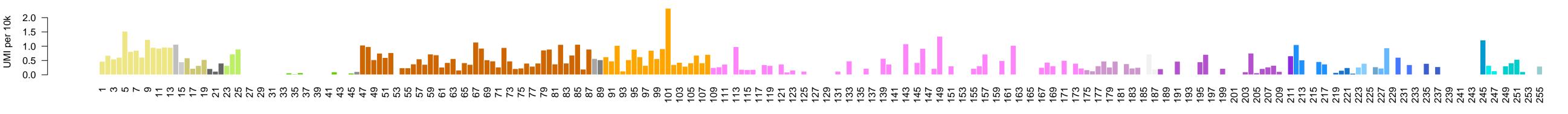
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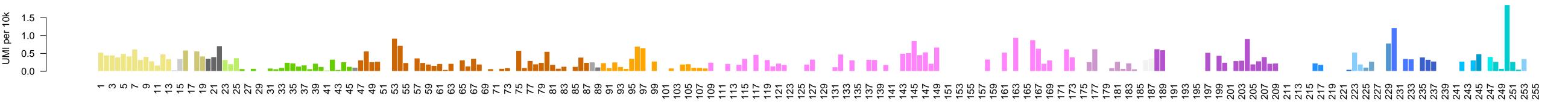
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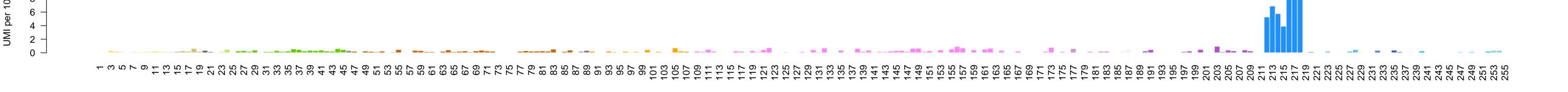


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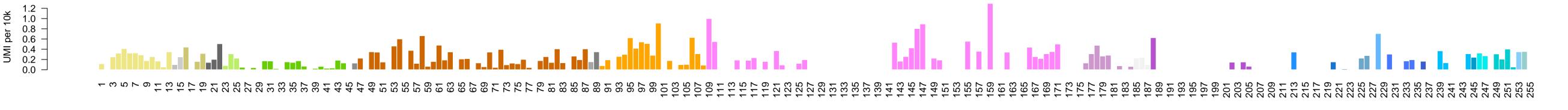
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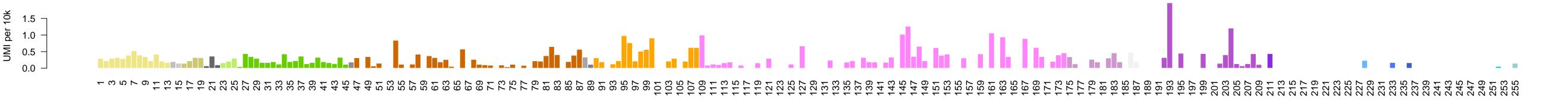
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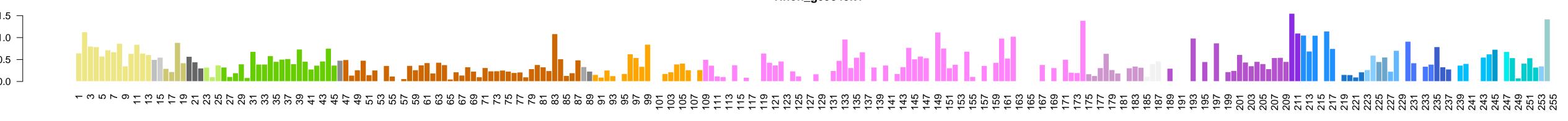
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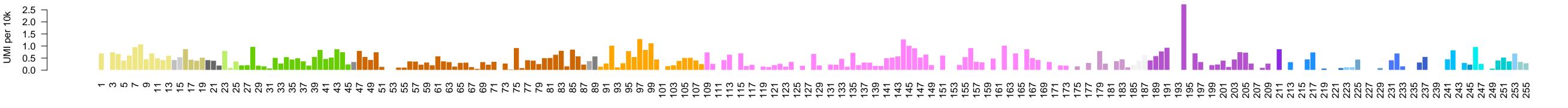
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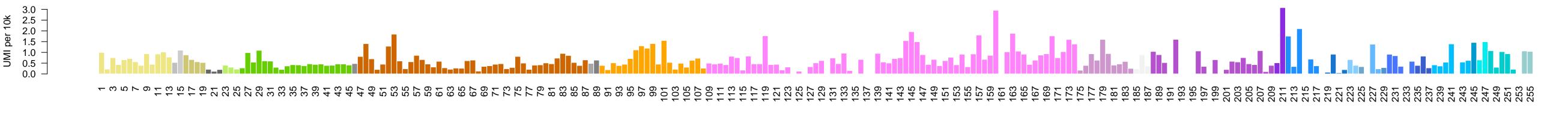
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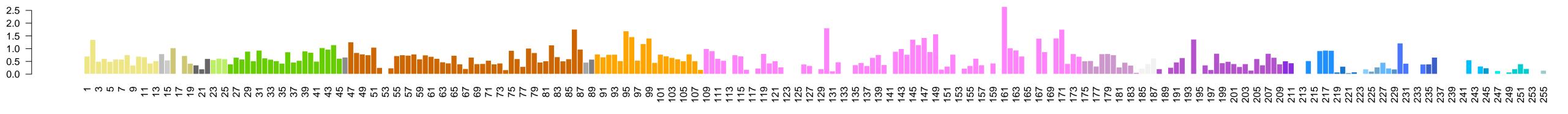
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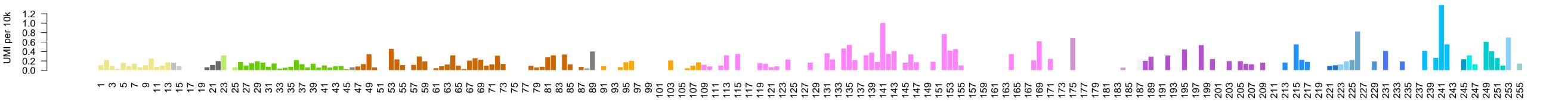
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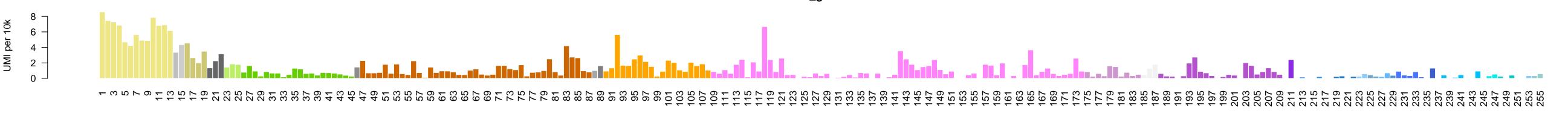
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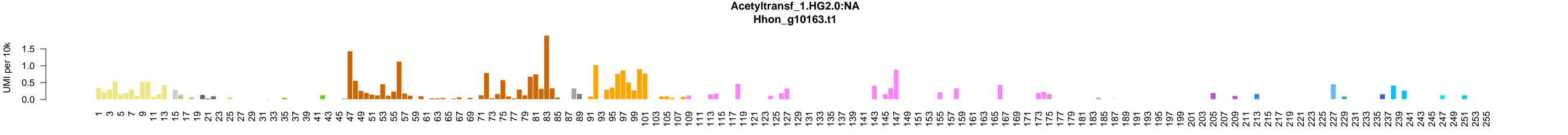


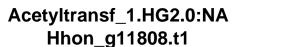
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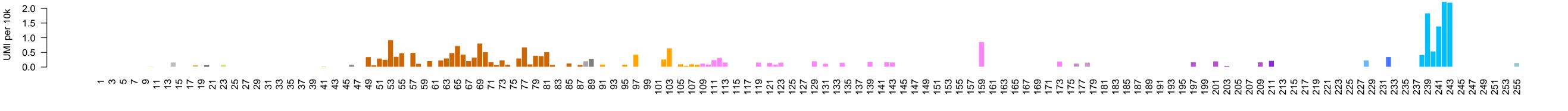


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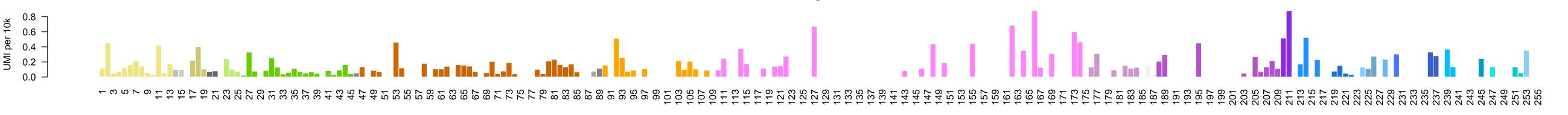




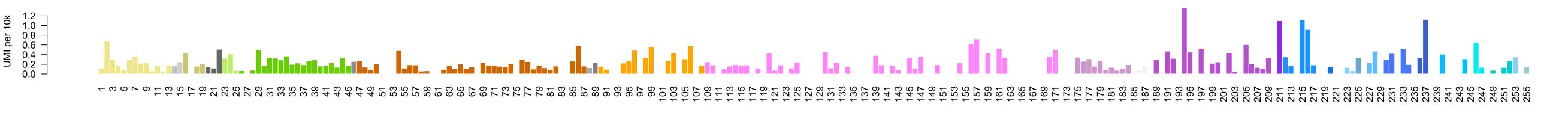




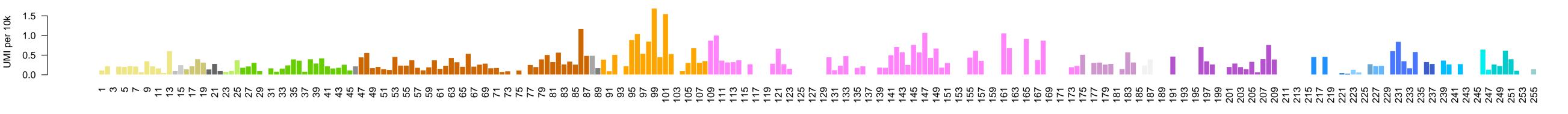
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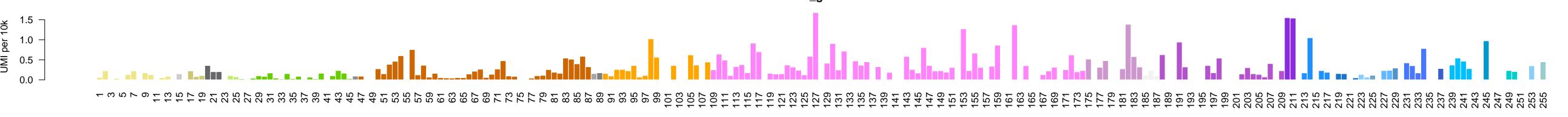
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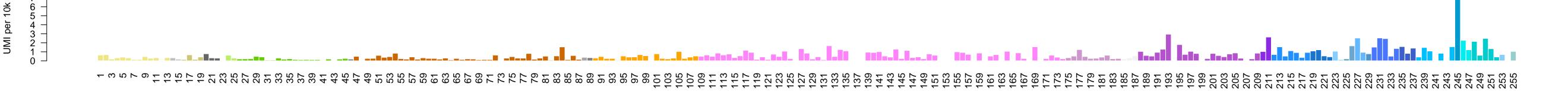
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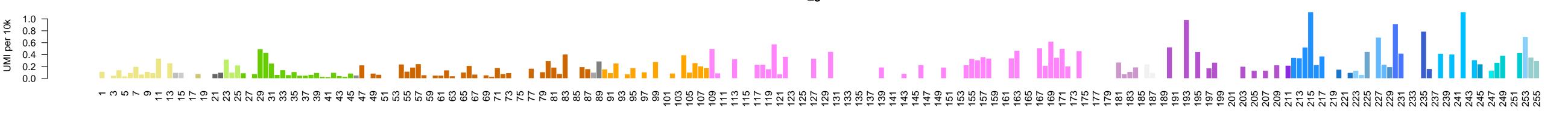
BAH.HG1.3:PBRM1;Bromodomain.HG1.29:PBRM1 Hhon_g09288.t1



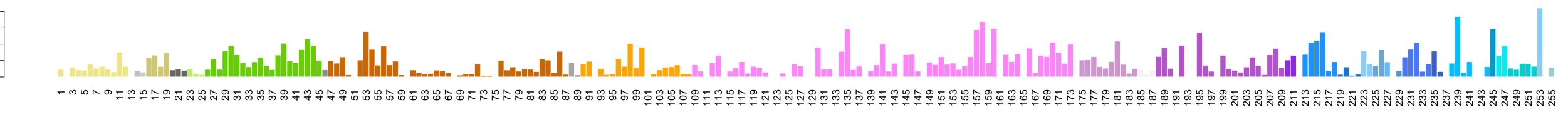




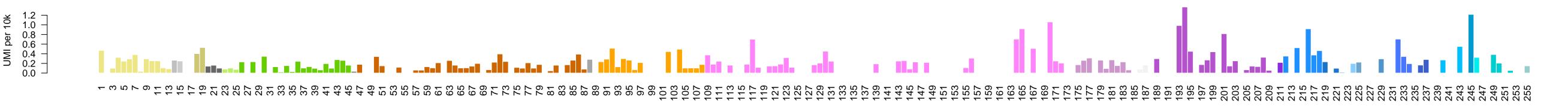
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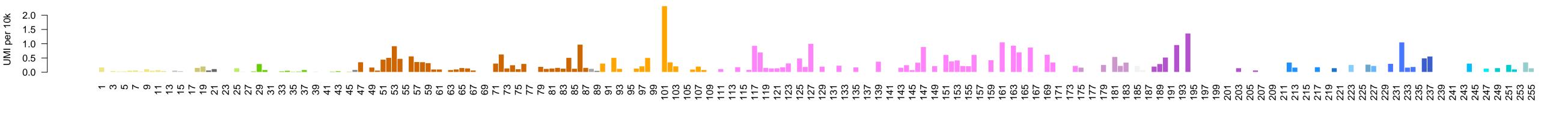
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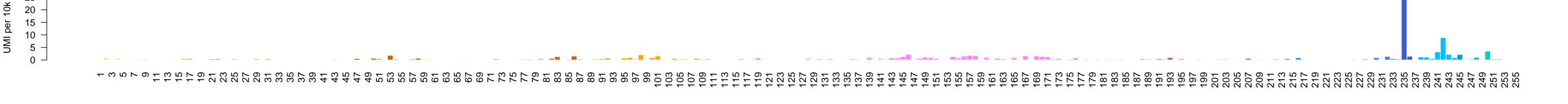
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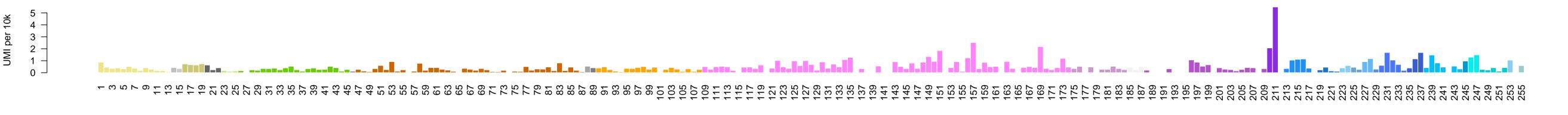




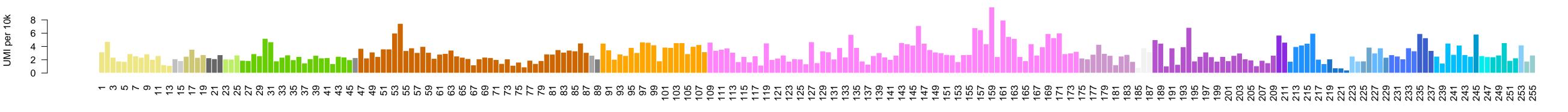




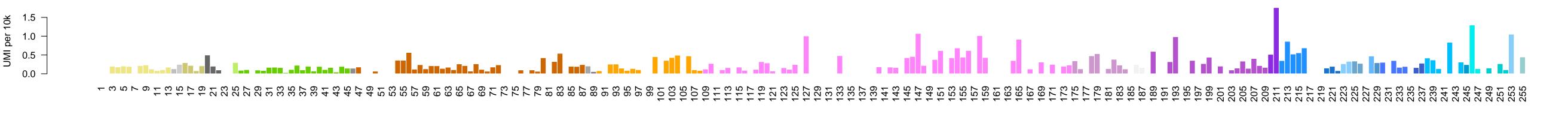
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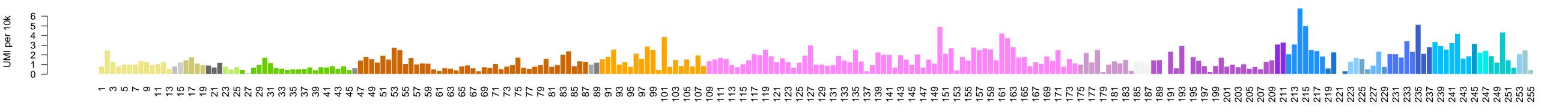
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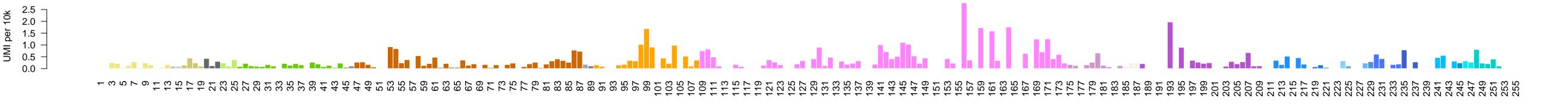
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Hhon_g08078.t1



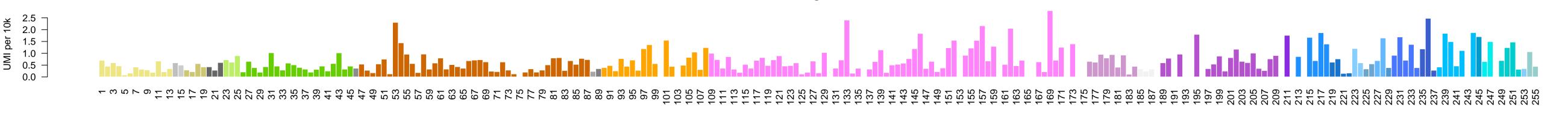
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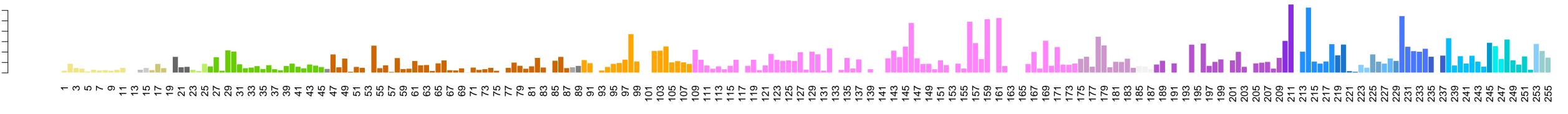
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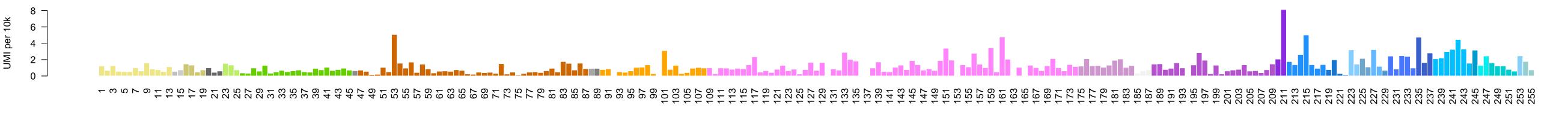
Bromodomain.HG1.30:SMARCA2/SMARCA4;SNF2_N.HG1.8:SMARCA2/SMARCA4 Hhon_g07719.t1



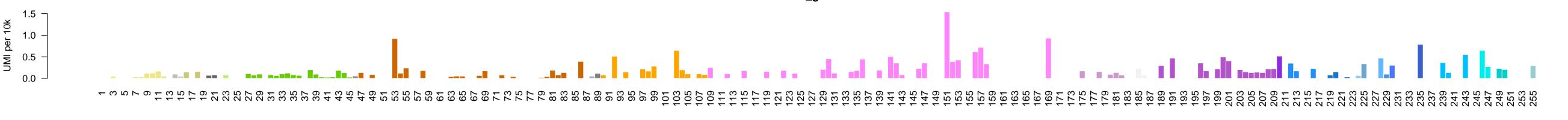
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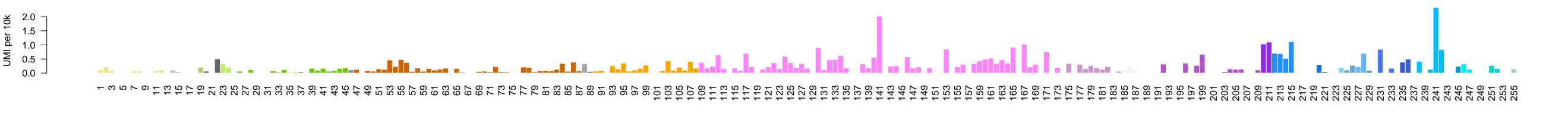
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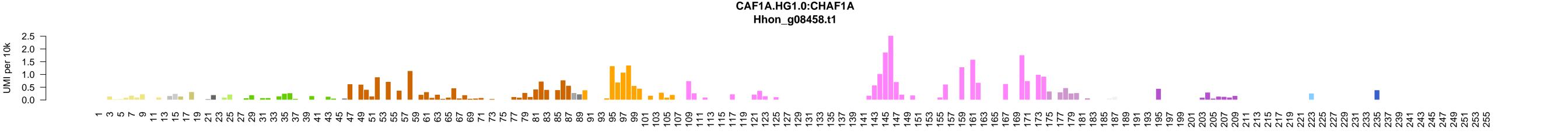


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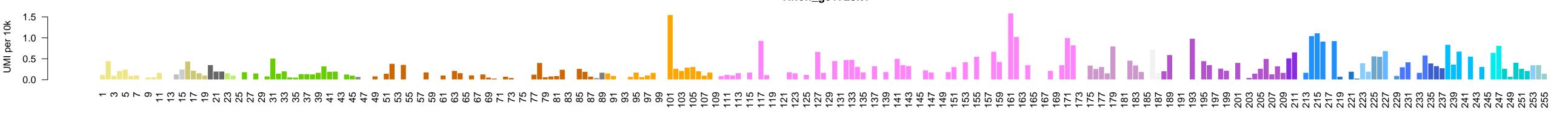


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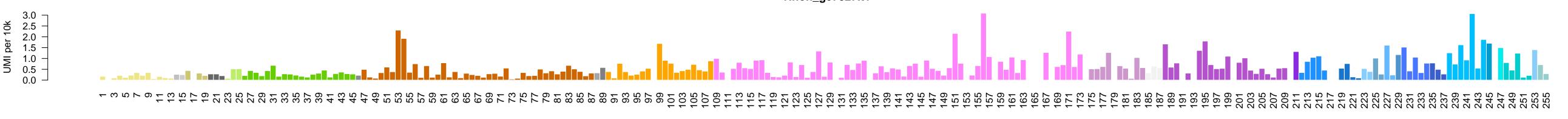




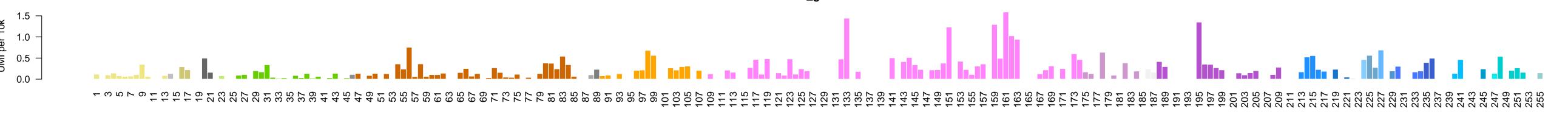
Chromo.HG1.12:AC006064.6/CHD3/CHD4/CHD5;SNF2_N.HG1.26:AC006064.6/CHD3/CHD4/CHD5 Hhon_g01723.t1



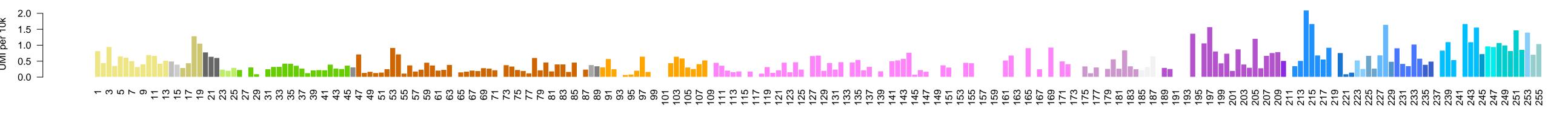
Chromo.HG1.13:CHD6/CHD7/CHD8/CHD9;SNF2_N.HG1.28:CHD6/CHD7/CHD8/CHD9 Hhon_g07827.t1



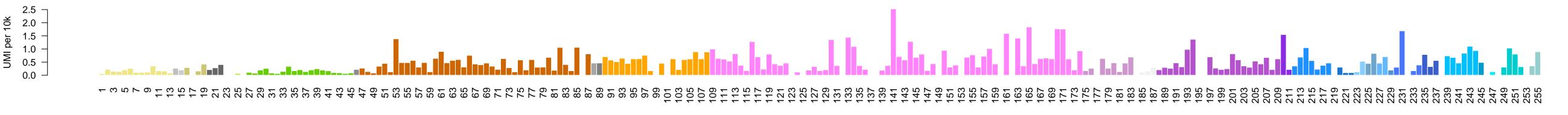
Chromo.HG1.14:CHD1/CHD2;SNF2_N.HG1.27:CHD1/CHD2 Hhon_g05913.t1



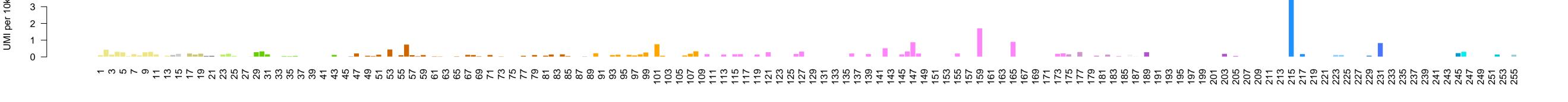
Chromo.HG1.21:CBX2/CBX4/CBX6/CBX7/CBX8
Hhon_g03880.t1



Chromo.HG1.6:SUV39H1/SUV39H2 Hhon_g09127.t1



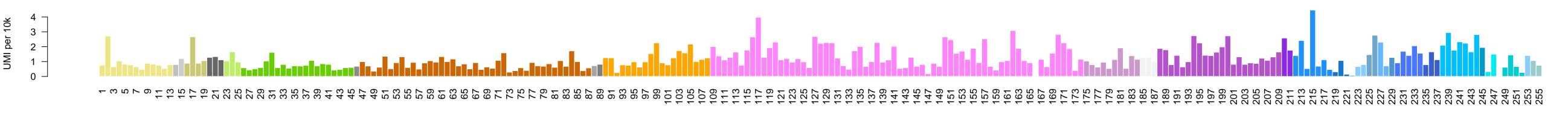
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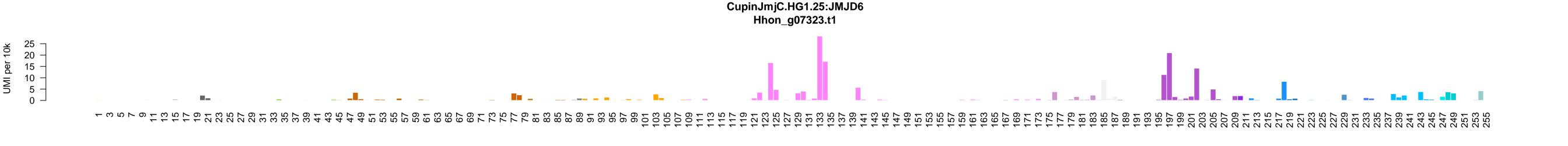


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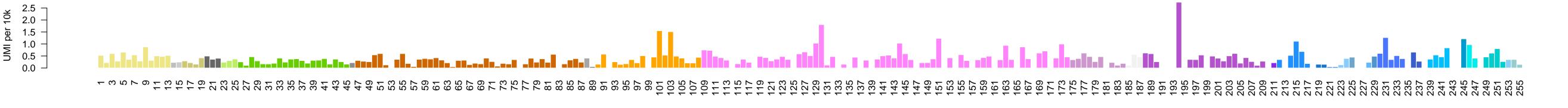


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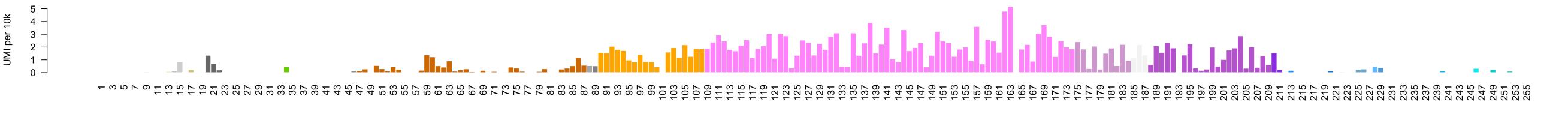




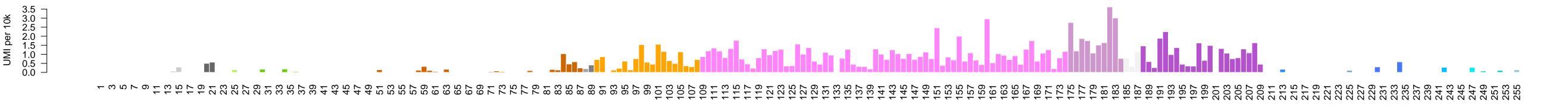
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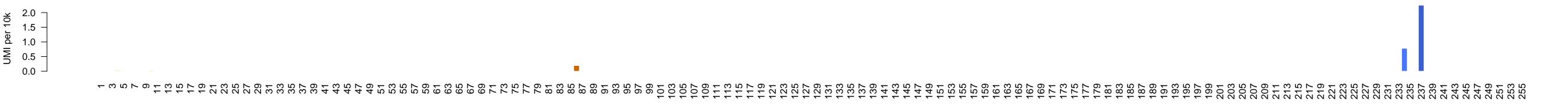
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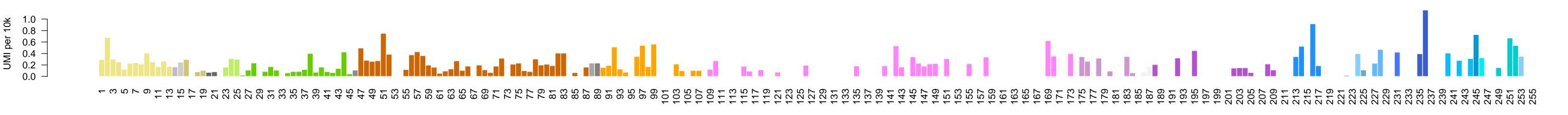
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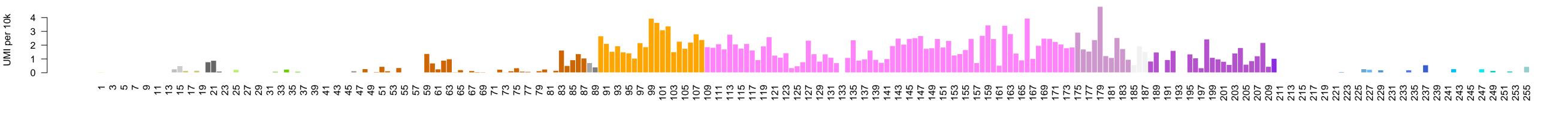
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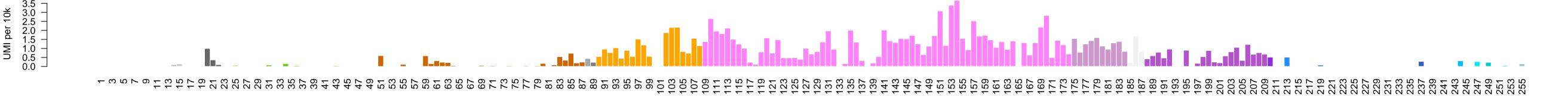
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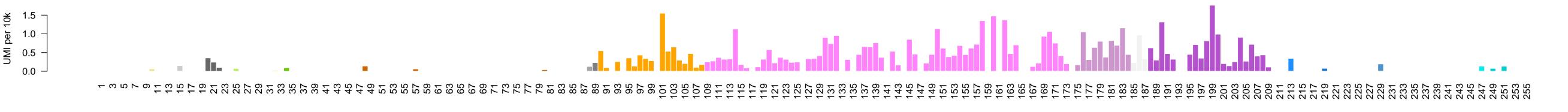
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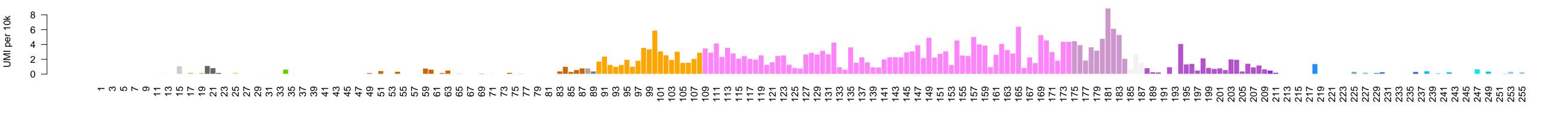
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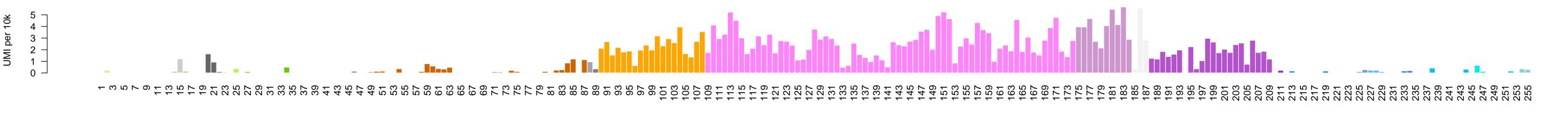
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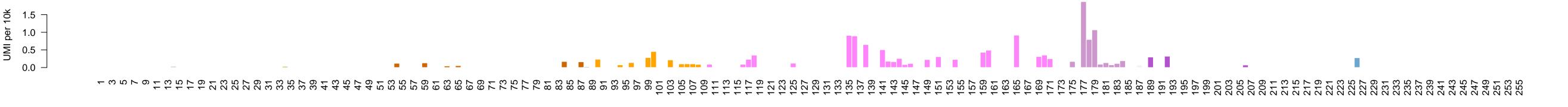
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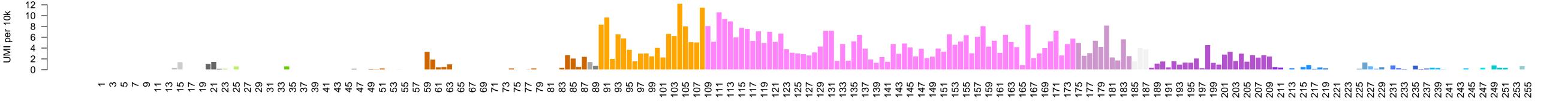
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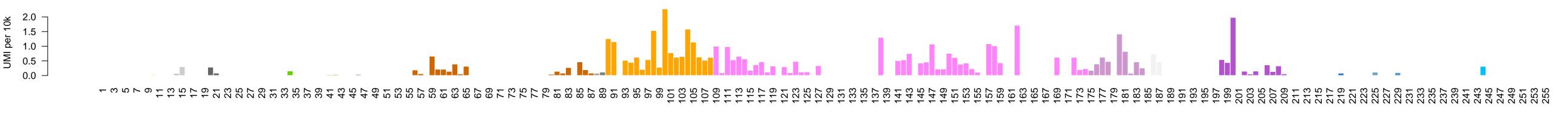
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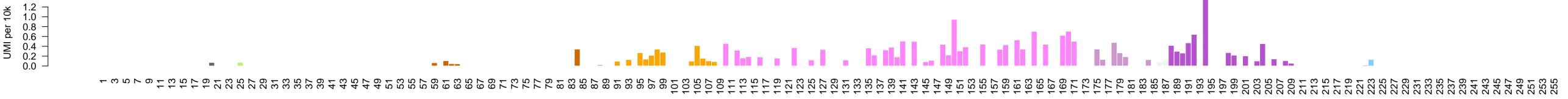
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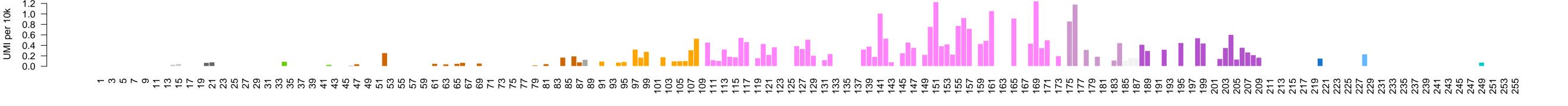
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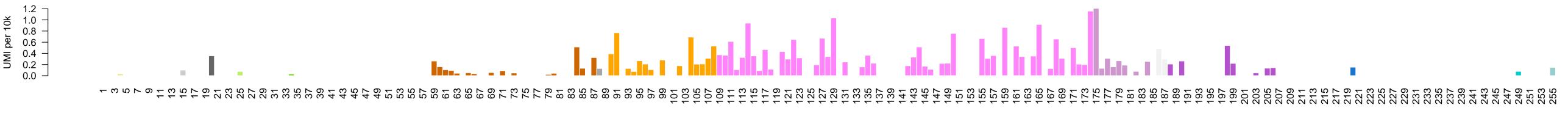
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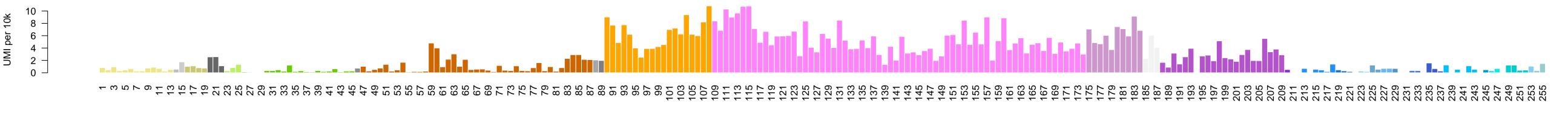
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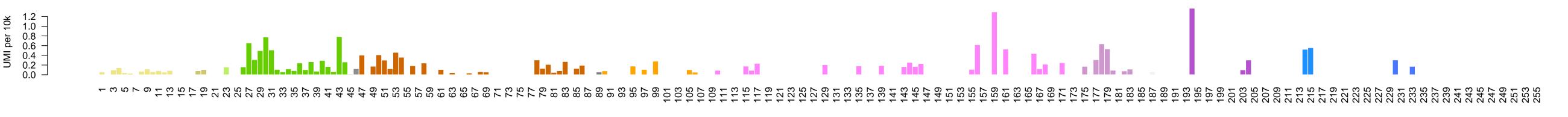
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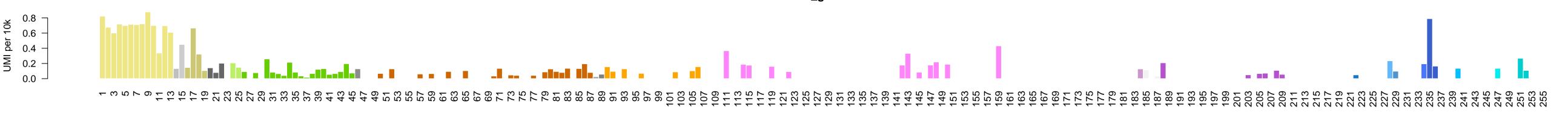


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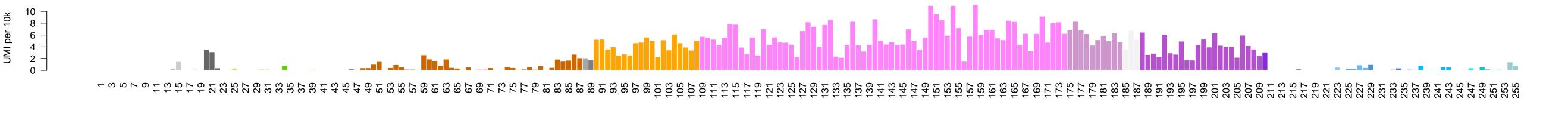


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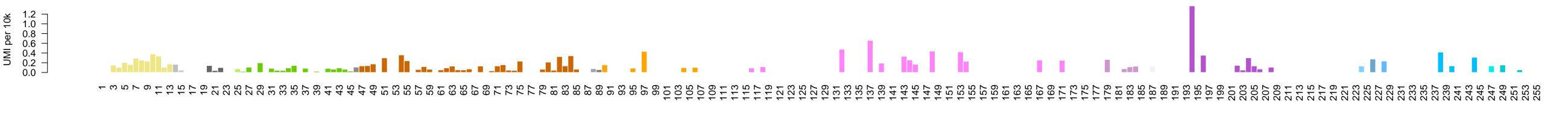
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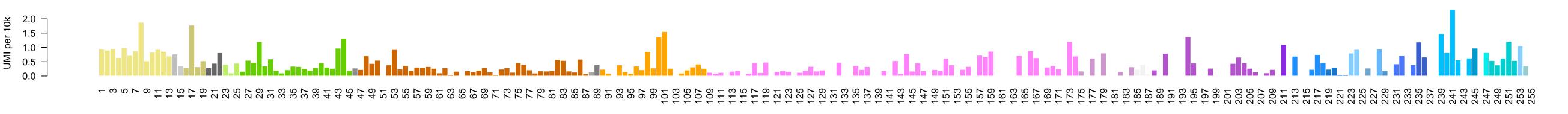
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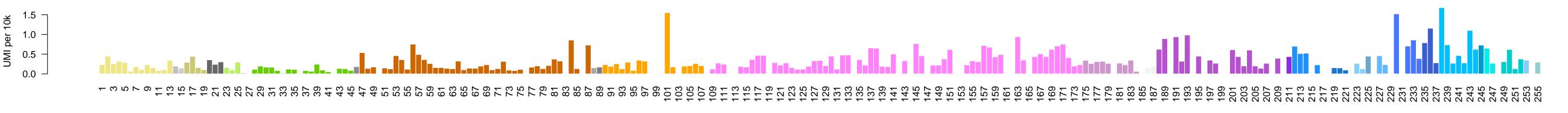
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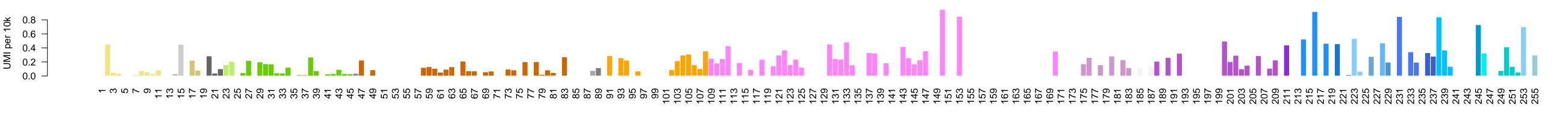
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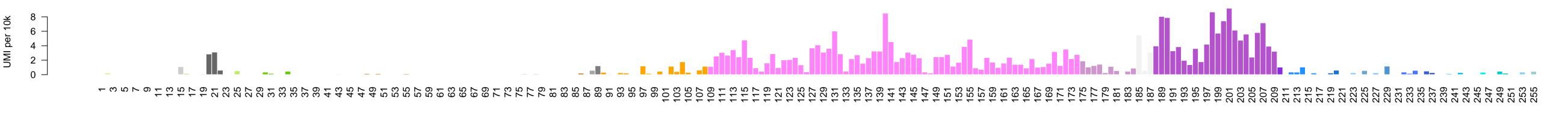
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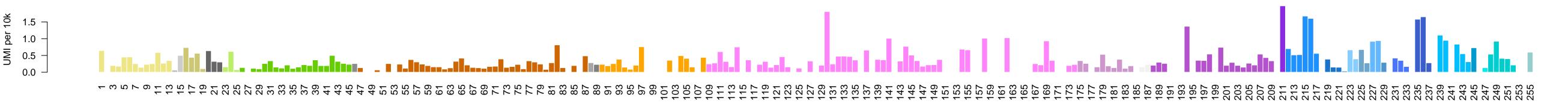
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DOT1.HG1.2:like:DOT1L Hhon_g06626.t1

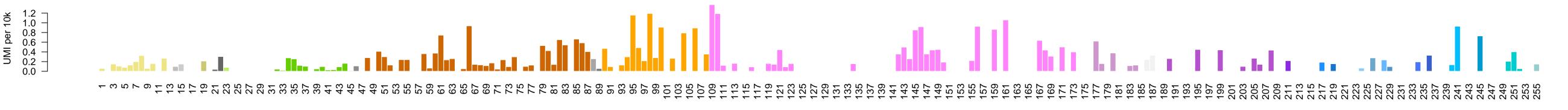


DOT1.HG1.3:DOT1L Hhon_g00168.t1

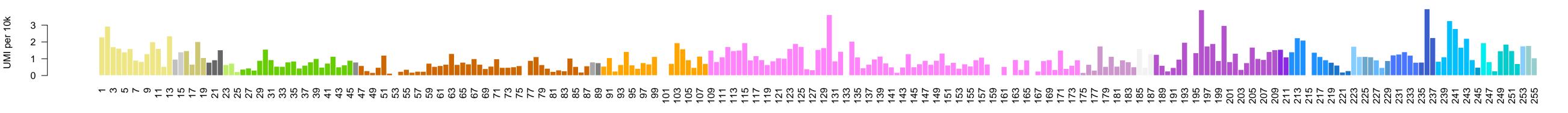


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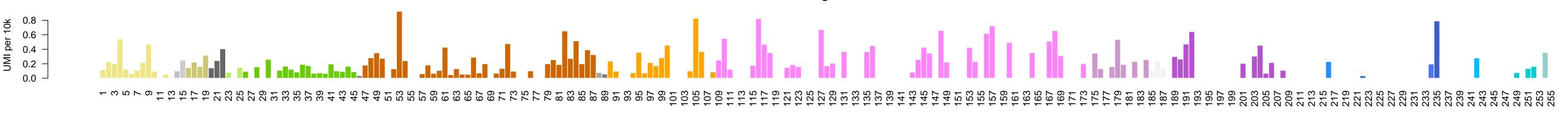
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Hist_deacetyl.HG1.1:HDAC11 Hhon_g04359.t1

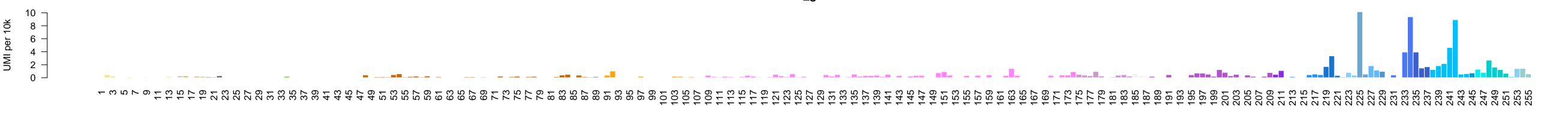


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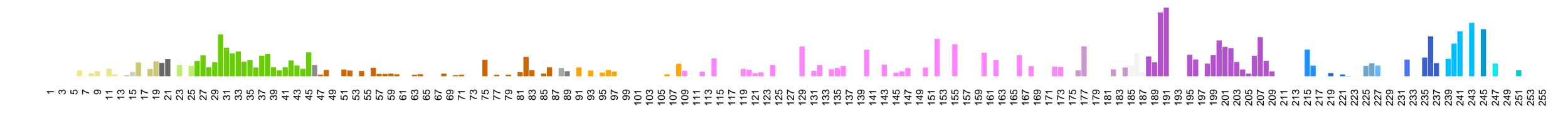


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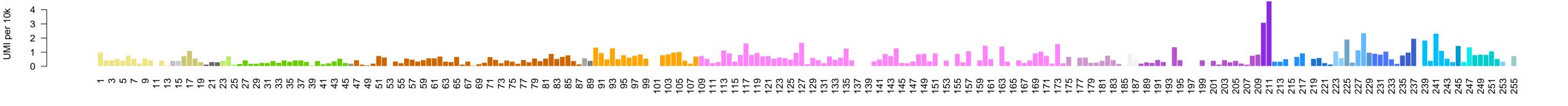


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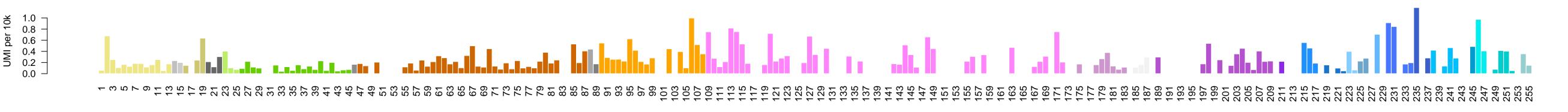


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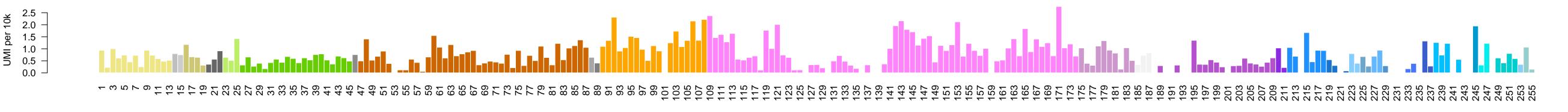
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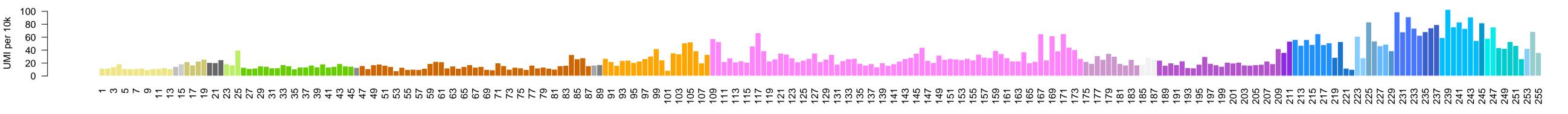
Hist_deacetyl.HG1.8:HDAC3 Hhon_g03531.t1



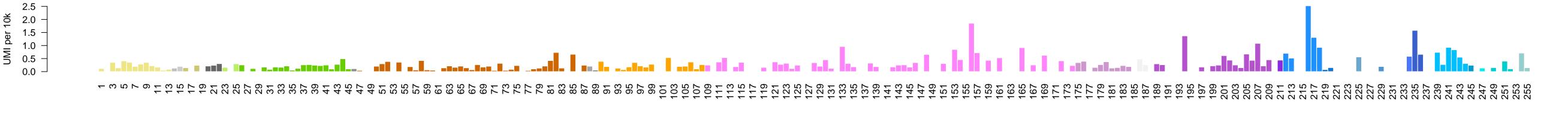
Histone.HG2.1:CENPA Hhon_g09811.t1



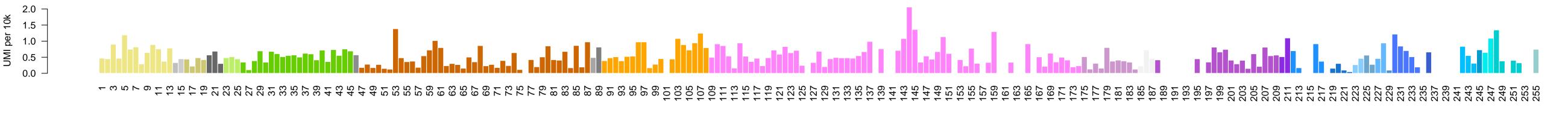
Histone.HG2.13:like:H3-3A/H3-3B/H3-5/H3Y1/H3Y2 Hhon_g06924.t1



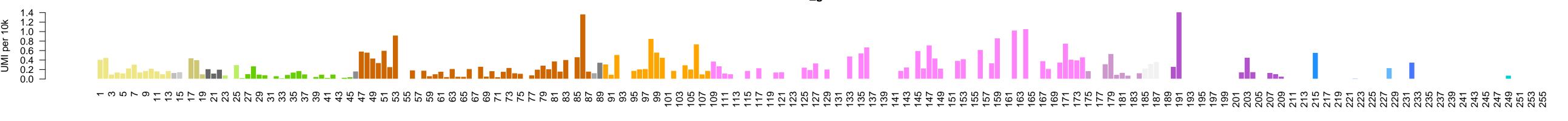
Histone.HG3.0:DRAP1 Hhon_g07246.t1



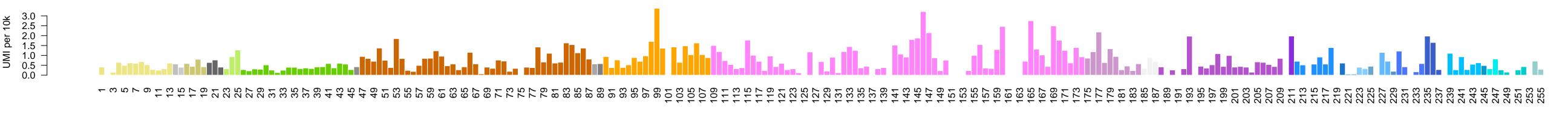
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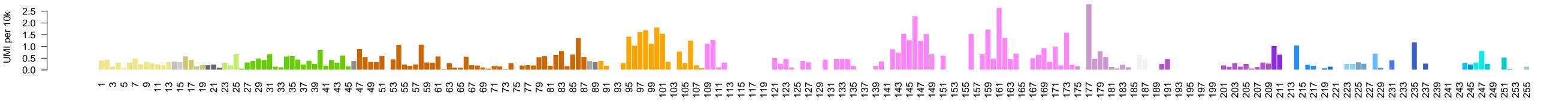
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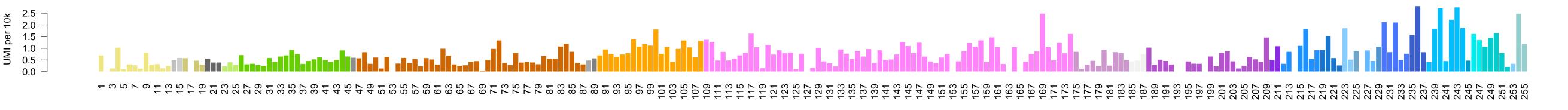
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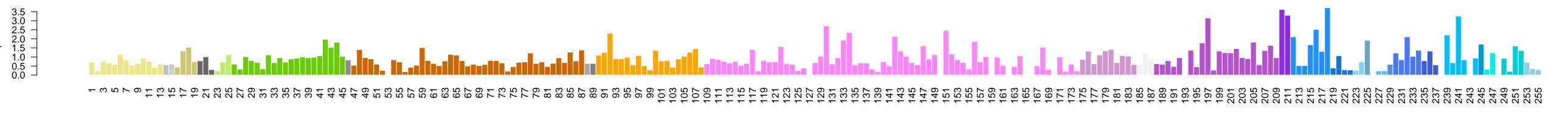
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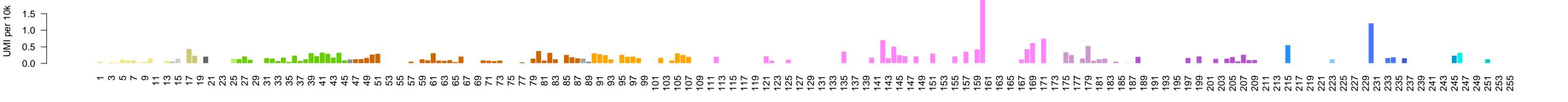
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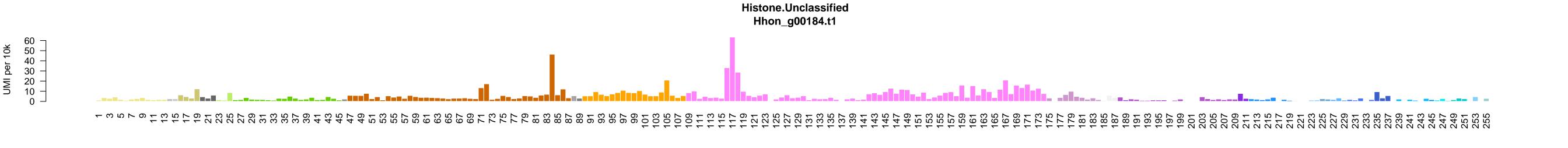


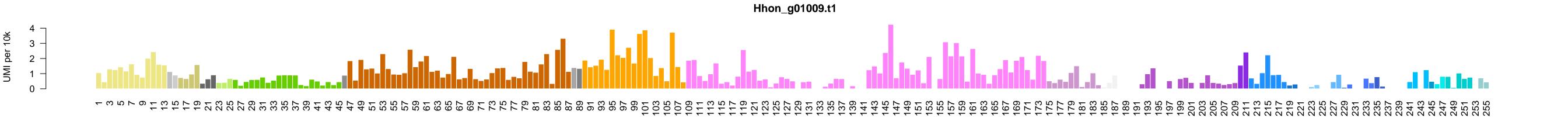
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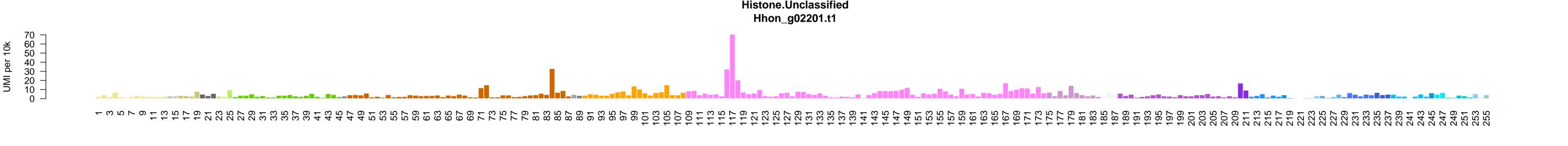


Histone.HG8.0:NA Hhon_g01462.t1

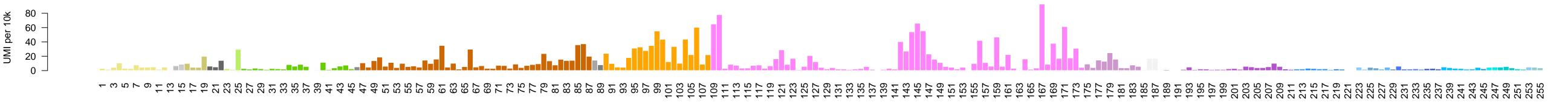


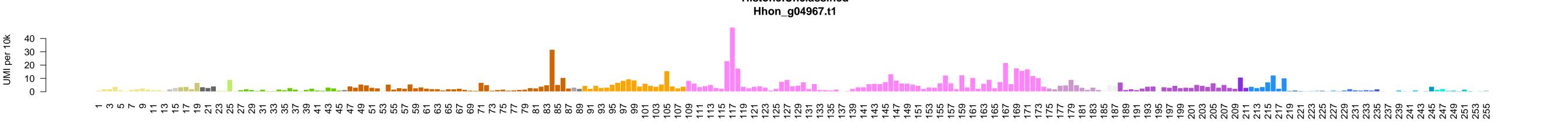




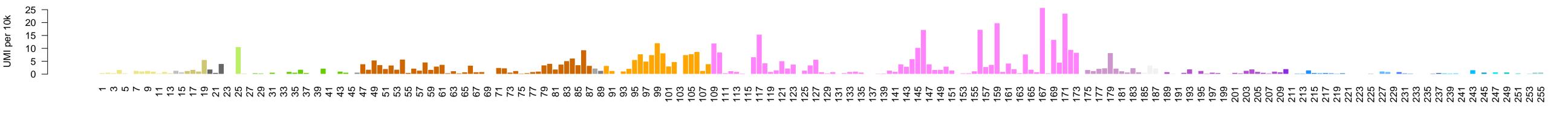


Histone.Unclassified Hhon_g02498.t1

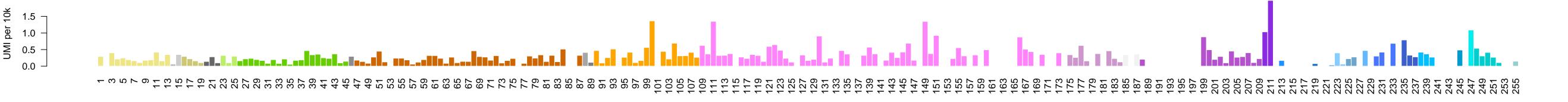




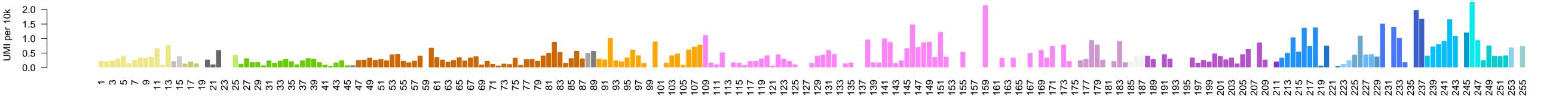
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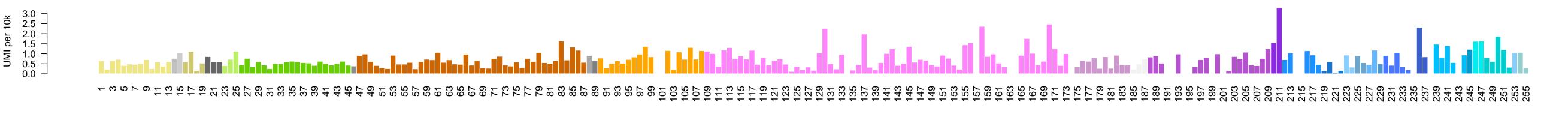
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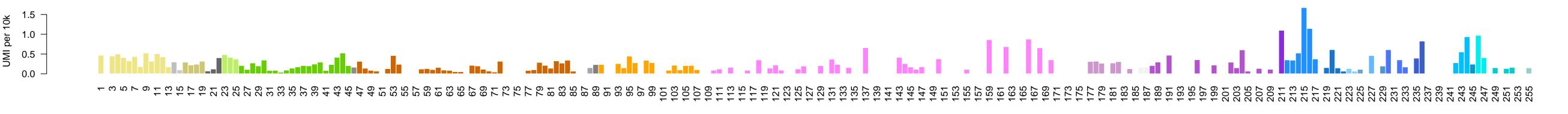
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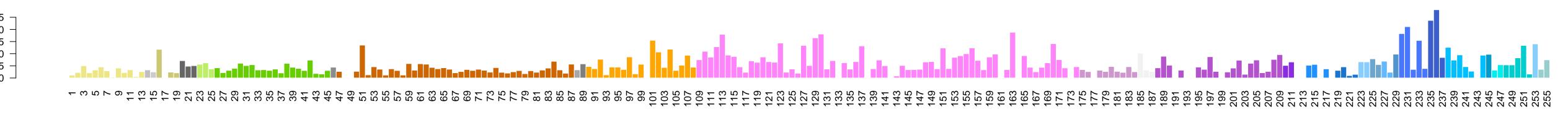
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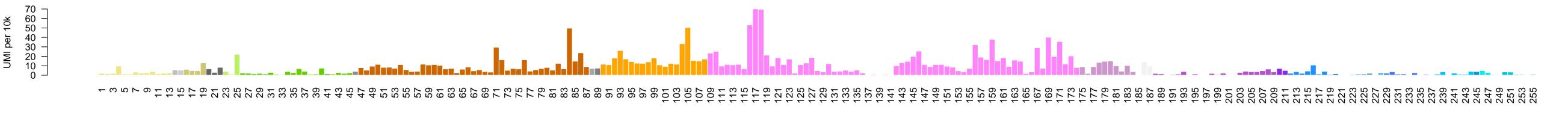
LinkerHistone.HG1.0:like:H1-0/H1-1/H1-2/H1-3/H1-4/H1-5/H1-6/H1-8/H1-10/HP1BP3/SHPRH Hhon_g00175.t1



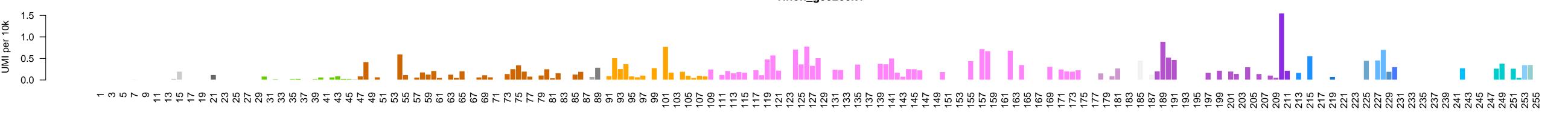
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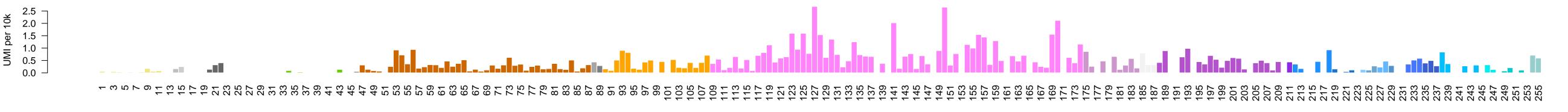


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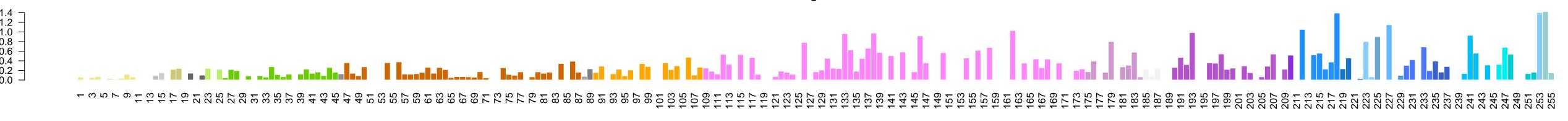


MBT.HG1.1:like:PHF20L1 Hhon_g00266.t1





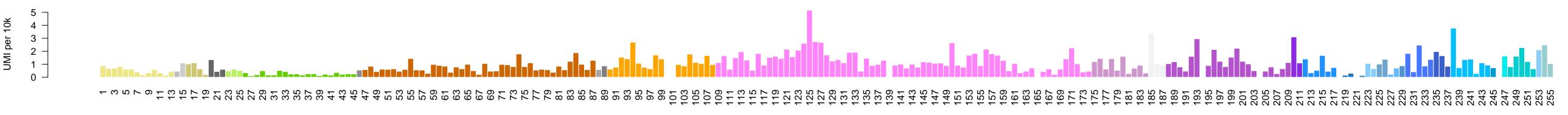
MBT.HG1.3:AL031681.2/L3MBTL1/L3MBTL3/L3MBTL4 Hhon_g00122.t1



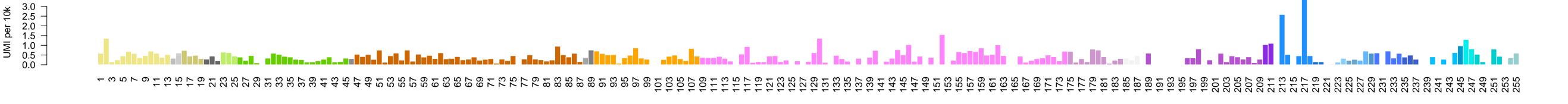
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MBT.HG1.4:SCMH1/SCML2

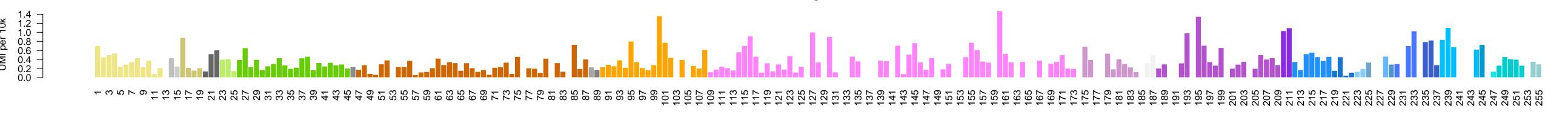


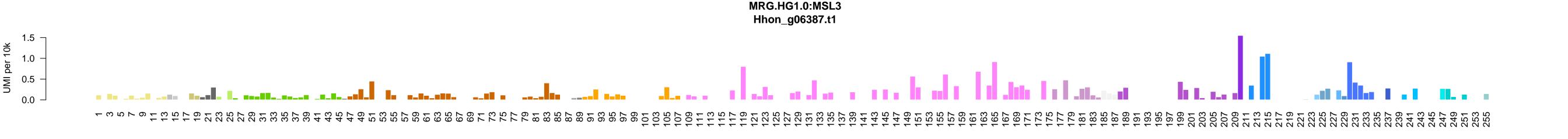




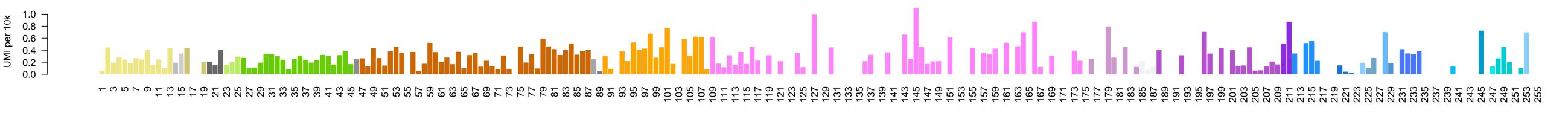


MOZ_SAS.HG1.7:KAT7 Hhon_g11664.t1



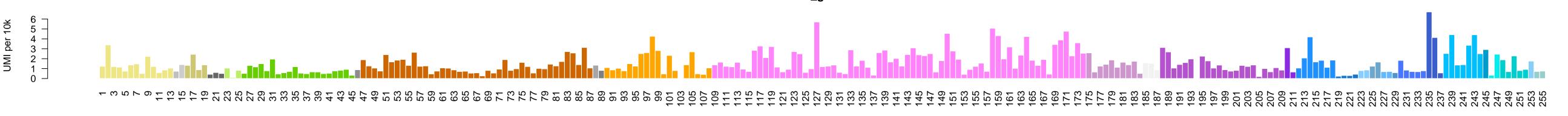


MRG.HG1.1:MORF4L1/MORF4L2 Hhon_g08447.t1



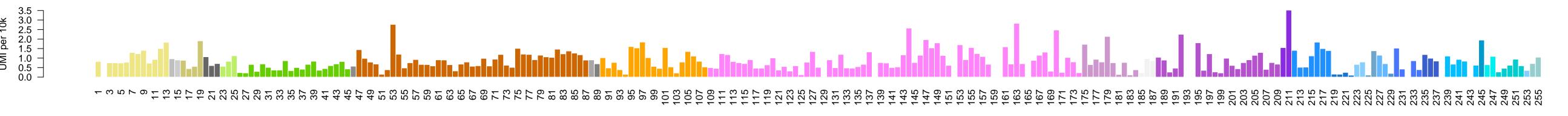
NAP.HG1.0:SET/SETSIP/TSPY1/TSPY2/TSPY3/TSPY4/TSPY8/TSPY9P/TSPY10/TSPYL1/TSPYL2/TSPYL4/TSPYL5/TSPYL6

Hhon_g09411.t1

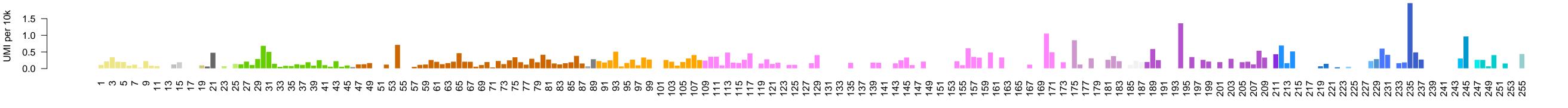




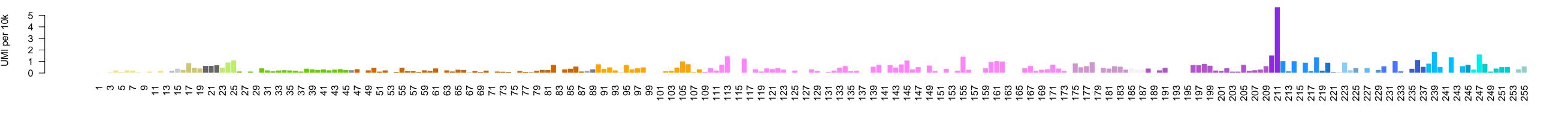
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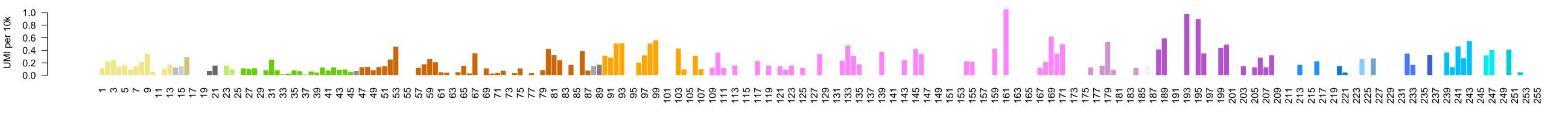
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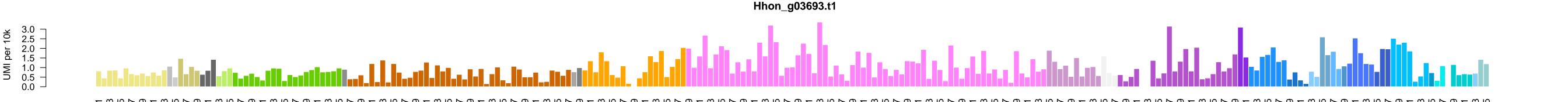




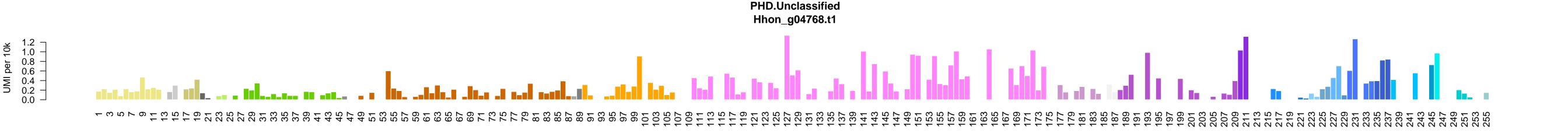


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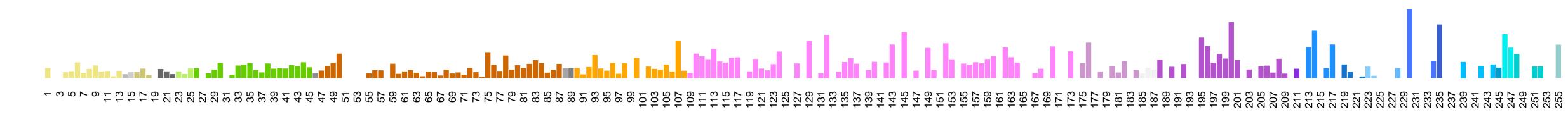




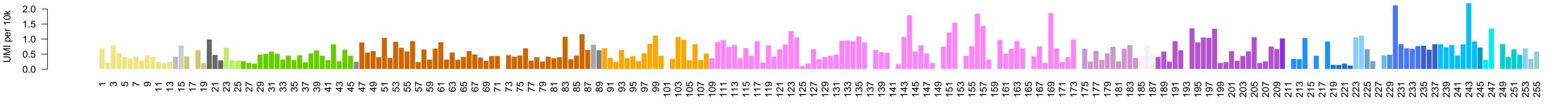
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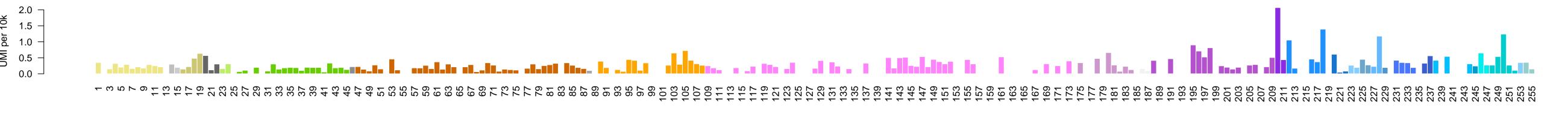
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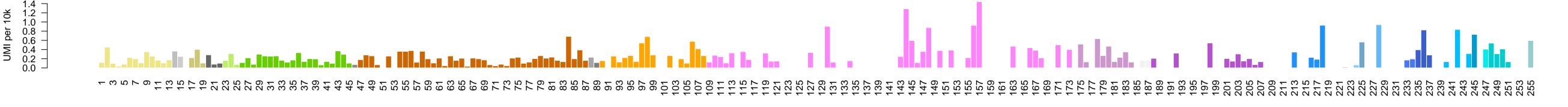


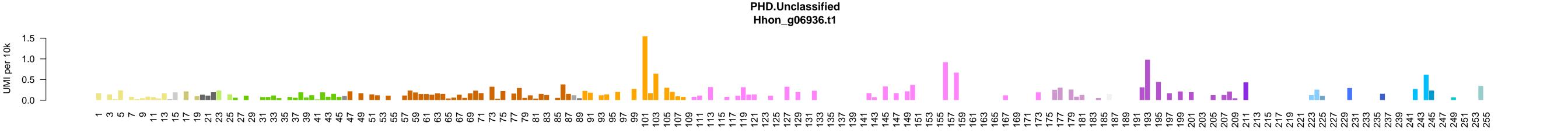




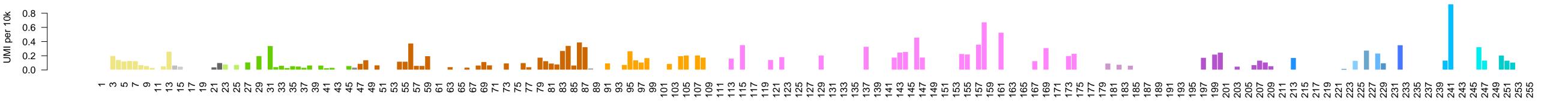


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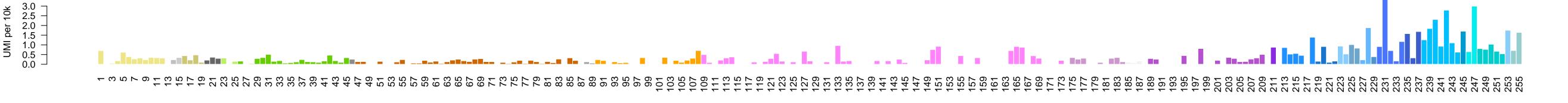




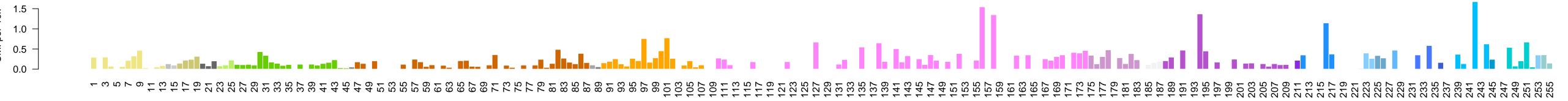
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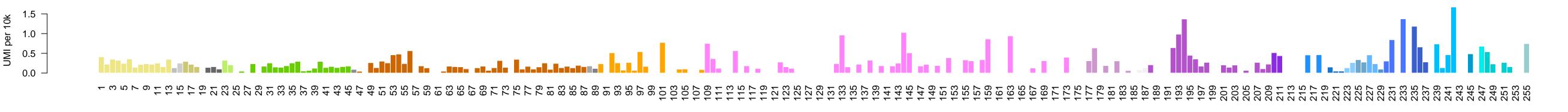




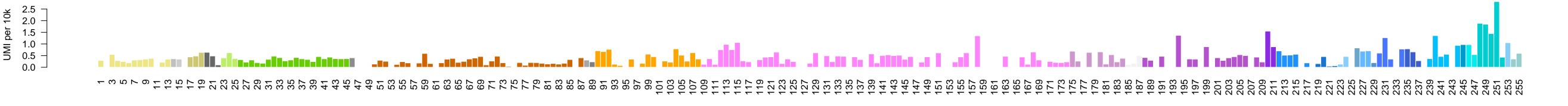




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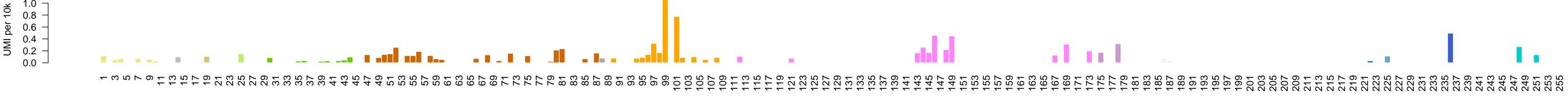




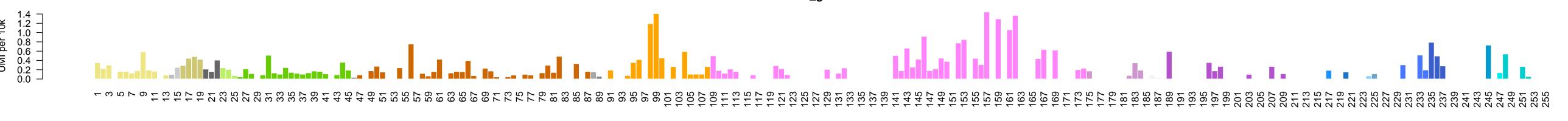
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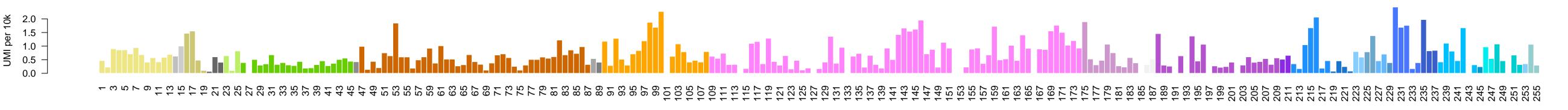
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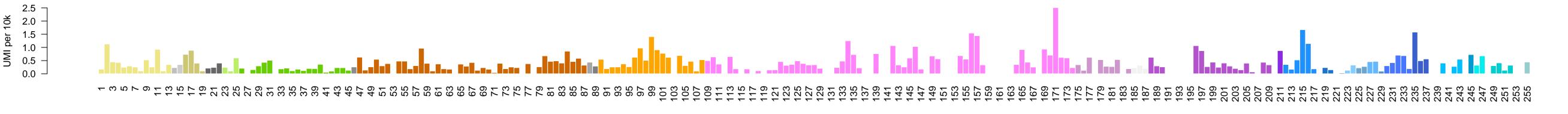
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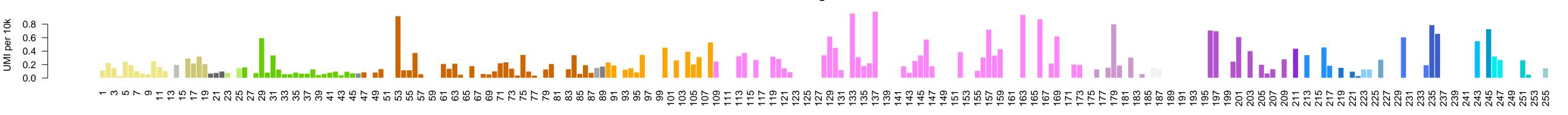
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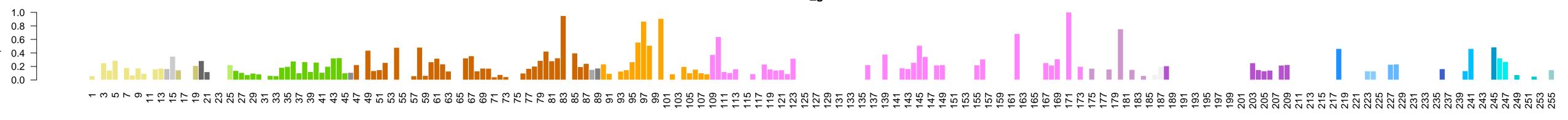
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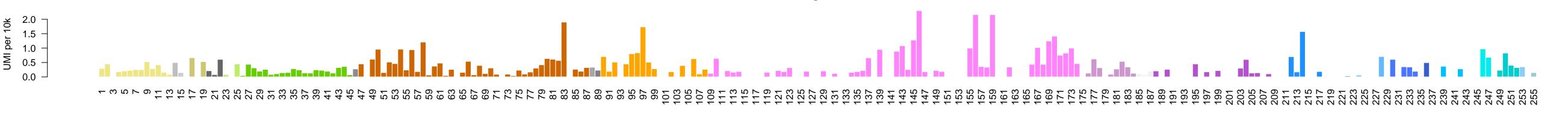
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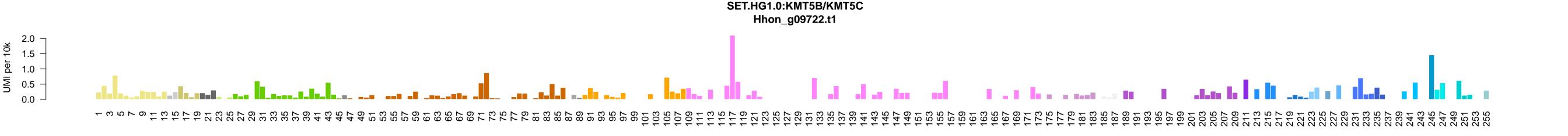


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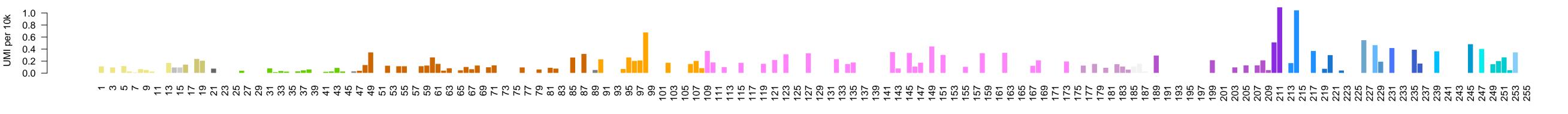


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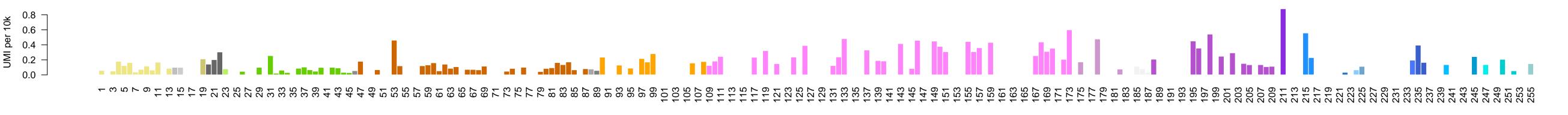




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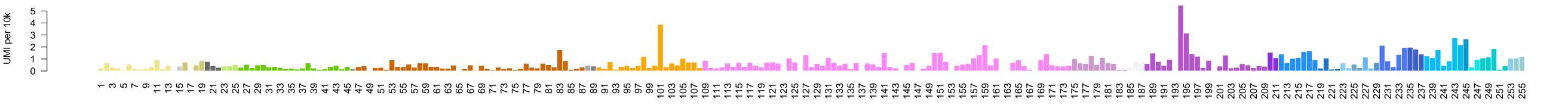


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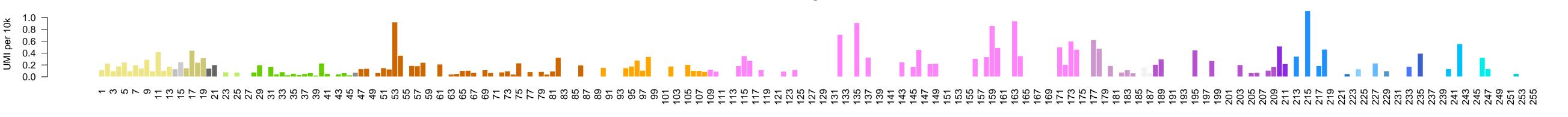


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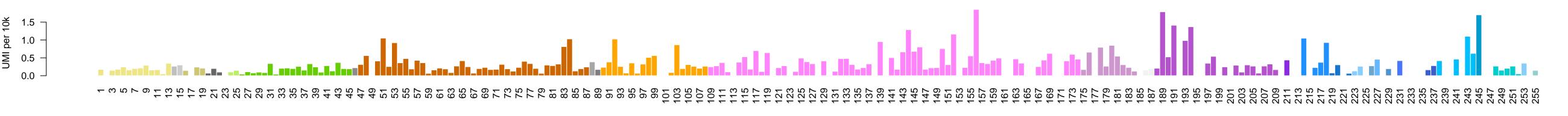
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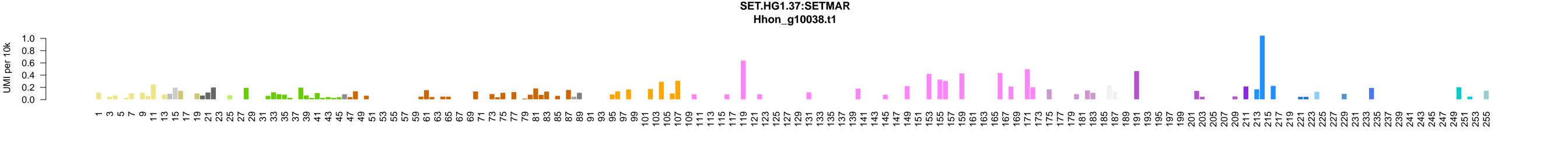


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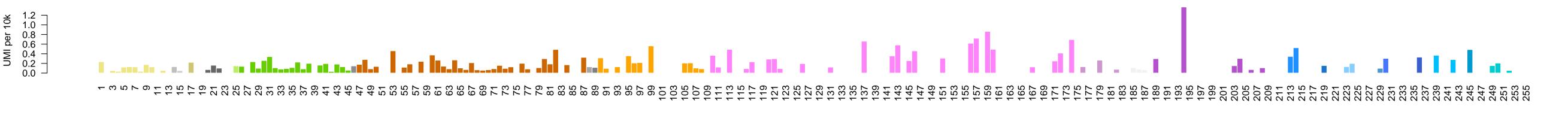


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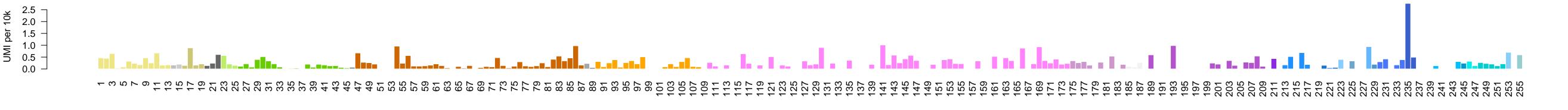




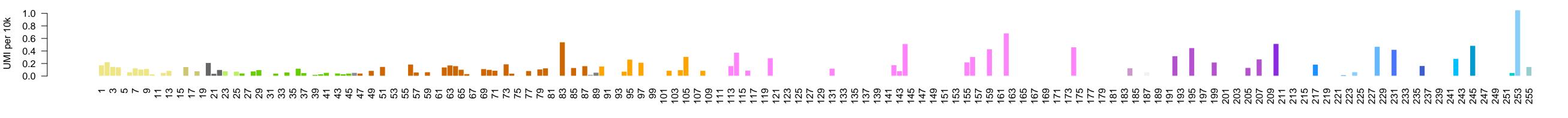
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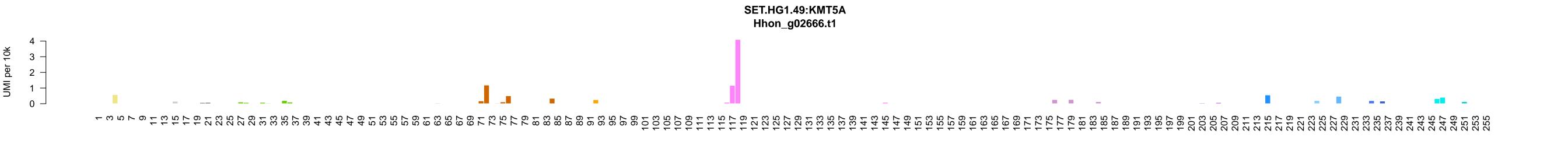


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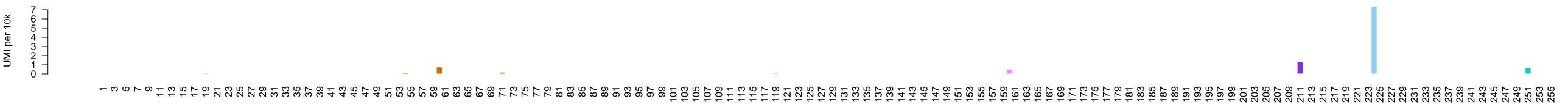


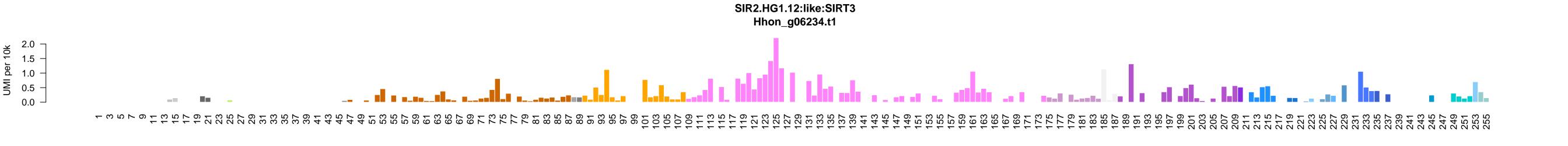
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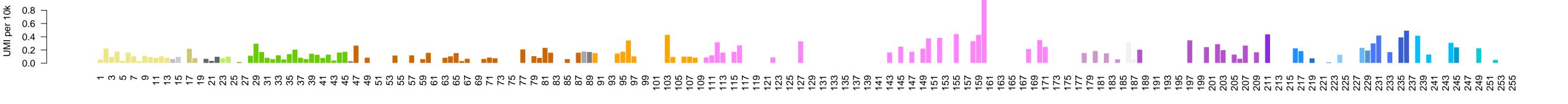
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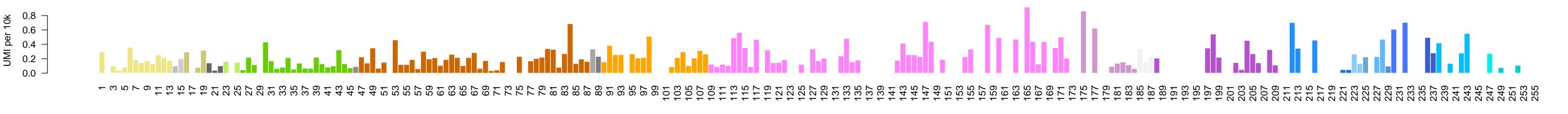




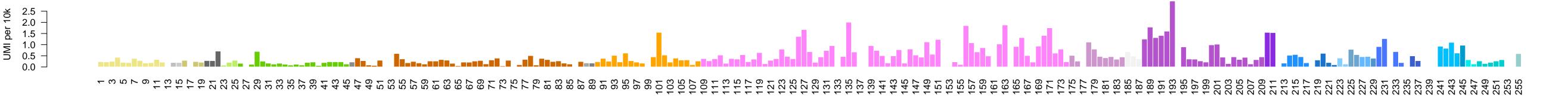
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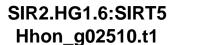


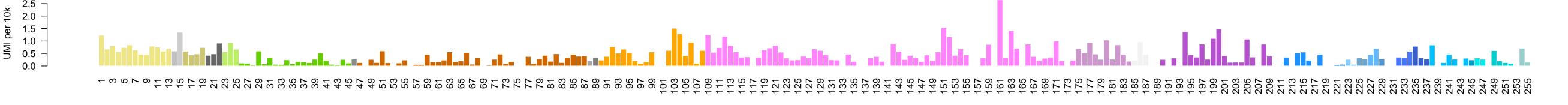
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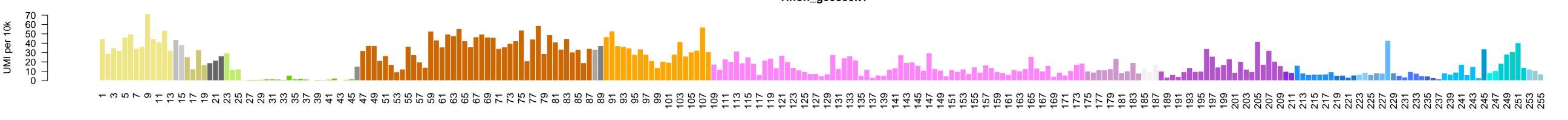




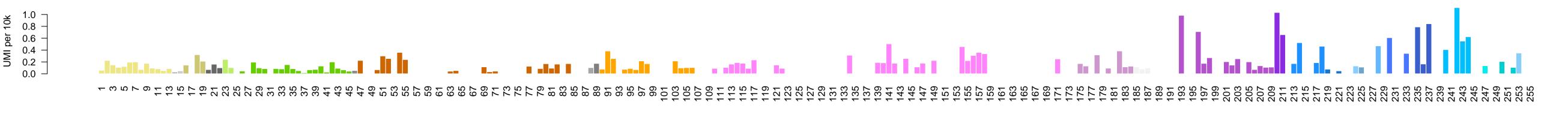
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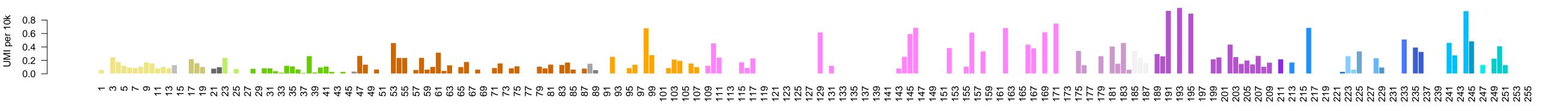
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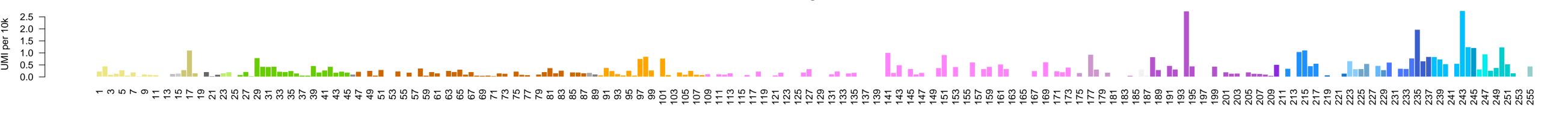
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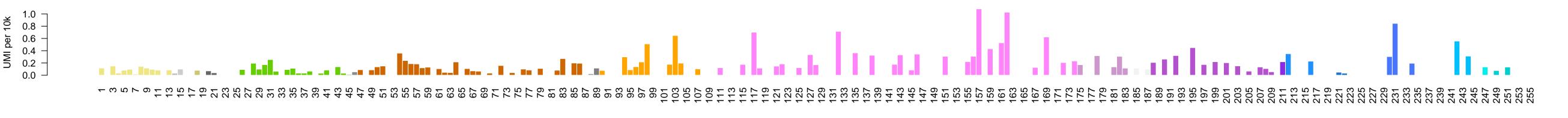
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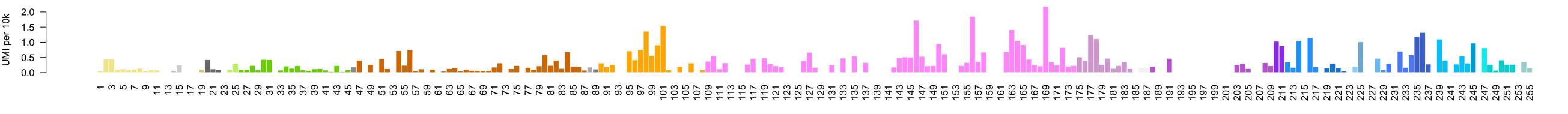
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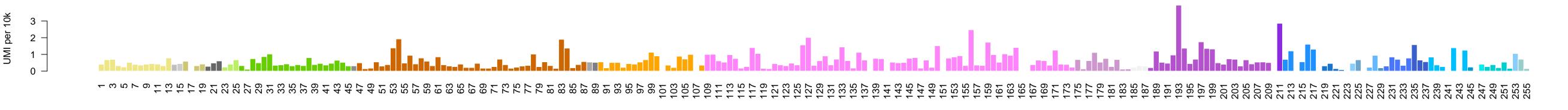
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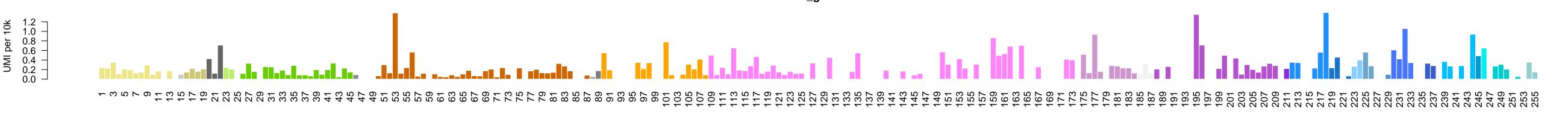
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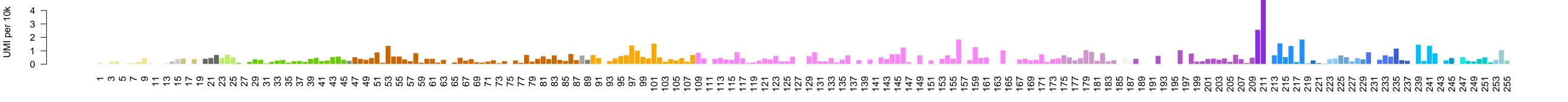
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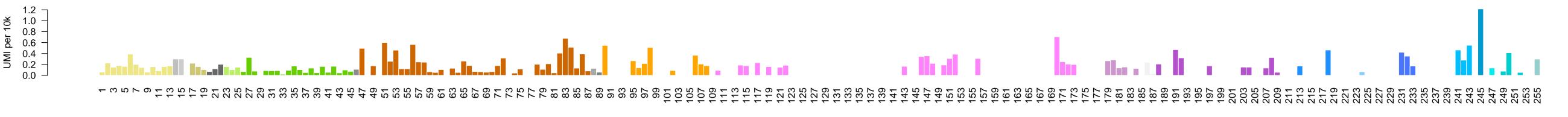
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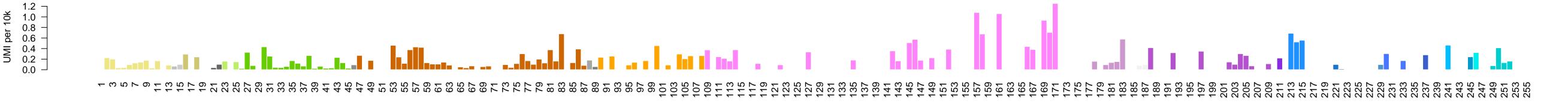
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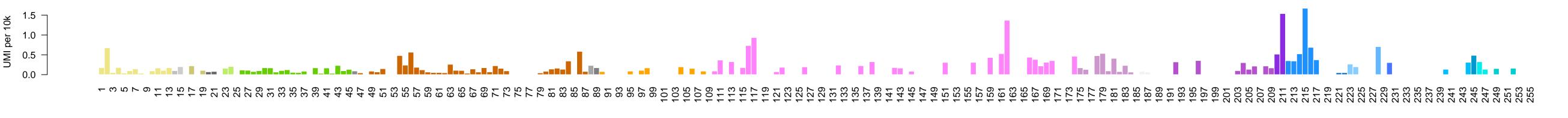
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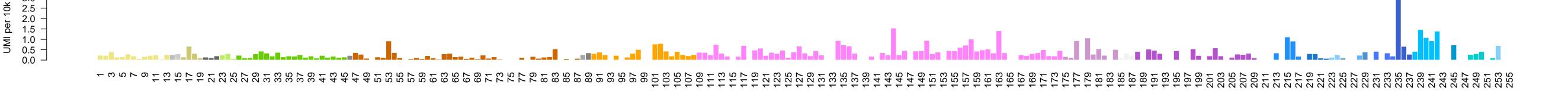
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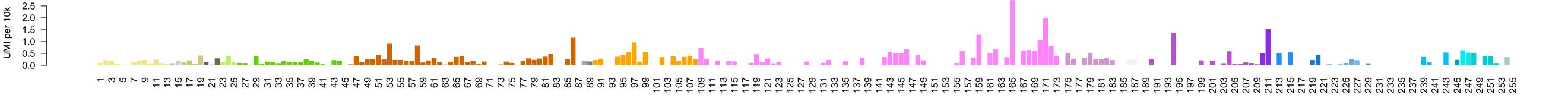
SWIRM.HG1.1:KDM1A Hhon_g10903.t1



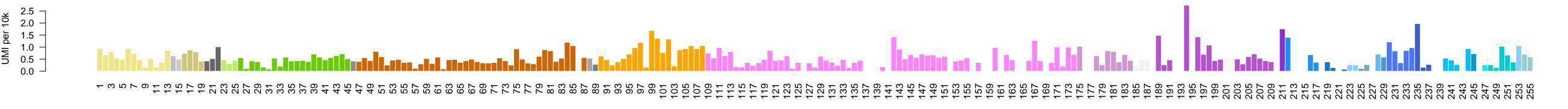


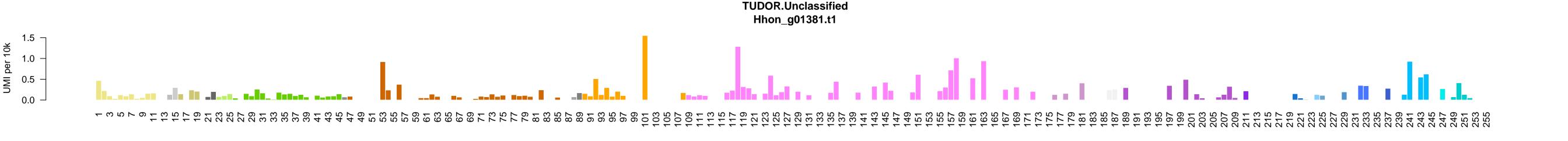


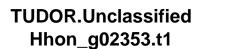
SWIRM.HG1.3:SMARCC1/SMARCC2 Hhon_g07010.t1

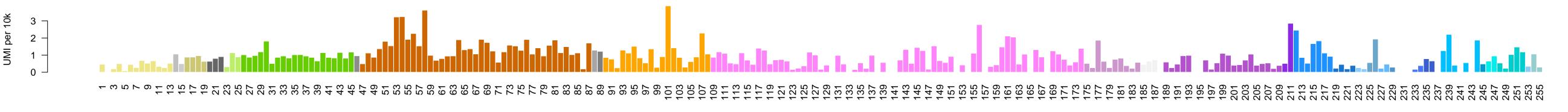




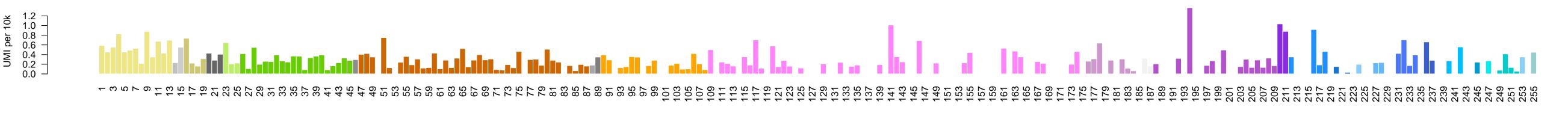




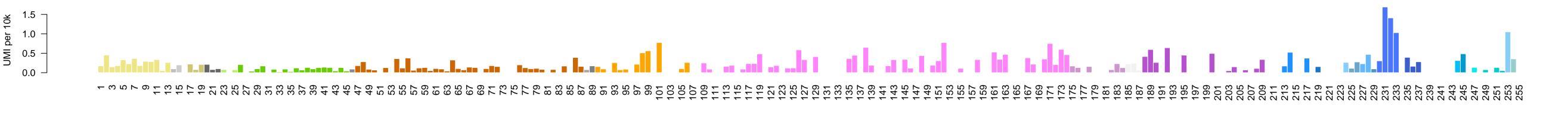




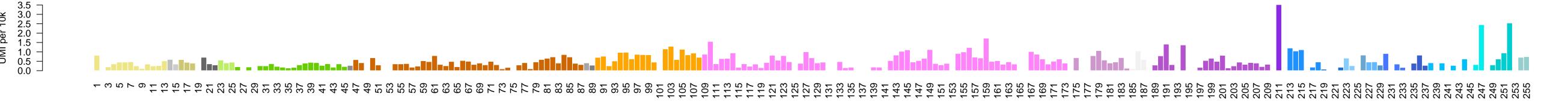
TUDOR.Unclassified Hhon_g04056.t1



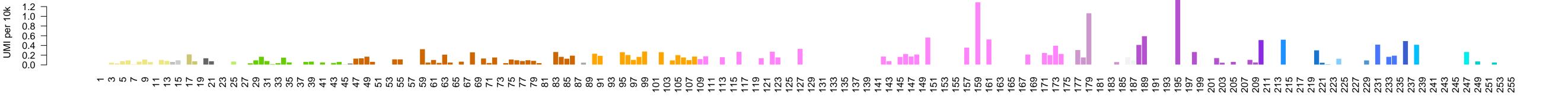
YEATS.HG1.0:MLLT1/MLLT3 Hhon_g05906.t1



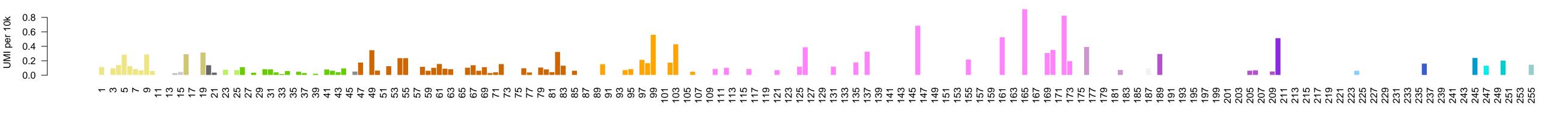
YEATS.HG1.1:YEATS4 Hhon_g01885.t1



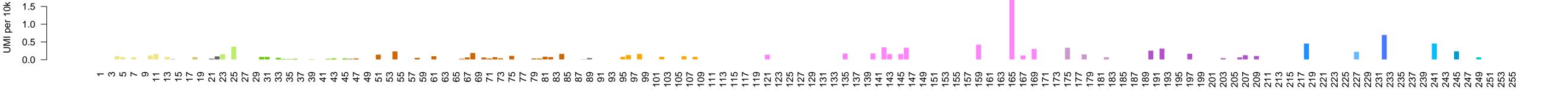
YEATS.HG1.2:YEATS2 Hhon_g04683.t1



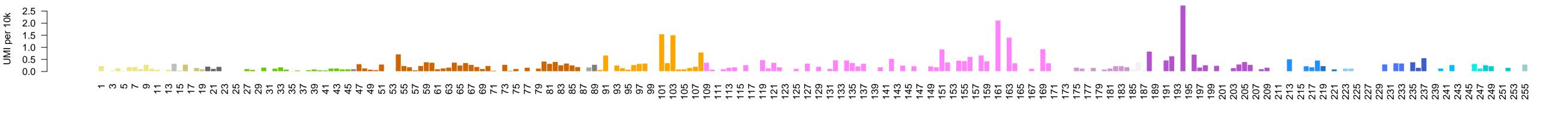
zf-CCHH.HG1.10:APLF Hhon_g07278.t1



zf-CCHH.HG1.2:like:APLF Hhon_g03071.t1



zf-CCHH.HG1.8:like:APLF Hhon_g06900.t1



zf-CCHH.HG2.0:CHFR Hhon_g02613.t1

