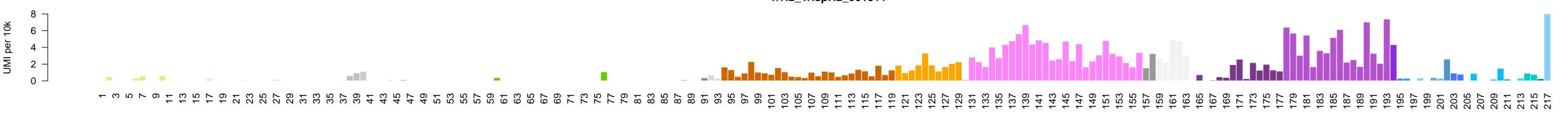
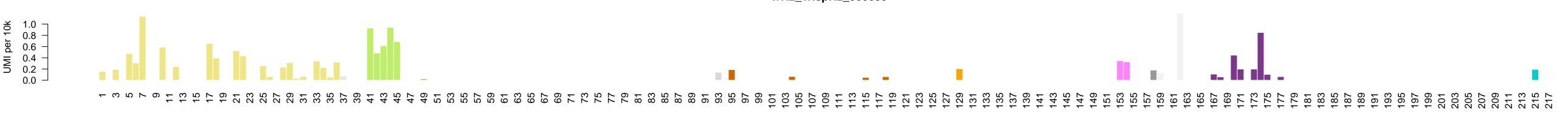
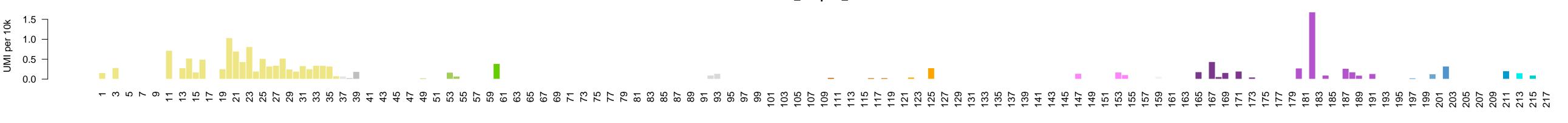
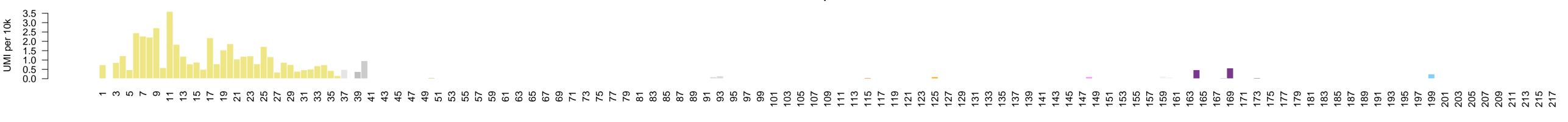
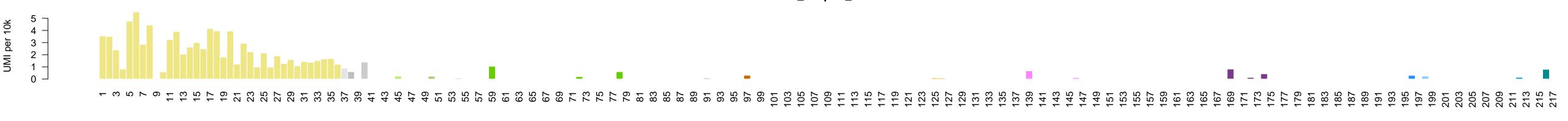
ASC.HG1.18:SCNN1A/SCNN1B/SCNN1D/SCNN1G TrH2_TrispH2_001311

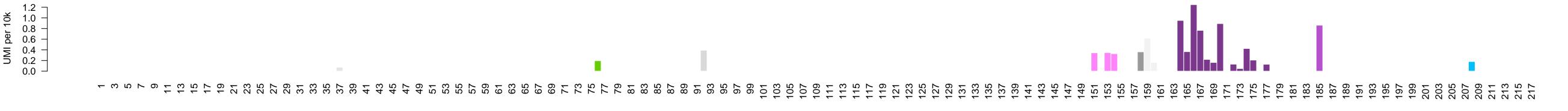




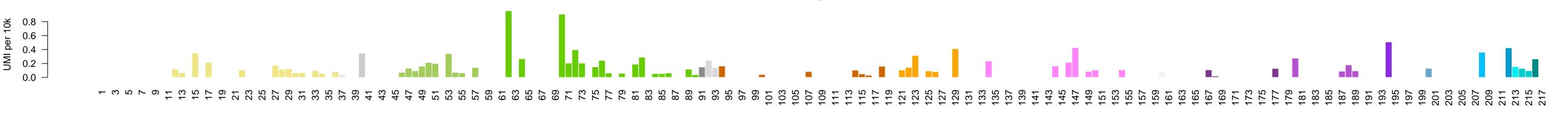




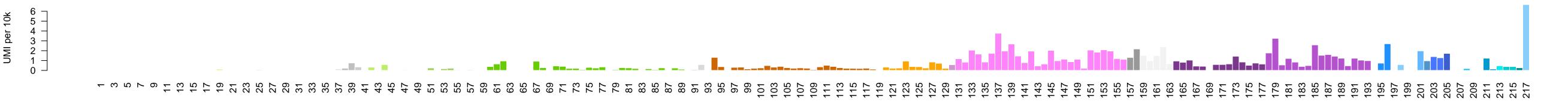


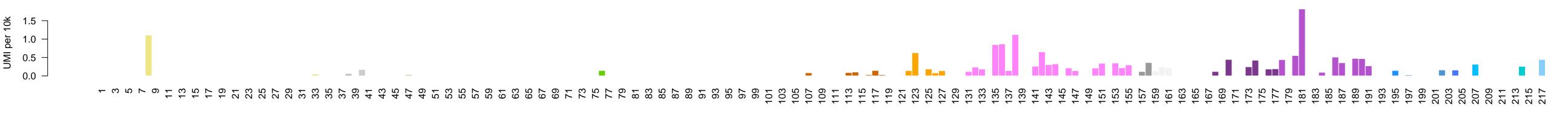


ASC.HG1.9:ASIC1/ASIC2/ASIC3/ASIC4/ASIC5
TrH2_TrispH2_000613

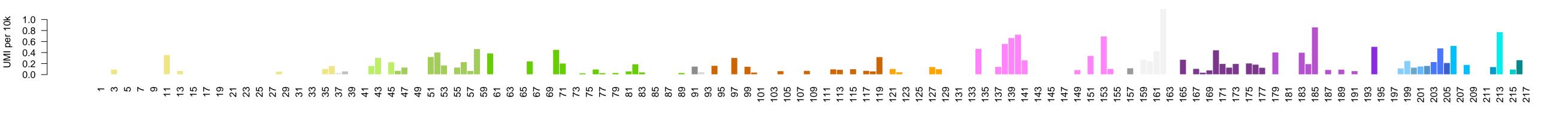


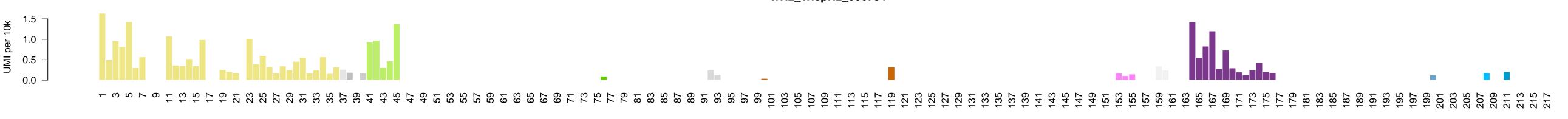
ASC.HG1.9:ASIC1/ASIC2/ASIC3/ASIC4/ASIC5
TrH2_TrispH2_001388



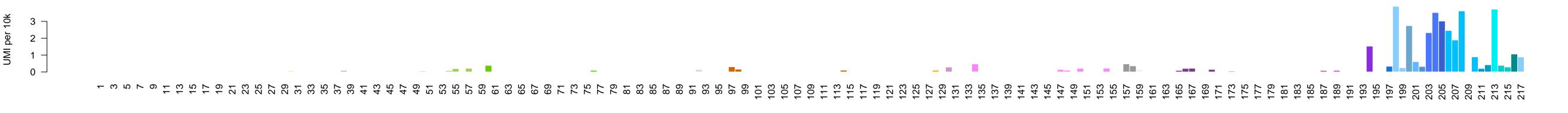


ASC.HG1.9:ASIC1/ASIC2/ASIC3/ASIC4/ASIC5
TrH2_TrispH2_001525

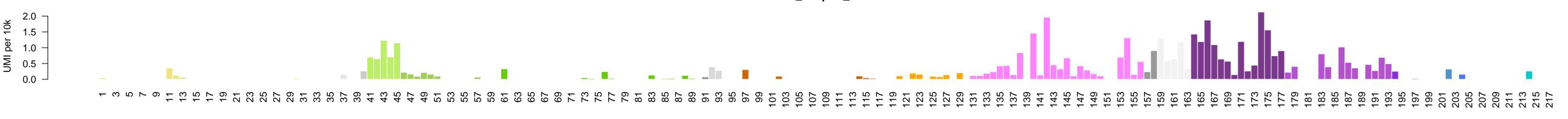




Ion_trans.HG1.1:KCNB1/KCNB2/KCNF1/KCNG1/KCNG2/KCNG3/KCNG4/KCNS1/KCNS2/KCNS3/KCNV1/KCNV2 TrH2_TrispH2_001435



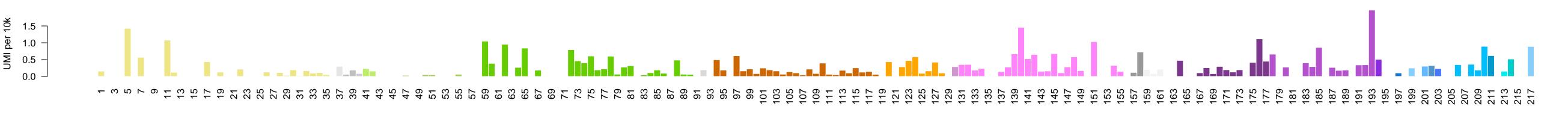
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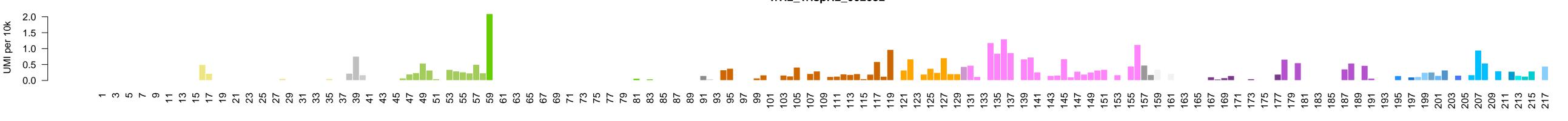
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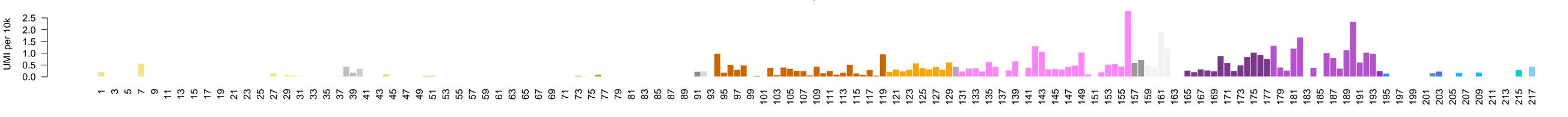
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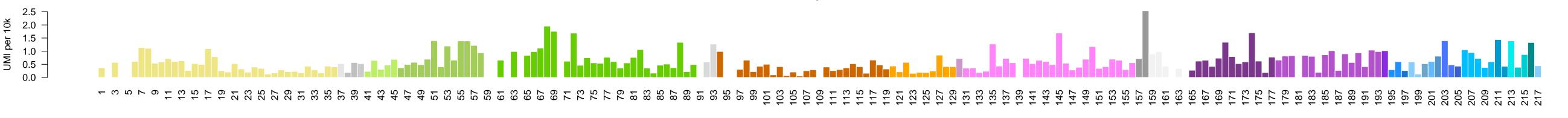
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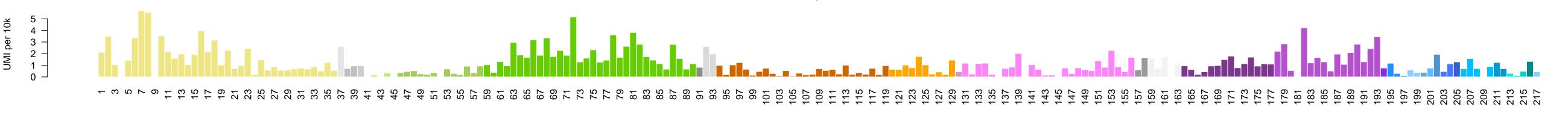
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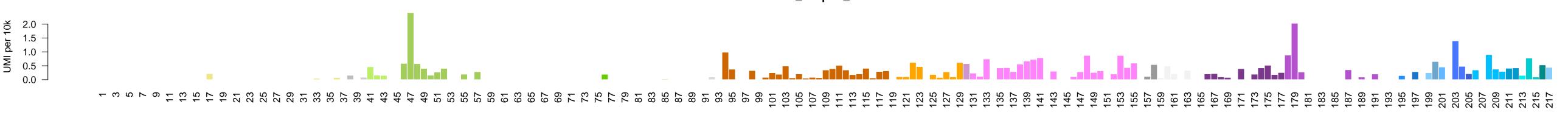
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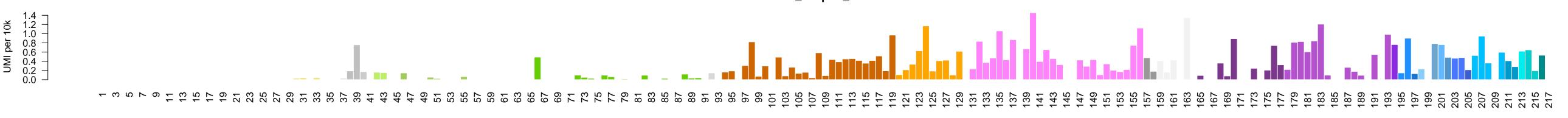
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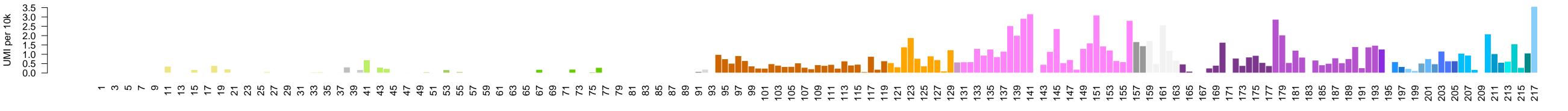
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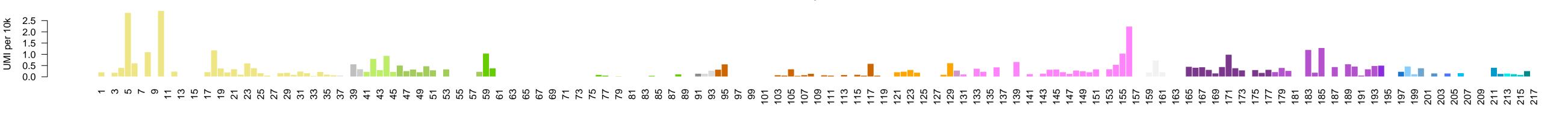
TrH2_TrispH2_002895

Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4

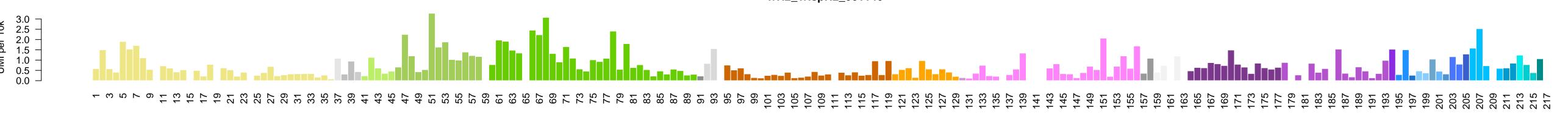
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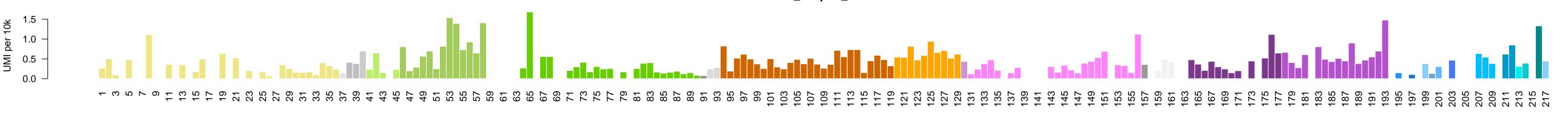
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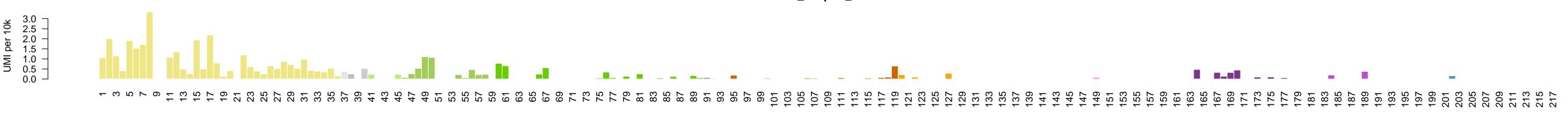
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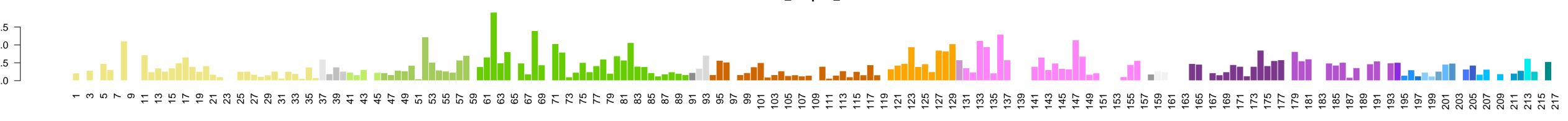
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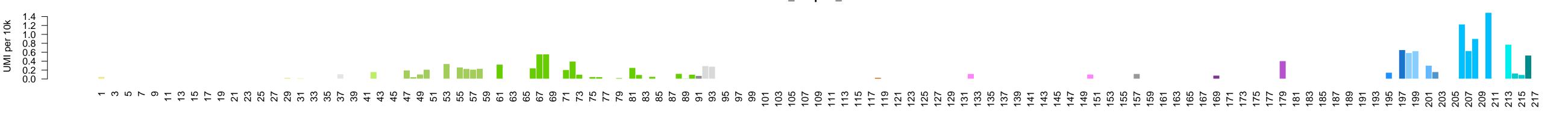
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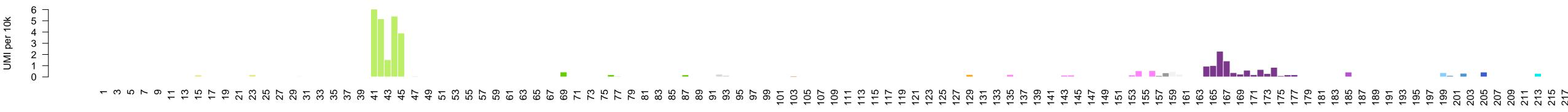


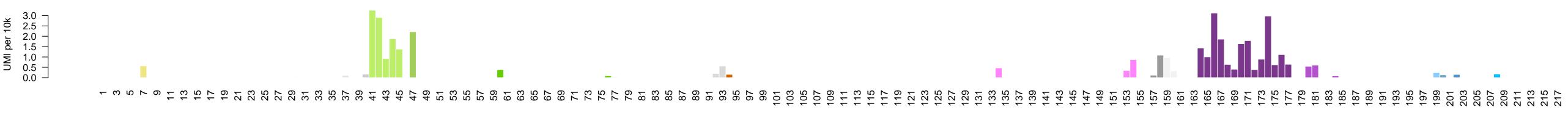
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TrH2_TrispH2_001401

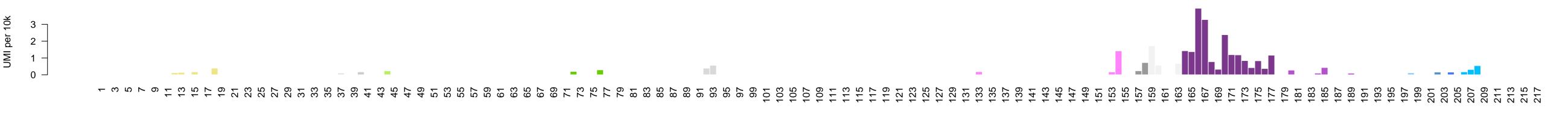


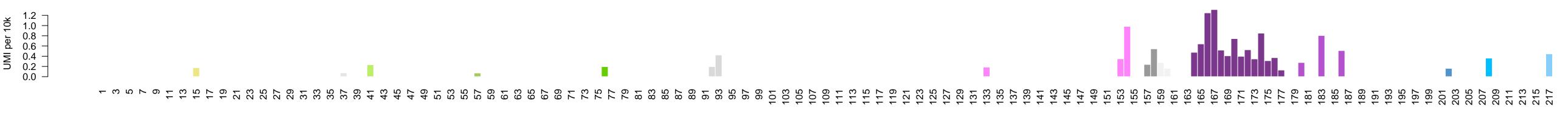
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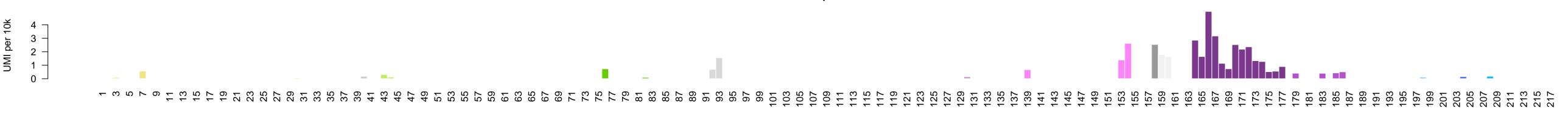


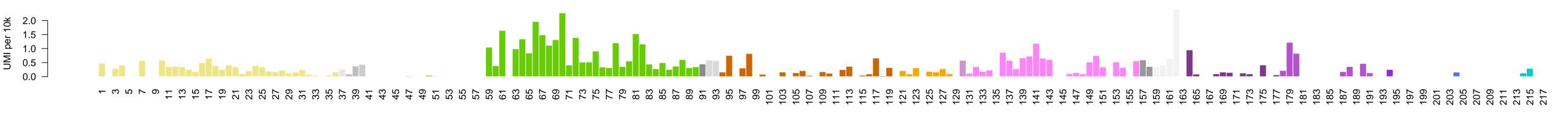


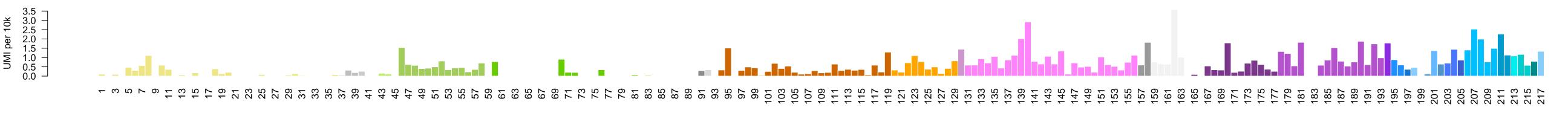




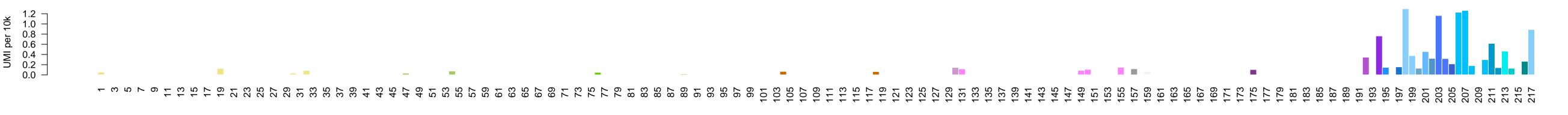




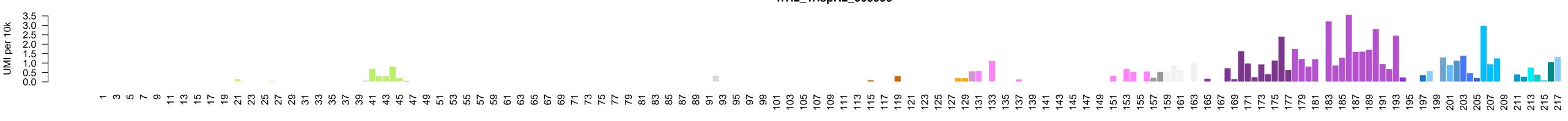


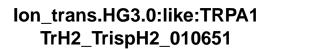


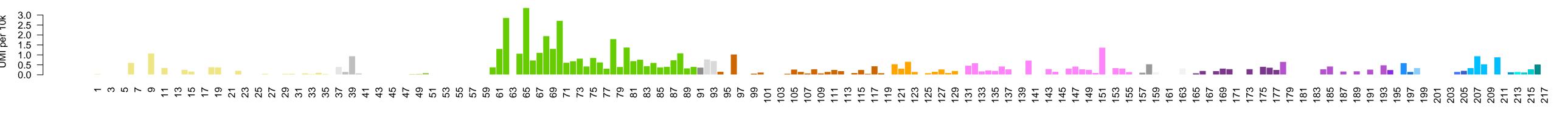
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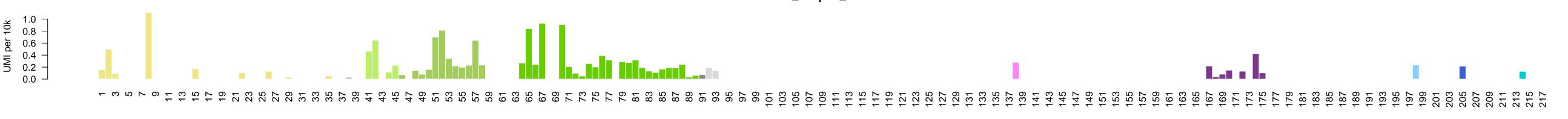
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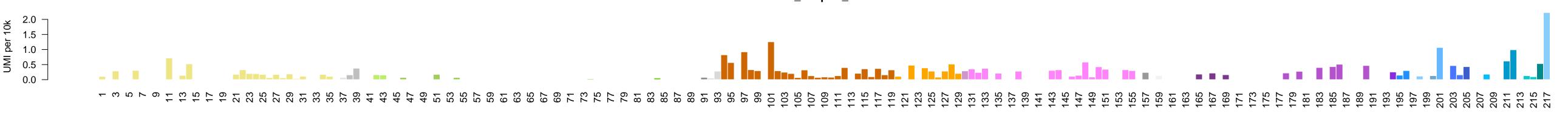


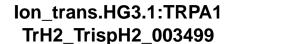


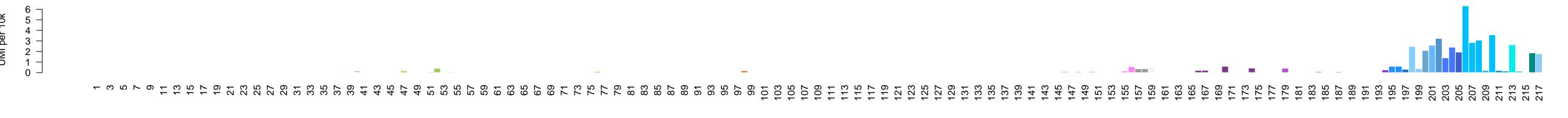
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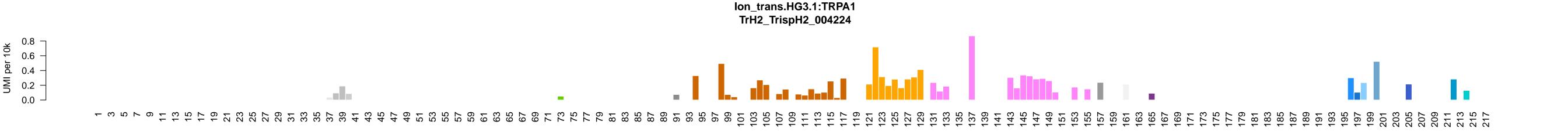


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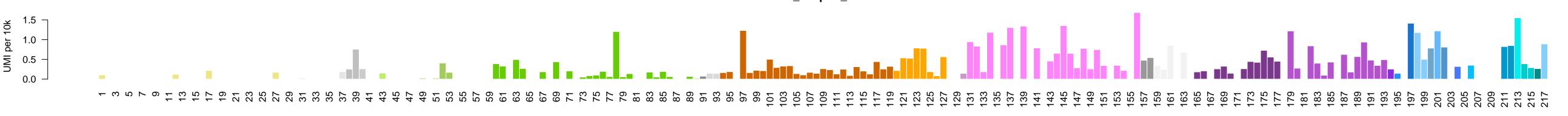




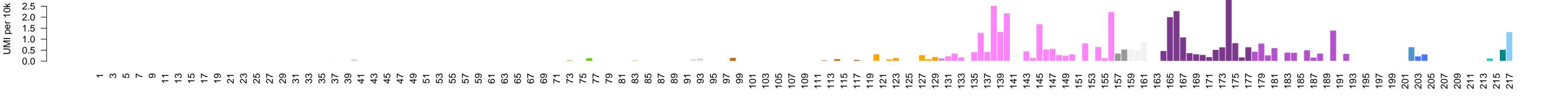


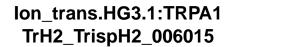


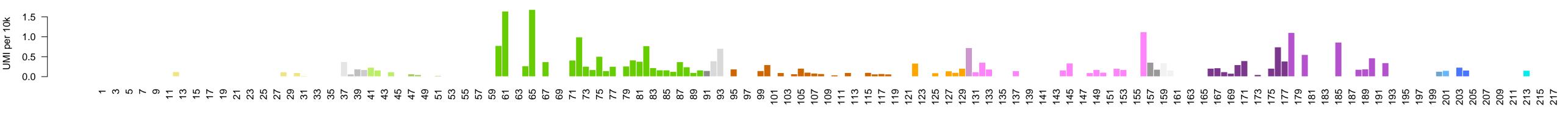
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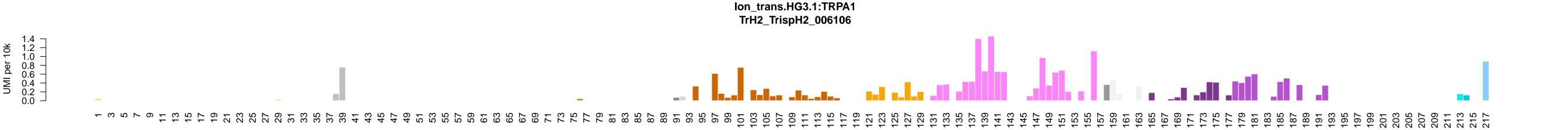


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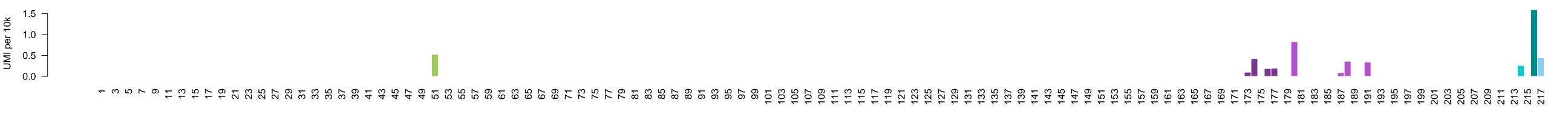




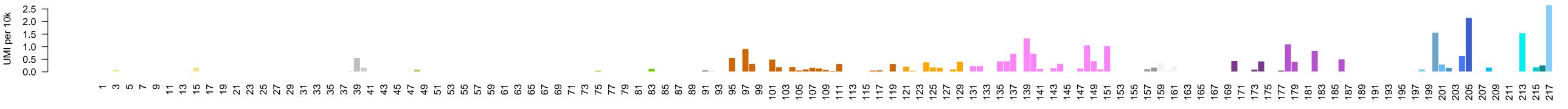




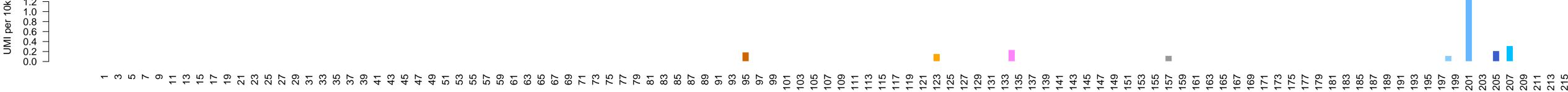
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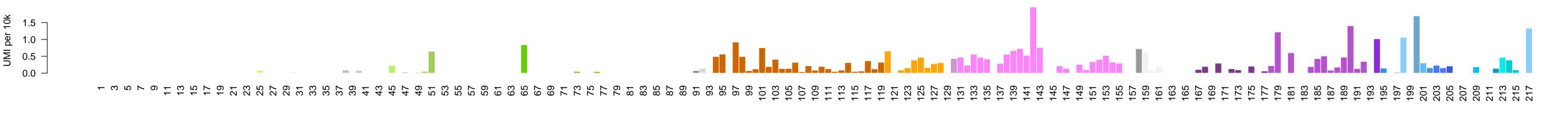




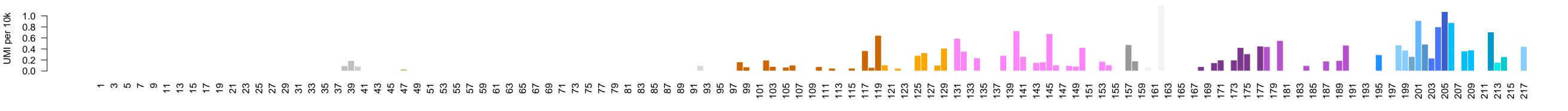




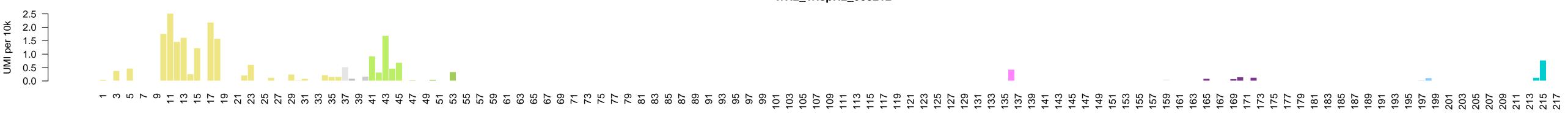
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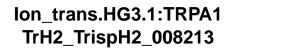


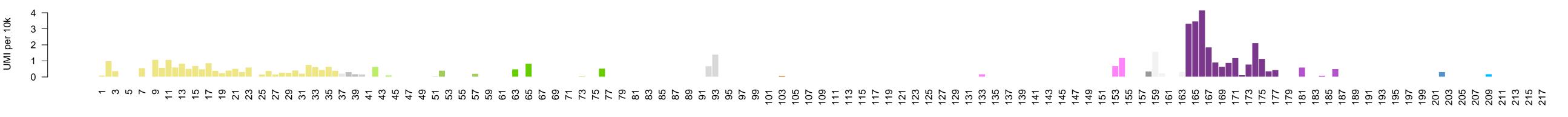
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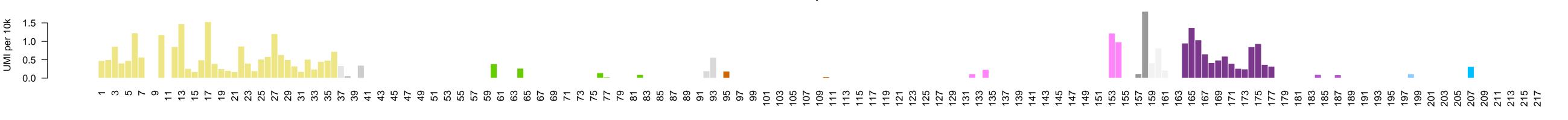
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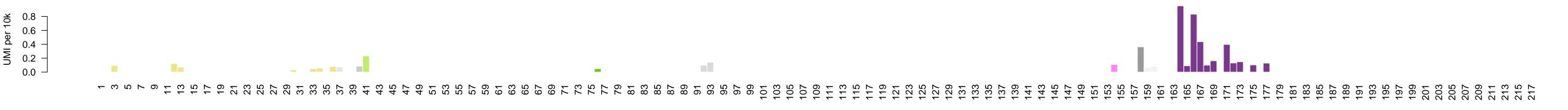




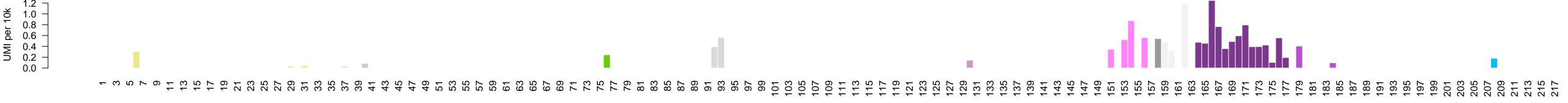
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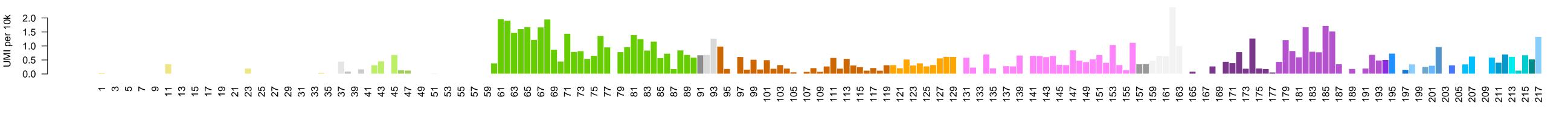
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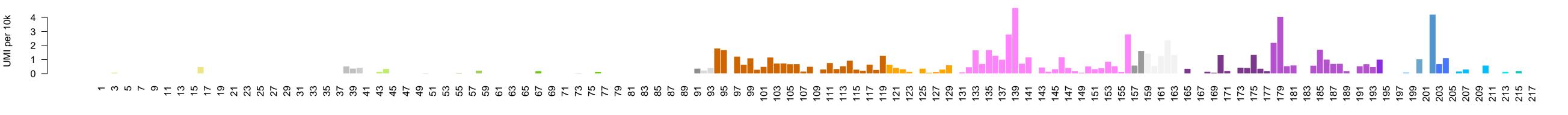




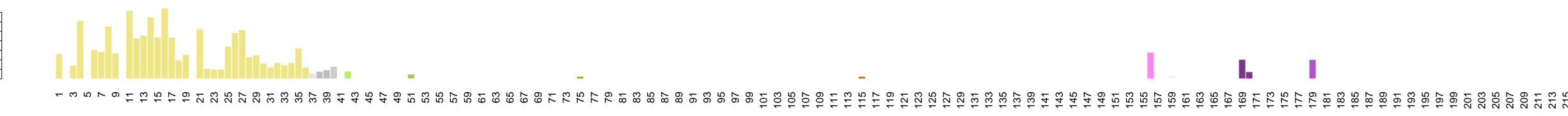
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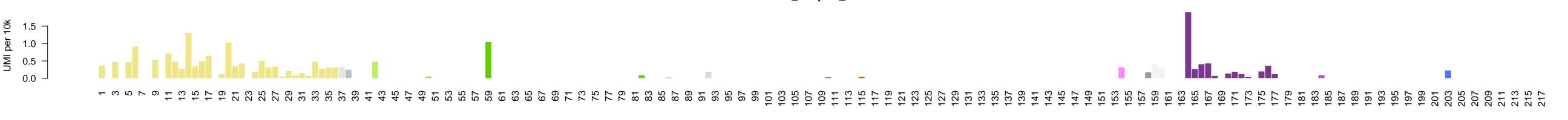
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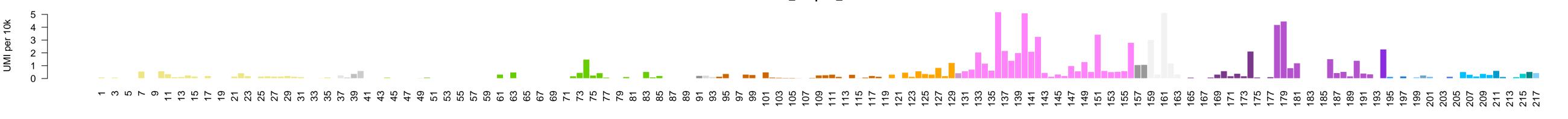
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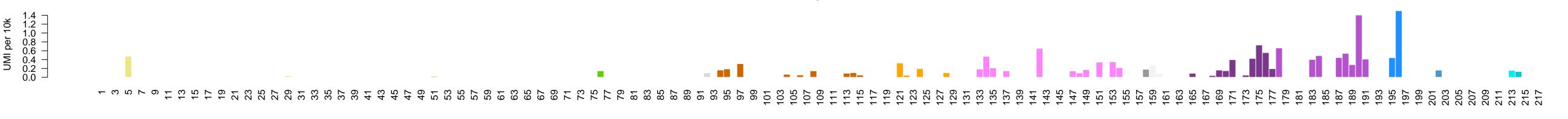
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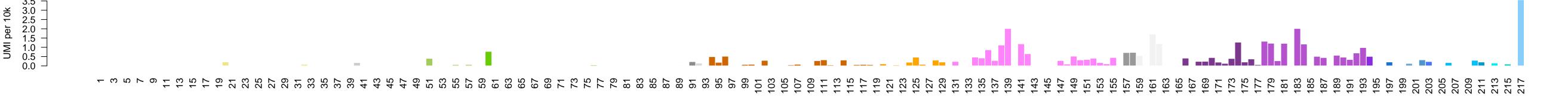
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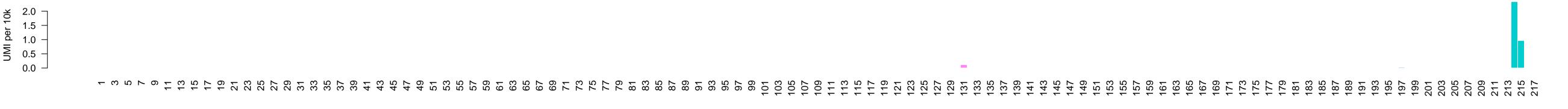
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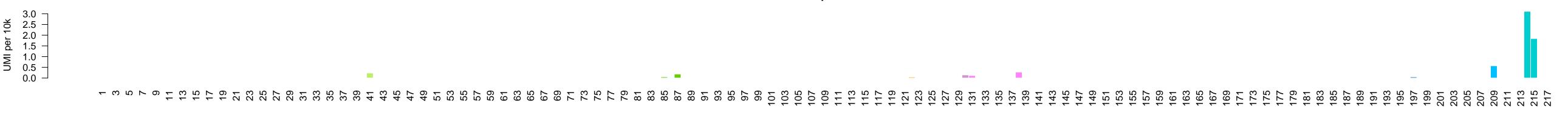
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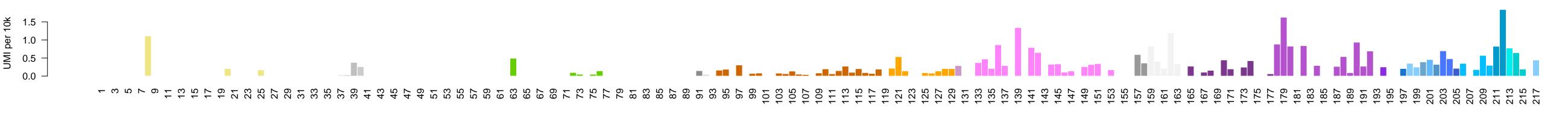
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Ion_trans.HG5.22:PKD2/PKD2L1/PKD2L2;PKD_channel.HG1.3:PKD2/PKD2L1/PKD2L2 TrH2_TrispH2_002072

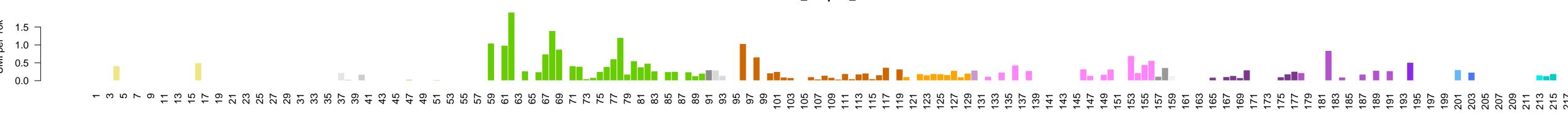


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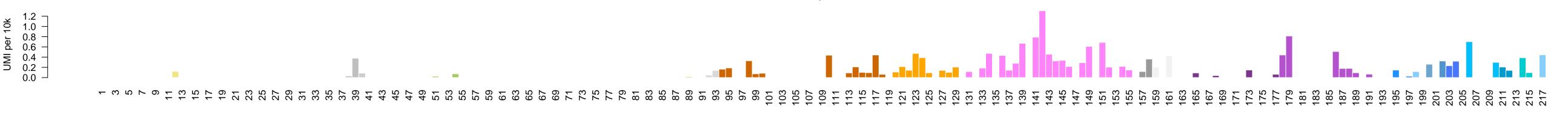


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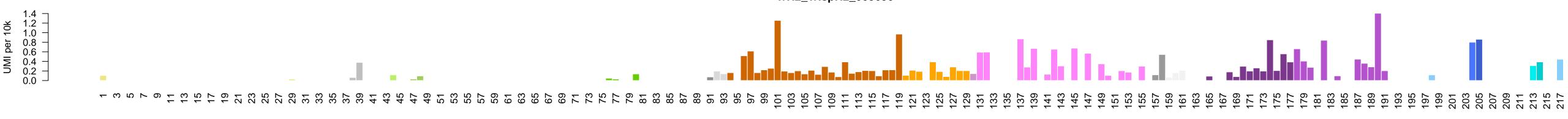
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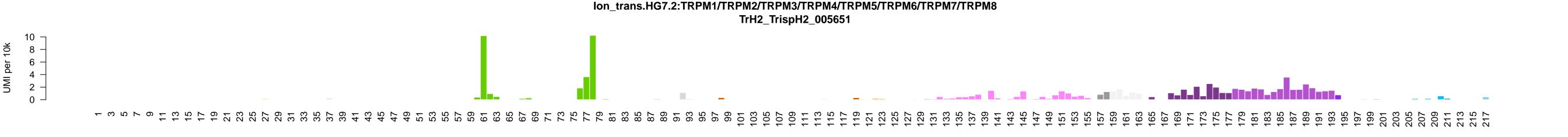


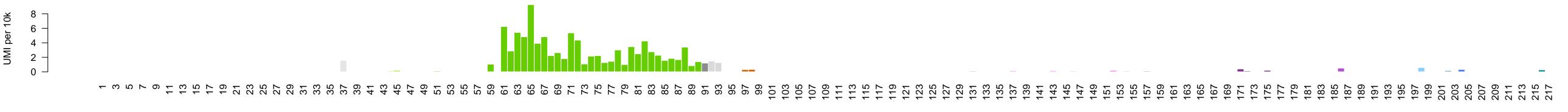
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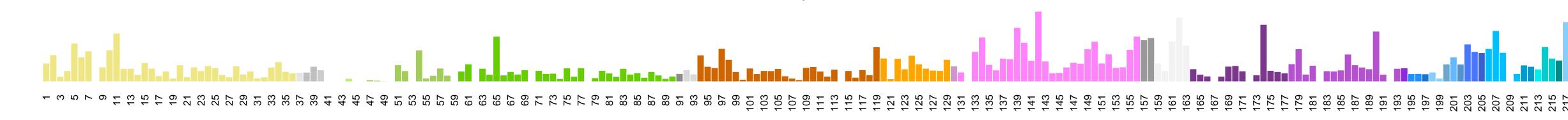


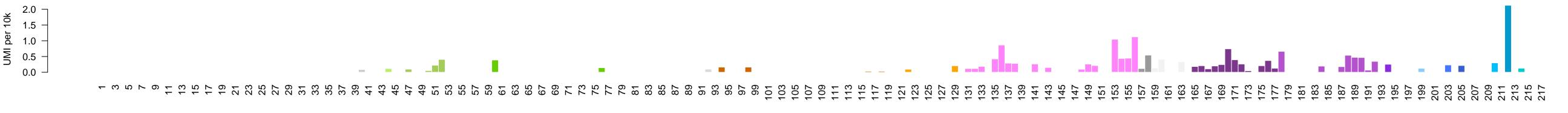
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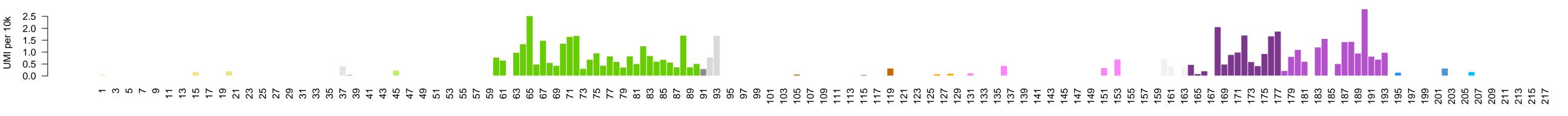


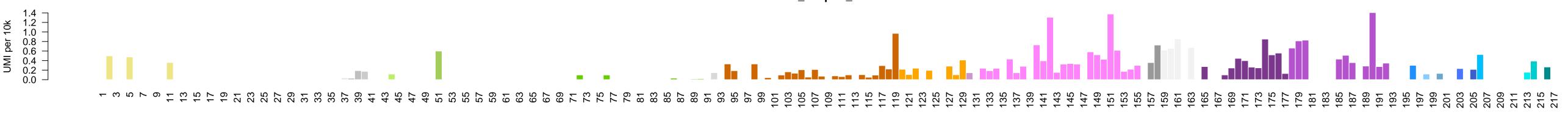


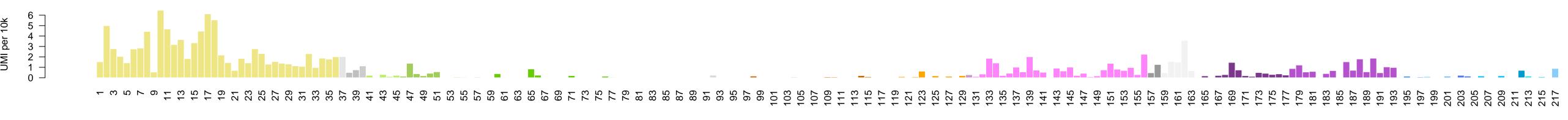


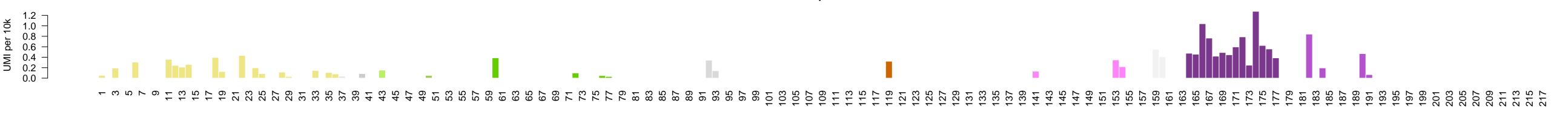


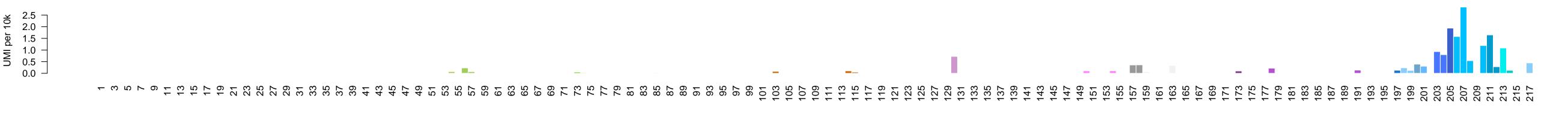


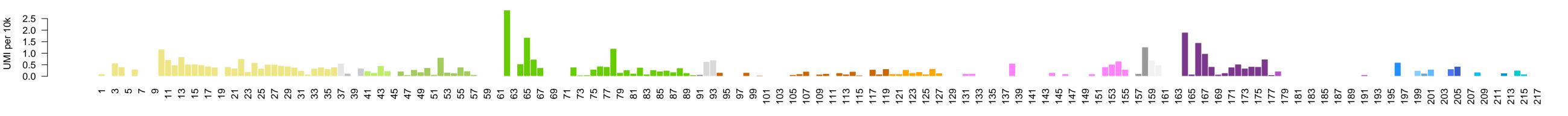


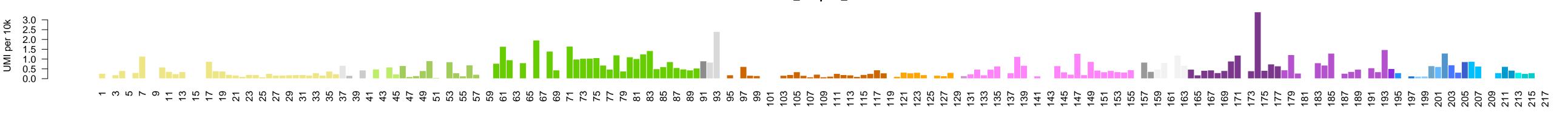


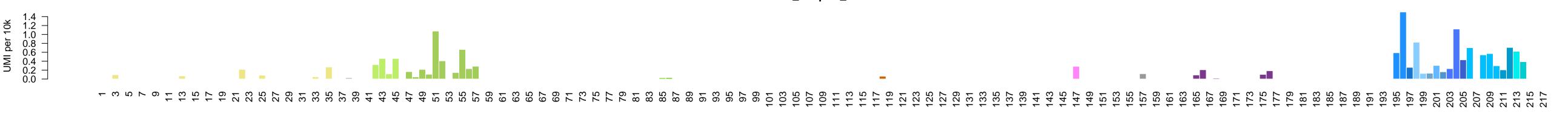






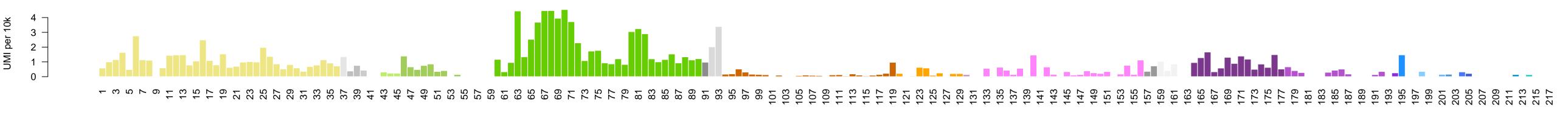


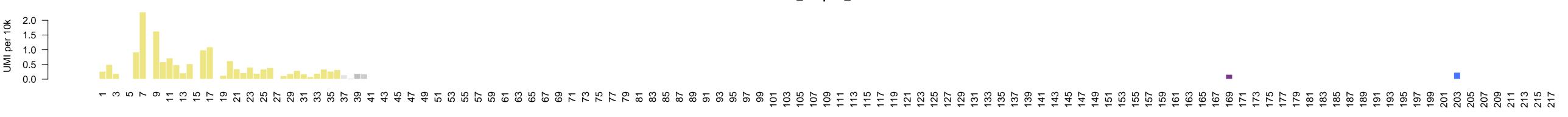




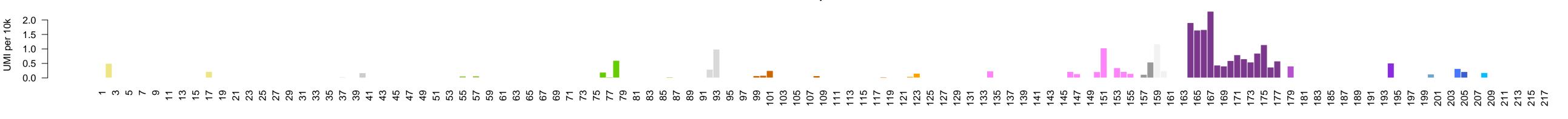
IRK.HG1.0:KCNJ1/KCNJ2/KCNJ3/KCNJ4/KCNJ5/KCNJ6/KCNJ8/KCNJ9/KCNJ10/KCNJ11/KCNJ12/KCNJ13/KCNJ14/KCNJ15/KCNJ16/KCNJ18

TrH2_TrispH2_008304

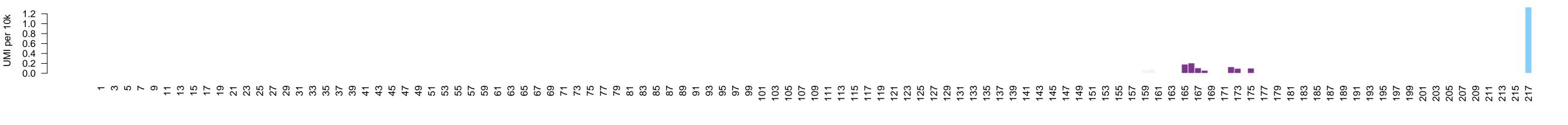




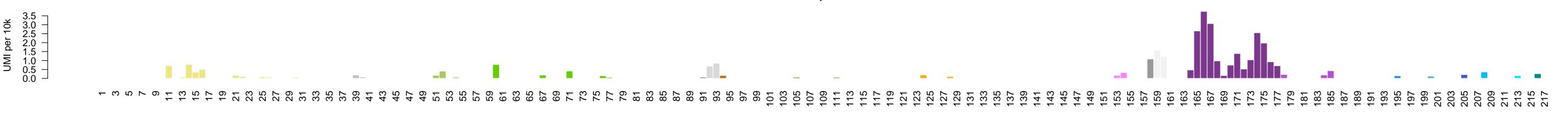
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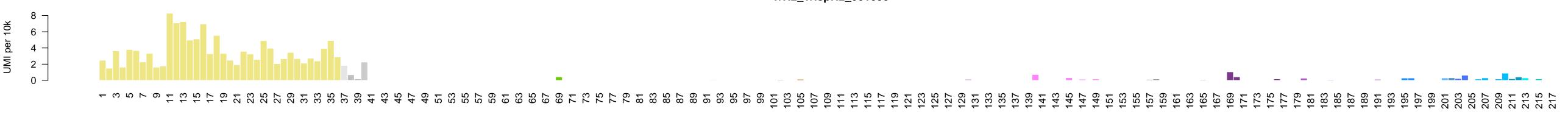
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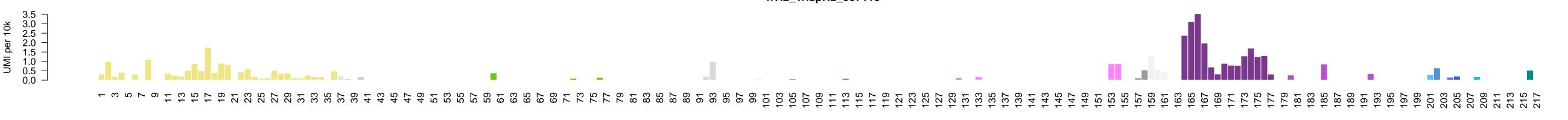
Lig_chan.HG1.4:like:GRIN1/GRIN2A/GRIN2B/GRIN2C/GRIN2D/GRIN3A/GRIN3B TrH2_TrispH2_001054



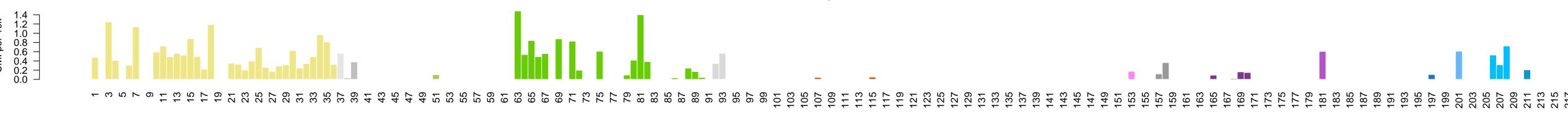
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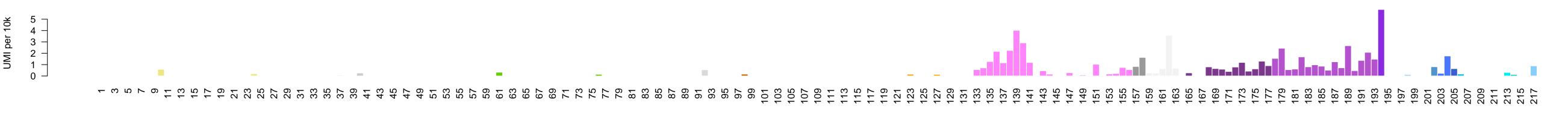
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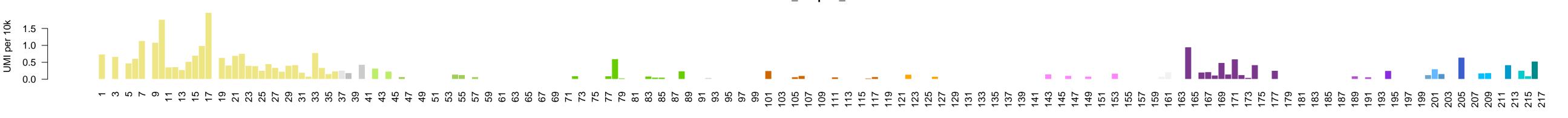


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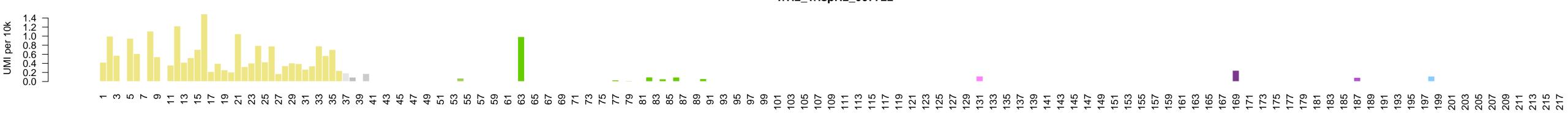
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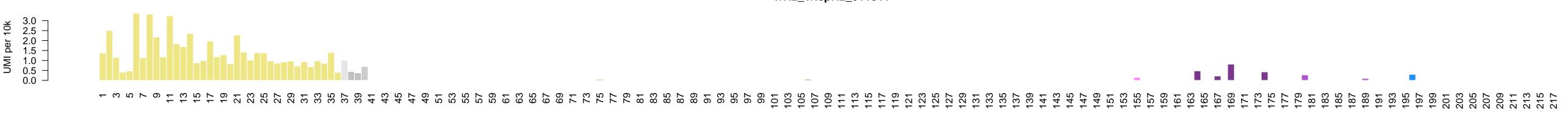




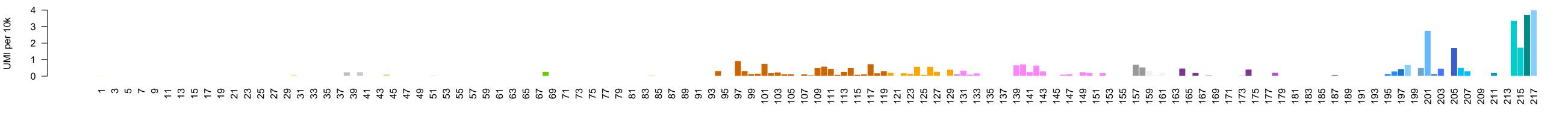
Lig_chan.HG1.6:like:GRIA1/GRIA2/GRIA3/GRIA4/GRID1/GRID2/GRIK1/GRIK2/GRIK3/GRIK4/GRIK5/GRIN1/GRIN2A/GRIN2B/GRIN2C/GRIN2D/GRIN3A/GRIN3B

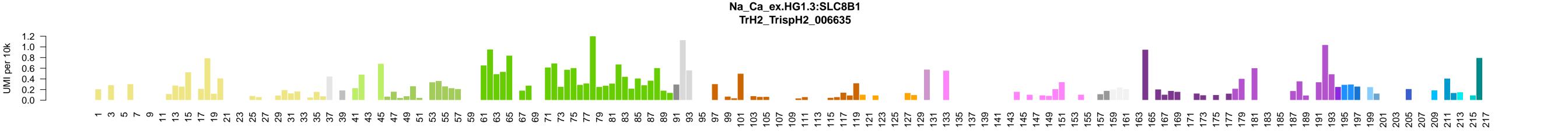
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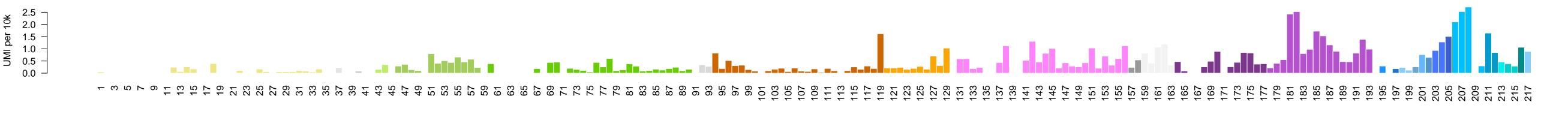


Na_Ca_ex.HG1.0:like:SLC24A1/SLC24A2/SLC24A3/SLC24A4/SLC24A5 TrH2_TrispH2_000976

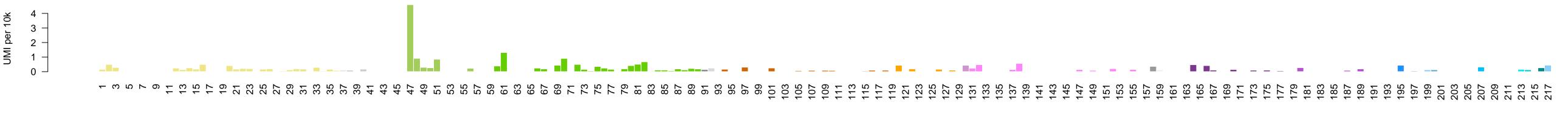




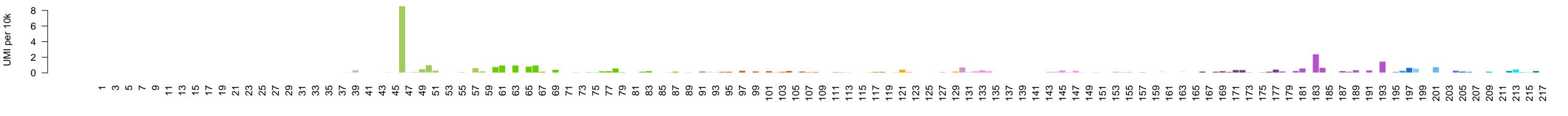
Na_Ca_ex.HG1.4:SLC24A1/SLC24A2 TrH2_TrispH2_001316



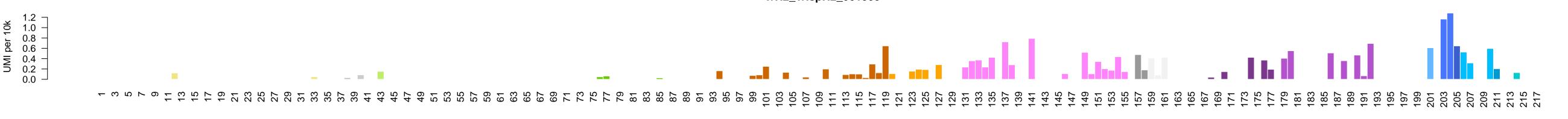
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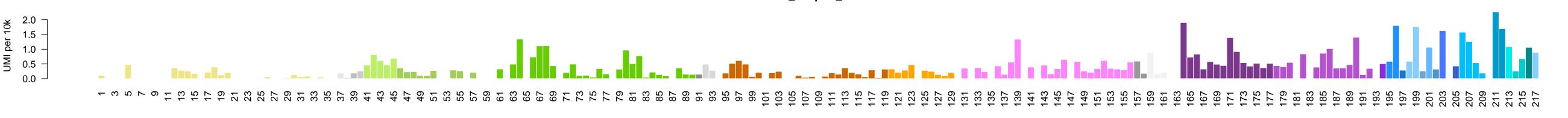
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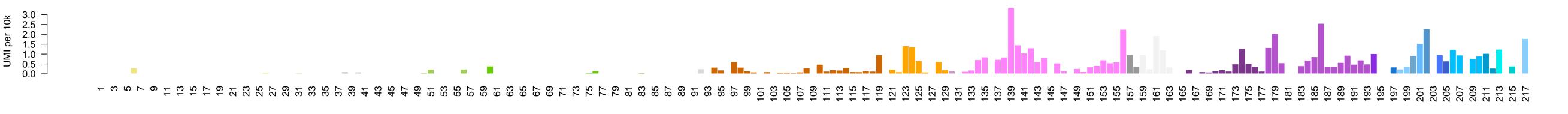
Na_Ca_ex.HG2.0:like:SLC8A1/SLC8A2/SLC8A3 TrH2_TrispH2_001305



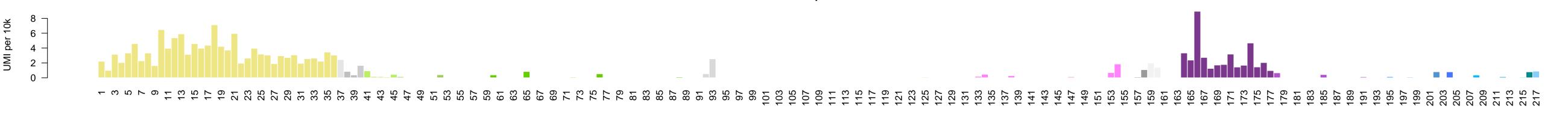
Na_Ca_ex.HG2.1:SLC8A1/SLC8A2/SLC8A3 TrH2_TrispH2_009772



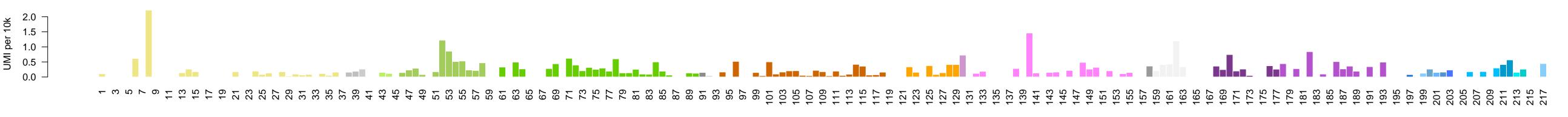
Na_Ca_ex.HG2.2:like:SLC8A1/SLC8A2/SLC8A3 TrH2_TrispH2_002321



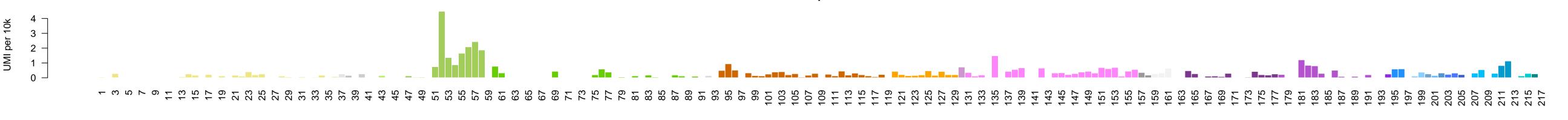
Na_Ca_ex.HG3.0:NA TrH2_TrispH2_001491



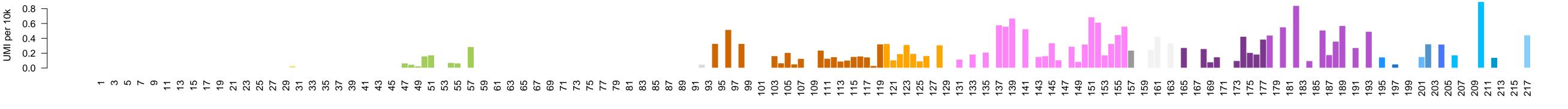
P2X_receptor.HG1.3:P2RX1/P2RX2/P2RX3/P2RX4/P2RX5/P2RX5-TAX1BP3/P2RX6/P2RX7
TrH2 TrispH2 000404



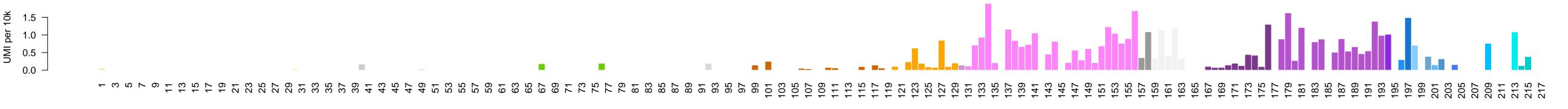
P2X_receptor.HG1.3:P2RX1/P2RX2/P2RX3/P2RX4/P2RX5/P2RX5-TAX1BP3/P2RX6/P2RX7 TrH2_TrispH2_006659



PKD_channel.HG1.13:like:TRPC3/TRPC6/TRPC7 TrH2_TrispH2_006118



VGCC_alpha2.HG1.0:CACNA2D3/CACNA2D4 TrH2_TrispH2_003762



VGCC_alpha2.HG1.1:CACNA2D1/CACNA2D2 TrH2_TrispH2_003756

