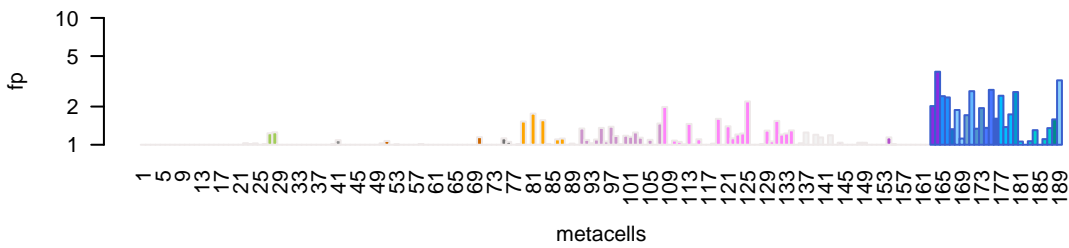
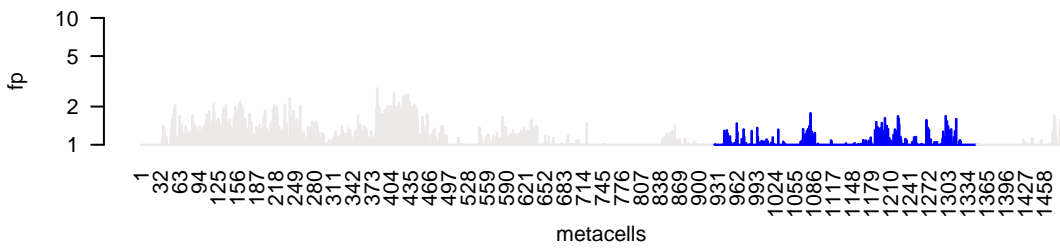


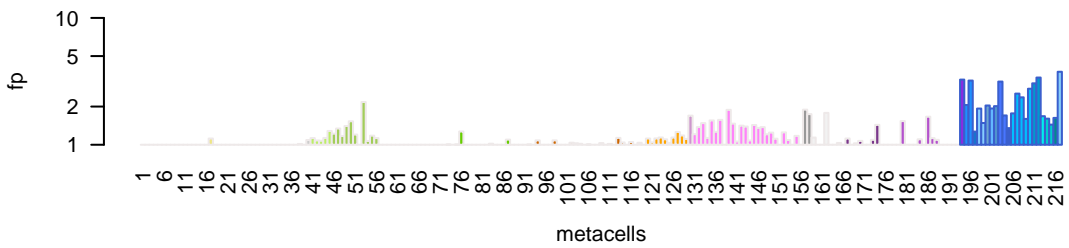
OG_1449:TCF3/TCF4/TCF12/Tcf4 Tadh | Tadh_TriadT8076



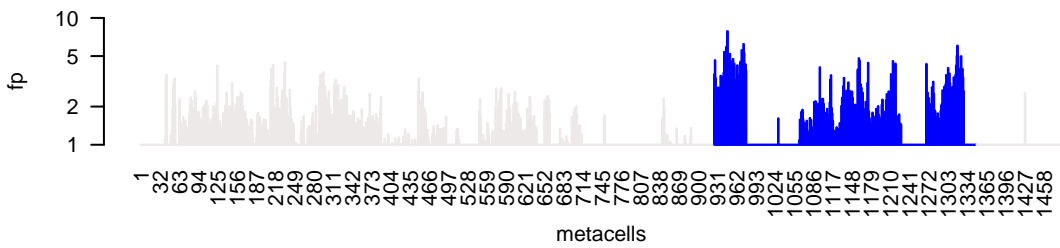
OG_1449:TCF3/TCF4/TCF12/Tcf4 Mmus | Mmus_ENSMUST00000183404



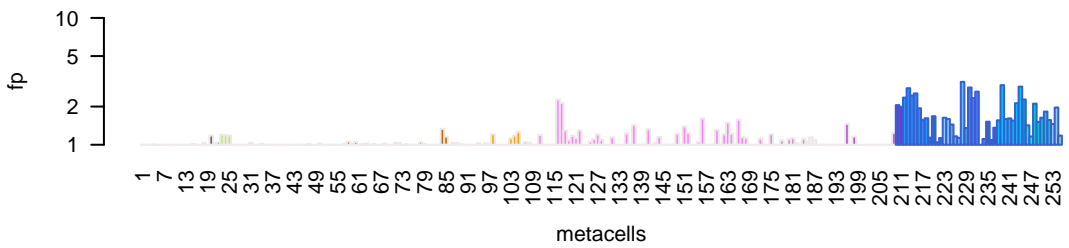
OG_1449:TCF3/TCF4/TCF12/Tcf4 TrH2 | TrH2_TrispH2_001049-RA



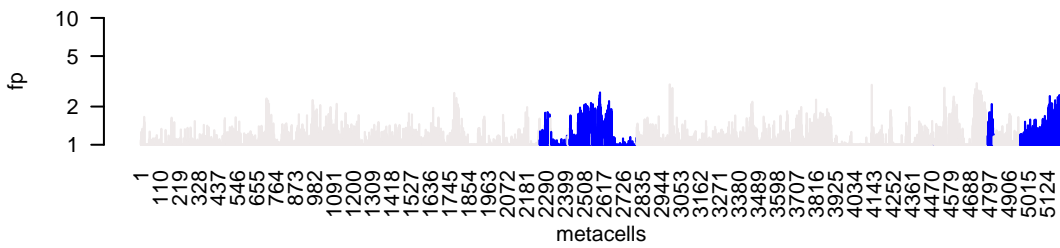
OG_1449:TCF3/TCF4/TCF12/Tcf4 Mmus | Mmus_ENSMUST00000066717



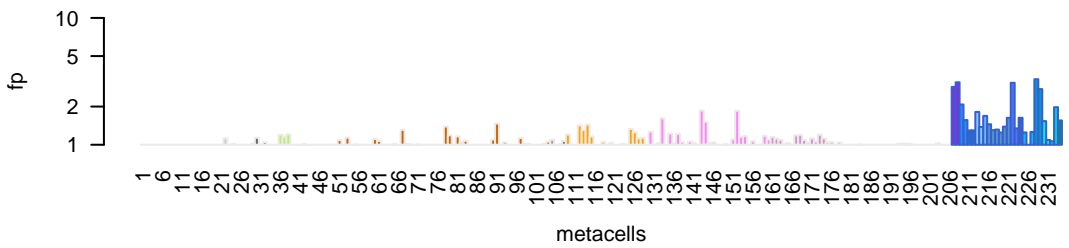
OG_1449:TCF3/TCF4/TCF12/Tcf4 Hhon | Hhon_g02873.t1



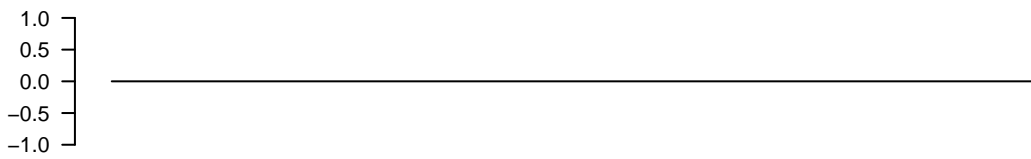
OG_1449:TCF3/TCF4/TCF12/Tcf4 Dmel | Dmel_FBtr0474232



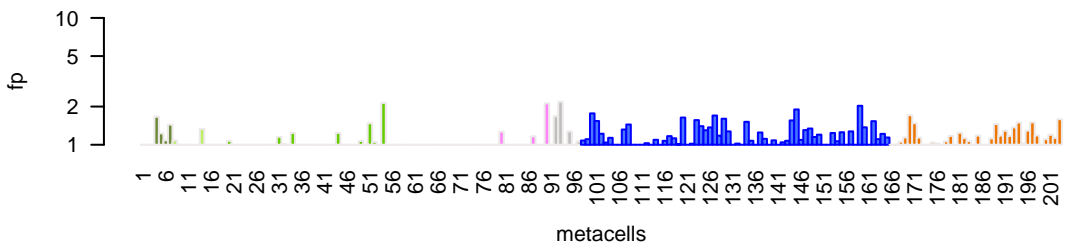
OG_1449:TCF3/TCF4/TCF12/Tcf4 HoiH23 | HoiH23_PIH23_009657-RA



OG_1449:TCF3/TCF4/TCF12/Tcf4 Mlei | no data



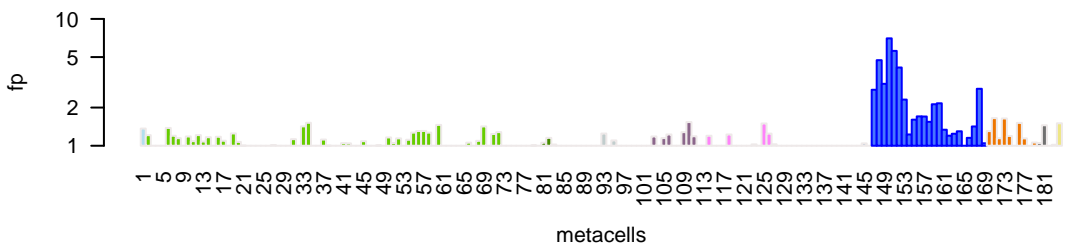
OG_1449:TCF3/TCF4/TCF12/Tcf4 Nvec | Nvec_v1g207162



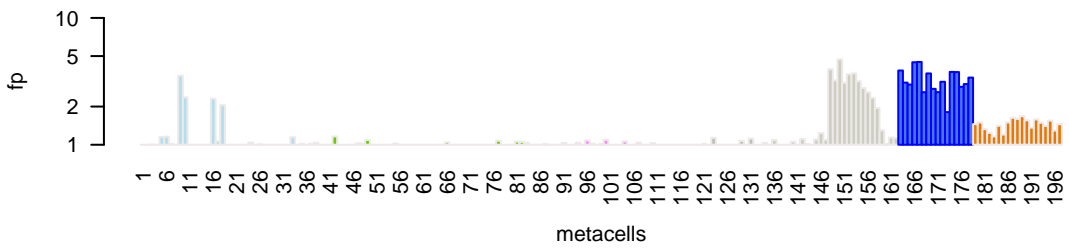
OG_1449:TCF3/TCF4/TCF12/Tcf4 Spolac | no data



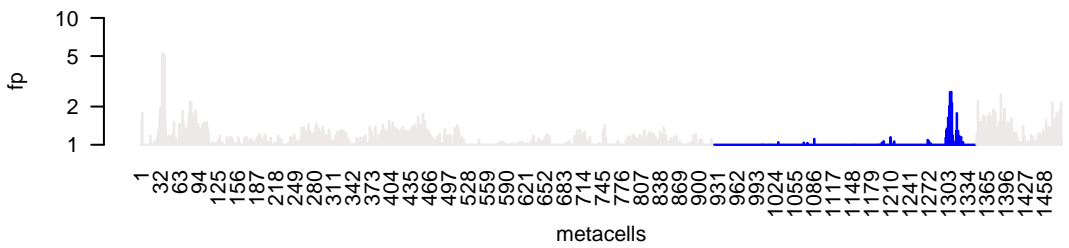
OG_1449:TCF3/TCF4/TCF12/Tcf4 Spis | Spis_XP_022797670_1

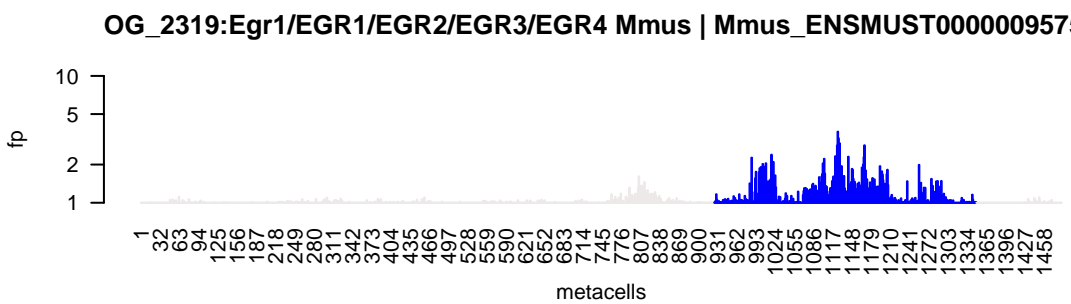
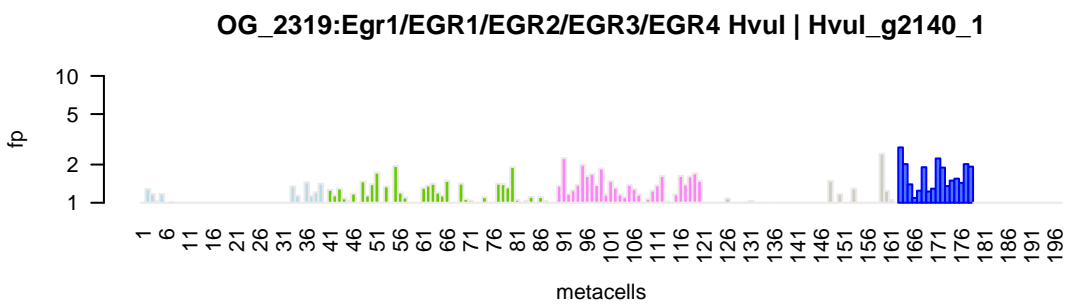
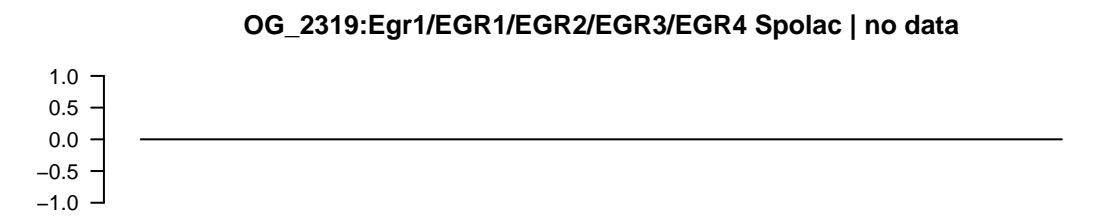
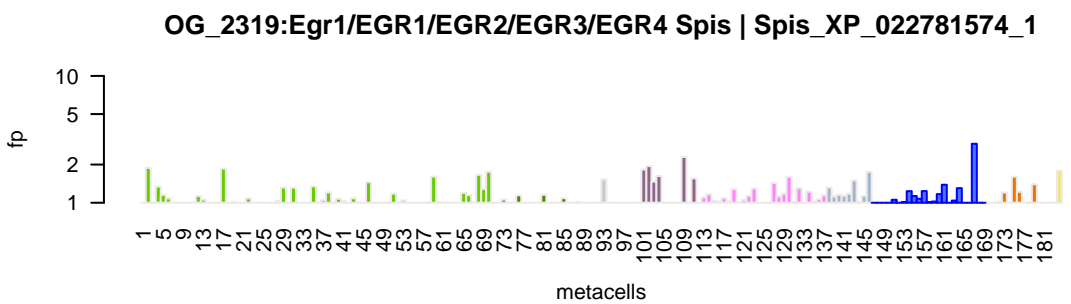
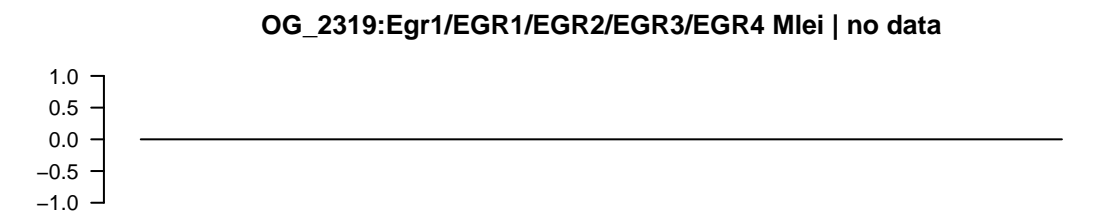
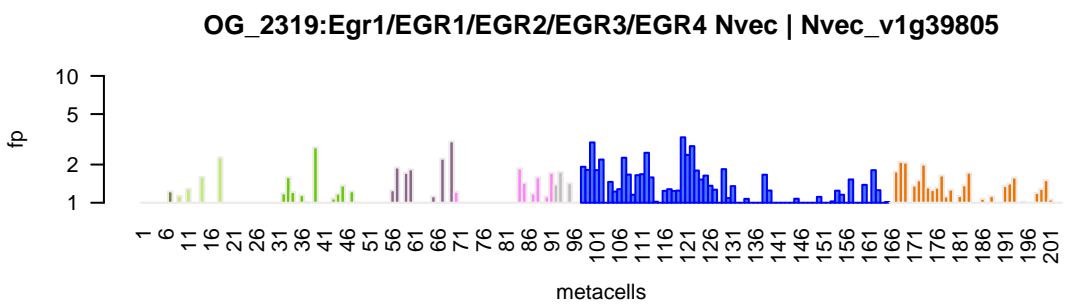
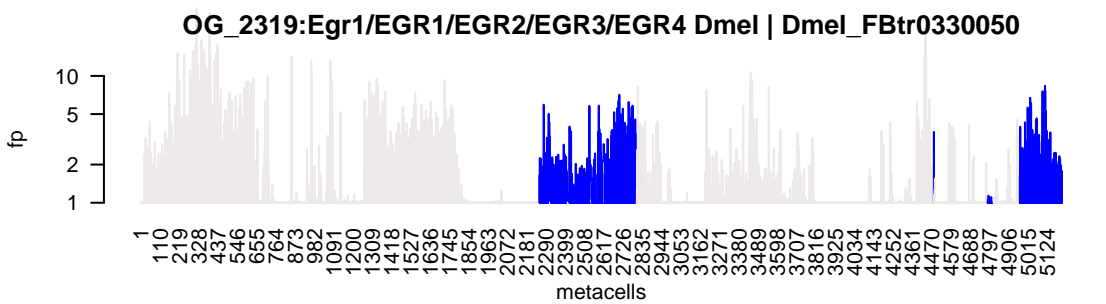
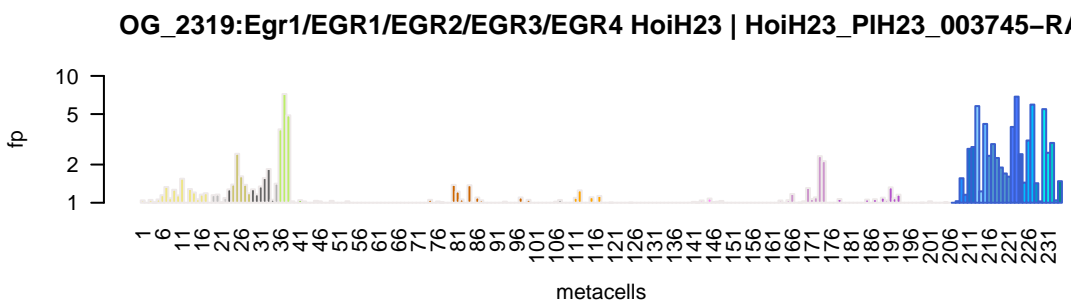
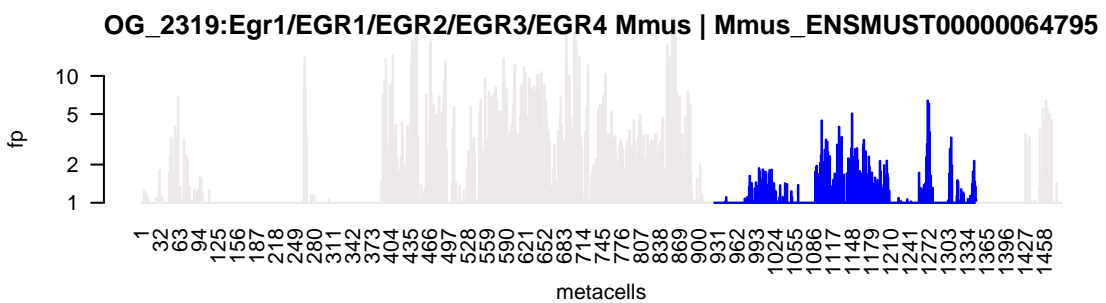
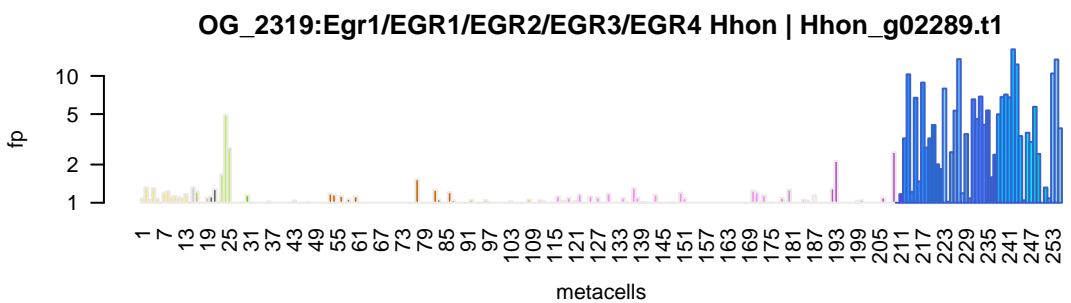
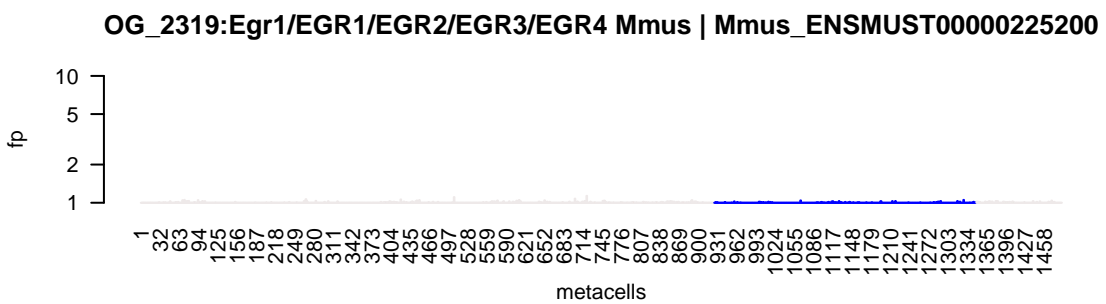
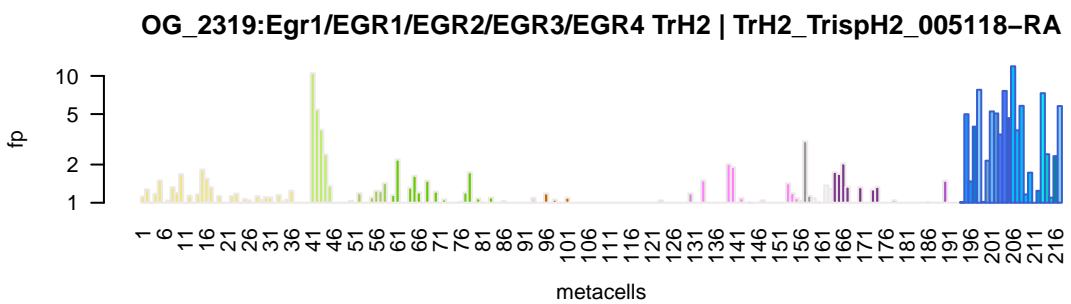
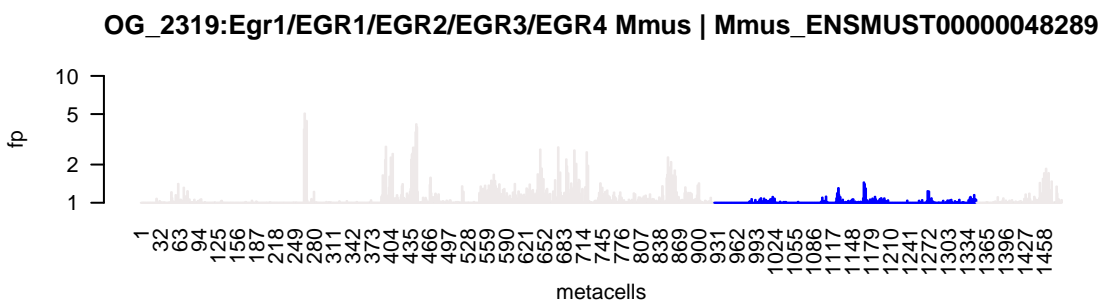
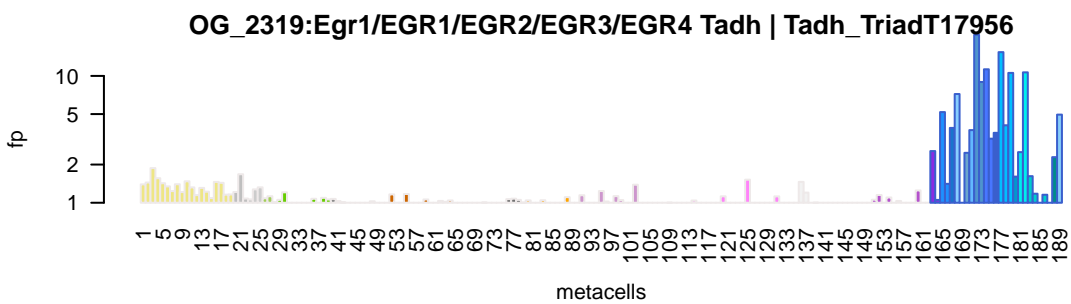


OG_1449:TCF3/TCF4/TCF12/Tcf4 Hvul | Hvul_g2666_1

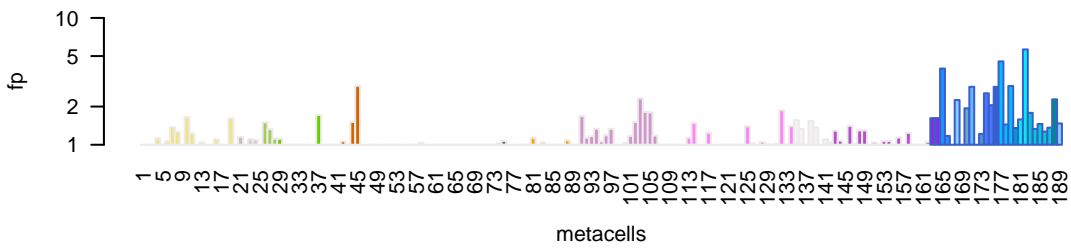


OG_1449:TCF3/TCF4/TCF12/Tcf4 Mmus | Mmus_ENSMUST00000105342

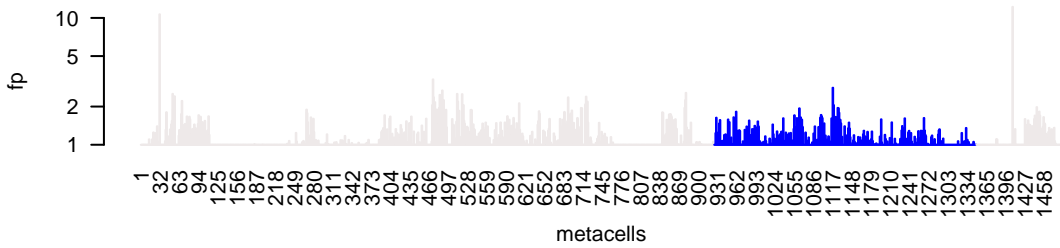




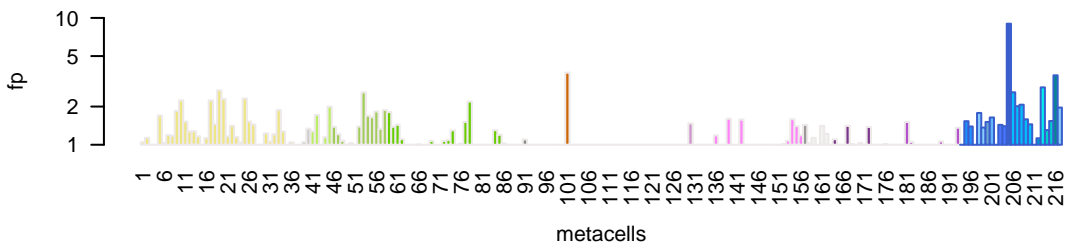
OG_3275:XPB1 Tadh | Tadh_TriadT64159



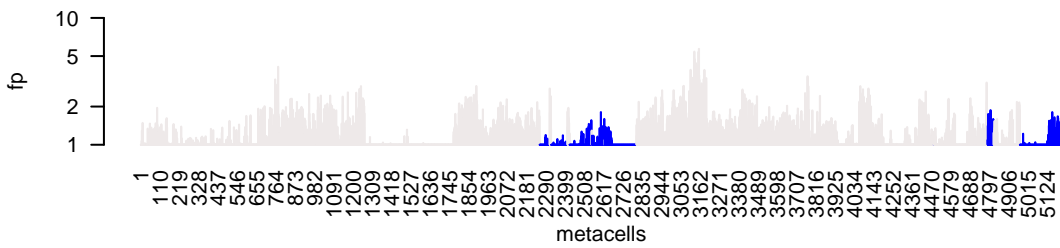
OG_3275:XPB1 Mmus | Mmus_ENSMUST00000239150



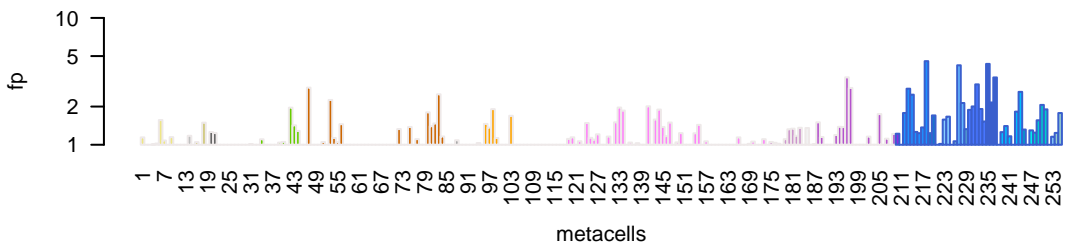
OG_3275:XPB1 TrH2 | TrH2_TrispH2_005934-RA



OG_3275:XPB1 Dmel | Dmel_FBtr0310045



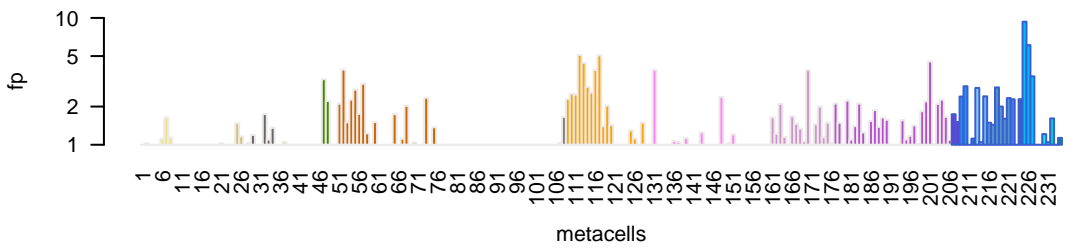
OG_3275:XPB1 Hhon | Hhon_g02674.t1



OG_3275:XPB1 Mlei | no data



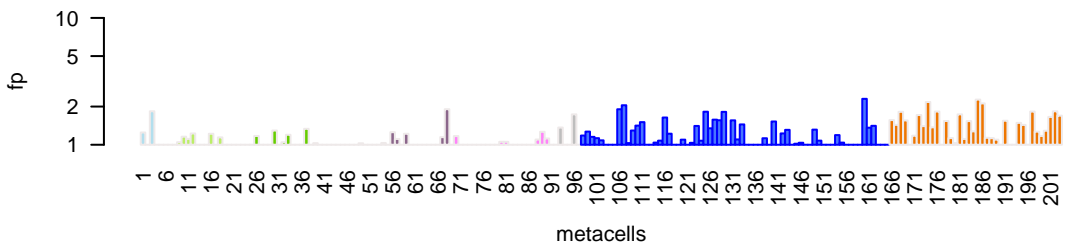
OG_3275:XPB1 HoiH23 | HoiH23_PIH23_000692-RA



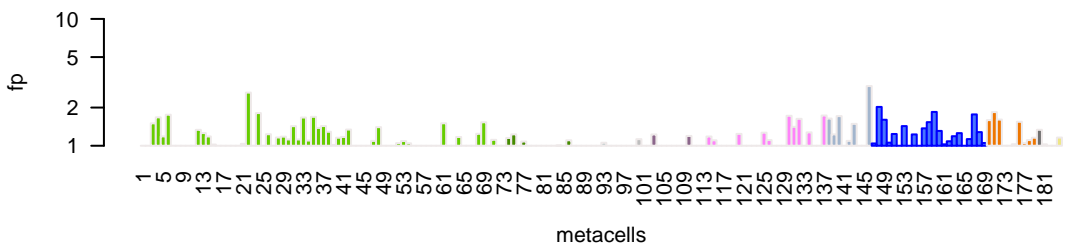
OG_3275:XPB1 Spolac | no data



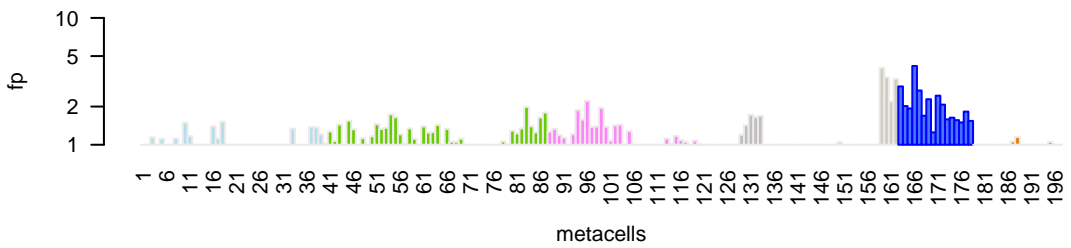
OG_3275:XPB1 Nvec | Nvec_v1g211292



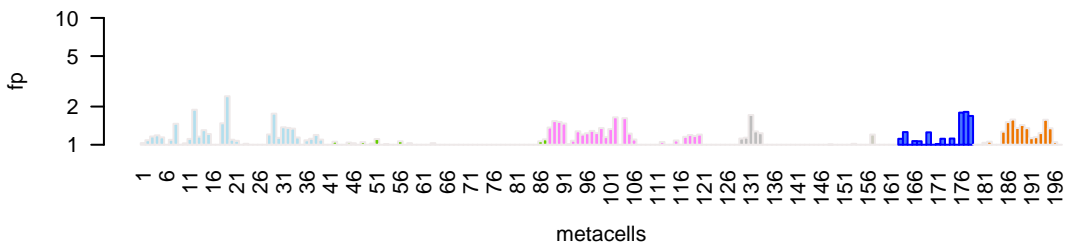
OG_3275:XPB1 Spis | Spis_XP_022794277_1

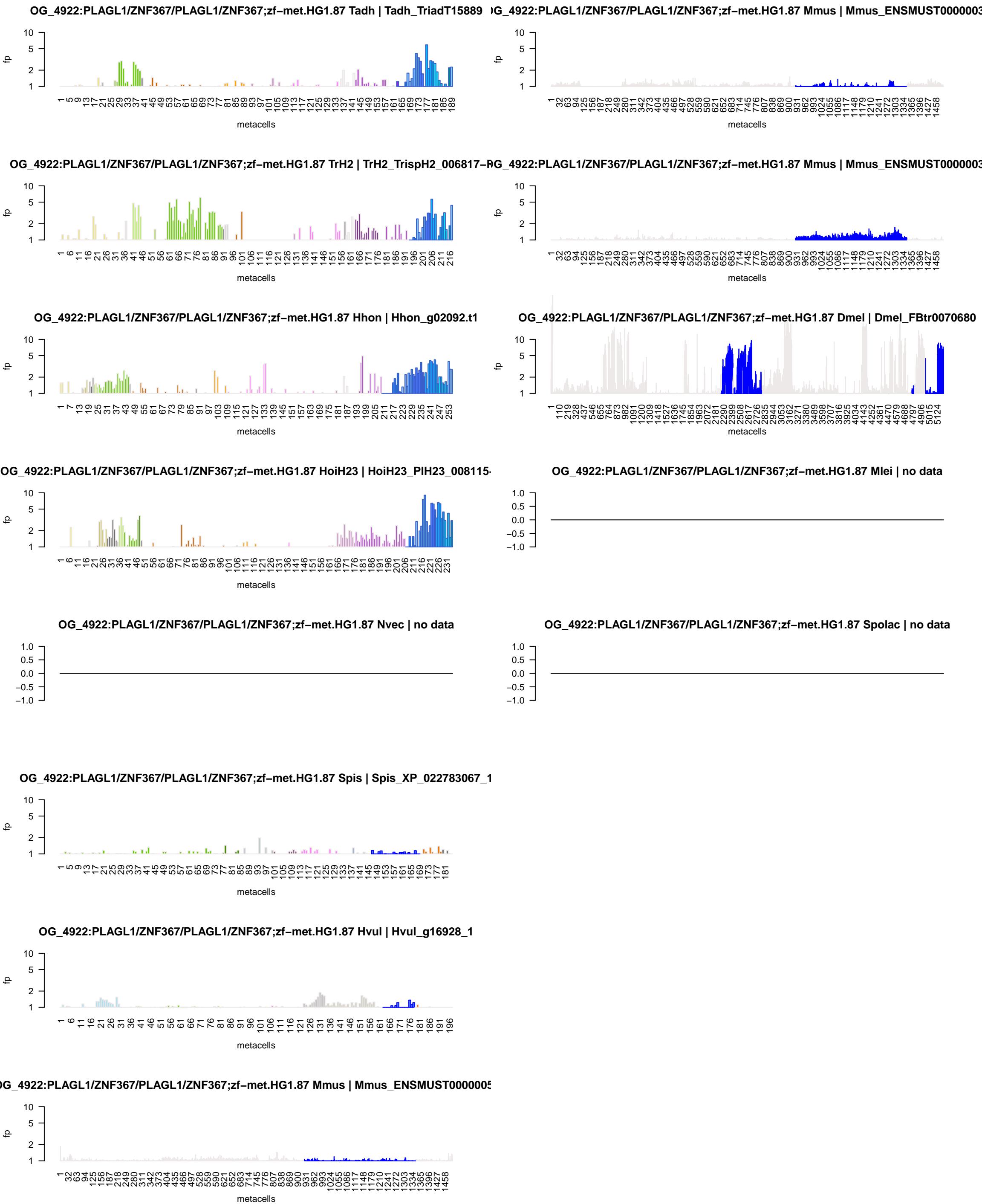


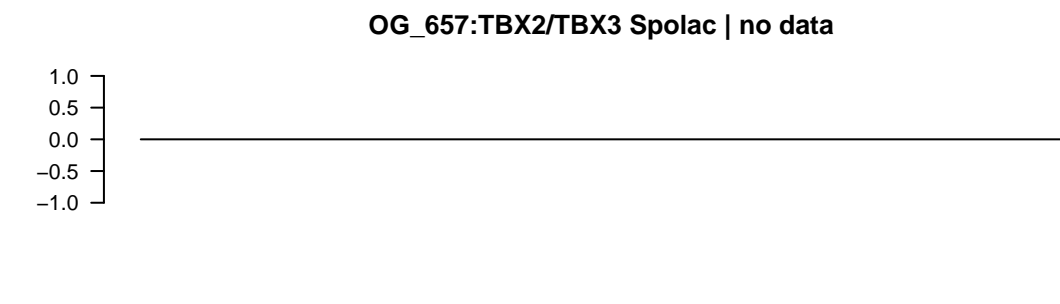
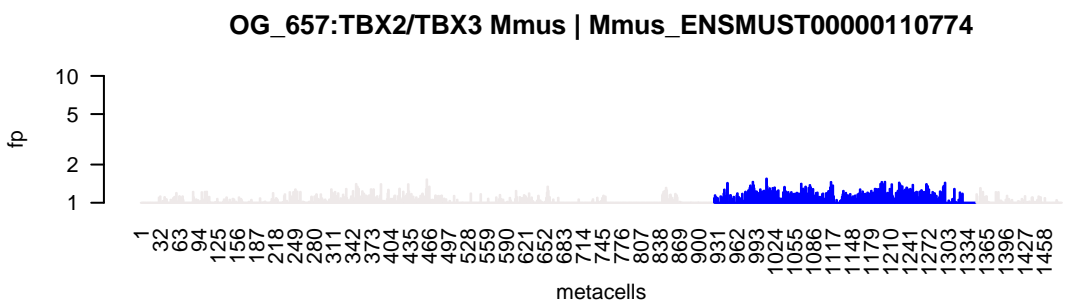
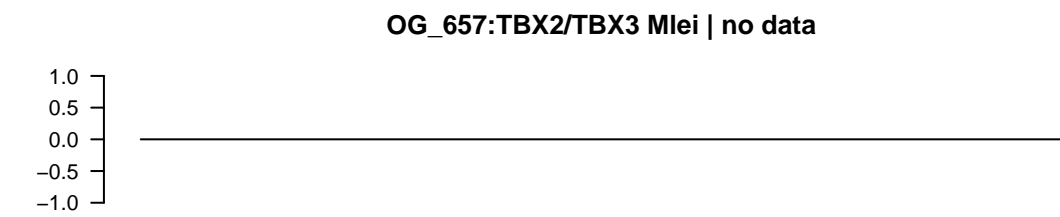
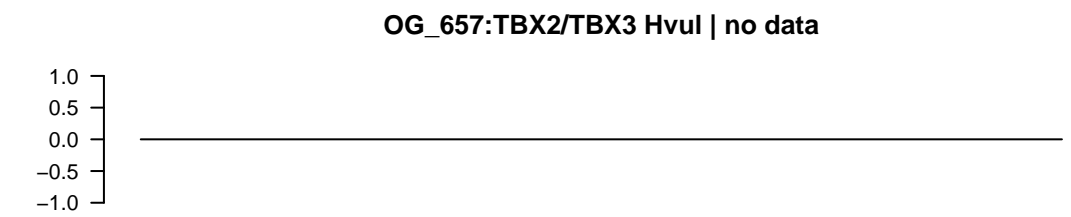
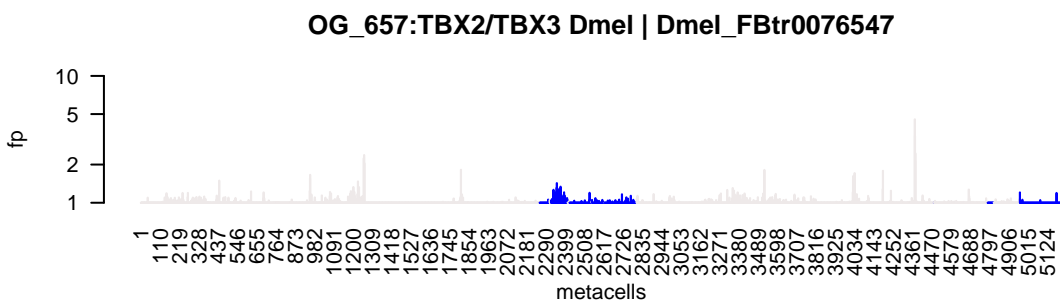
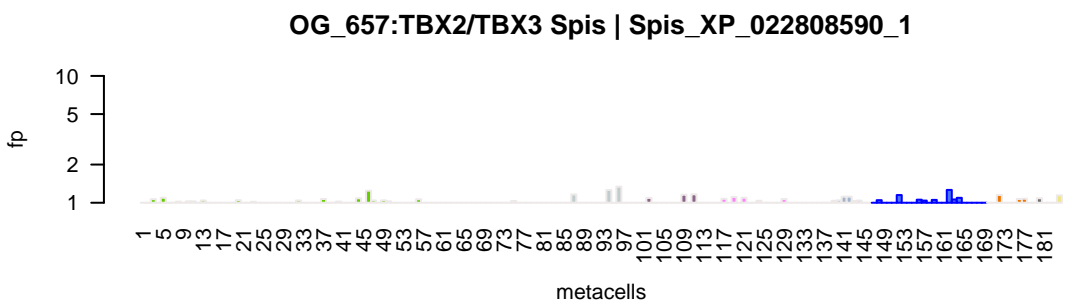
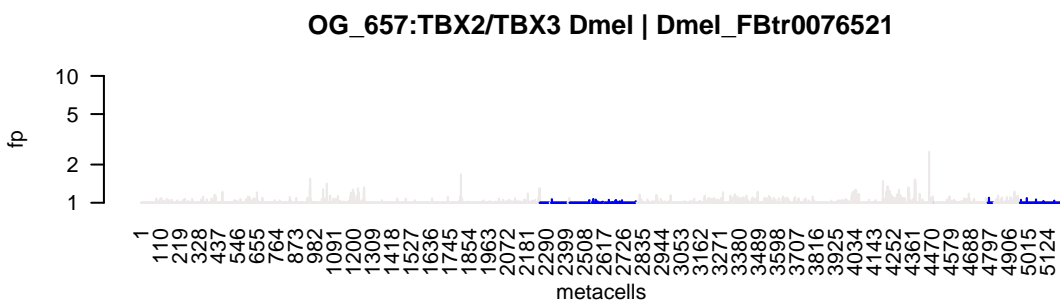
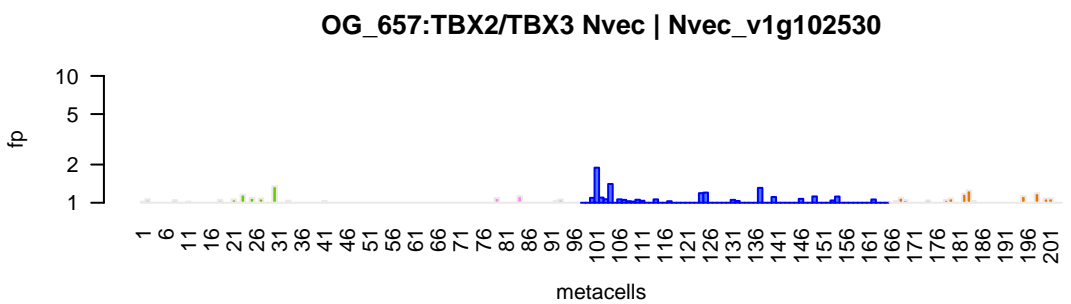
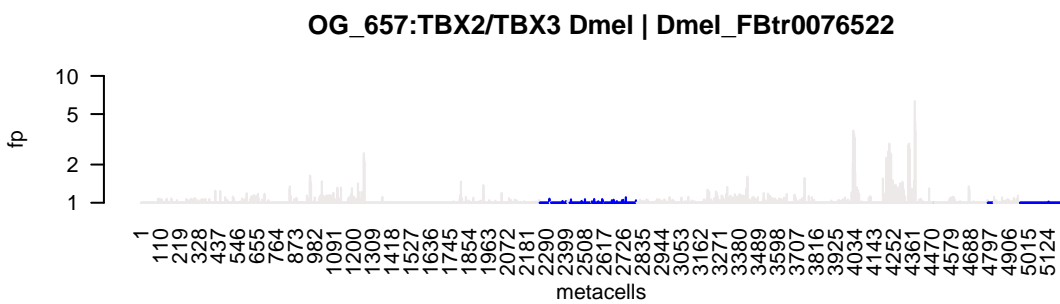
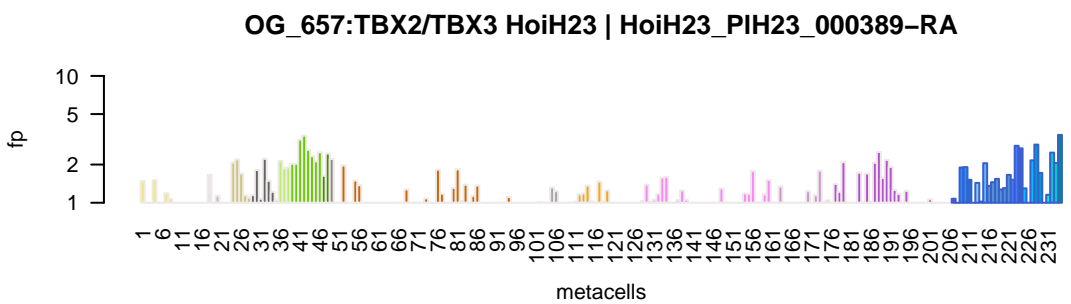
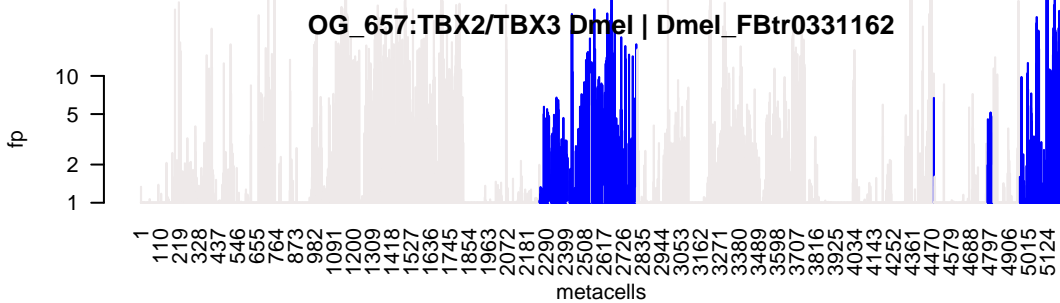
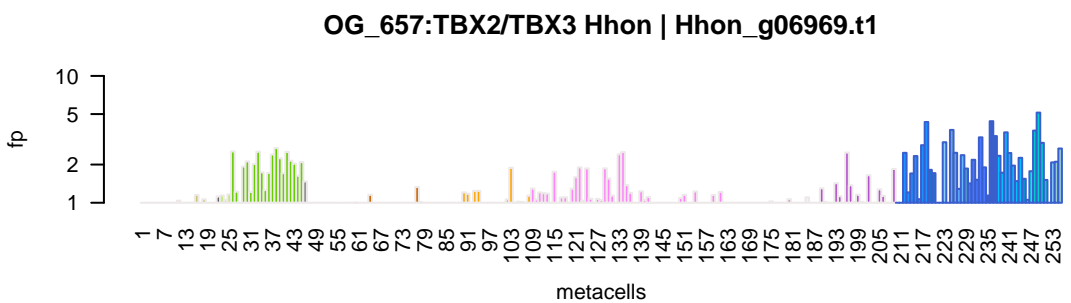
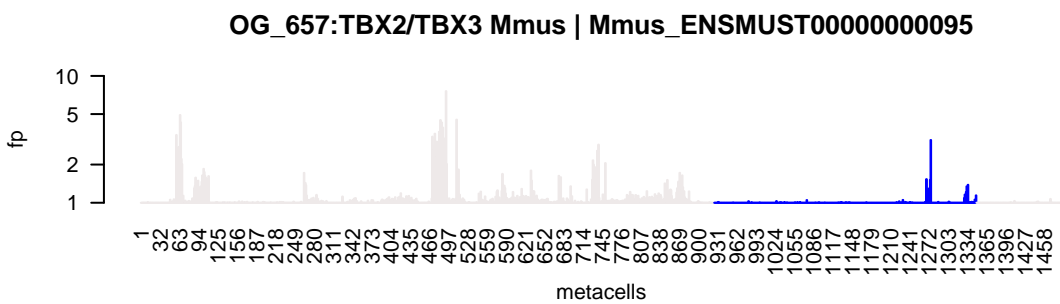
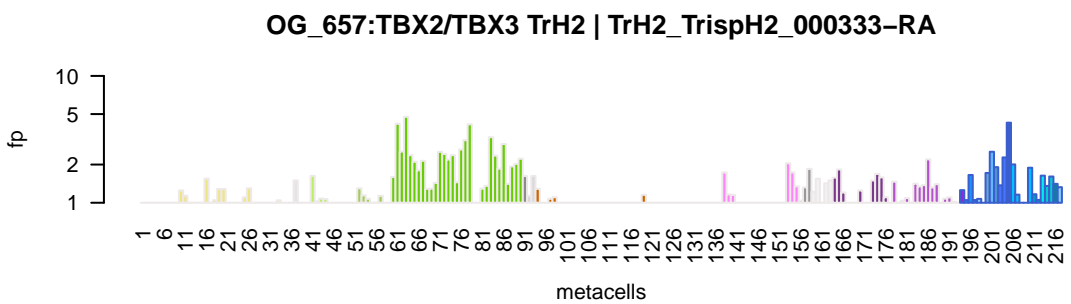
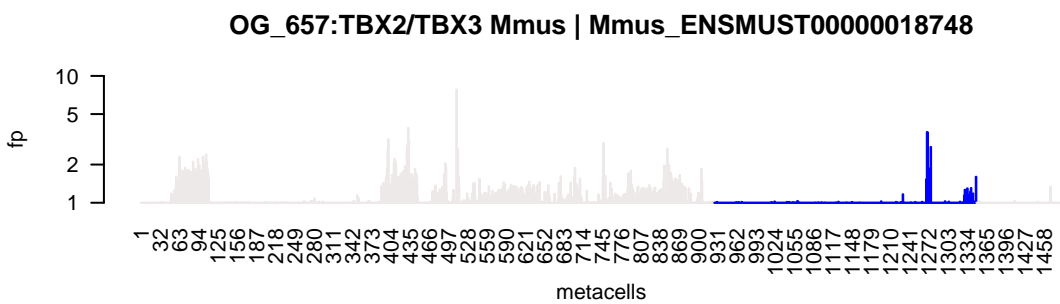
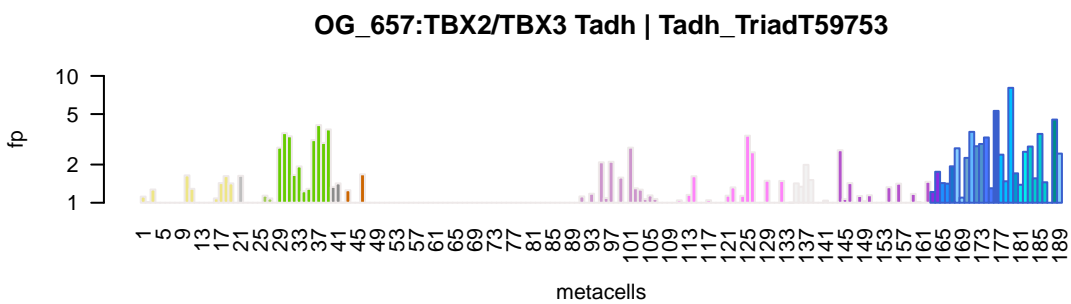
OG_3275:XPB1 Hvul | Hvul_g10594_1

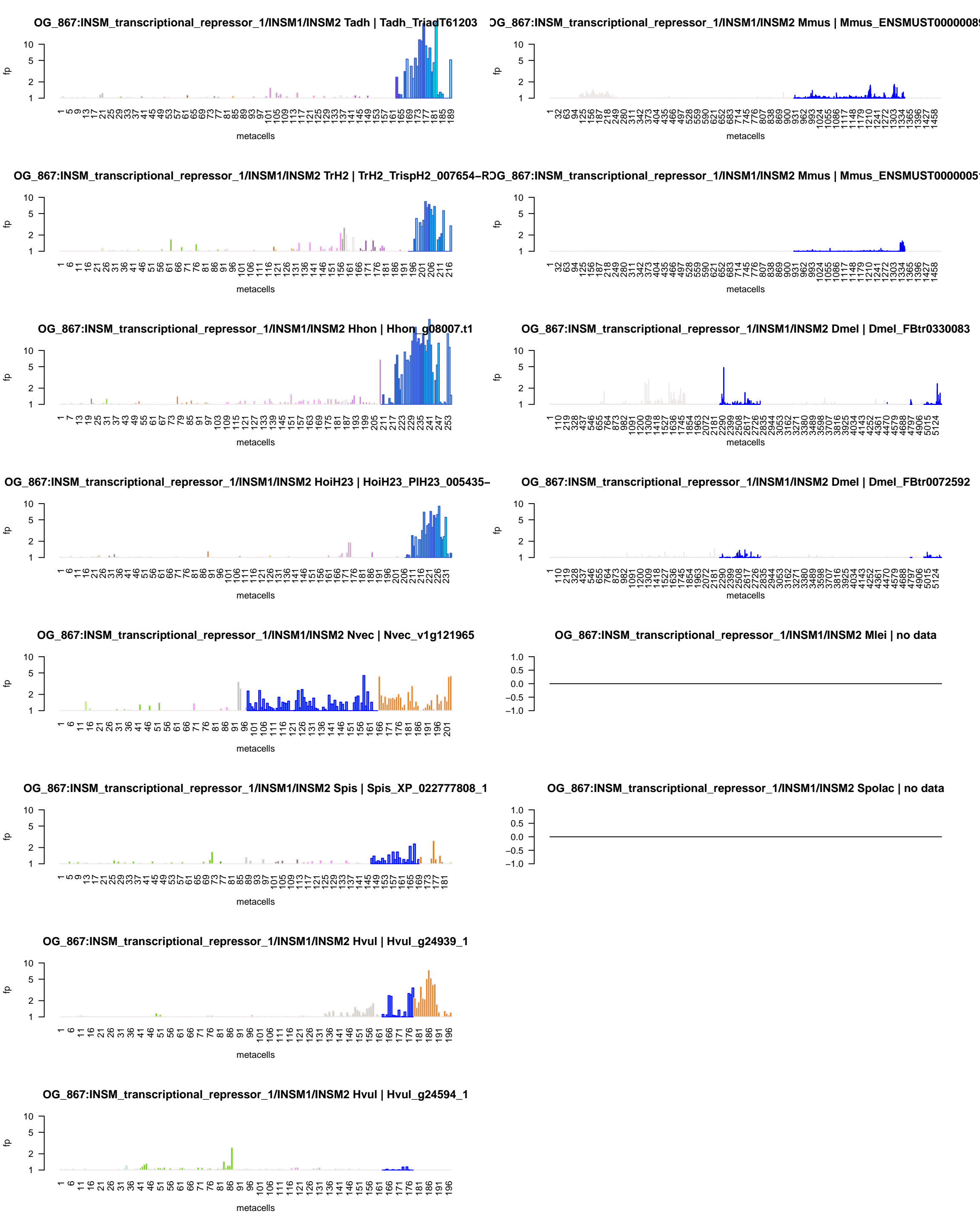


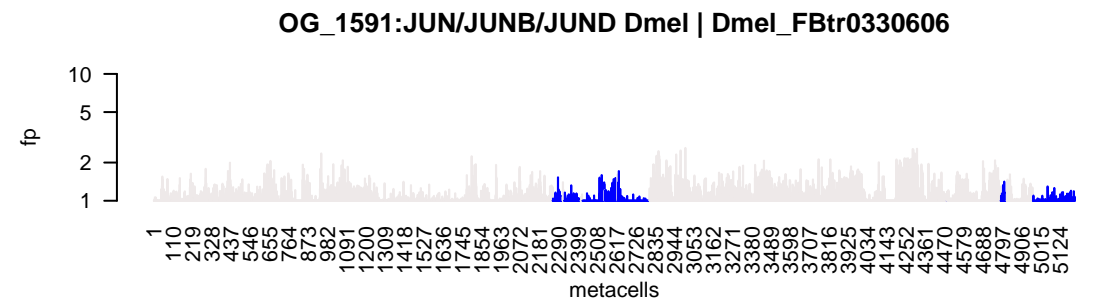
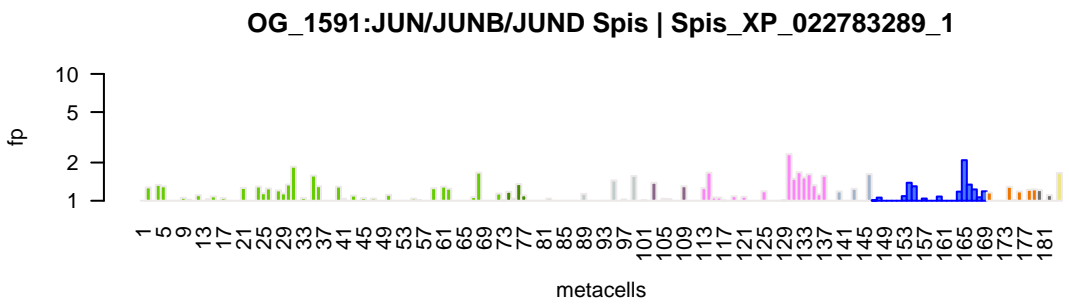
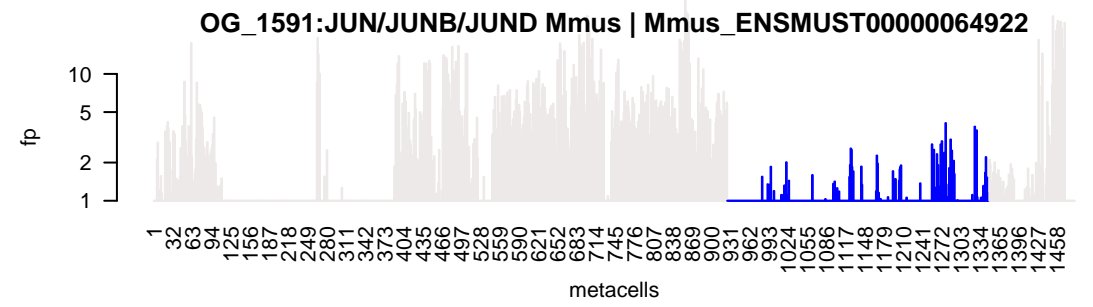
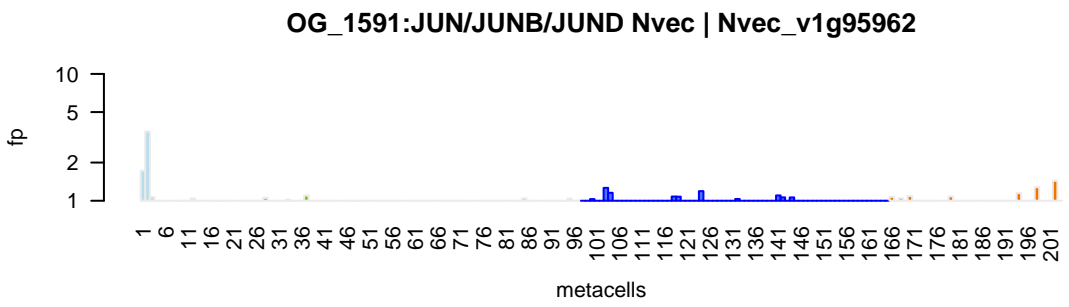
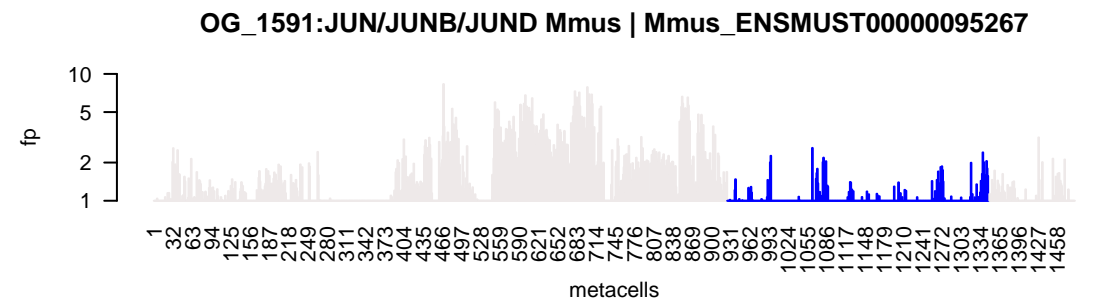
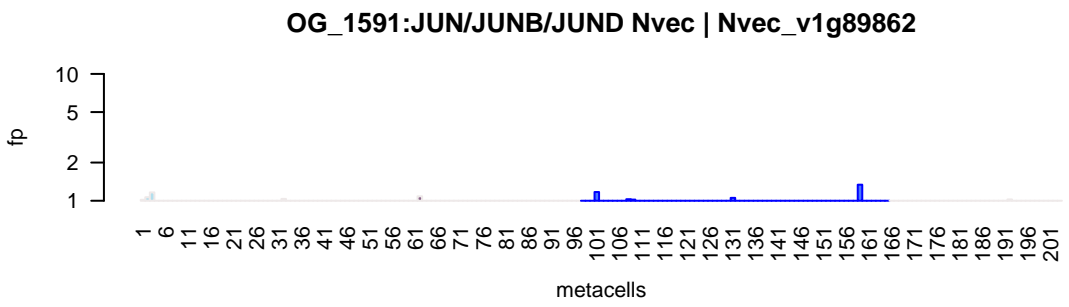
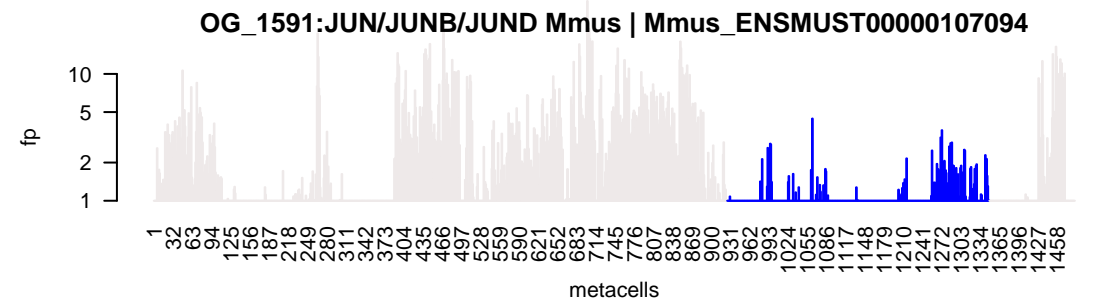
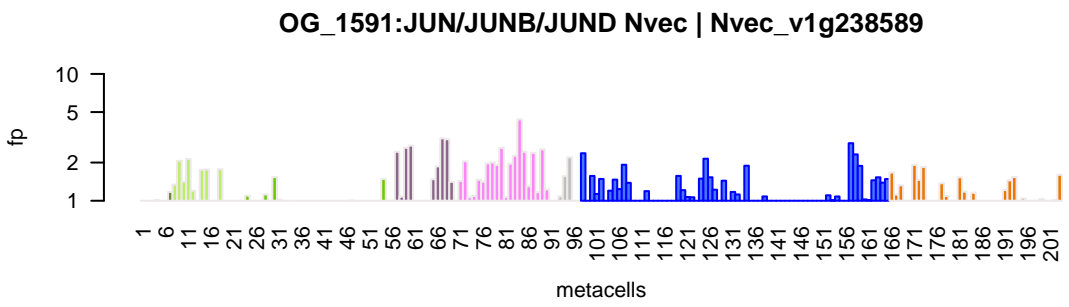
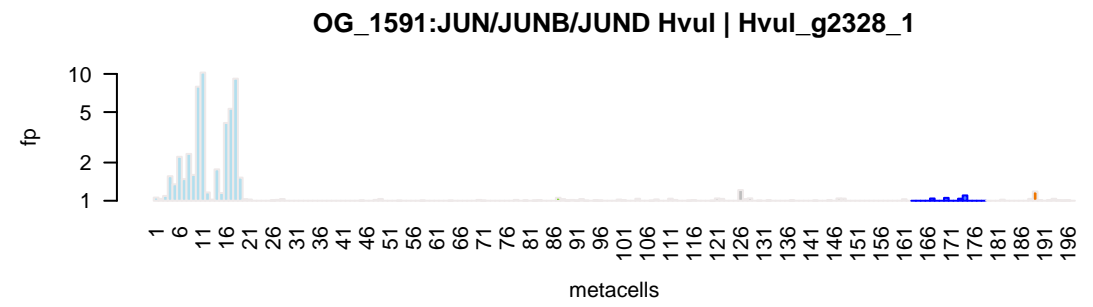
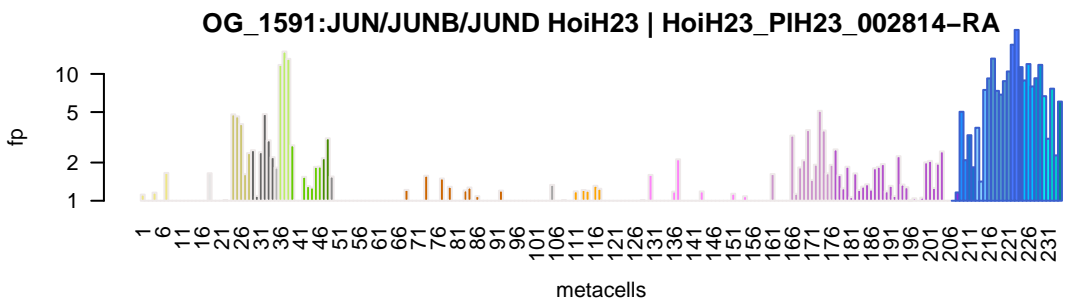
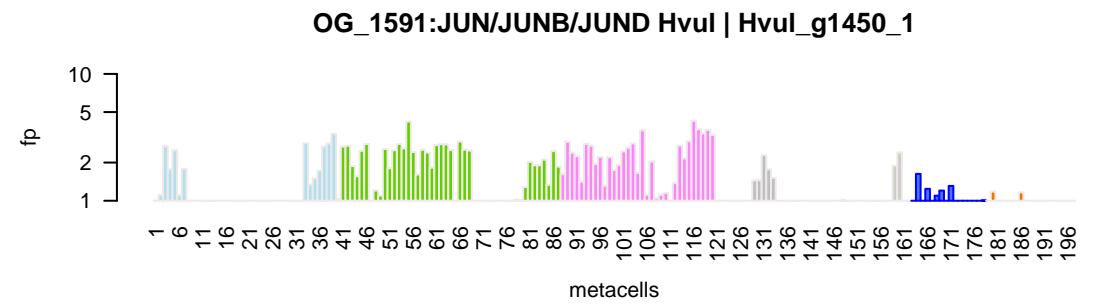
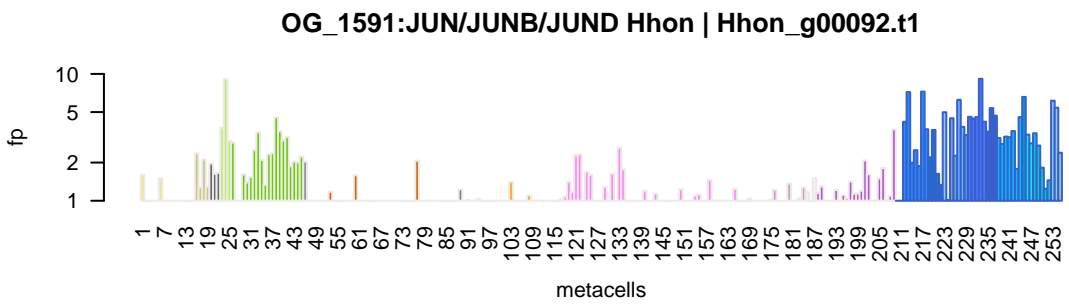
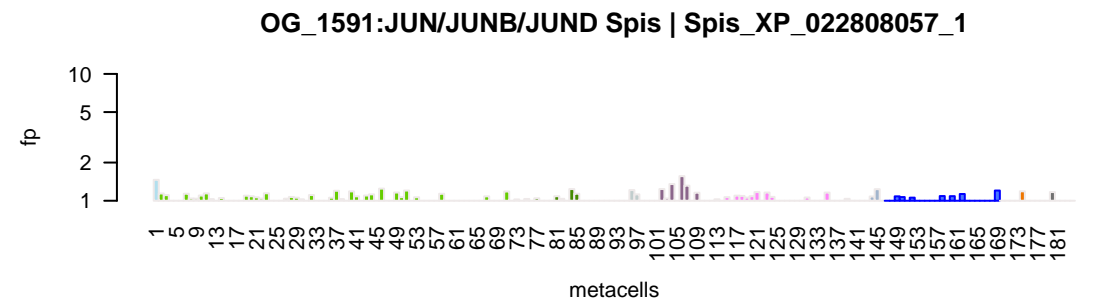
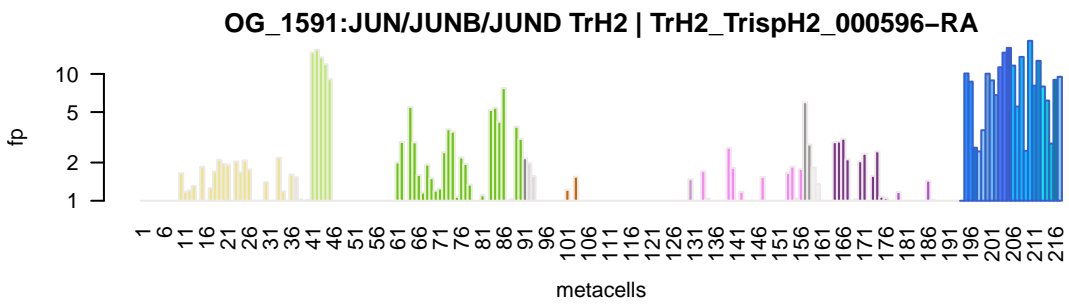
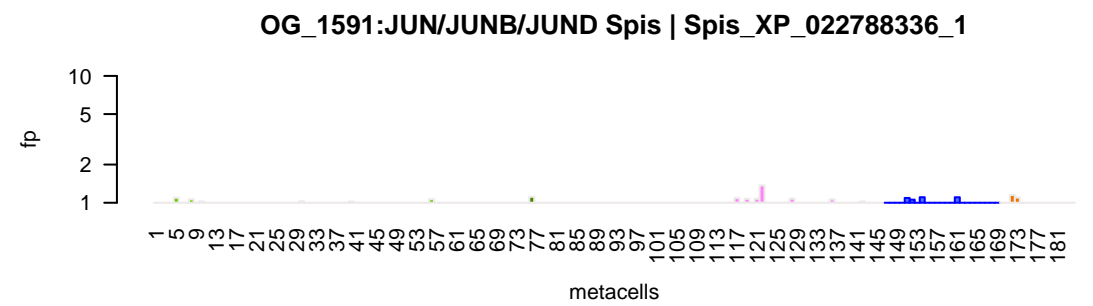
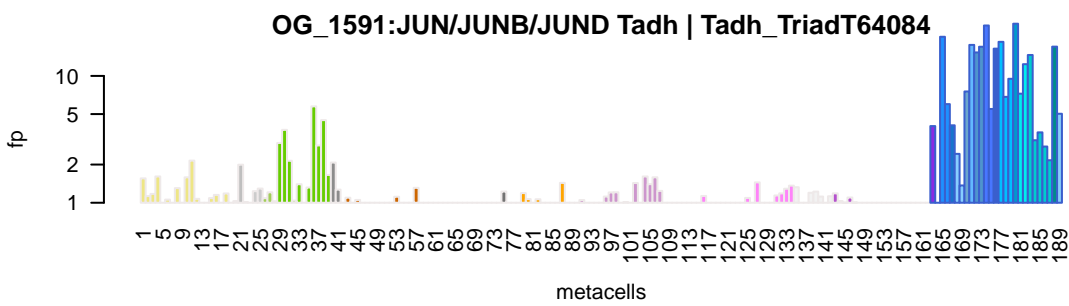
OG_3275:XPB1 Hvul | Hvul_g31026_1











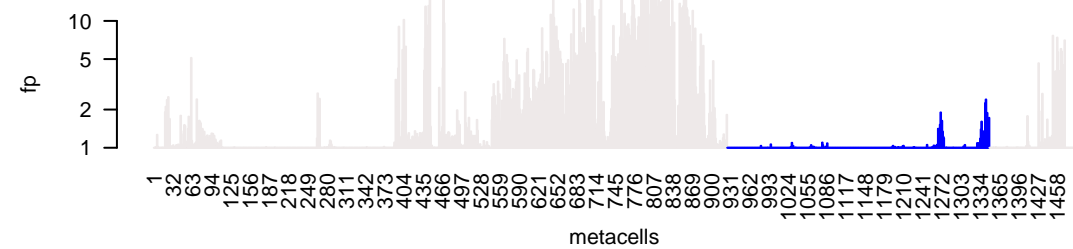
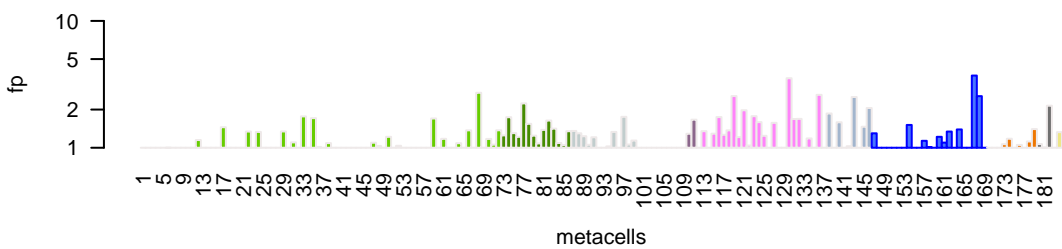
OG_1591:JUN/JUNB/JUND Mlei | no data



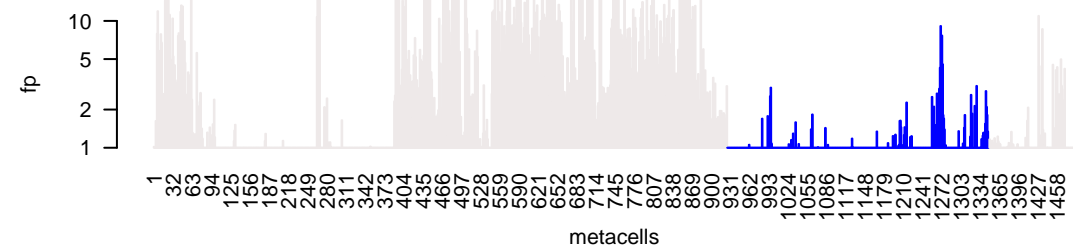
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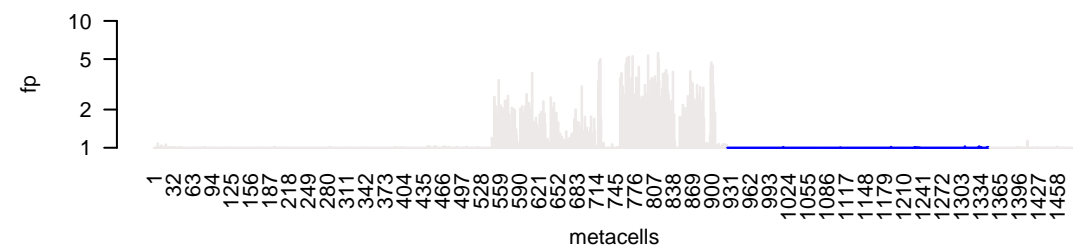
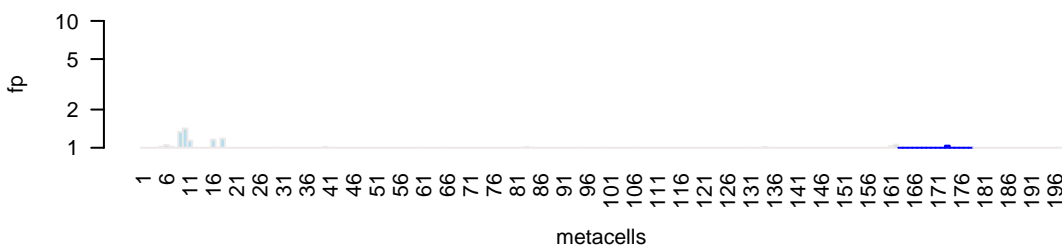
proto_oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 Spis | Spis.o oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 Mmus | Mmus.o



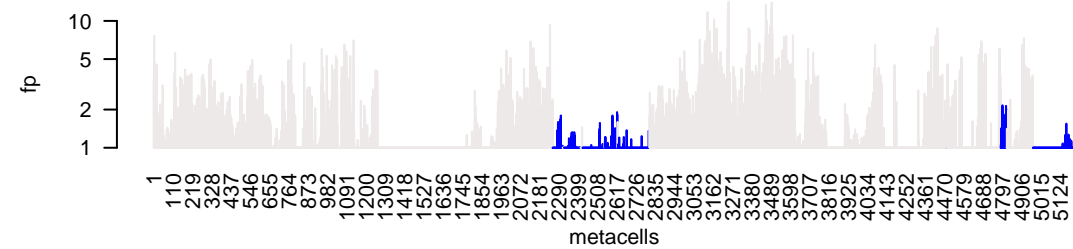
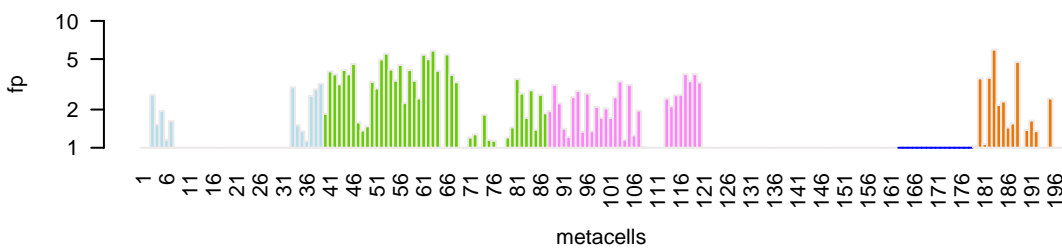
Fos_proto_oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 Hvu | fo_oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 Mmus | Mmus_1



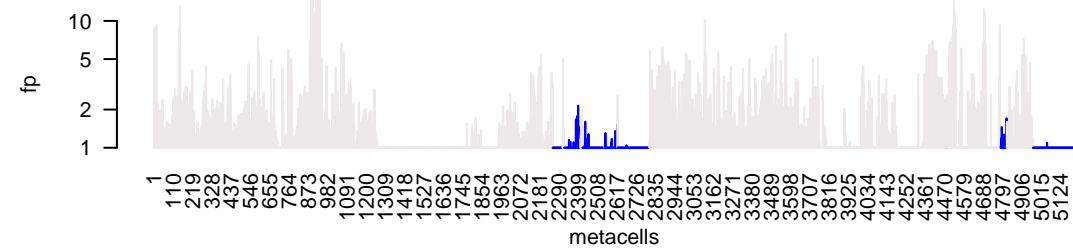
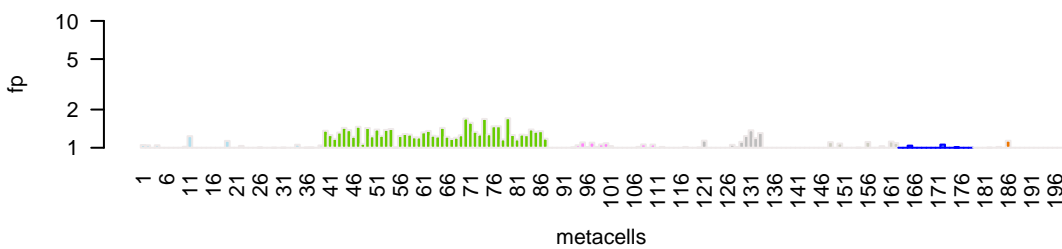
Fos_proto_oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 Hvu | fo_oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 Mmus | Mmus_1



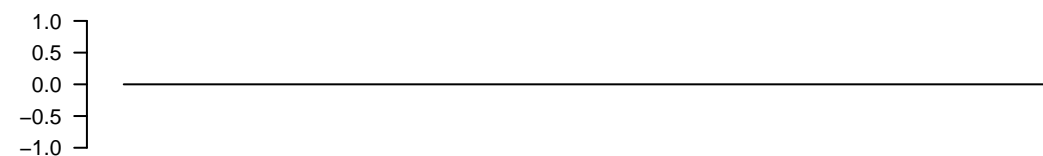
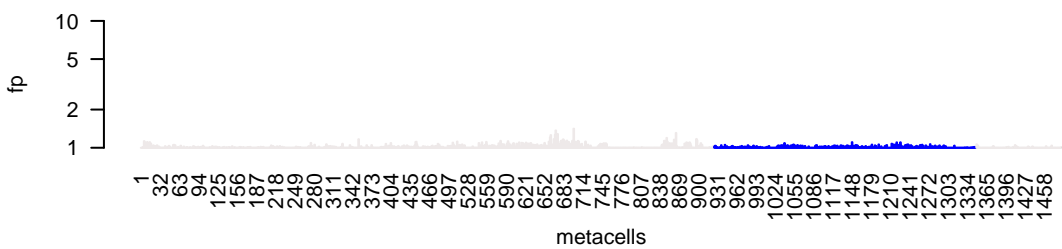
Fos_proto_oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 HvuI | Is_proto_oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 DmeI | Dr



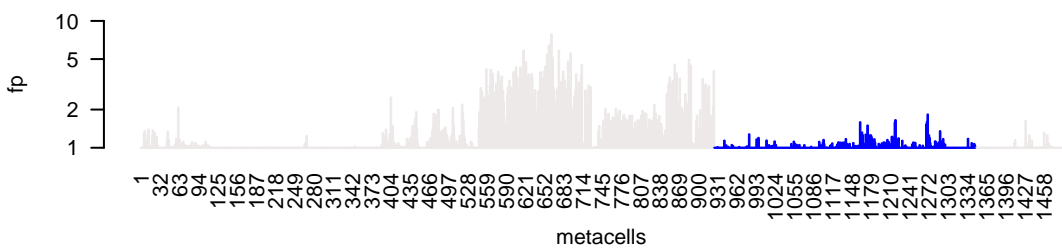
Fos_proto_oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 HvuI | Is_proto_oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 DmeI | Dr



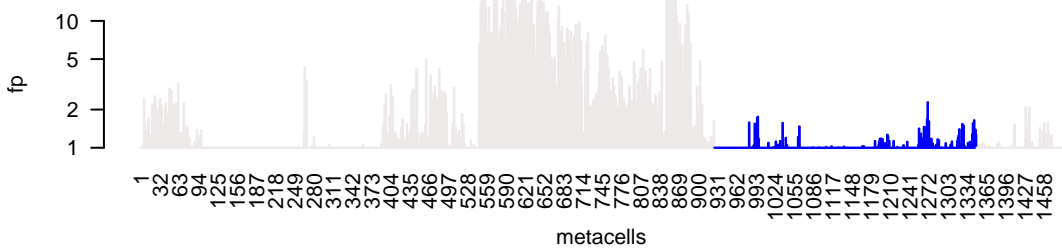
o oncogene AP_1 transcription factor subunit/FOS/FOSB/FOSL1/FOSL2 Mmus | Mmus.155:Fos_proto oncogene AP_1 transcription factor subunit/FOS/FOSB/FOSL1/FOSL2 Mle



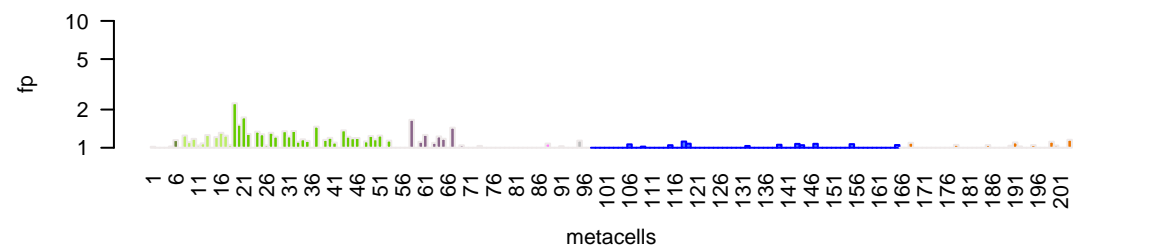
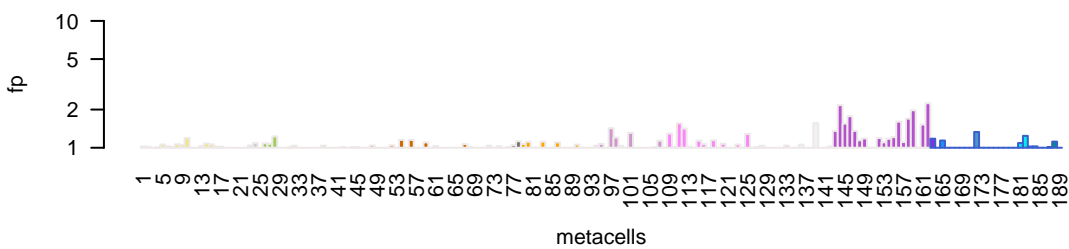
o oncogene AP_1 transcription factor subunit/FOS/FOSB/FOSL1/FOSL2 Mmus | Mmus_l5:Fos_proto oncogene AP_1 transcription factor subunit/FOS/FOSB/FOSL1/FOSL2 Spol



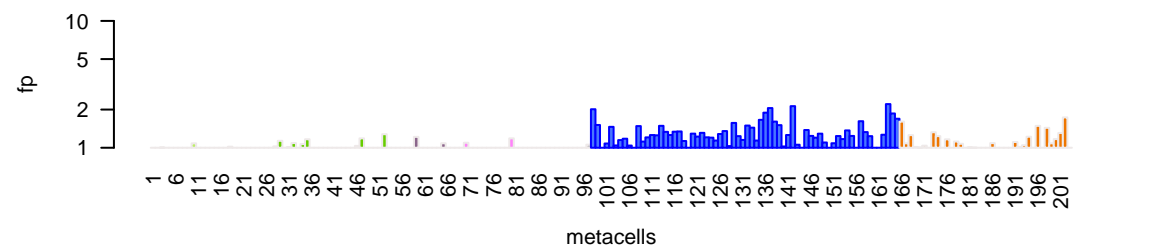
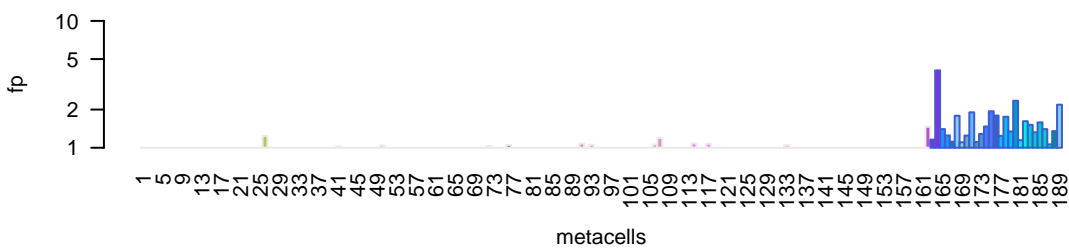
o oncogene_AP_1_transcription_factor_subunit/FOS/FOSB/FOSL1/FOSL2 Mmus | Mmus_I



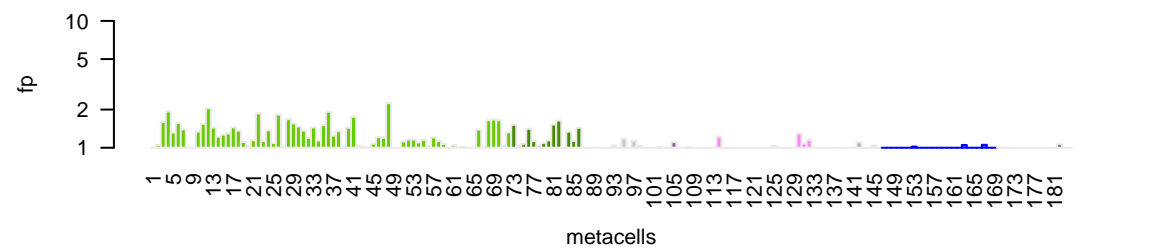
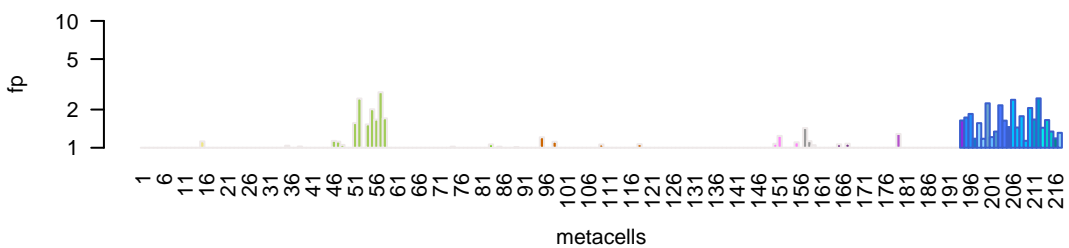
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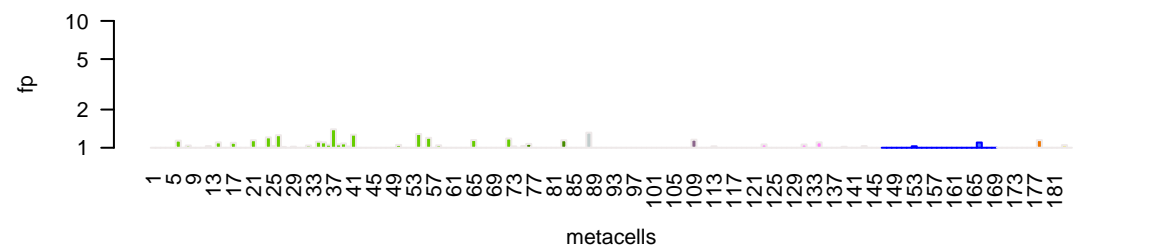
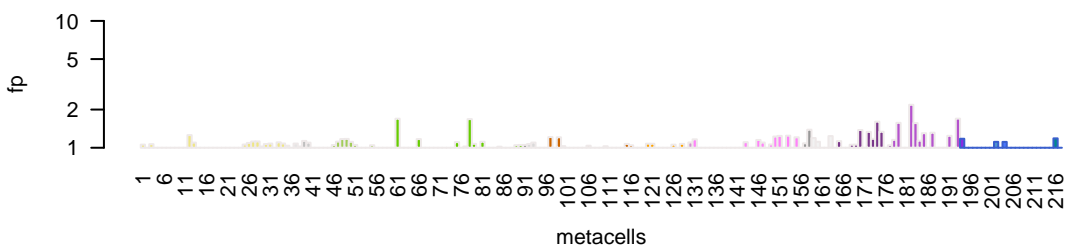
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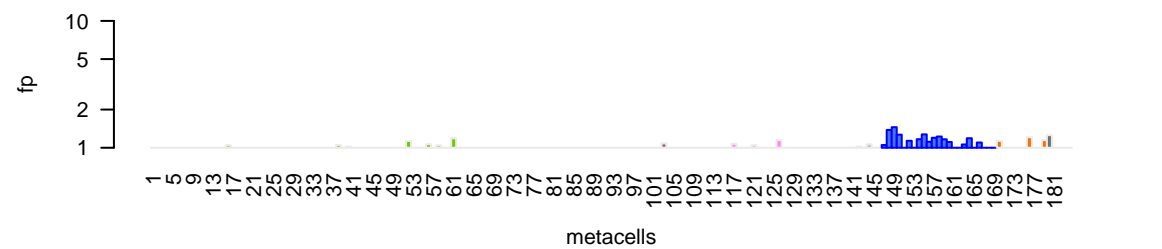
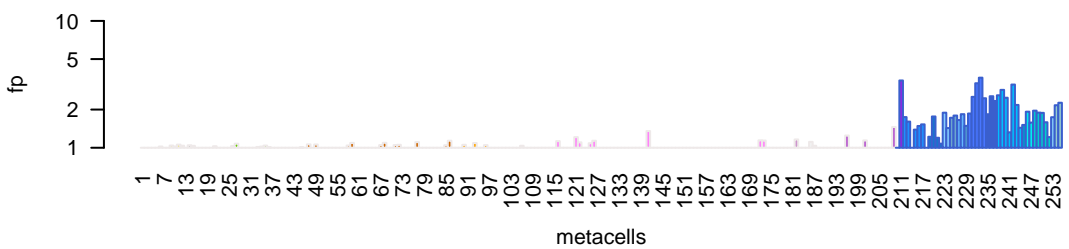
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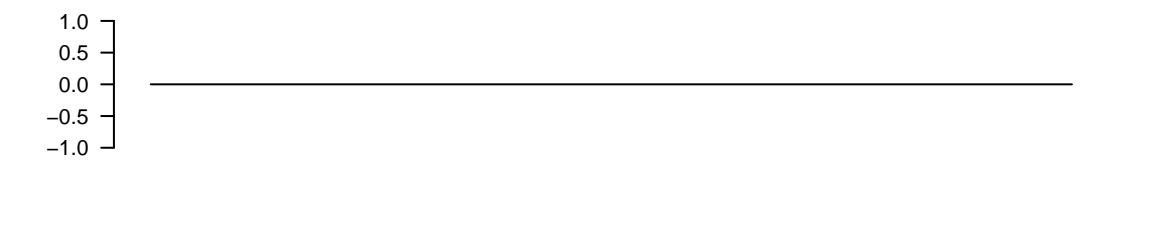
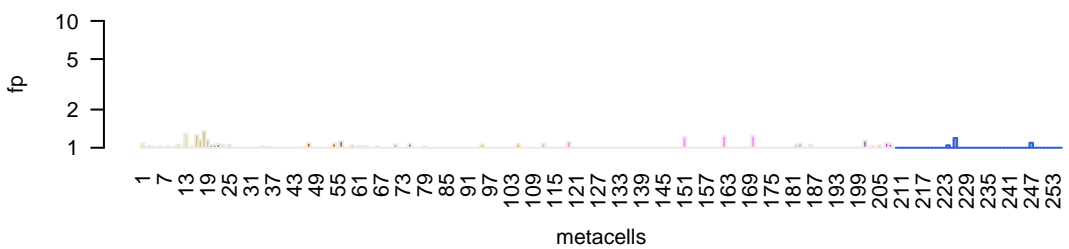
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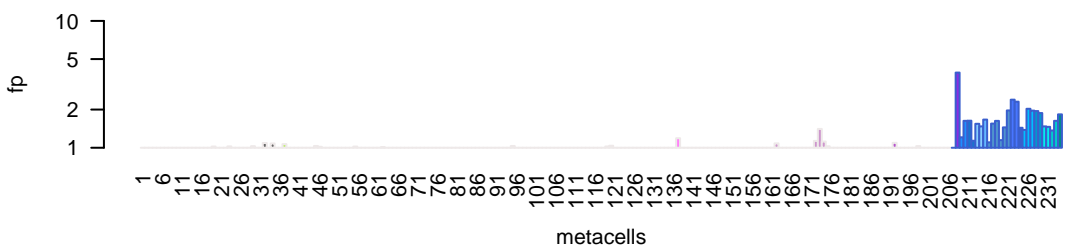
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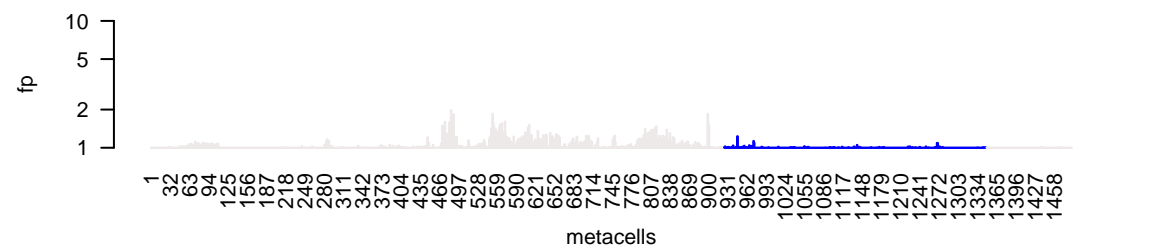
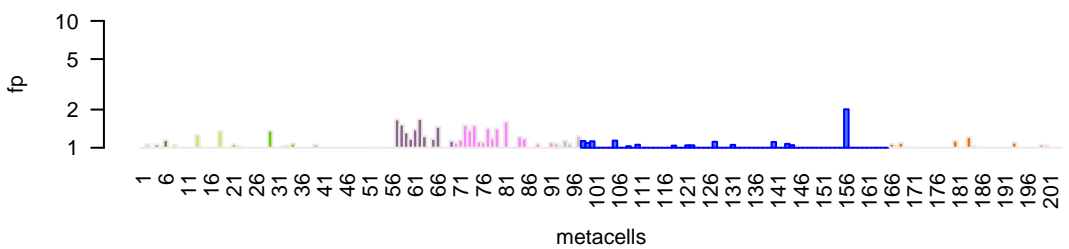
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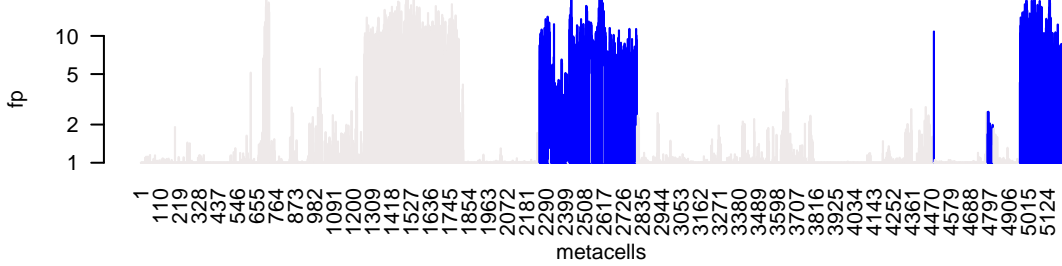
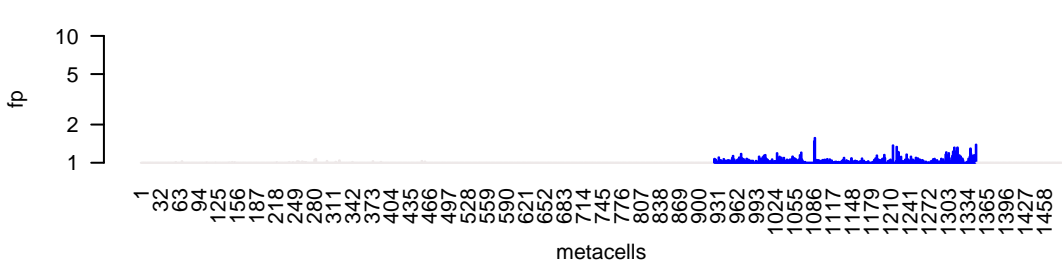
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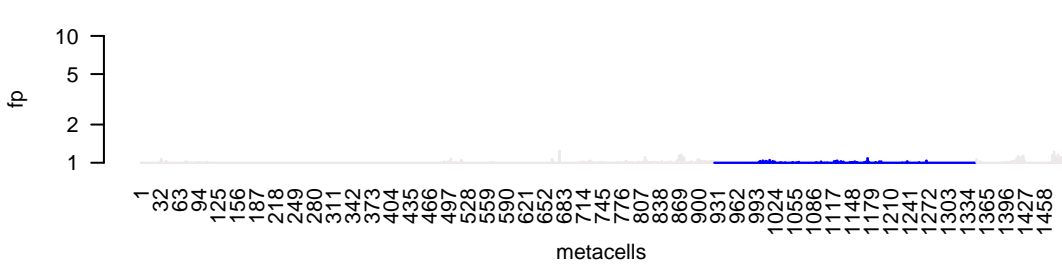
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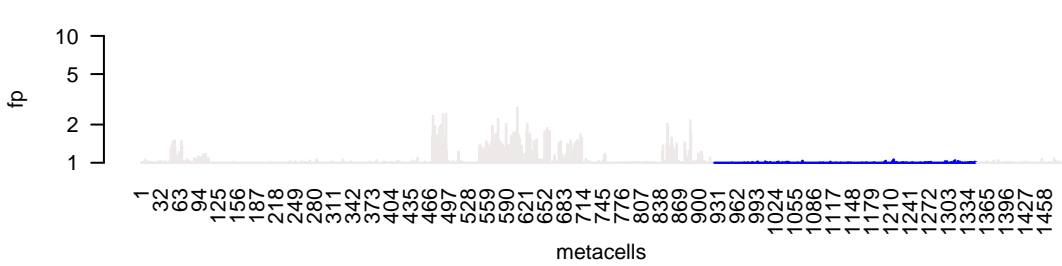
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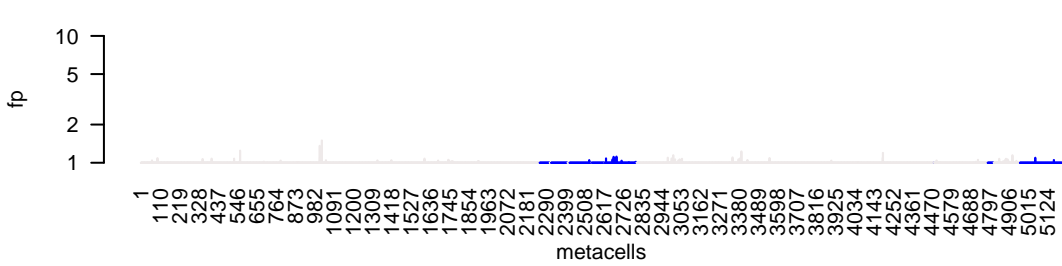
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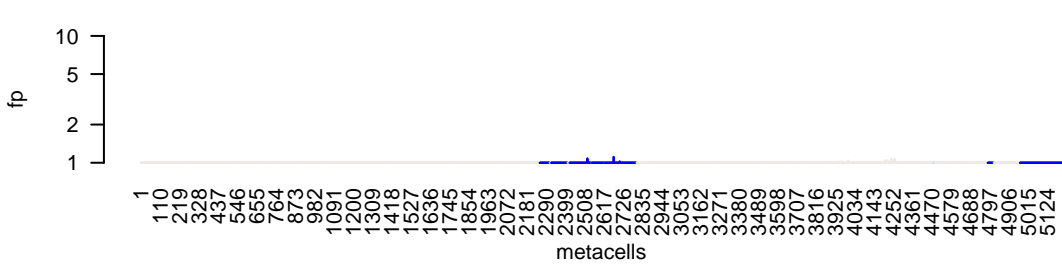
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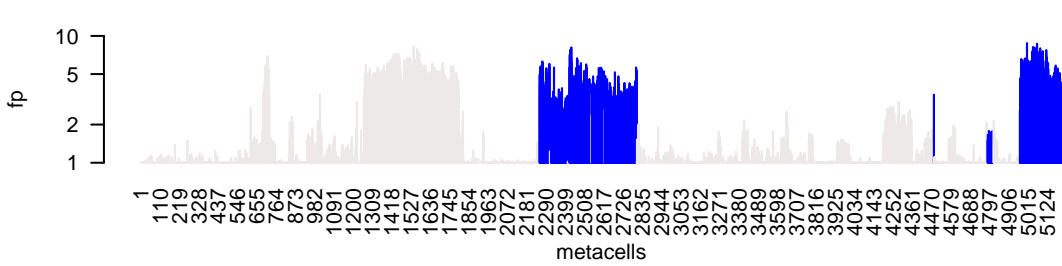
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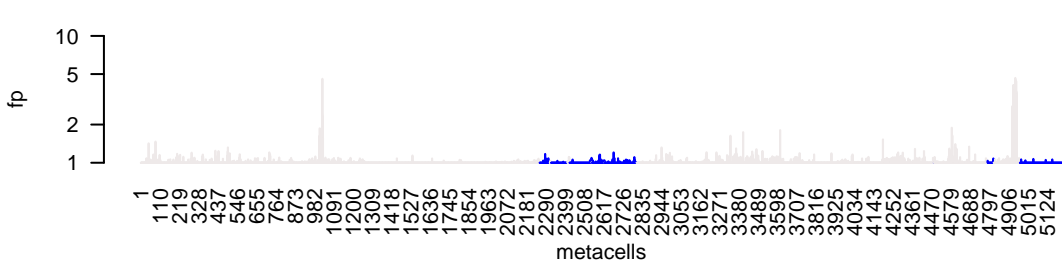
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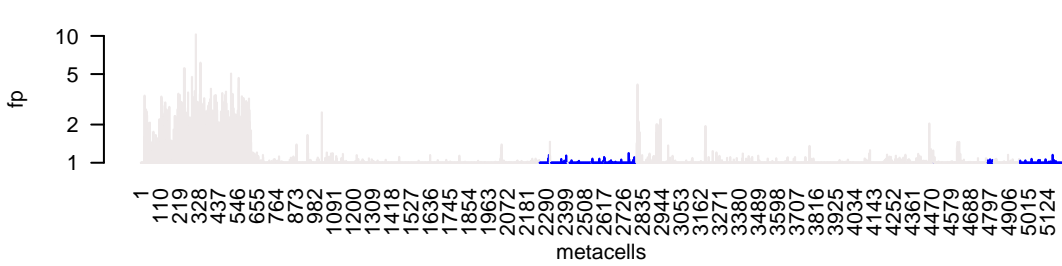
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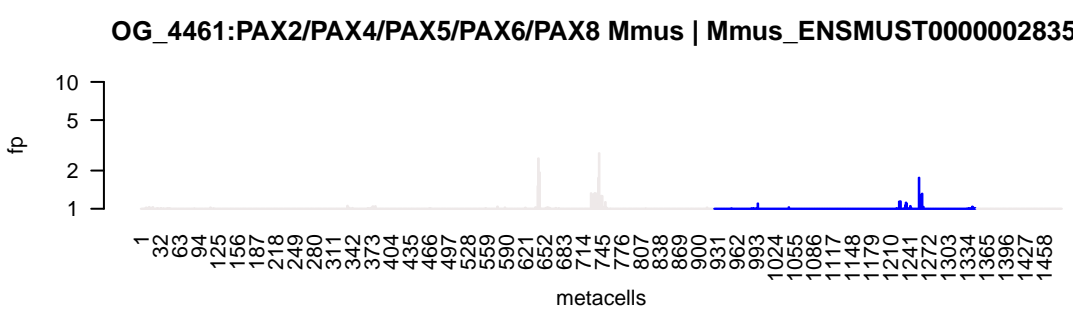
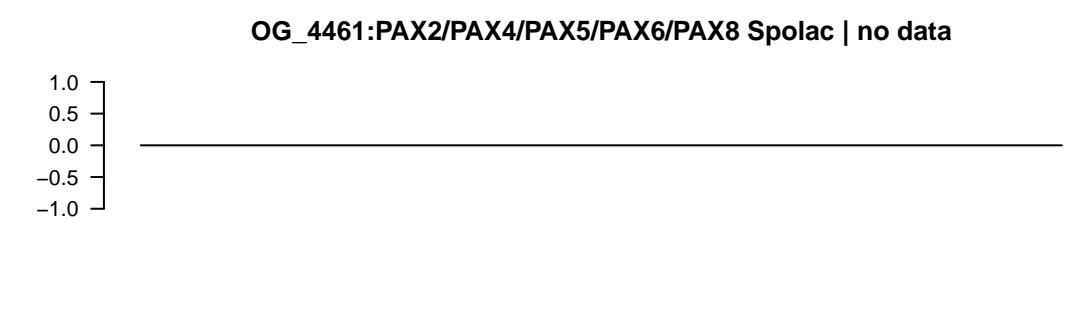
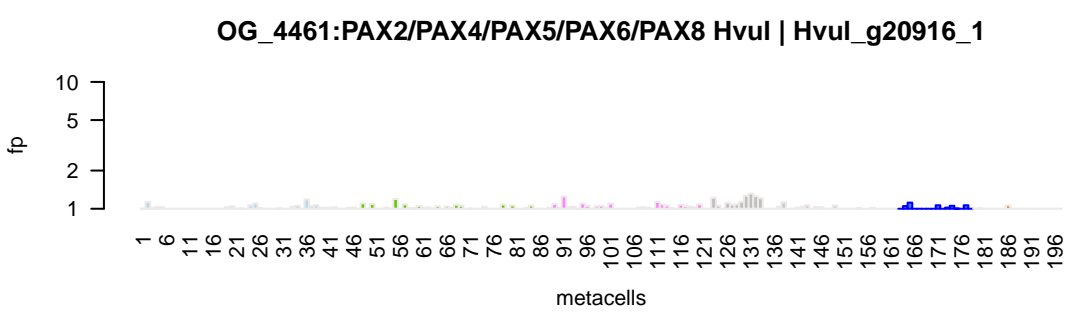
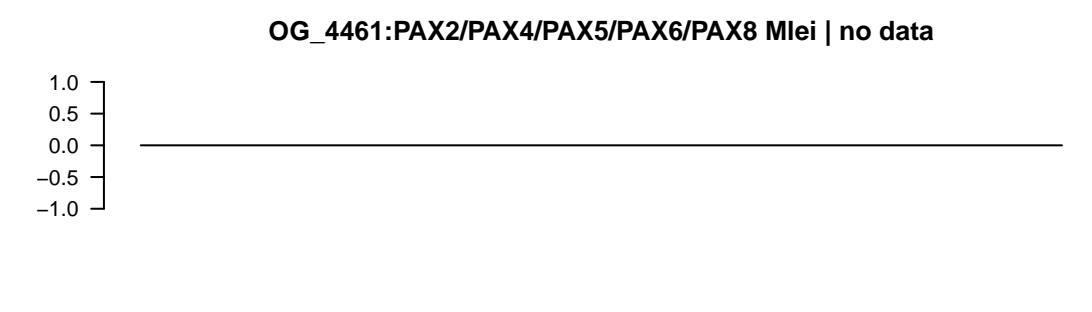
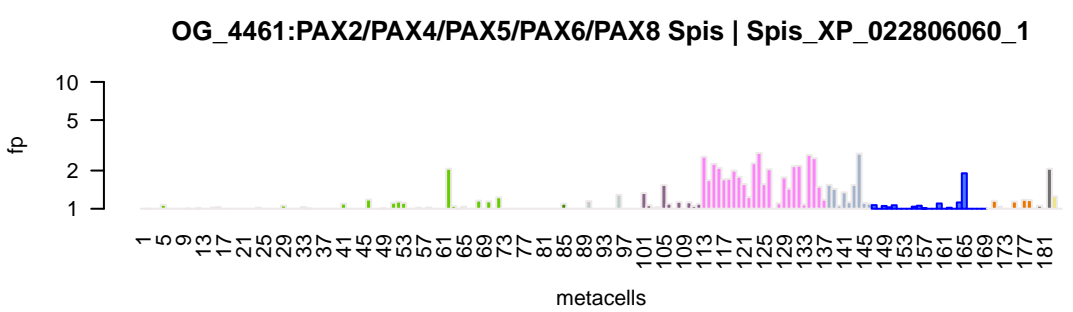
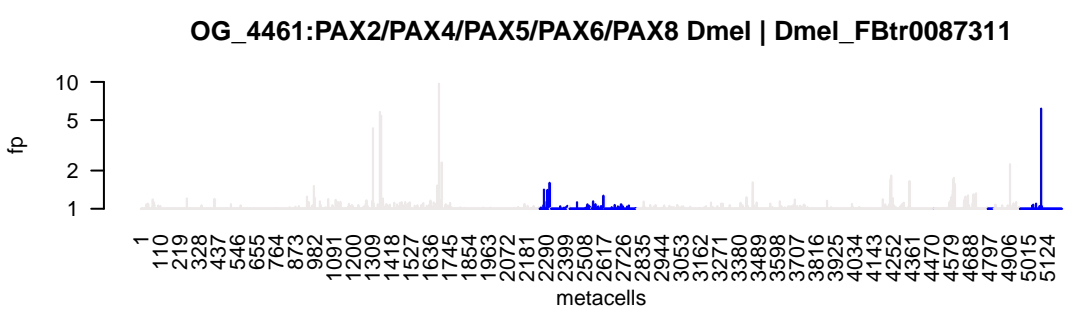
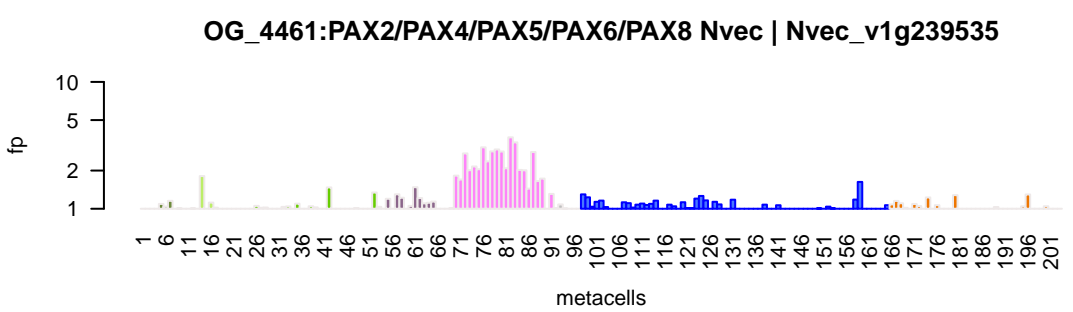
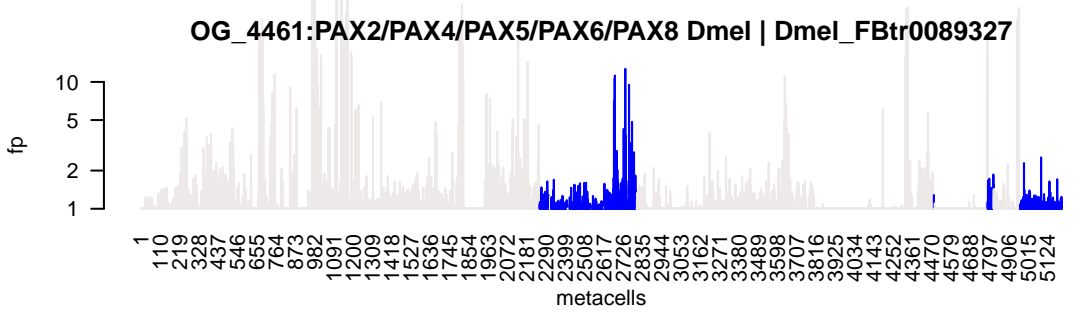
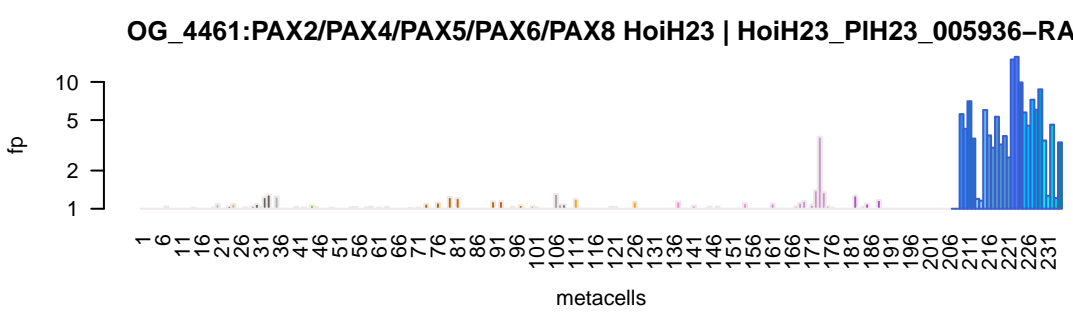
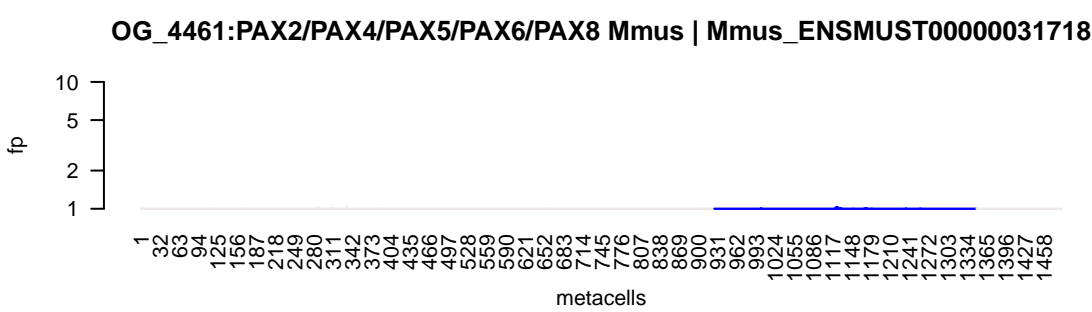
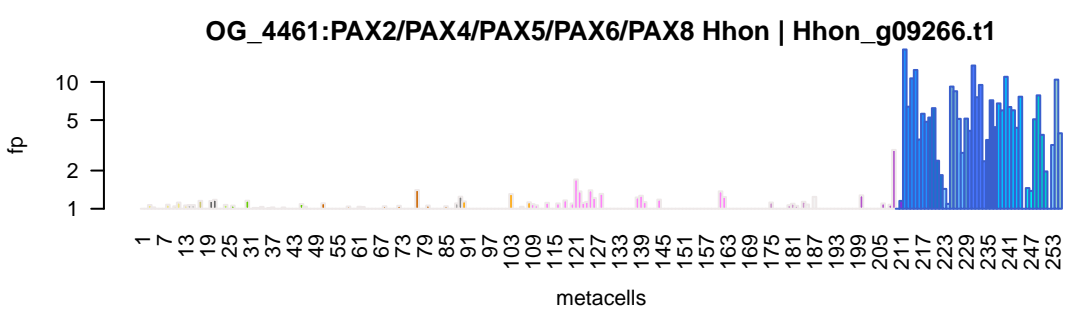
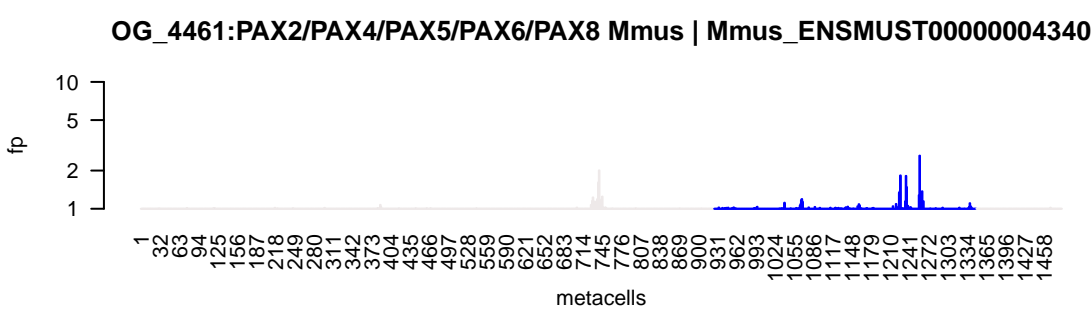
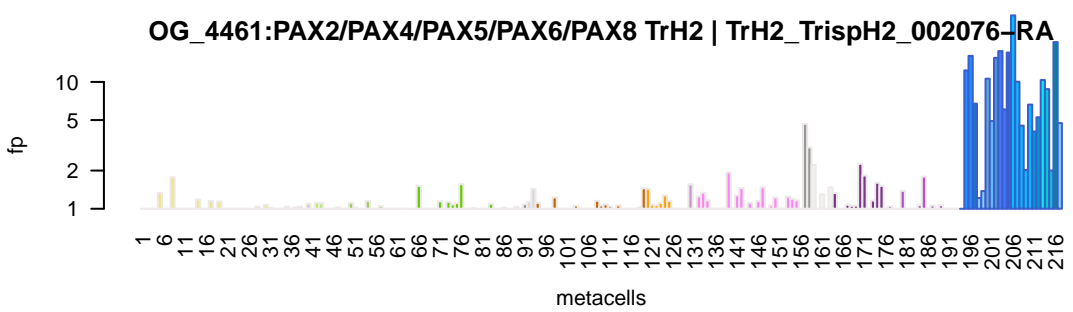
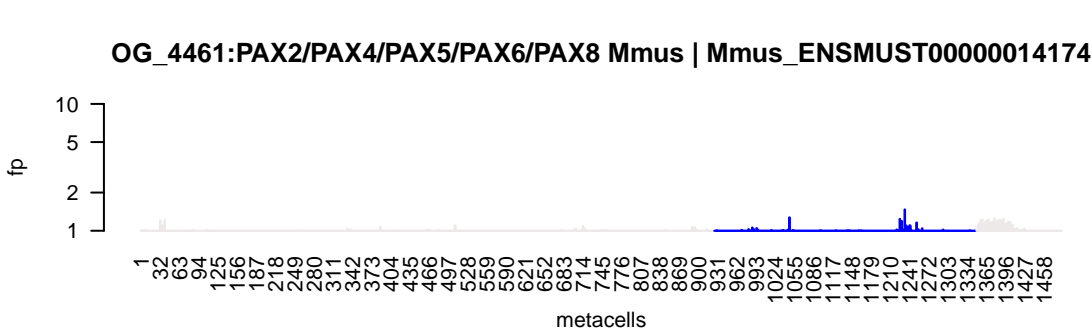
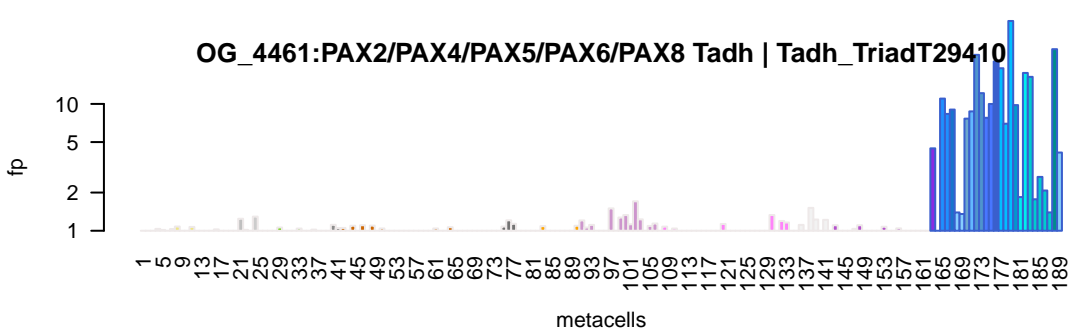


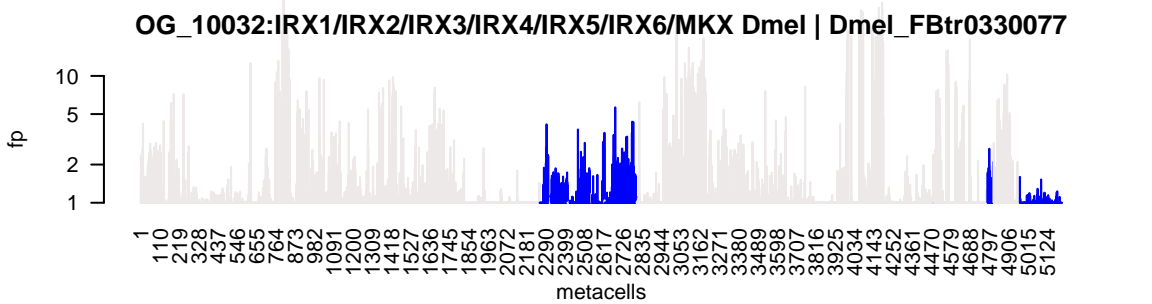
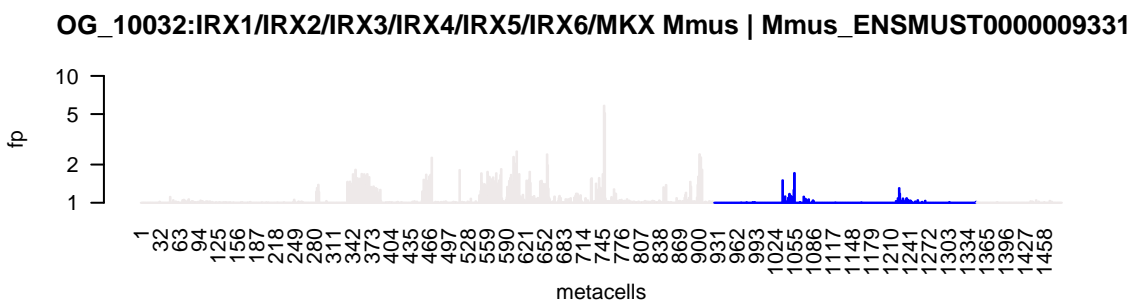
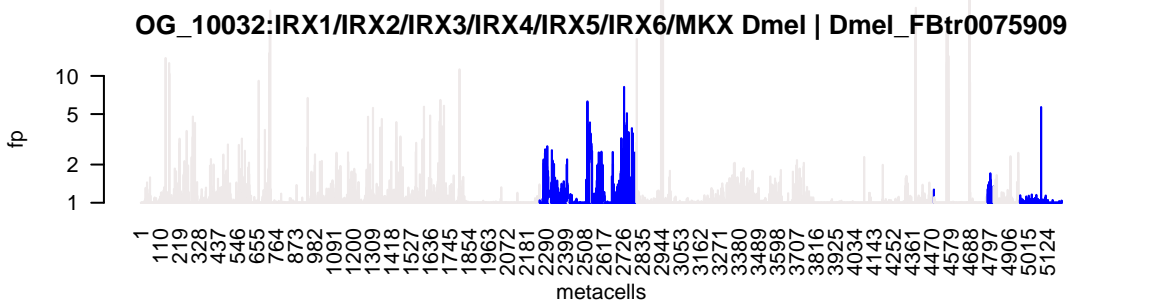
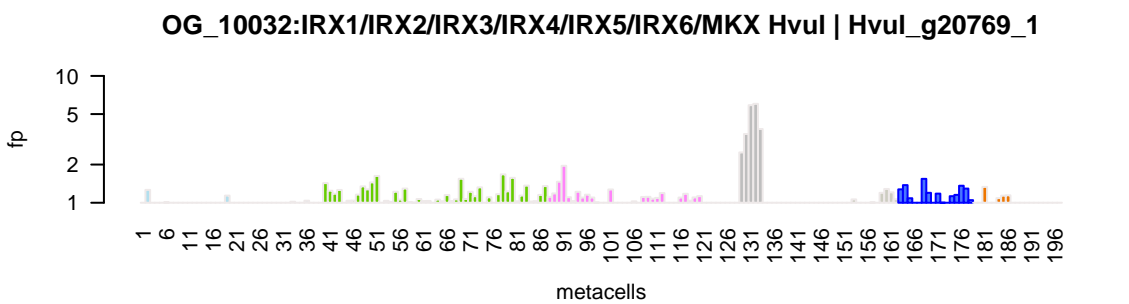
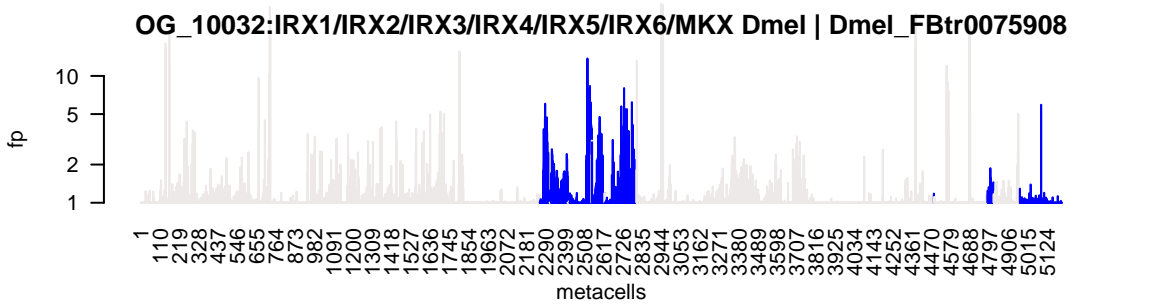
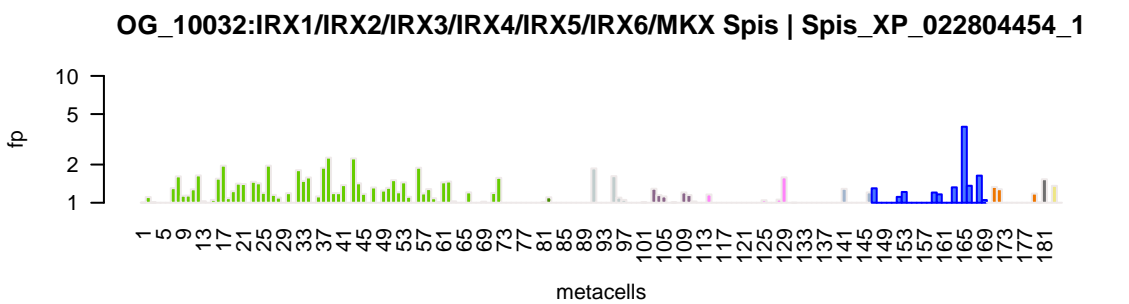
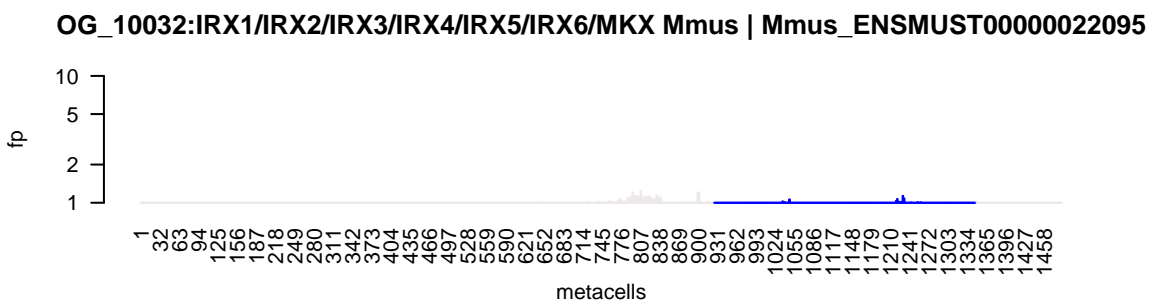
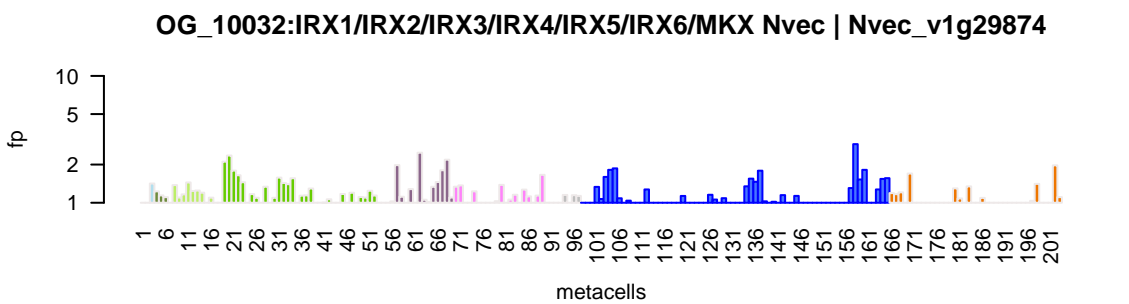
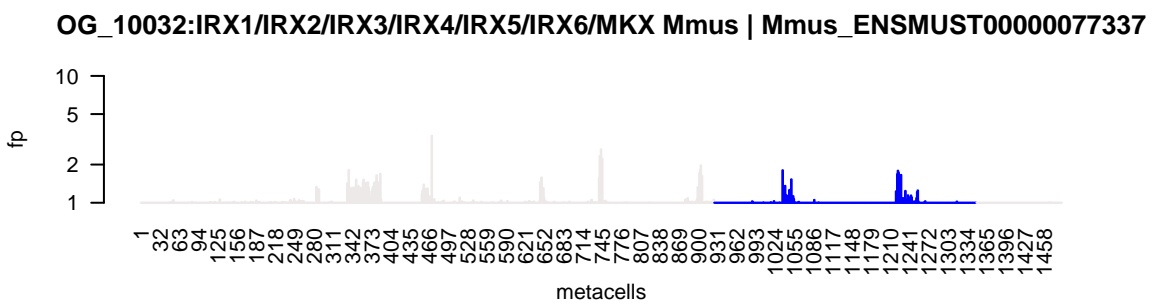
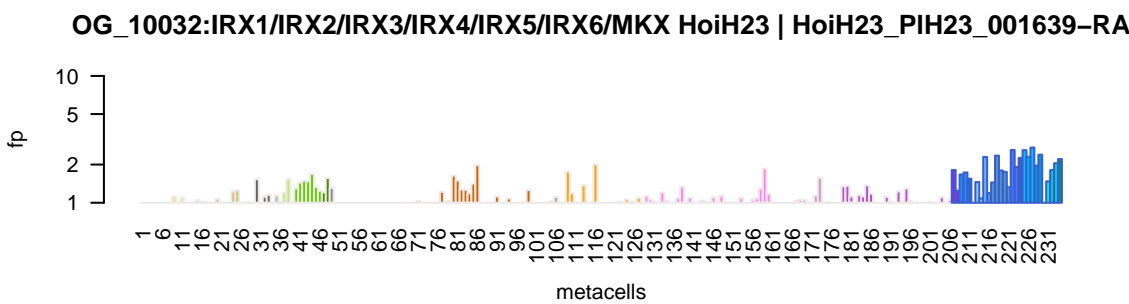
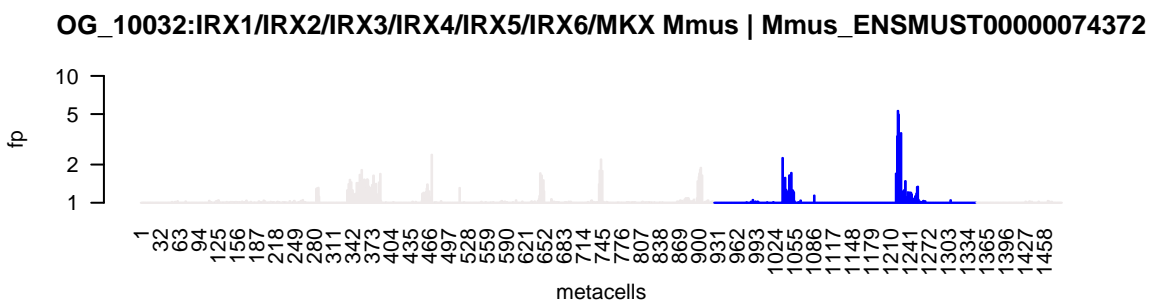
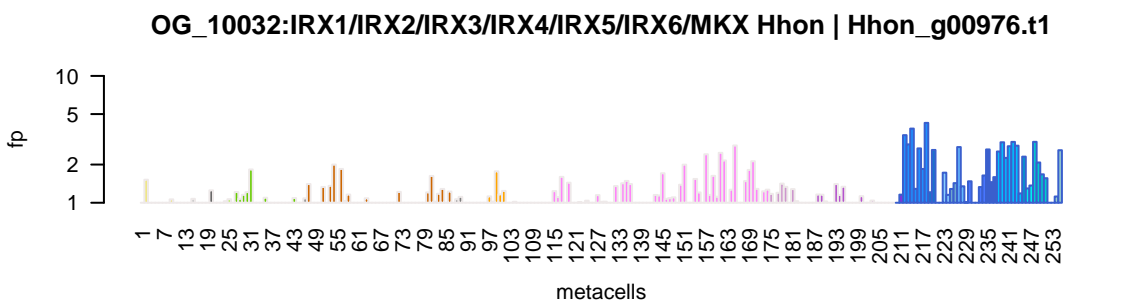
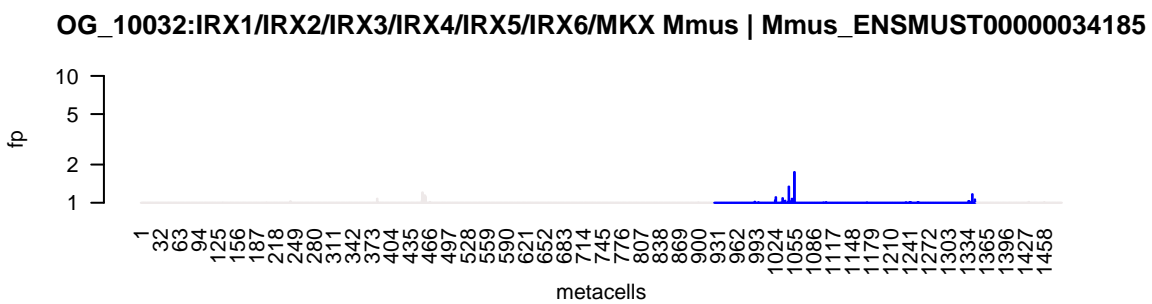
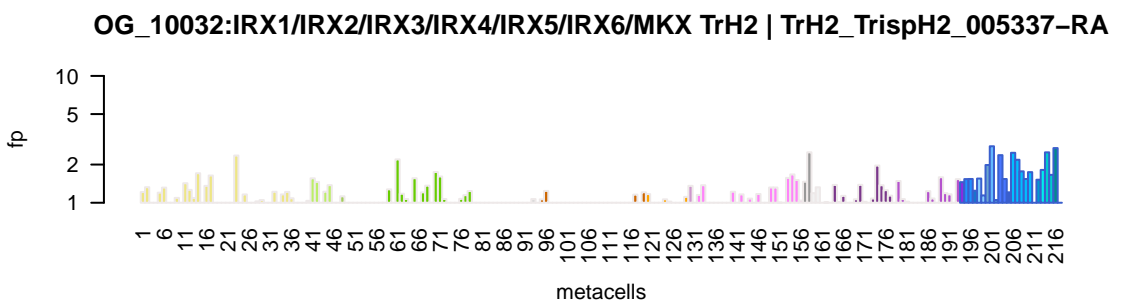
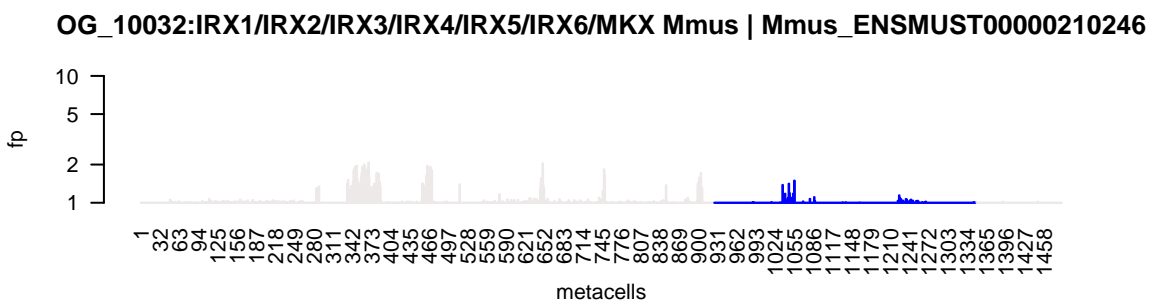
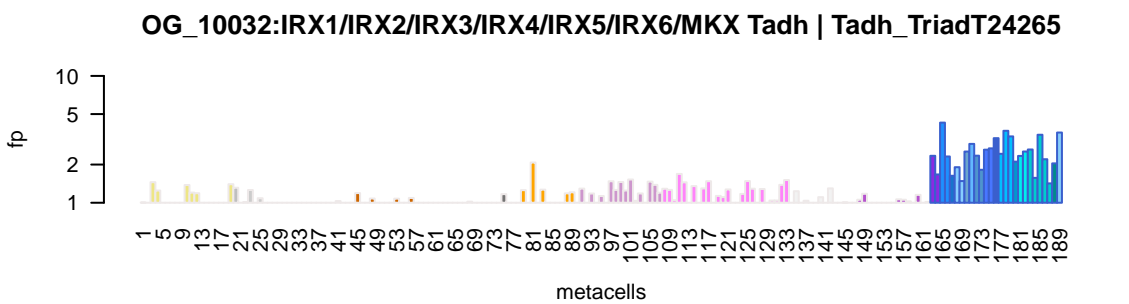
T2/SNAI1/SNAI2/SNAI3/ZNF526/SCRT1/SCRT2/SNAI1/SNAI2/SNAI3/ZNF526;zf-met.HG1.40 I



T2/SNAI1/SNAI2/SNAI3/ZNF526/SCRT1/SCRT2/SNAI1/SNAI2/SNAI3/ZNF526;zf-met.HG1.40 I







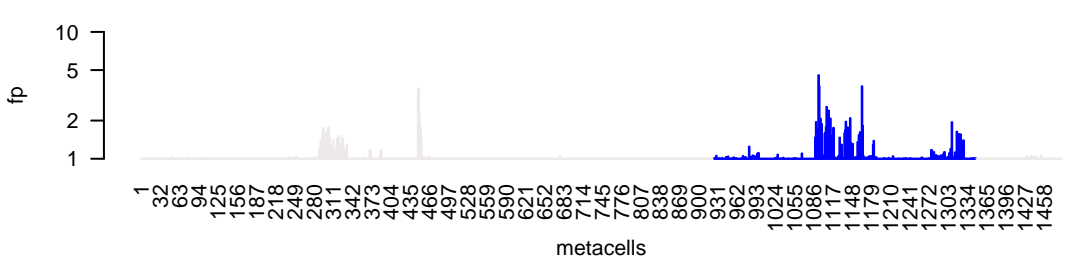
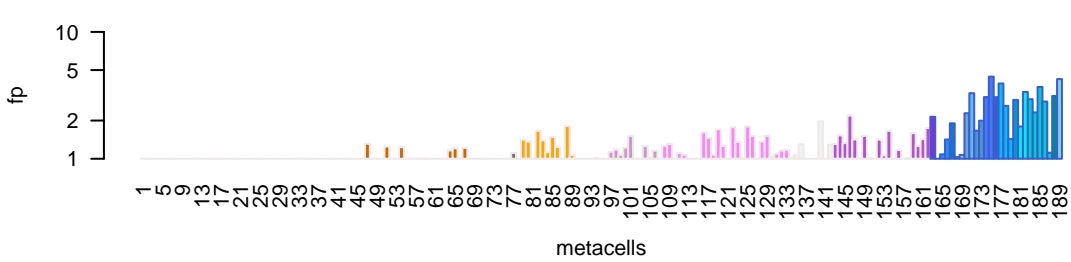
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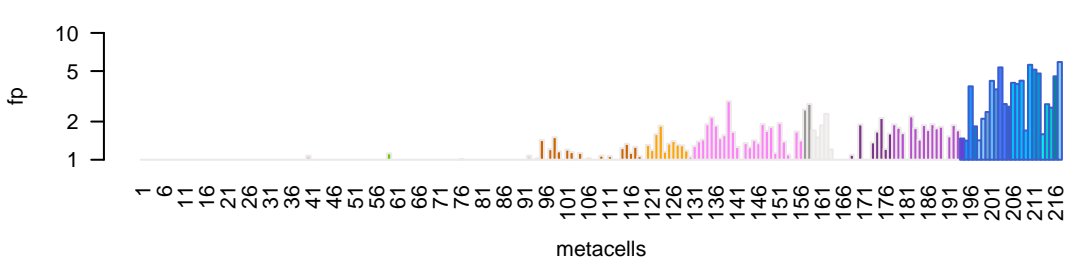
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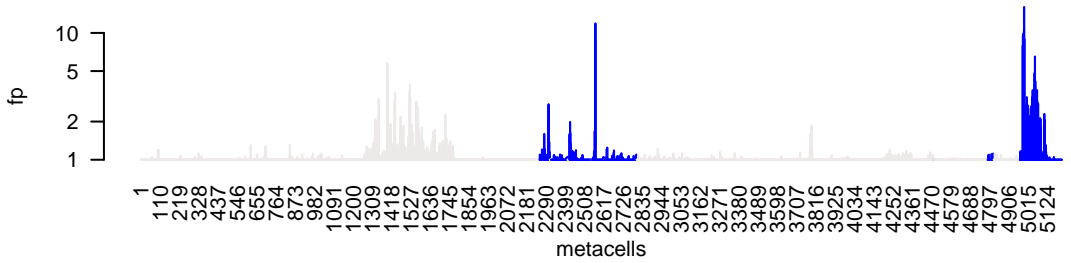
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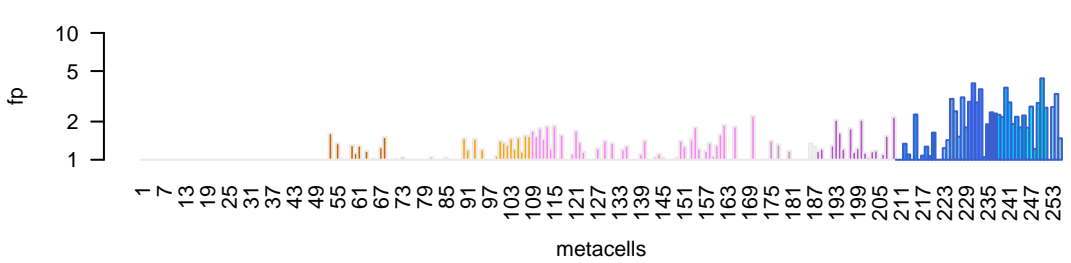
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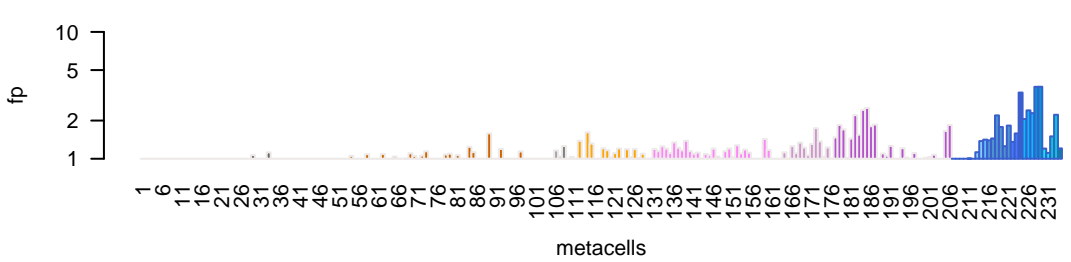
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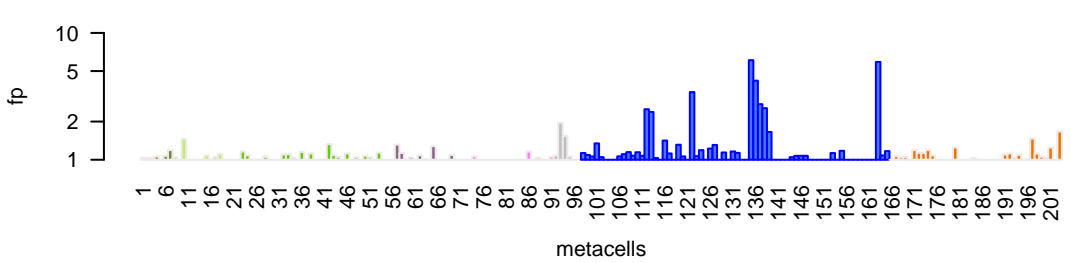
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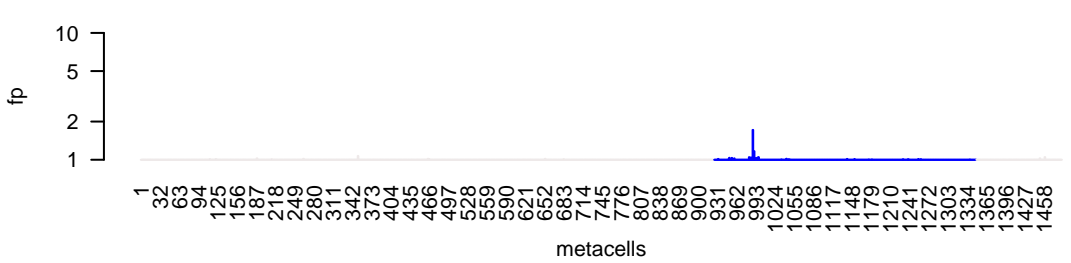
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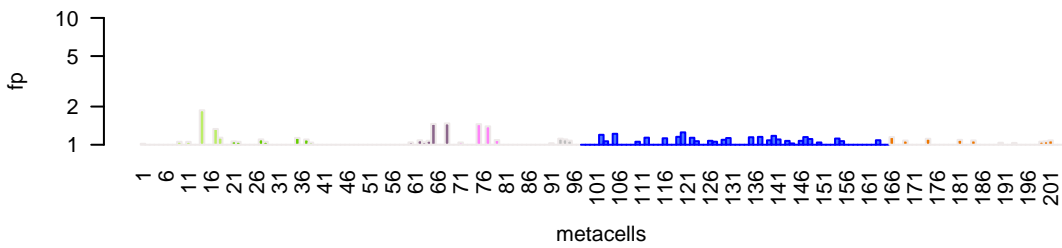
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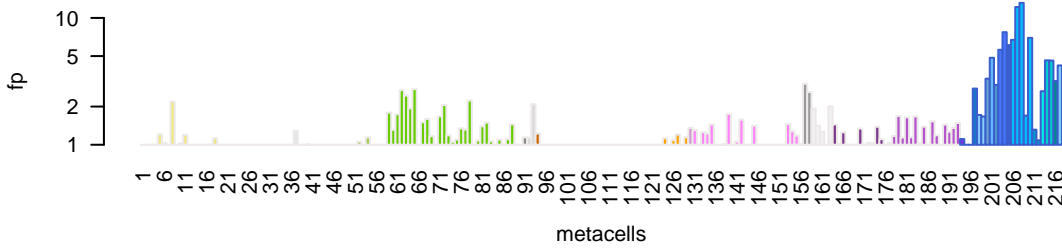
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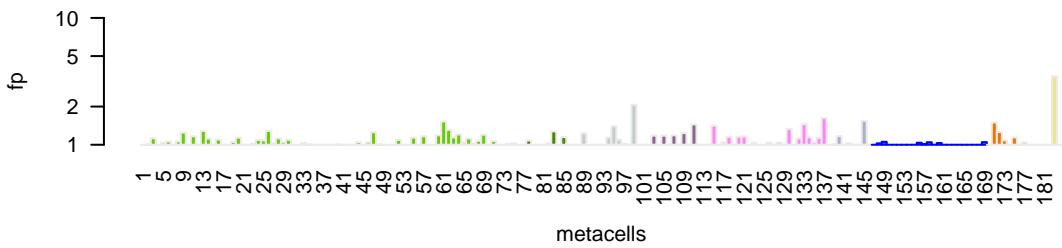
OG_3081:IRF1/IRF2 Nvec | Nvec_v1g86955



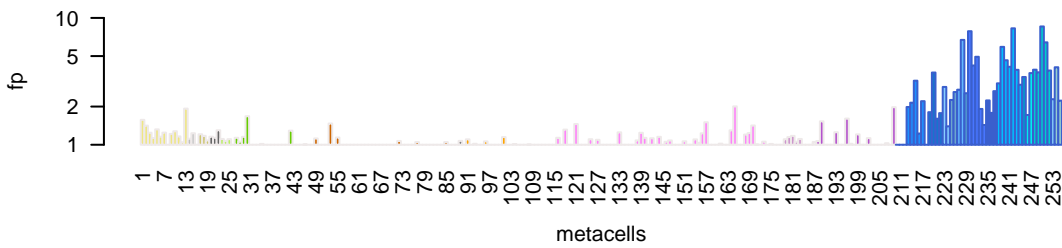
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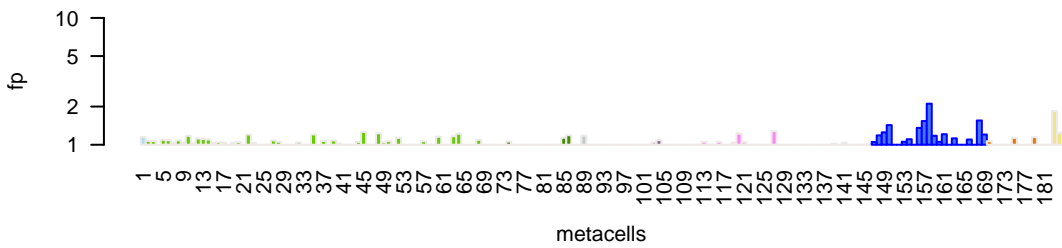
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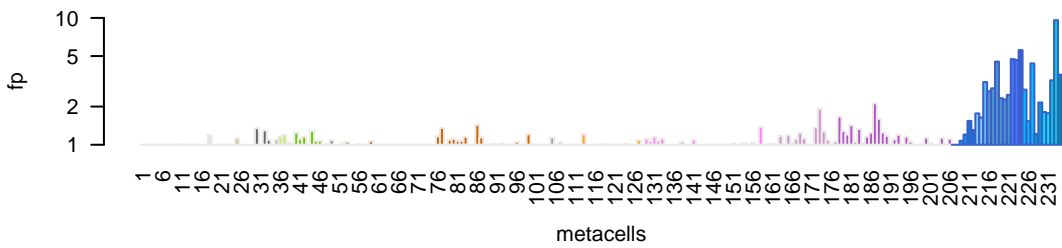
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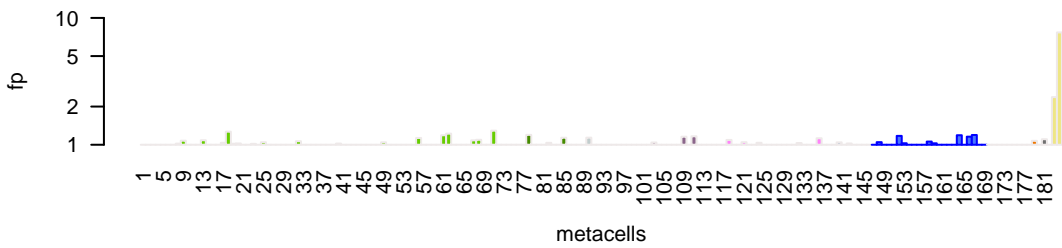
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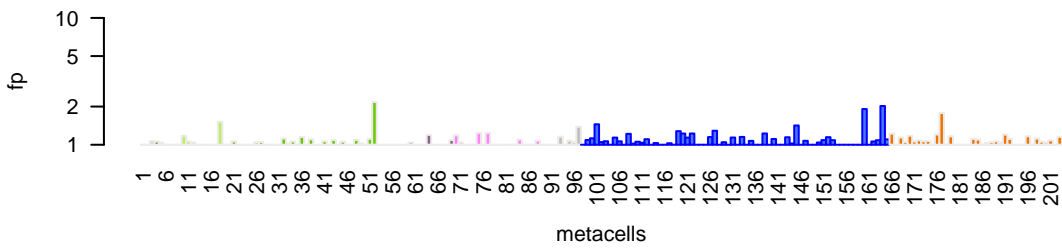
OG_3081:IRF1/IRF2 HoiH23 | HoiH23_PIH23_008602-RA



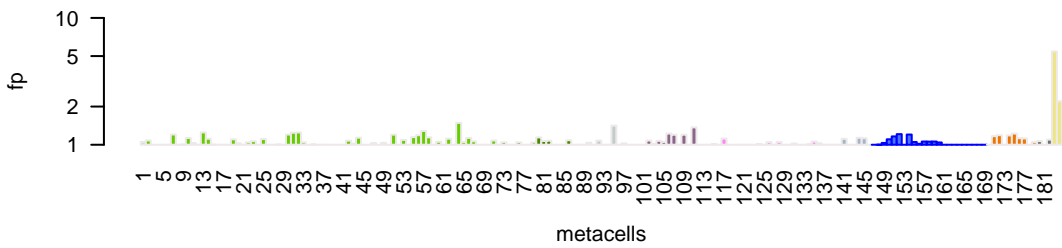
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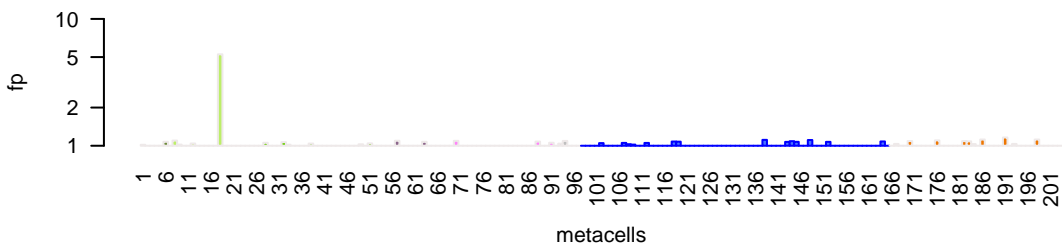
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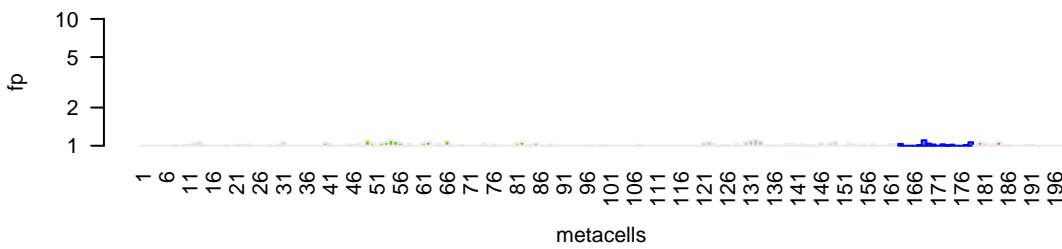
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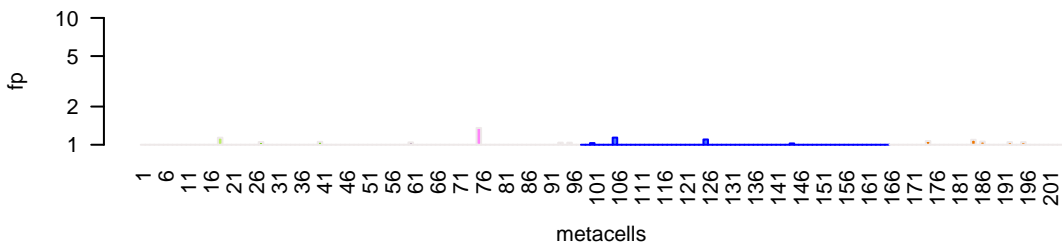
OG_3081:IRF1/IRF2 Nvec | Nvec_v1g141683



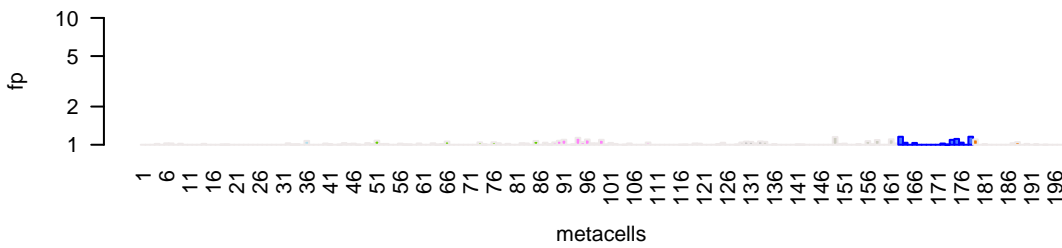
OG_3081:IRF1/IRF2 Hvul | Hvul_g12909_1



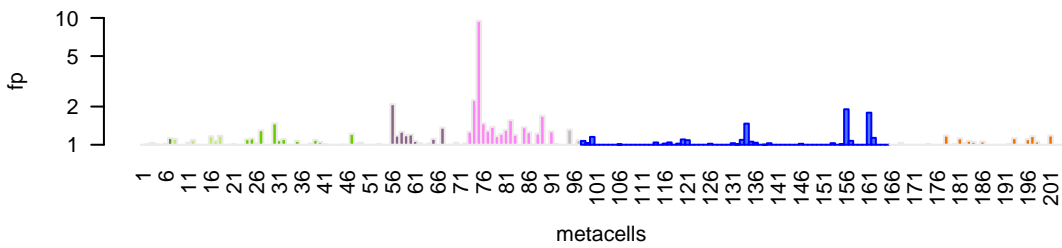
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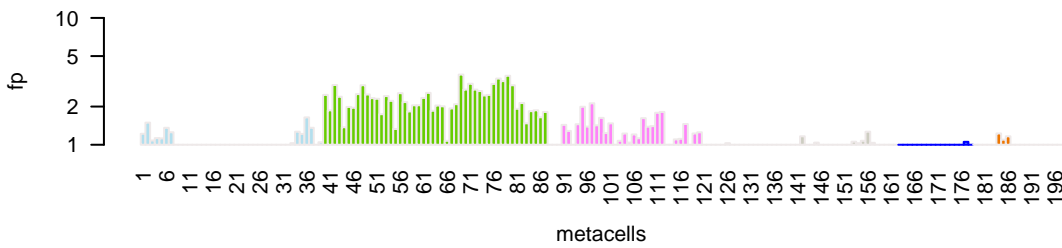
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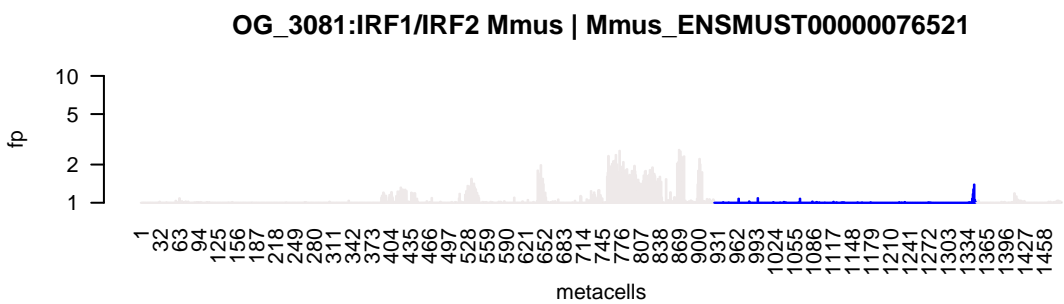
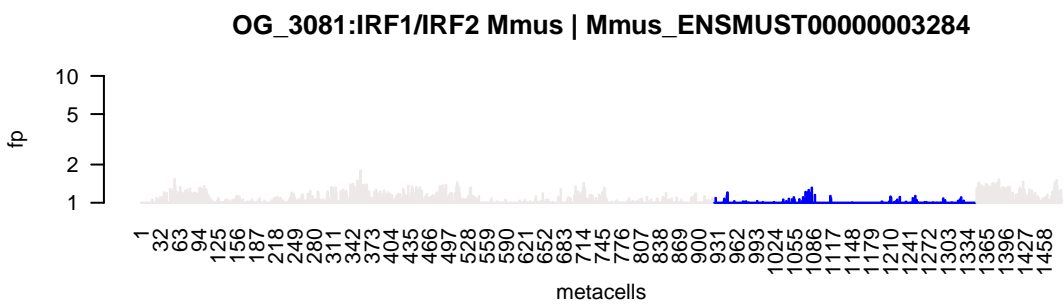
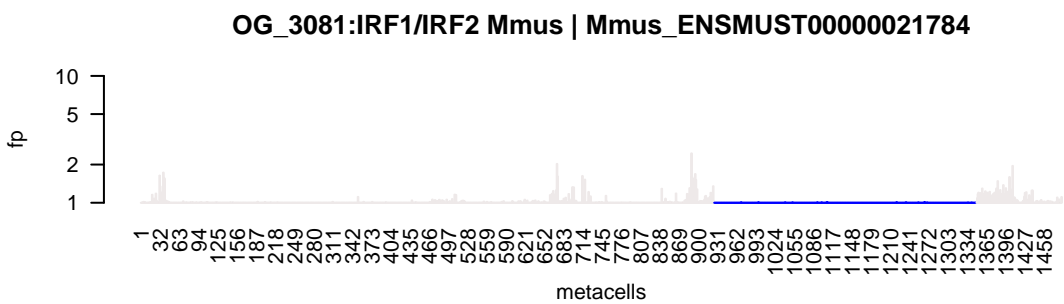
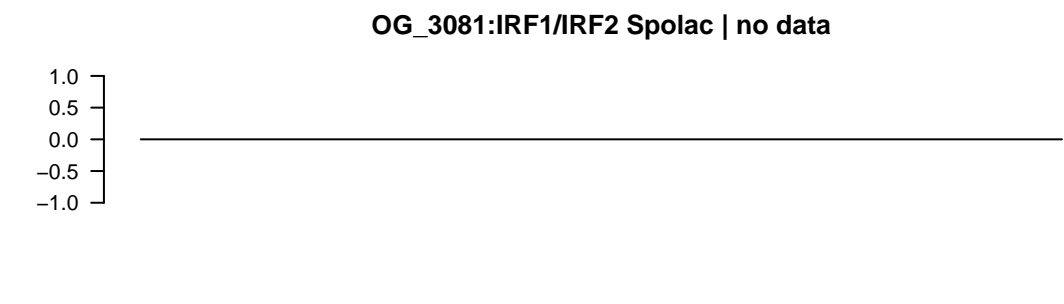
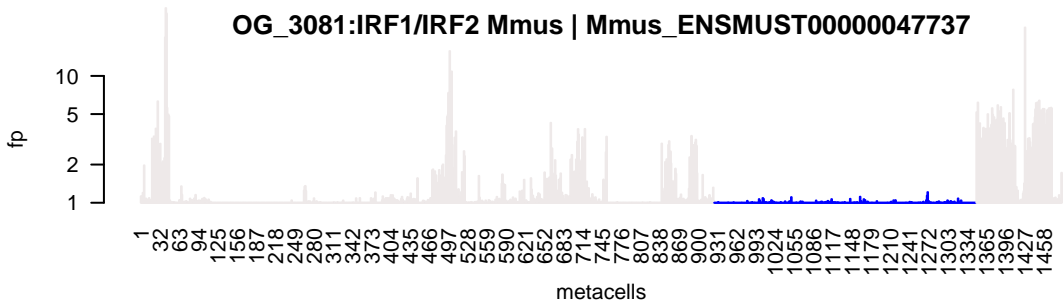
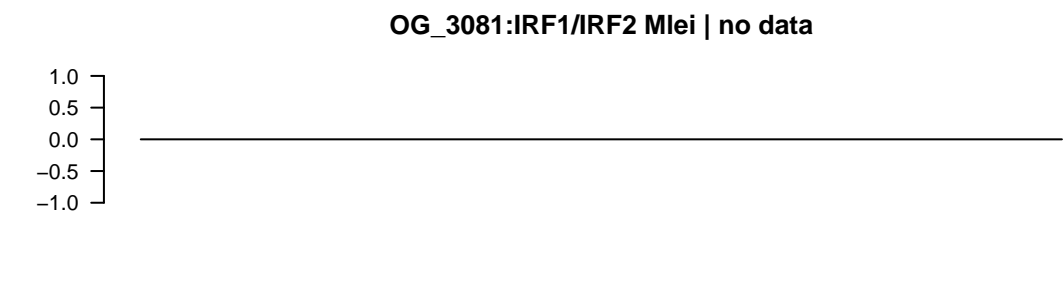
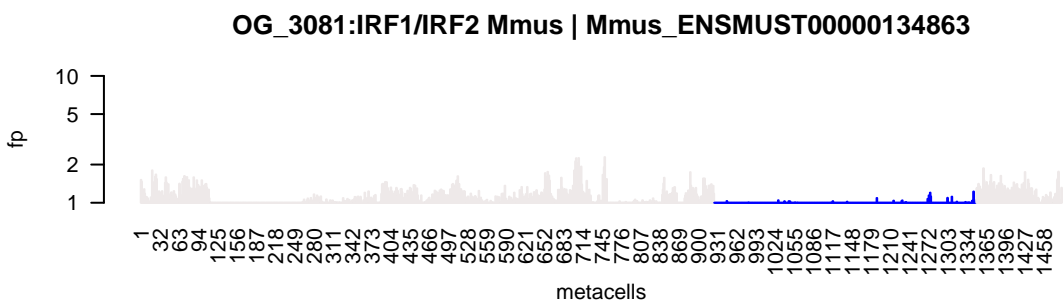
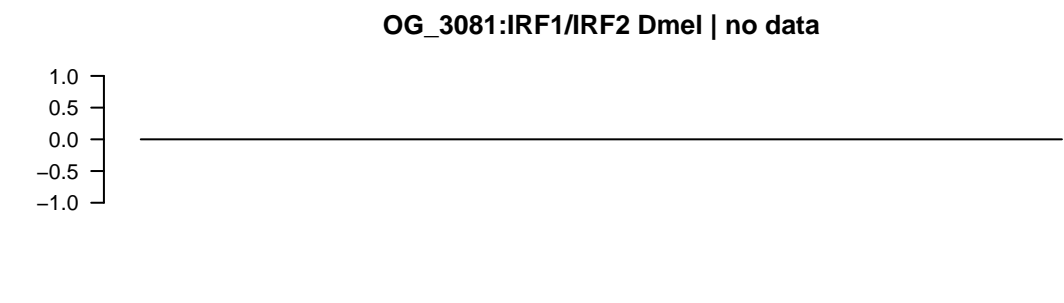
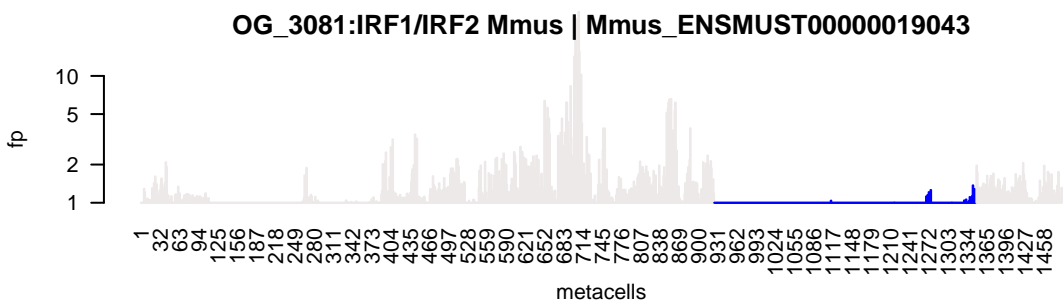
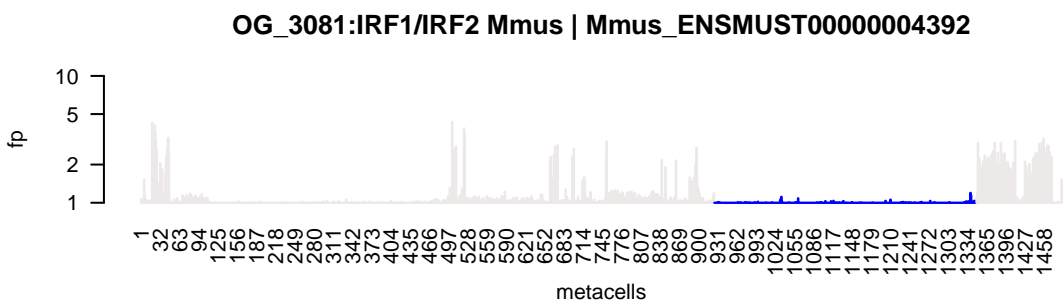
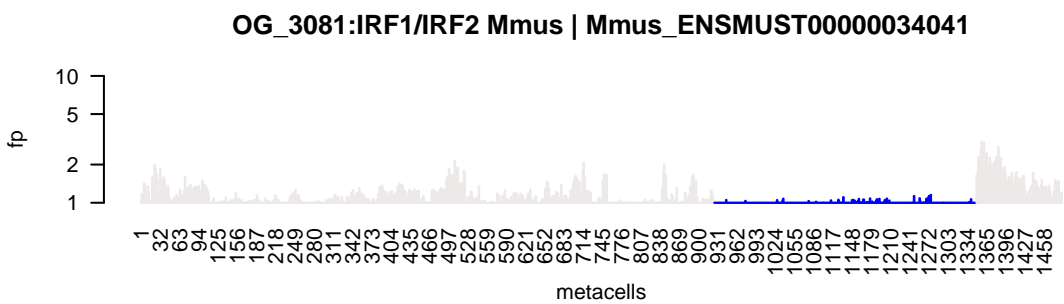
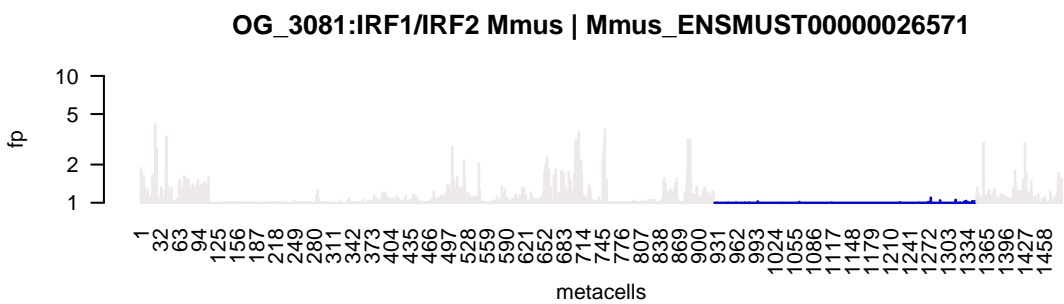
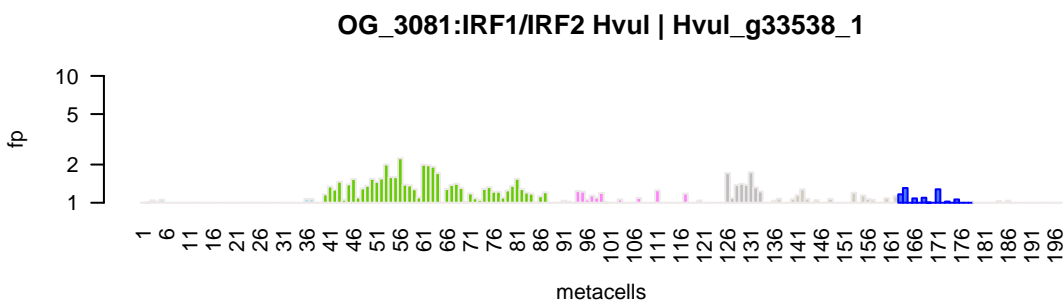


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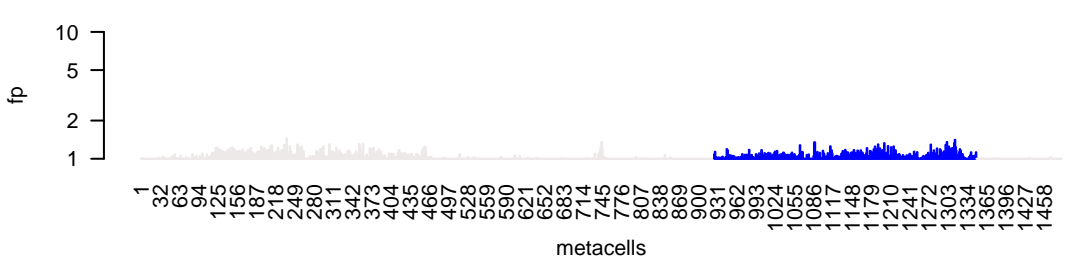
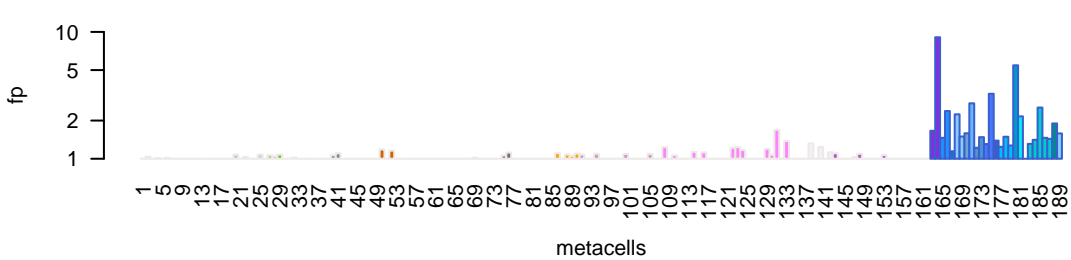


OG_3081:IRF1/IRF2 Hvul | Hvul_g32638_1

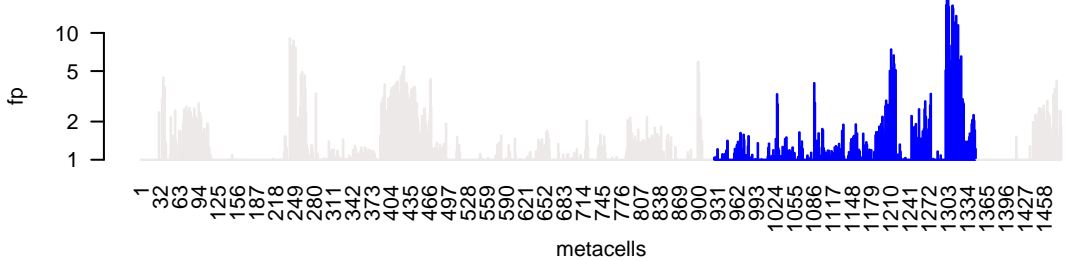
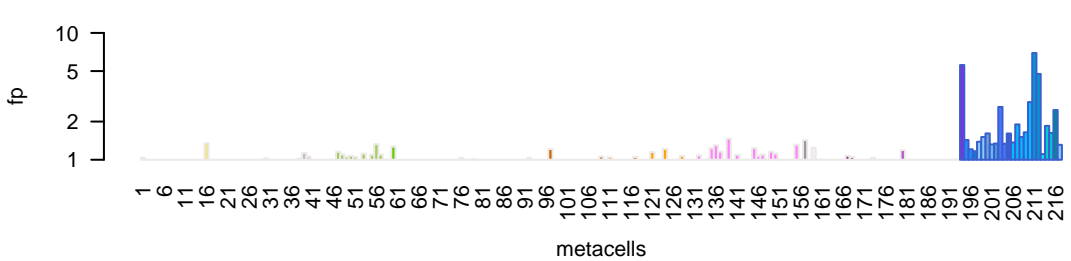




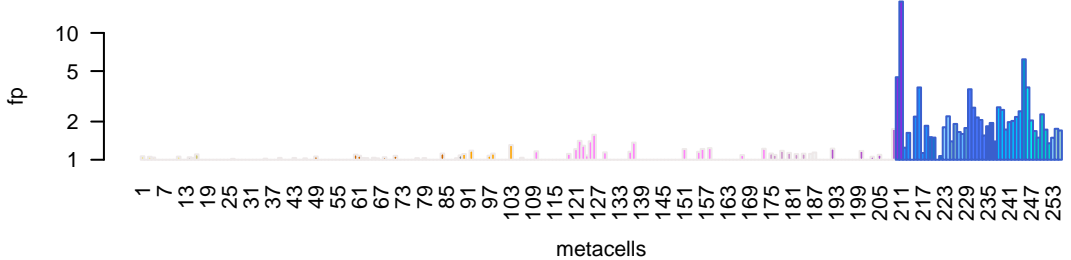
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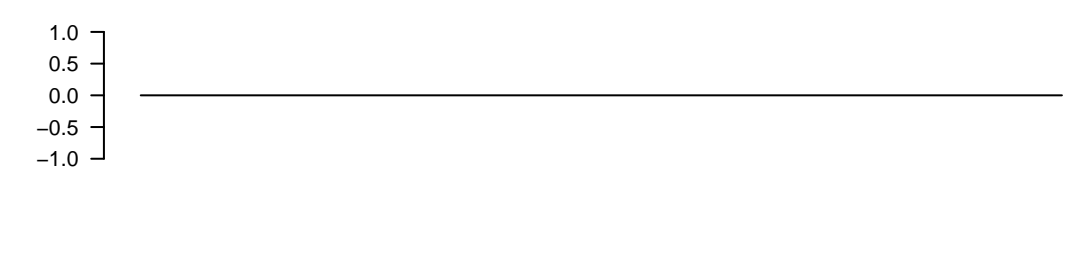
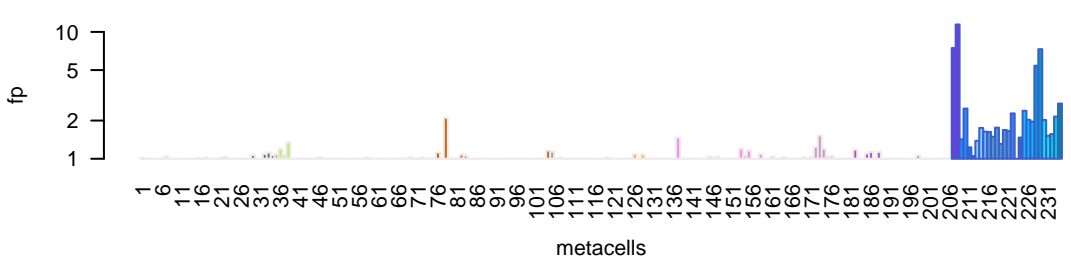
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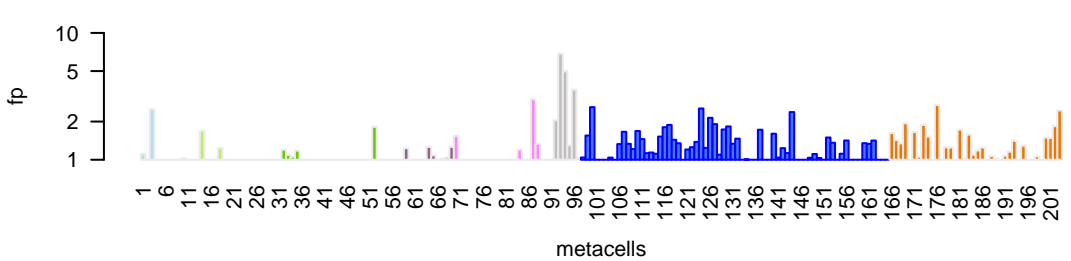
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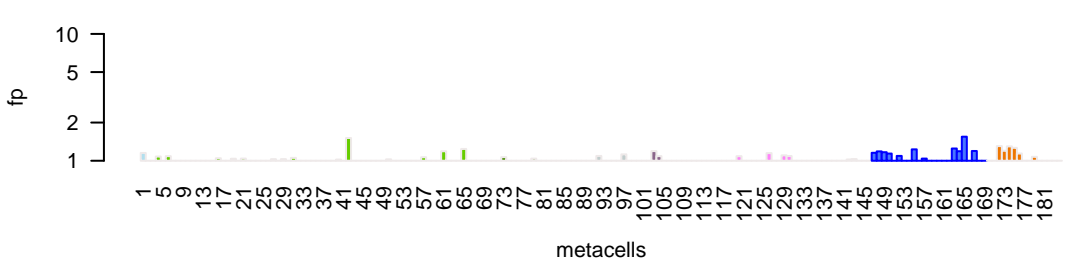
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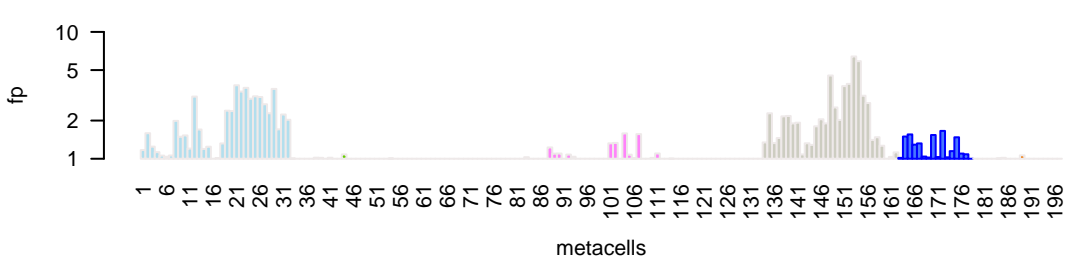
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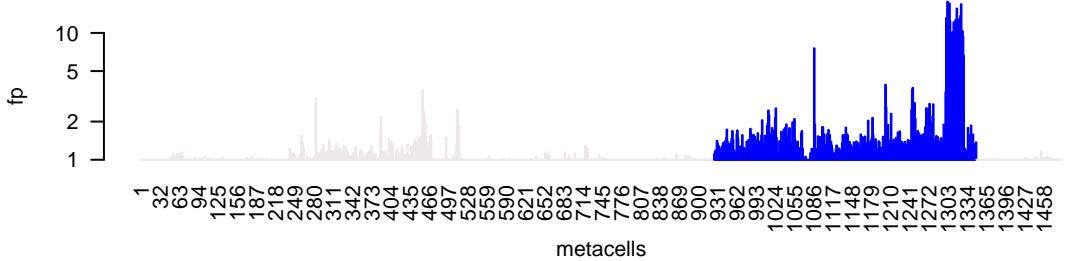
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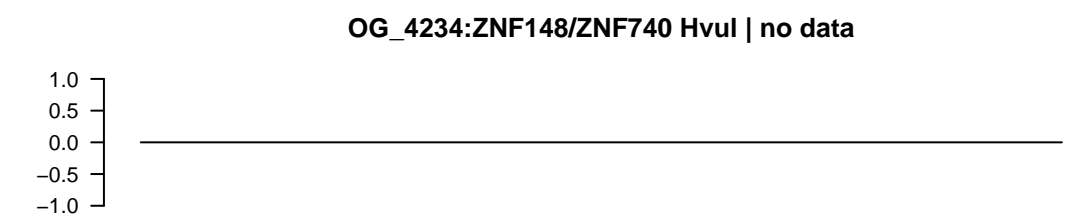
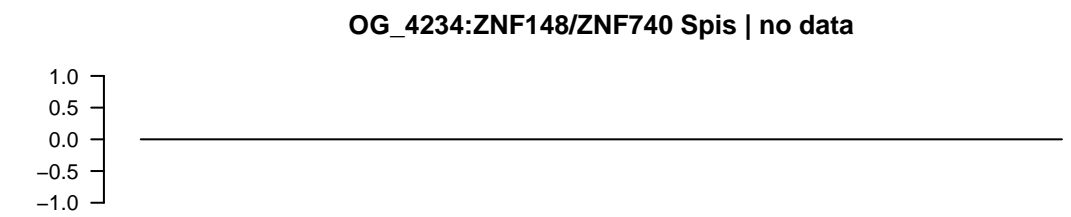
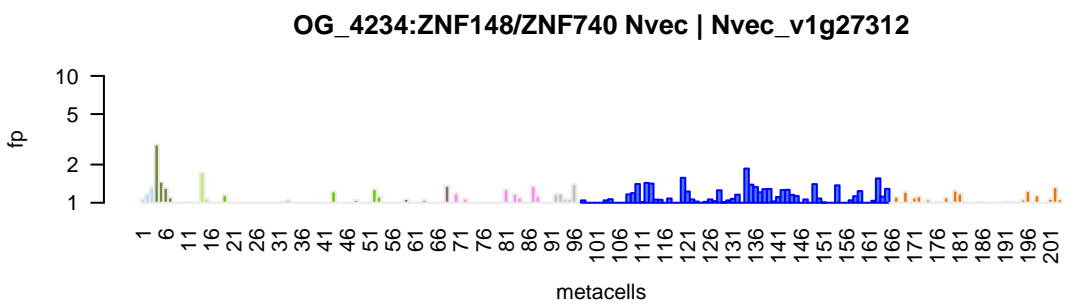
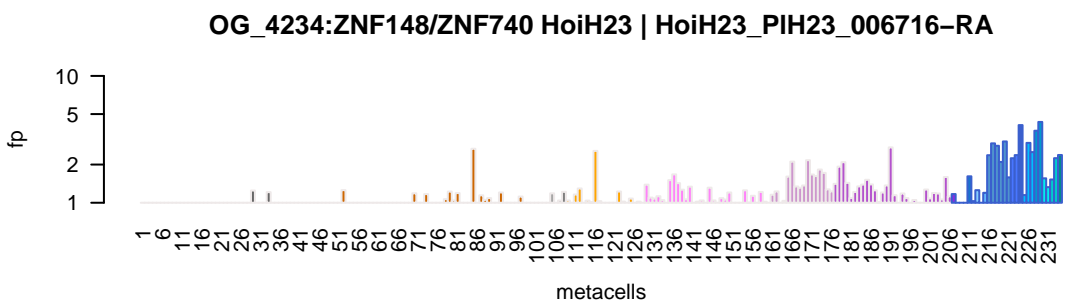
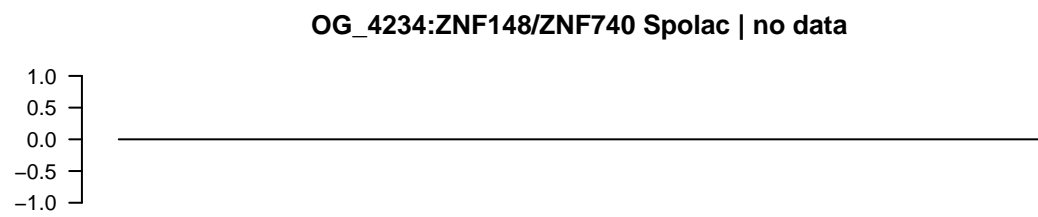
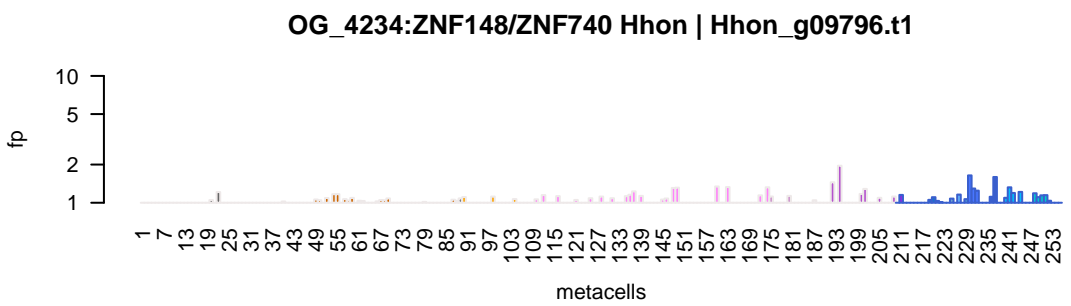
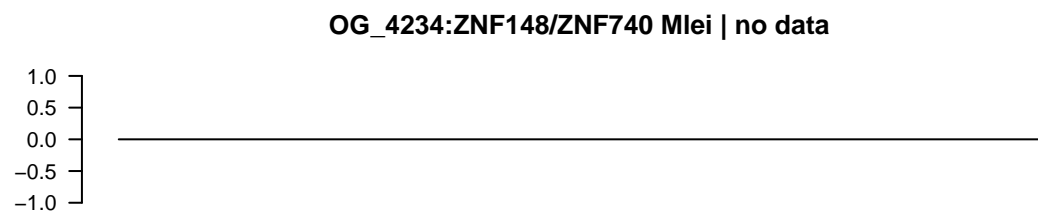
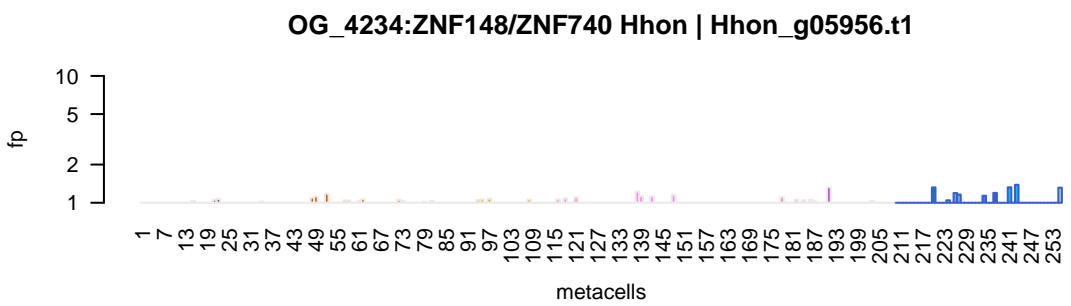
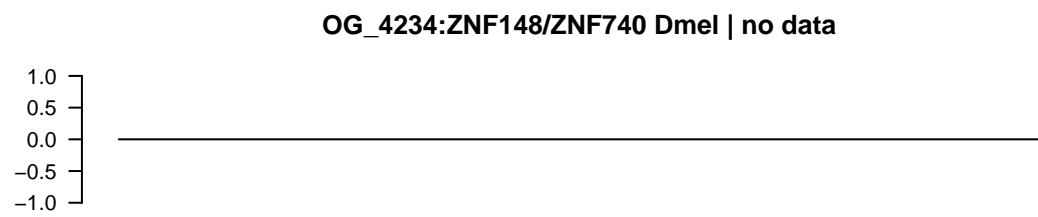
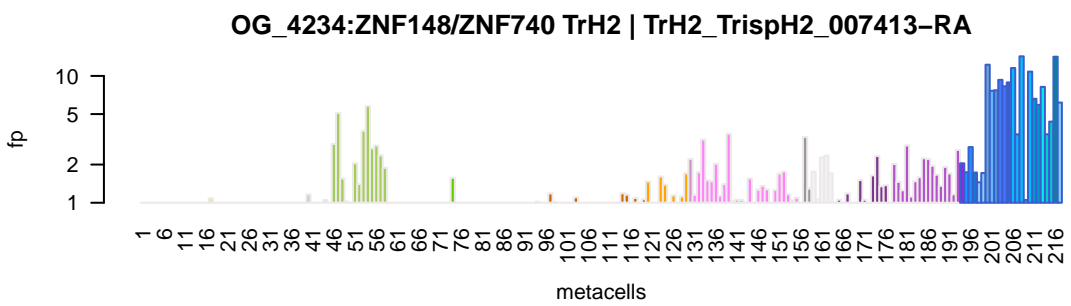
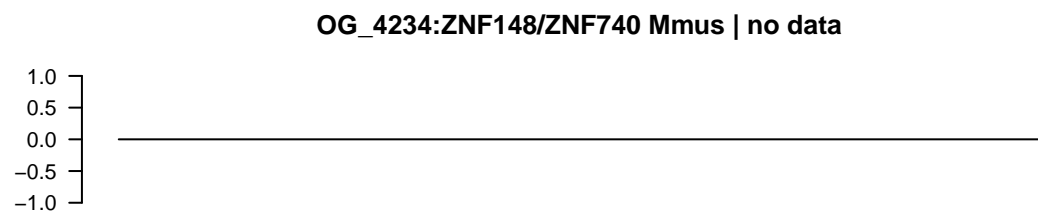
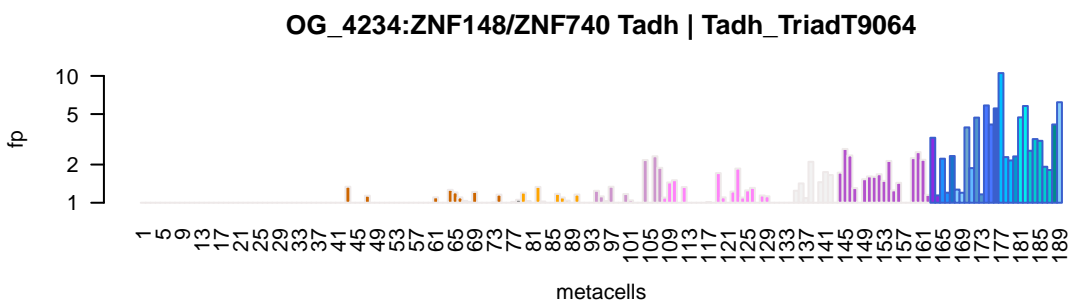


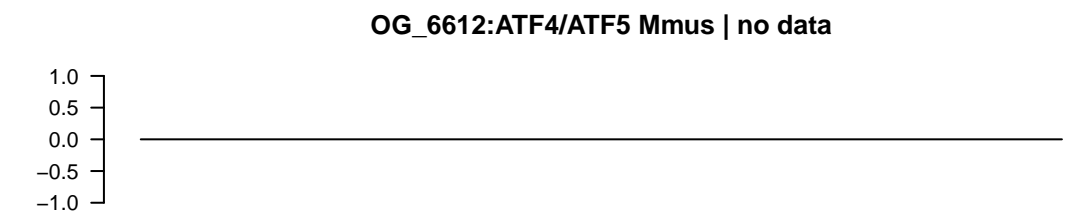
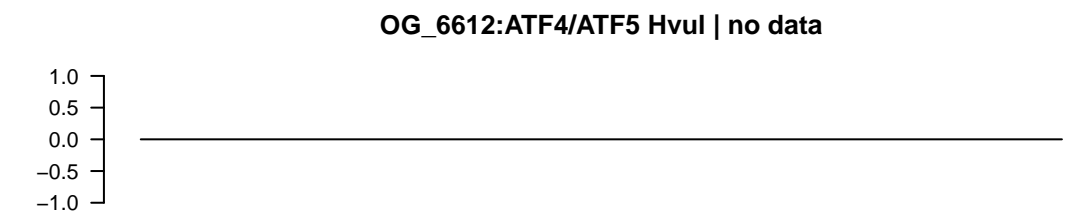
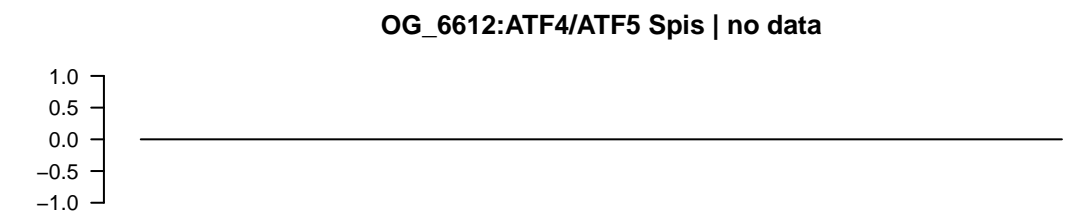
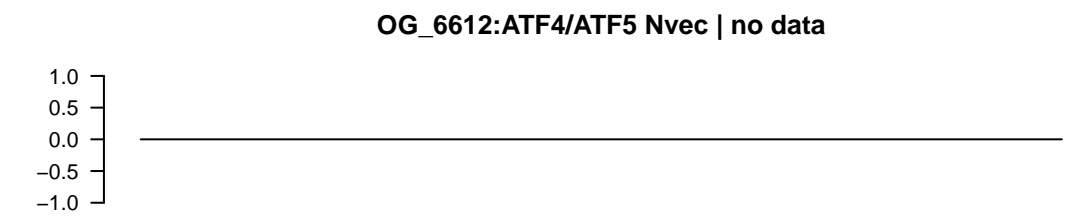
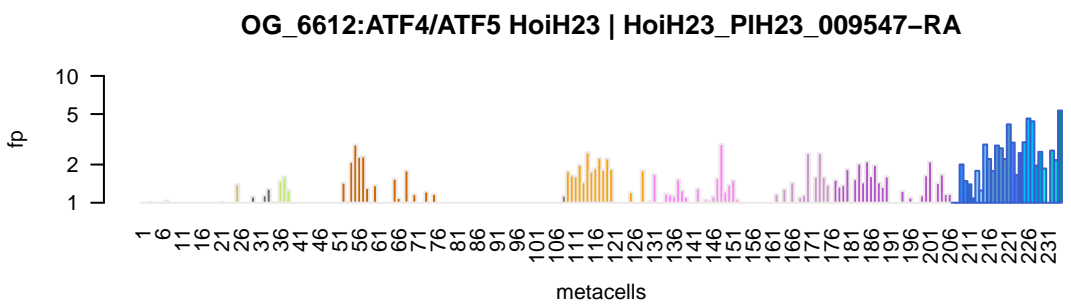
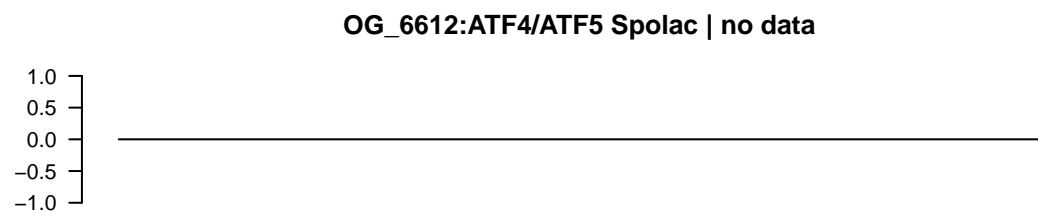
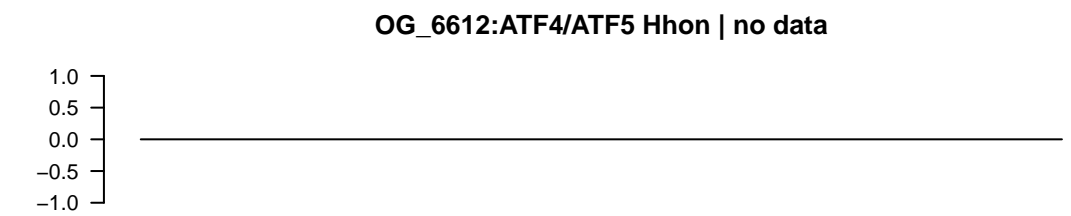
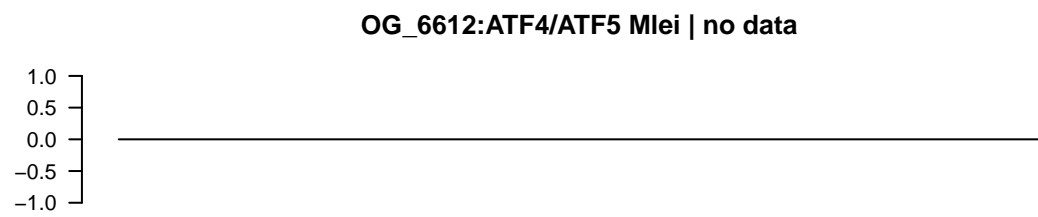
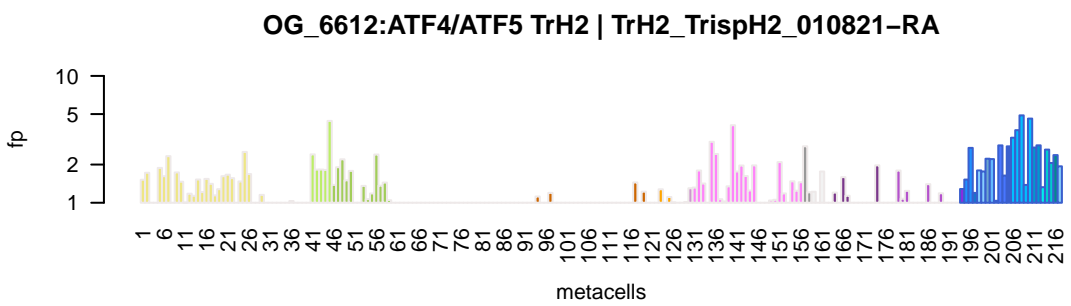
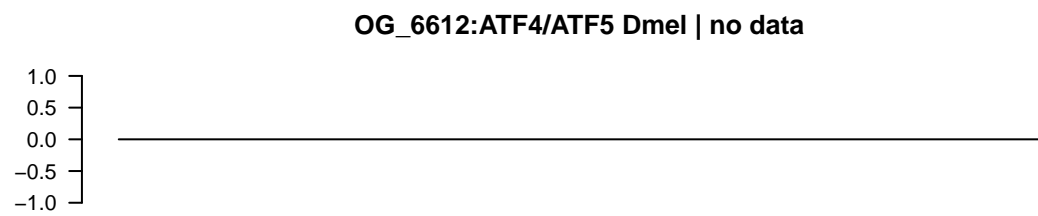
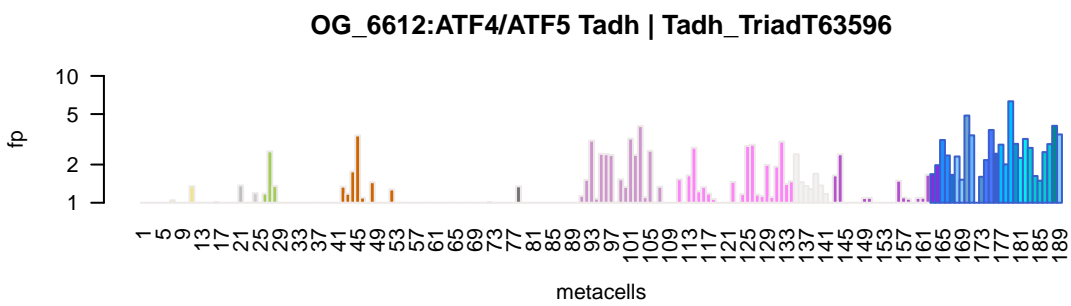
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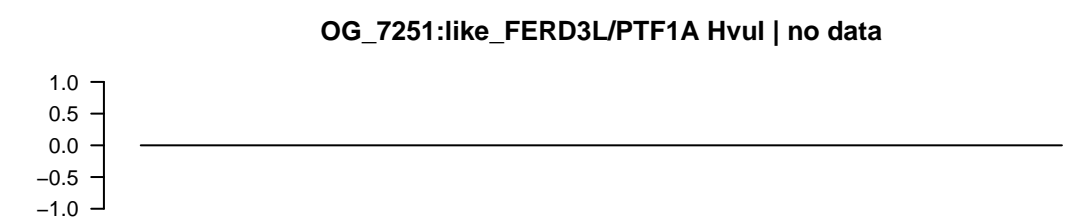
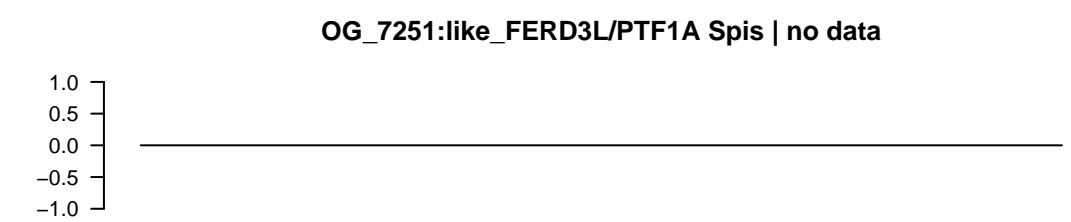
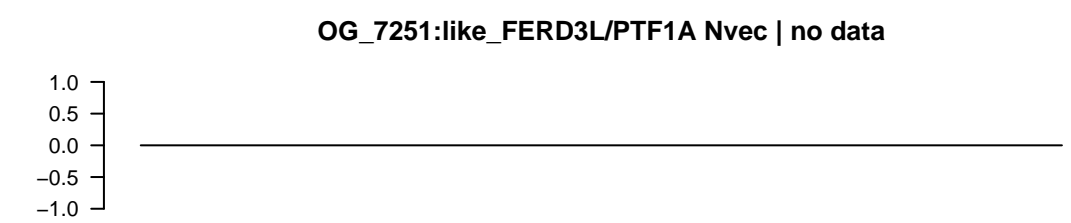
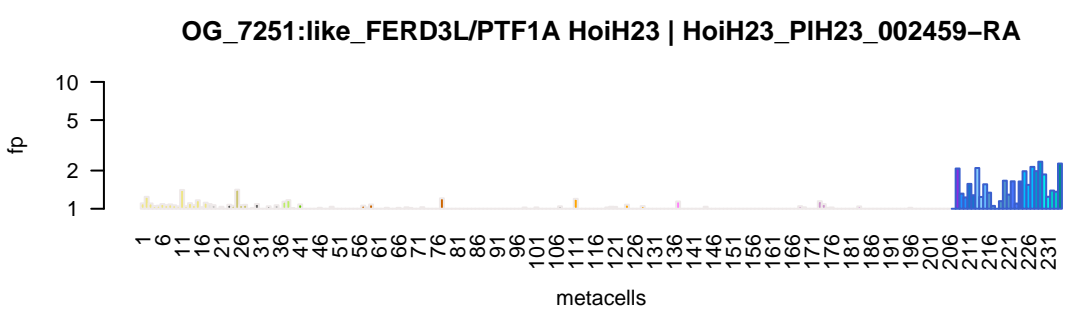
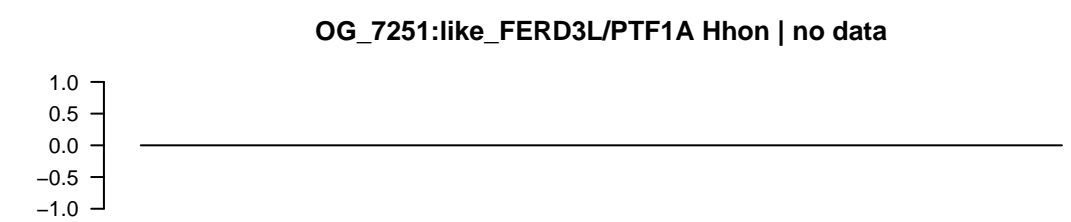
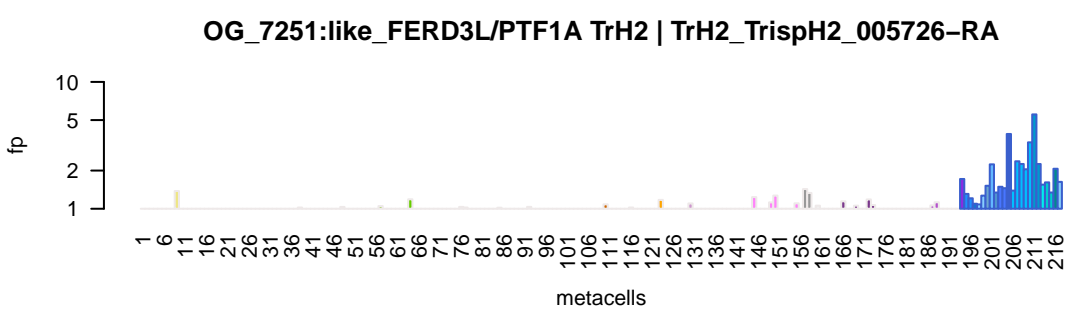
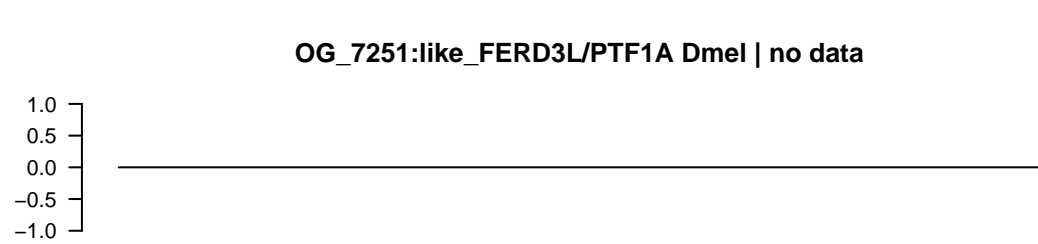
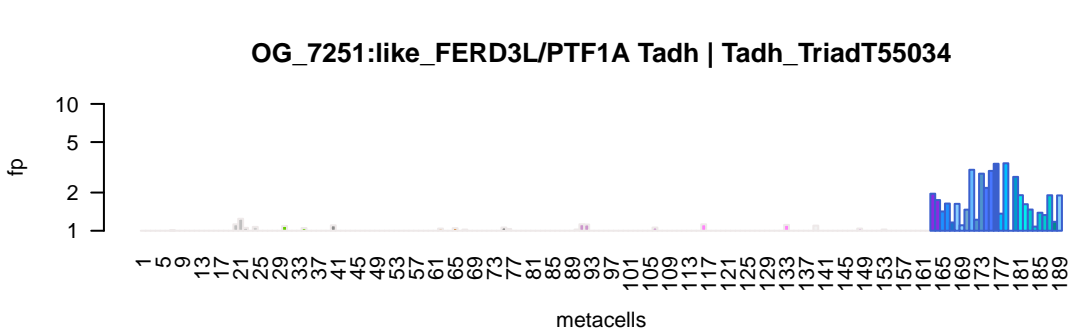


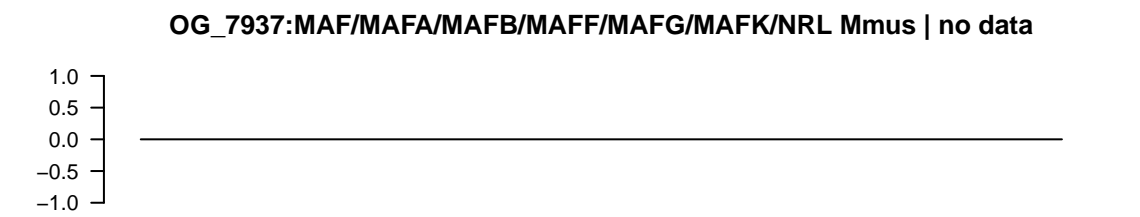
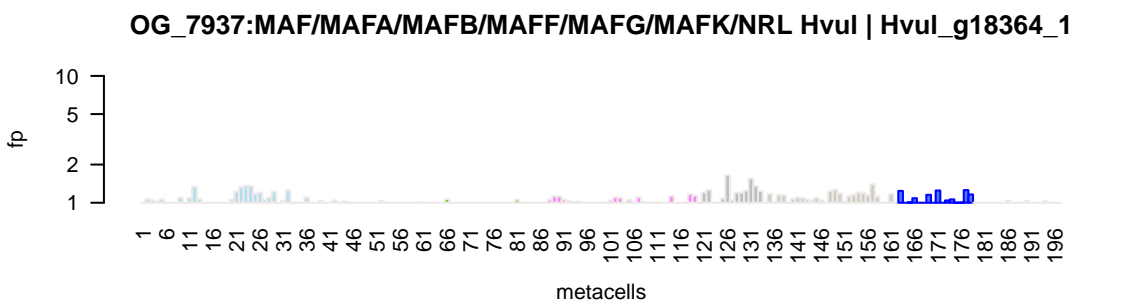
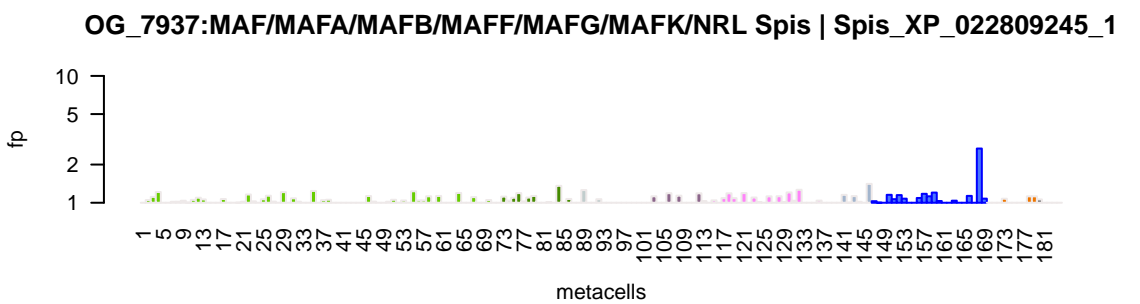
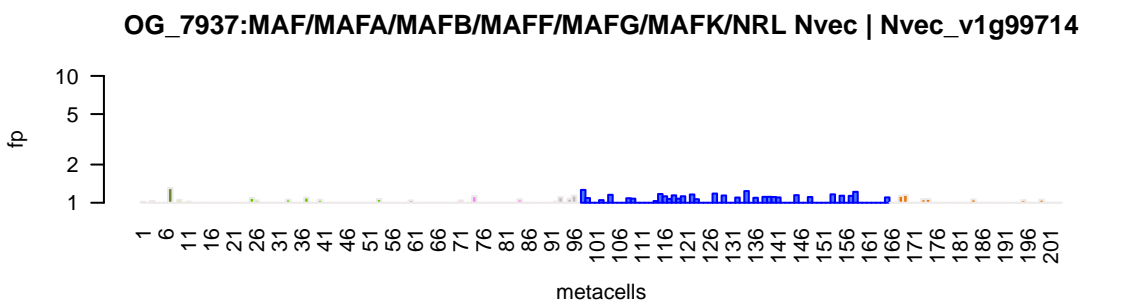
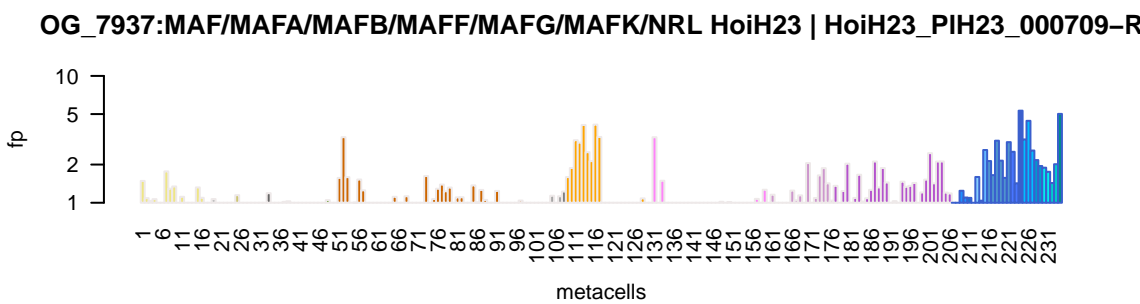
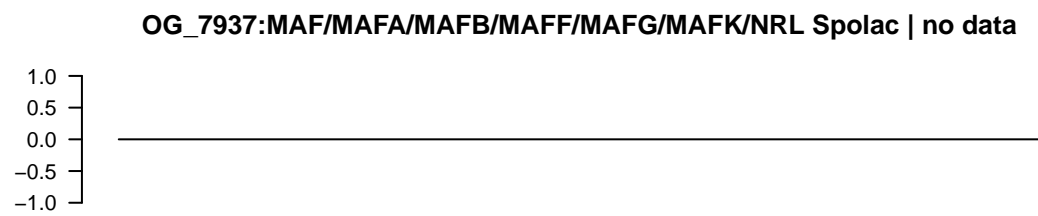
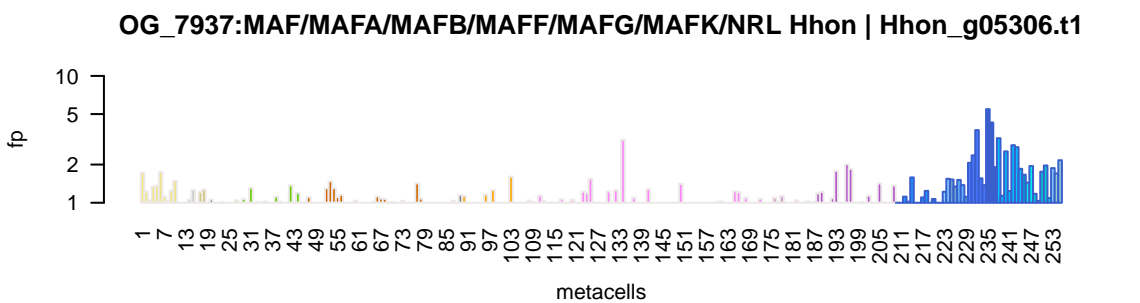
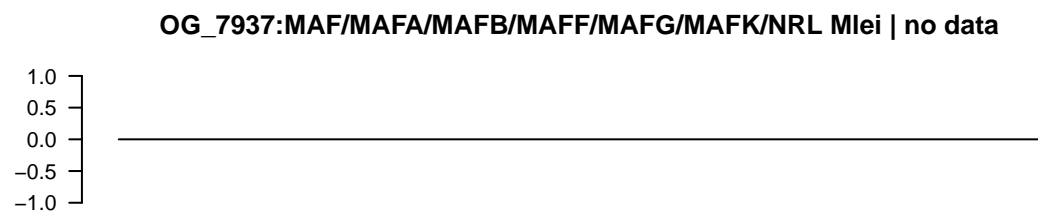
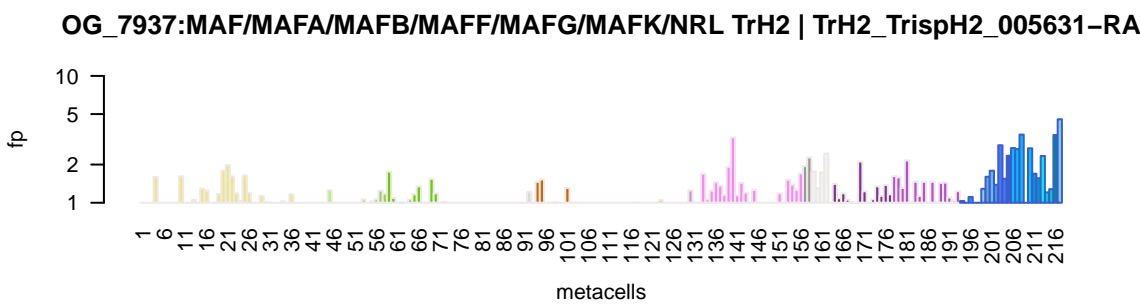
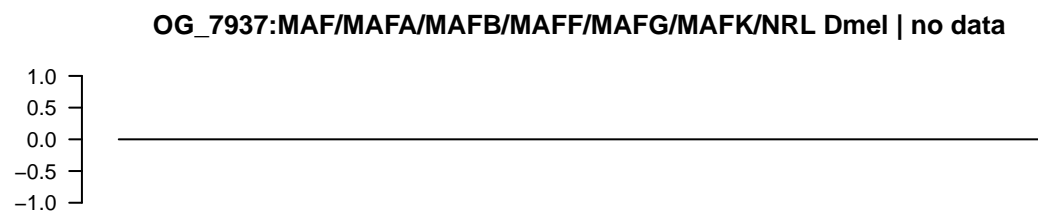
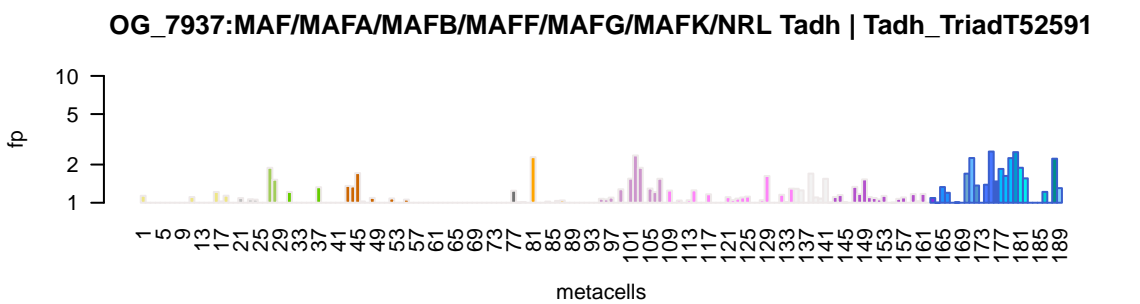
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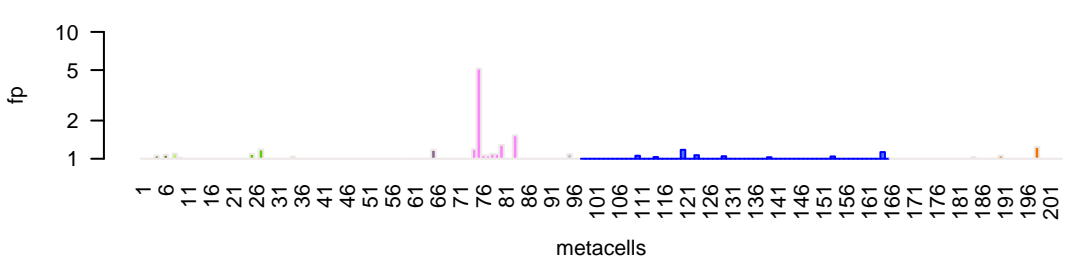
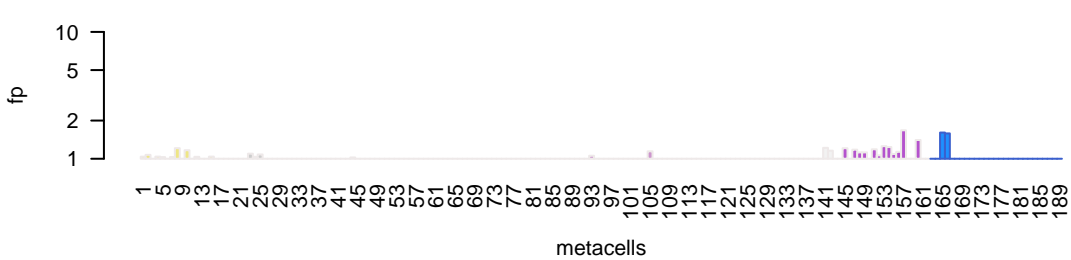




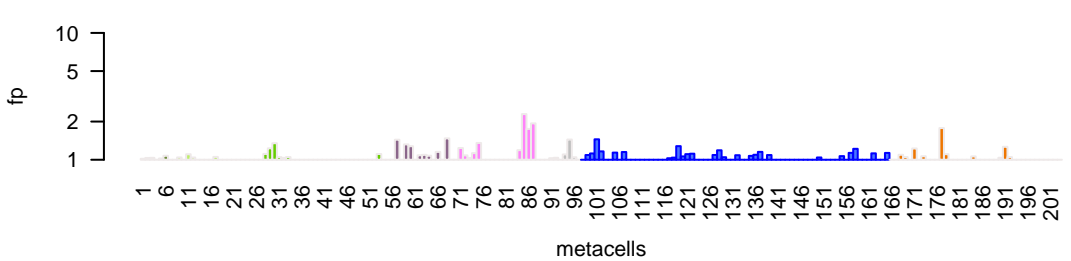
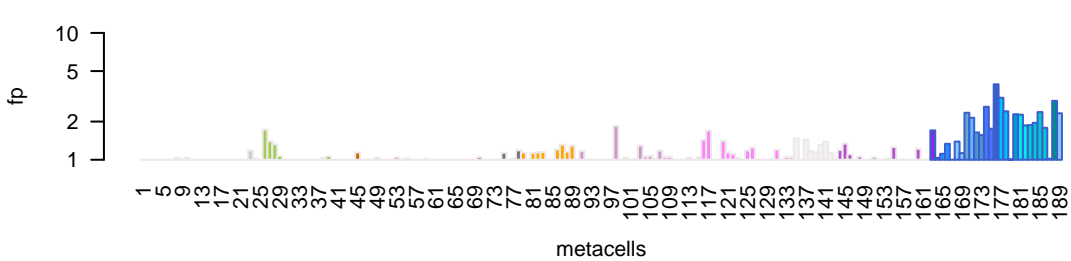




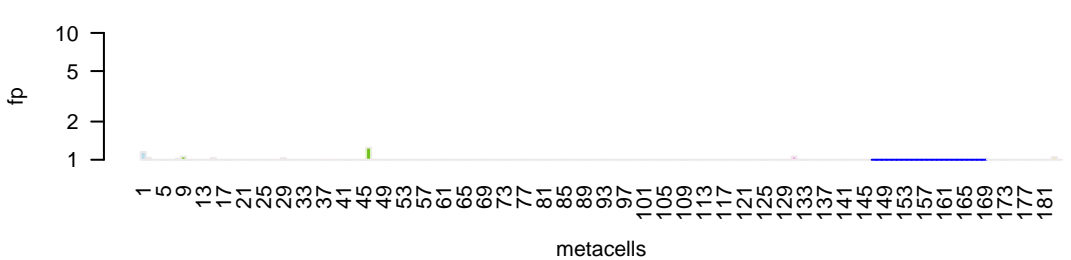
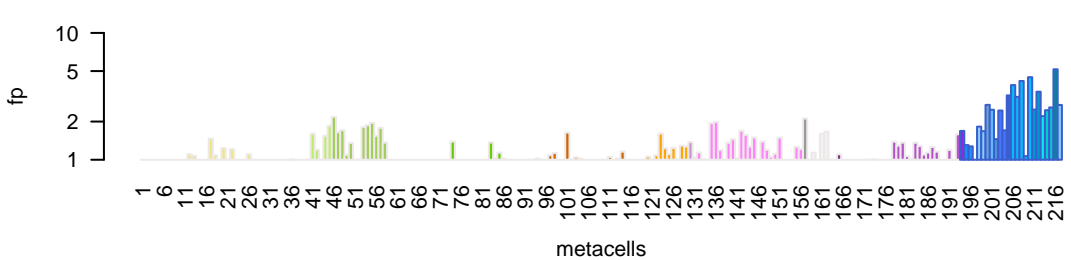
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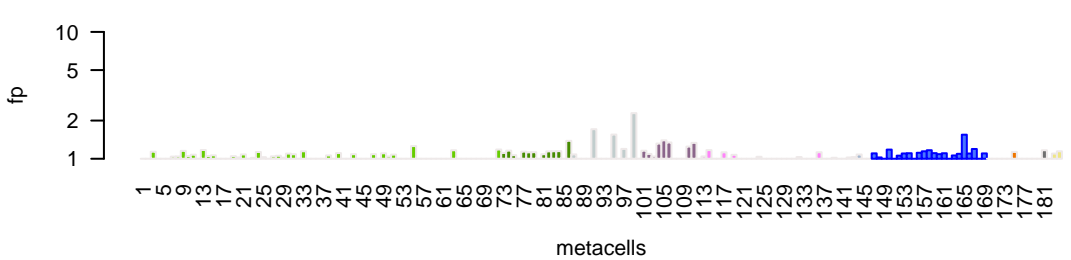
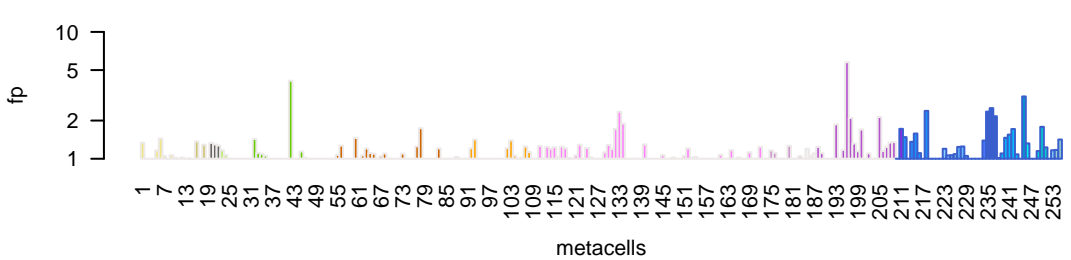
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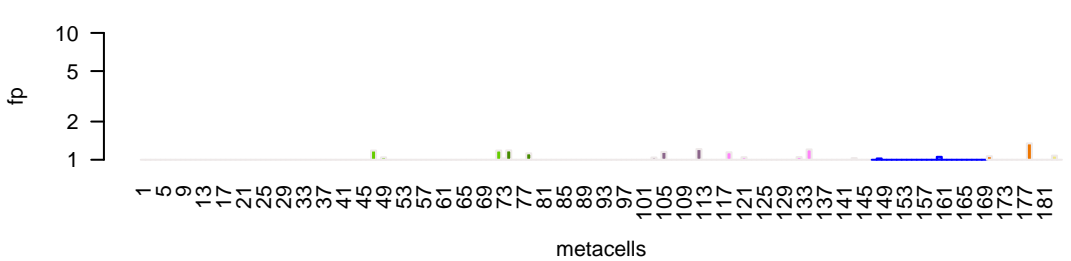
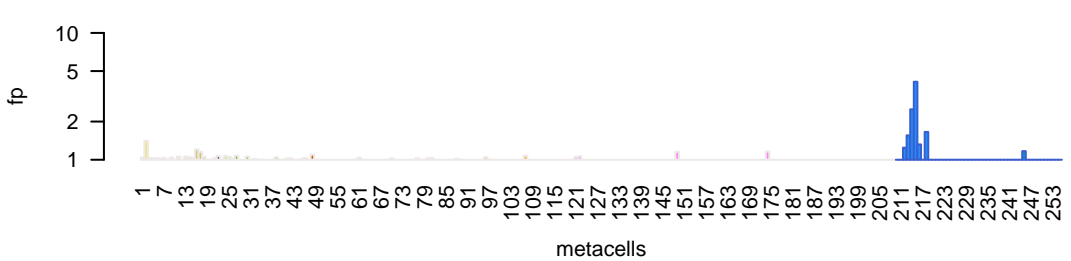
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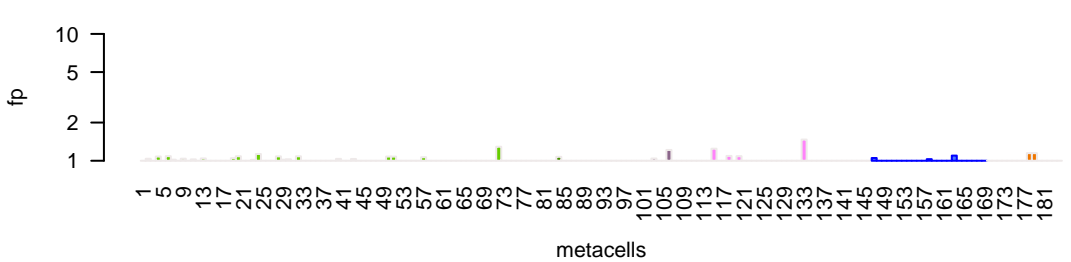
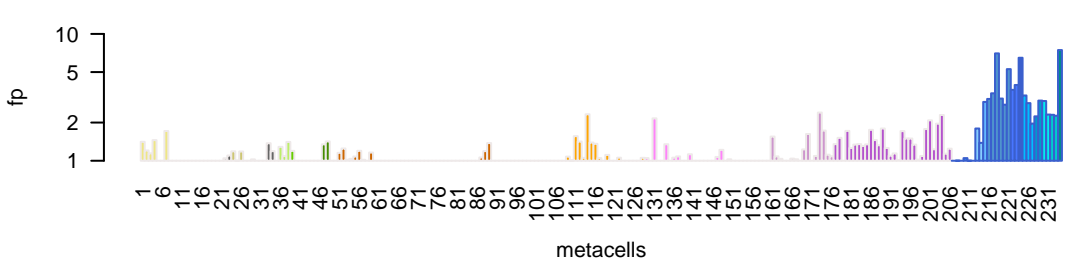
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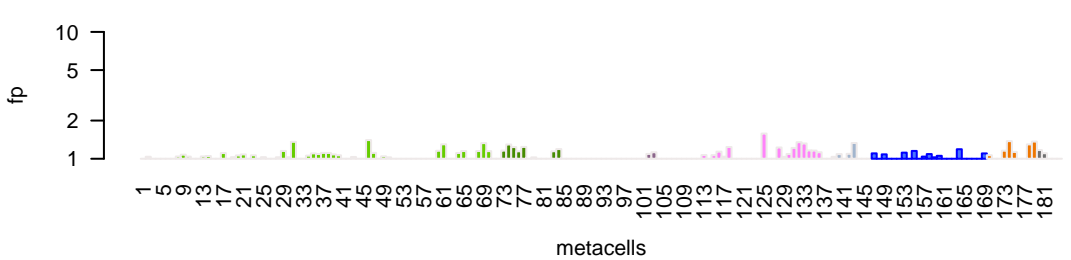
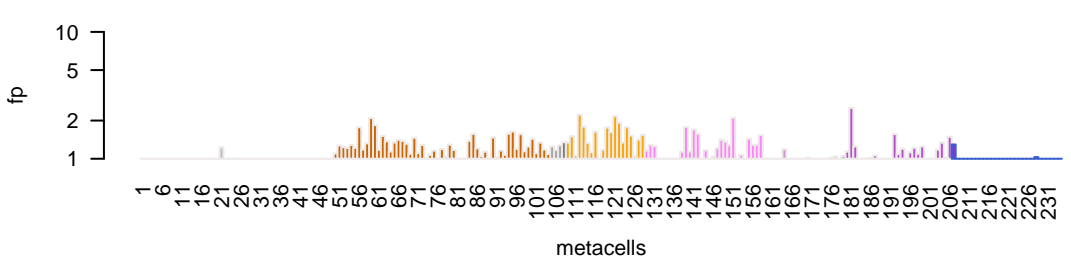
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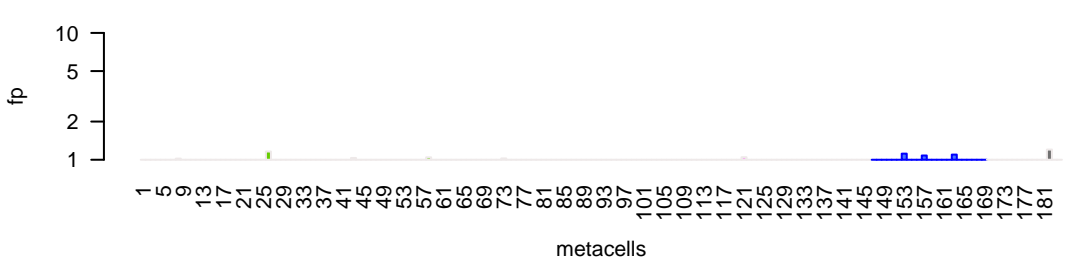
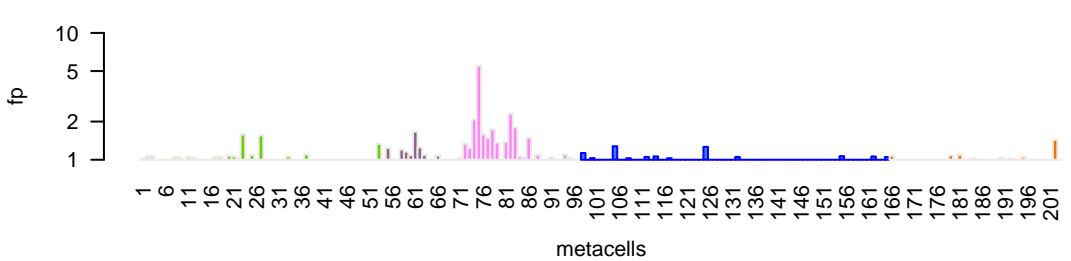
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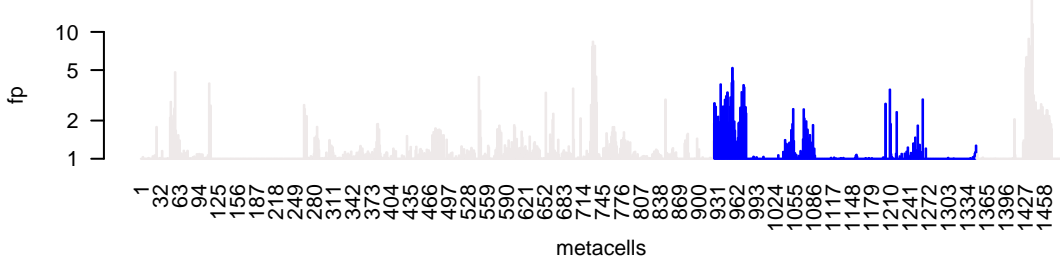
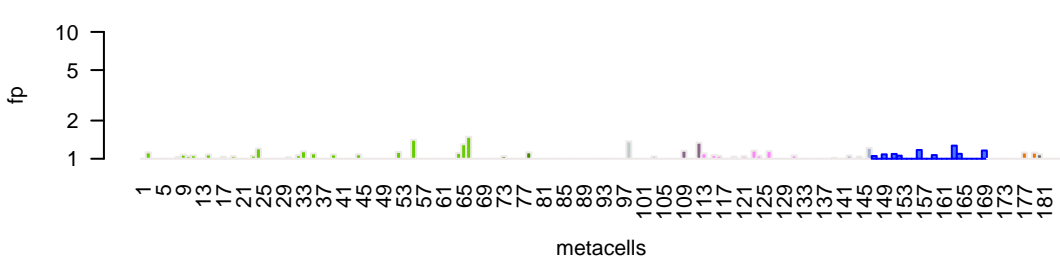
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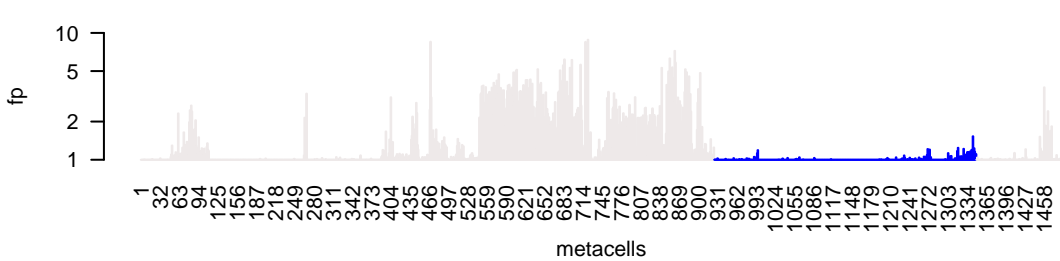
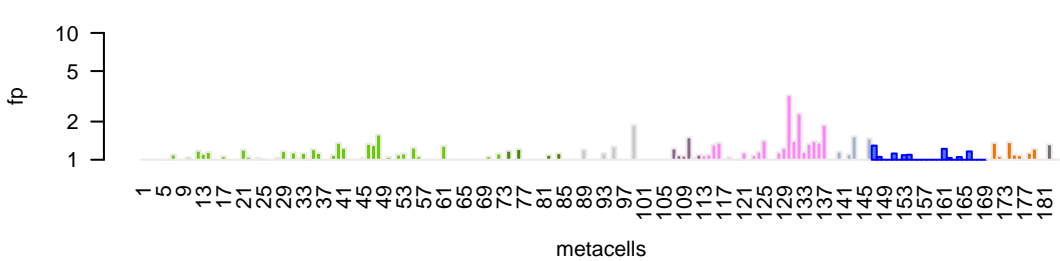
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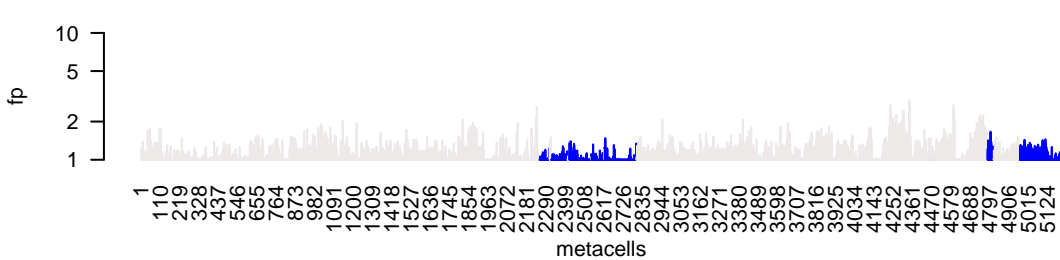
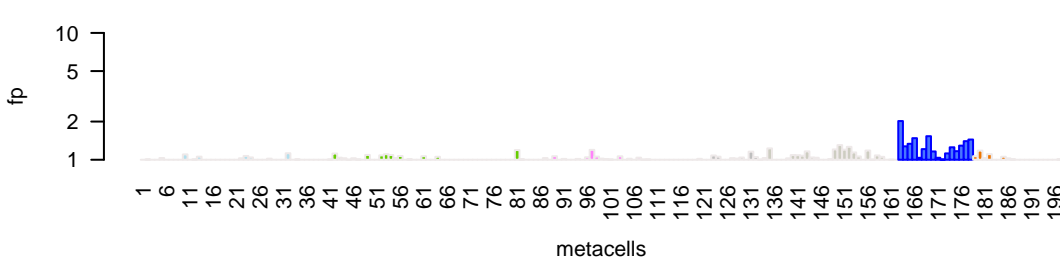
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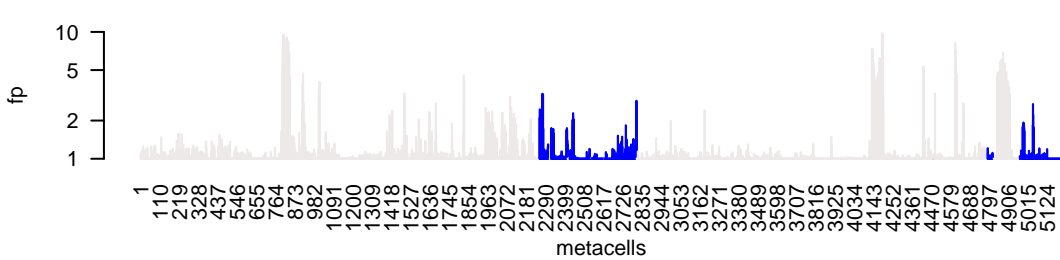
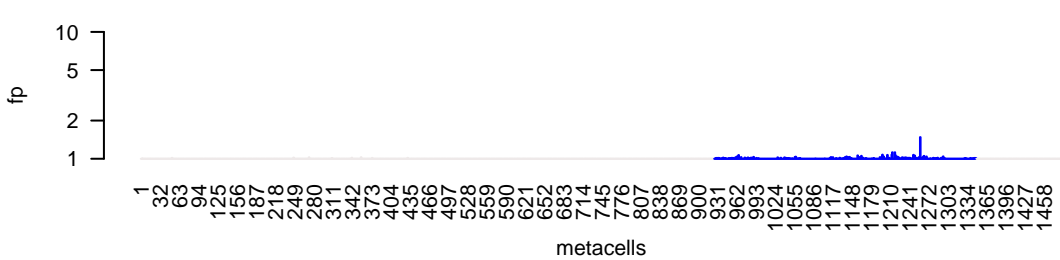
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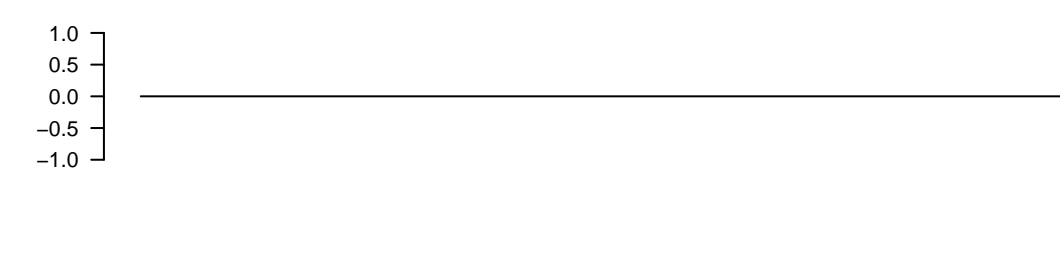
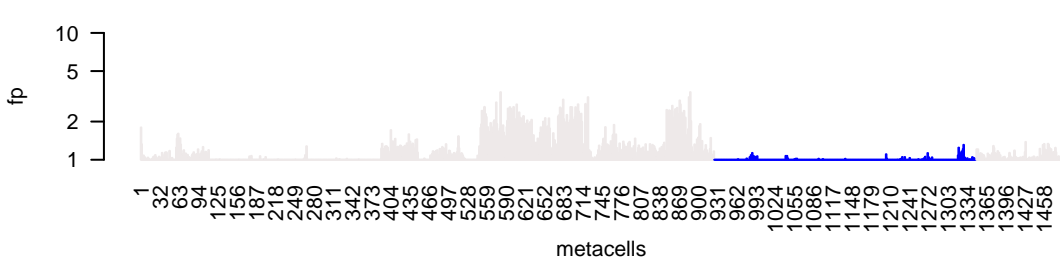
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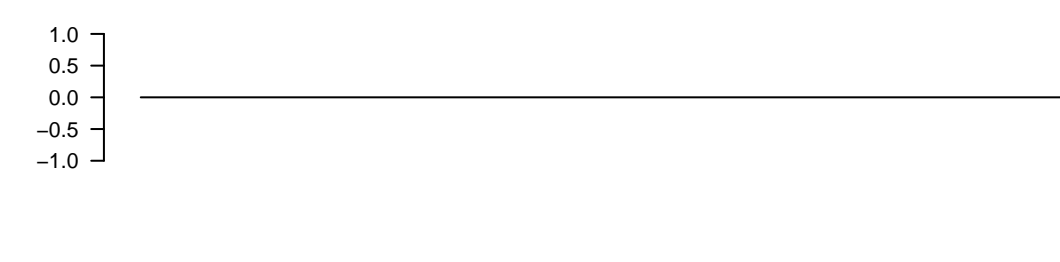
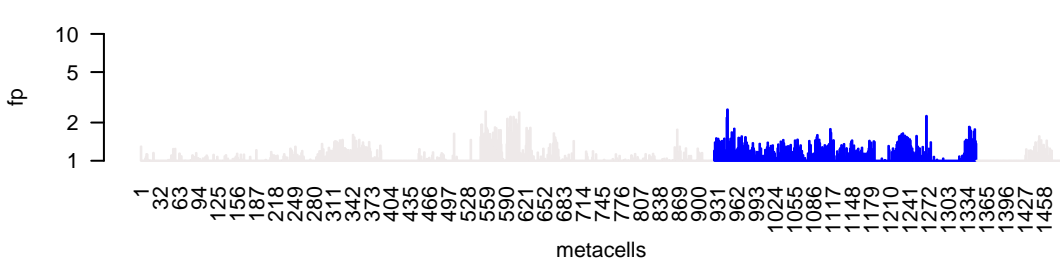
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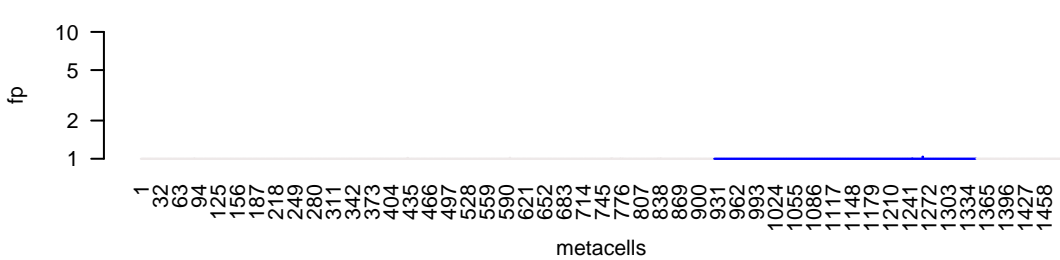
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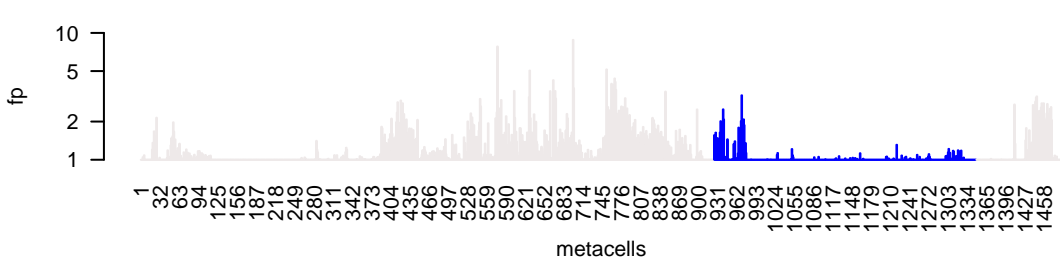
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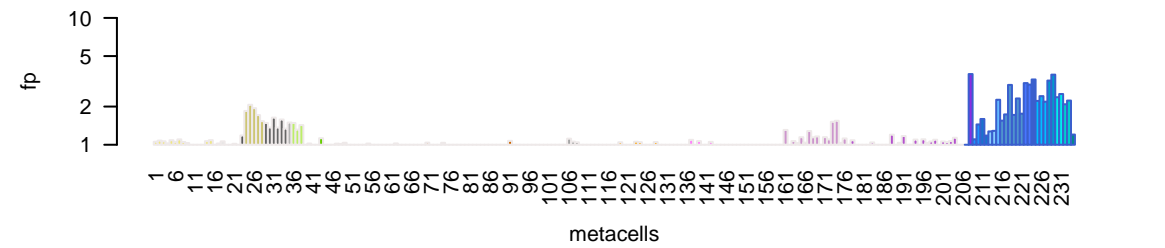
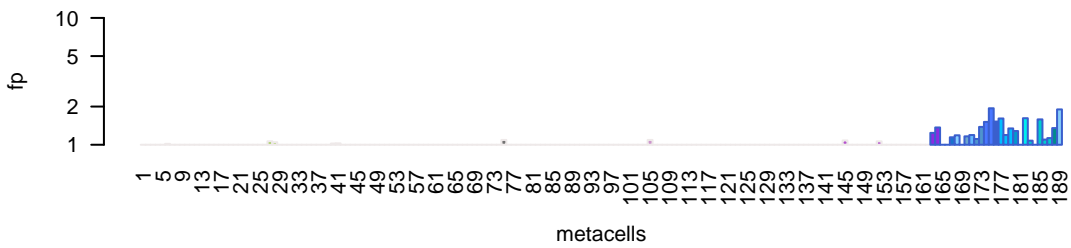
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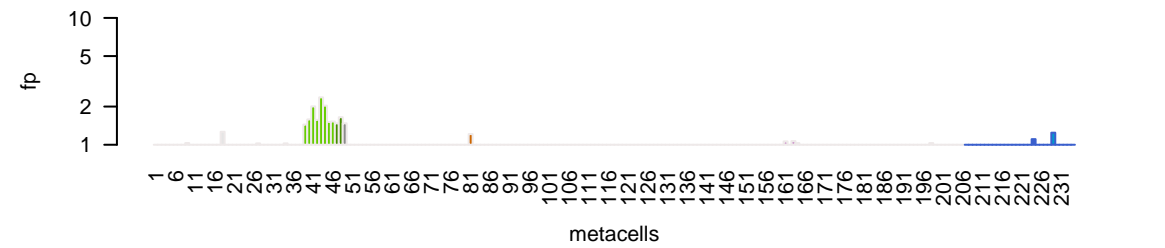
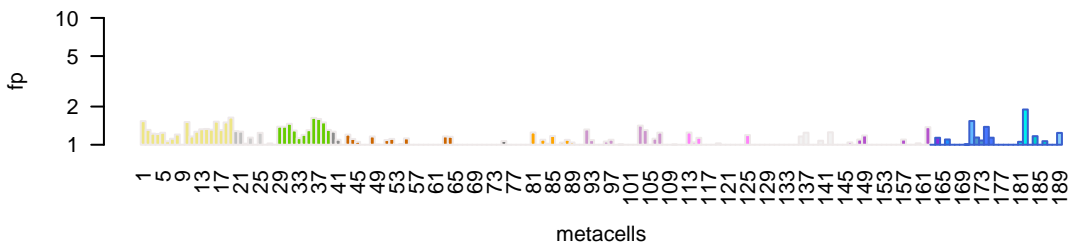
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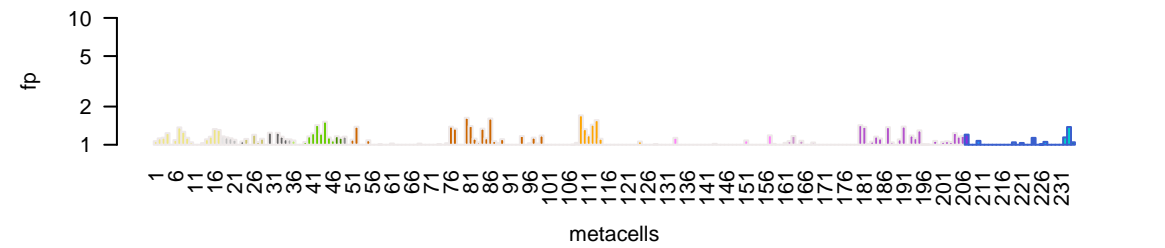
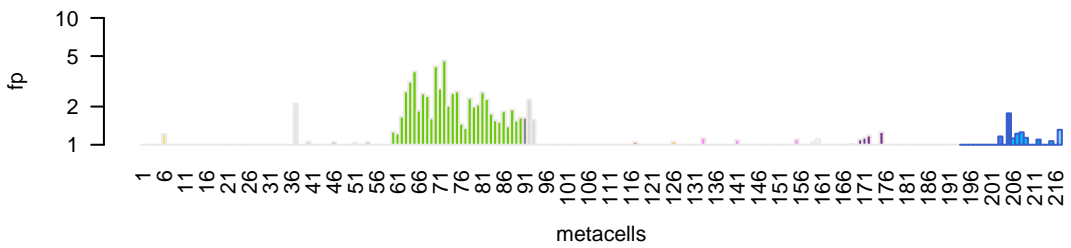
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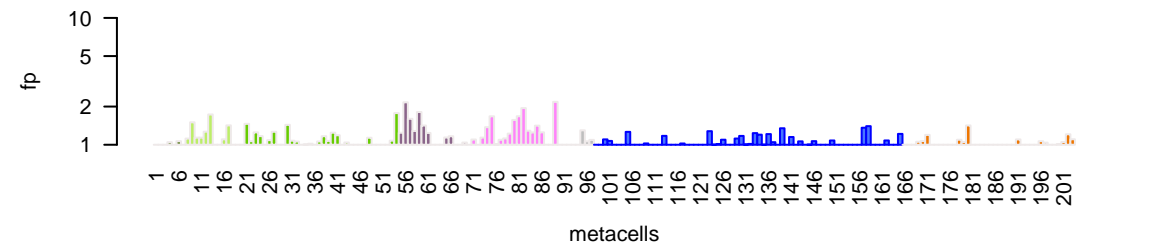
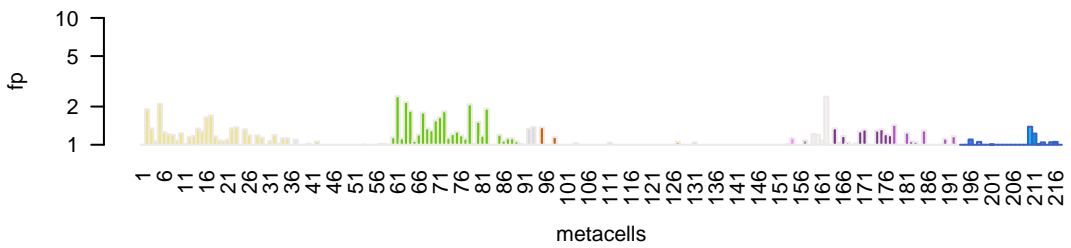
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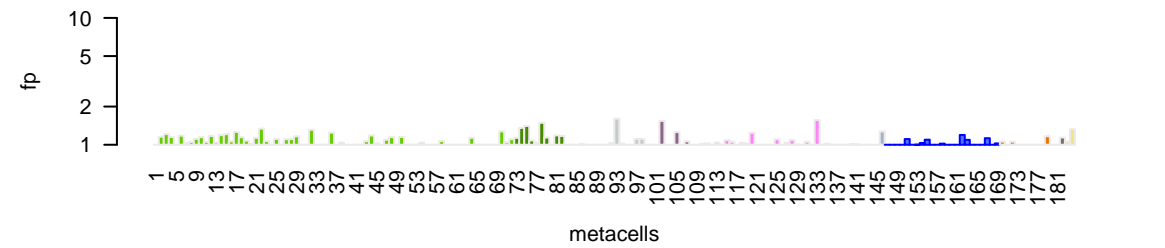
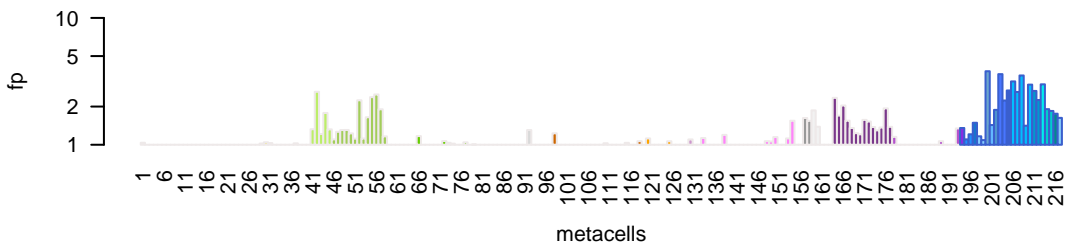
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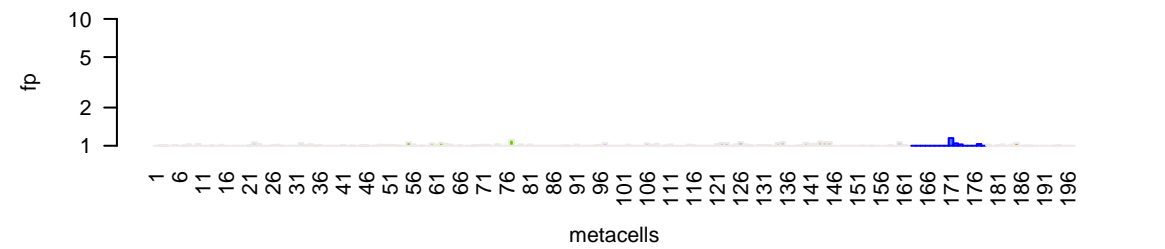
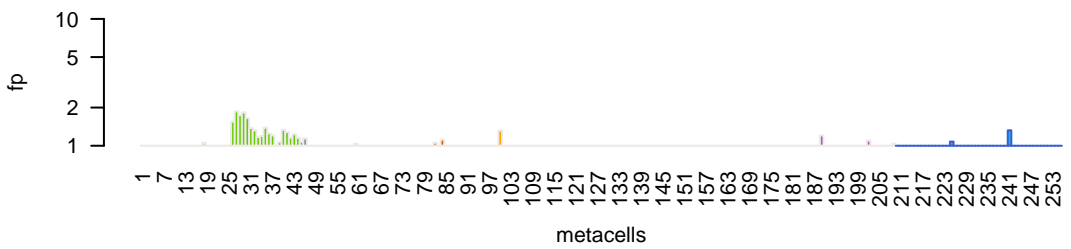
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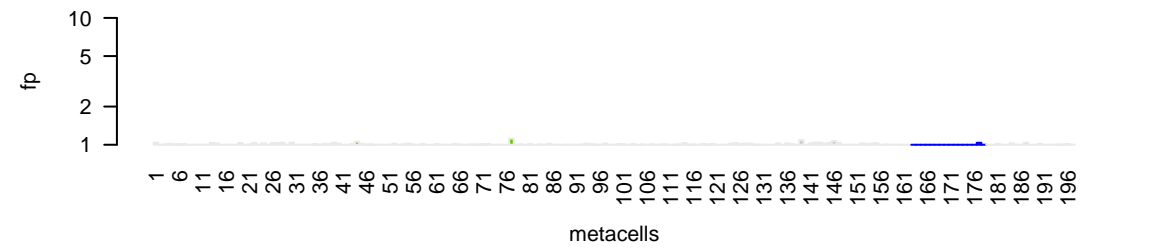
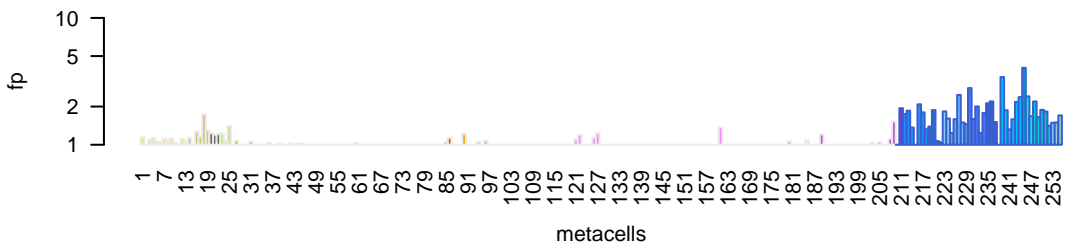
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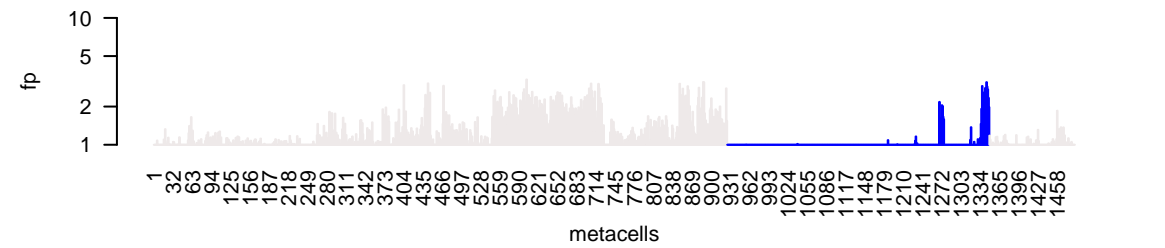
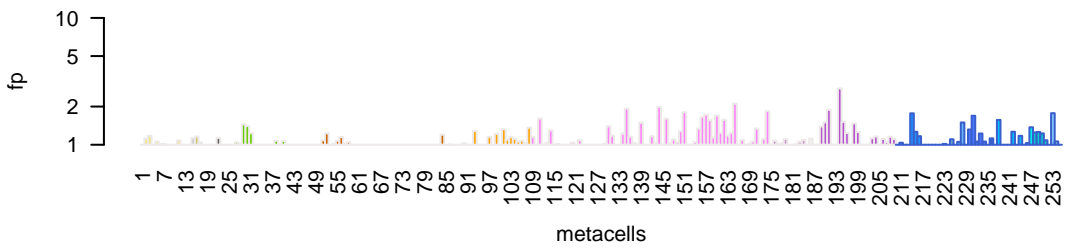
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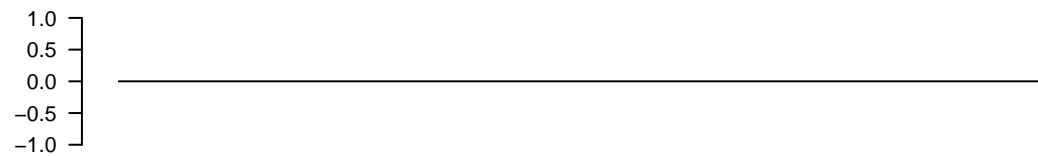
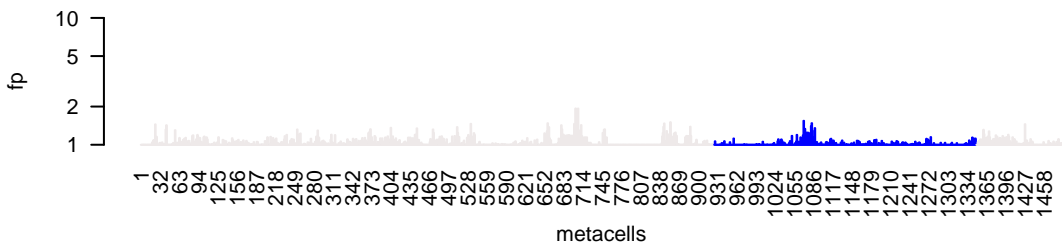
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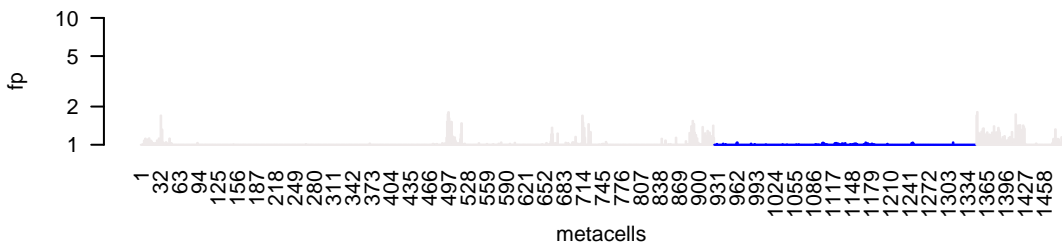
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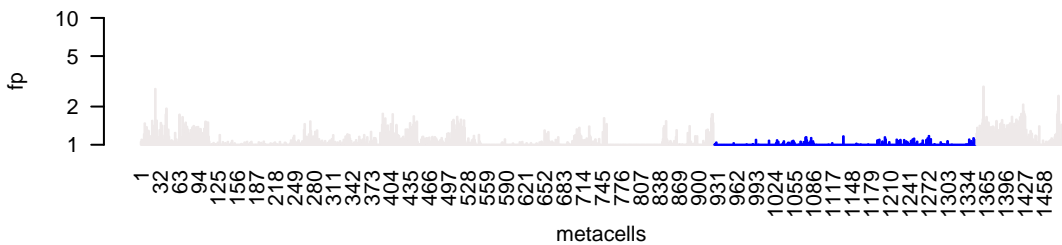
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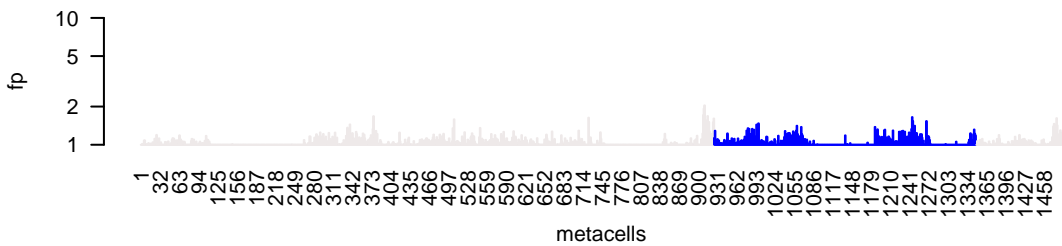
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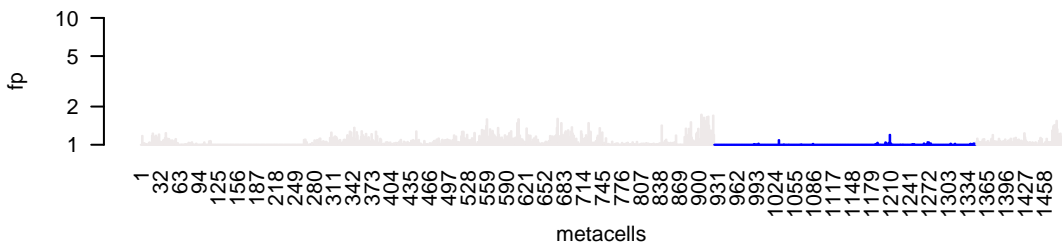
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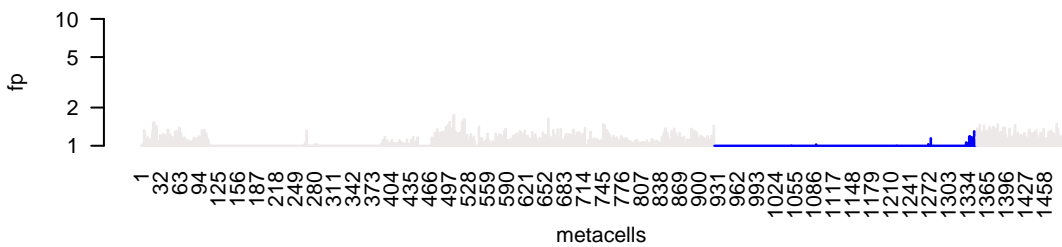
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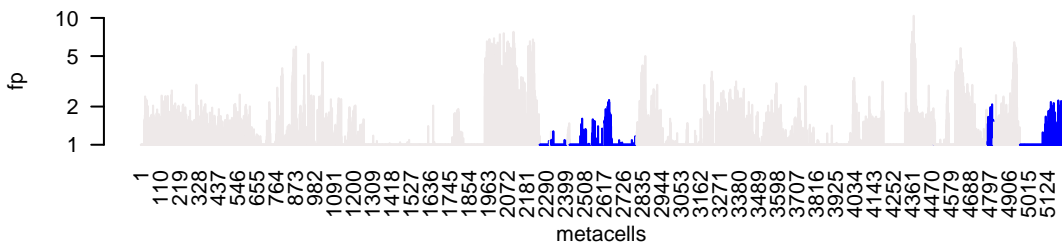
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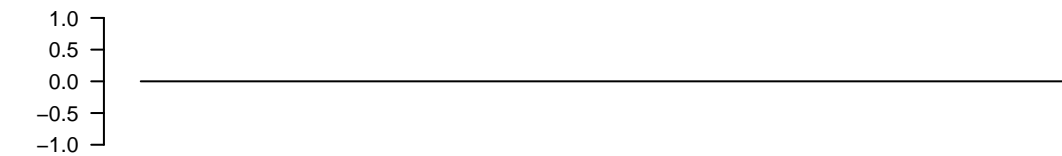
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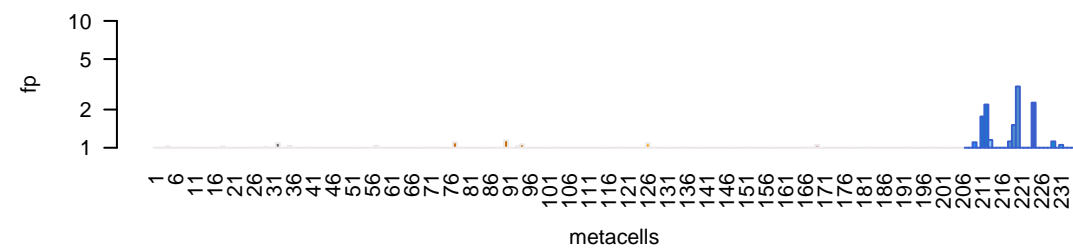
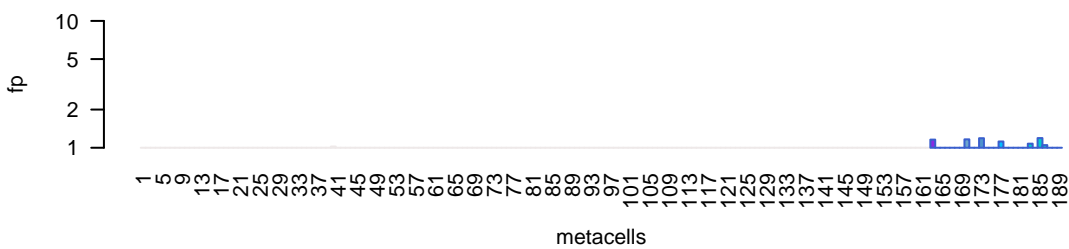
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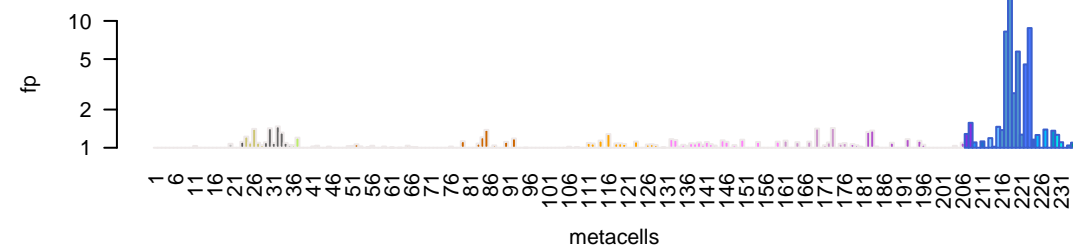
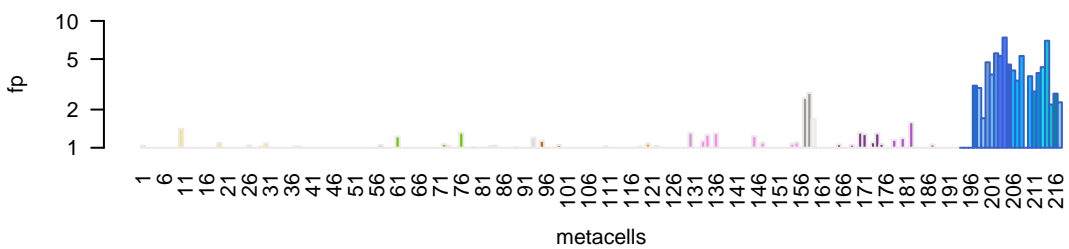
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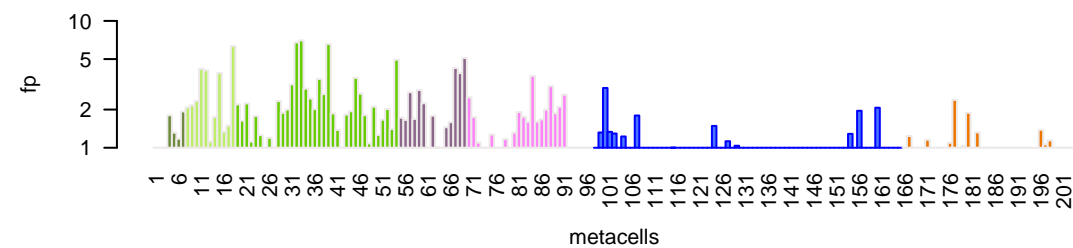
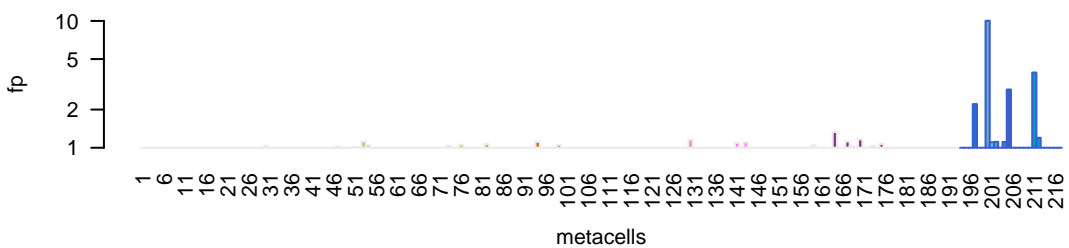
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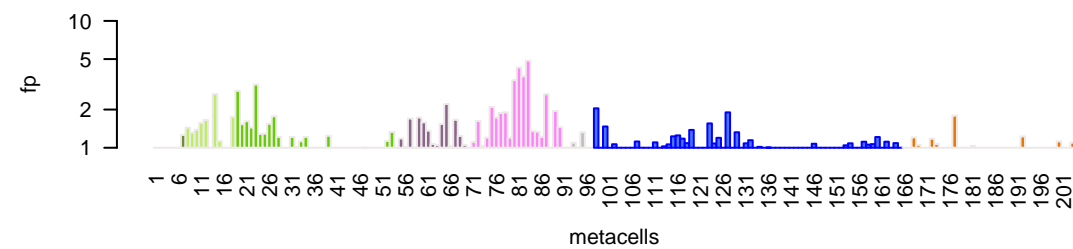
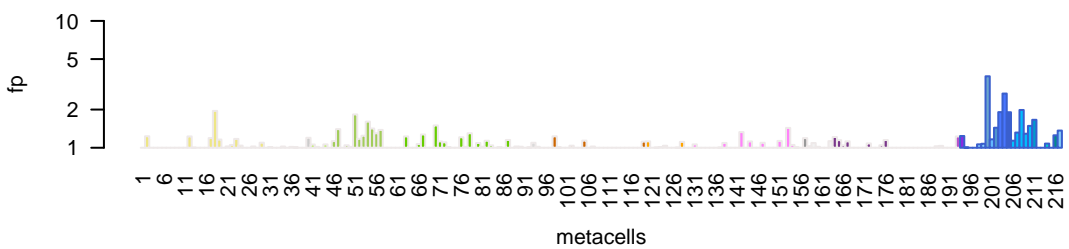
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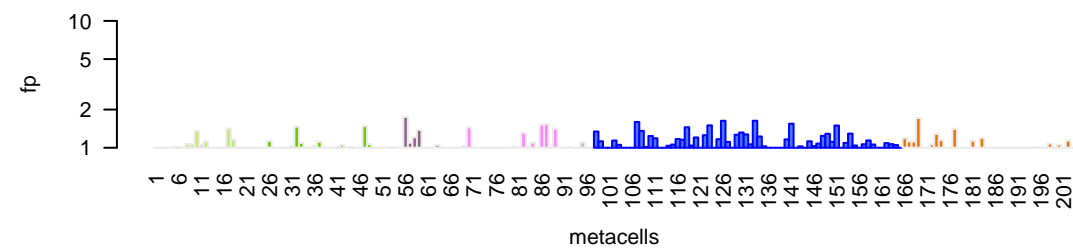
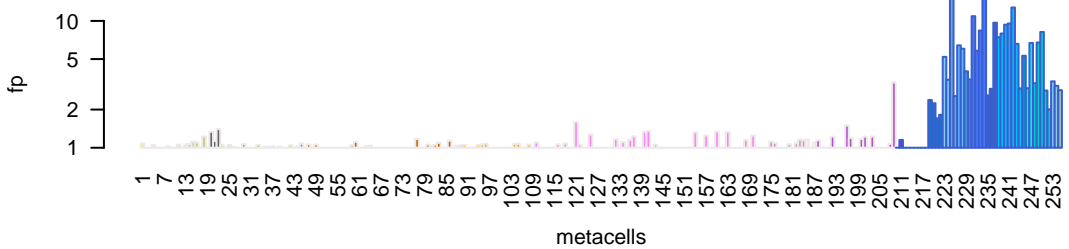
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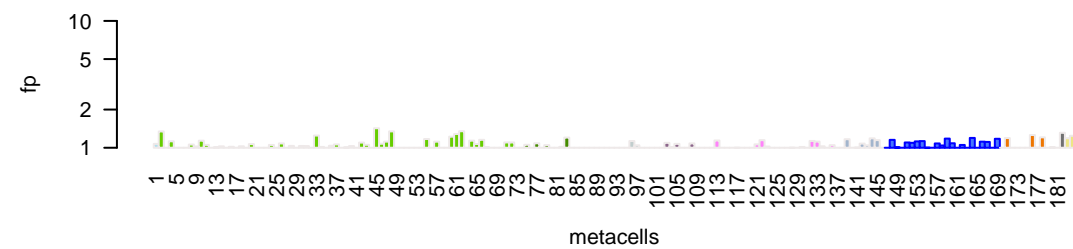
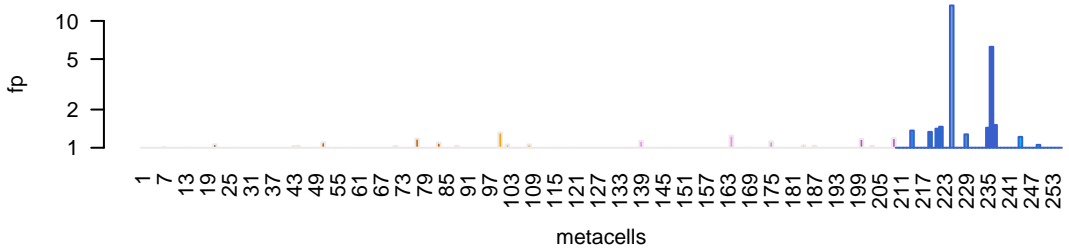
V3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GABPA G/ETV3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GA



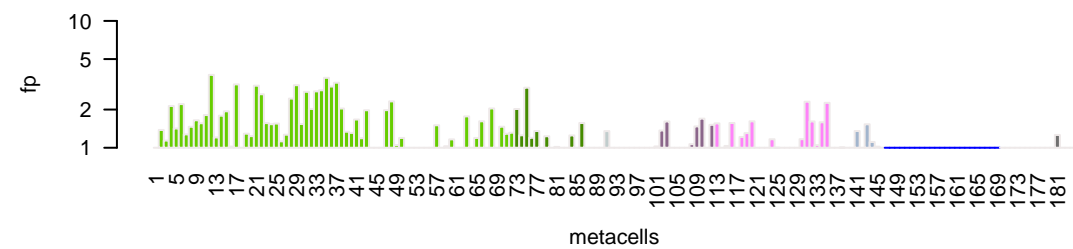
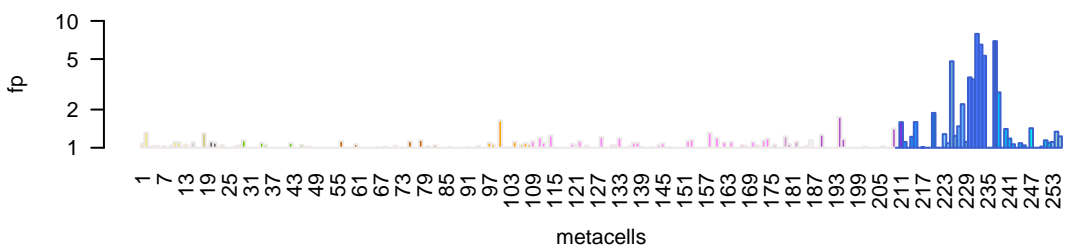
G/ETV3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GAG/ETV3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GA



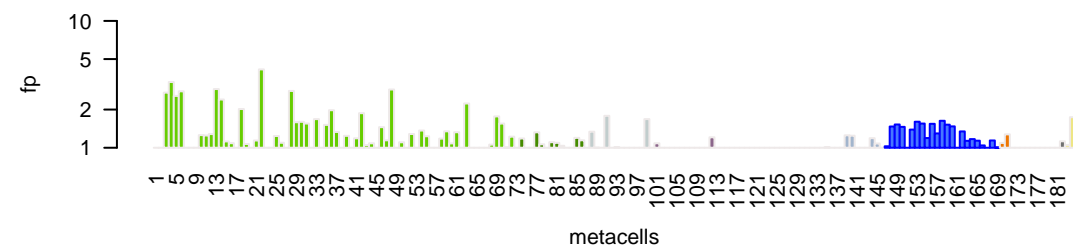
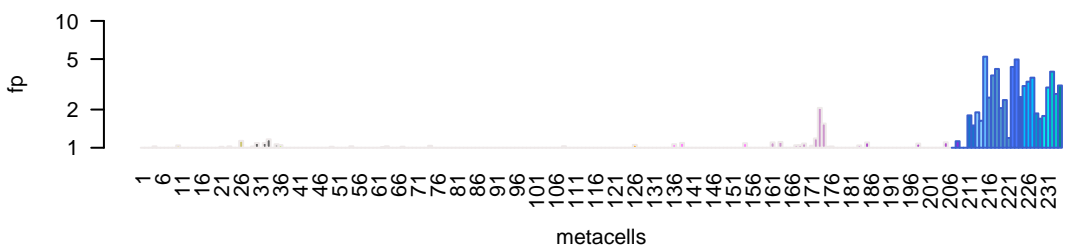
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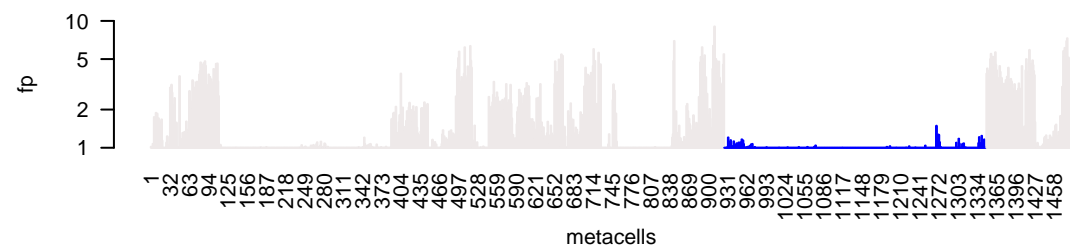
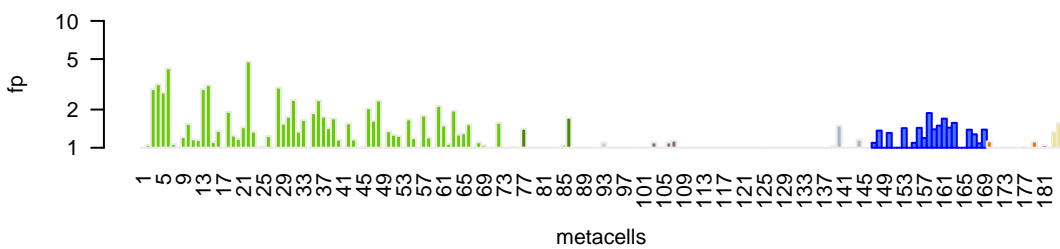
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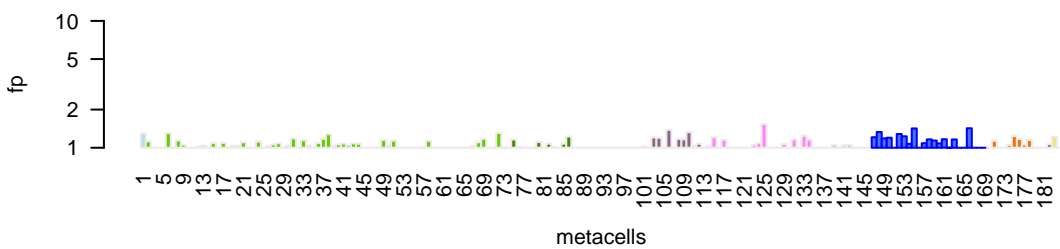
3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GABPA H:TV3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GABP



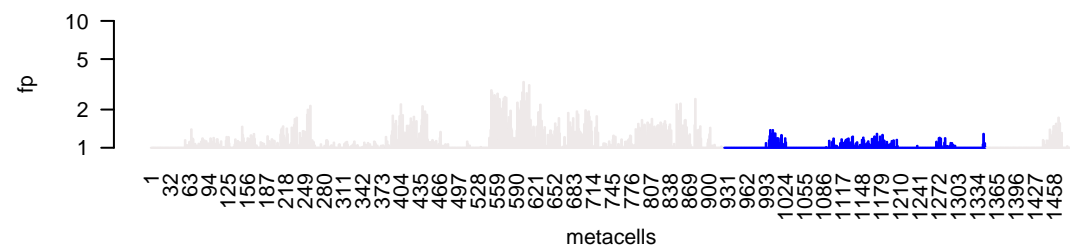
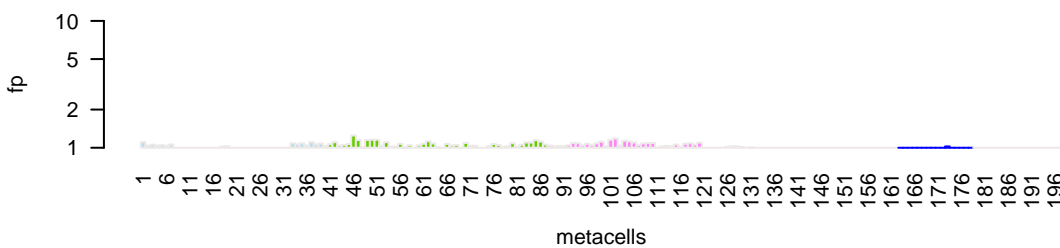
ETV3/ETV3L/FEV/FLI1/ETS proto oncogene 1 transcription factor/ETS1/ETS2/ETV2/GABP



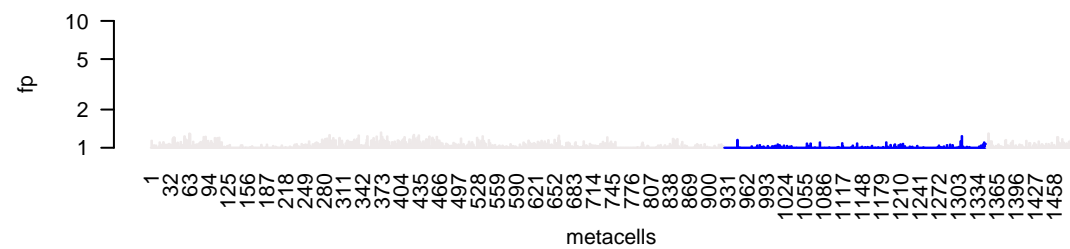
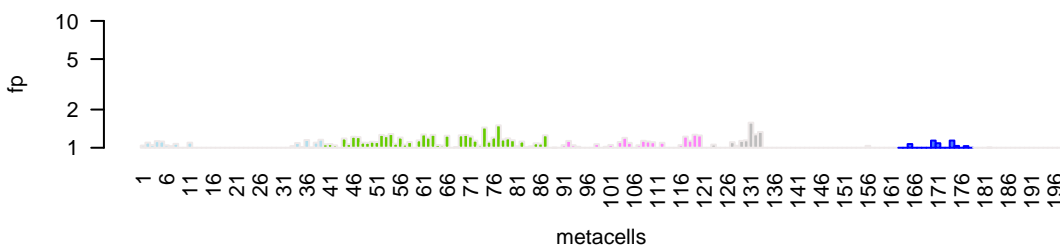
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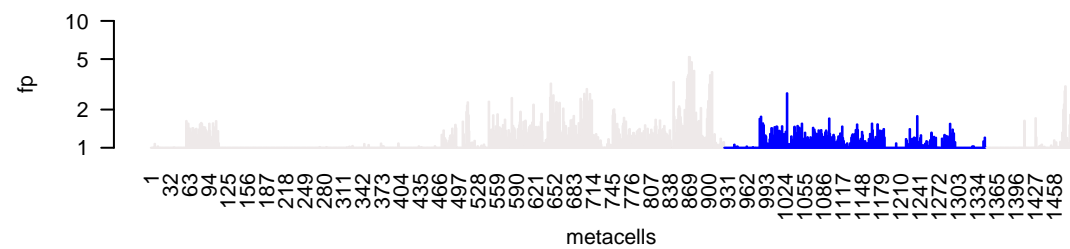
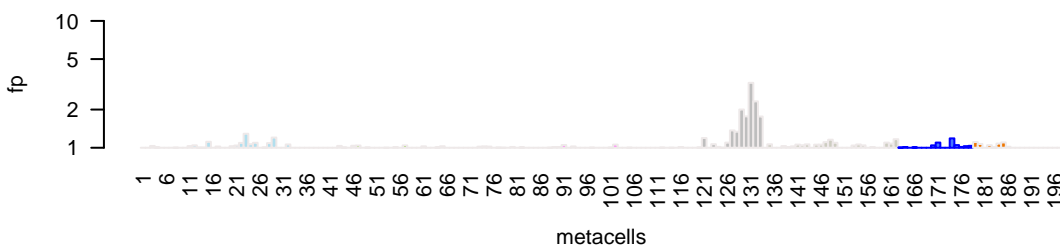
RG/ETV3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/G/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GABPA Mn



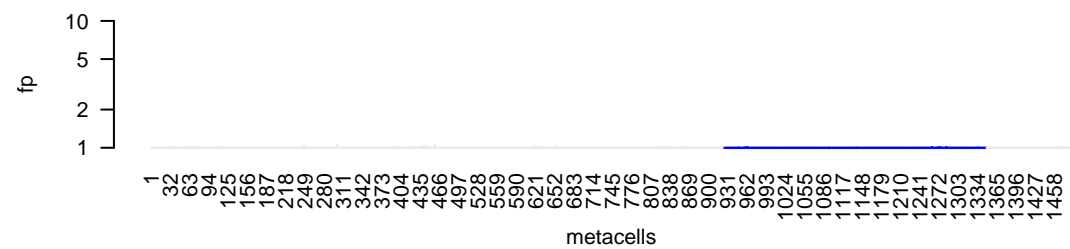
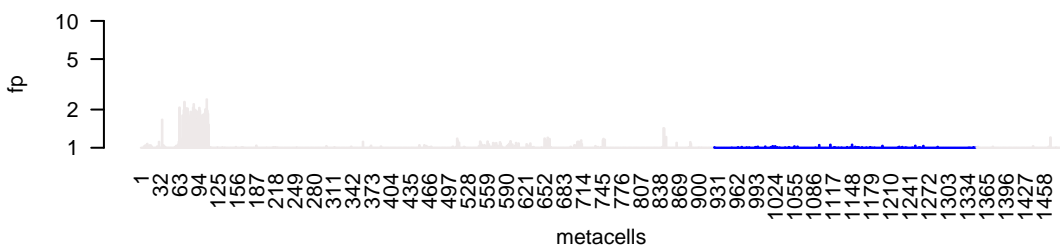
G/ETV3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/G/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GABPA Mm



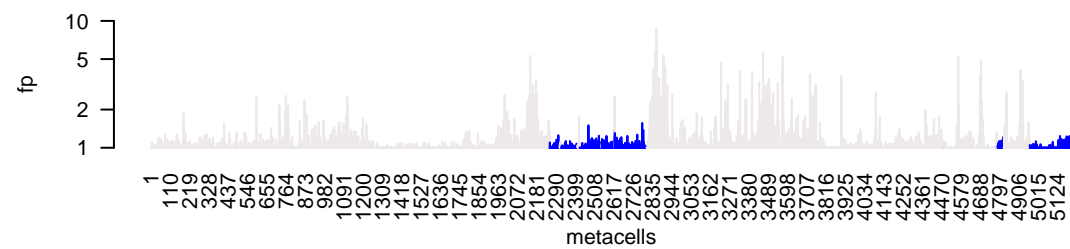
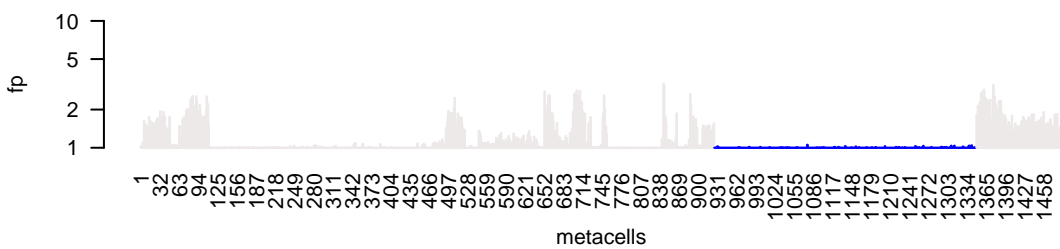
RG/ETV3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/G/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GABPA Mn



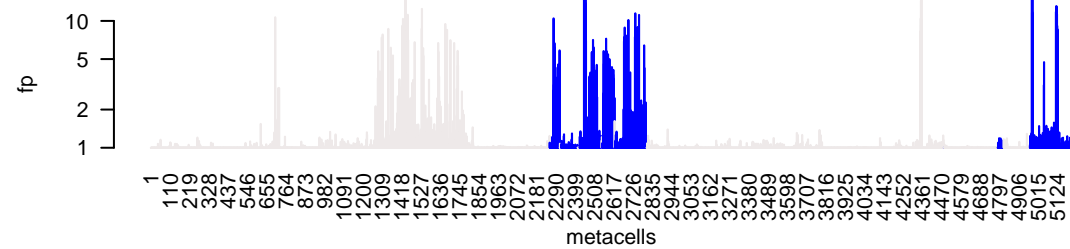
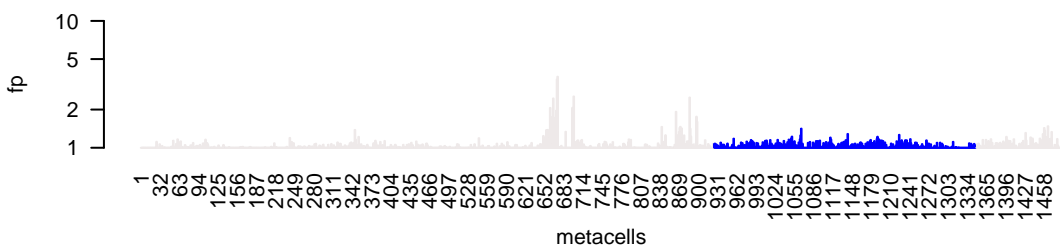
ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GABPA MrETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GABPA Mr



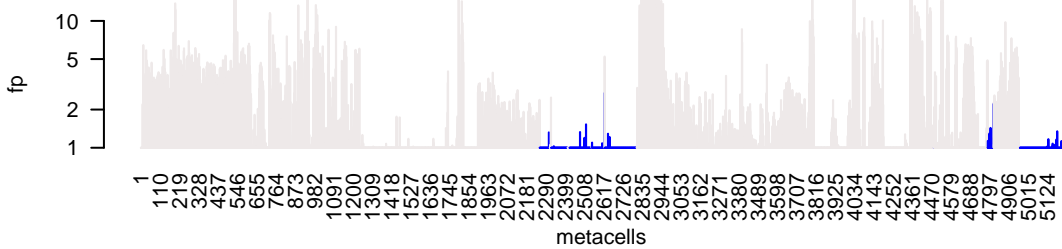
ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GABPA Mn/ETV3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV2/GAB



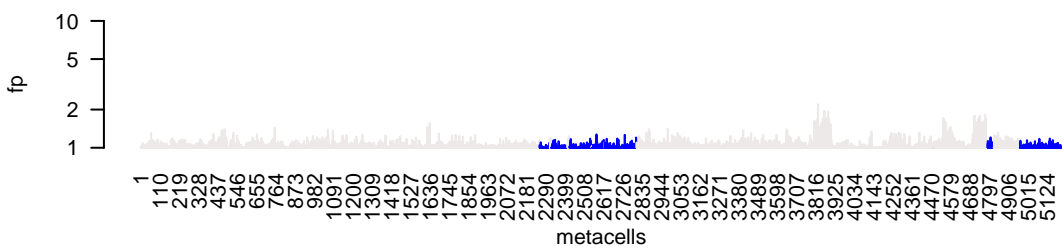
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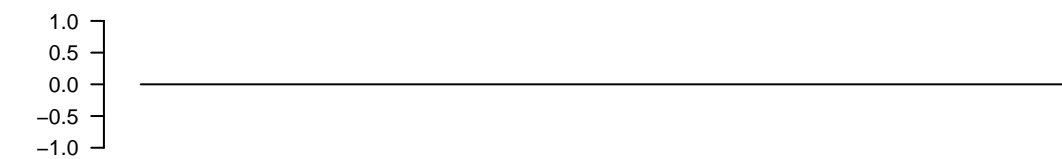
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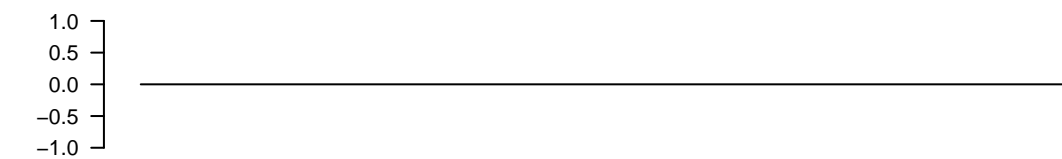
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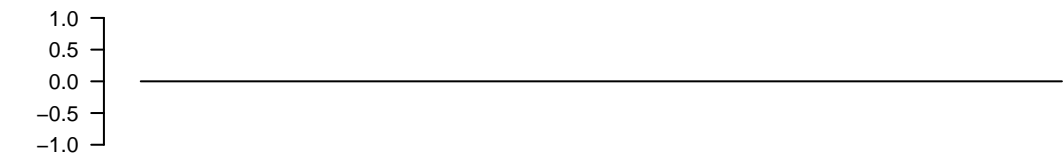
L/ERG/ETV3/ETV3L/FEV/FLI1/ETS_proto_oncogene_1_transcription_factor/ETS1/ETS2/ETV



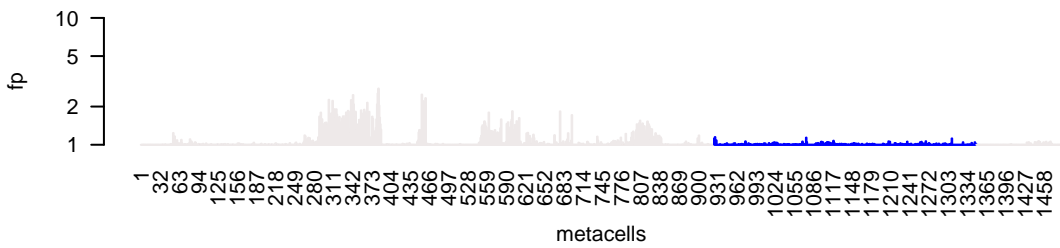
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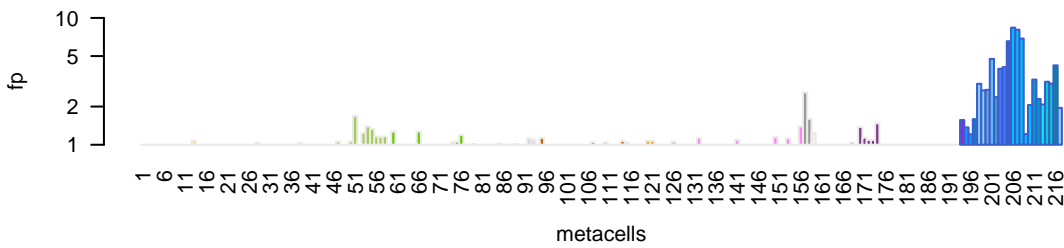
OG_10092:Etv1/ETV1/ETV4/ETV5 Tadh | no data



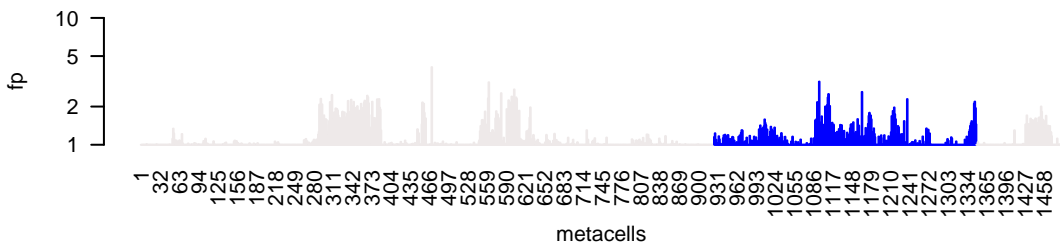
OG_10092:Etv1/ETV1/ETV4/ETV5 Mmus | Mmus_ENSMUST00000017868



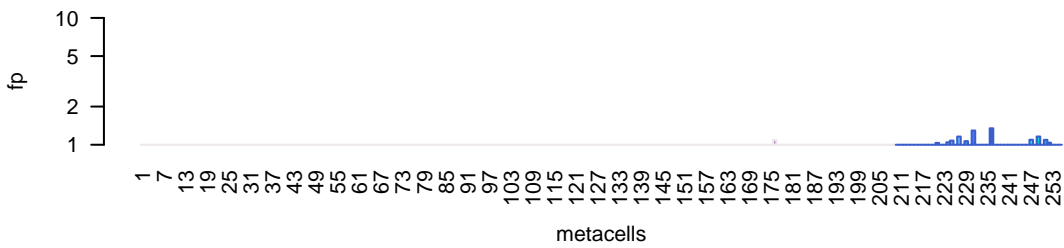
OG_10092:Etv1/ETV1/ETV4/ETV5 TrH2 | TrH2_TrispH2_002769-RA



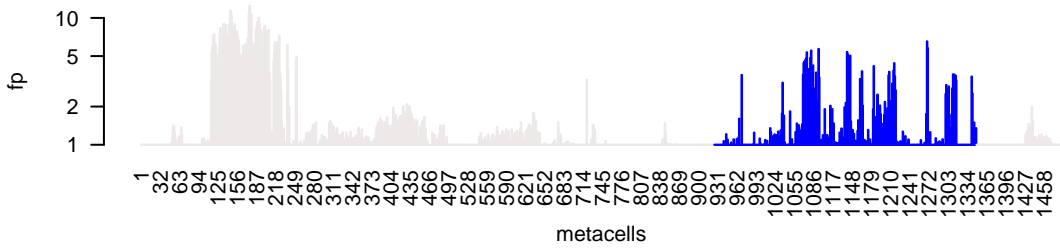
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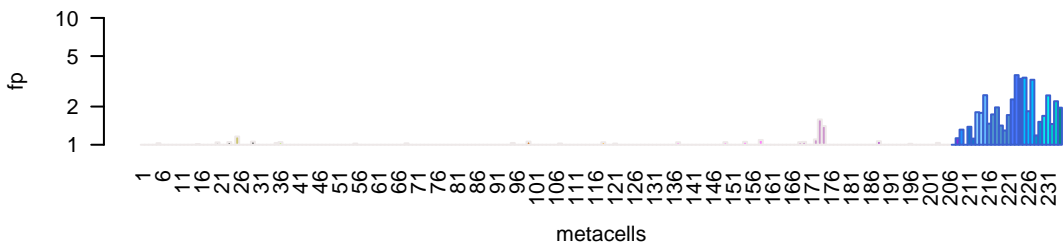
OG_10092:Etv1/ETV1/ETV4/ETV5 Hhon | Hhon_g00417.t1



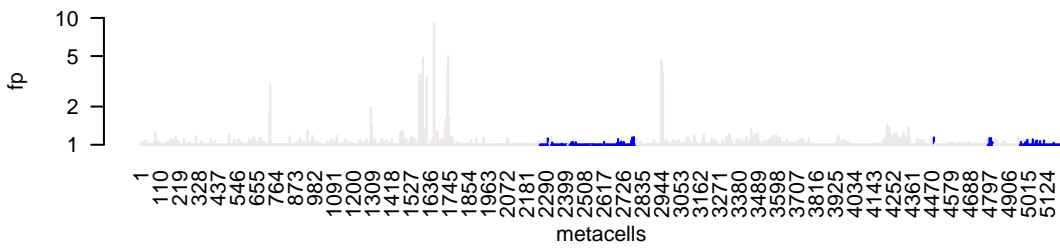
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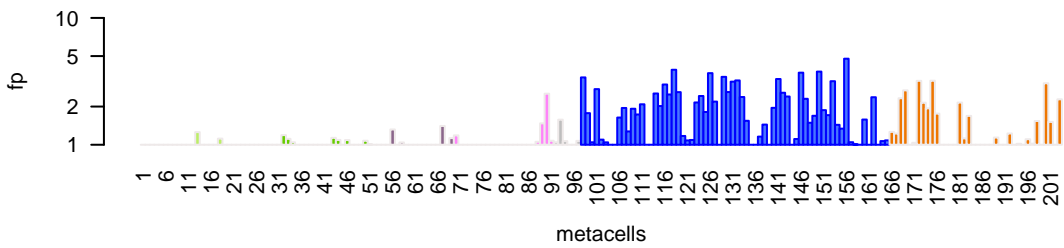
OG_10092:Etv1/ETV1/ETV4/ETV5 HoiH23 | HoiH23_PIH23_001793-RA



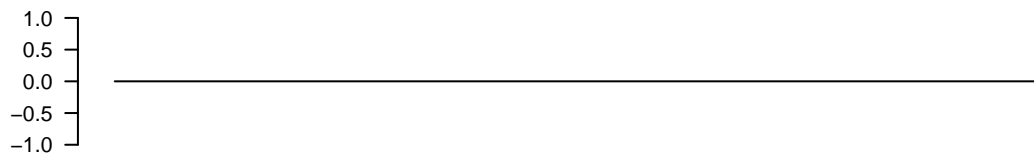
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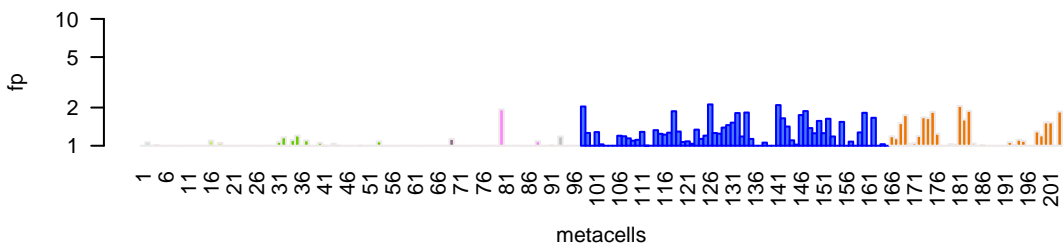
OG_10092:Etv1/ETV1/ETV4/ETV5 Nvec | Nvec_v1g16815



OG_10092:Etv1/ETV1/ETV4/ETV5 Mlei | no data



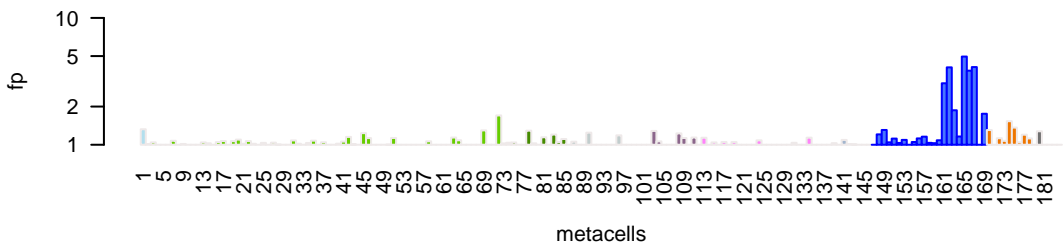
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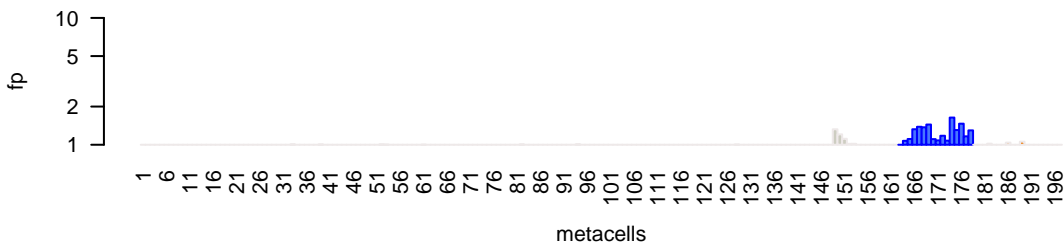
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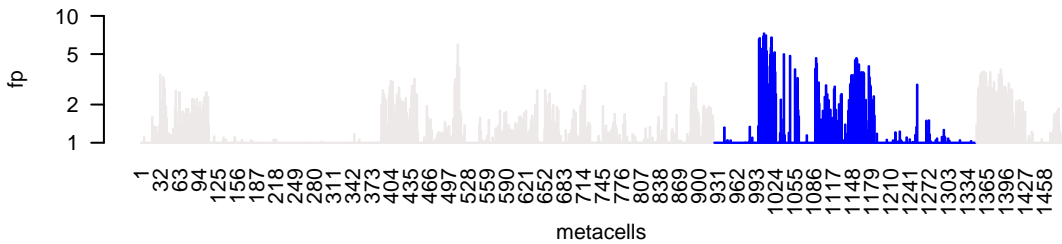
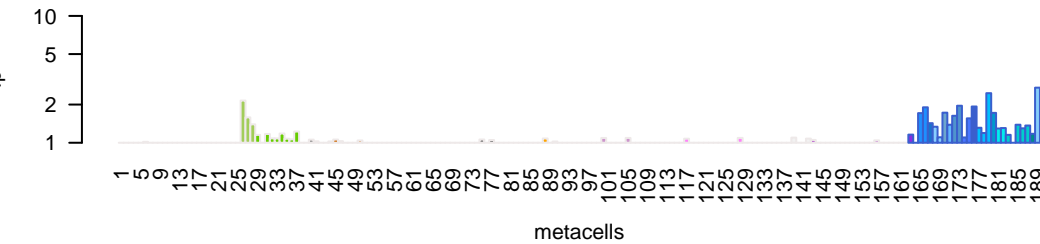
OG_10092:Etv1/ETV1/ETV4/ETV5 Spis | Spis_XP_022794506_1



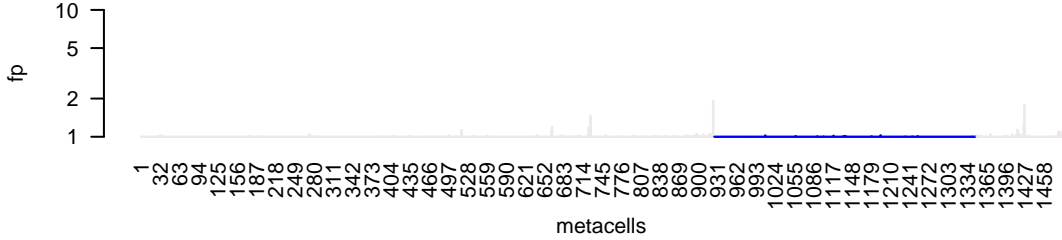
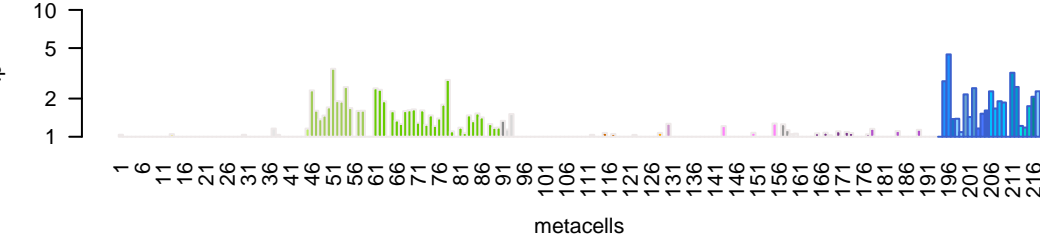
OG_10092:Etv1/ETV1/ETV4/ETV5 Hvul | Hvul_g28310_1



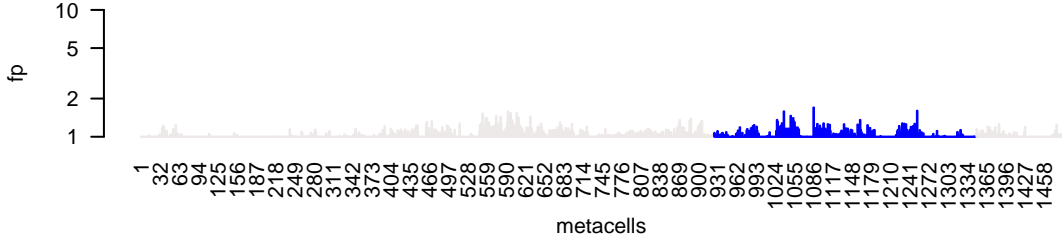
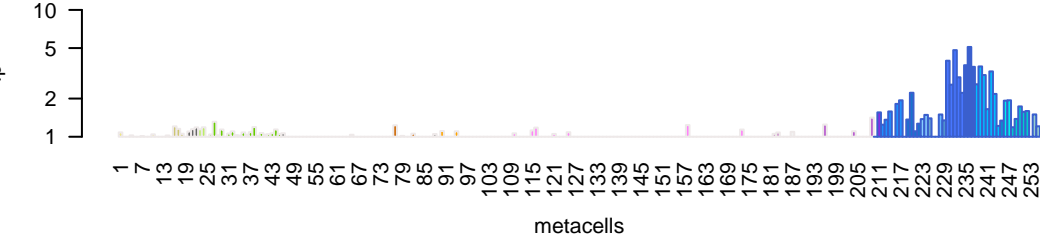
OG_10429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 Tadh | Tadh_TriadT24510429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 Mmus | Mmus_ENSMUST0000



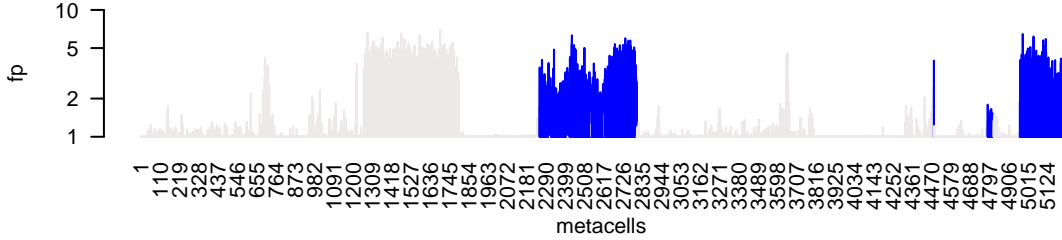
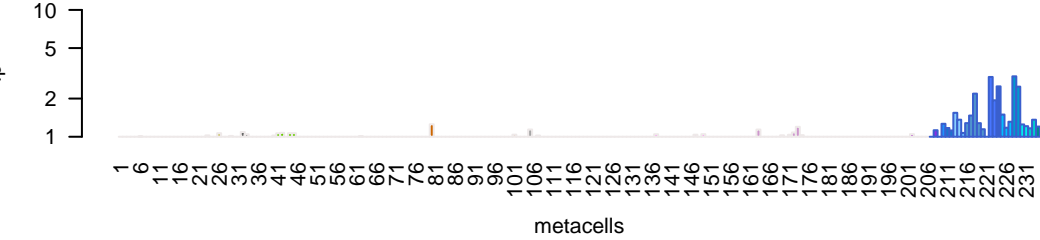
G_10429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 TrH2 | TrH2_TrispH2_0018010429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 Mmus | Mmus_ENSMUST0000



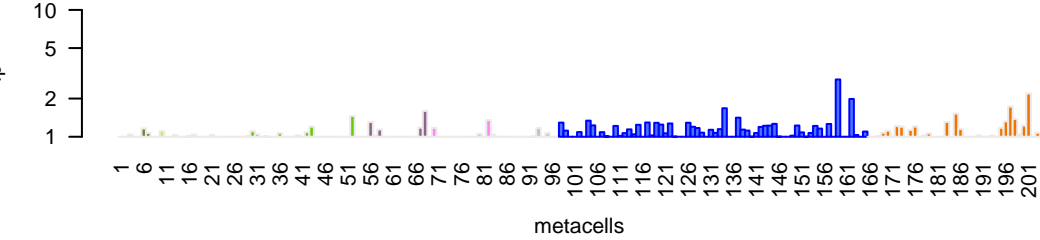
OG_10429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 Hhon | Hhon_g02158.t10429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 Mmus | Mmus_ENSMUST0000



_10429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 HoiH23 | HoiH23_PIH23_002: OG_10429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 Dmel | Dmel_FBtr03048



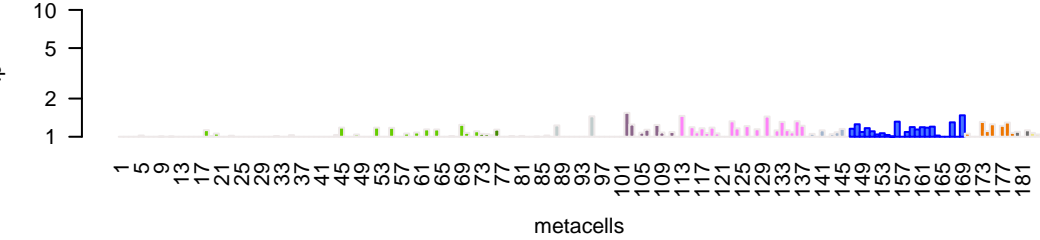
OG_10429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 Nvec | Nvec_v1g16100



OG_10429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 Mlei | no data



OG_10429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 Spis | Spis_XP_02280116



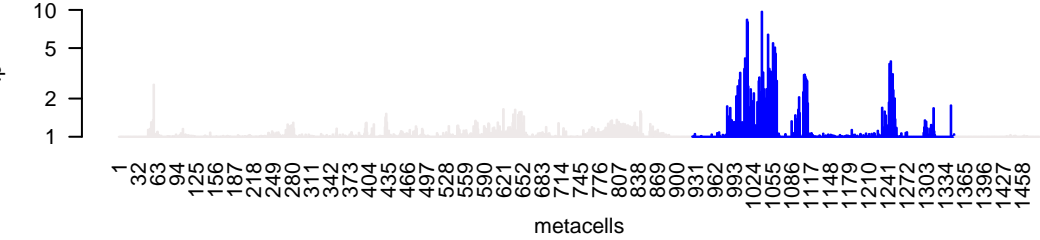
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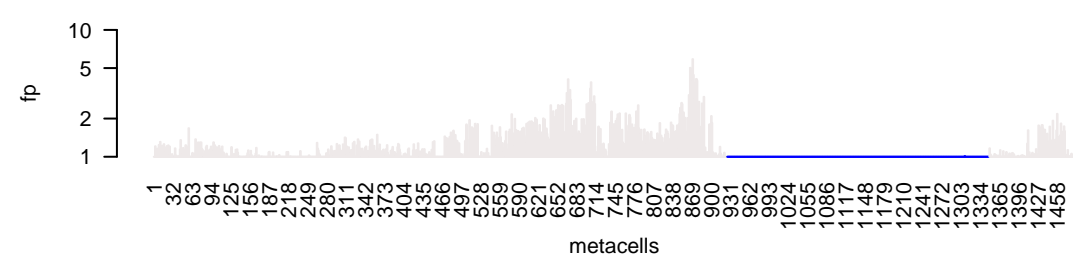
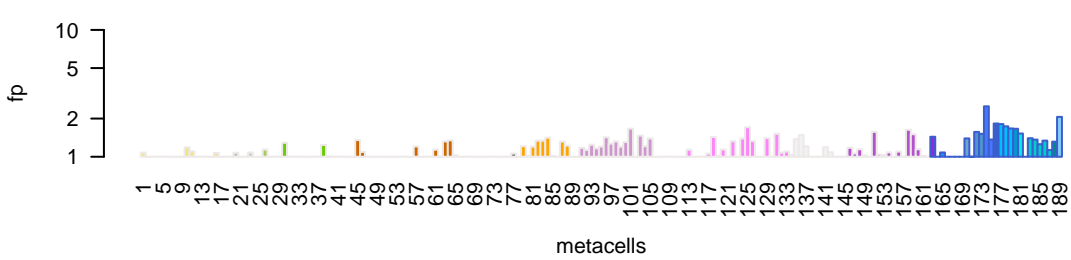
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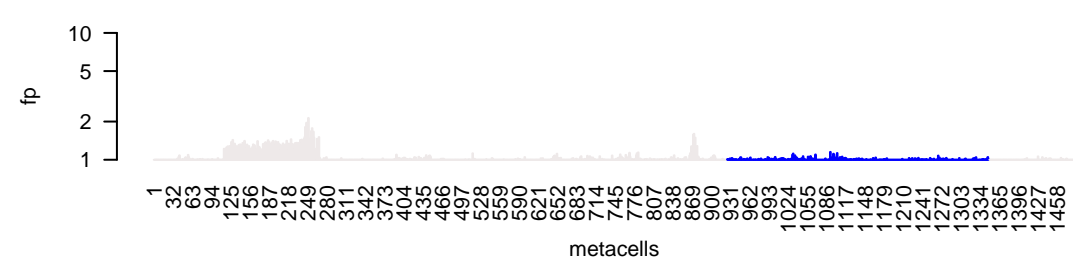
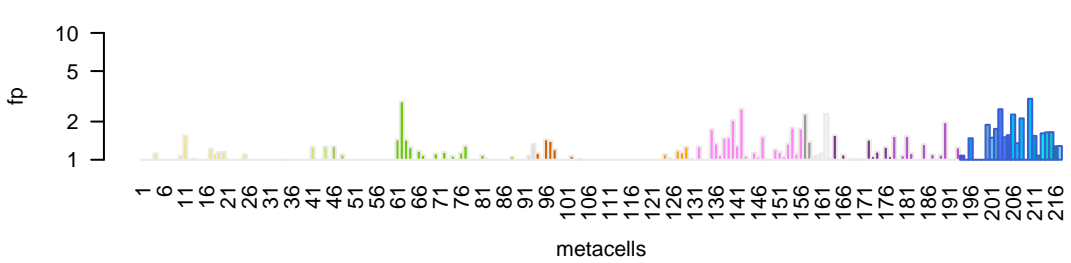
10429:AC097634.4/FOXP1/FOXP2/FOXP3/FOXP4/Foxp1/Foxp2 Mmus | Mmus_ENSMUST0000



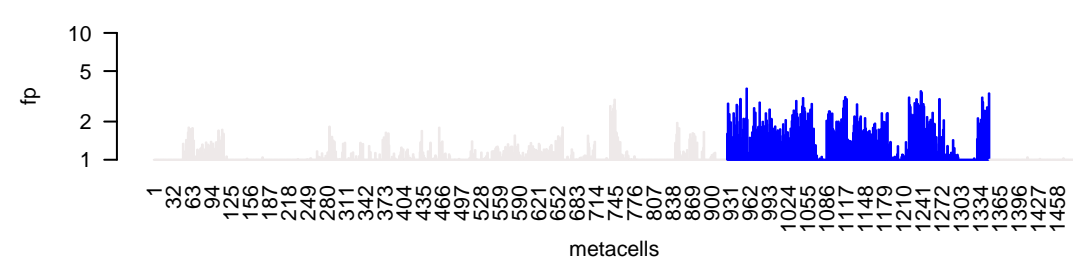
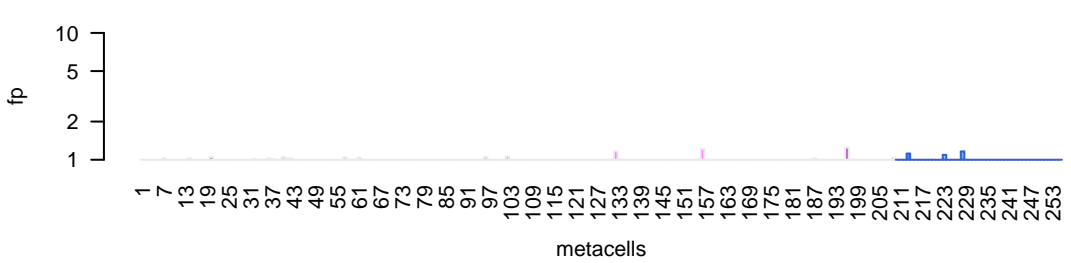
OG_749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Tadh | Tadh_TriadT60616; _749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Mmus | Mmus_ENSMUST00000



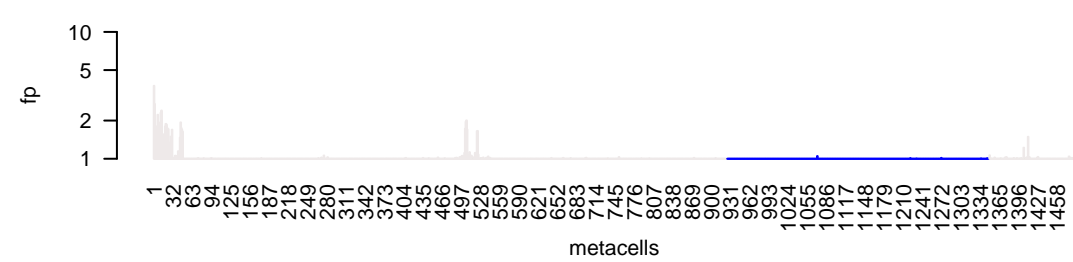
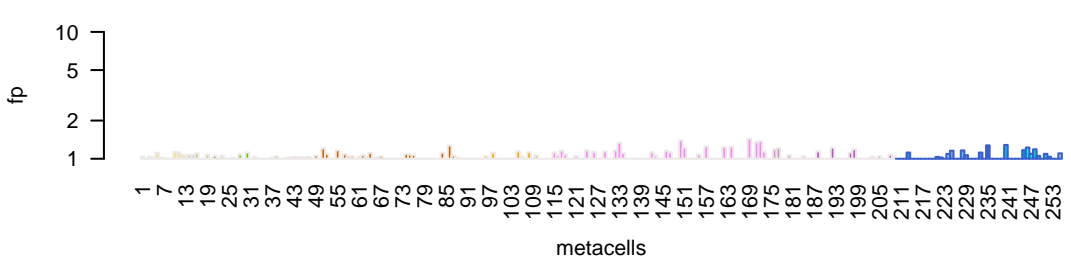
OG_749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 TrH2 | TrH2_TrispH2_010892; _749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Mmus | Mmus_ENSMUST00000



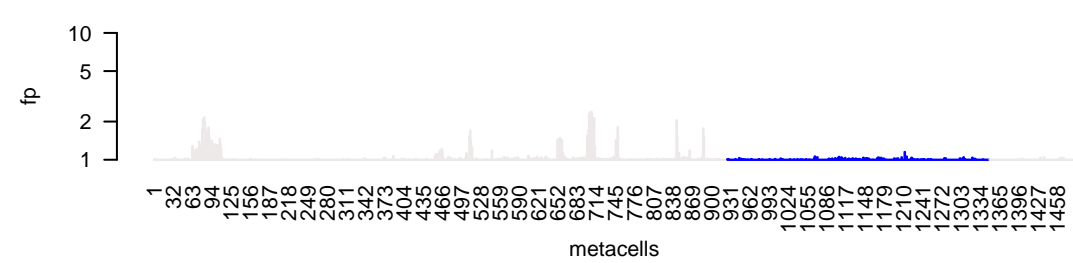
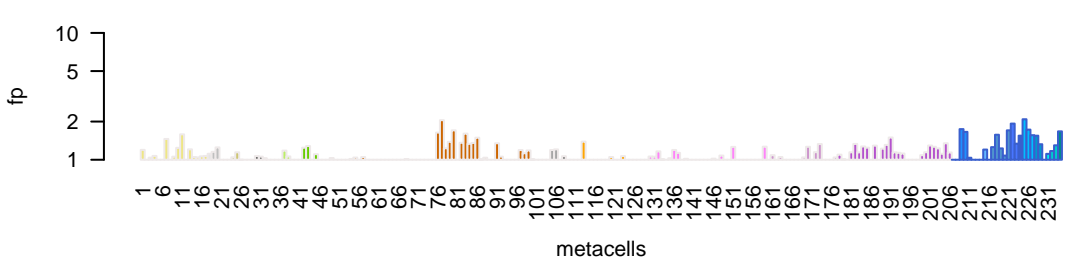
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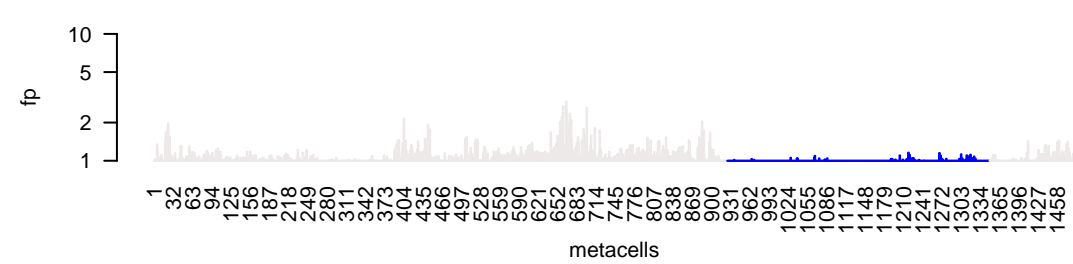
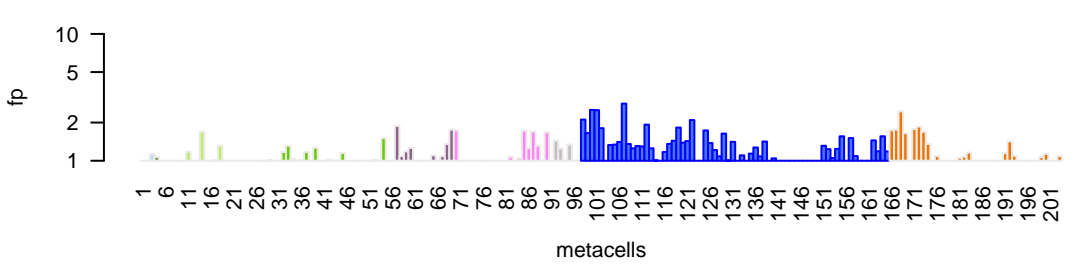
OG_749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Hhon | Hhon_g09126.t1; _749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Mmus | Mmus_ENSMUST00000



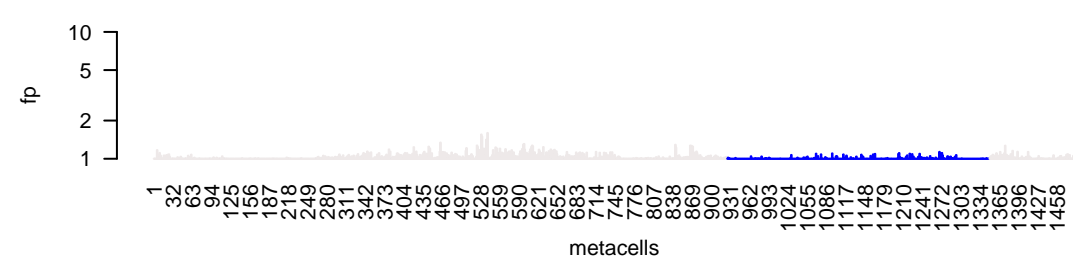
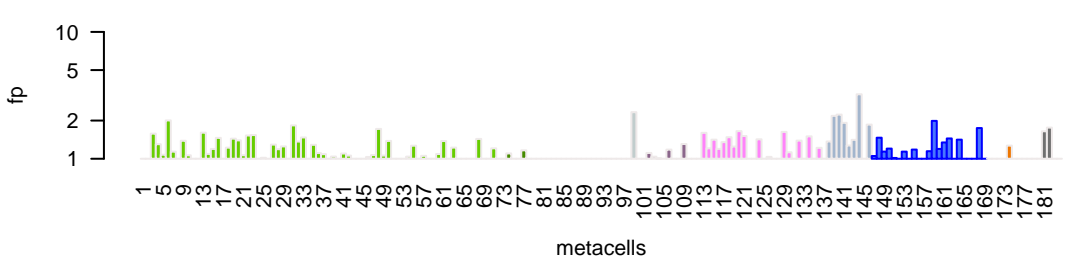
G_749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 HoiH23 | HoiH23_PIH23_00449; _749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Mmus | Mmus_ENSMUST00000



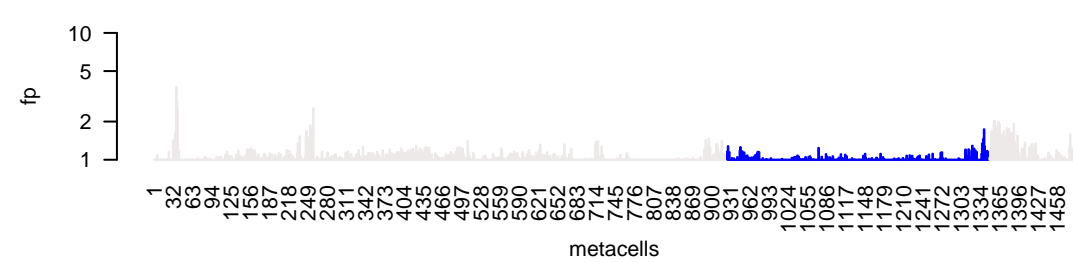
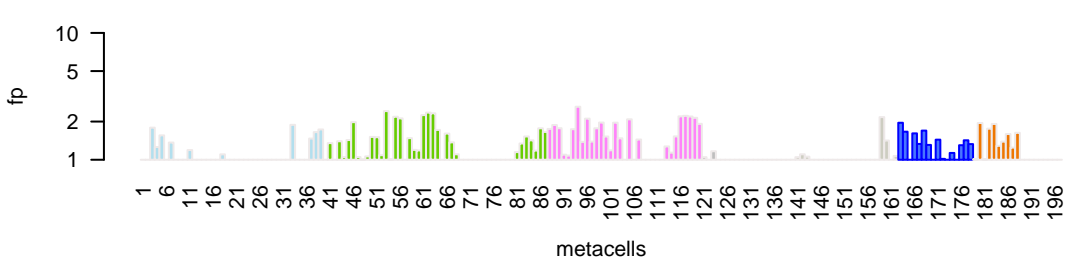
OG_749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Nvec | Nvec_v1g127893; _749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Mmus | Mmus_ENSMUST00000



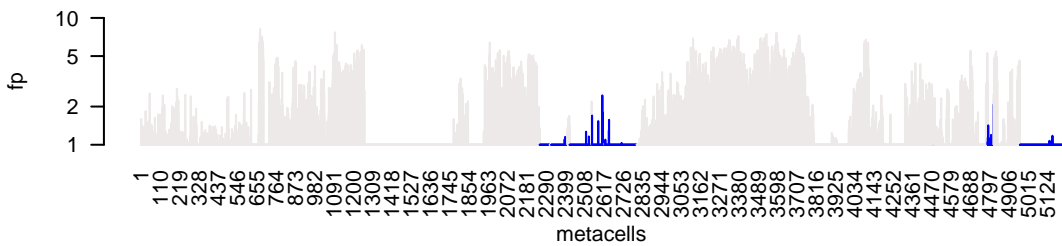
OG_749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Spis | Spis_XP_022779261; _749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Mmus | Mmus_ENSMUST00000



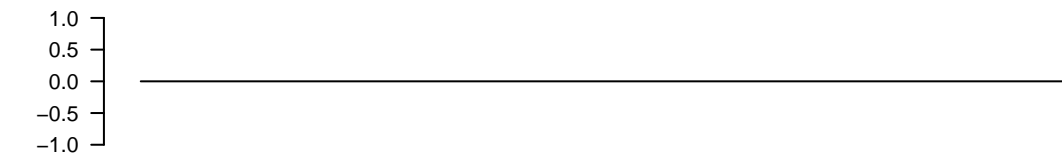
OG_749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 HvuI | HvuI_g28368_1; _749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Mmus | Mmus_ENSMUST00000



OG_749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Dmel | Dmel_FBtr0330372

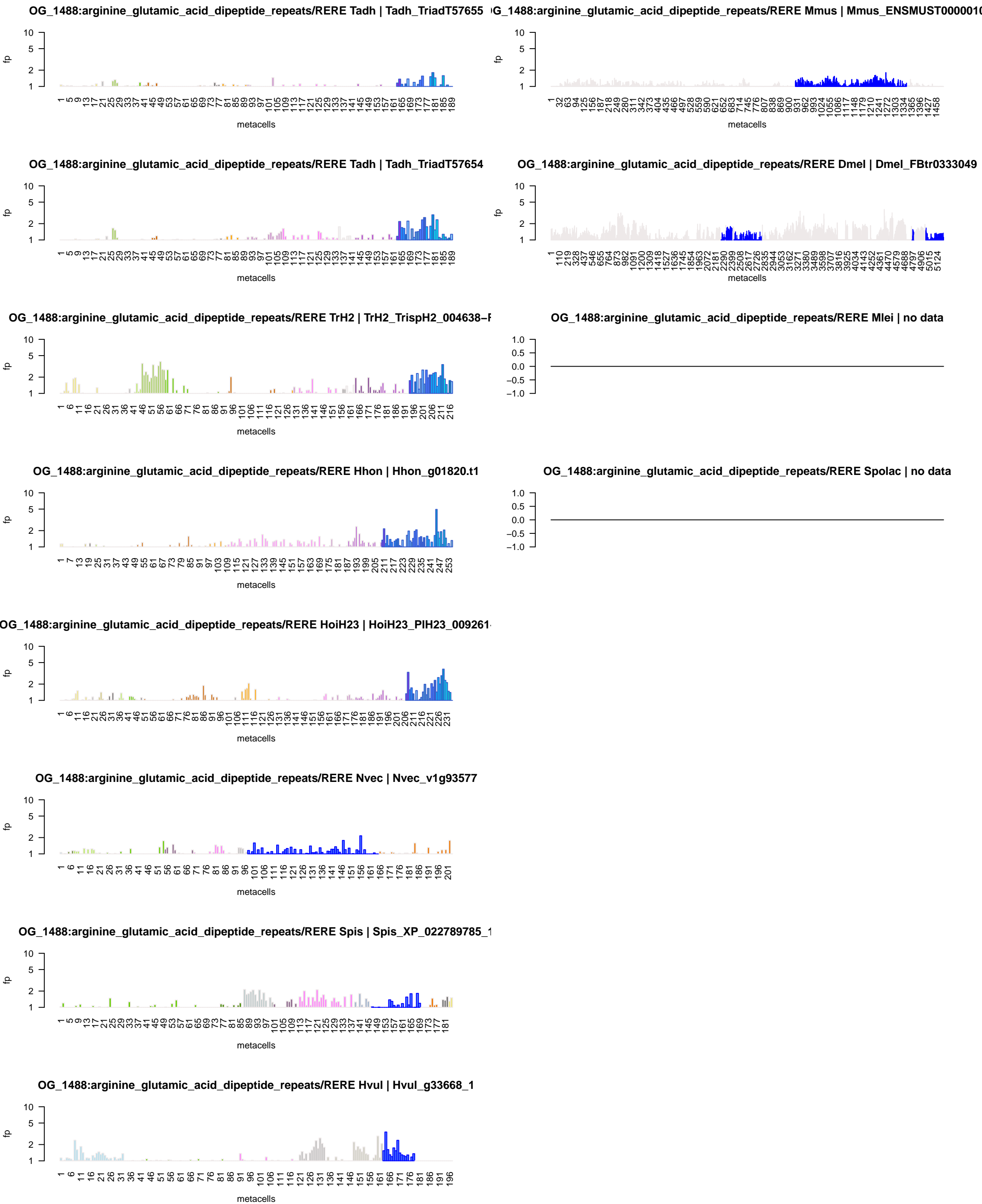


OG_749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Mlei | no data

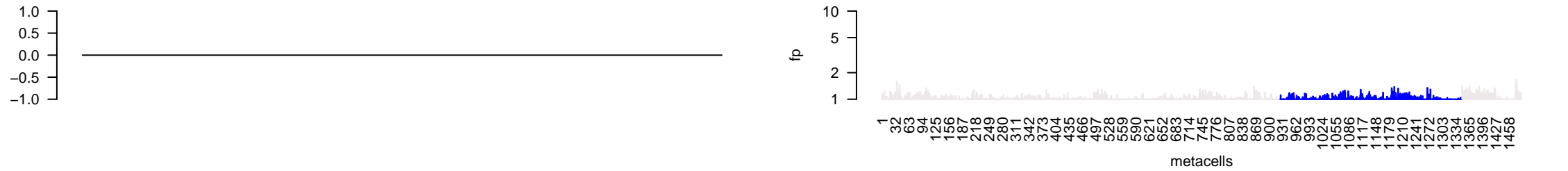


OG_749:BACH1/BACH2/NFE2/NFE2L1/NFE2L2/NFE2L3/Nfe2l1 Spolac | no data

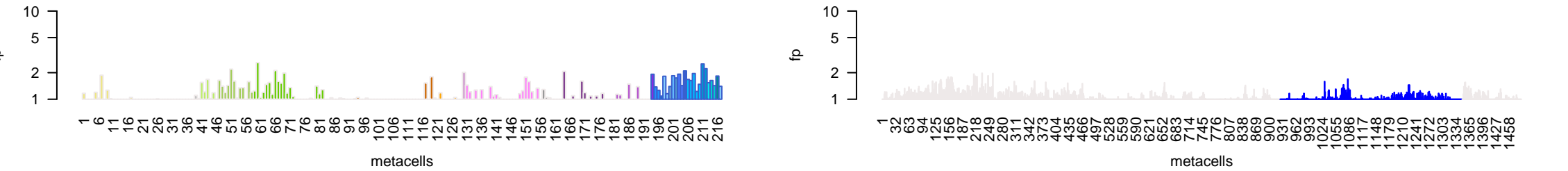




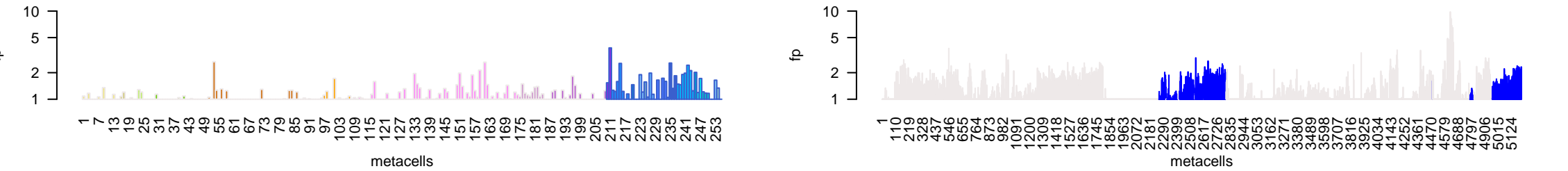
OG_3560:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Tadh | no daBAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Mmus | Mmus_ENSMUS



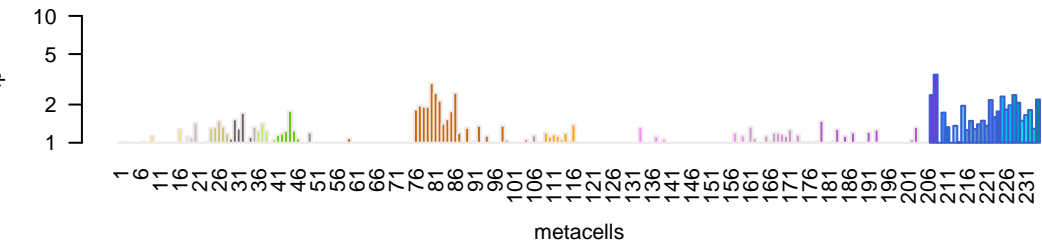
60:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B TrH2 | TrH2_TrispH2_BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Mmus | Mmus_ENSMUS



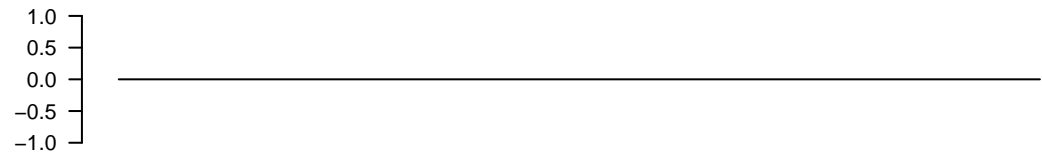
3560:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Hhon | Hhon_g0_3560:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Dmel | Dmel_FBtr



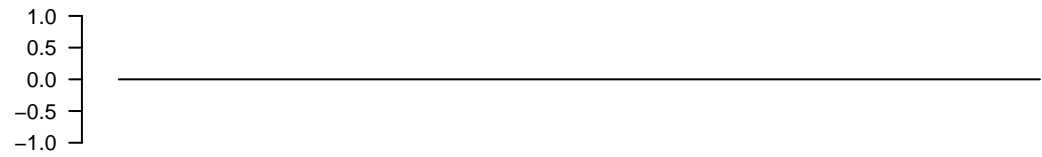
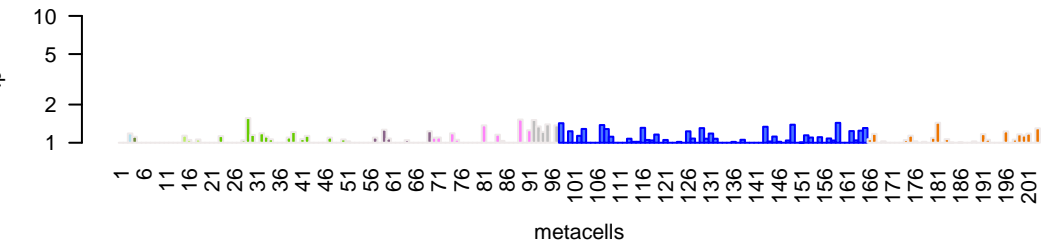
:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B HoiH23 | HoiH23_PIH23



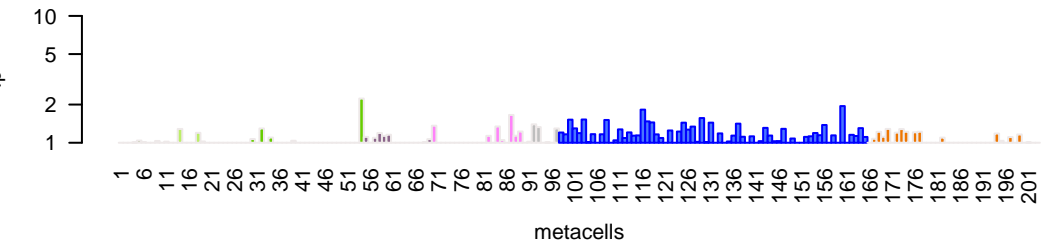
OG_3560:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Mlei | no da



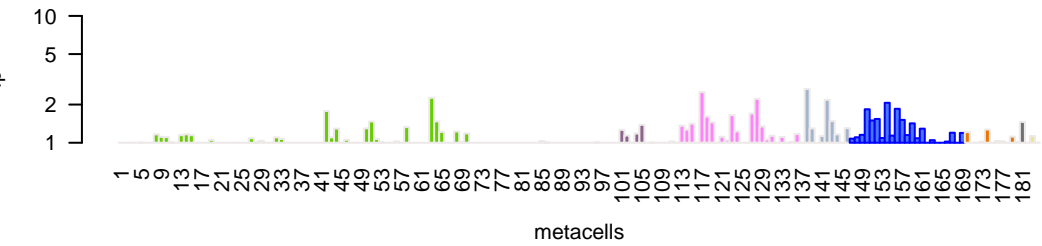
3560:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Nvec | Nvec_v13560:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Spolac | no d



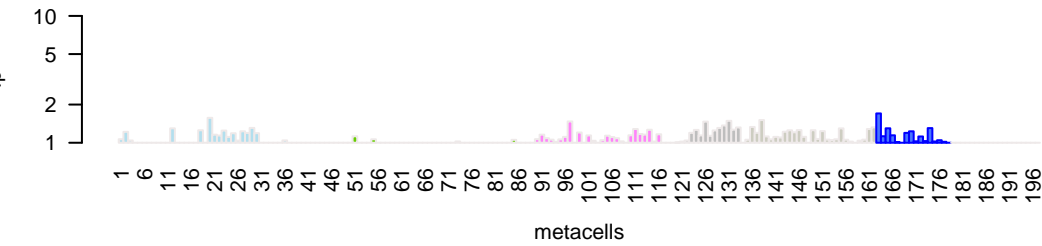
3560:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Nvec | Nvec_v13560:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Spis | Spis_XP_022



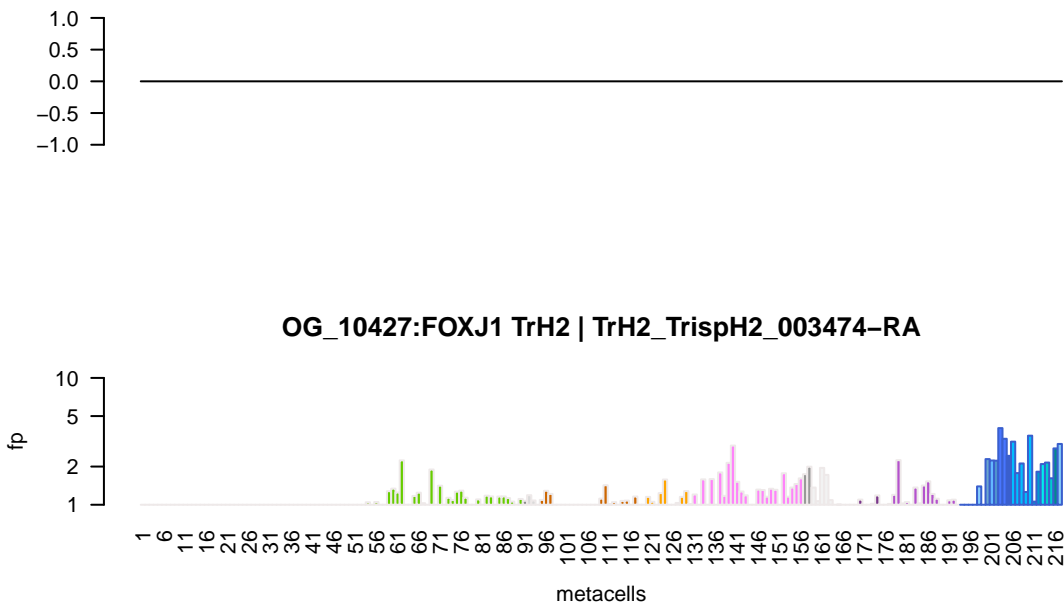
560:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Spis | Spis_XP_022



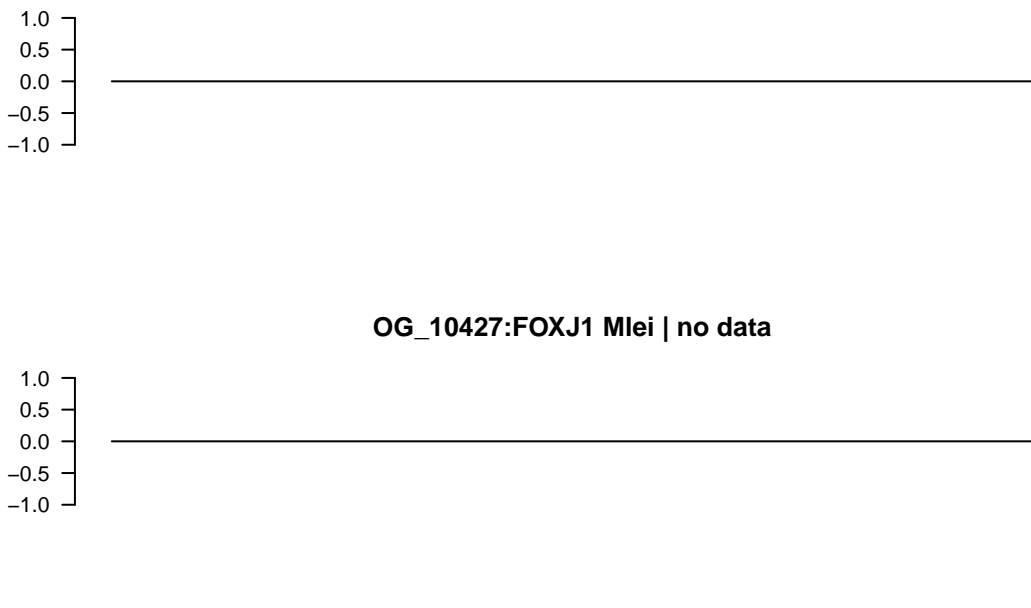
3560:BAZ2A/BAZ2B/bromodomain_adjacent_to_zinc_finger_domain_2B Hvul | Hvul_g84



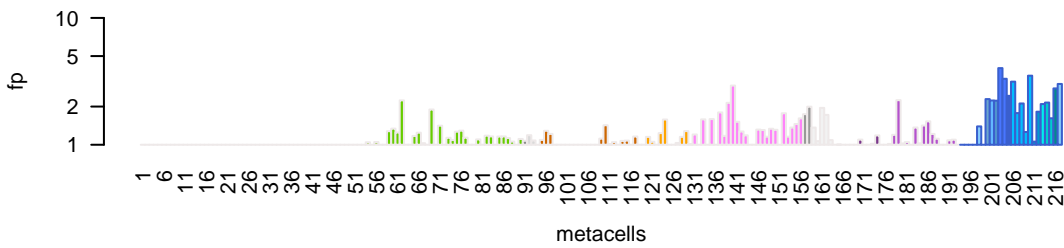
OG_10427:FOXJ1 Tadh | no data



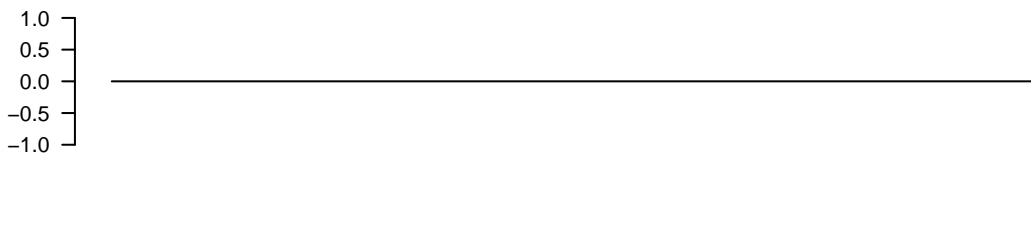
OG_10427:FOXJ1 Dmel | no data



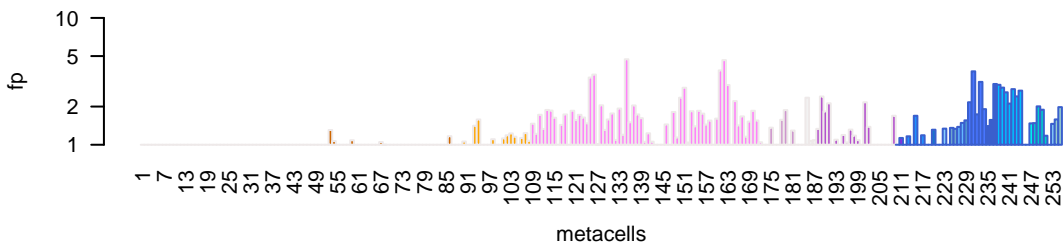
OG_10427:FOXJ1 TrH2 | TrH2_TrispH2_003474-RA



OG_10427:FOXJ1 Mlei | no data



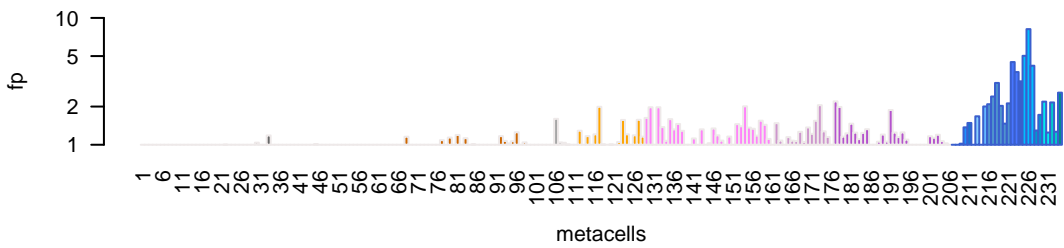
OG_10427:FOXJ1 Hhon | Hhon_g03786.t1



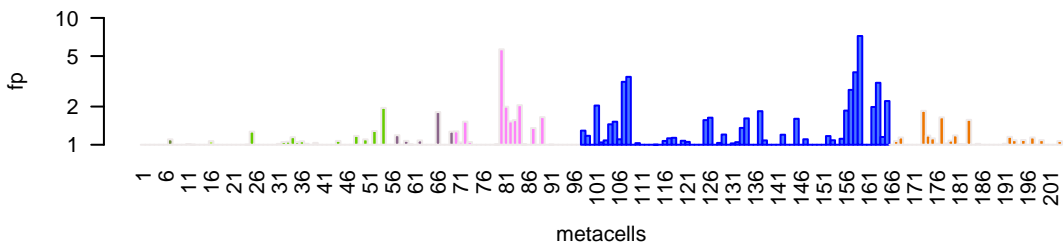
OG_10427:FOXJ1 Spolac | no data



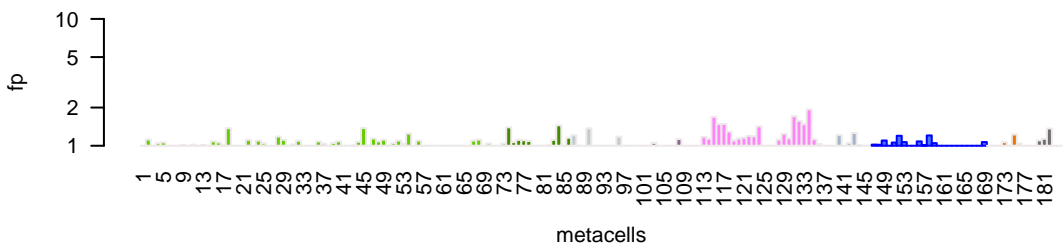
OG_10427:FOXJ1 HoiH23 | HoiH23_PIH23_000096-RA



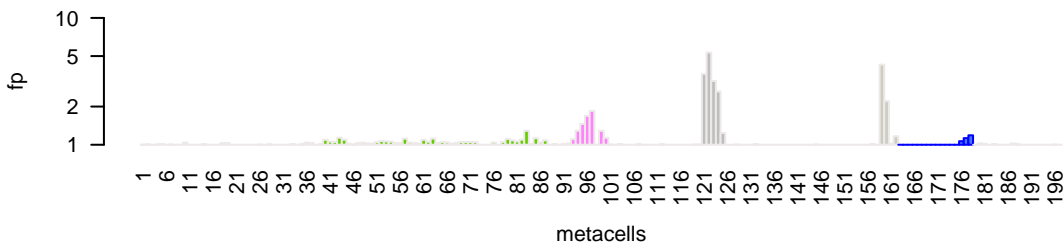
OG_10427:FOXJ1 Nvec | Nvec_v1g65438



OG_10427:FOXJ1 Spis | Spis_XP_022784179_1



OG_10427:FOXJ1 Hvul | Hvul_g8279_1



OG_10427:FOXJ1 Mmus | Mmus_ENSMUST00000036215

