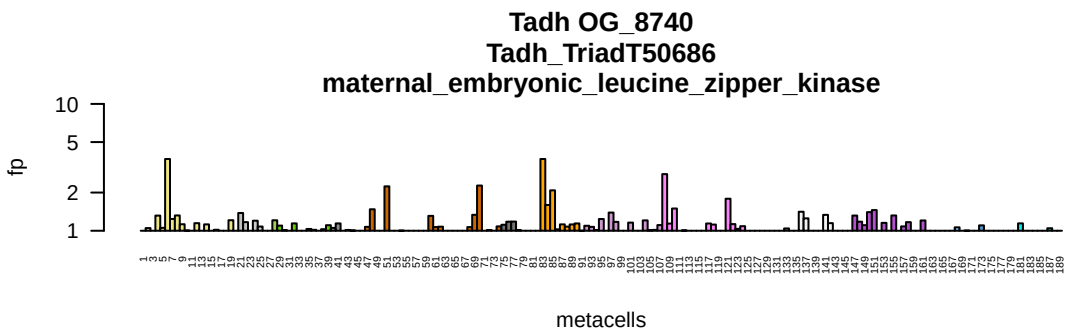
HoiH23_PlH23_009529−RA
NIMA_related_kinase_1
fp
metacells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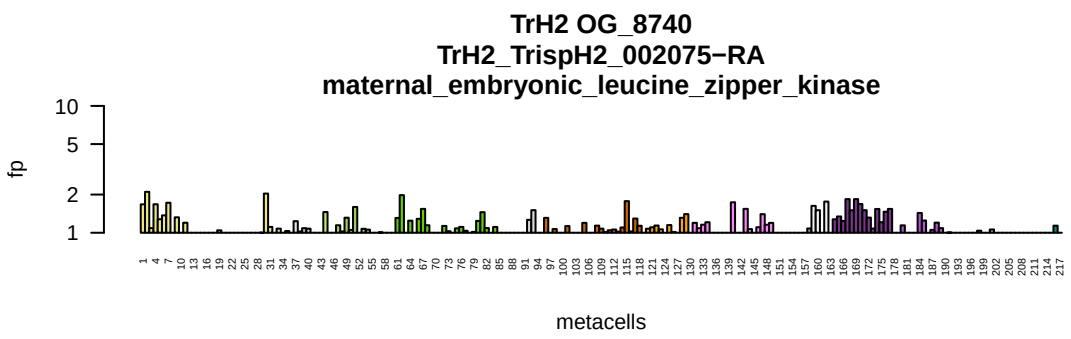
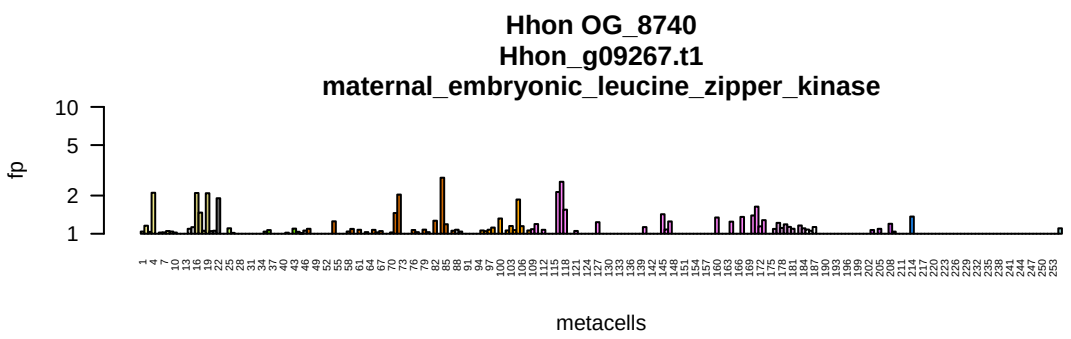
Tadh_TriadT50686
maternal_embryonic_leucine_zipper_kinase
fp
metacells


TrH2 OG_8740
TrH2_TrispH2_002075−RA
maternal_embryonic_leucine_zipper_kinase
fp
metacells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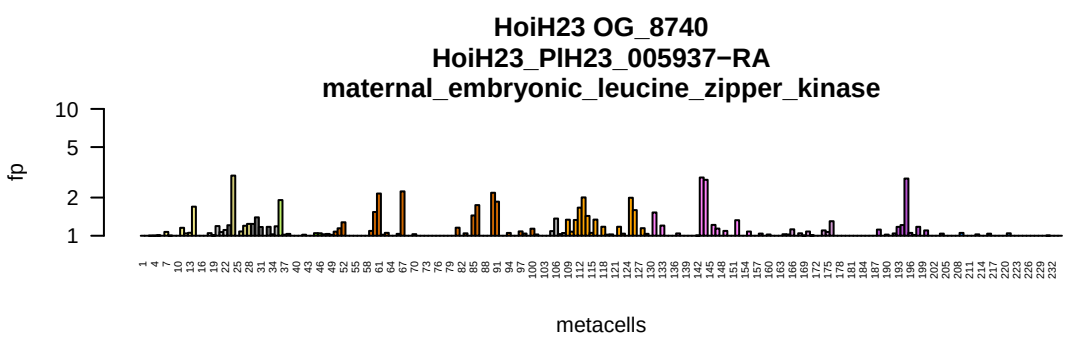
maternal_embryonic_leucine_zipper_kinase
fp
metacells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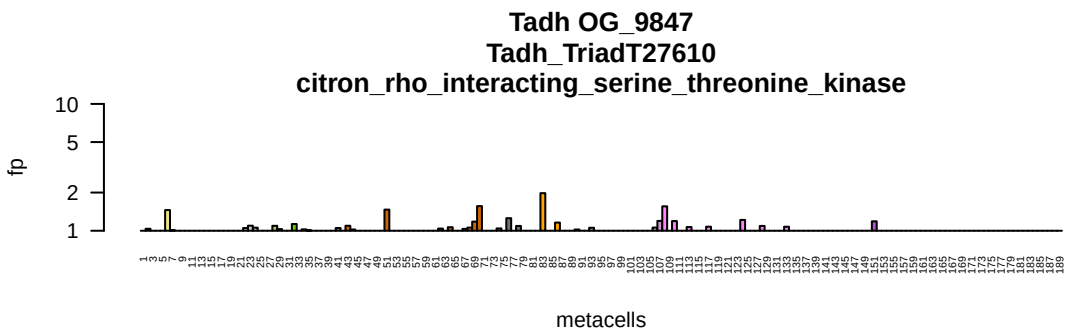
HoiH23_PlH23_005937−RA
maternal_embryonic_leucine_zipper_kinase
fp
metacells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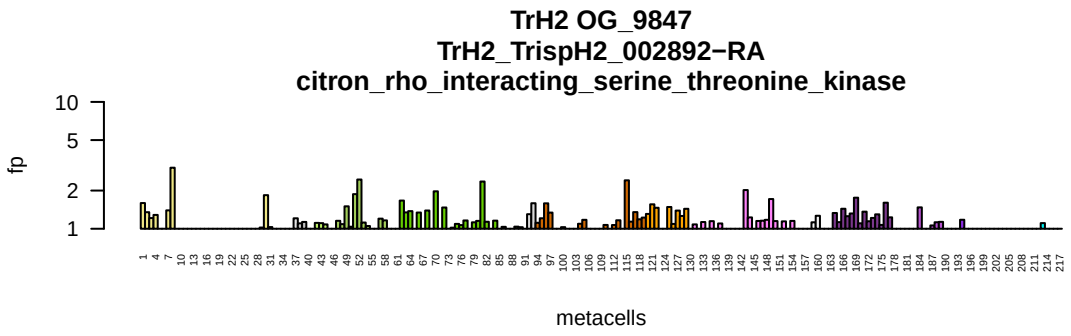
citron_rho_interacting_serine_threonine_kinase
fp
metacells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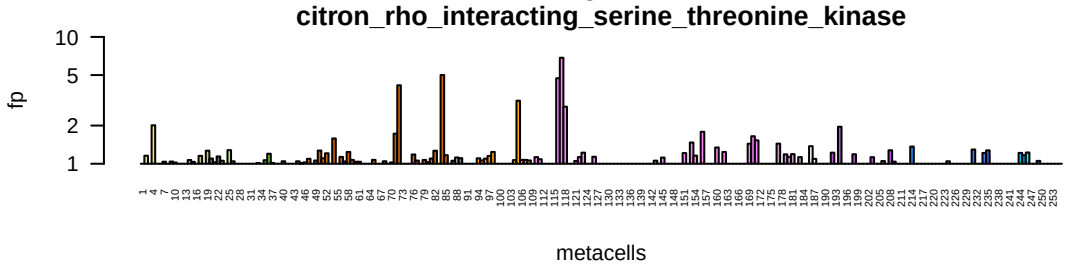
citron_rho_interacting_serine_threonine_kinase
fp
metacells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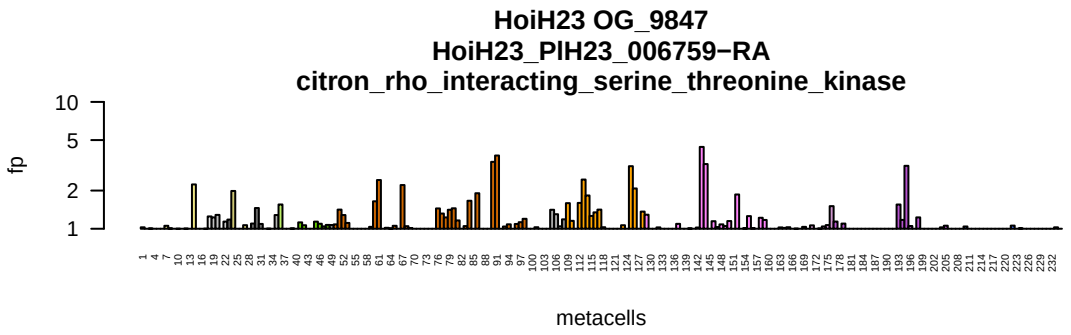
Hhon_g06660.t1
citron_rho_interacting_serine_threonine_kinase
fp
metacells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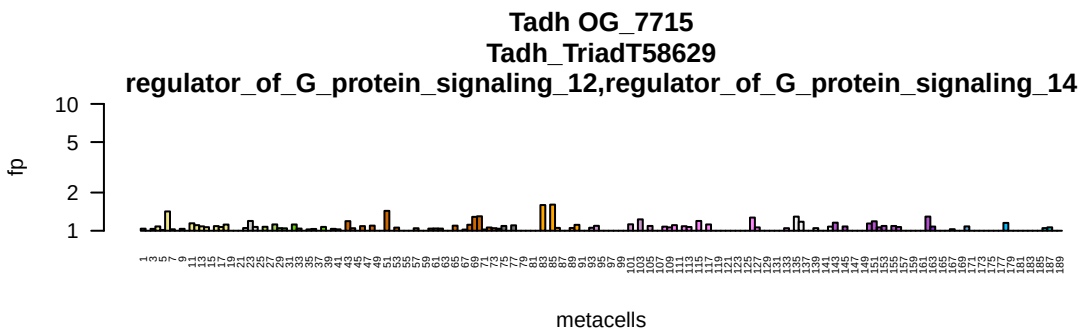
citron_rho_interacting_serine_threonine_kinase
fp
metacells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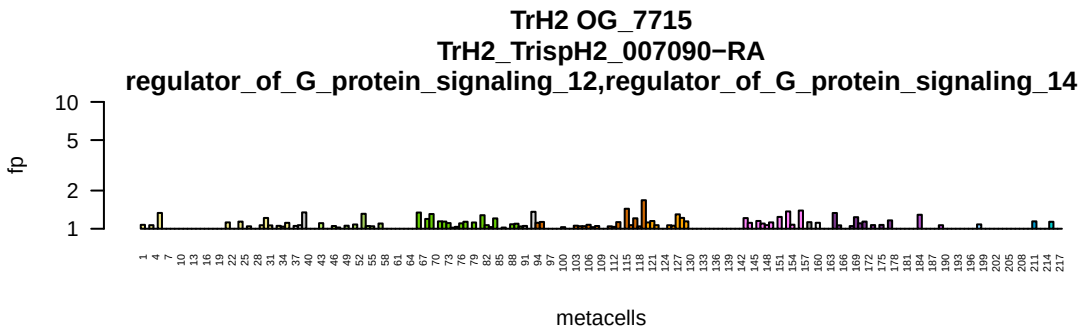
fp
metacells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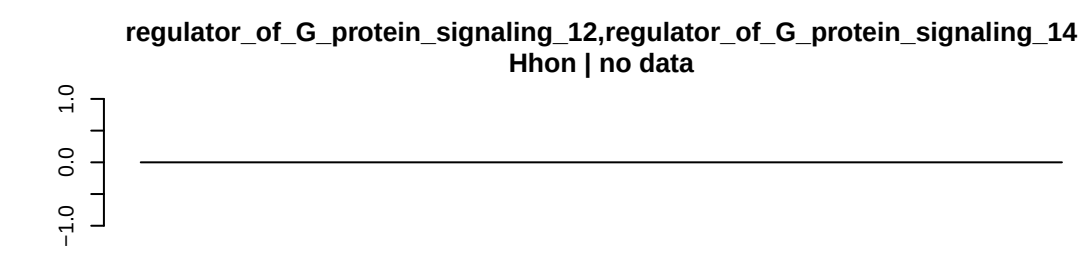
fp
metacells


regulator_of_G_protein_signaling_12,regulator_of_G_protein_signaling_14
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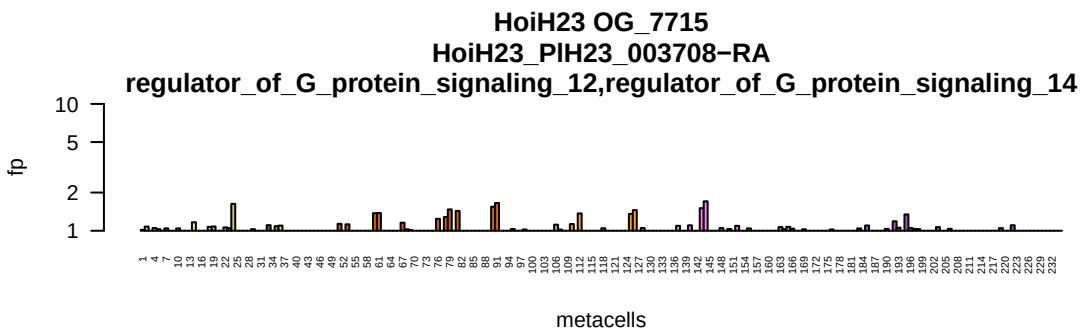
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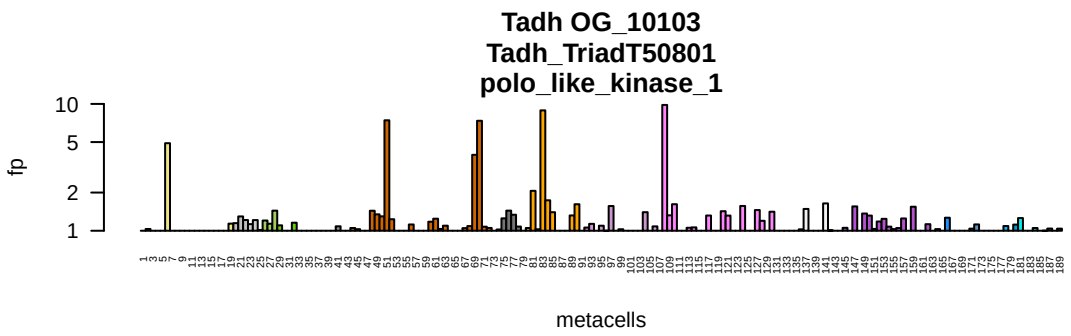
HoiH23_PlH23_003708−RA
regulator_of_G_protein_signaling_12,regulator_of_G_protein_signaling_14
fp
metacells


Tadh OG_10103
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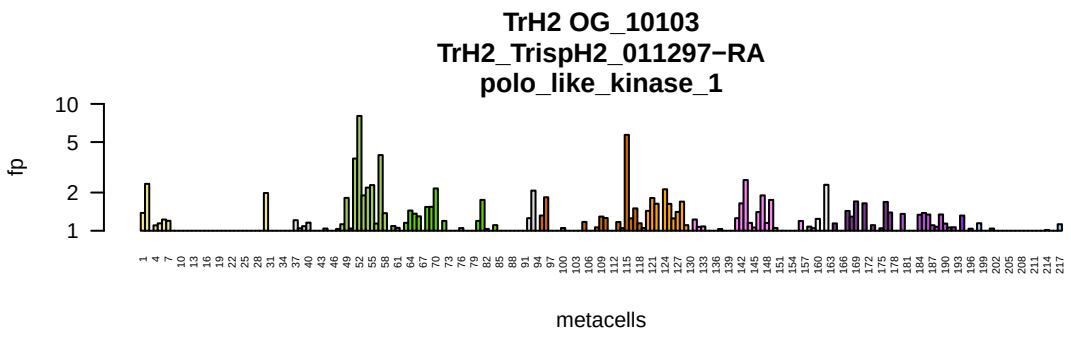
polo_like_kinase_1
fp
10
5
2
1
metacells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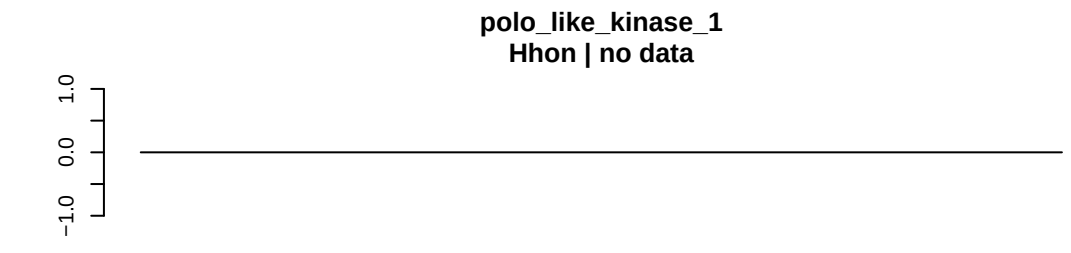
TrH2_TrispH2_011297−RA
polo_like_kinase_1
fp
10
5
2
1
metacells


polo_like_kinase_1
Hhon | no data
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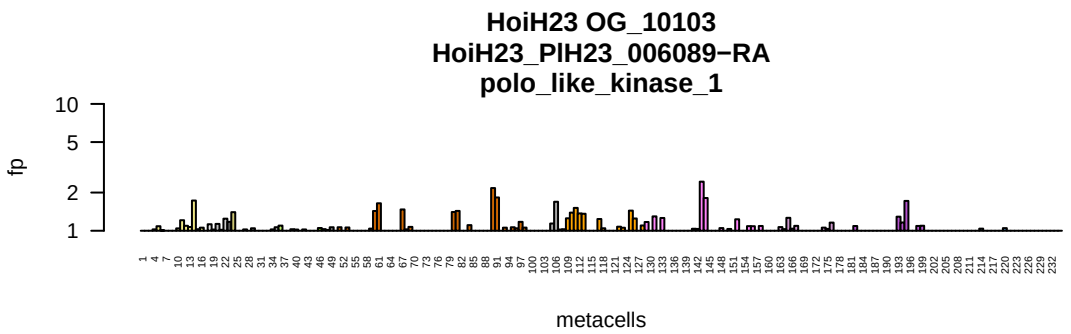
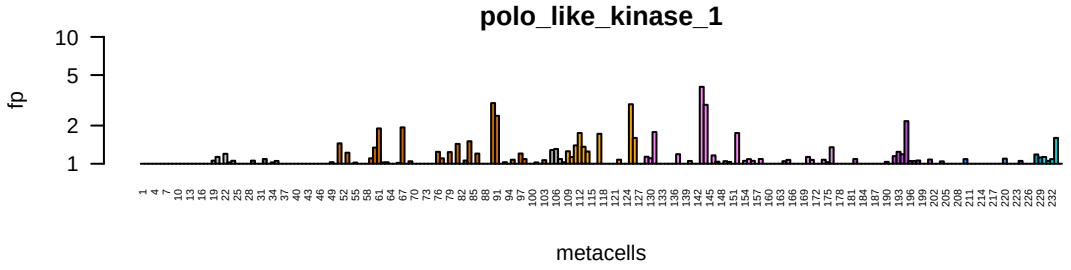
0.0
−1.0


HoiH23 OG_10103
HoiH23_PlH23_006089−RA
polo_like_kinase_1
fp
10
5
2
1
metacells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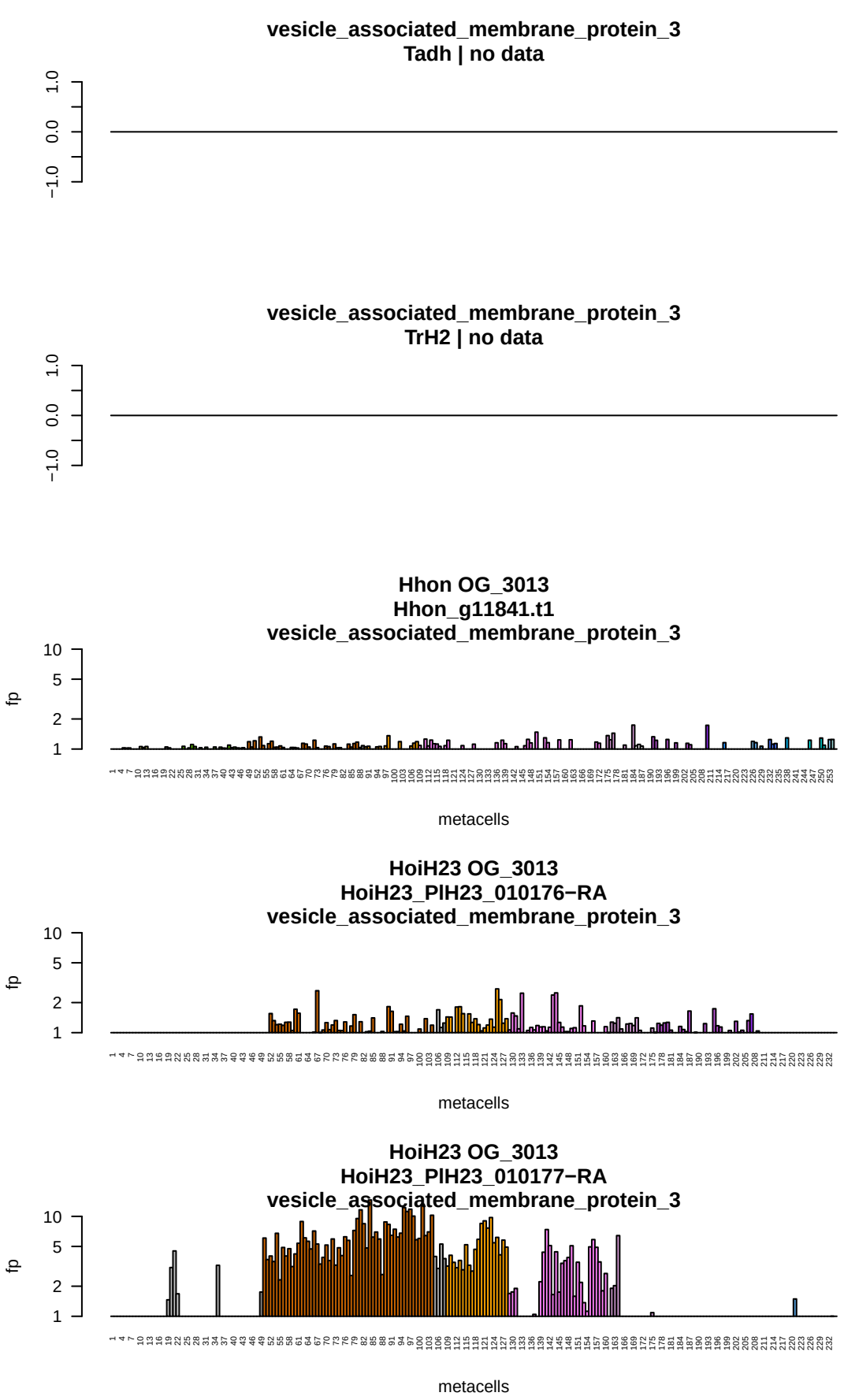
HoiH23_PlH23_006088−RA
polo_like_kinase_1
fp
10
5
2
1
metacells


vesicle_associated_membrane_protein_3
Tadh | no data
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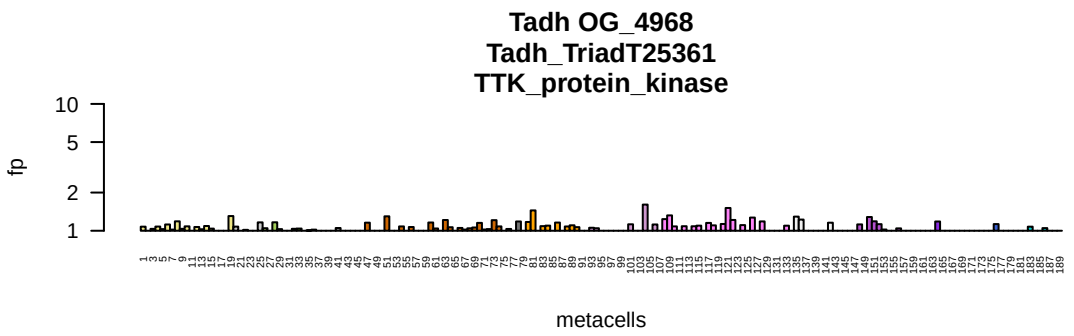
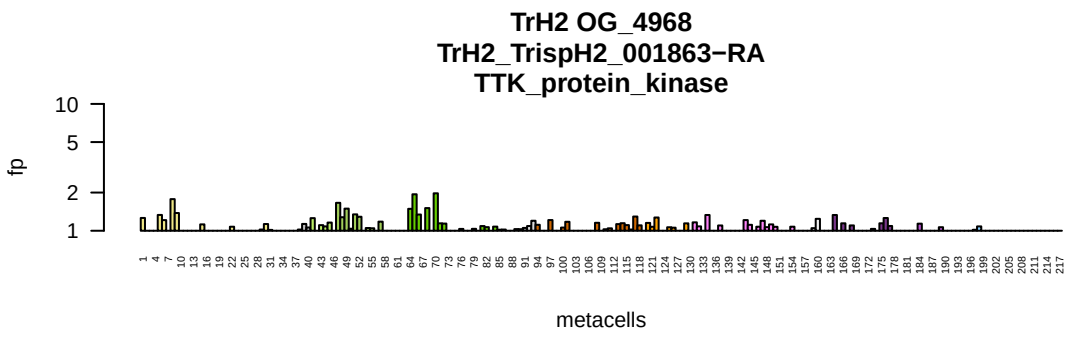
0.0
−1.0
vesicle_associated_membrane_protein_3
TrH2 | no data
1.0
0.0
−1.0

Hhon OG_3013
Hhon_g11841.t1
vesicle_associated_membrane_protein_3
fp
10
5
2
1
metacells

HoiH23 OG_3013
HoiH23_PlH23_010176−RA
vesicle_associated_membrane_protein_3
fp
10
5
2
1
metacells

HoiH23 OG_3013
HoiH23_PlH23_010177−RA
vesicle_associated_membrane_protein_3
fp
10
5
2
1
metacells


Tadh OG_4968
Tadh_TriadT25361
TTK_protein_kinase
fp
metacells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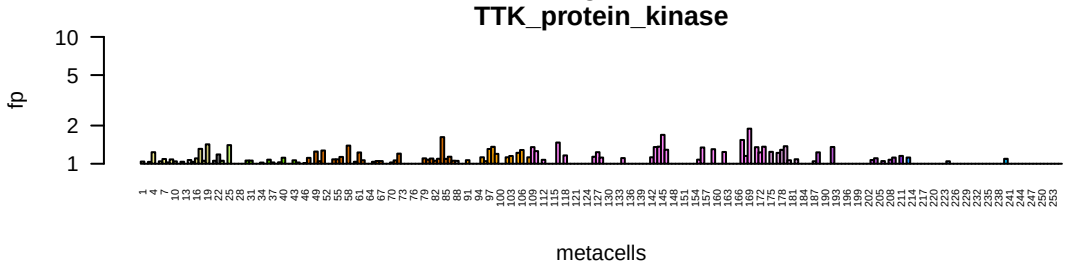
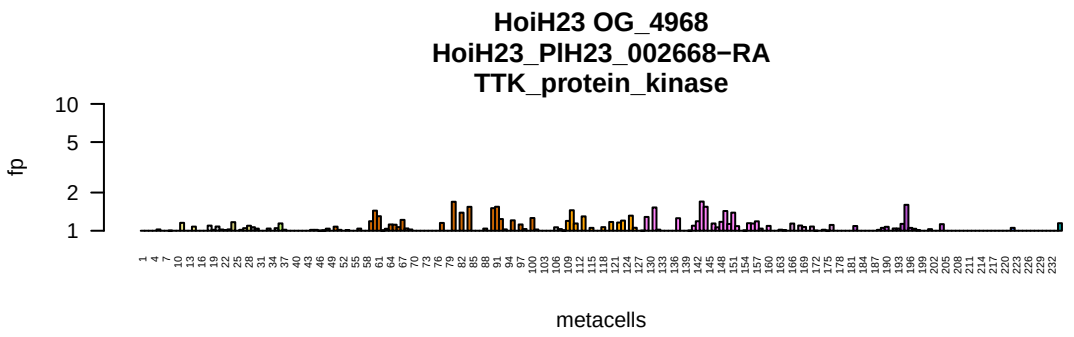
TrH2_TrispH2_001863−RA
TTK_protein_kinase
fp
metacells


Hhon OG_4968
Hhon_g05873.t1
TTK_protein_kinase
fp
metacells


HoiH23 OG_4968
HoiH23_PlH23_002668−RA
TTK_protein_kinase
fp
metacells