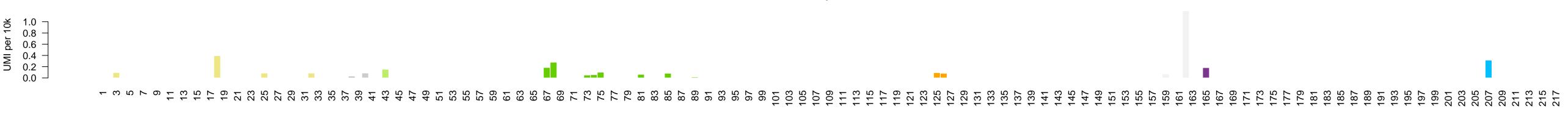
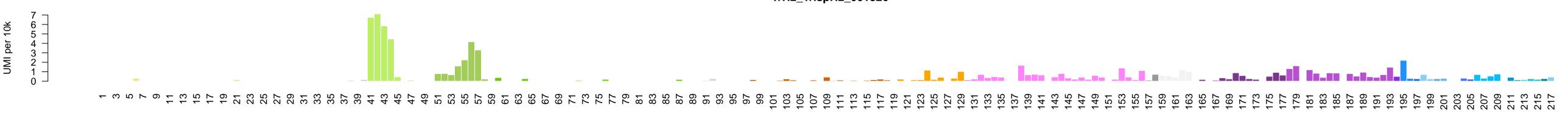


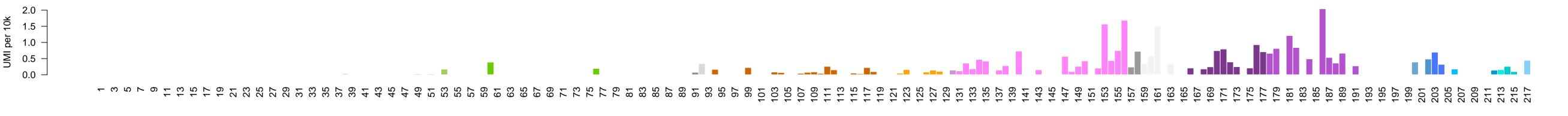
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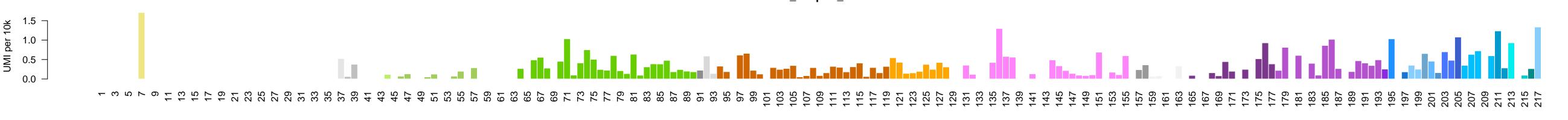
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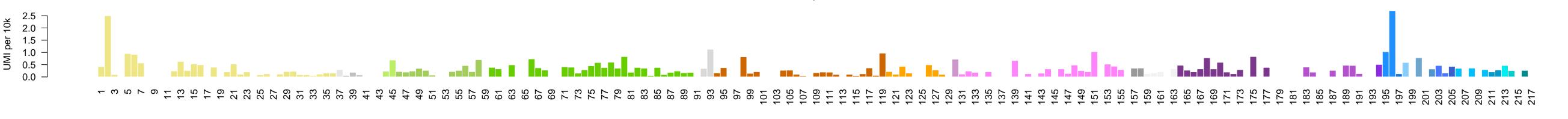
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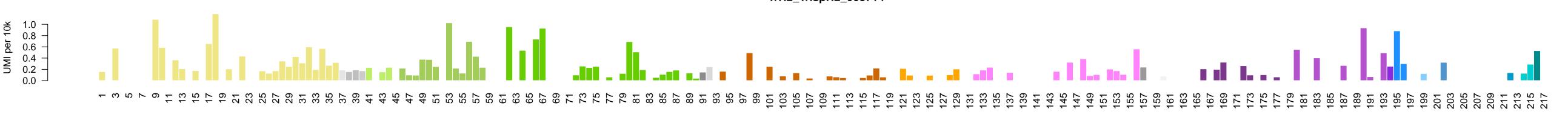
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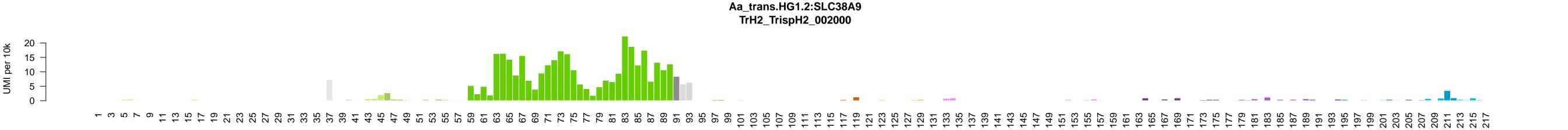


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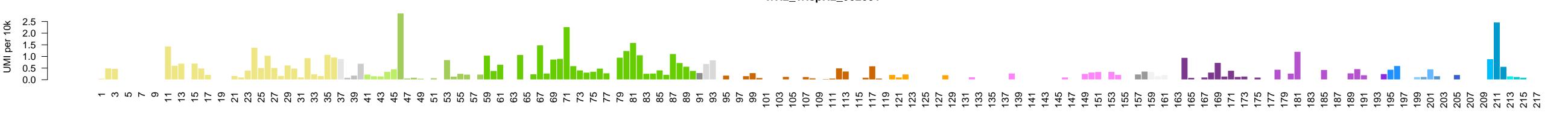


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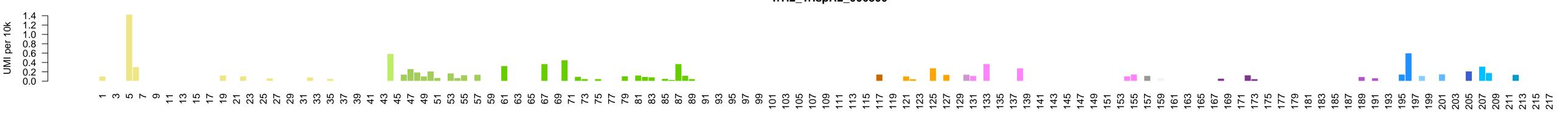




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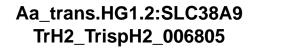


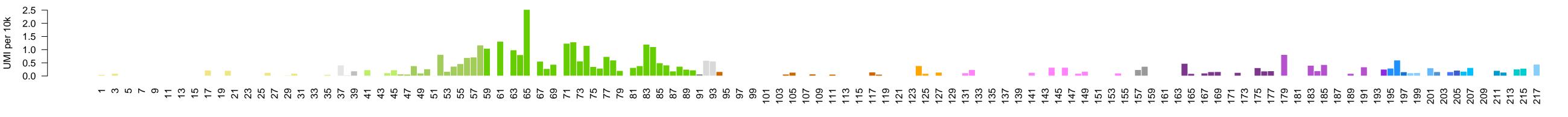
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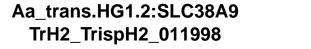


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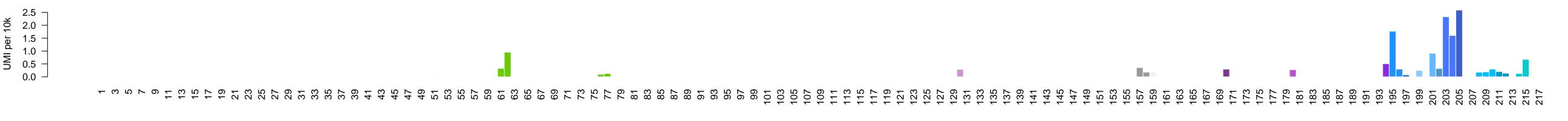




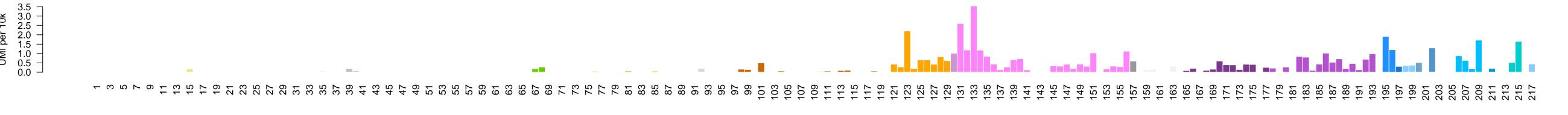




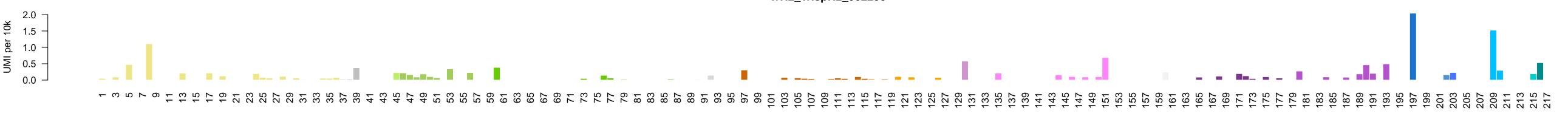
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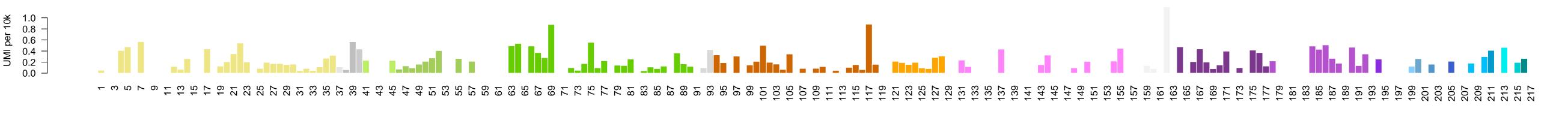
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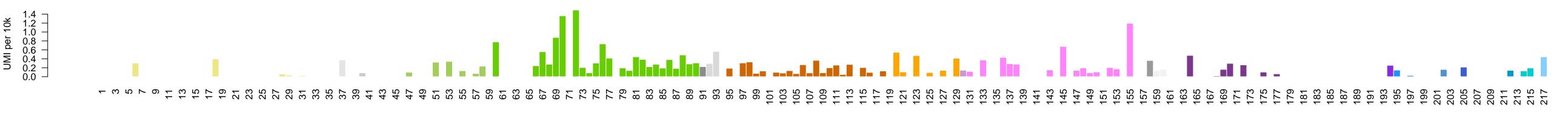
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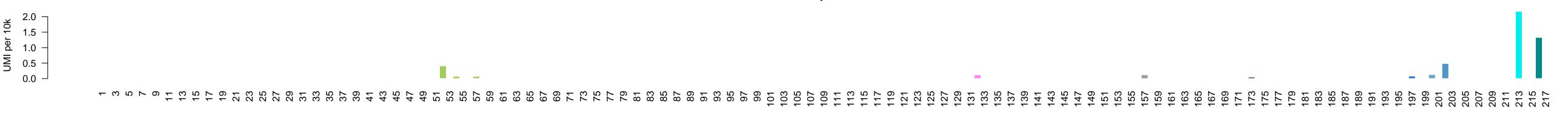
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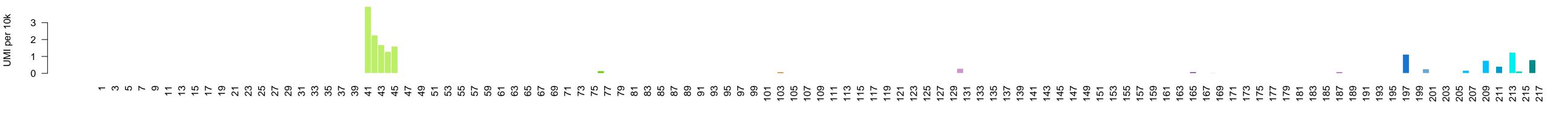
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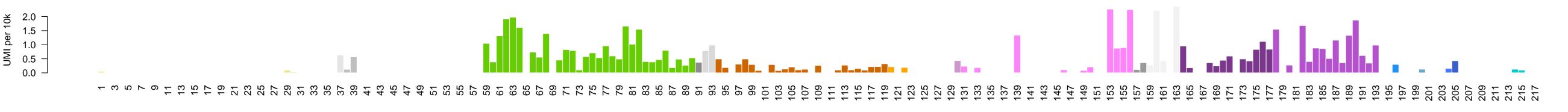
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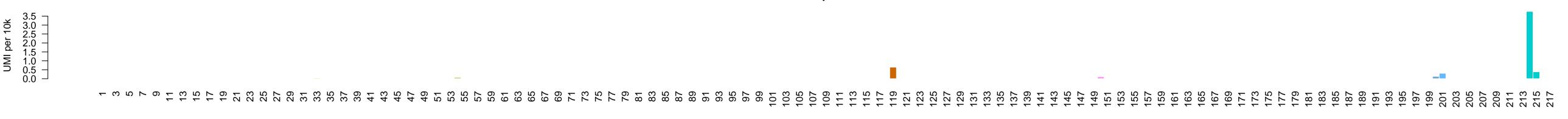
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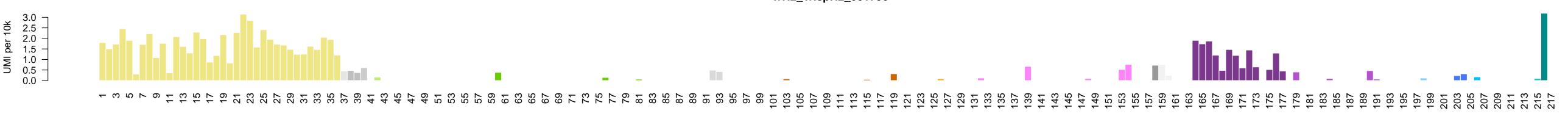
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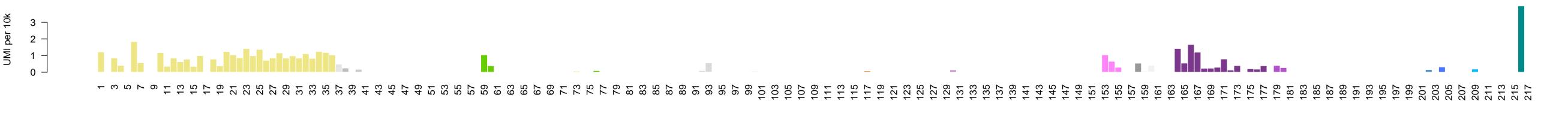
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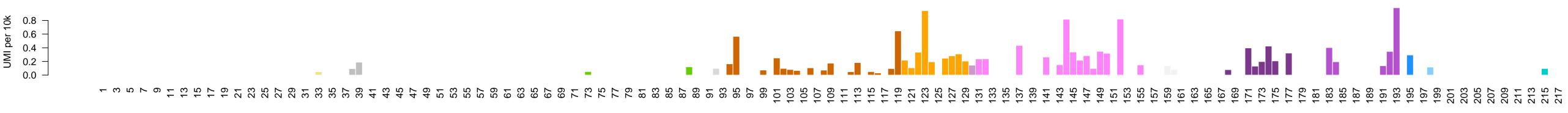
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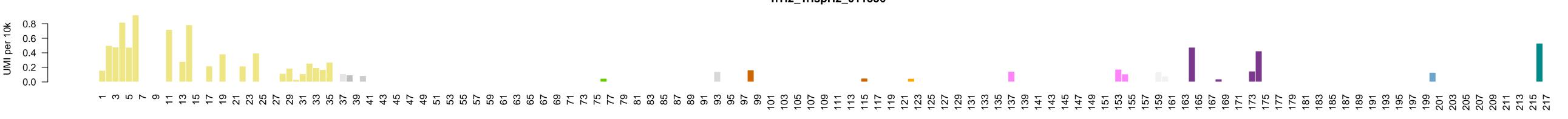
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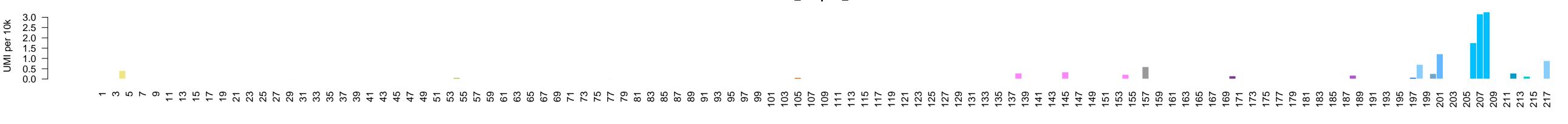
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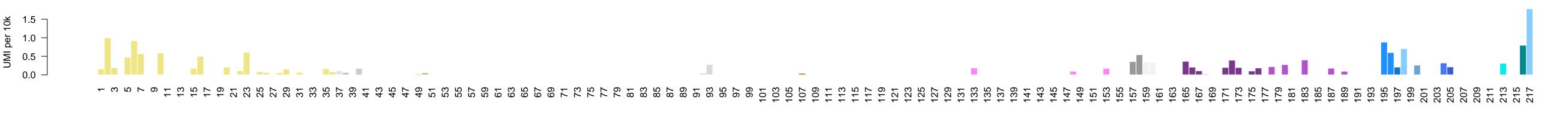
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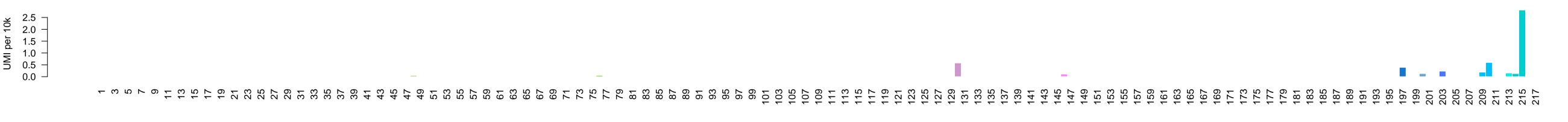
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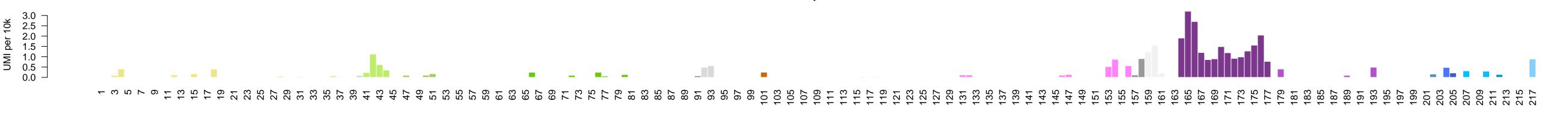
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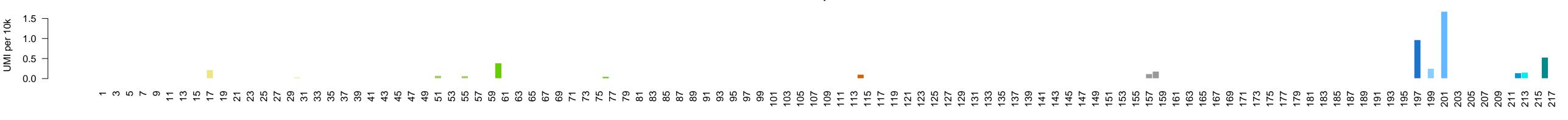
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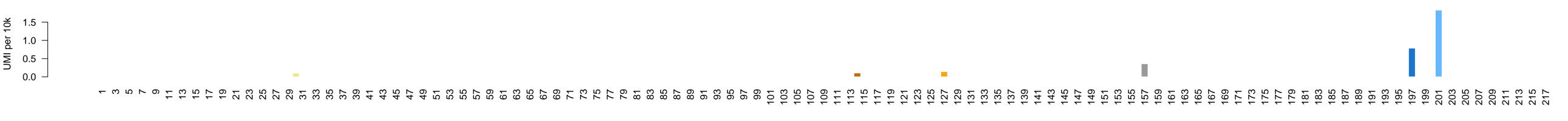
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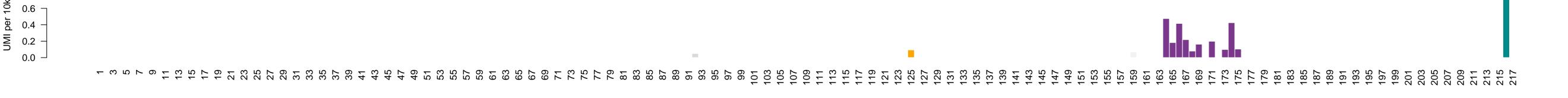
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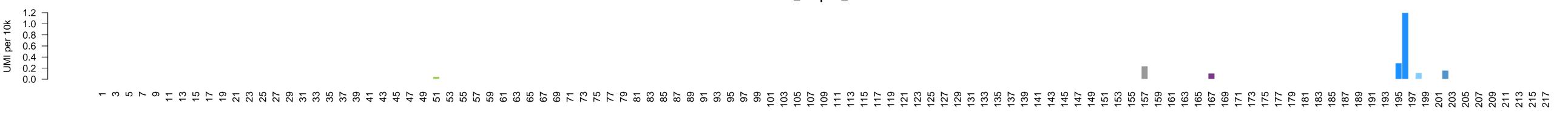
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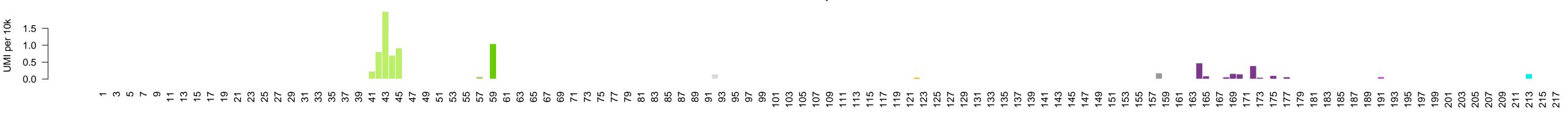
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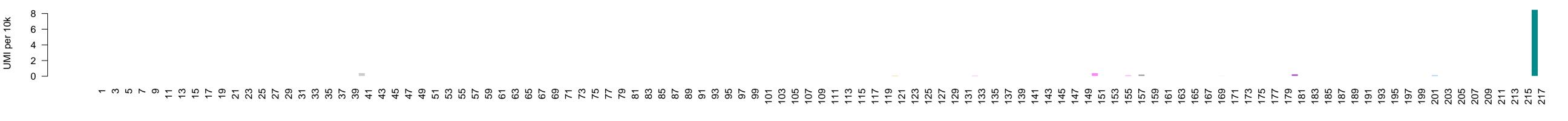
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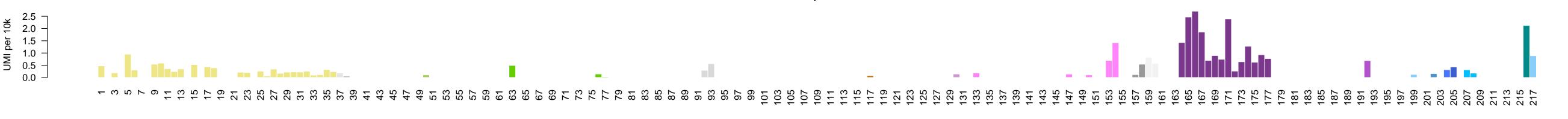
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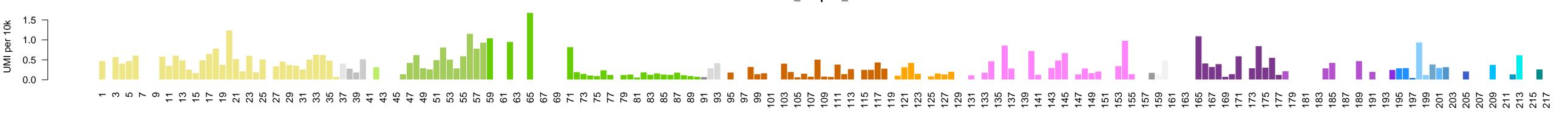
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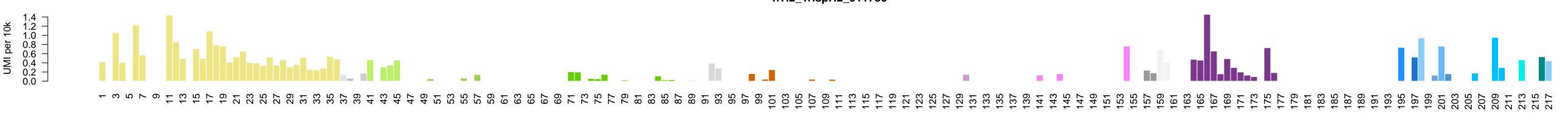
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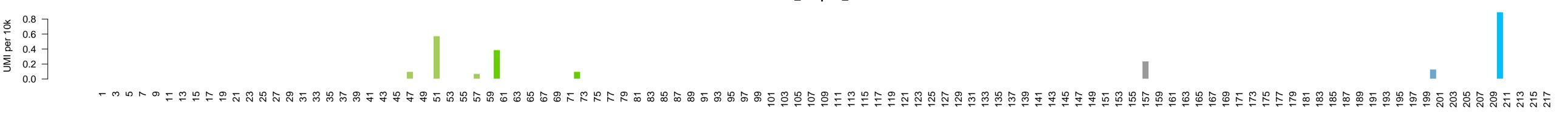
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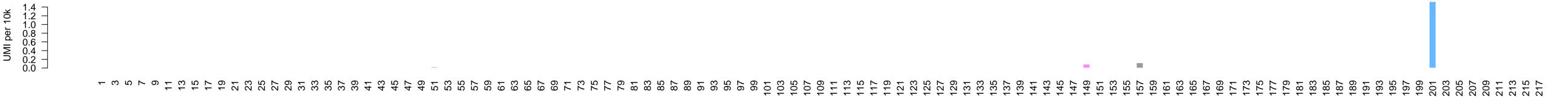
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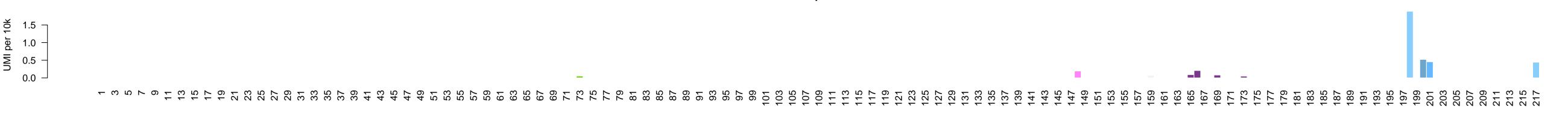
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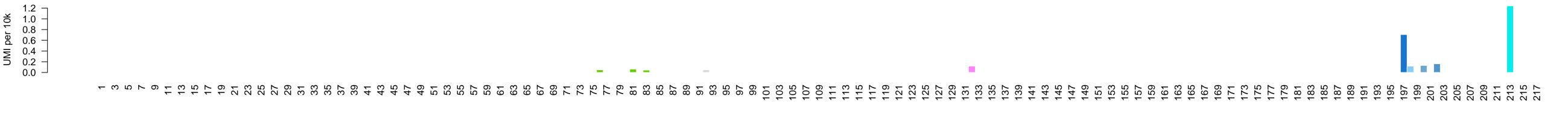
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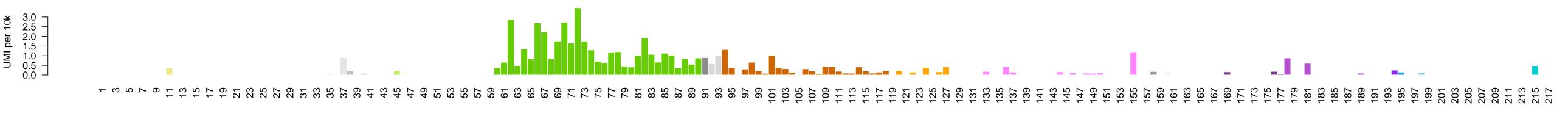
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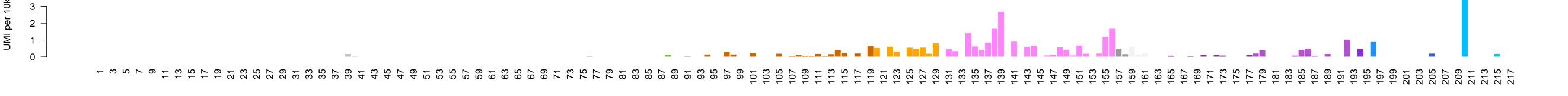
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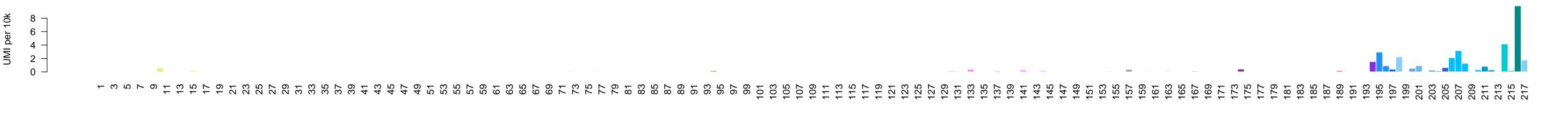


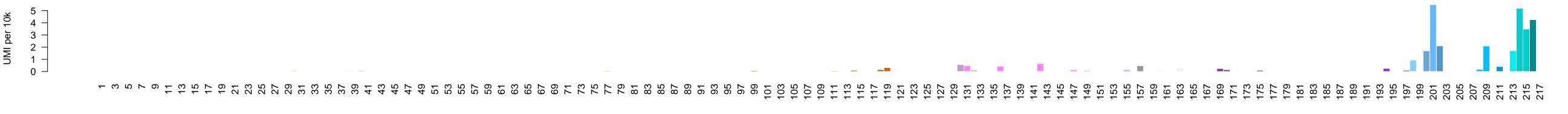
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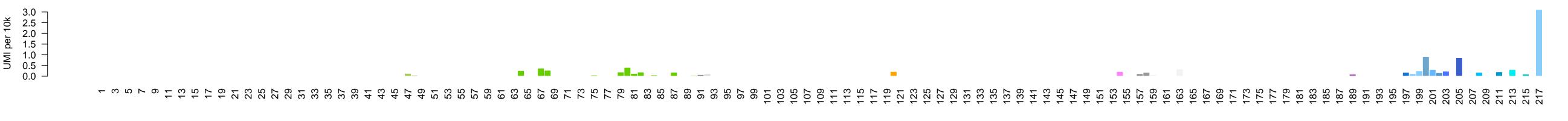


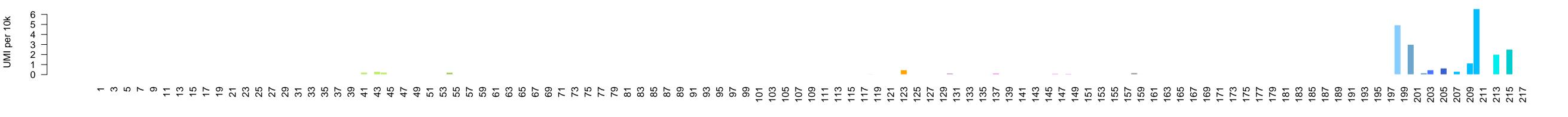
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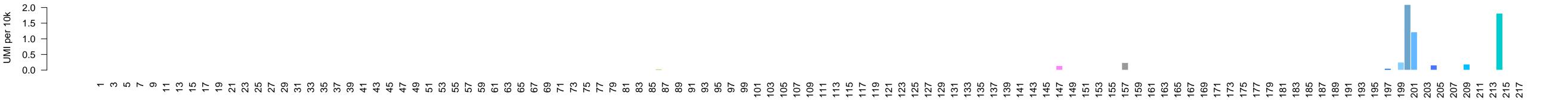


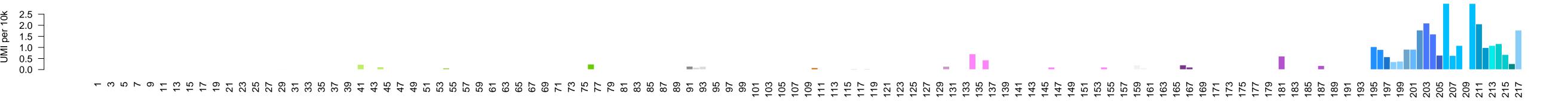


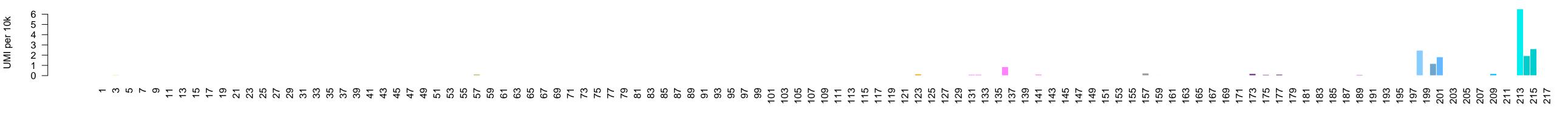


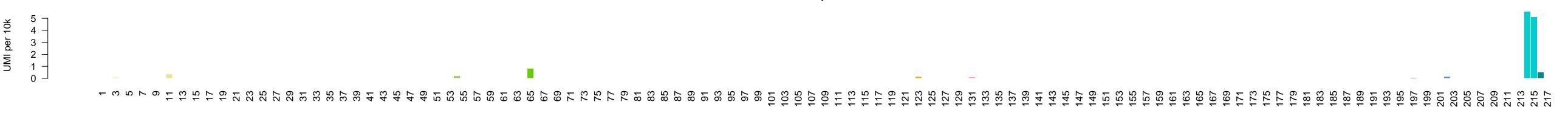


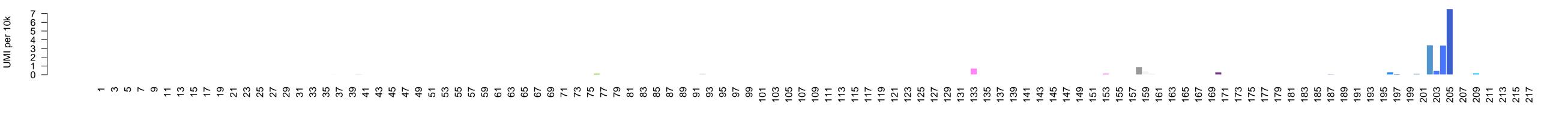


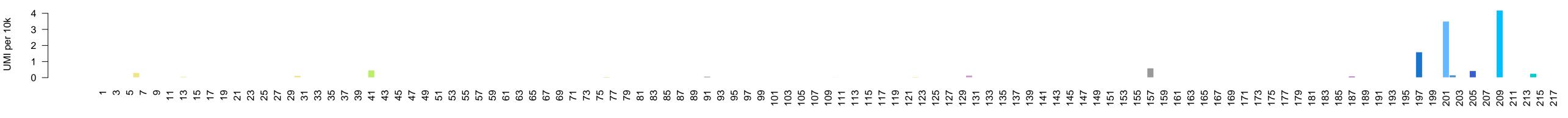


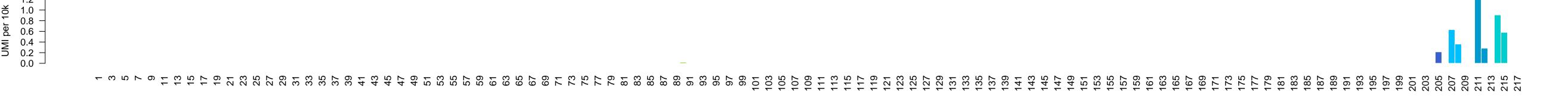




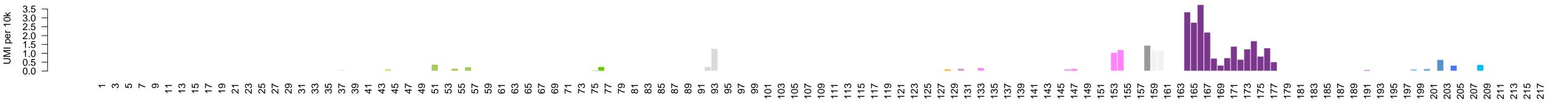


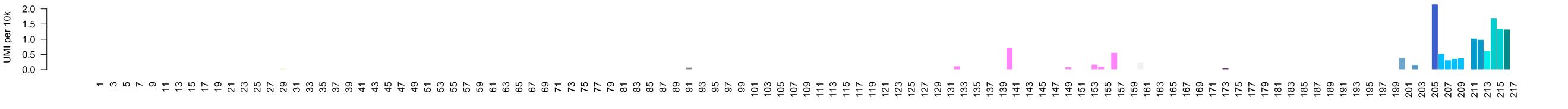


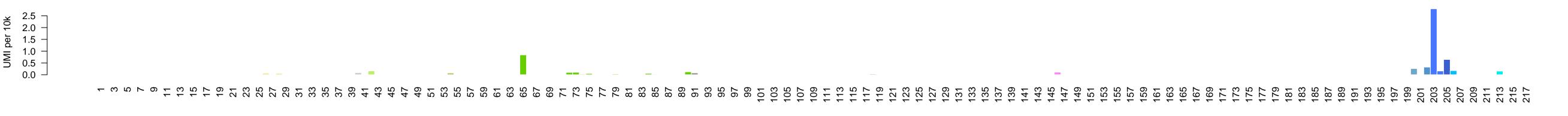


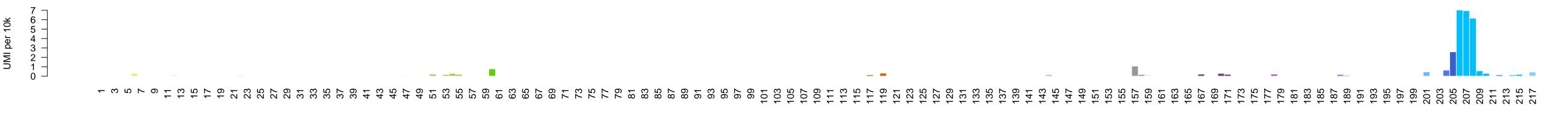


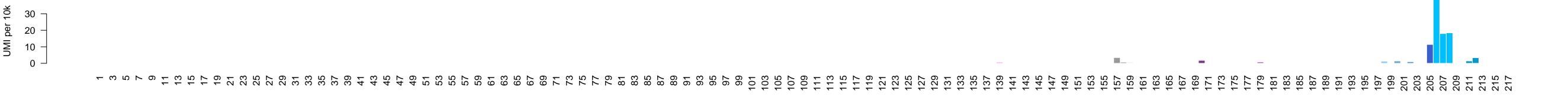
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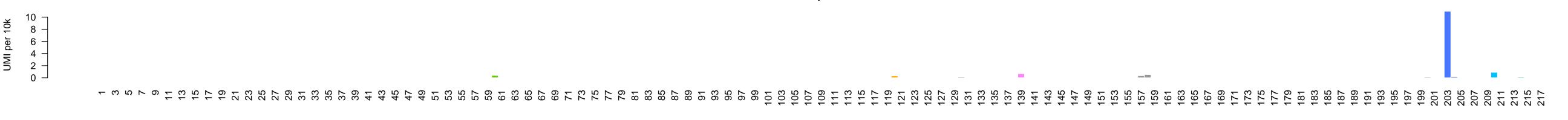


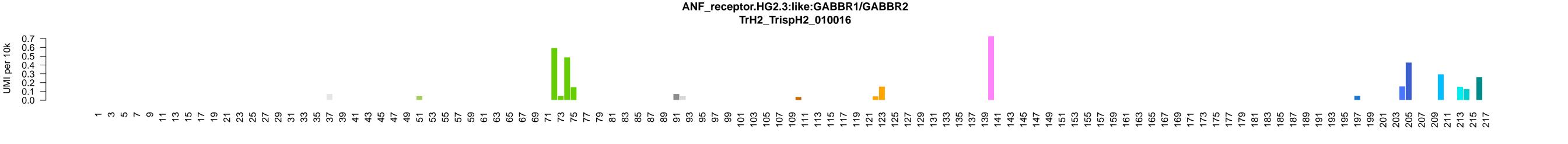


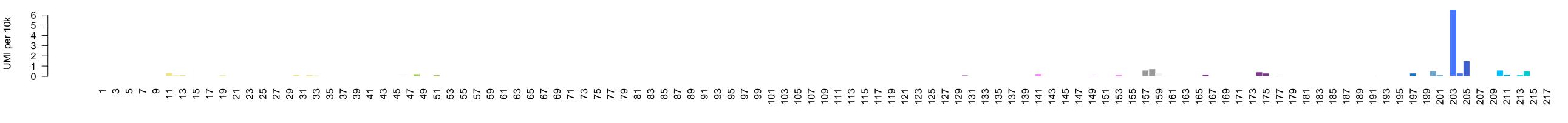


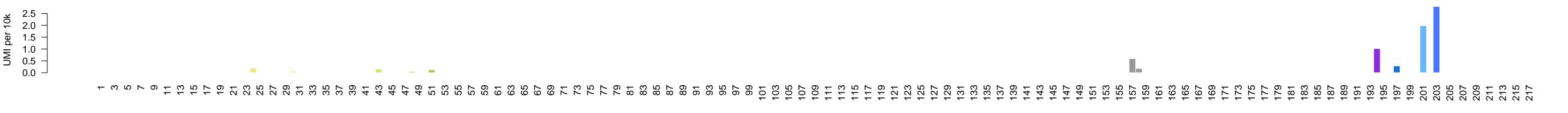


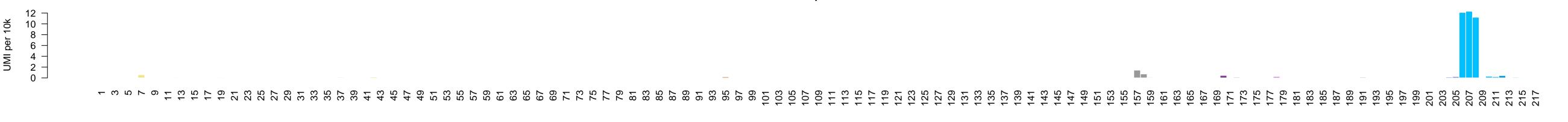


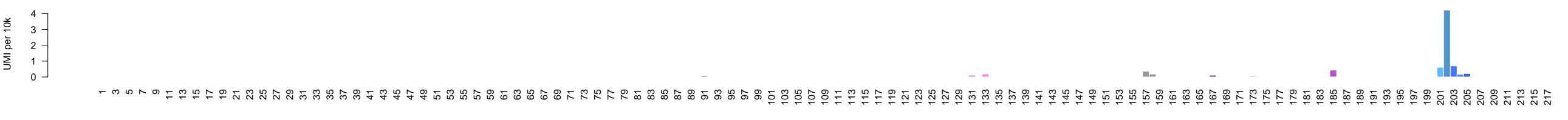


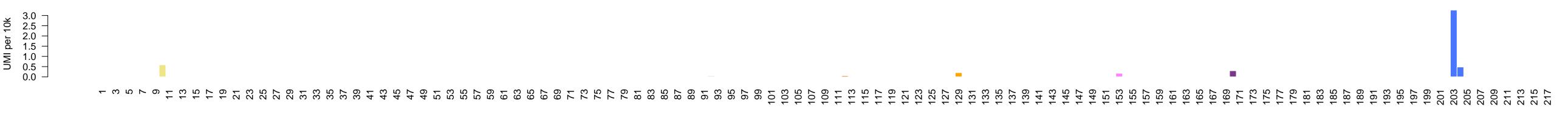


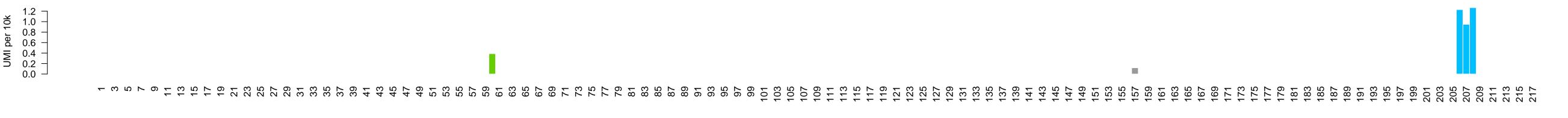




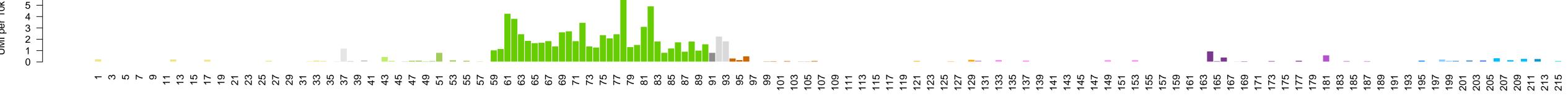


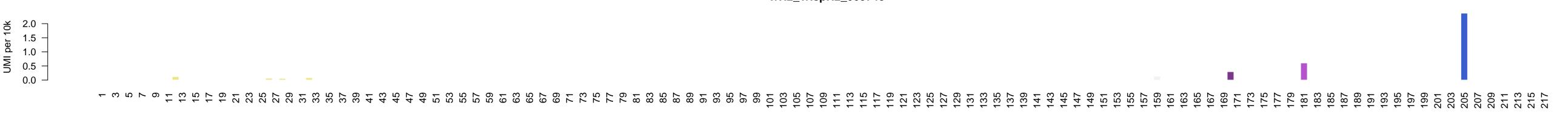


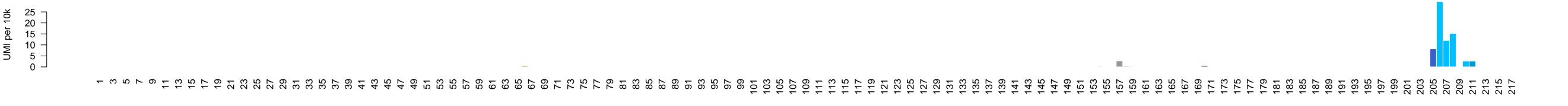


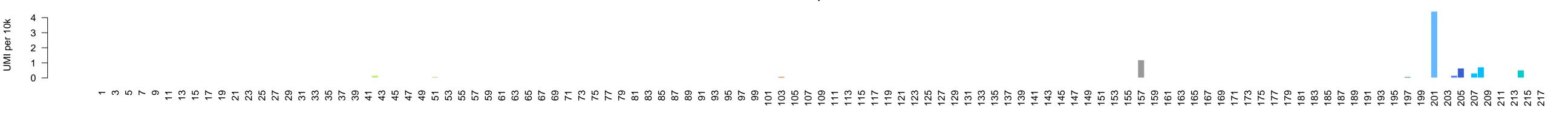


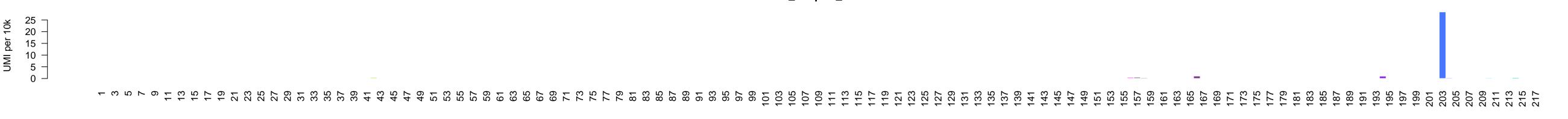
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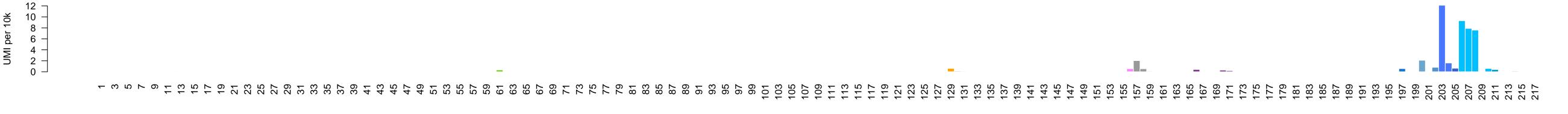


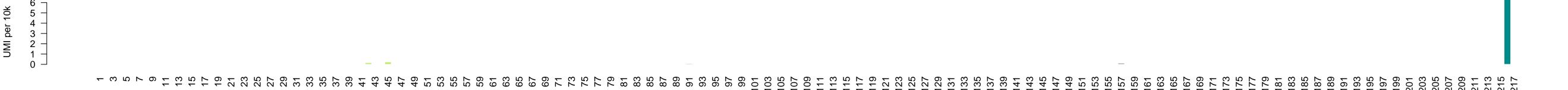


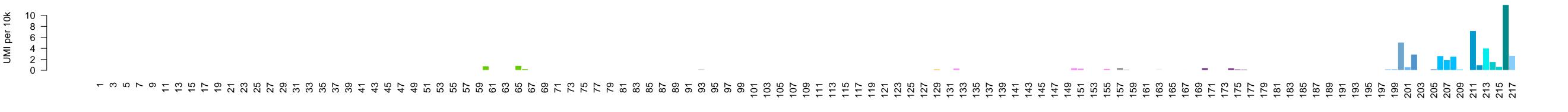


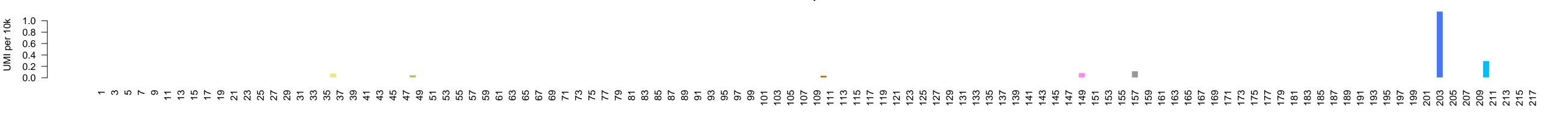


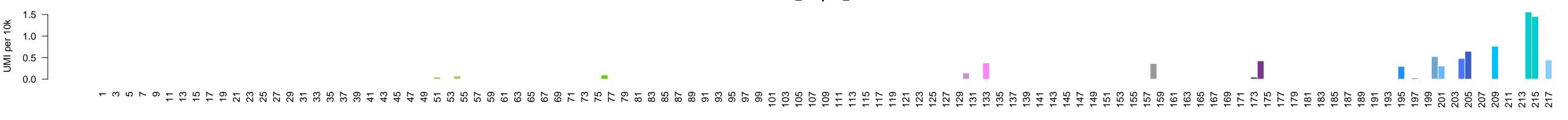


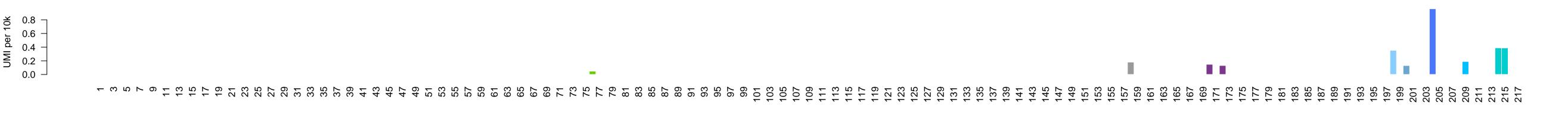


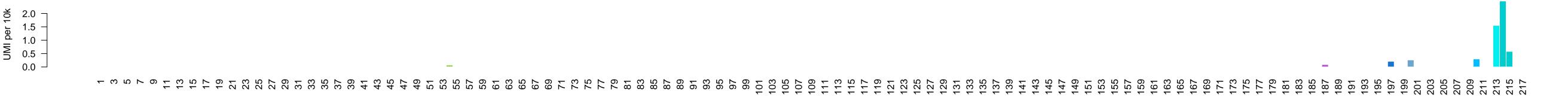


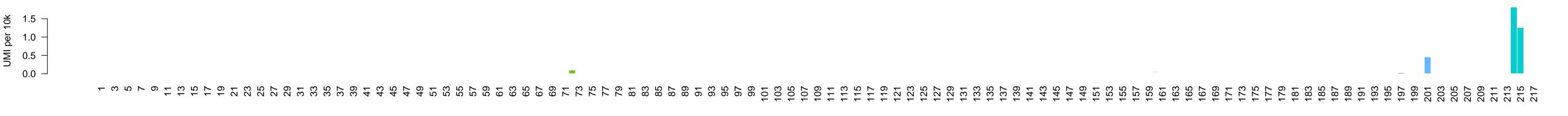


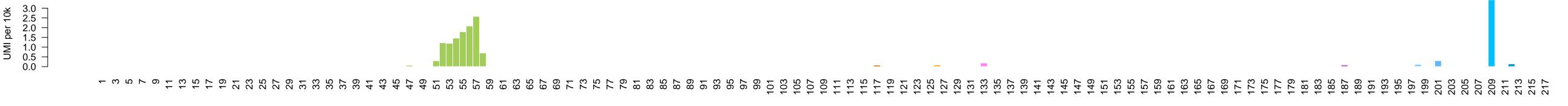


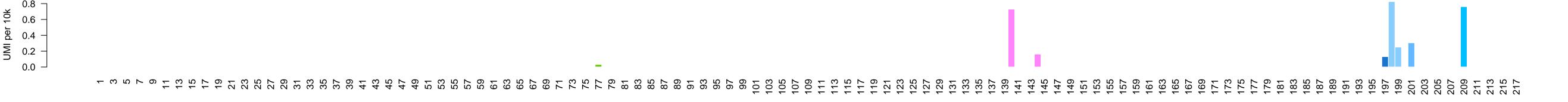


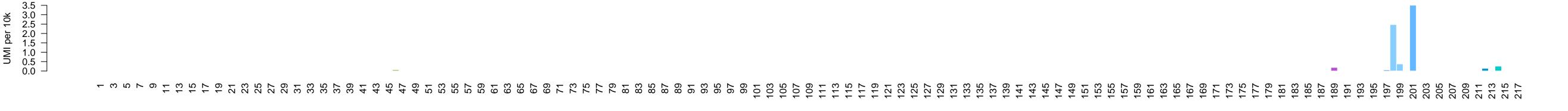


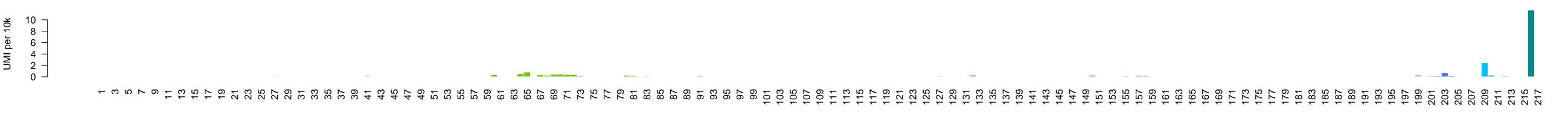


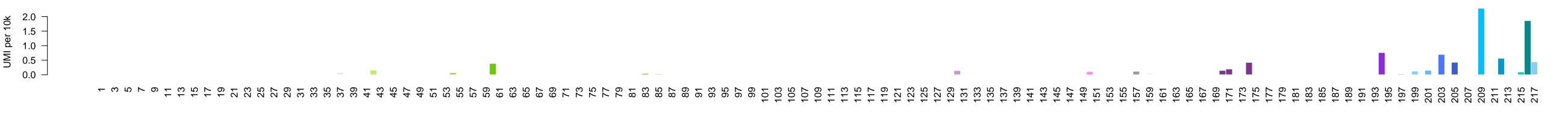


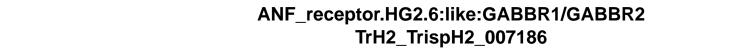






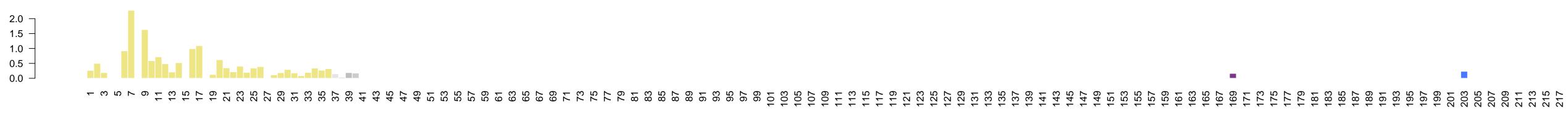


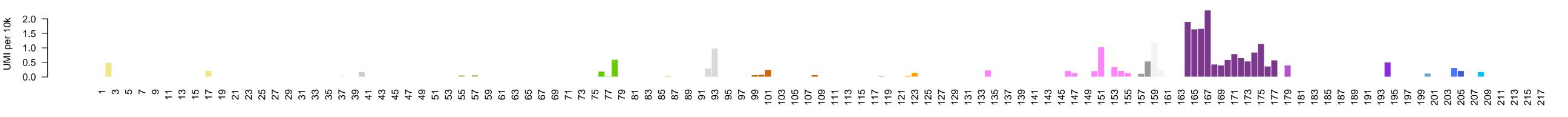


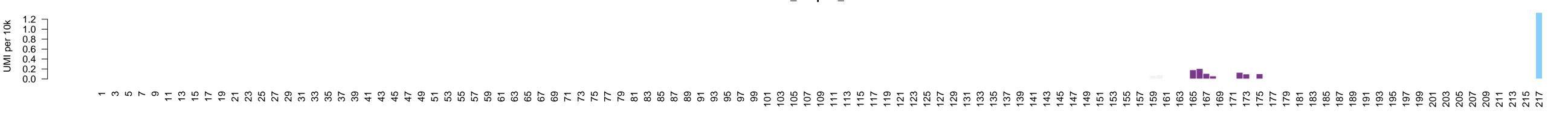




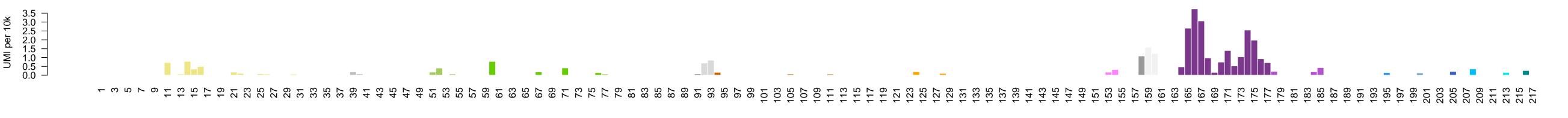
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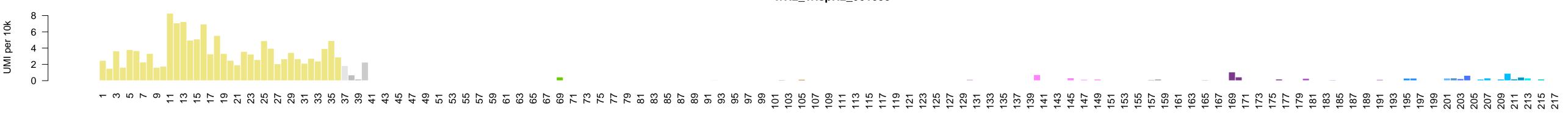


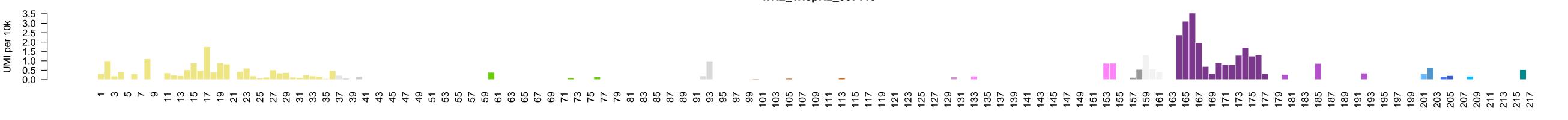




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TrH2_TrispH2_001054

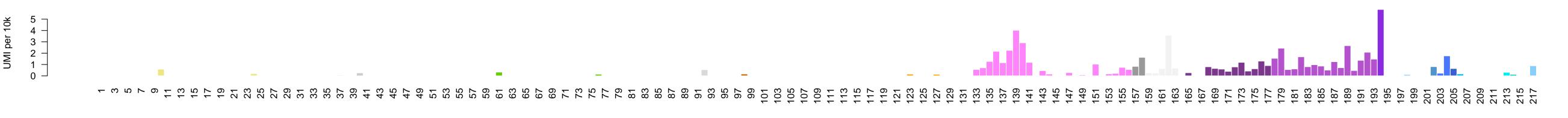








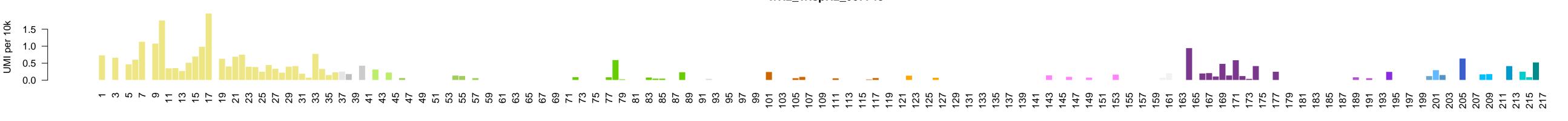
ANF_receptor.HG3.3:like:GRIN1/GRIN2A/GRIN2B/GRIN2C/GRIN2D
TrH2_TrispH2_010734



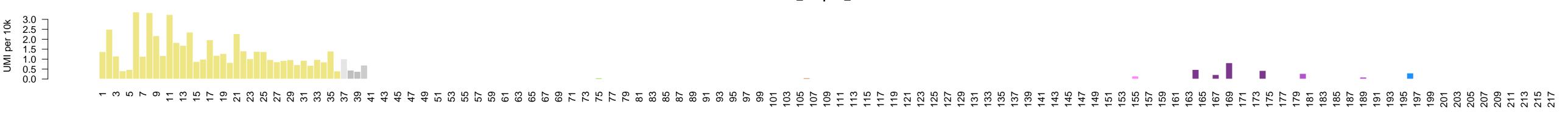
ANF_receptor.HG3.4:like:GRIA1/GRIA2/GRIA3/GRIA4/GRIK1/GRIK2/GRIK3/GRIK4/GRIK5 TrH2_TrispH2_007722



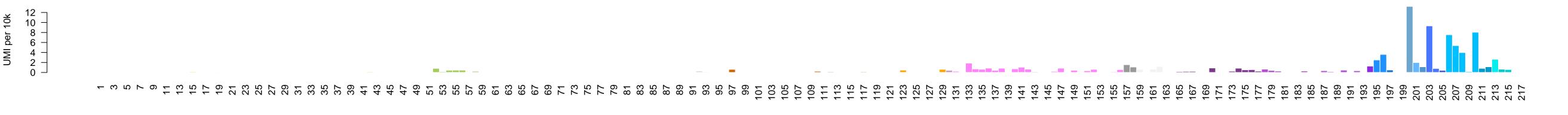
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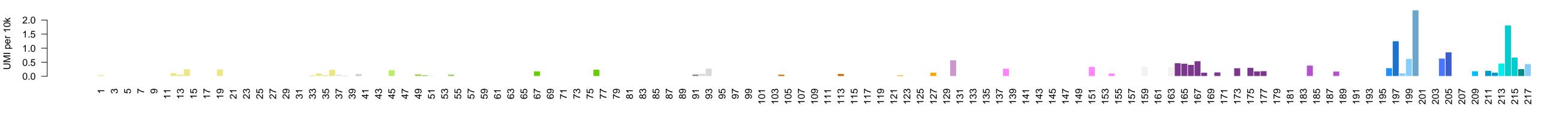
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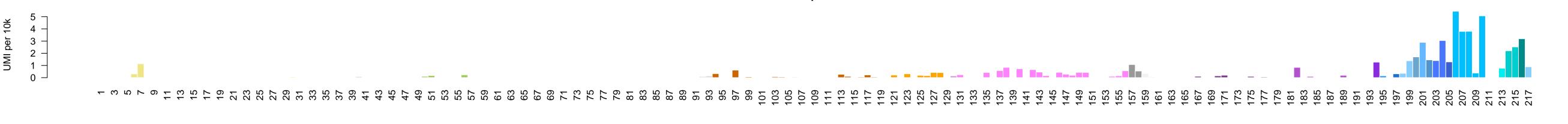
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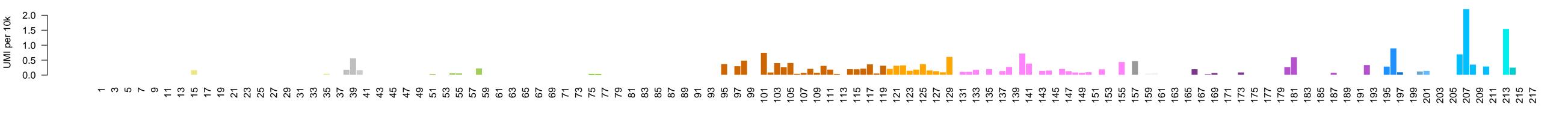
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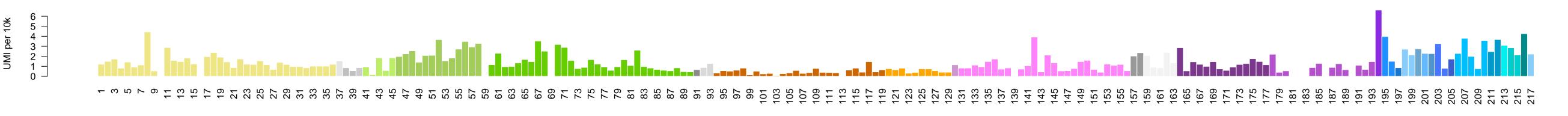
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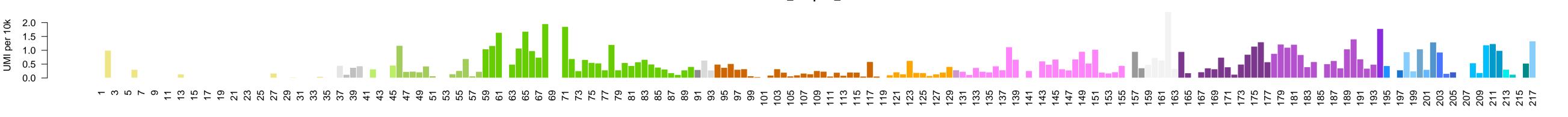
ANF_receptor.HG4.9:like:GUCY2D/GUCY2F/NPR1/NPR2/NPR3
TrH2_TrispH2_002264



CaMKII.HG1.1:CAMK2A/CAMK2B/CAMK2D/CAMK2G
TrH2_TrispH2_000247

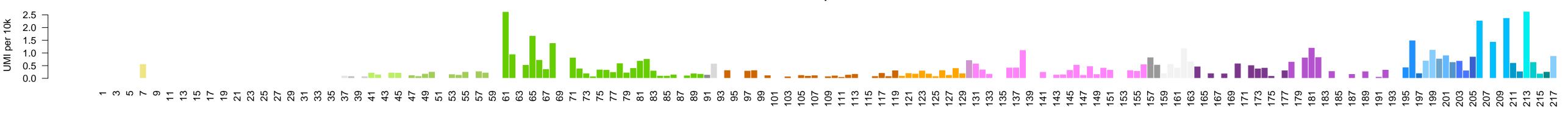


Caps.HG1.0:CADPS/CADPS2
TrH2 TrispH2 004874

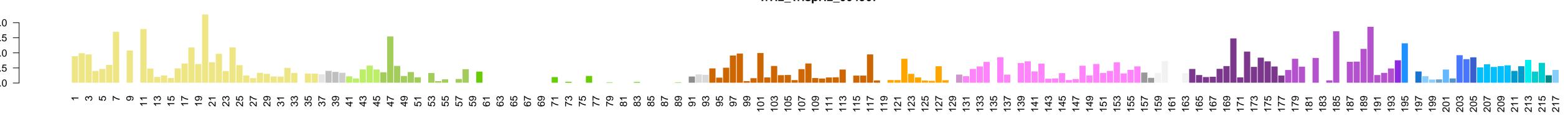


Caps.HG1.1:UNC13A/UNC13B/UNC13C;Synaptotagmin.HG1.66:UNC13A/UNC13C;Unc13.HG1.1:UNC13A/UNC13B/UNC13C

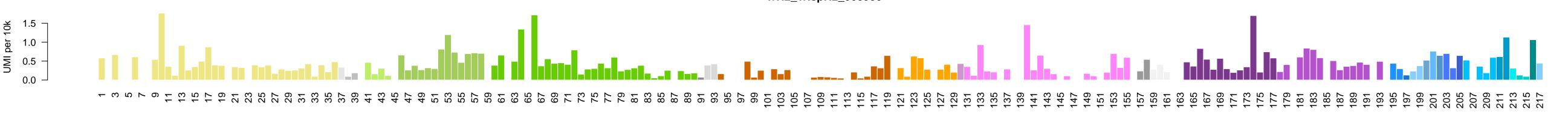
TrH2_TrispH2_007769



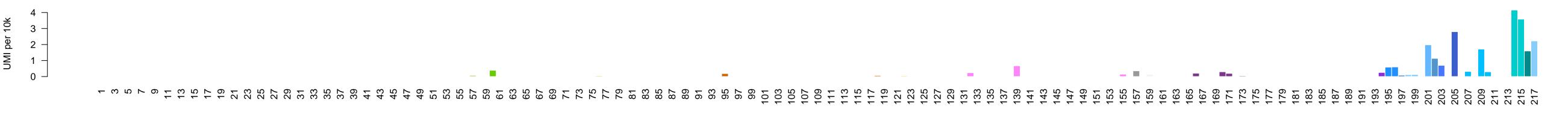
Citron.HG1.1:CDC42BPA/CDC42BPB/CDC42BPG TrH2_TrispH2_004967



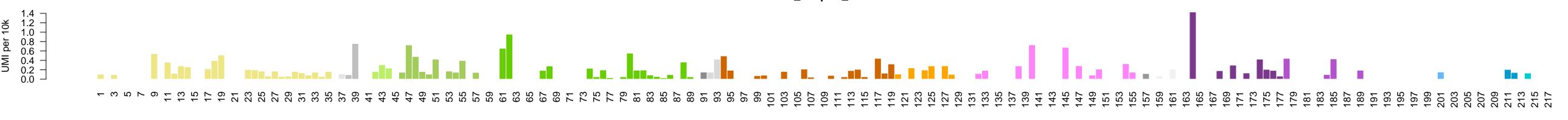
Citron.HG1.3:MAP4K4/MINK1/TNIK TrH2_TrispH2_000980



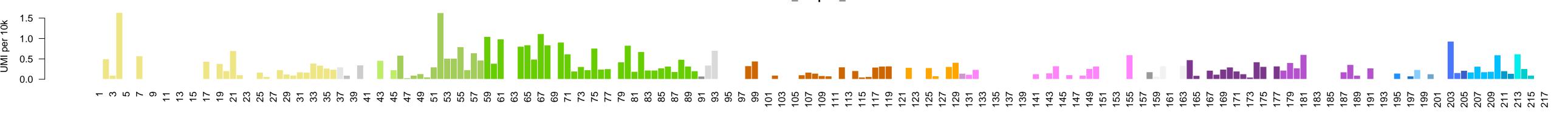
Citron.HG1.4:GARNL3 TrH2_TrispH2_009344



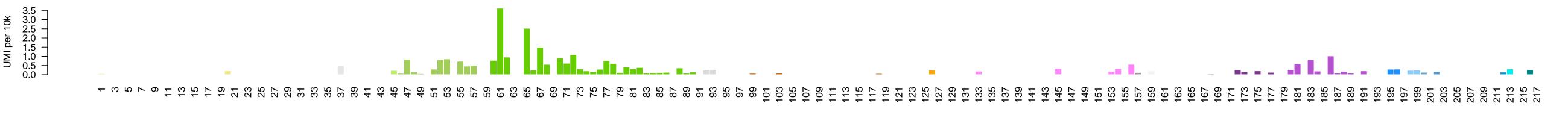
Citron.HG1.5:TGFBRAP1
TrH2_TrispH2_005198



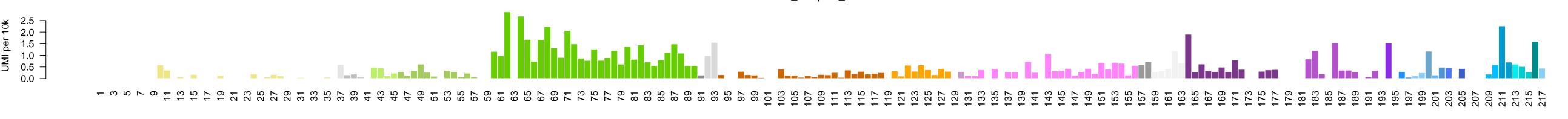
Citron.HG1.6:VPS39 TrH2_TrispH2_004960



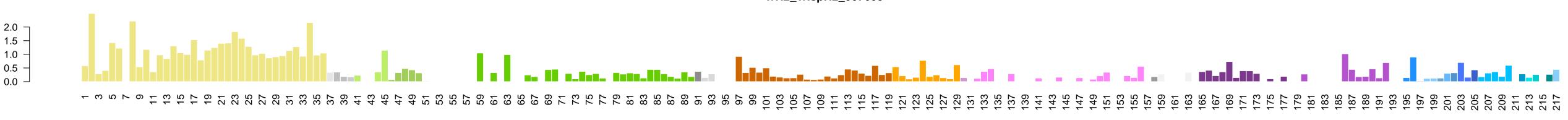
Citron.HG2.0:MAP4K1/MAP4K2/MAP4K3/MAP4K5
TrH2_TrispH2_001823

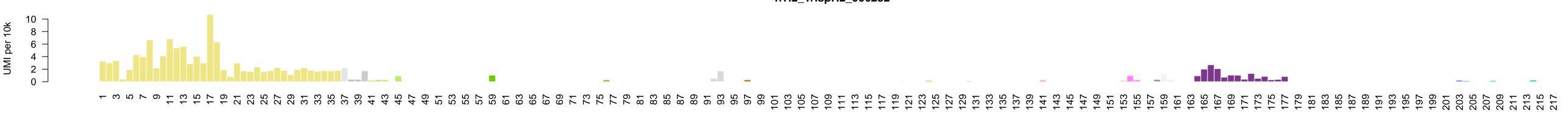


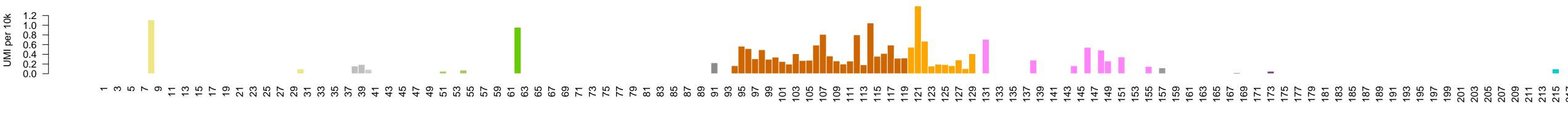
Citron.HG2.0:MAP4K1/MAP4K2/MAP4K3/MAP4K5 TrH2_TrispH2_010131



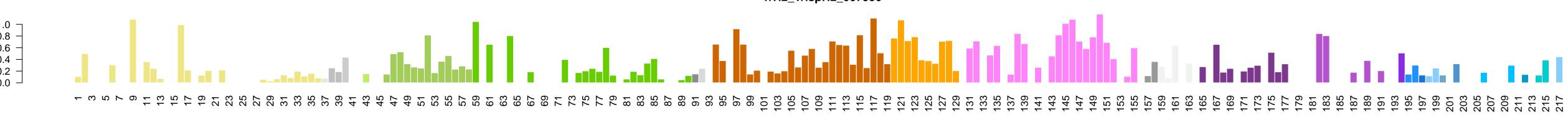
COesterase.HG1.1:like:AADAC/AADACL2/ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/NLGN1/NLGN2/NLGN3/NLGN4X/NLGN4Y/TG TrH2_TrispH2_007605

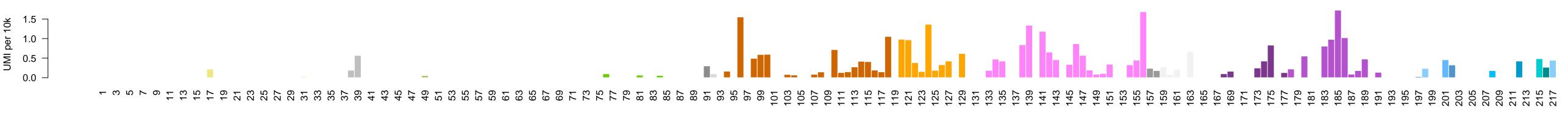


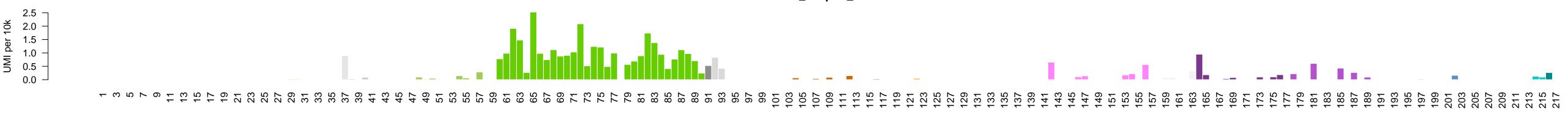


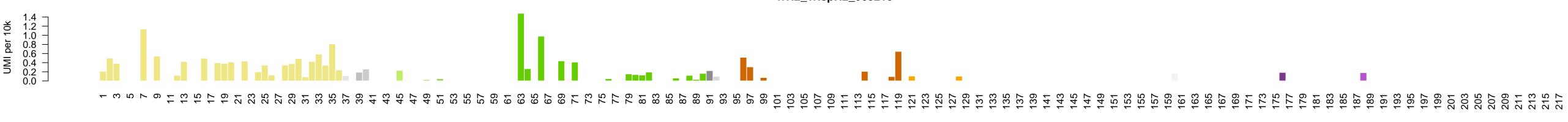


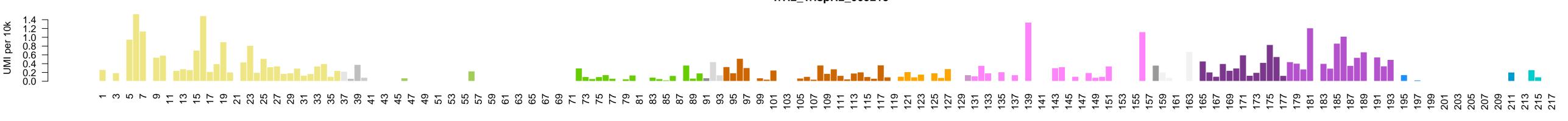
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TrH2_TrispH2_007950



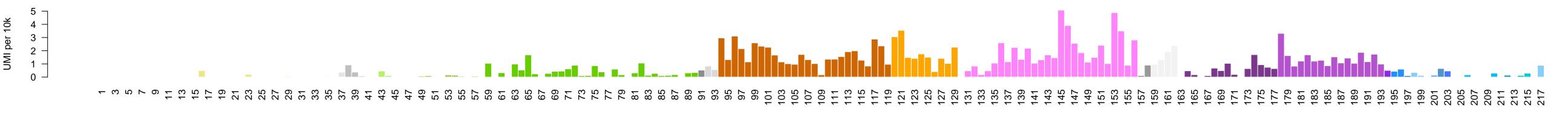


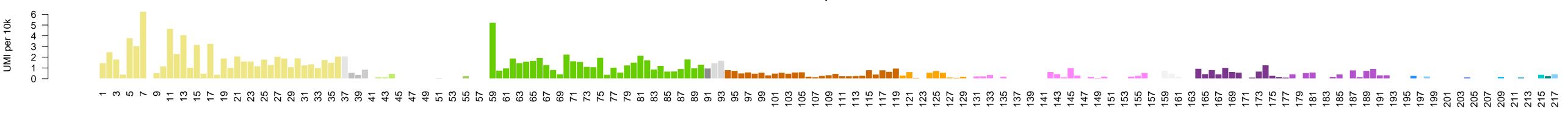


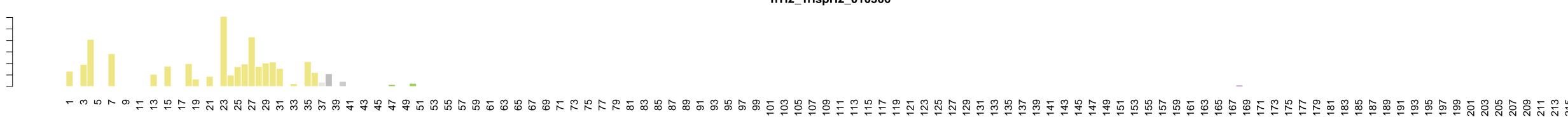


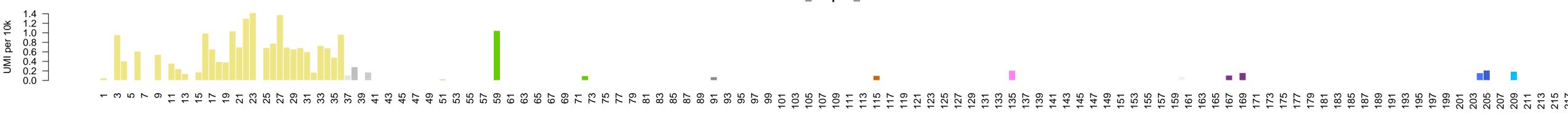


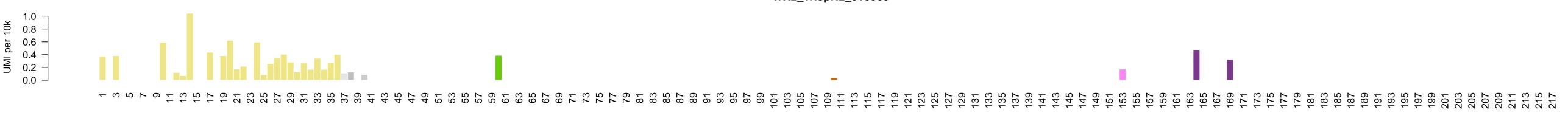
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TrH2_TrispH2_010564

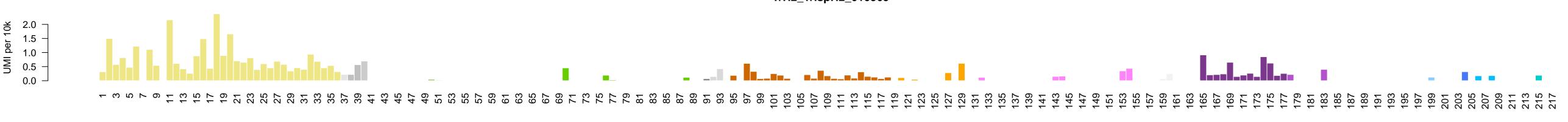


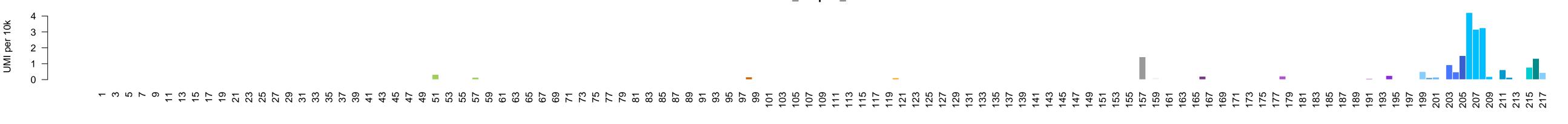


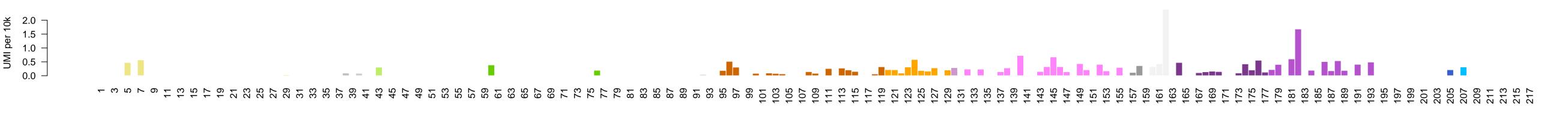


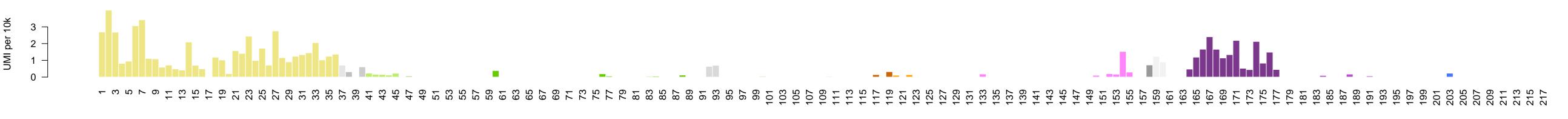


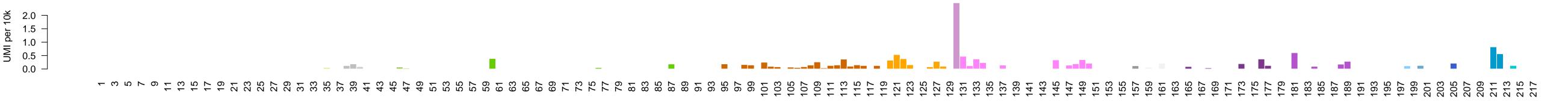


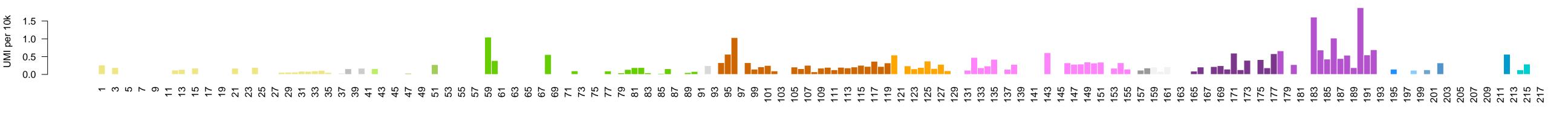


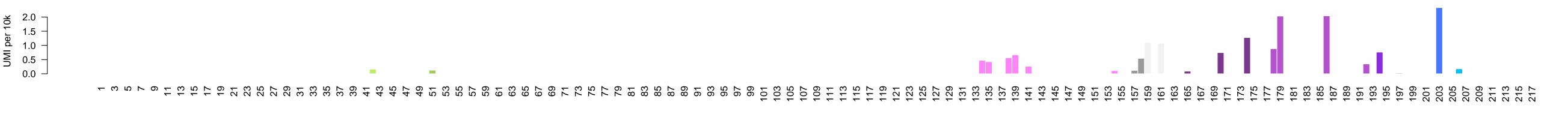


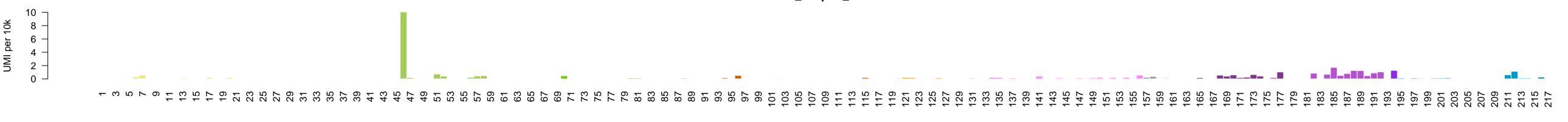


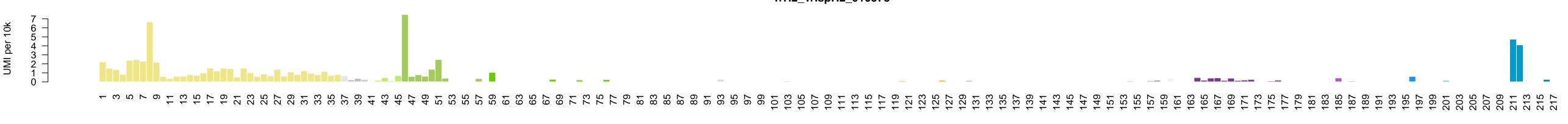


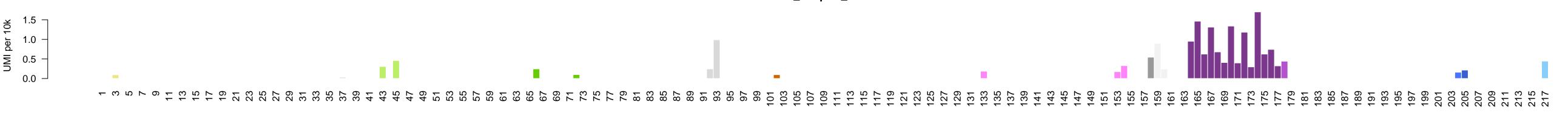




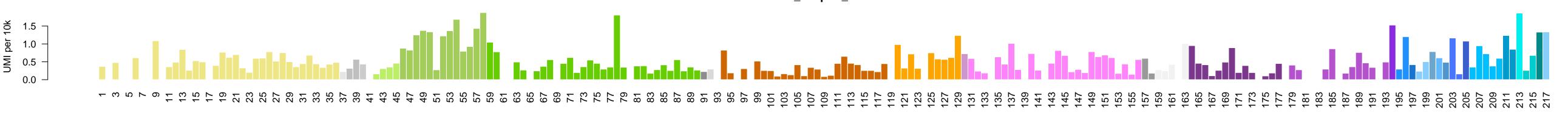




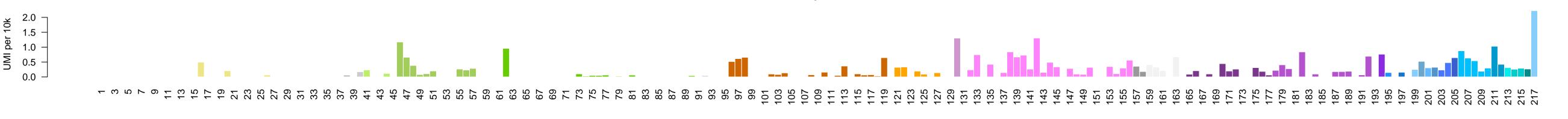




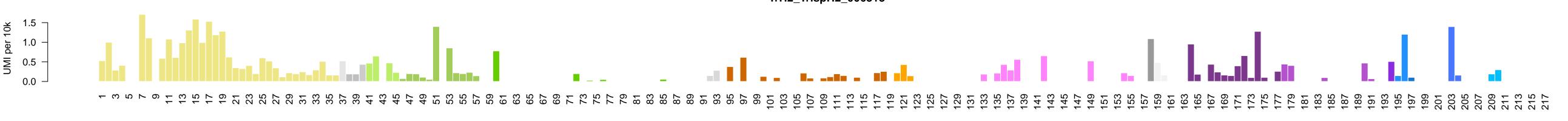
Cript.HG1.0:CRIPT
TrH2 TrispH2 000364



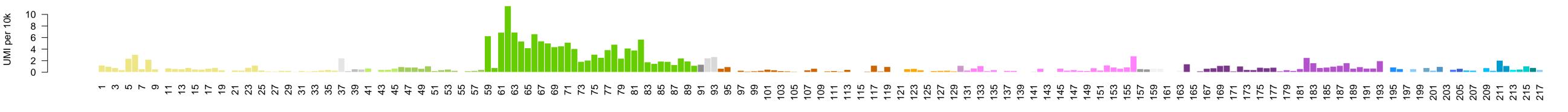
FERM_M.HG1.1:like:MYO10/PLEKHH1/PLEKHH2/PLEKHH3 TrH2_TrispH2_000438



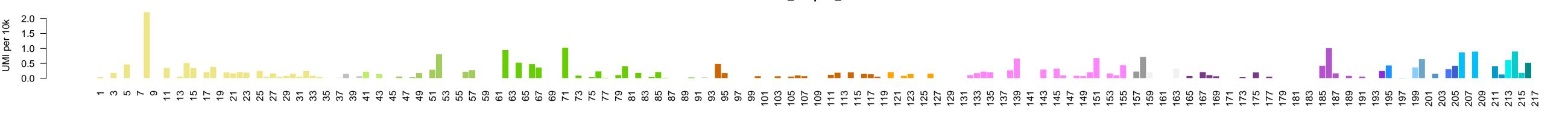
FERM_M.HG1.14:like:MYO7A/MYO7B TrH2_TrispH2_006313



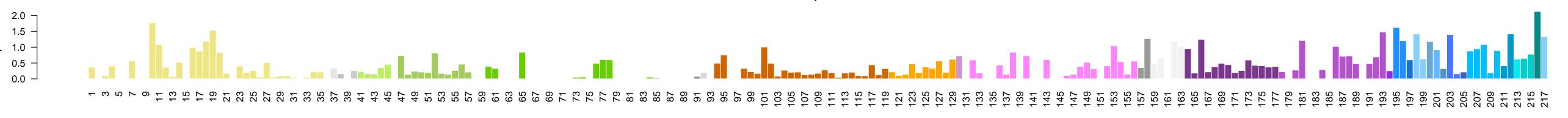
FERM_M.HG1.18:TLN1/TLN2 TrH2_TrispH2_007006



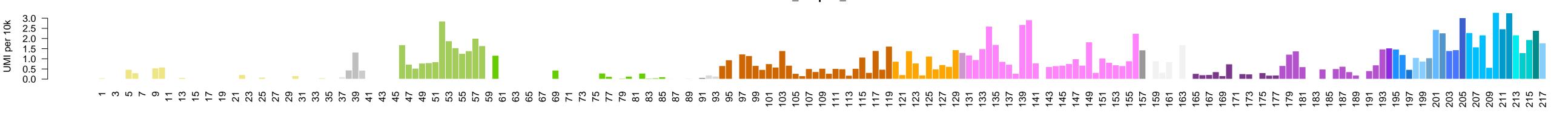
FERM_M.HG1.22:PTPN14/PTPN21;Y_phosphatase.HG1.37:PTPN14/PTPN21
TrH2 TrispH2 004976



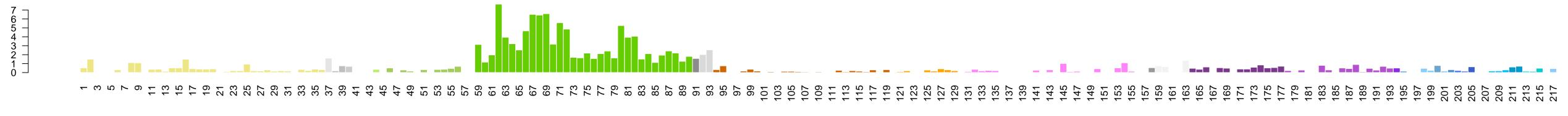
FERM_M.HG1.24:FRMD1/FRMD6/FRMPD2/PTPN13
TrH2_TrispH2_009055



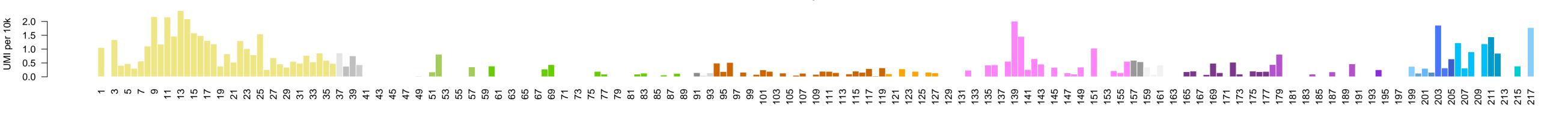
FERM_M.HG1.24:FRMD1/FRMD6/FRMPD2/PTPN13;Y_phosphatase.HG1.21:PTPN13/PTPN20 TrH2_TrispH2_000245



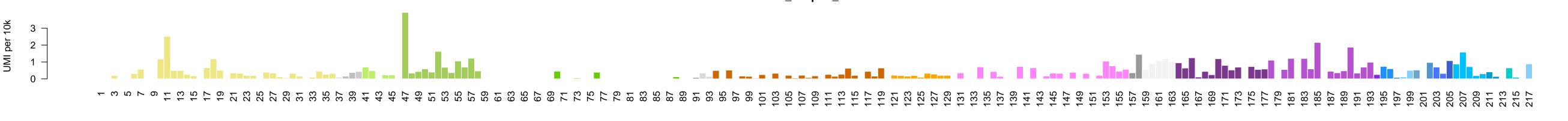
FERM_M.HG1.26:FERMT1/FERMT2/FERMT3
TrH2_TrispH2_009554



FERM_M.HG1.3:PLEKHH1/PLEKHH2 TrH2_TrispH2_010036

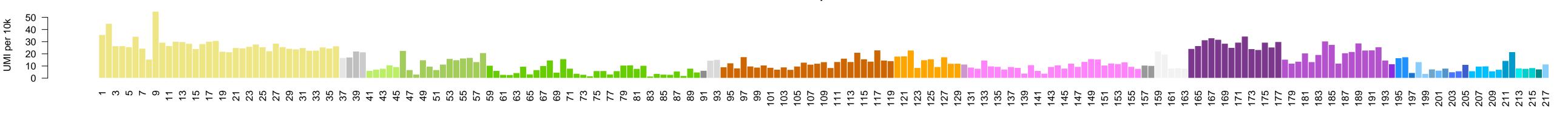


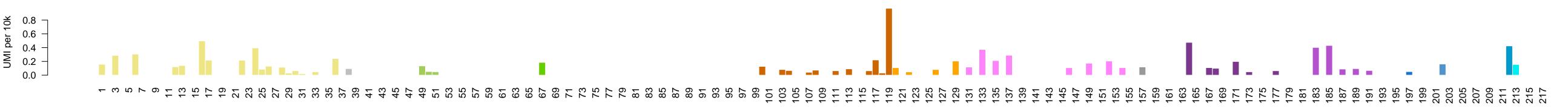
FERM_M.HG1.30:FRMD4A/FRMD4B TrH2_TrispH2_003765



FERM_M.HG1.32:EPB41/EPB41L1/EPB41L2/EPB41L3/EPB41L4A/EPB41L4B/EPB41L5/FARP1/FARP2/FRMD3/FRMD5/FRMD7/PTPN3/PTPN4

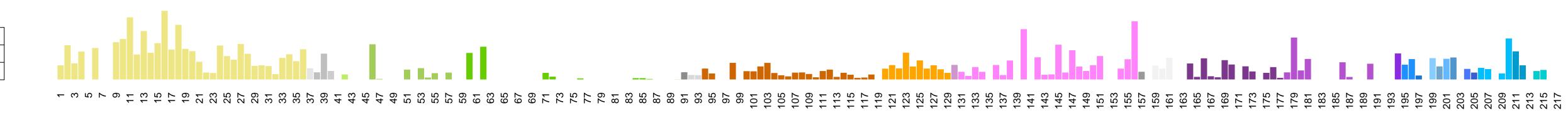
TrH2_TrispH2_001493

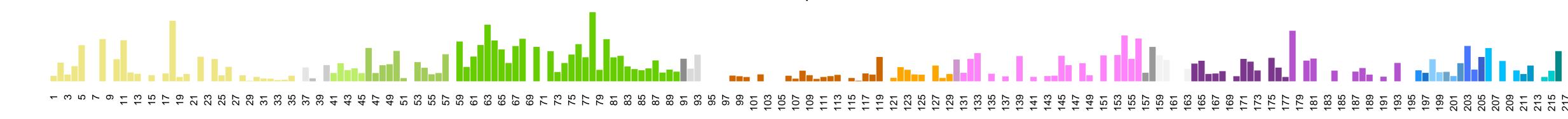


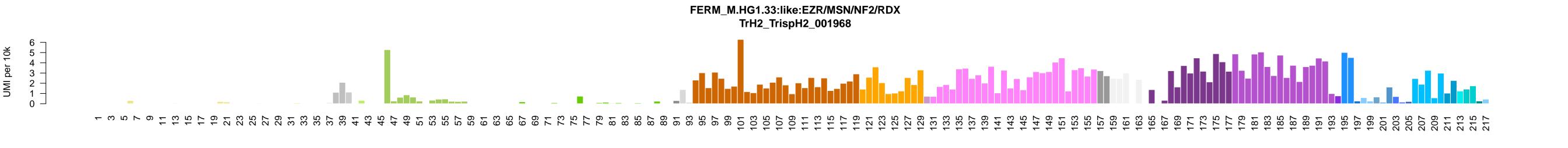


FERM_M.HG1.32:EPB41/EPB41L1/EPB41L2/EPB41L3/EPB41L4A/EPB41L4B/EPB41L5/FARP1/FARP2/FRMD3/FRMD5/FRMD7/PTPN3/PTPN4

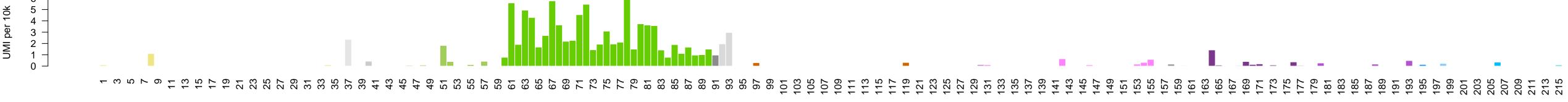
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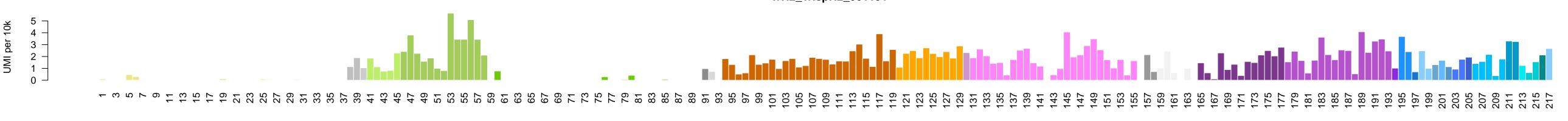




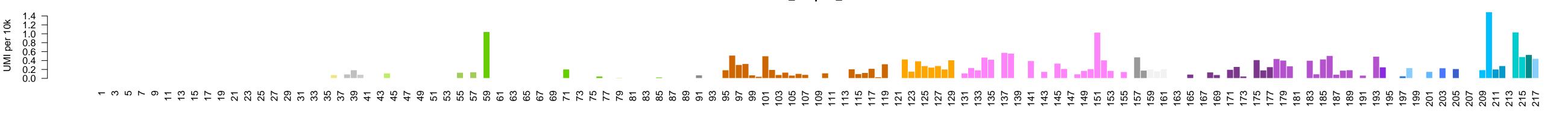
FERM_M.HG1.4:MYO10/PLEKHH3 TrH2_TrispH2_004795



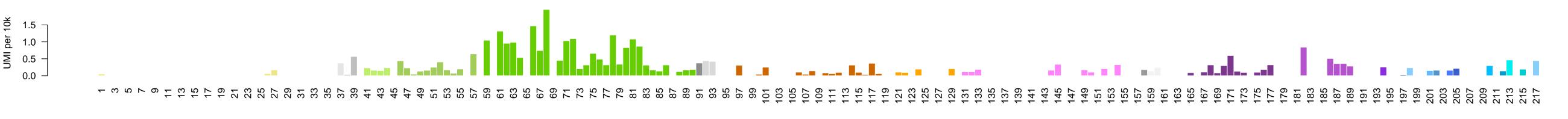
FERM_M.HG1.40:EZR/MSN/RDX TrH2_TrispH2_001191



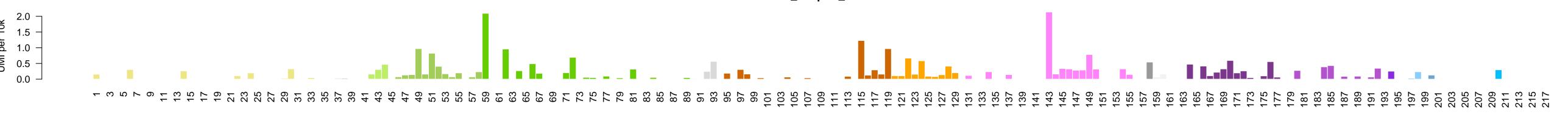
FERM_M.HG1.7:like:FERMT1/FERMT2/FERMT3/TLN1/TLN2 TrH2_TrispH2_006459



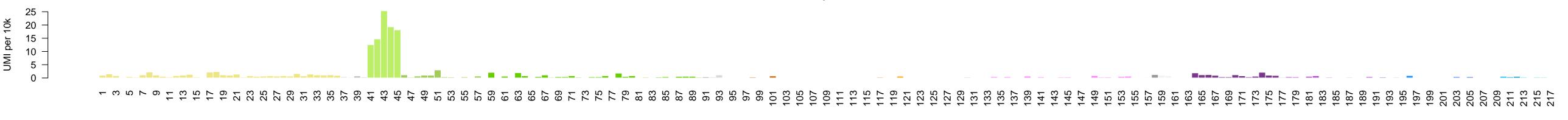
FERM_M.HG2.1:FRMPD1/FRMPD3/FRMPD4 TrH2_TrispH2_004472



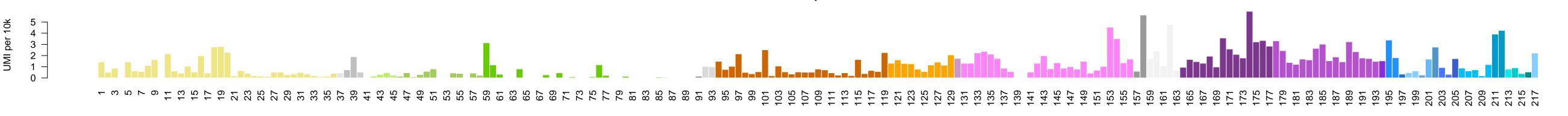
GKAP.HG1.0:DLGAP5 TrH2 TrispH2 001466



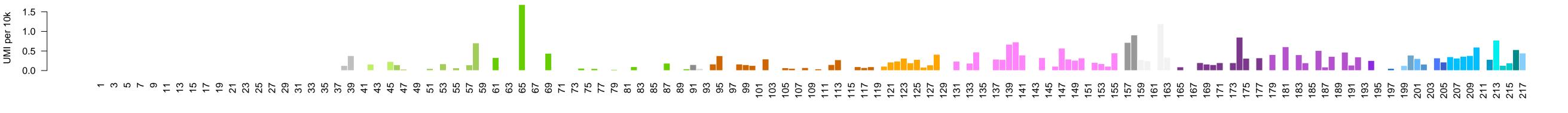
GKAP.HG1.1:DLGAP1/DLGAP2/DLGAP3/DLGAP4 TrH2_TrispH2_009500



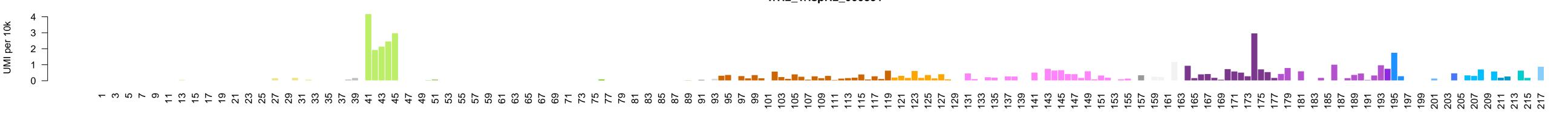
Guanylate_kin.HG1.0:MAGI1/MAGI2/MAGI3 TrH2_TrispH2_001762



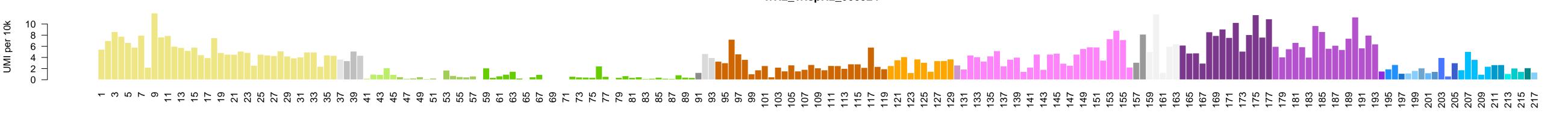
Guanylate_kin.HG1.10:like:GUK1/LRGUK
TrH2 TrispH2 005156



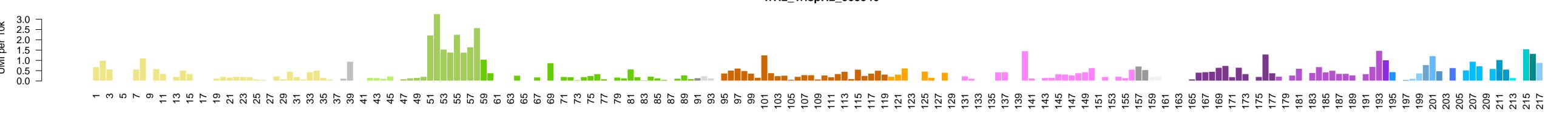
Guanylate_kin.HG1.11:AL358113.1/DLG5/TJP1/TJP2/TJP3 TrH2_TrispH2_000851

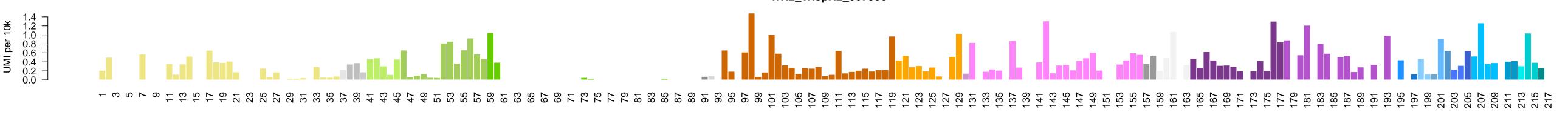


Guanylate_kin.HG1.14:DLG1/DLG2/DLG3/DLG4 TrH2_TrispH2_000924

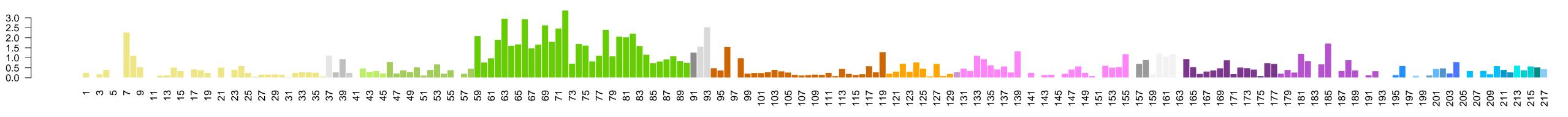


Guanylate_kin.HG1.16:GUK1 TrH2 TrispH2 003940

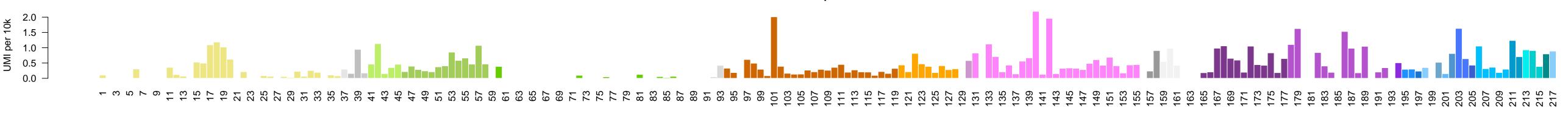




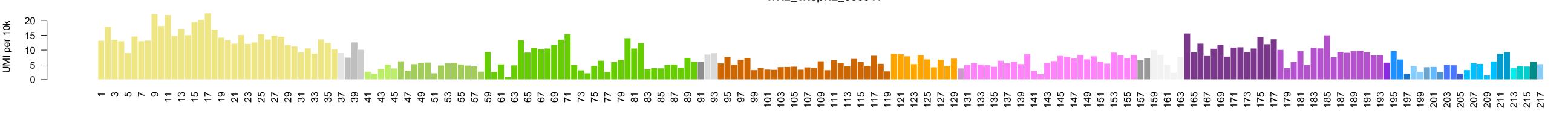
Guanylate_kin.HG1.20:CASK/MPP1;PDZ.HG2.8:CASK TrH2_TrispH2_001744



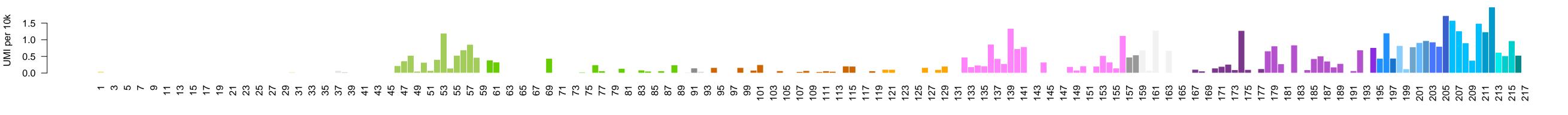
Guanylate_kin.HG1.3:MPP3/MPP4/MPP7;PDZ.HG2.10:like:MPP5 TrH2_TrispH2_006045



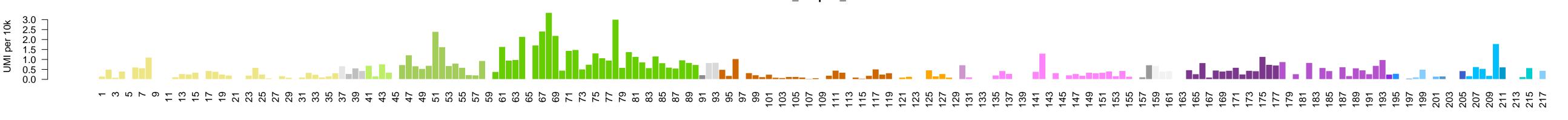
Guanylate_kin.HG1.9:like:CASK/MPP1/MPP2/MPP6;PDZ.HG2.5:MPP2/MPP6
TrH2_TrispH2_006041



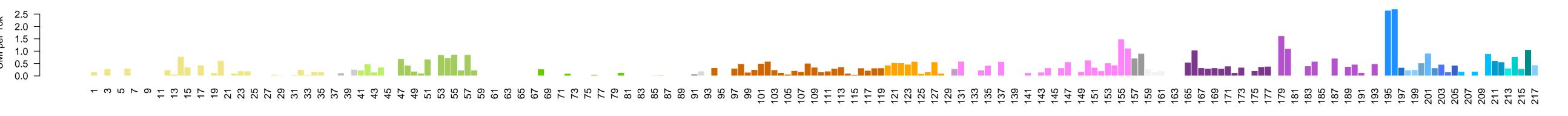
Guanylate_kin.HG2.0:CACNB1/CACNB2/CACNB3/CACNB4 TrH2_TrispH2_007753



Homer.HG1.1:ENAH/EVL/SPRED1/SPRED2/SPRED3/VASP
TrH2_TrispH2_002471

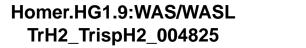


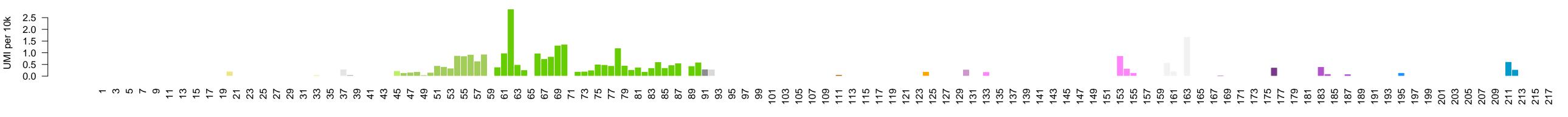
Homer.HG1.2:HOMER1/HOMER2/HOMER3 TrH2_TrispH2_001315



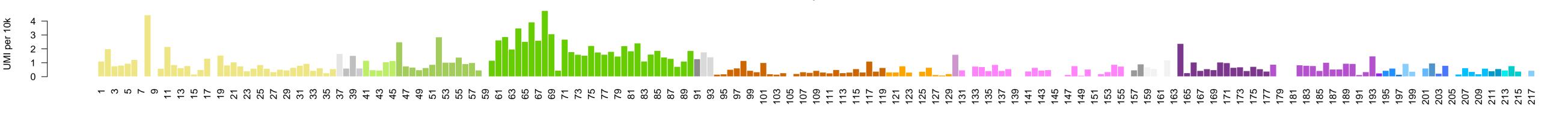
Homer.HG1.6:like:WAS/WASL TrH2_TrispH2_001223



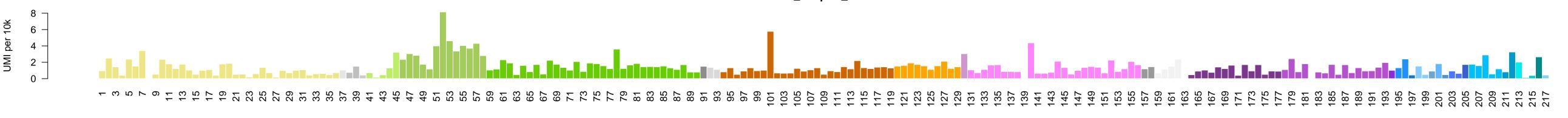




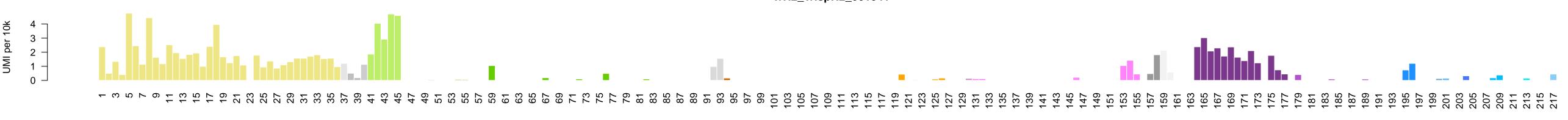
Homer.HG1.9:WAS/WASL TrH2_TrispH2_004827



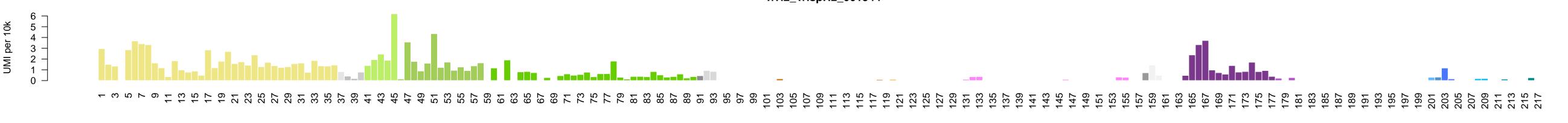
MARVEL.HG1.13:SYNPR/SYP/SYPL1/SYPL2 TrH2_TrispH2_000100



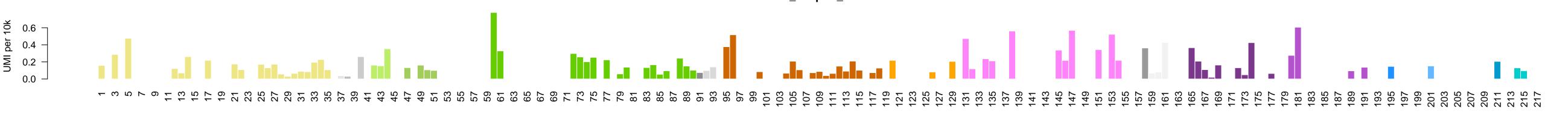
MARVEL.HG1.7:like:OCLN/SYNPR/SYP/SYPL1/SYPL2 TrH2_TrispH2_001541

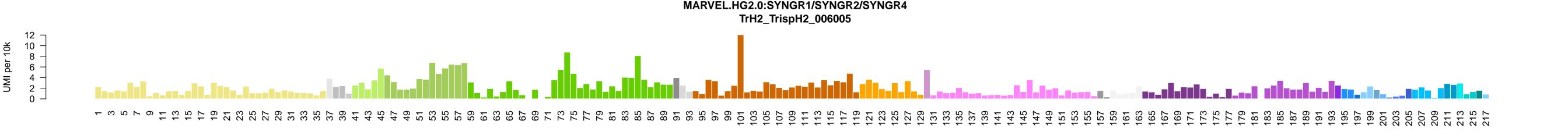


MARVEL.HG1.8:like:OCLN/SYNPR/SYP/SYPL1/SYPL2 TrH2_TrispH2_001544

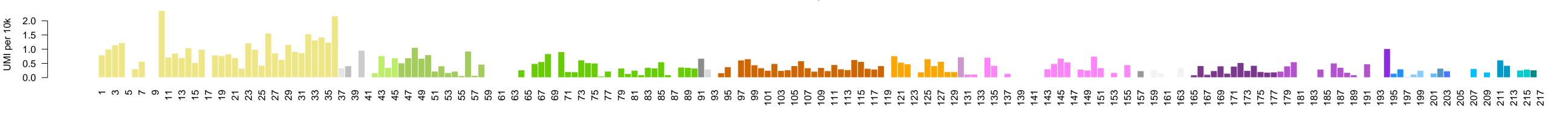


MARVEL.HG2.0:SYNGR1/SYNGR2/SYNGR4 TrH2 TrispH2 004250

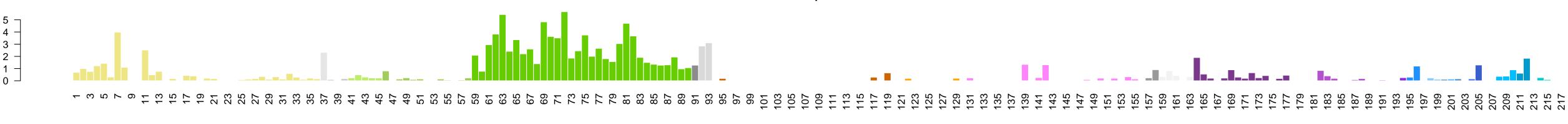




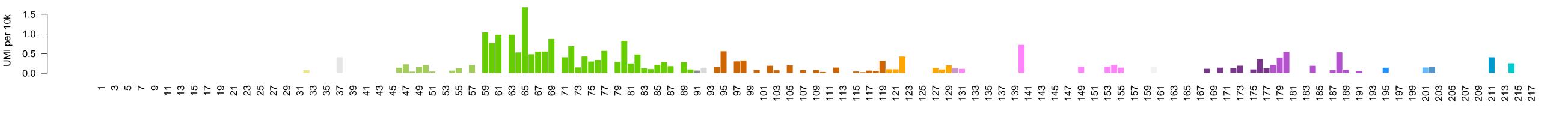
EL.HG2.0:SYNGR1/SYNGR2/SYNGR4 TrH2_TrispH2_006006



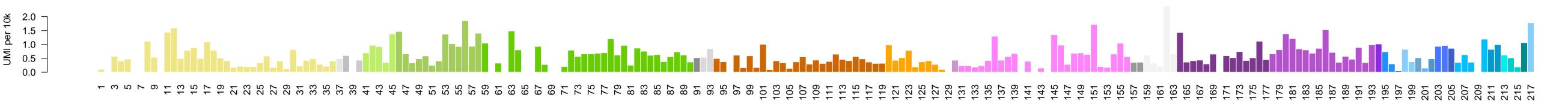
MFS_1.HG1.16:SLC16A2/SLC16A10 TrH2_TrispH2_001018



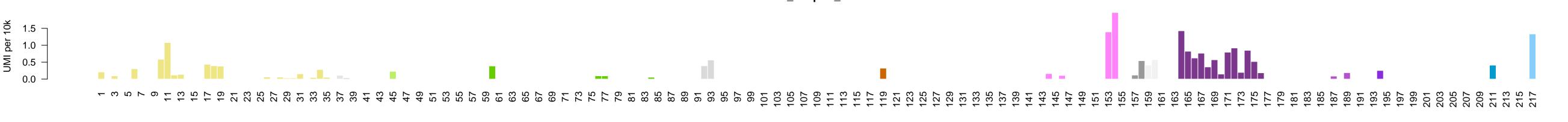
MFS_1.HG10.0:FLVCR1/FLVCR2 TrH2_TrispH2_010114



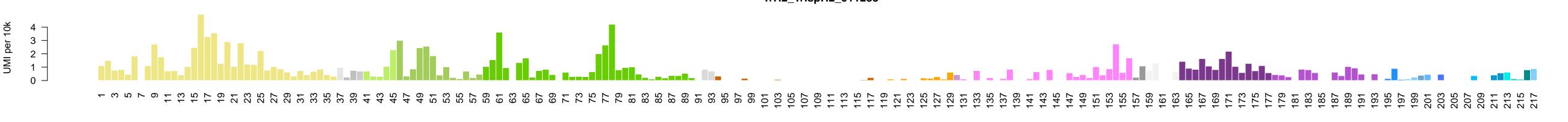
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MFS_1.HG10.1:SLC49A3 TrH2_TrispH2_008197



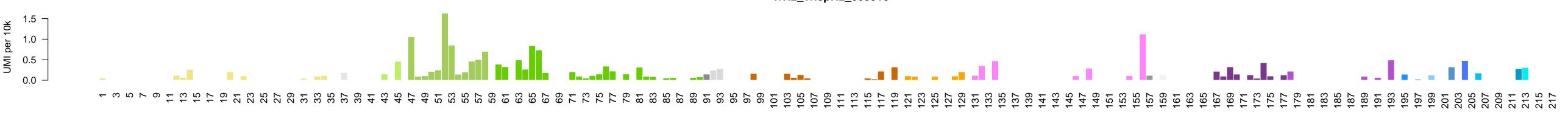
MFS_1.HG10.3:like:SLC49A3 TrH2_TrispH2_011283



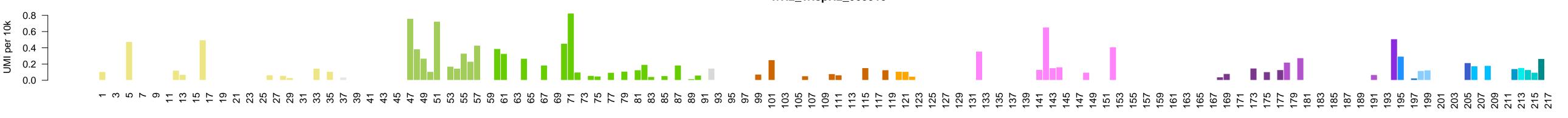
TrH2_TrispH2_003185

MFS_1.HG11.1:SLC37A4

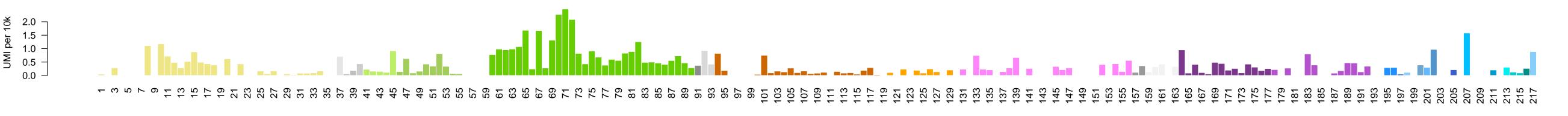
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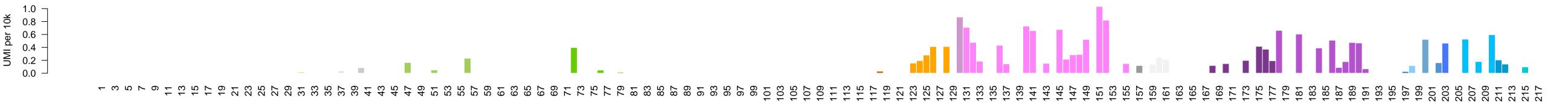
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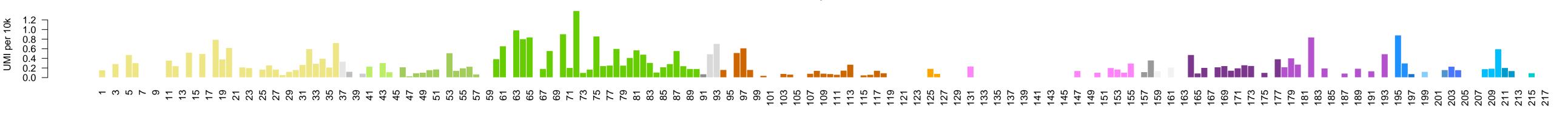
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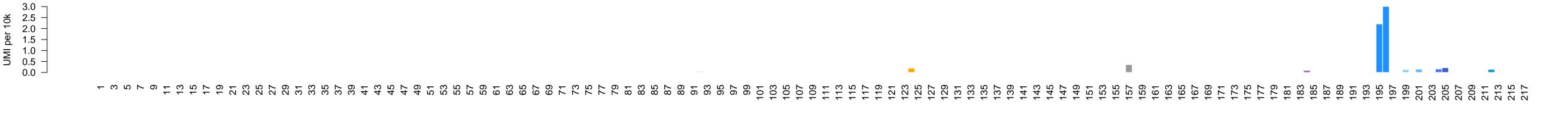
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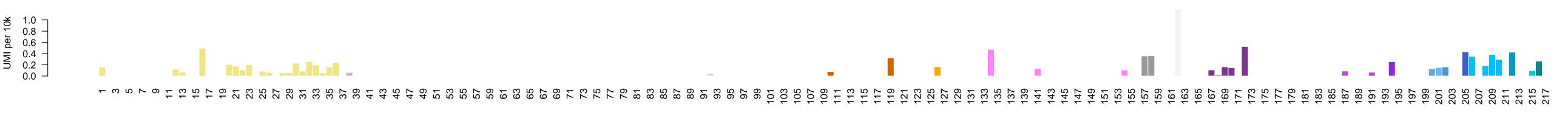
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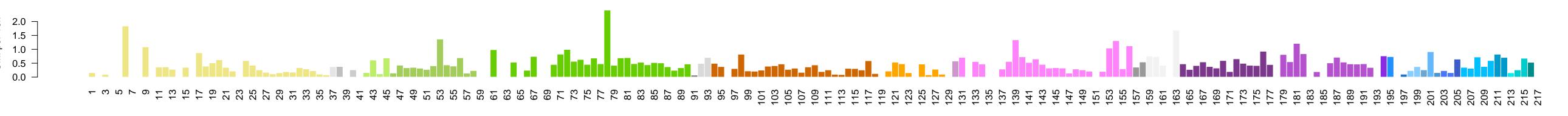
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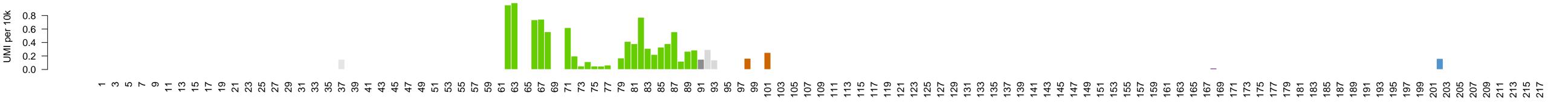




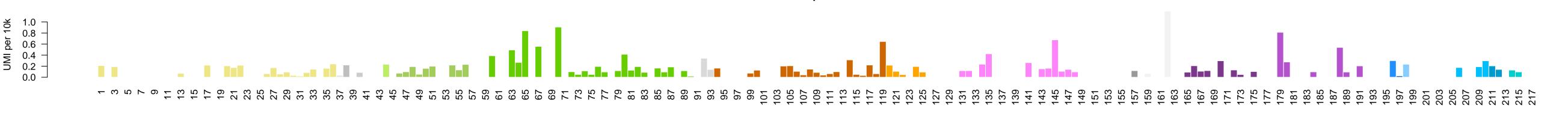
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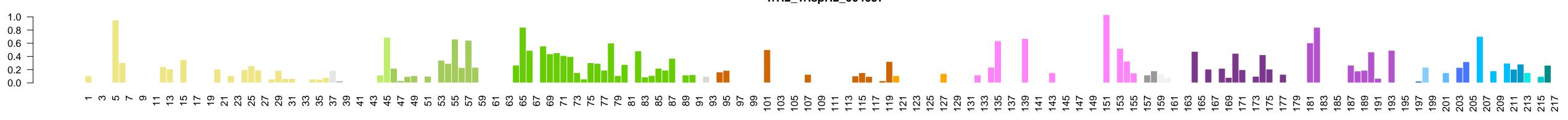
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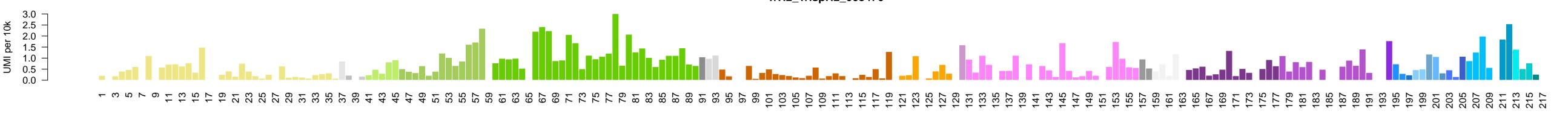
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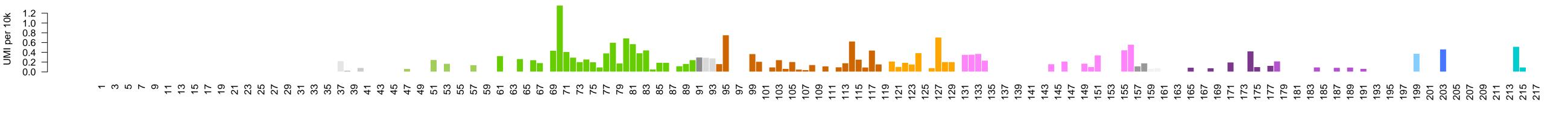
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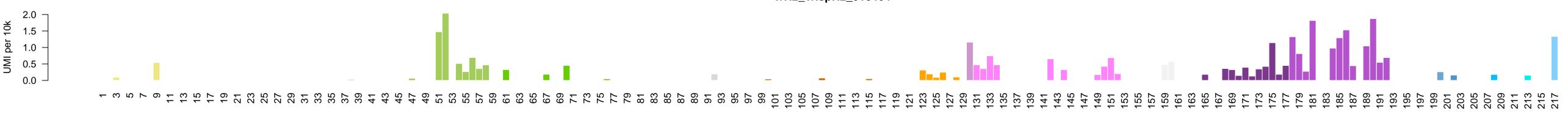
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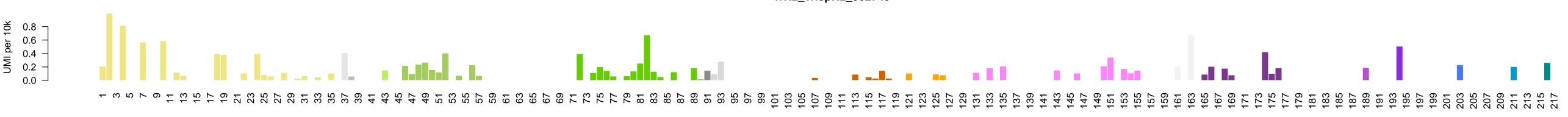
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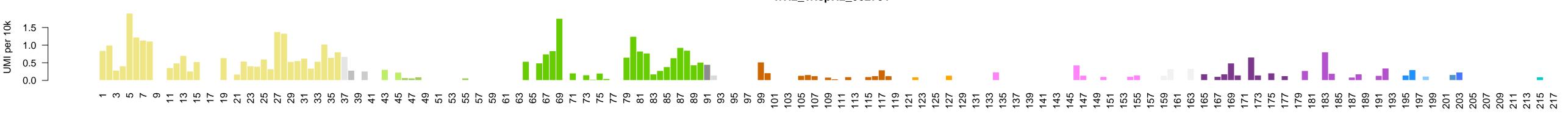
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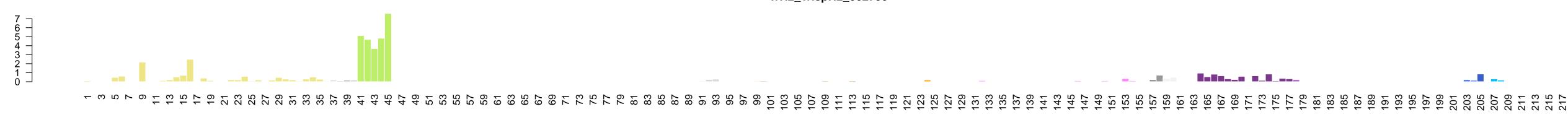
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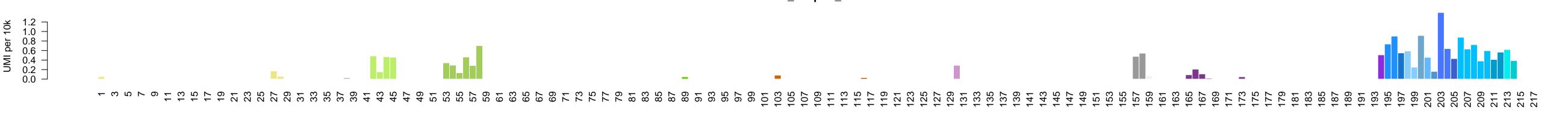
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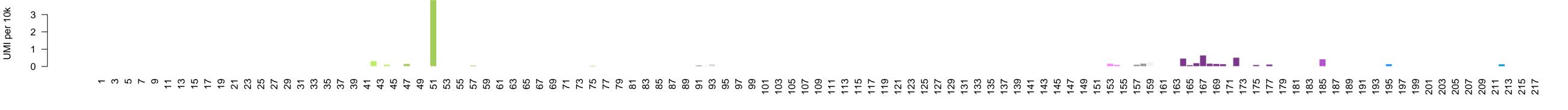
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MFS_1.HG15.1:SLC45A1/SLC45A2/SLC45A3 TrH2_TrispH2_002756

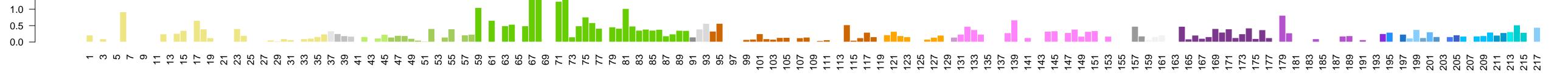


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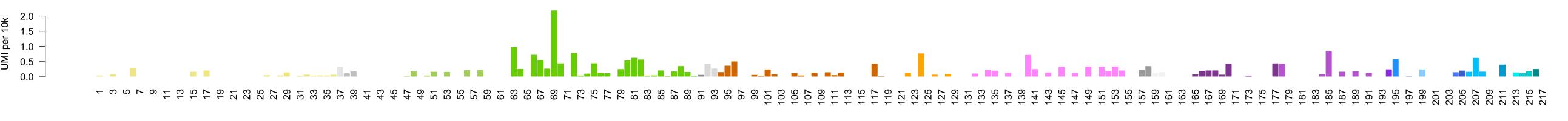
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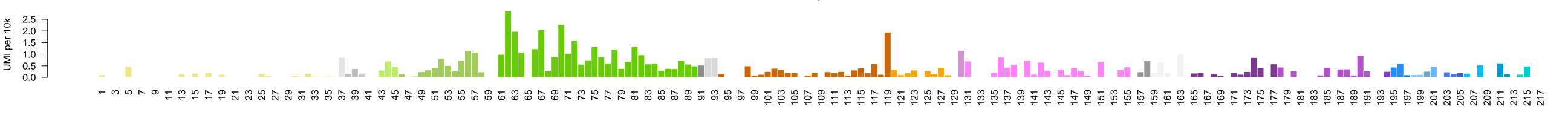
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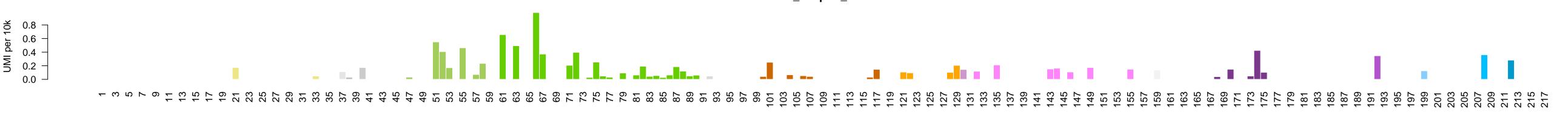
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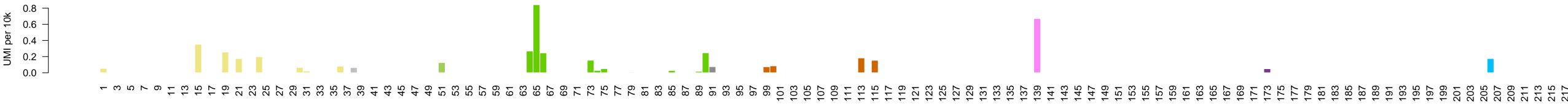
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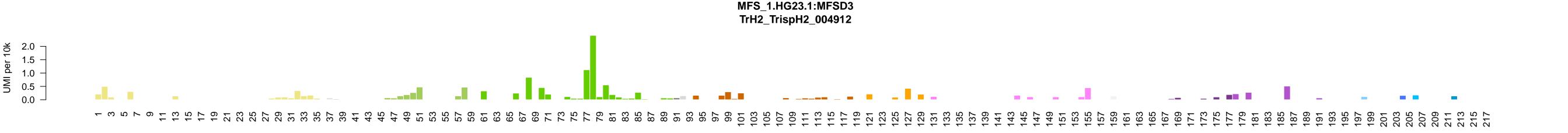


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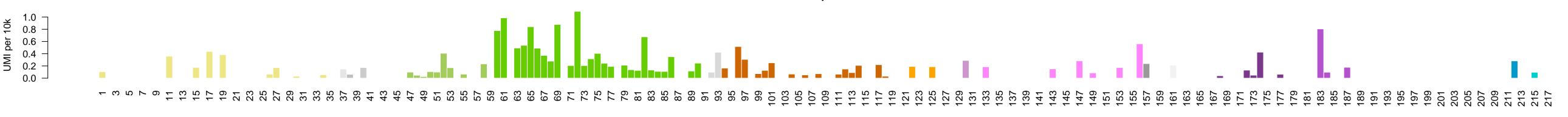


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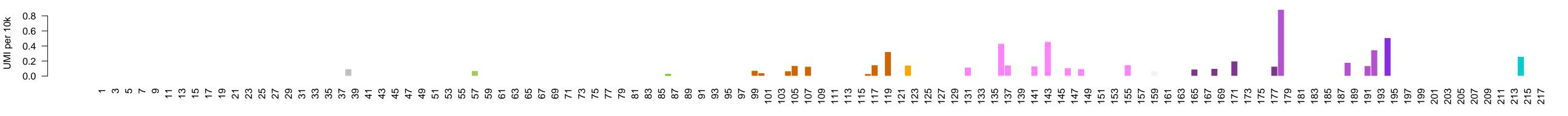




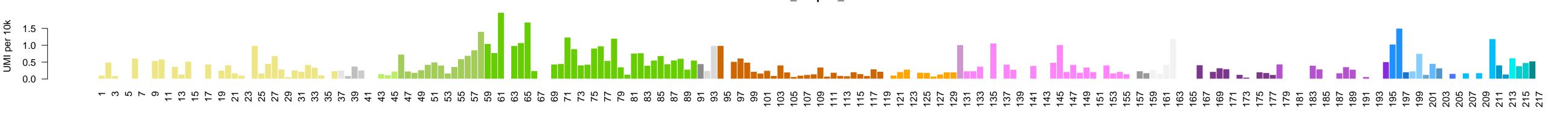
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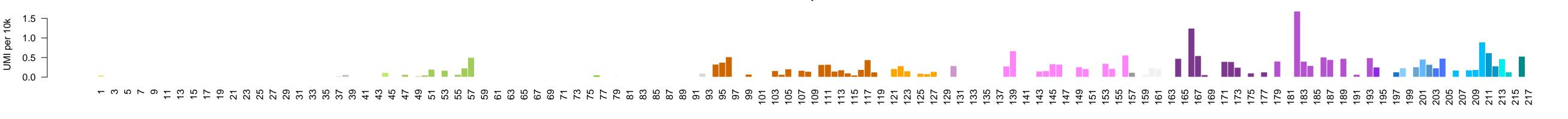
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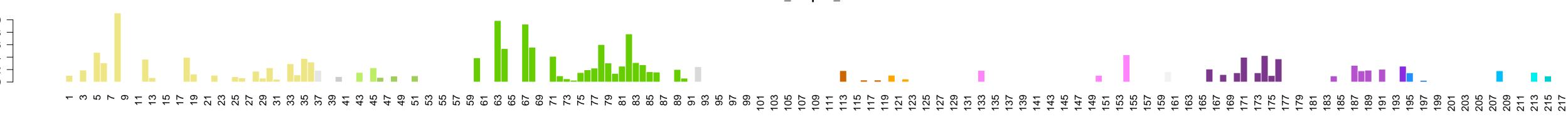
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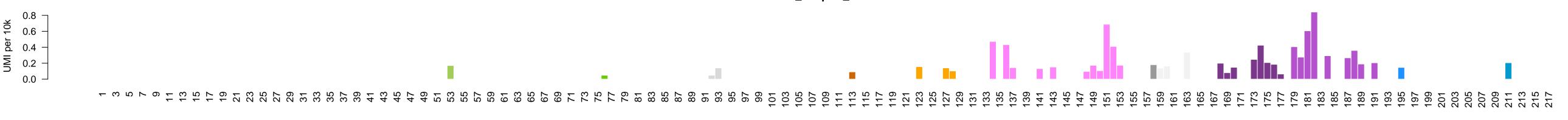
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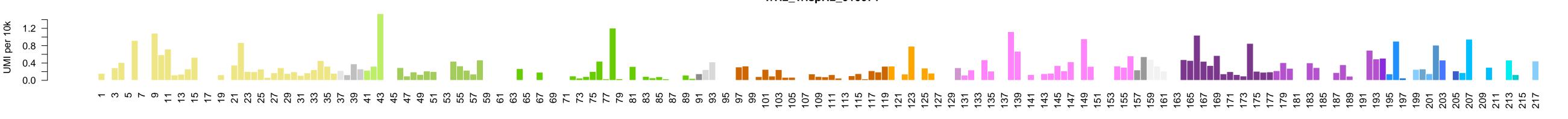
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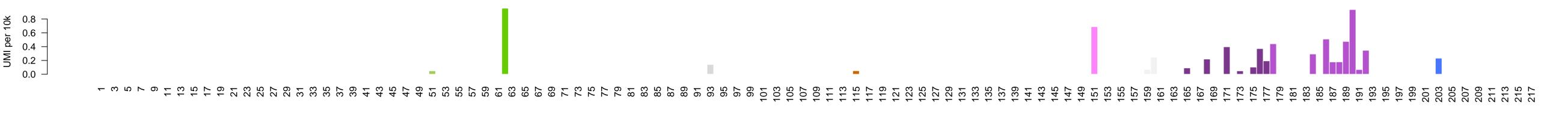
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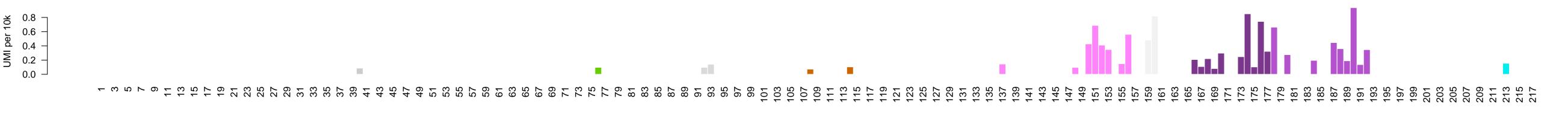
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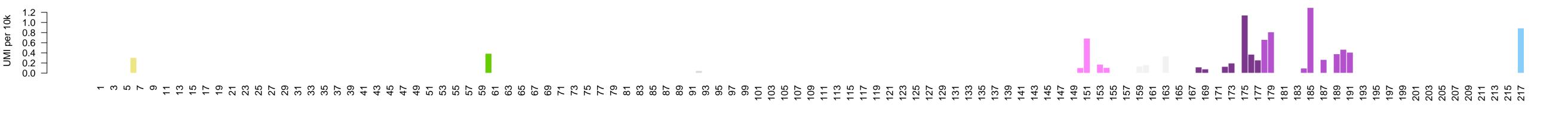
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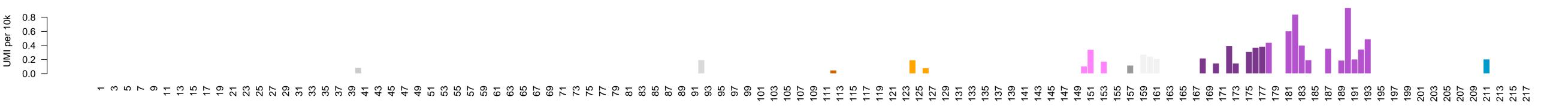
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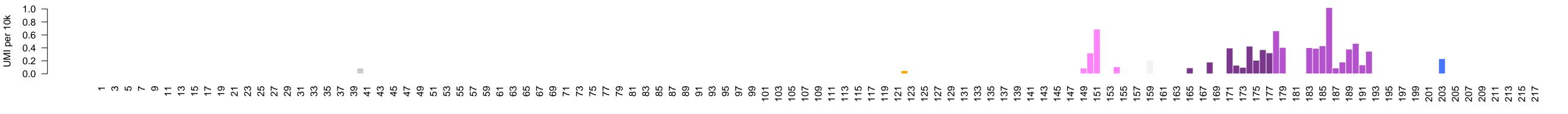
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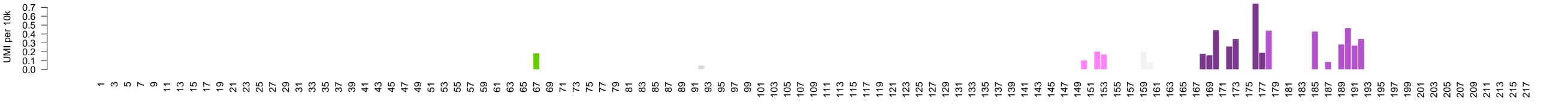
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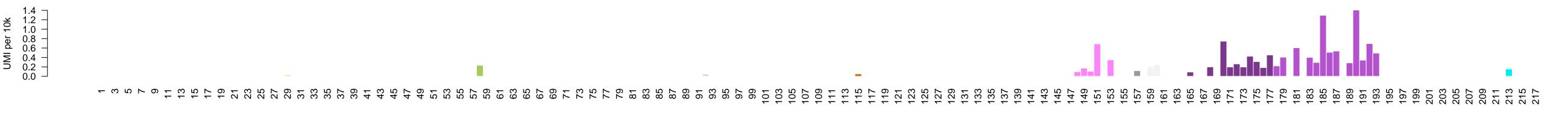
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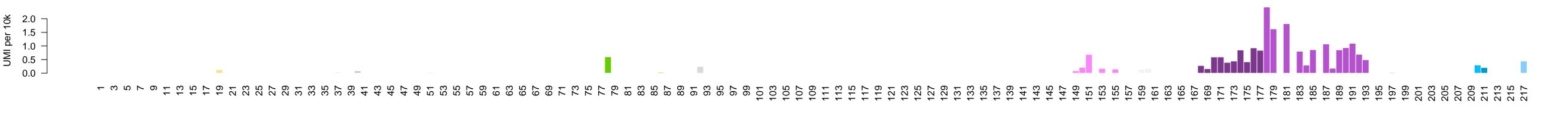
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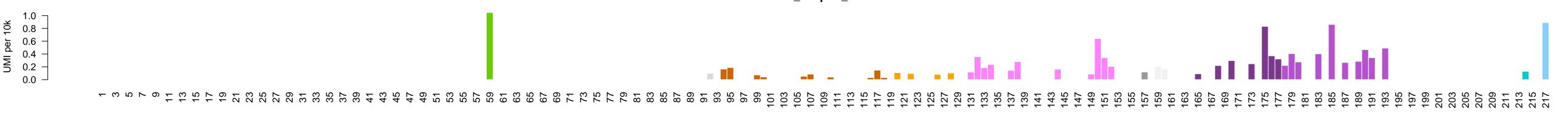
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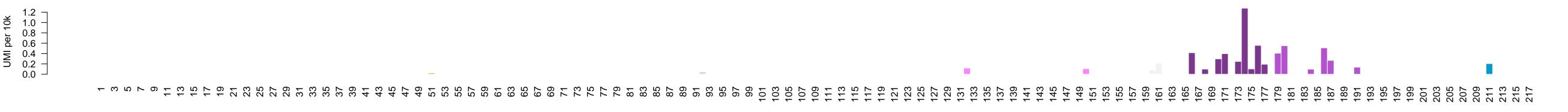
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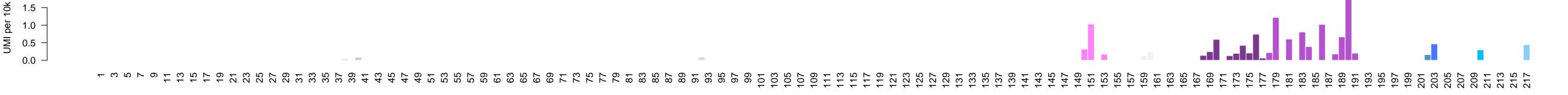
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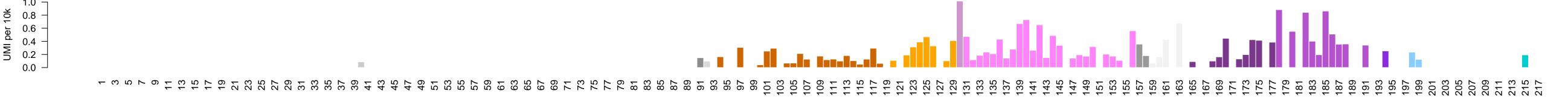
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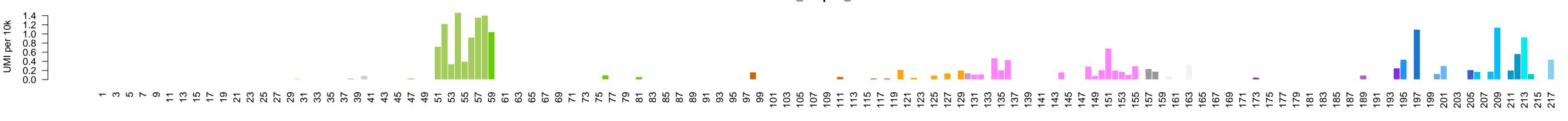
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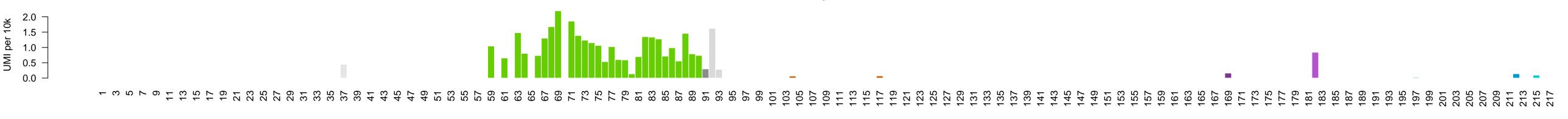
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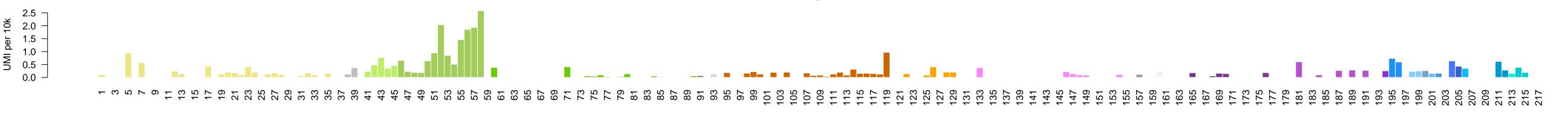


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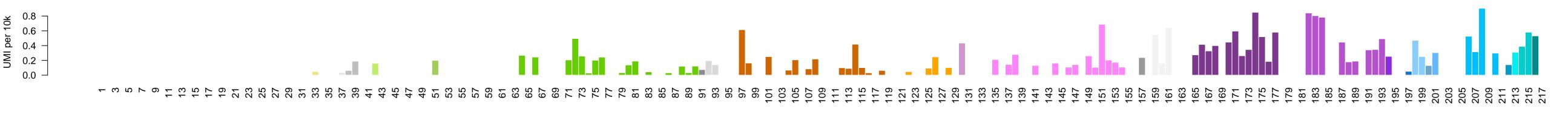


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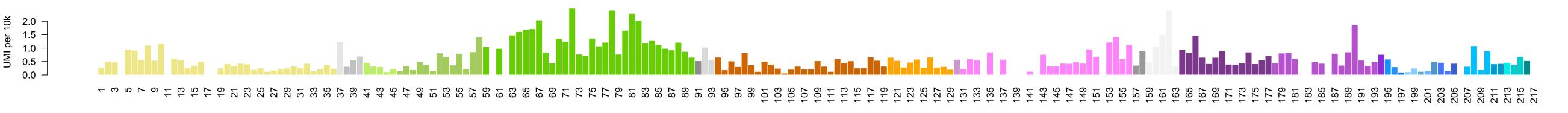
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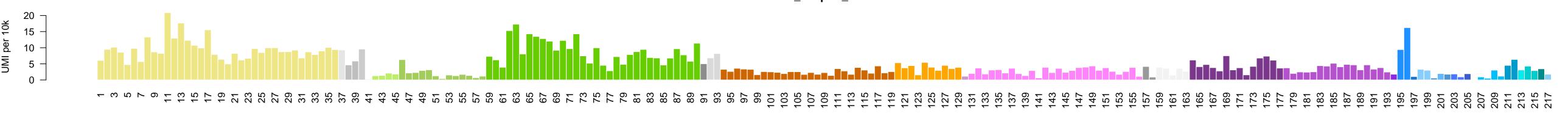
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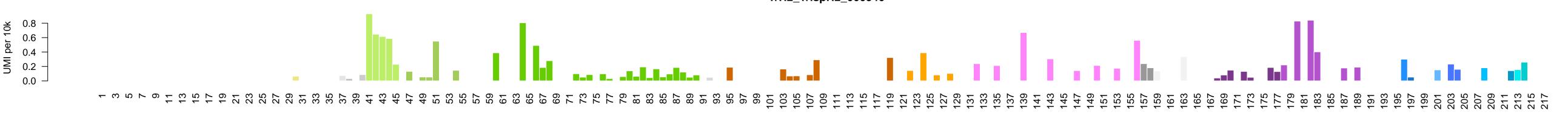
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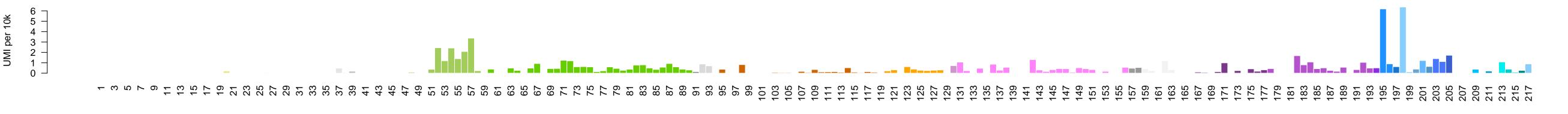
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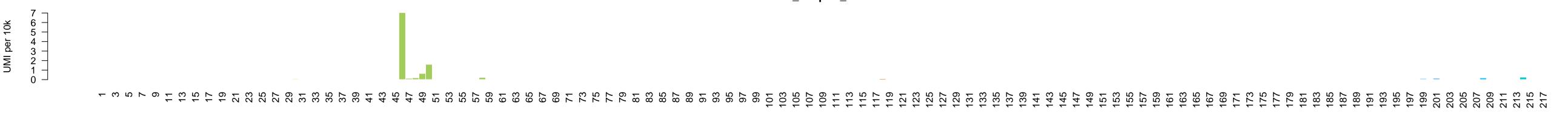
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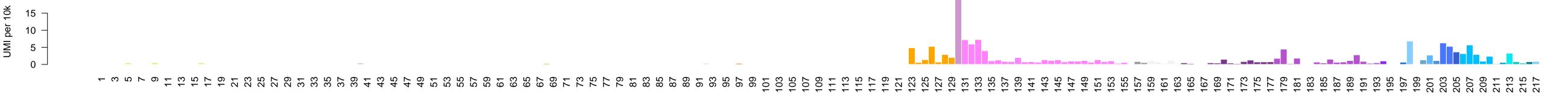
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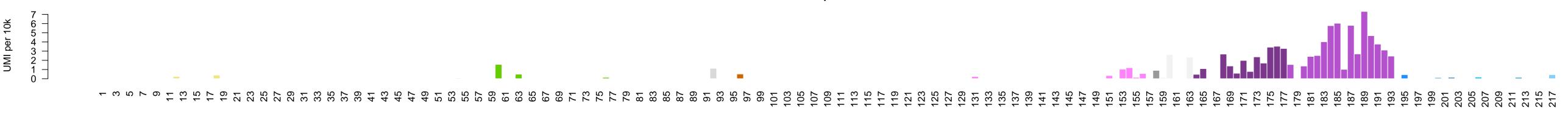


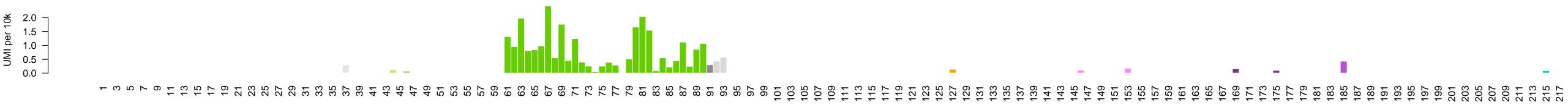
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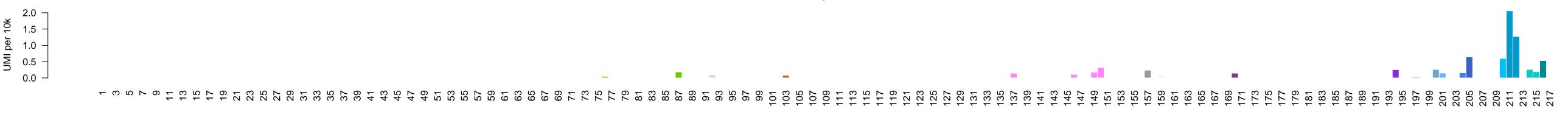


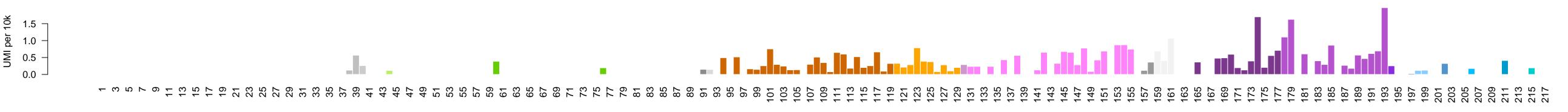
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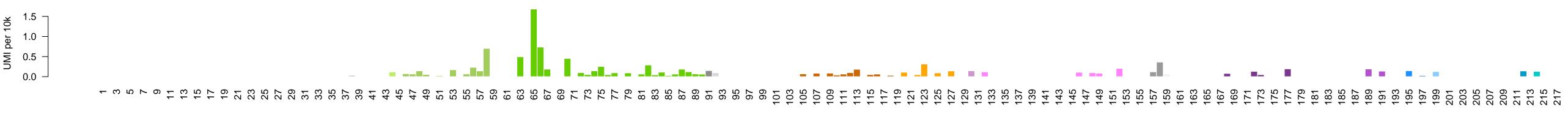


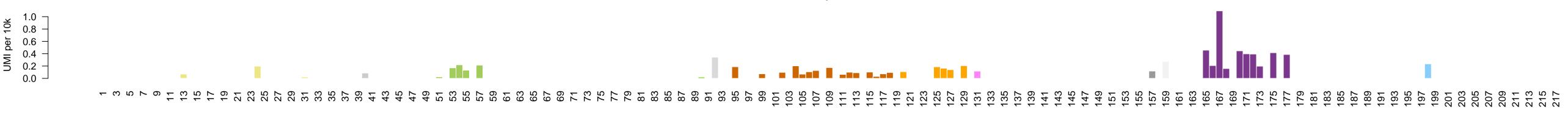


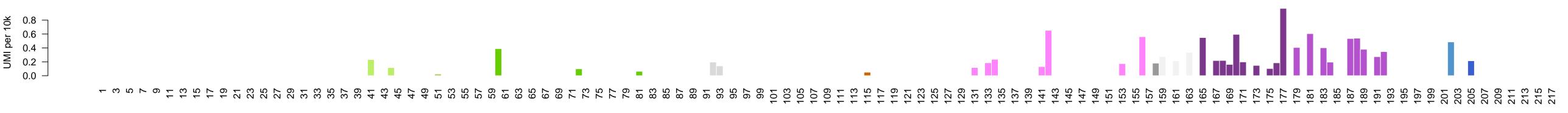


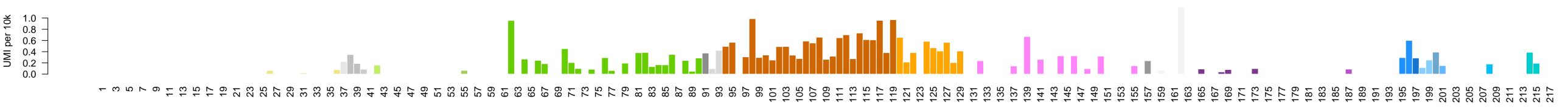






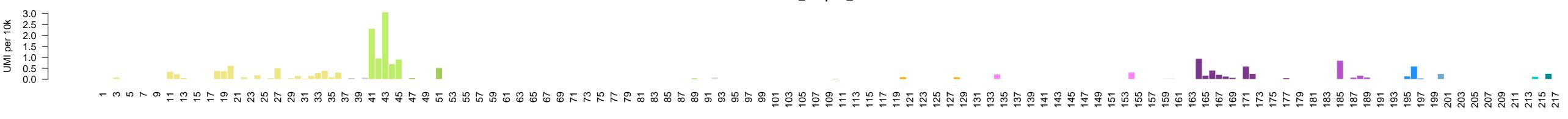






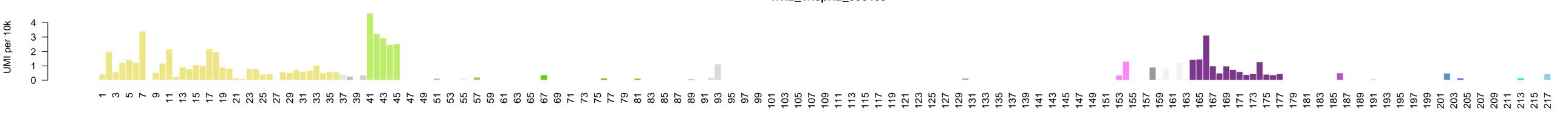
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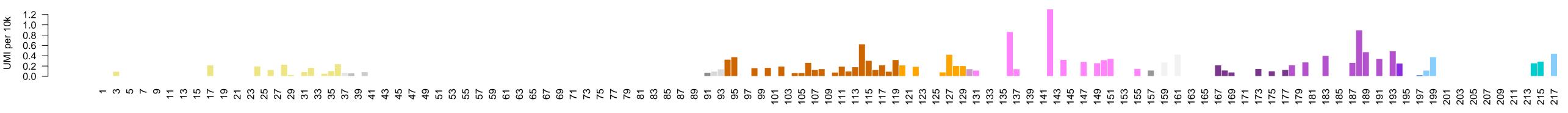
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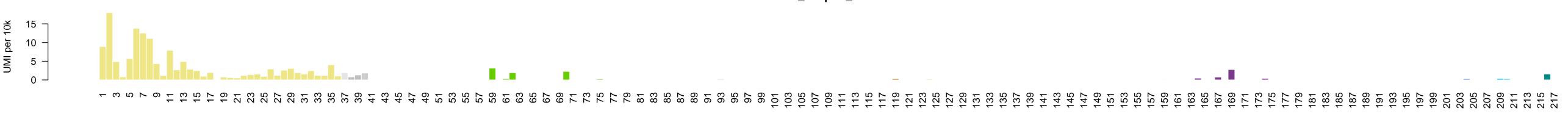


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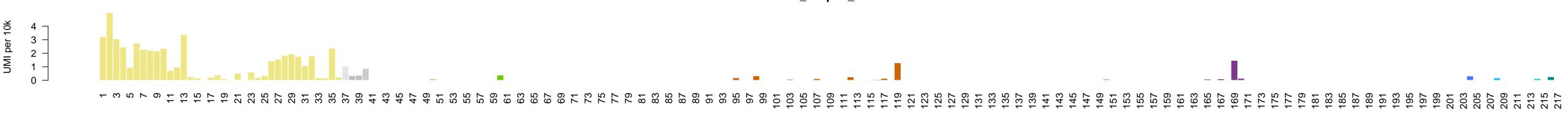






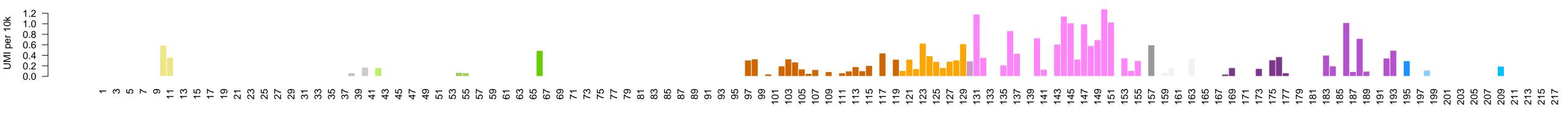
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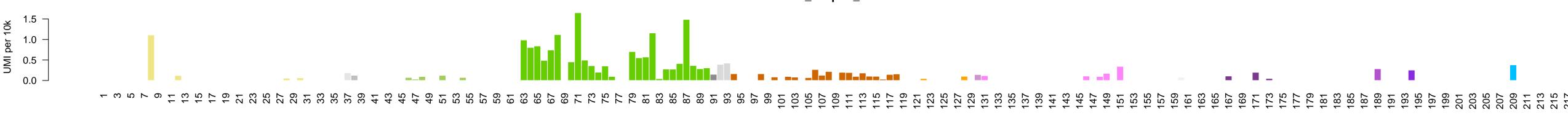
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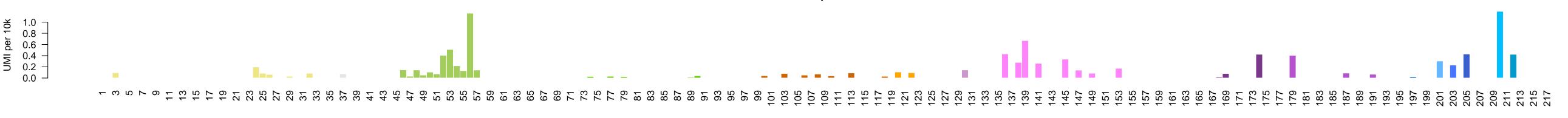
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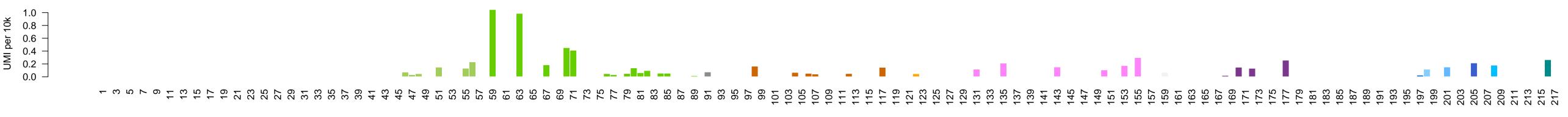


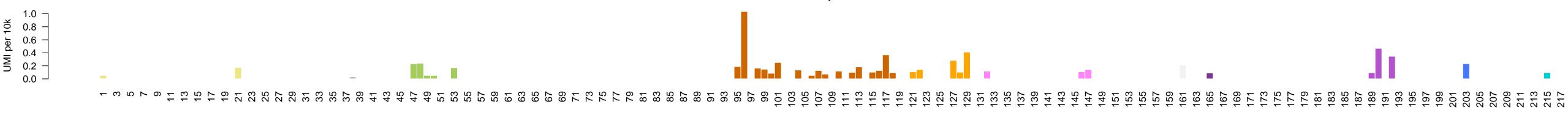


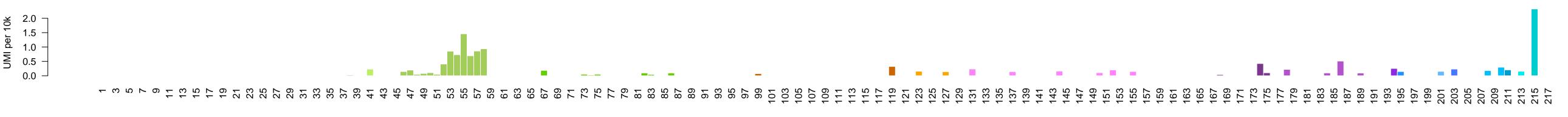
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TrH2_TrispH2_008906



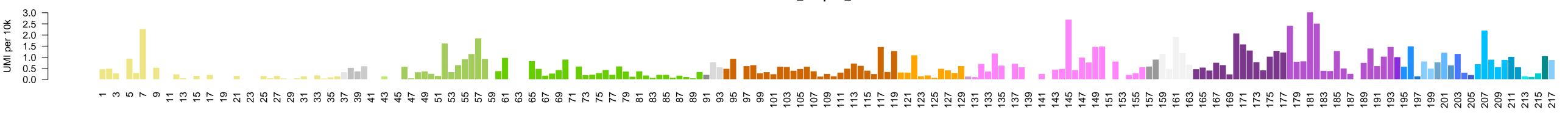


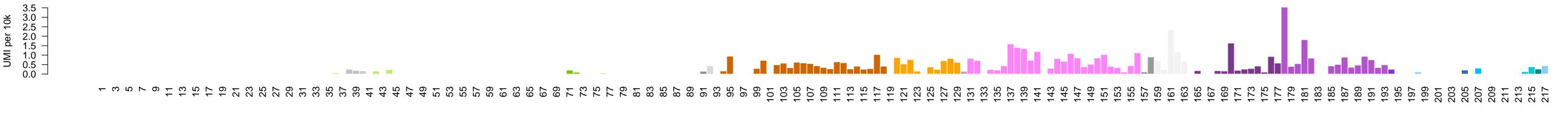




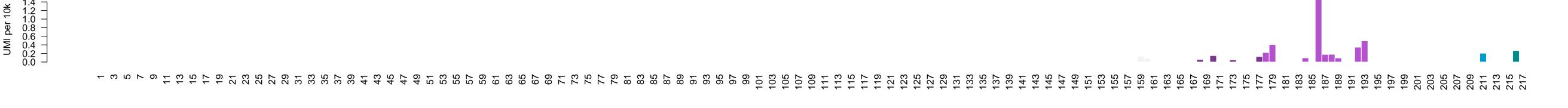
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TrH2_TrispH2_011183

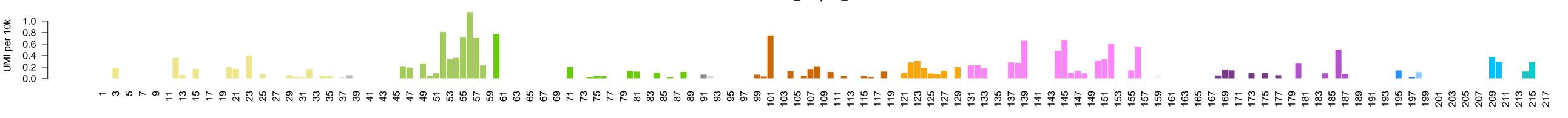




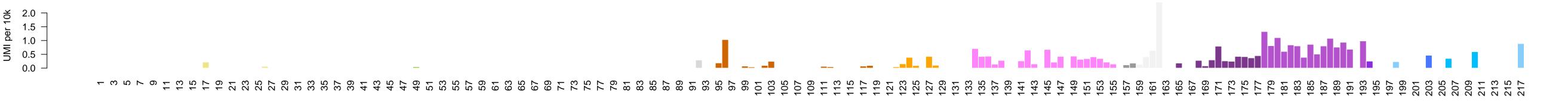
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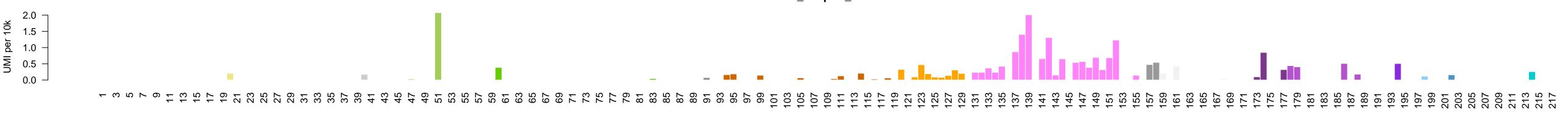
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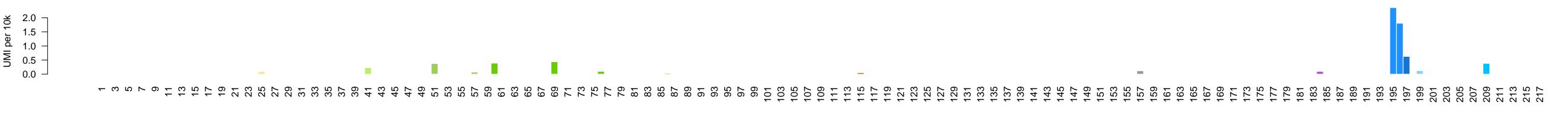
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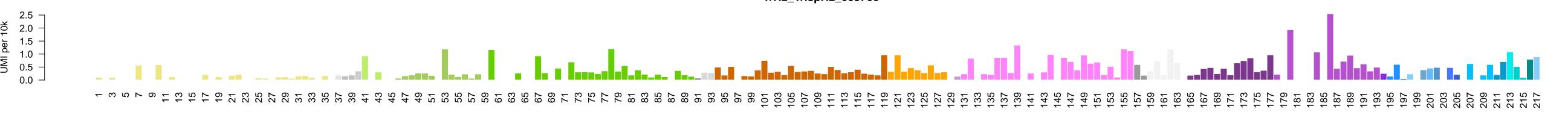
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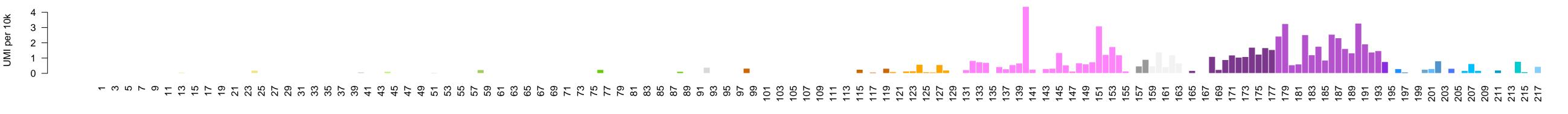
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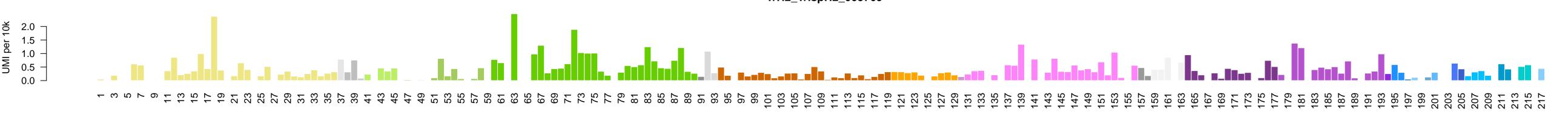
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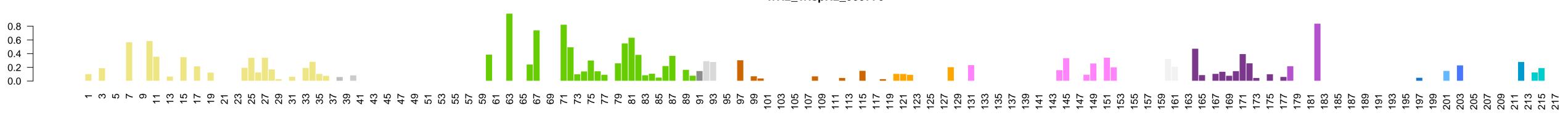
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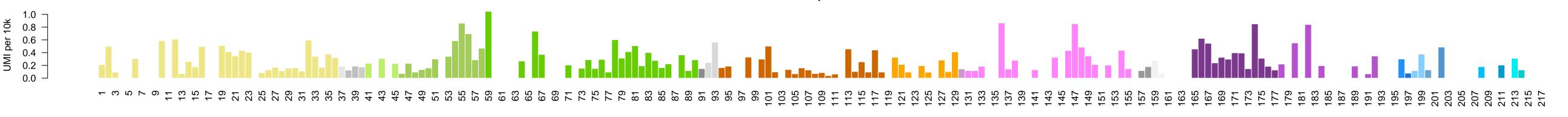
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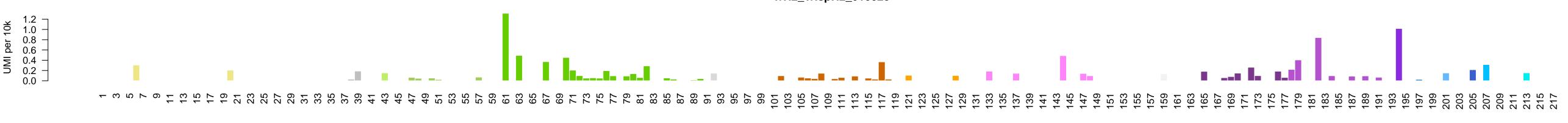
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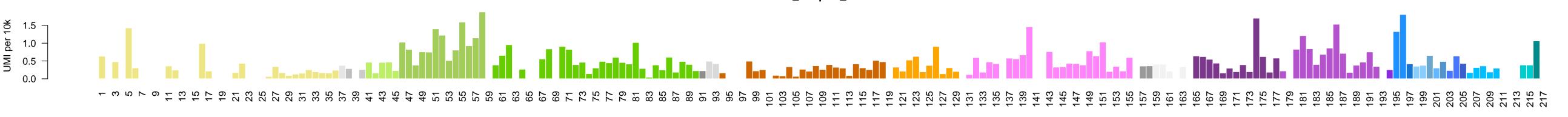
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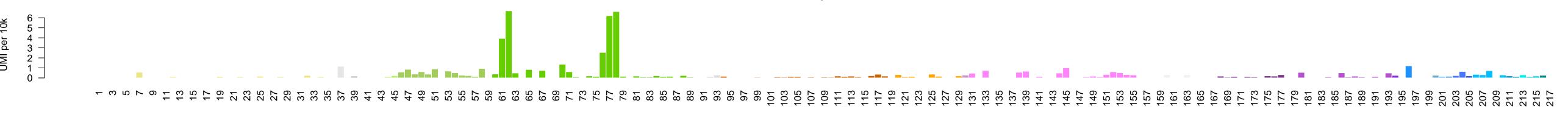


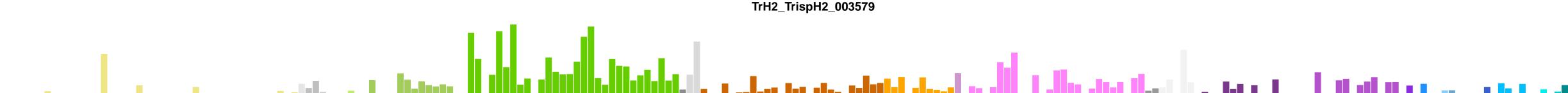
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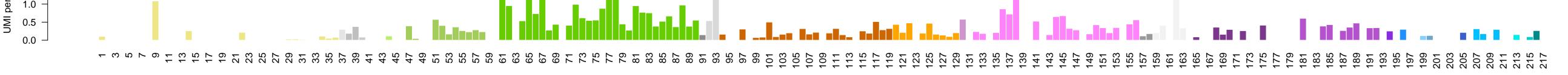
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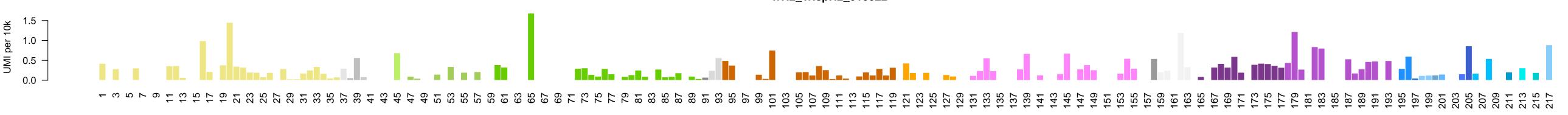




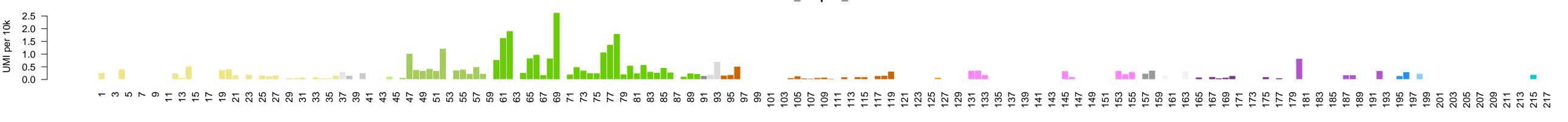
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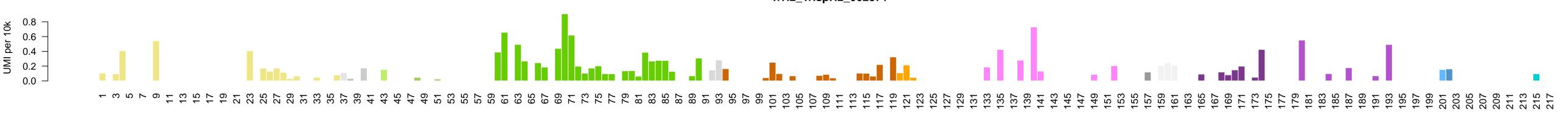
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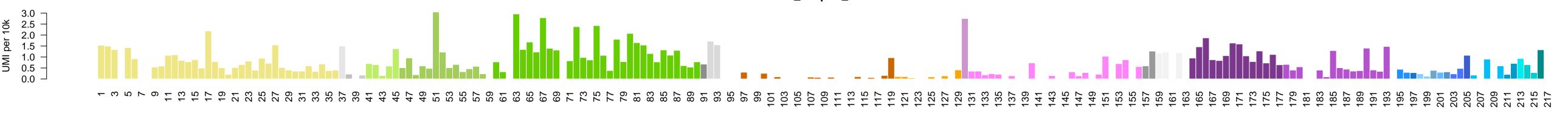
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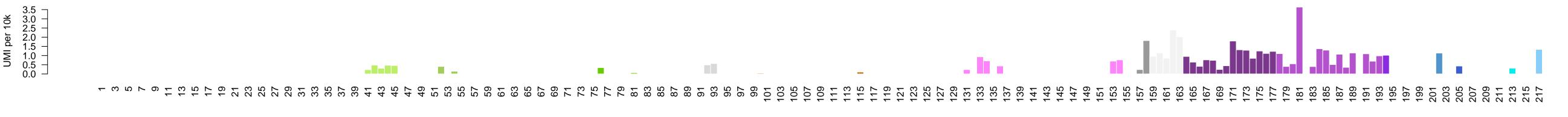
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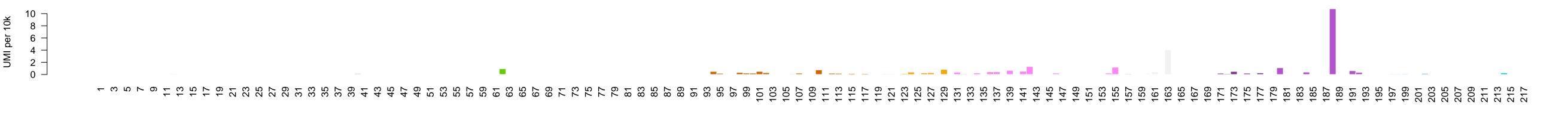
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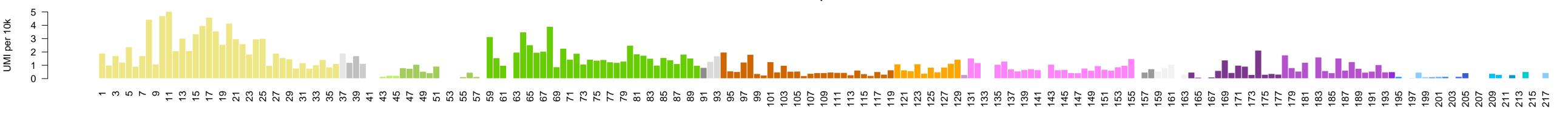
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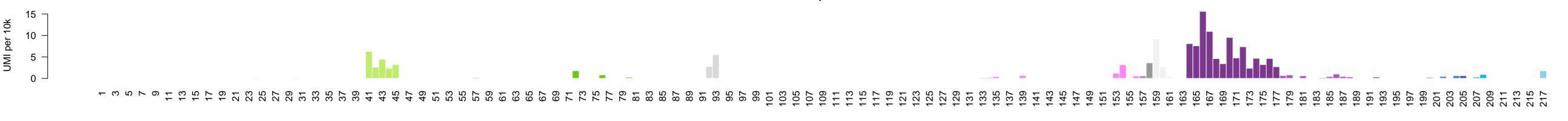
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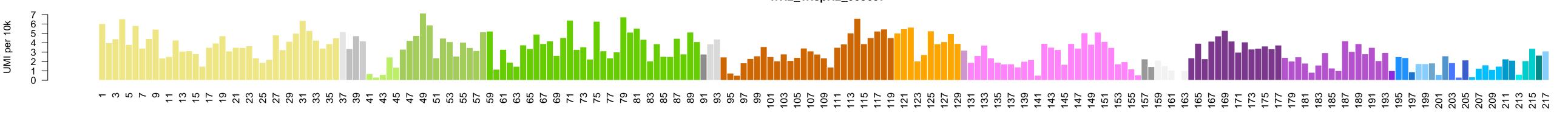
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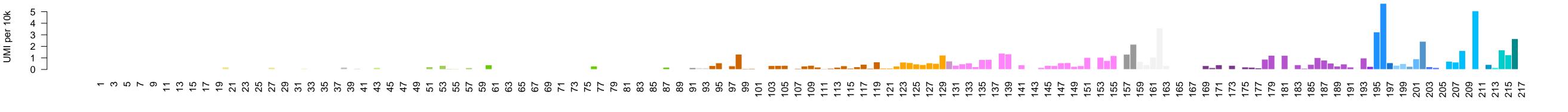
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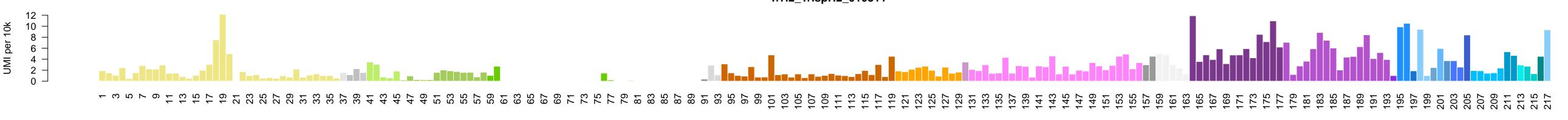
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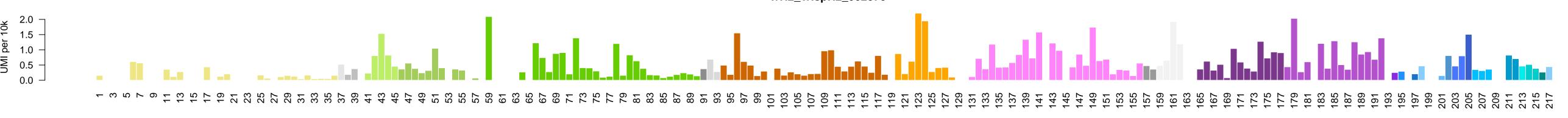
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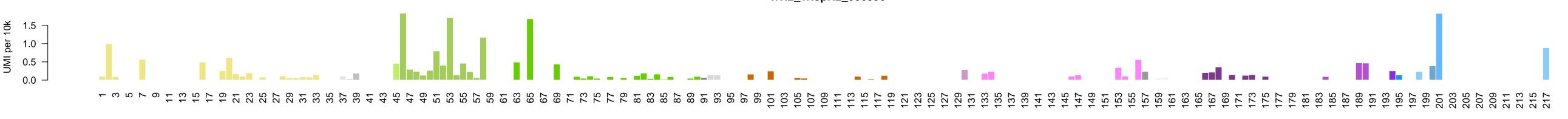
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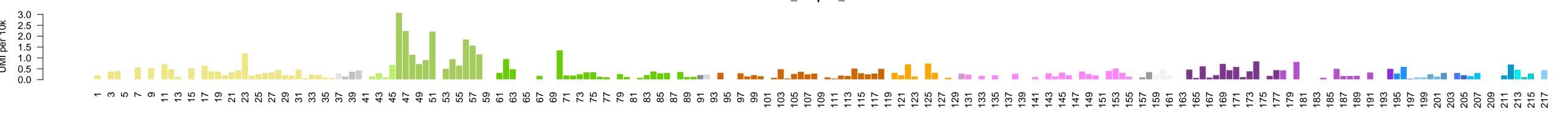
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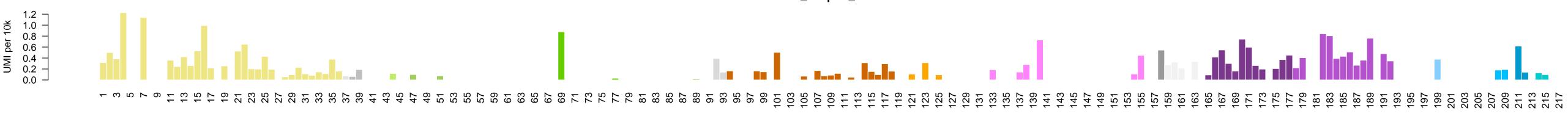
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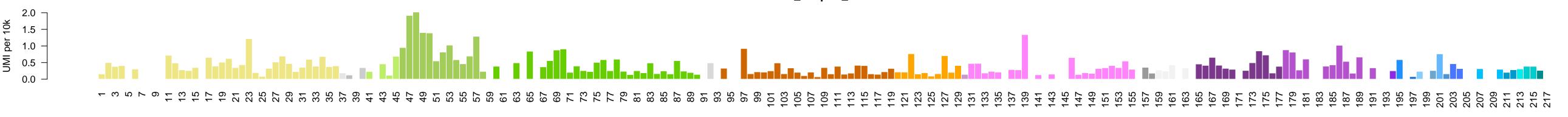
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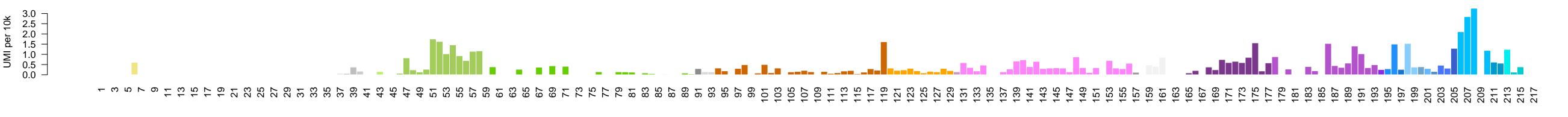
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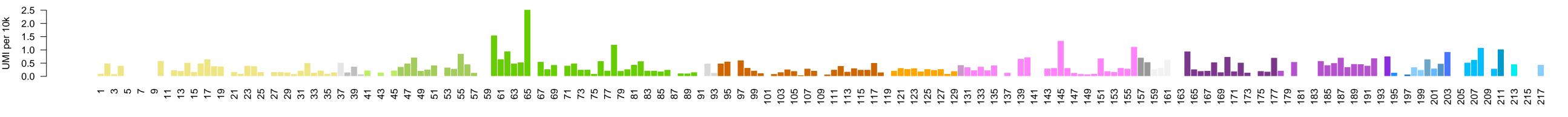


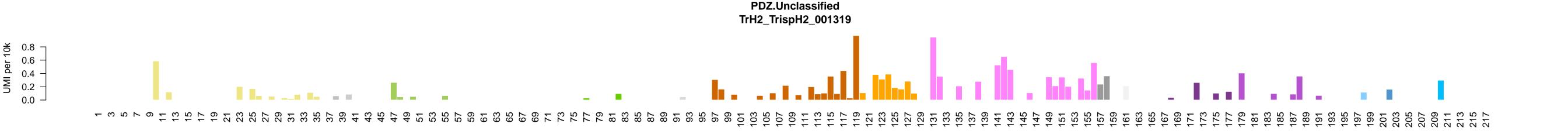
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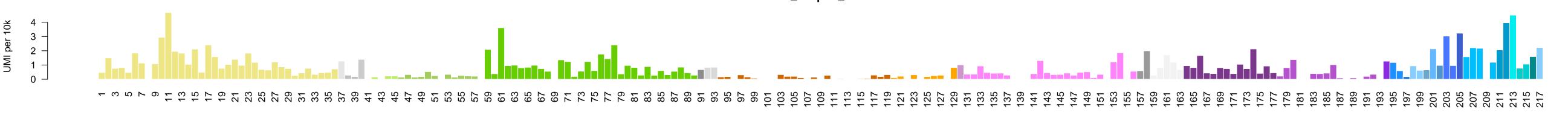
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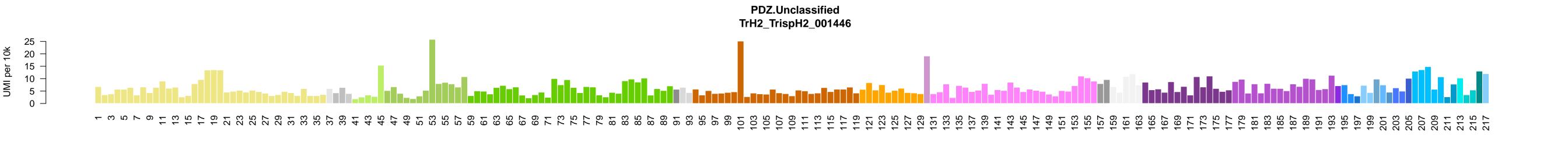


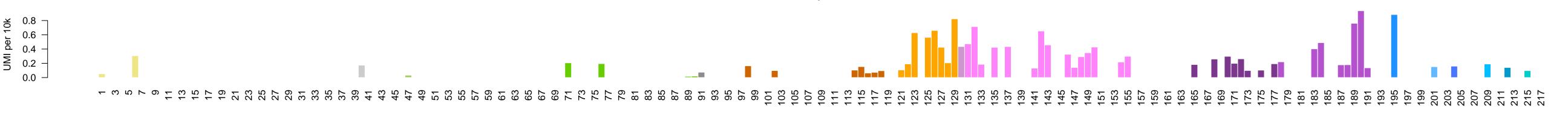




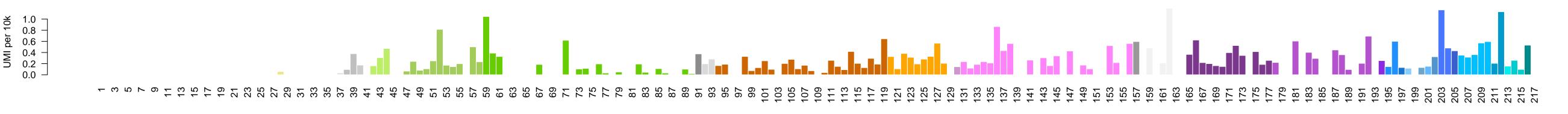
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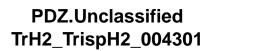


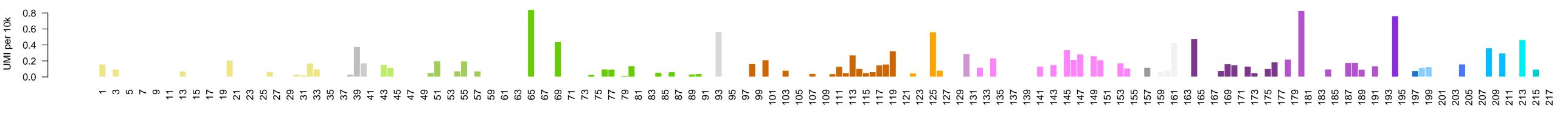




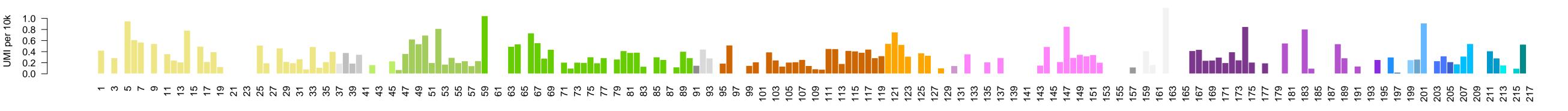
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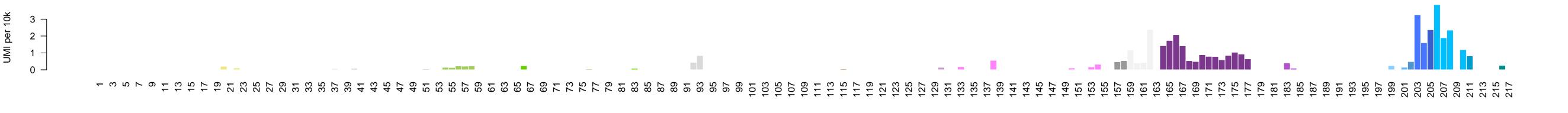


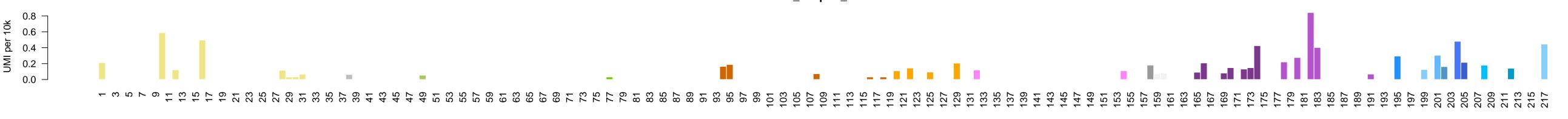


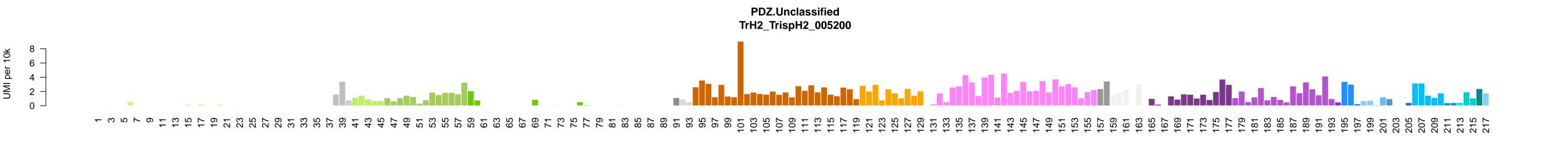
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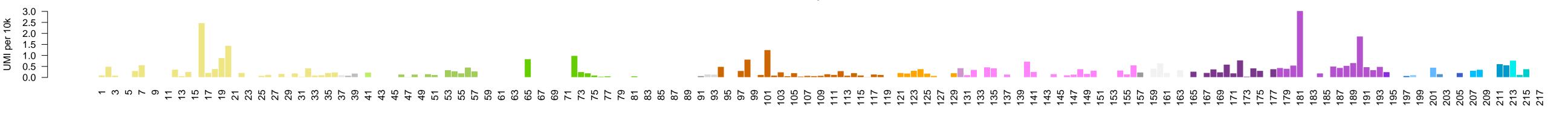
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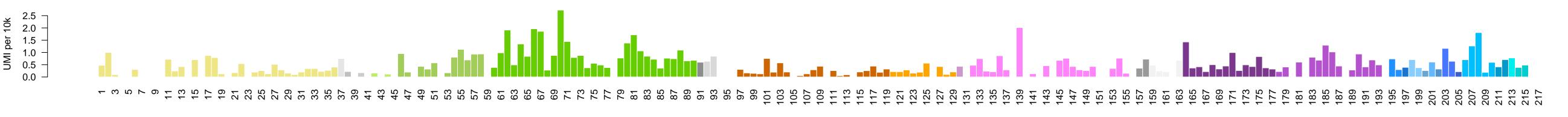




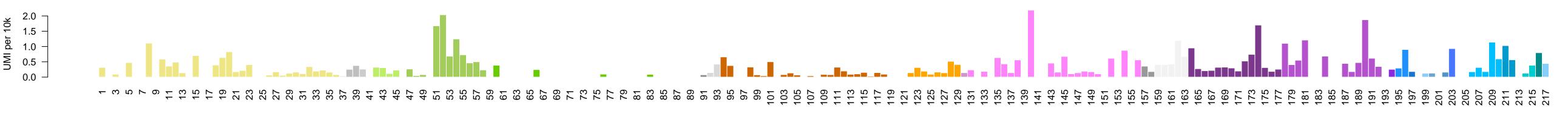


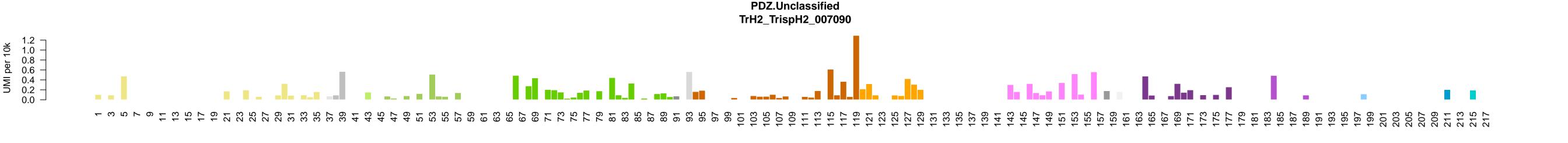
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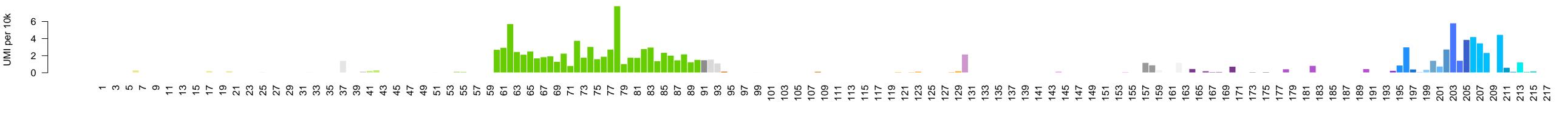


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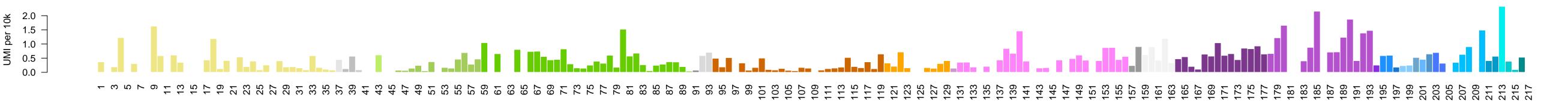




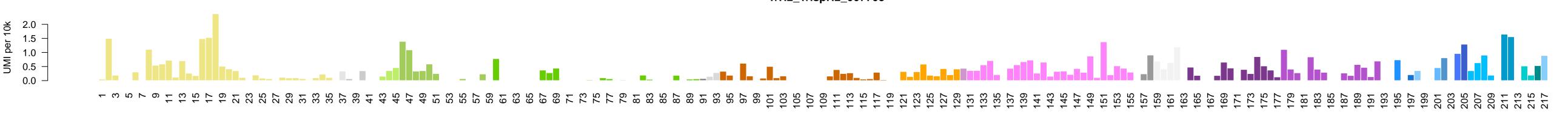




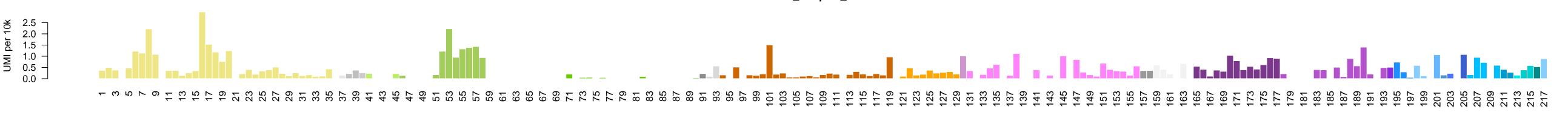
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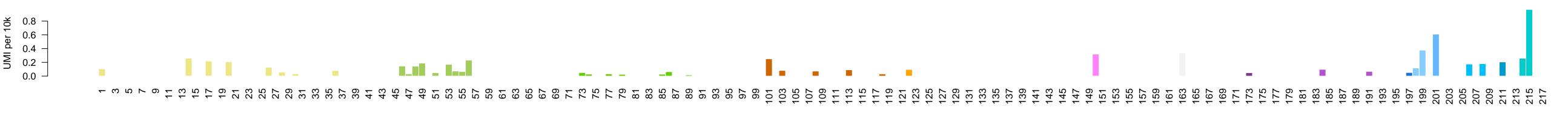


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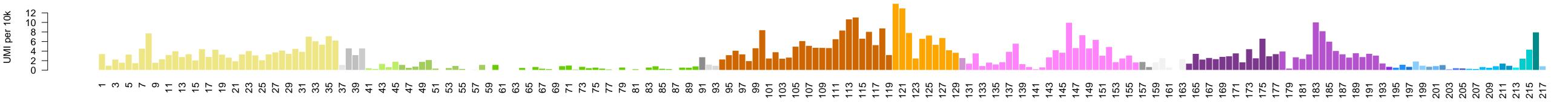


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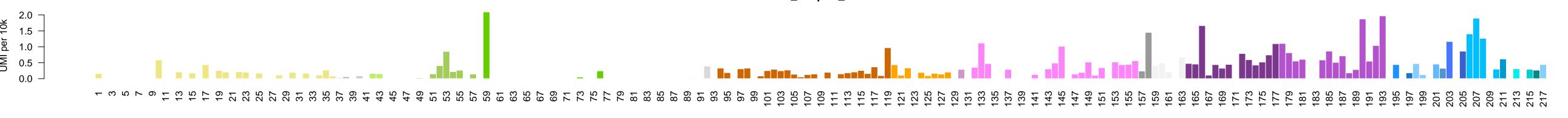




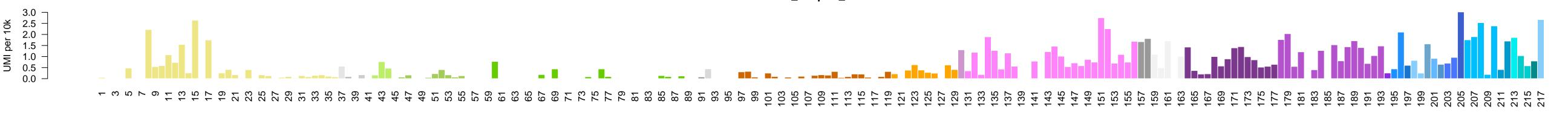




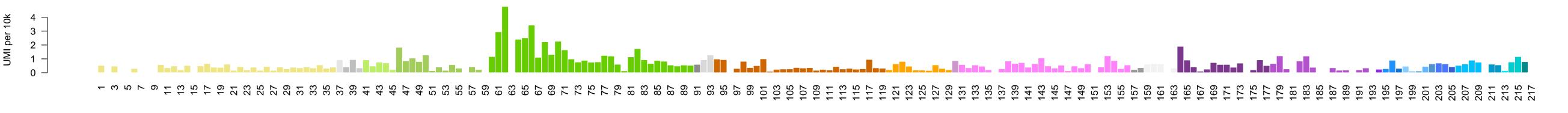
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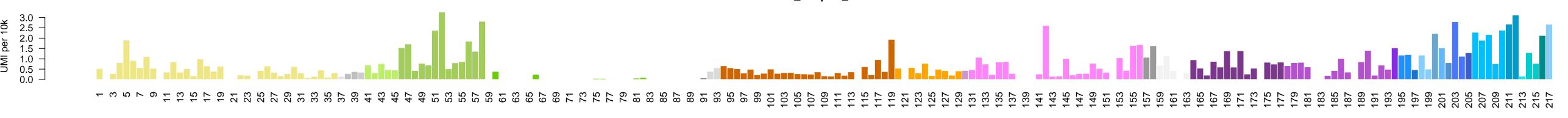
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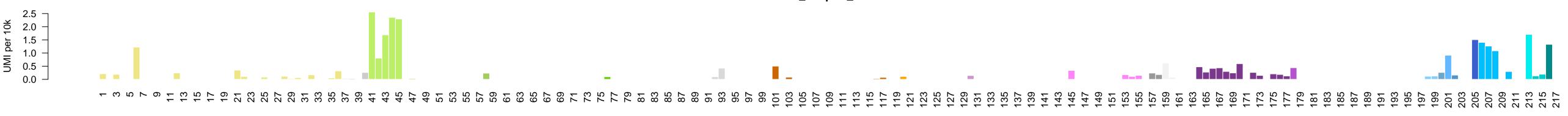


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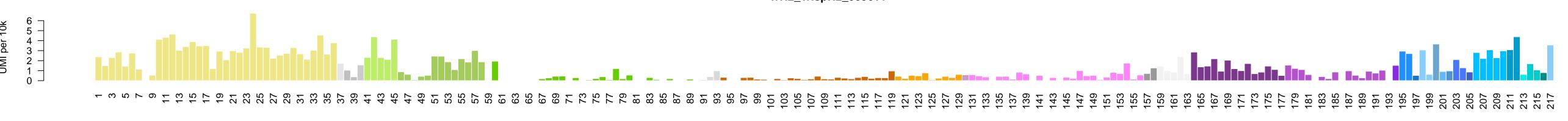


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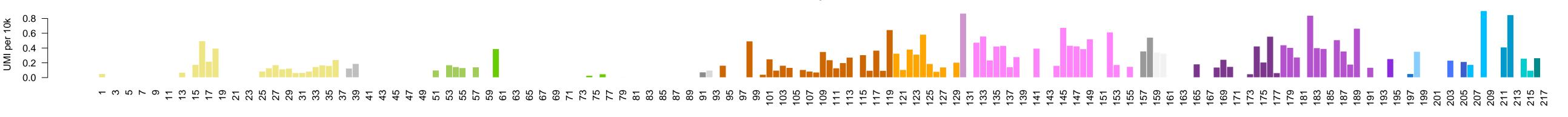


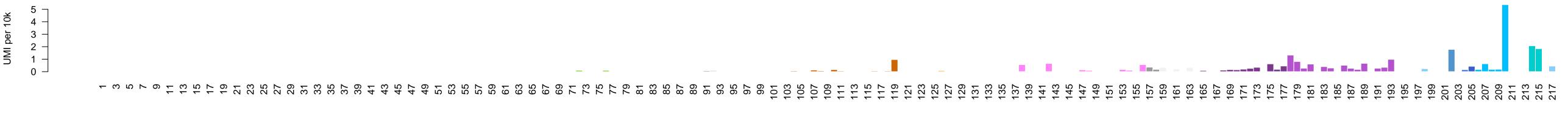


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TrH2 TrispH2 009611

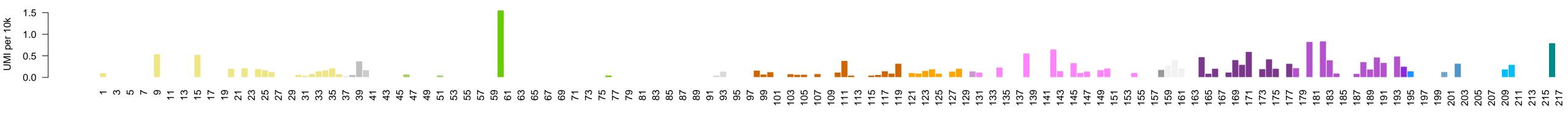


PDZ.Unclassified
TrH2_TrispH2_010021

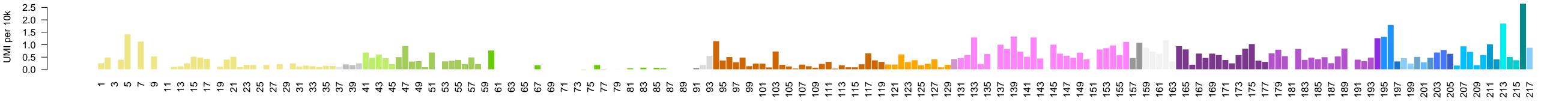


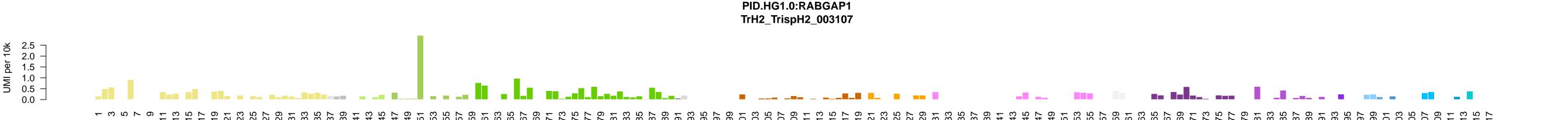




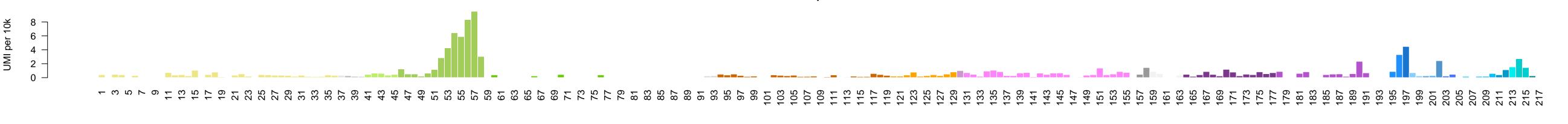


PDZ.Unclassified
TrH2 TrispH2 011429

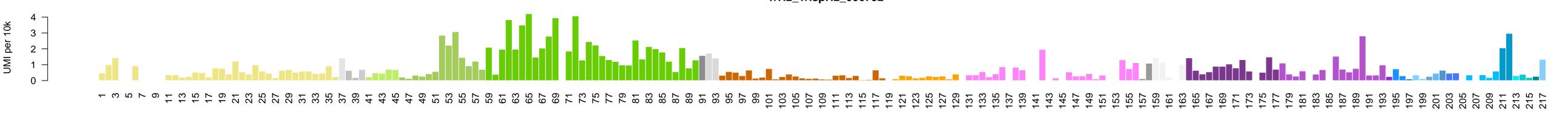




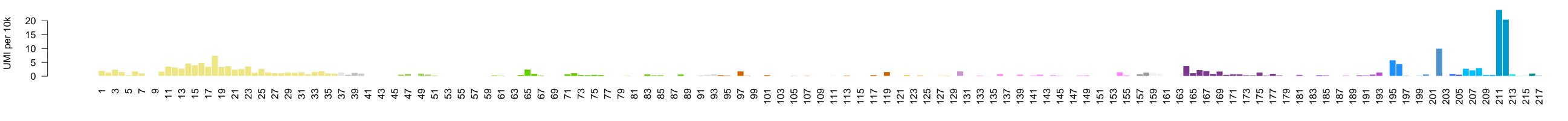
PID.HG1.1:TBC1D1/TBC1D4 TrH2_TrispH2_007739



PID.HG1.12:SHC1/SHC2/SHC3/SHC4 TrH2_TrispH2_006752

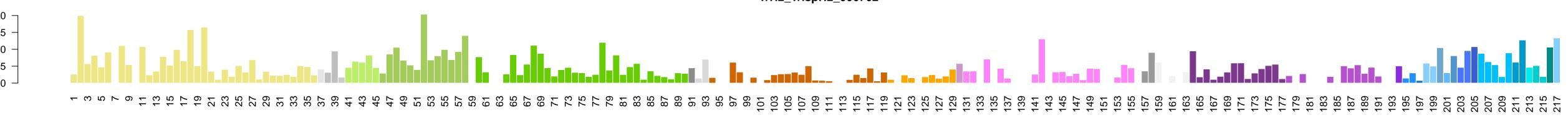


PID.HG1.14:NUMB/NUMBL TrH2_TrispH2_006441

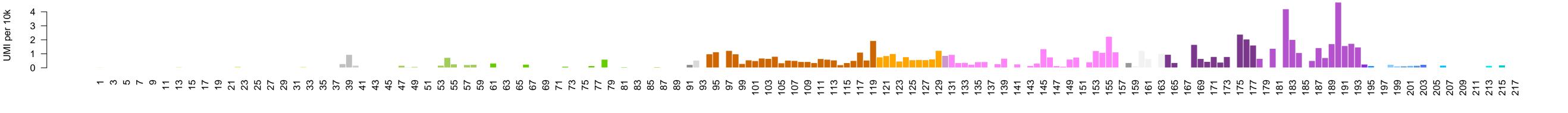


PID.HG1.16:MAPK8IP1/MAPK8IP2 TrH2_TrispH2_003875

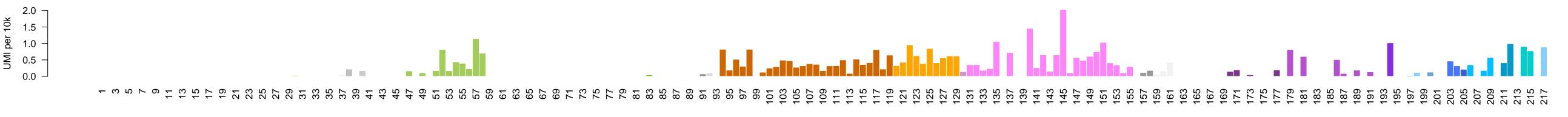
PID.HG1.19:APBA1/APBA2/APBA3 TrH2_TrispH2_006762



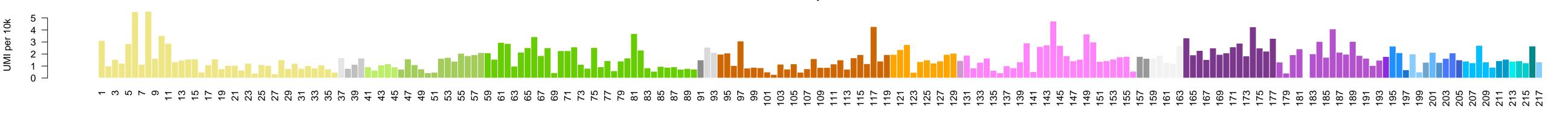
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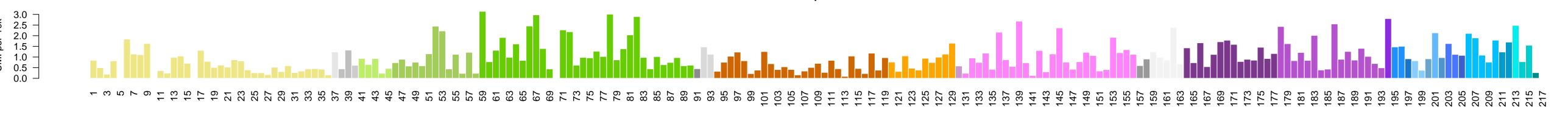
PID.HG1.8:like:LDLRAP1/PID1 TrH2_TrispH2_001443



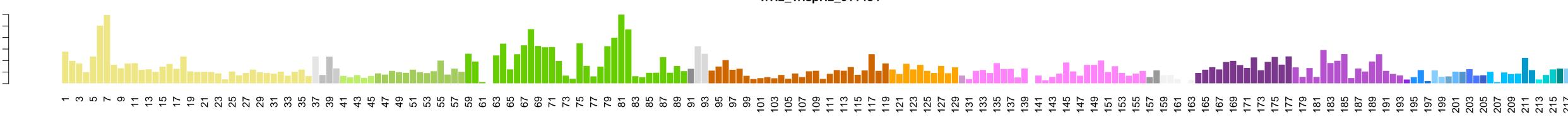
PMCA.HG1.1:ATP2A1/ATP2A2/ATP2A3 TrH2_TrispH2_006864



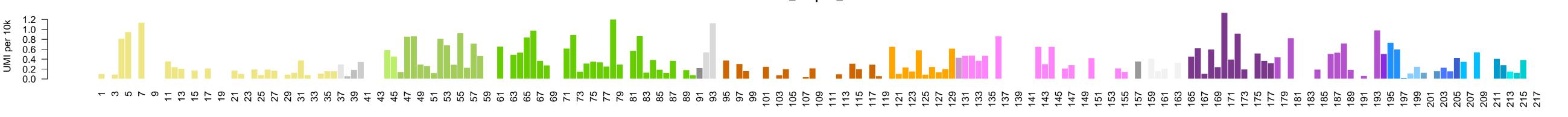
PMCA.HG1.3:ATP2B1/ATP2B2/ATP2B3/ATP2B4 TrH2_TrispH2_001165



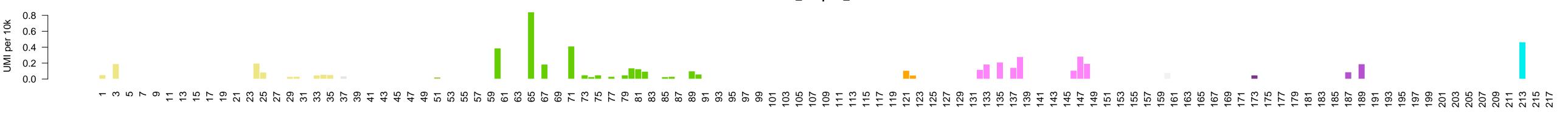
PMCA.HG1.5:AC010616.1/ATP1A1/ATP1A2/ATP1A3/ATP1A4/ATP4A/ATP12A TrH2_TrispH2_011484



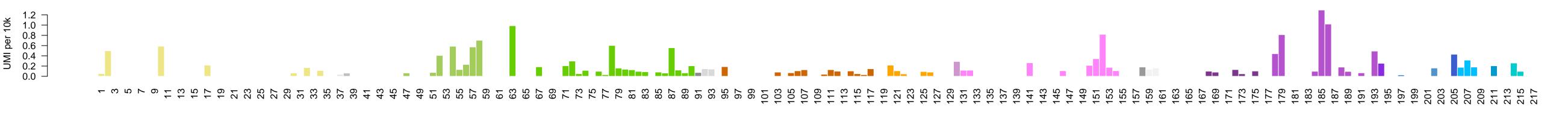
Pyridoxal_deC.HG1.2:PDXDC1 TrH2_TrispH2_002324



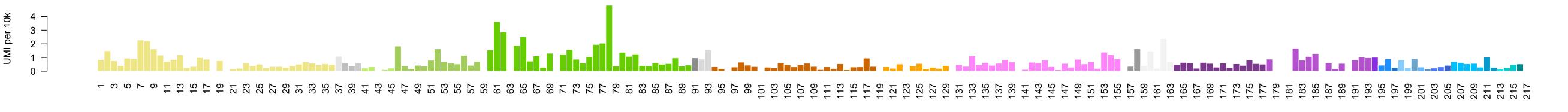
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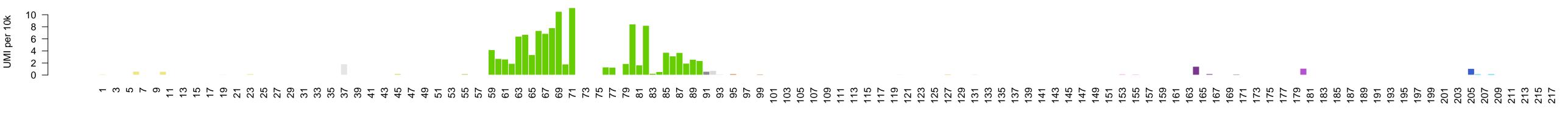
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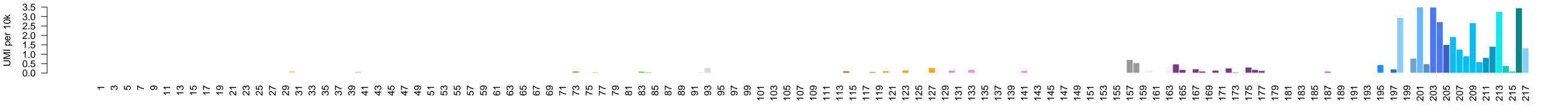


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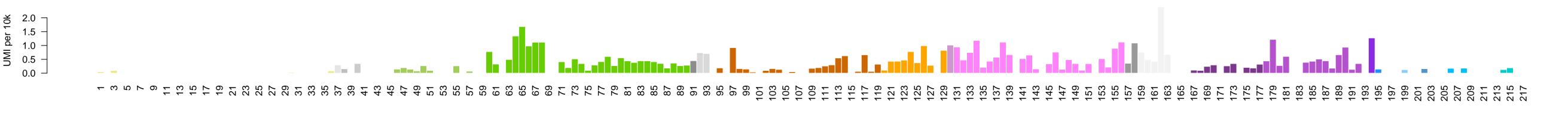


Pyridoxal_deC.HG1.8:DDC/HDC TrH2_TrispH2_007894

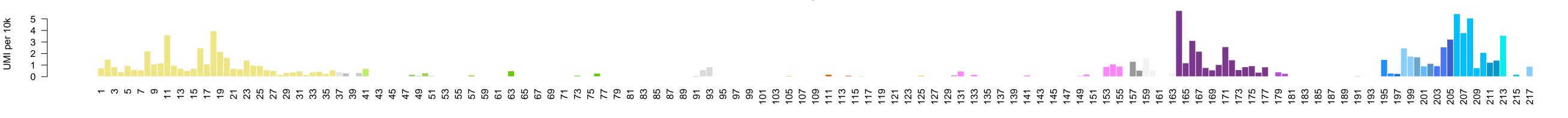




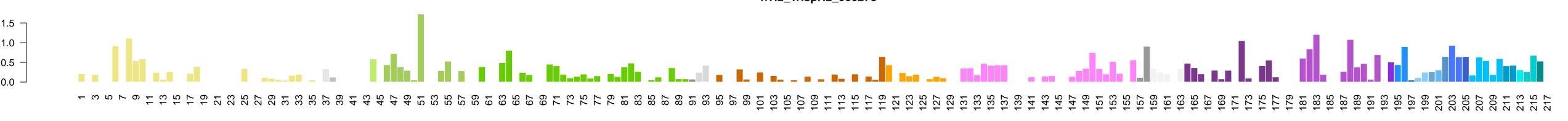
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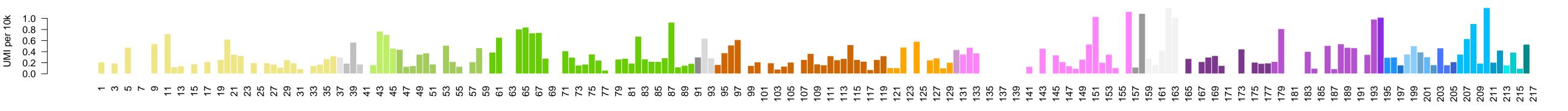
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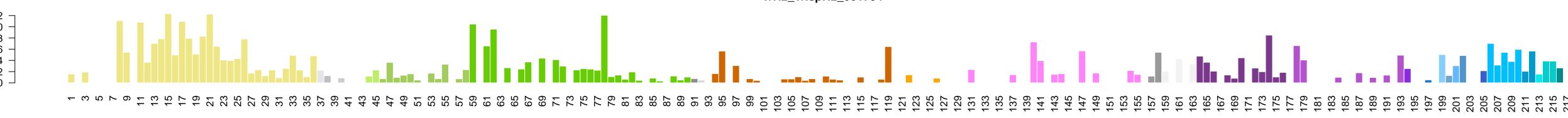
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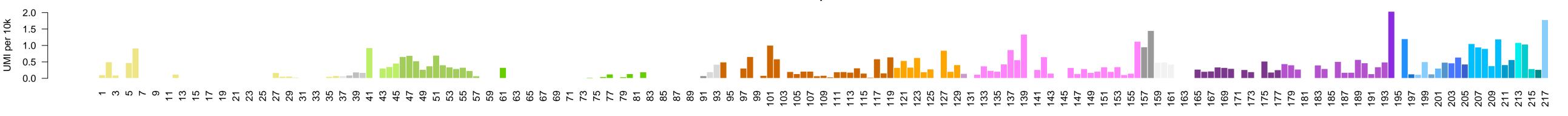
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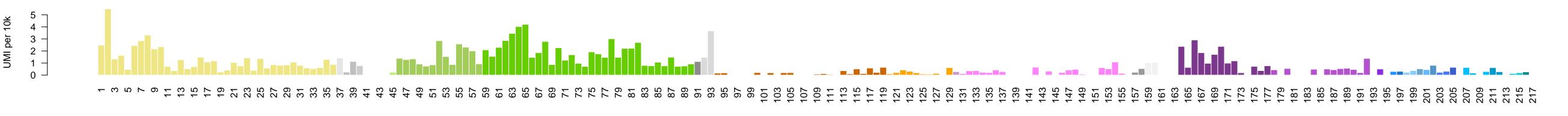
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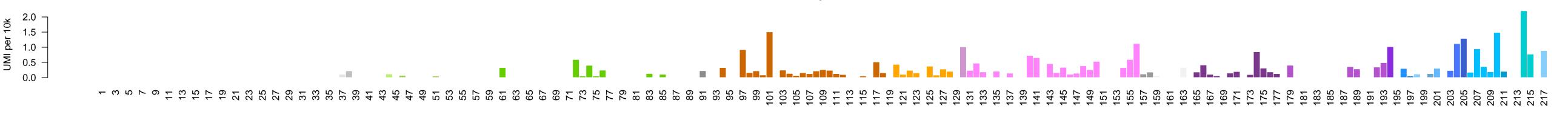
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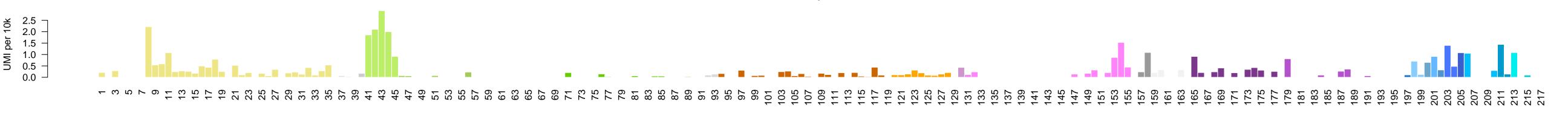
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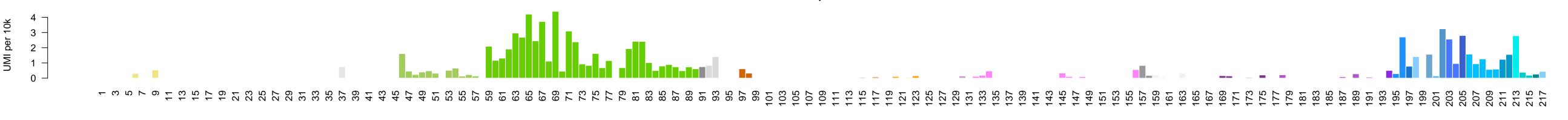
RasGAP.HG1.5:DAB2IP/RASAL2/RASAL3/SYNGAP1;SynGAP.HG1.2:like:DAB2IP/RASAL2/SYNGAP1 TrH2_TrispH2_003065



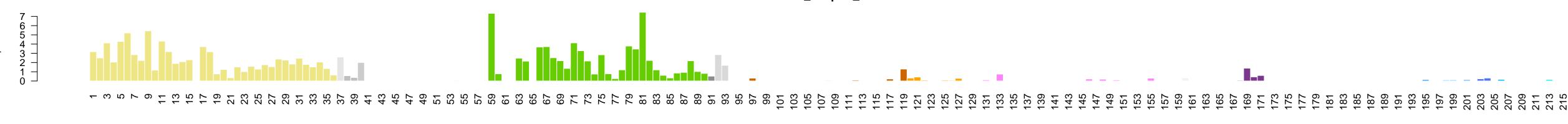
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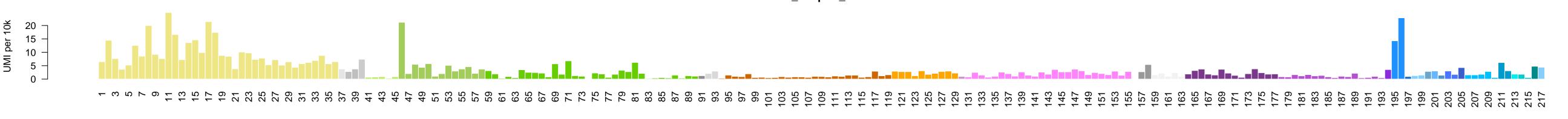
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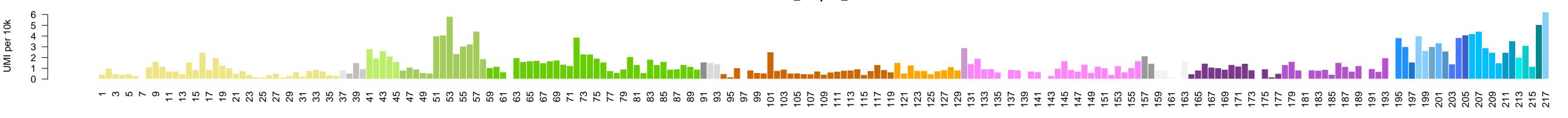
SDF.HG1.1:SLC1A1/SLC1A2/SLC1A3/SLC1A4/SLC1A5/SLC1A6/SLC1A7 TrH2_TrispH2_006787



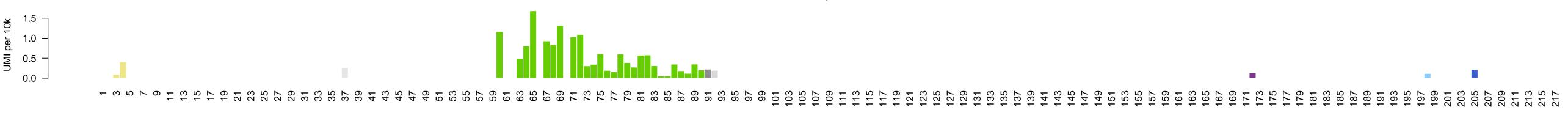
SDF.HG1.1:SLC1A1/SLC1A2/SLC1A3/SLC1A4/SLC1A5/SLC1A6/SLC1A7 TrH2_TrispH2_006789



SNAP-25.HG1.2:SNAP25 TrH2_TrispH2_003129

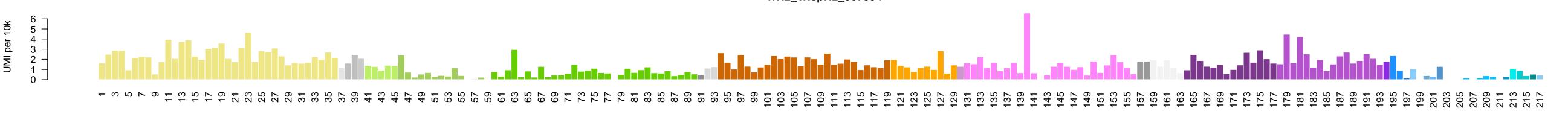


SNAP-25.HG1.2:SNAP25 TrH2_TrispH2_003130



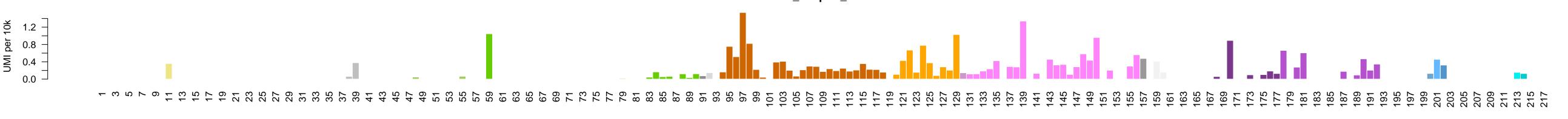
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TrH2_TrispH2_007554



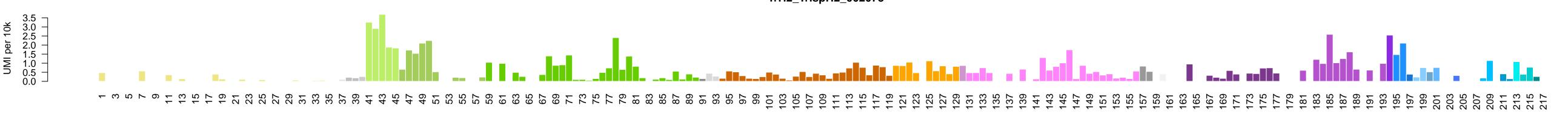
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TrH2_TrispH2_002071



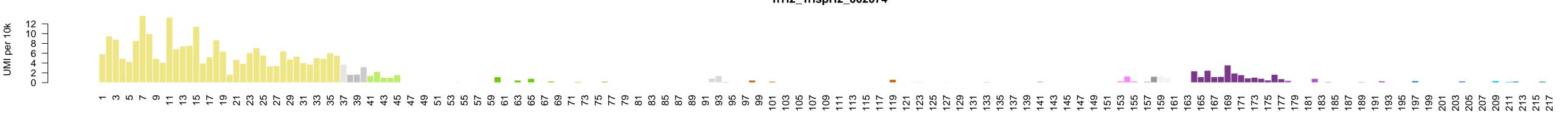
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TrH2_TrispH2_002073



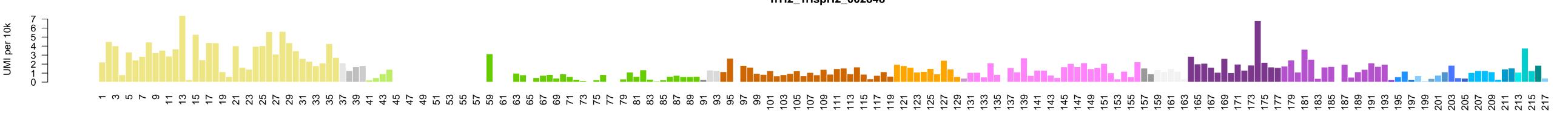
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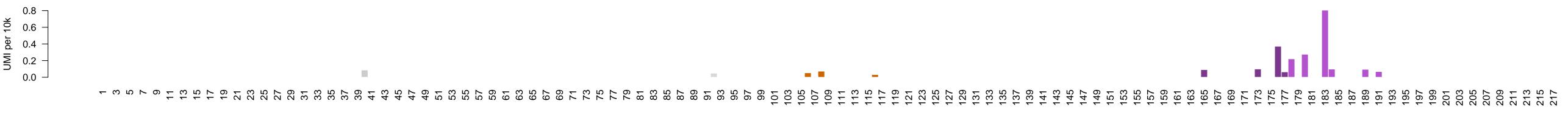
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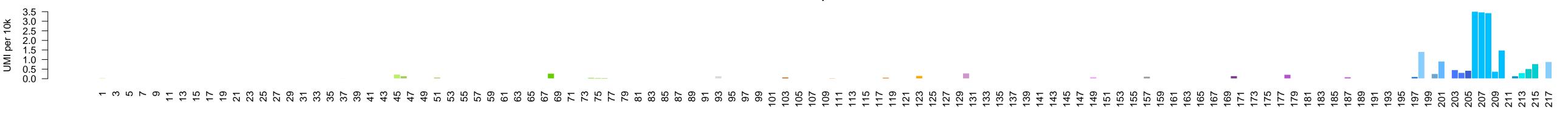


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TrH2_TrispH2_002848

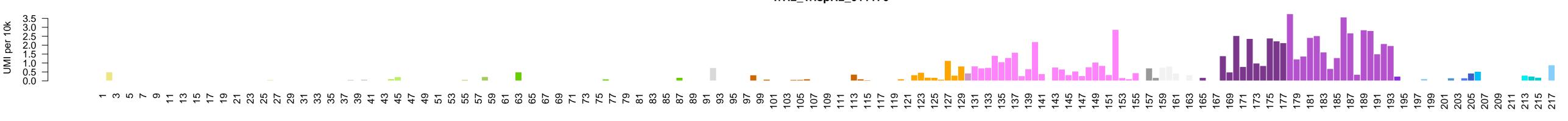






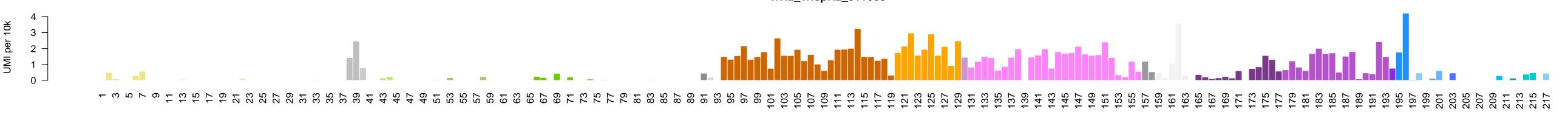
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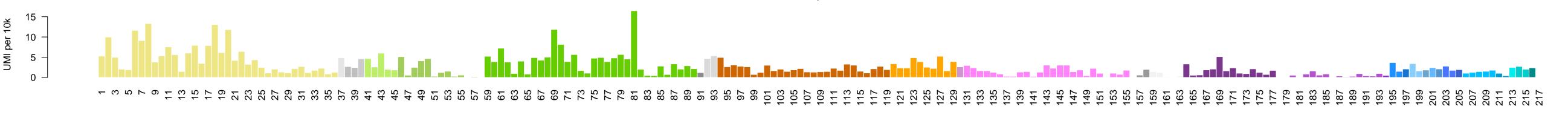


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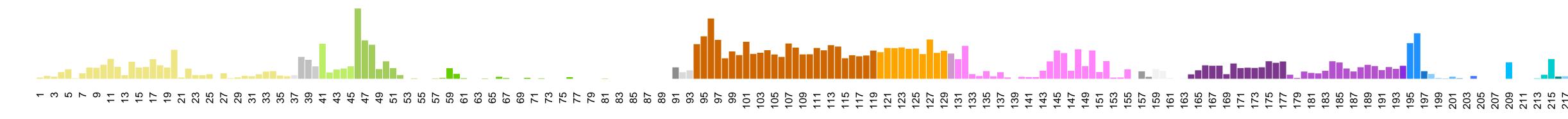
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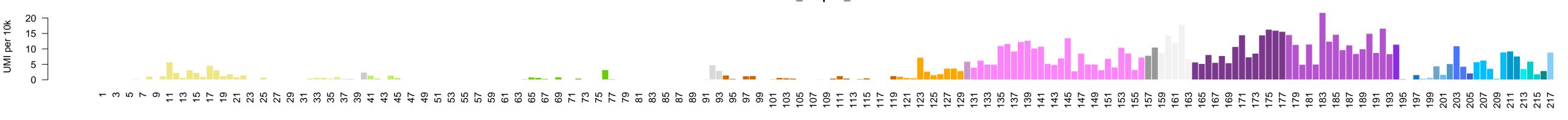
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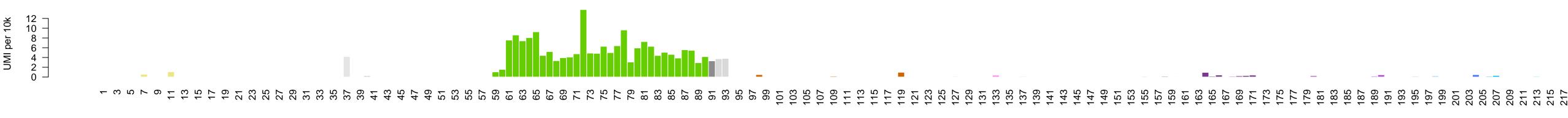
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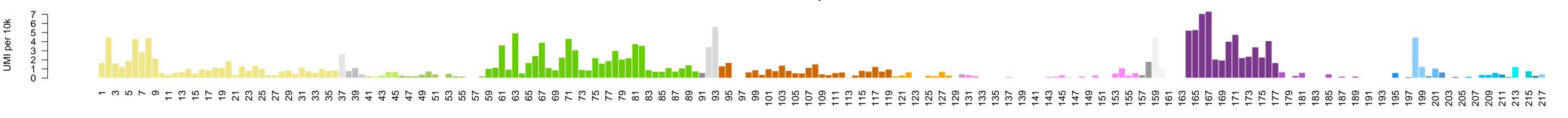
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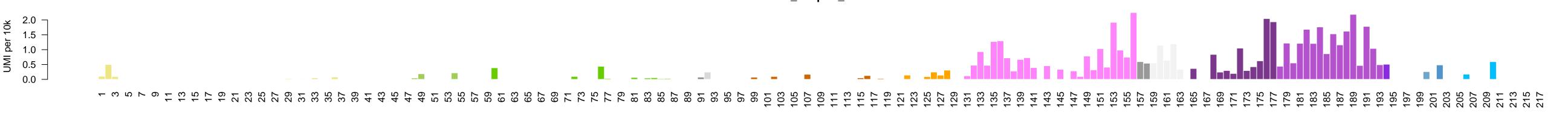
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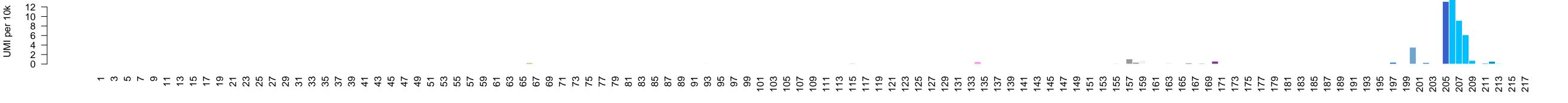
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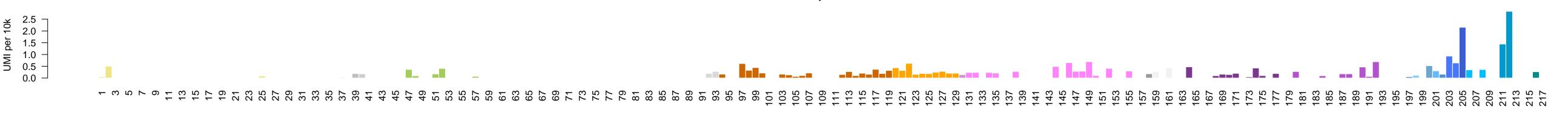
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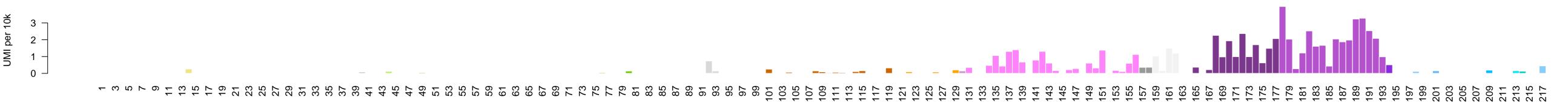


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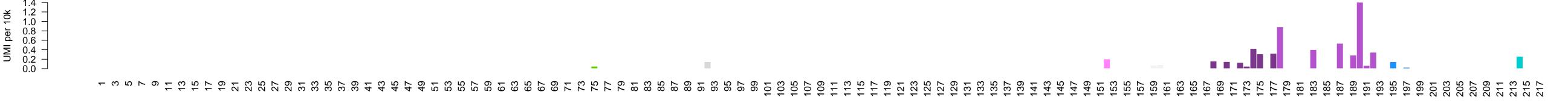
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TrH2_TrispH2_009352

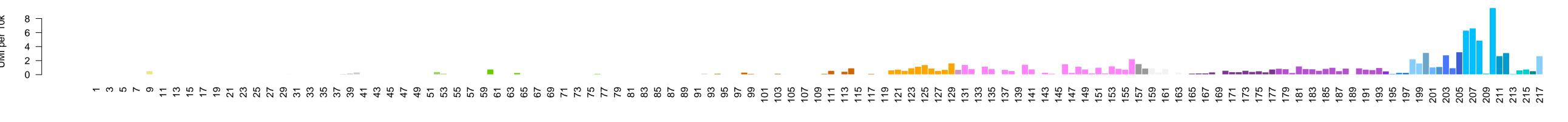




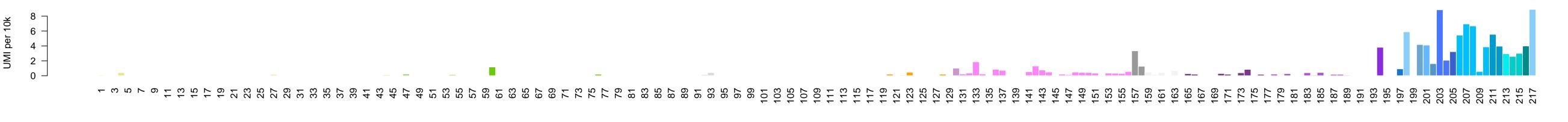
Sugar_tr.HG2.59:SVOP TrH2_TrispH2_010089



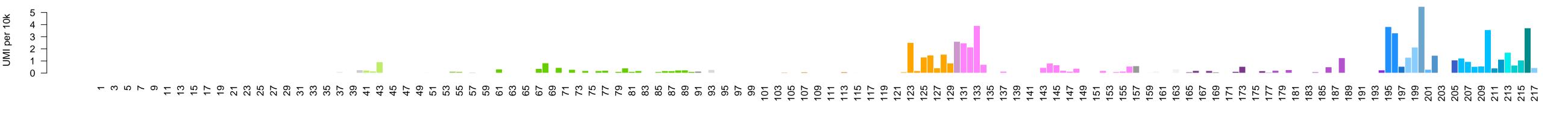
Synaphin.HG2.0:NA TrH2_TrispH2_005012



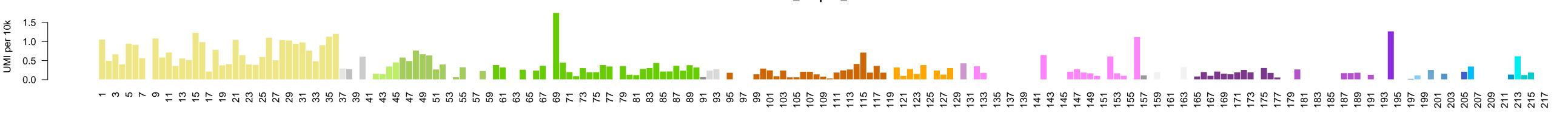
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aptotagmin.HG1.18:FO681492.1/SYT15 TrH2_TrispH2_008612

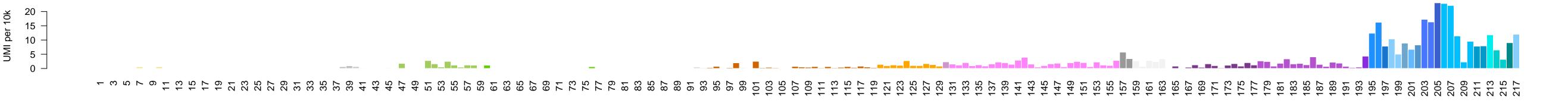


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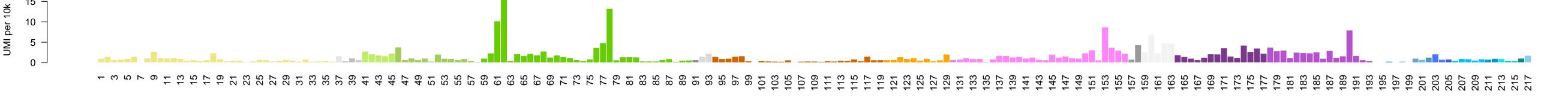


Synaptotagmin.HG1.2:SYT1/SYT2/SYT5/SYT8

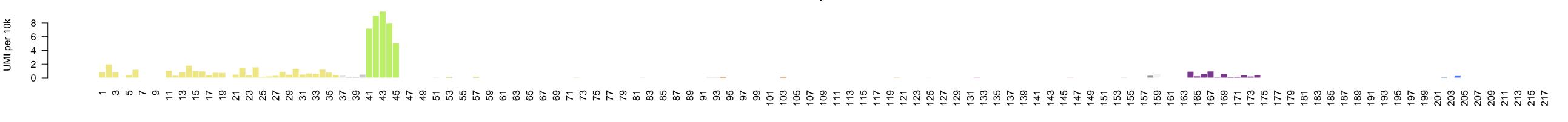
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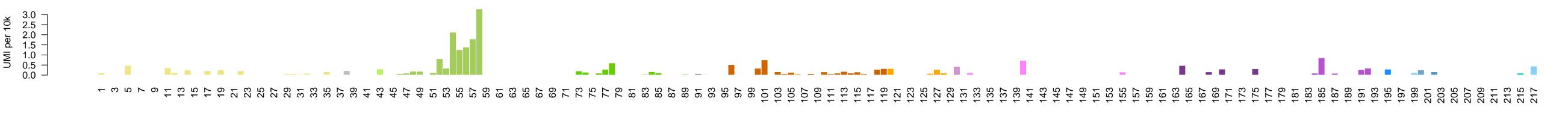
BAIAP3/C2CD5/DOC2A/DOC2B/DYSF/ESYT1/ESYT2/ESYT3/ITSN2/JMJD7-PLA2G4B/MCTP1/MCTP2/MYOF/OTOF/PIK3C2B/PLA2G4B



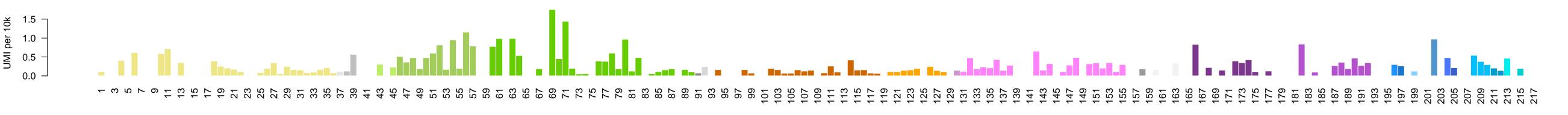
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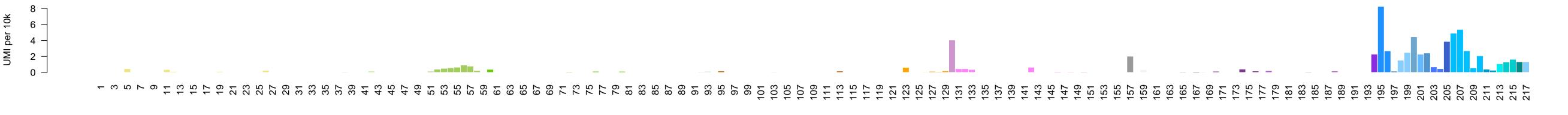
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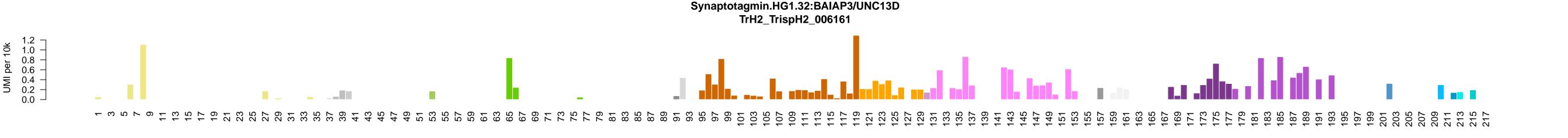


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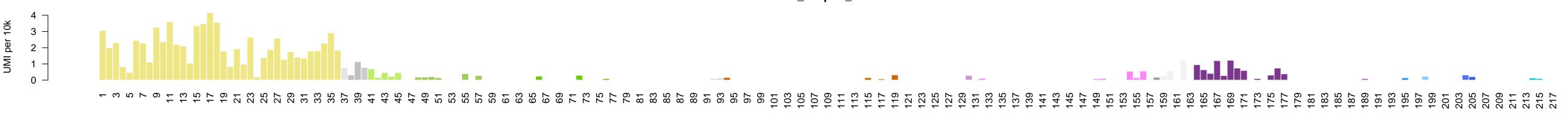


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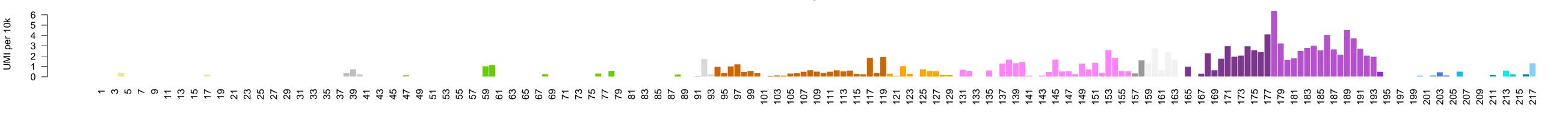




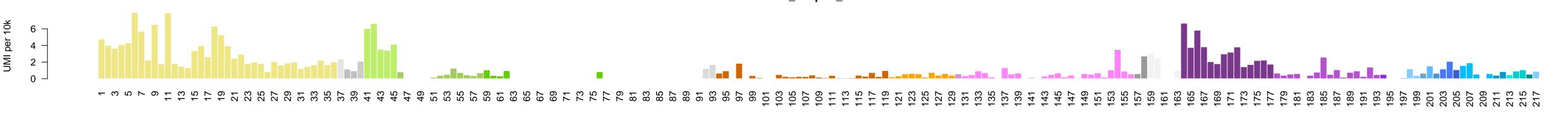
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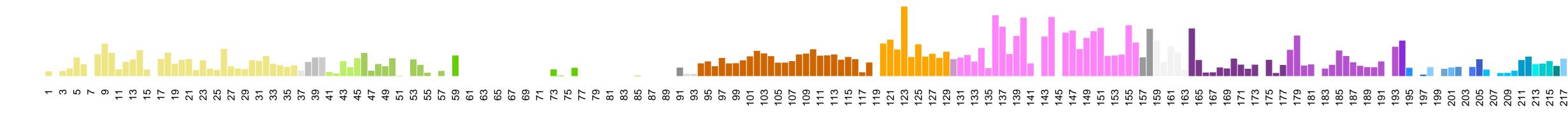
Synaptotagmin.HG1.35:DYSF/MYOF TrH2_TrispH2_005138



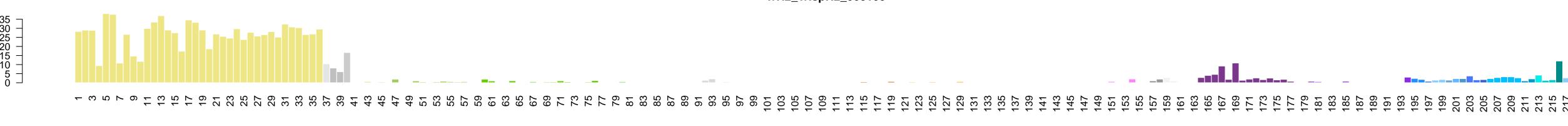
naptotagmin.HG1.42:MCTP1 TrH2_TrispH2_002925



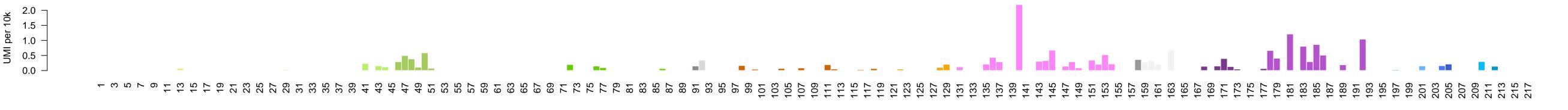
Synaptotagmin.HG1.43:like:ESYT1/ESYT2/ESYT3
TrH2_TrispH2_001163



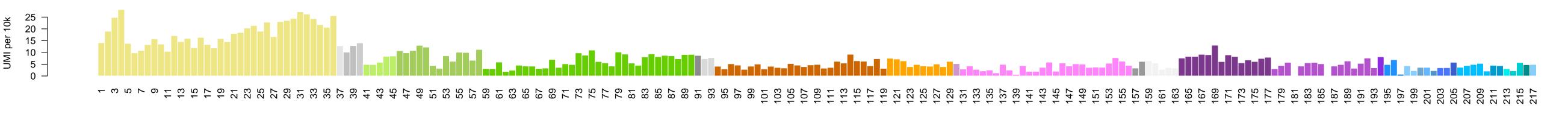
Synaptotagmin.HG1.49:ESYT1/ESYT2/ESYT3 TrH2_TrispH2_008105



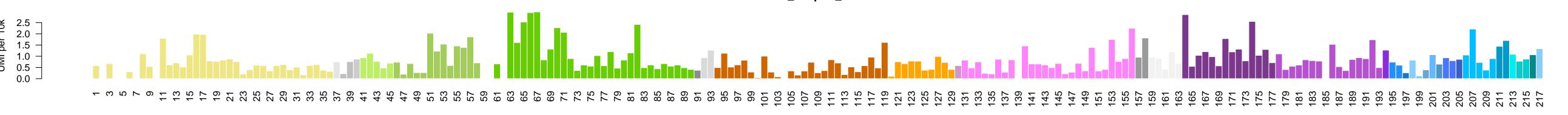
Synaptotagmin.HG1.5:like:FO681492.1/SYT12/SYT13/SYT15 TrH2_TrispH2_003054



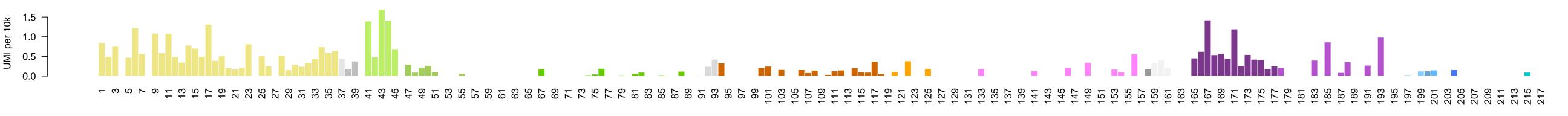
Synaptotagmin.HG1.52:like:MCTP2 TrH2_TrispH2_007539



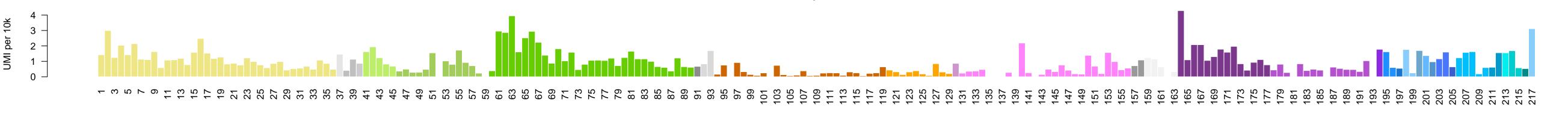
Synaptotagmin.HG1.59:like:ITSN2/UNC13A/UNC13C TrH2_TrispH2_003908



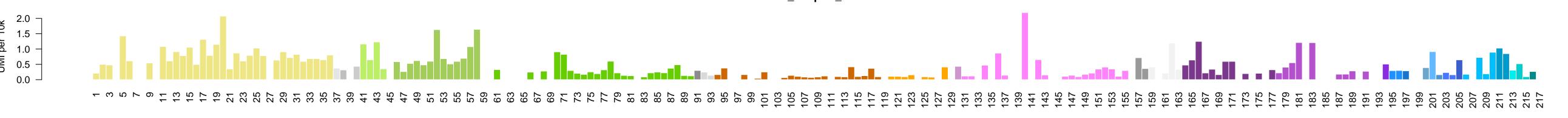
Synaptotagmin.HG1.60:like:ITSN2/UNC13A/UNC13C TrH2_TrispH2_006333



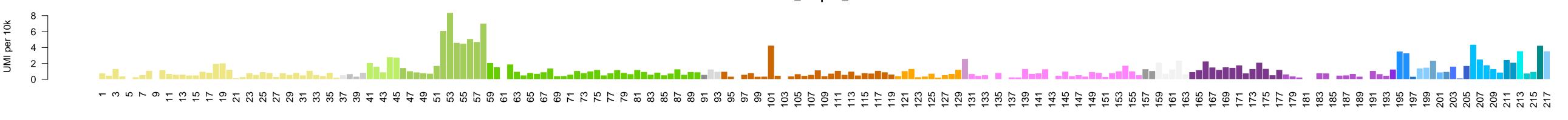
Synaptotagmin.HG1.62:PRKCA/PRKCB/PRKCG TrH2_TrispH2_007855



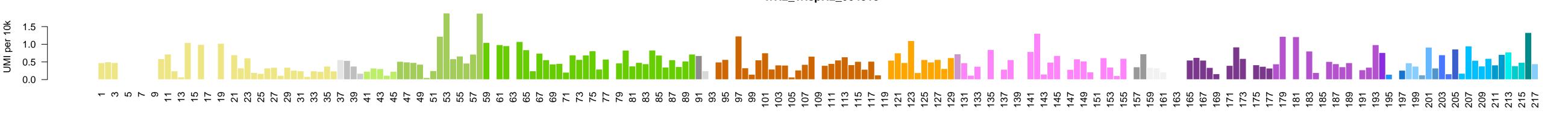
Syntaxin.HG1.1:STX16/STX16-NPEPL1 TrH2_TrispH2_000160



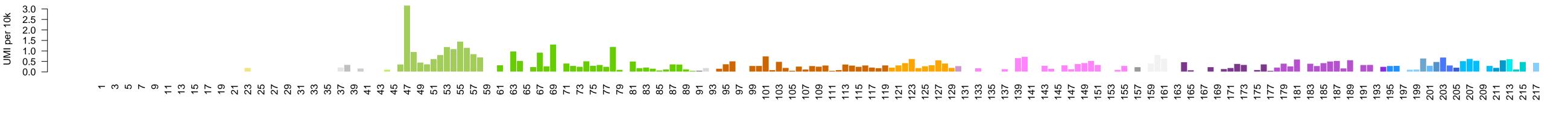
Syntaxin.HG1.2:STX7/STX12 TrH2_TrispH2_000530



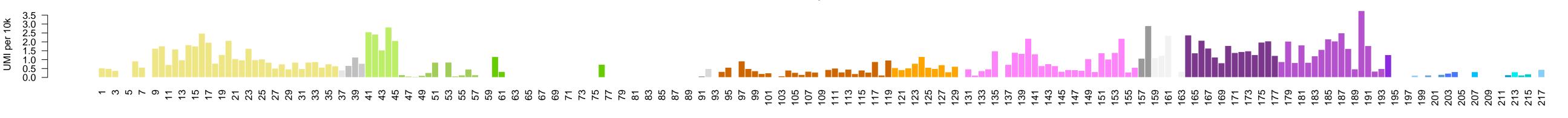
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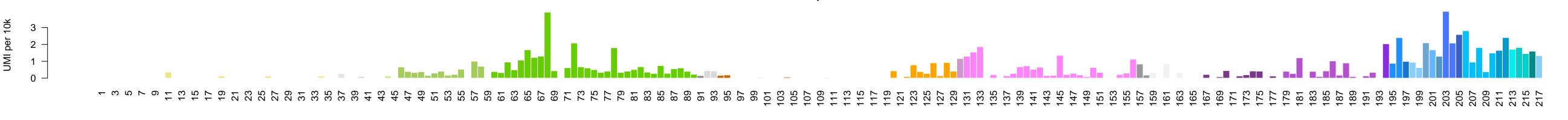
Syntaxin.HG1.4:STX1A/STX1B/STX2/STX3/STX4 TrH2 TrispH2 004549



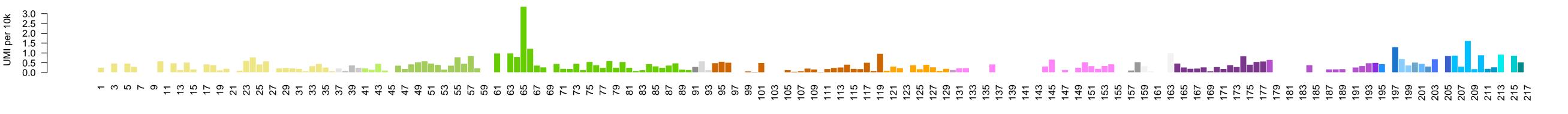
Syntaxin.HG1.4:STX1A/STX1B/STX2/STX3/STX4
TrH2_TrispH2_006861



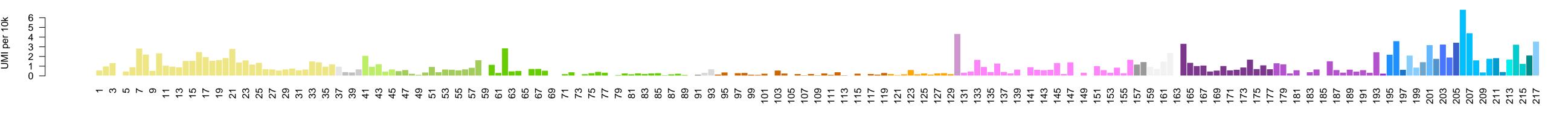
ntaxin.HG1.4:STX1A/STX1B/STX2/STX3/STX4



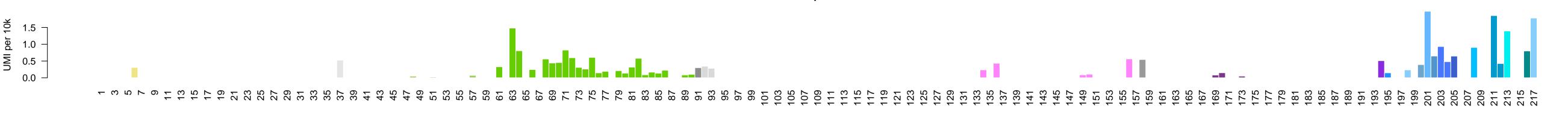
Y_phosphatase.HG1.13:PTPN6/PTPN11 TrH2 TrispH2 005444



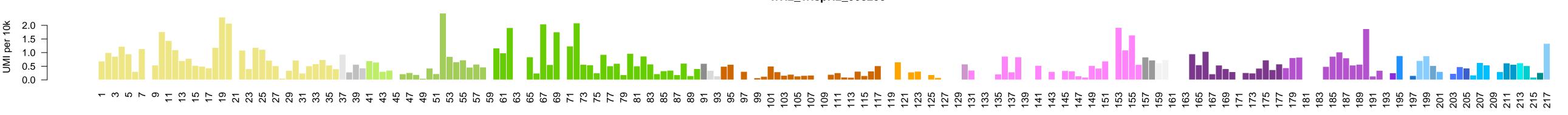
Y_phosphatase.HG1.14:PTPN5/PTPN7/PTPRR TrH2_TrispH2_005229



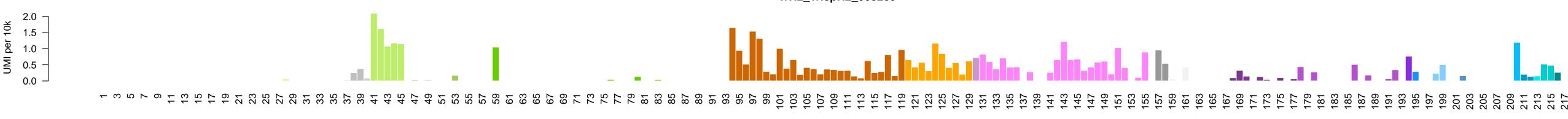
Y_phosphatase.HG1.16:like:PTPRK/PTPRM/PTPRT/PTPRU TrH2_TrispH2_001036



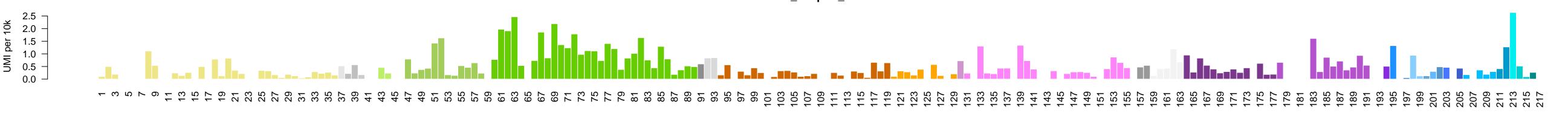
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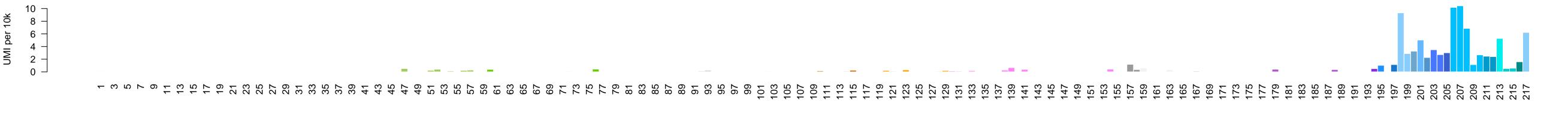
Y_phosphatase.HG1.24:like:PTPRA/PTPRC/PTPRD/PTPRE/PTPRF/PTPRG/PTPRS/PTPRZ1 TrH2_TrispH2_005289



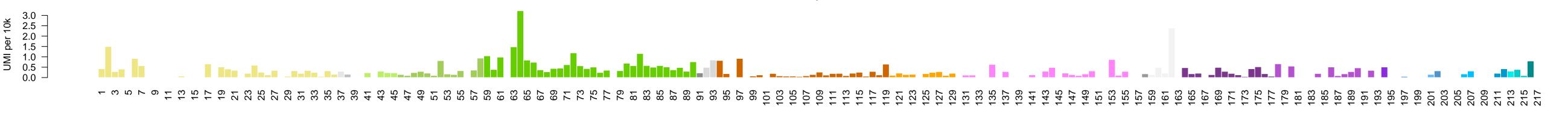
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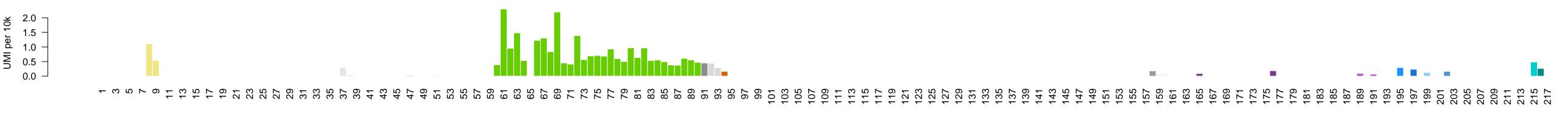
Y_phosphatase.HG1.29:PTPRN/PTPRN2 TrH2_TrispH2_005739



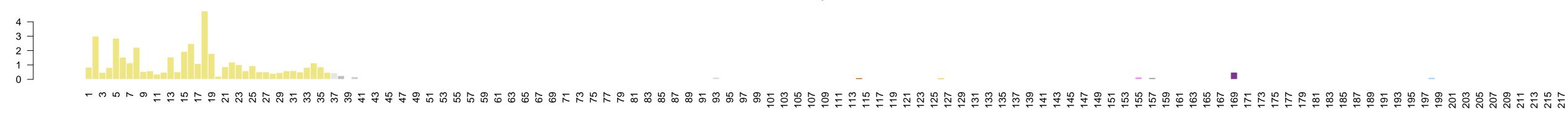
Y_phosphatase.HG1.30:PTPN1/PTPN2 TrH2_TrispH2_000150



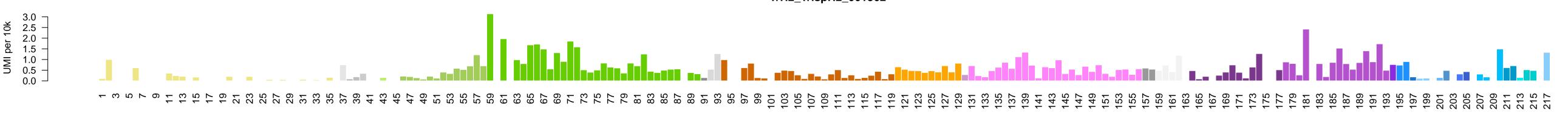
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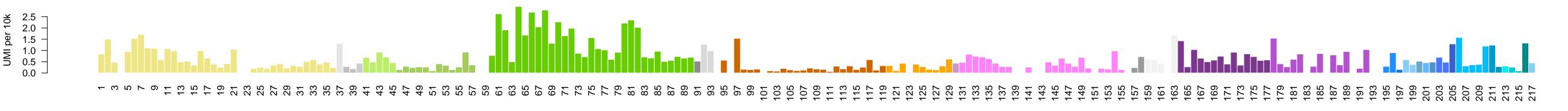
Y_phosphatase.HG1.41:like:PTPN5/PTPN7/PTPRR TrH2_TrispH2_002994



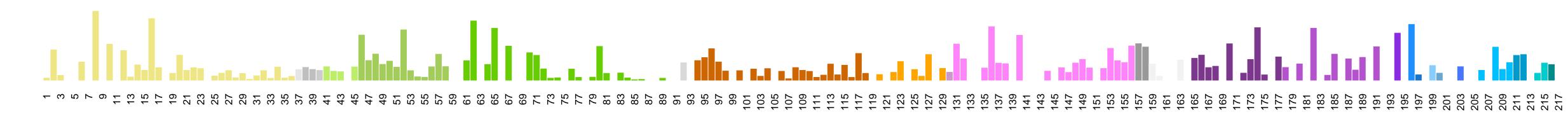
Y_phosphatase.HG1.45:like:PTPRA/PTPRC/PTPRD/PTPRE/PTPRF/PTPRG/PTPRS/PTPRZ1 TrH2_TrispH2_001362



Y_phosphatase.HG1.52:PTPRG/PTPRZ1 TrH2_TrispH2_000073

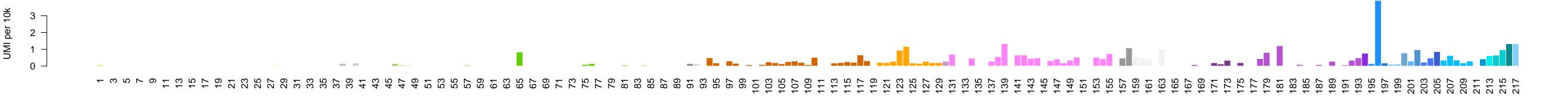


Y_phosphatase.HG2.3:like:CDC14A/CDC14B/CDC14C/DUSP23/PTP4A1/PTP4A2/PTP4A3/PTPDC1 TrH2_TrispH2_001234

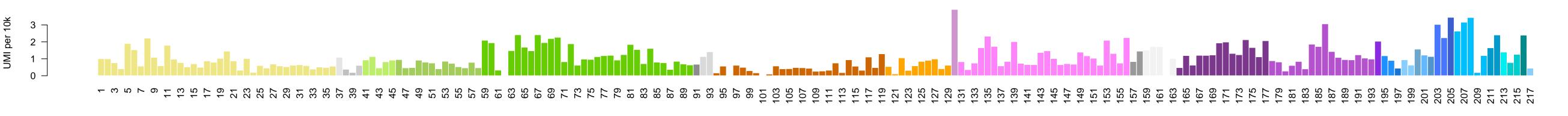


1.0 - 0.8 - 0.6 - 0.4 - 0.2 -

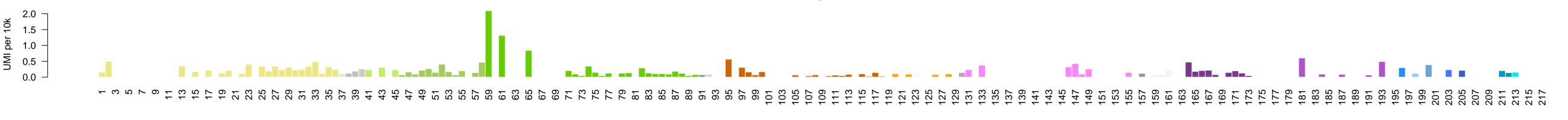
Y_phosphatase.HG2.6:PTPDC1 TrH2_TrispH2_004243



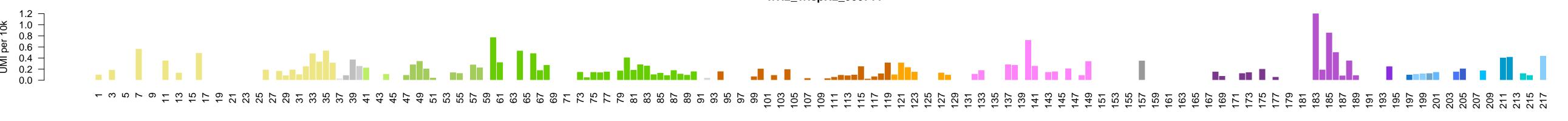
Y_phosphatase.HG2.8:PTP4A1/PTP4A2/PTP4A3 TrH2_TrispH2_010534



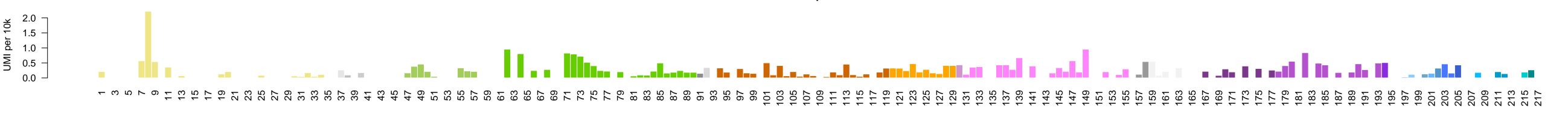
Y_phosphatase.HG3.1:like:DUSP1/DUSP4/DUSP5 TrH2_TrispH2_004372



/_phosphatase.HG3.4:like:DUSP1/DUSP4/DUSP5 TrH2_TrispH2_000711



Y_phosphatase.HG3.4:like:DUSP1/DUSP4/DUSP5 TrH2 TrispH2 000712



TrH2_TrispH2_005304

Y_phosphatase.HG6.0:NA