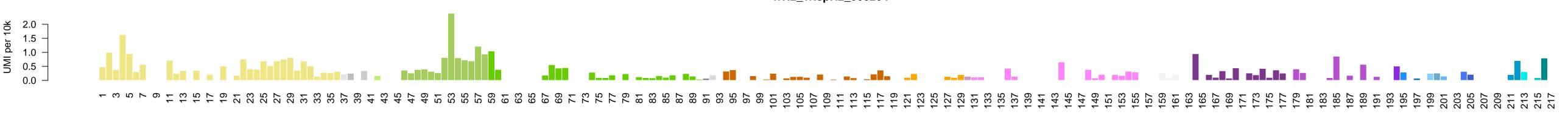
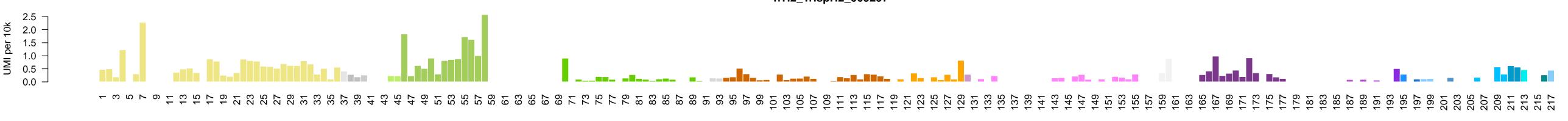


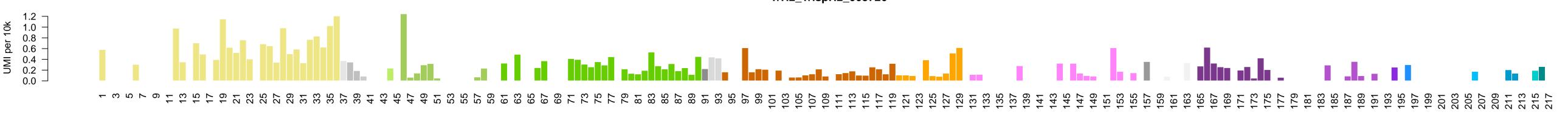
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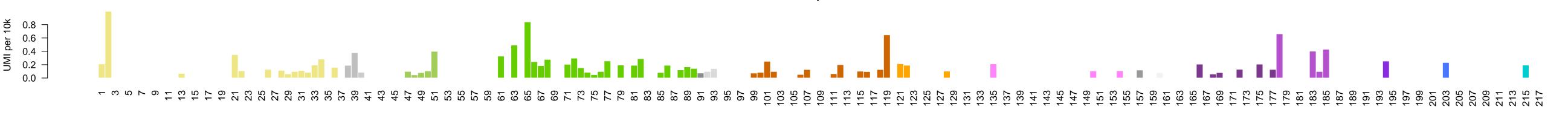
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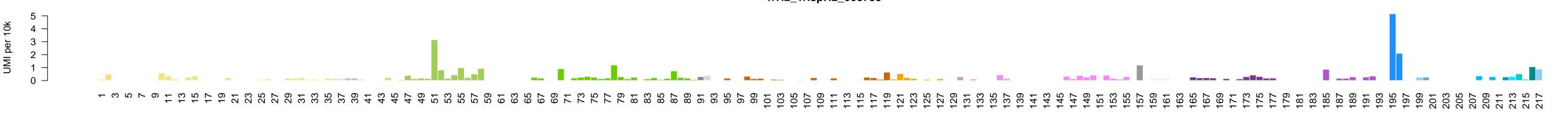
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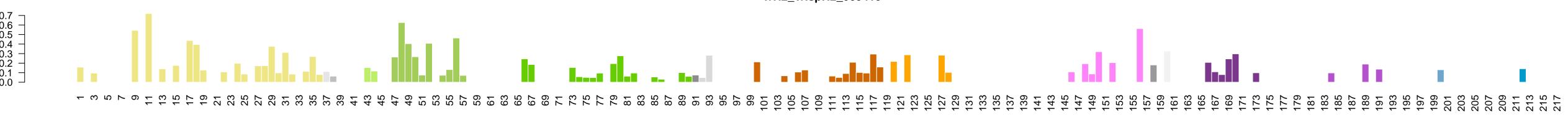
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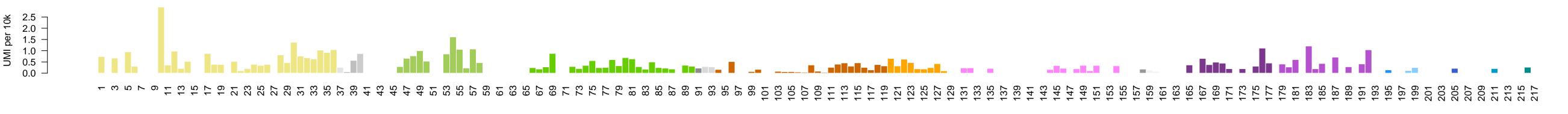
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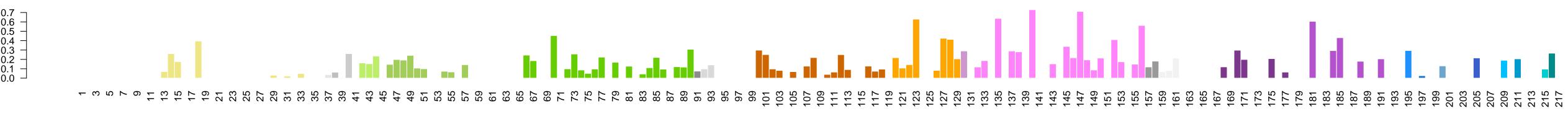
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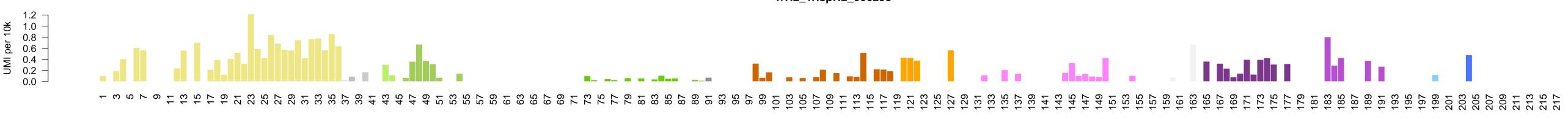
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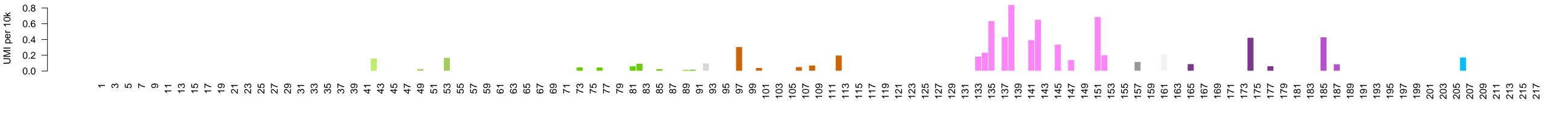
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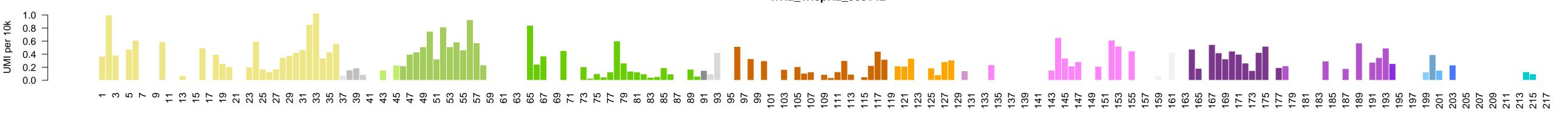
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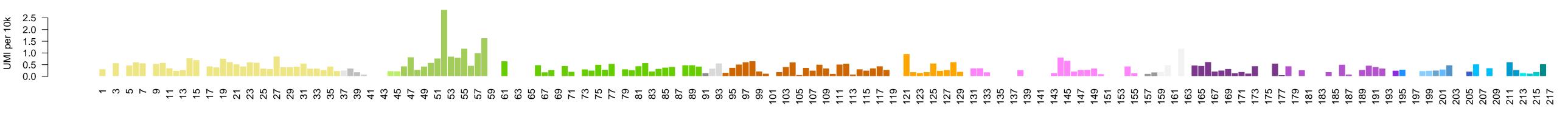
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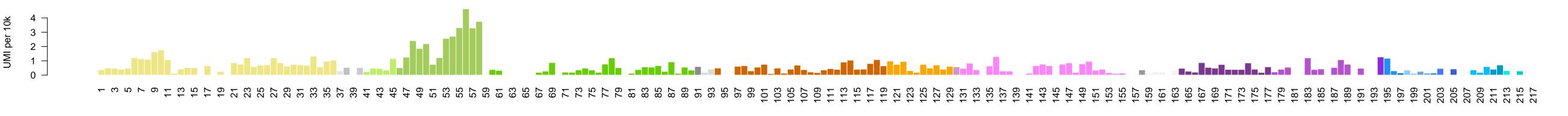
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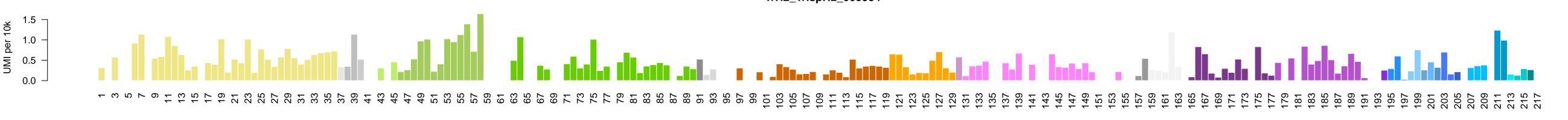
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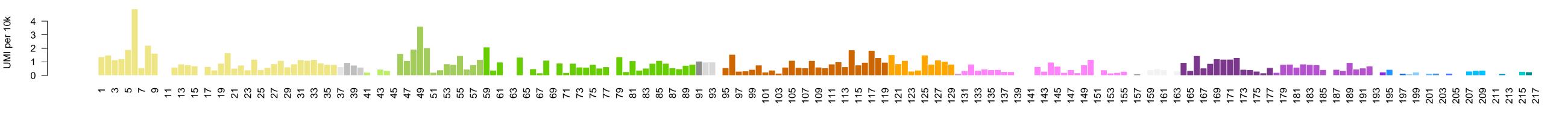
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Acetyltransf_1.HG1.8:NAA20 TrH2 TrispH2 005584

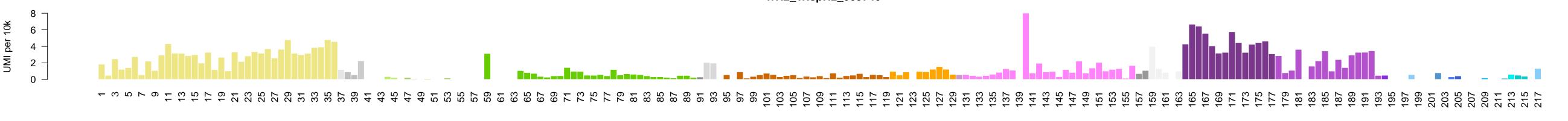


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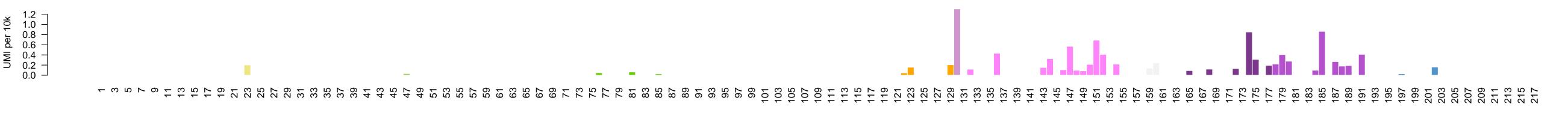


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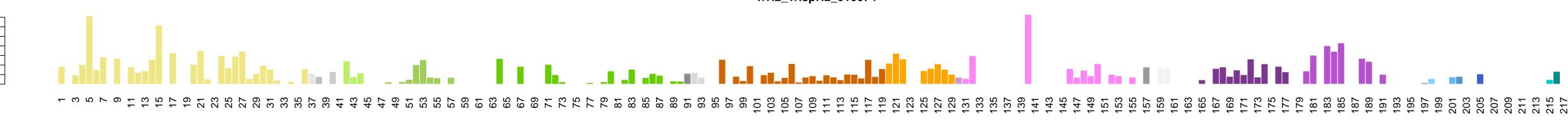
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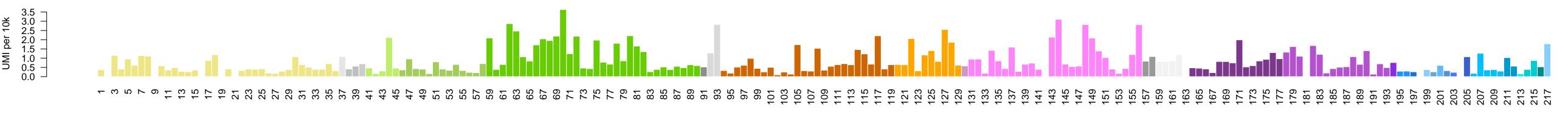
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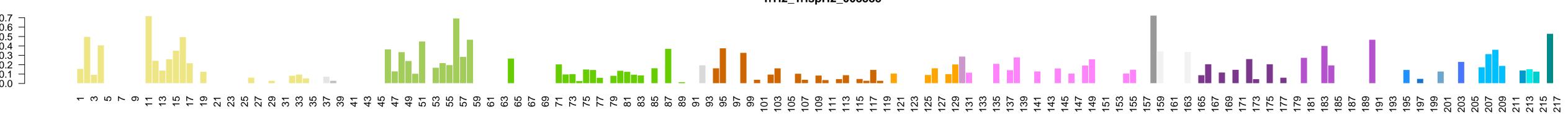
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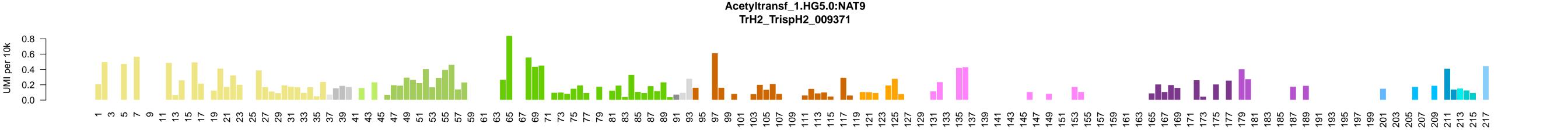


Acetyltransf_1.HG3.0:like:KAT2A/KAT2B TrH2_TrispH2_001370

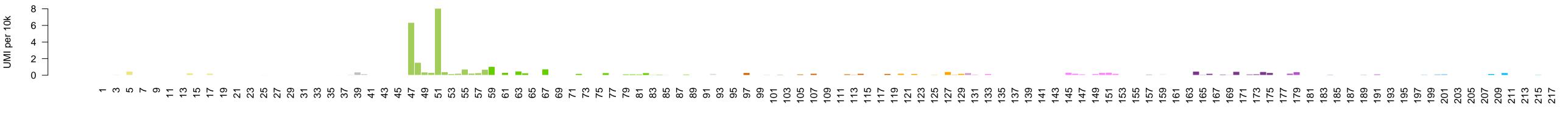


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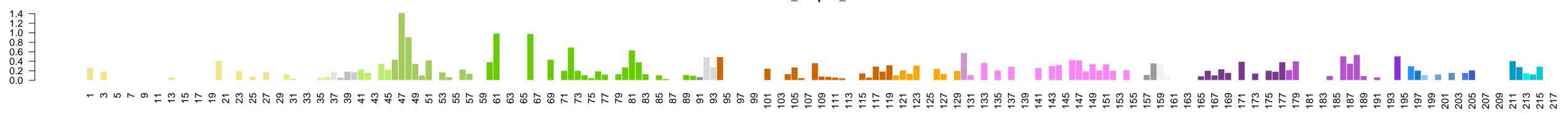




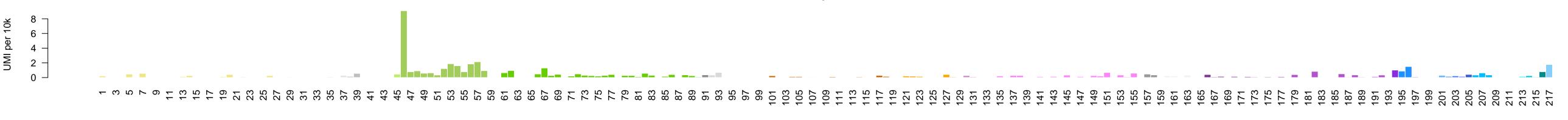
ASF1_hist_chap.HG1.0:ASF1A/ASF1B TrH2_TrispH2_006795



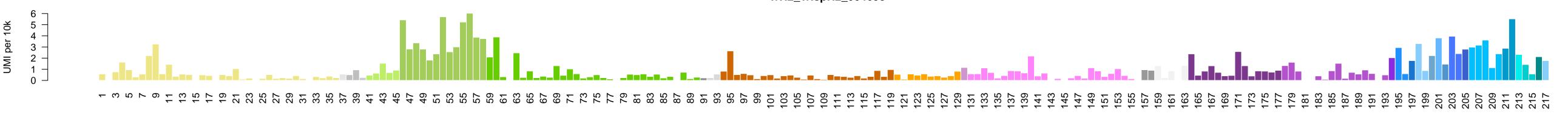
BAH.HG1.3:PBRM1;Bromodomain.HG1.29:PBRM1 TrH2_TrispH2_008263



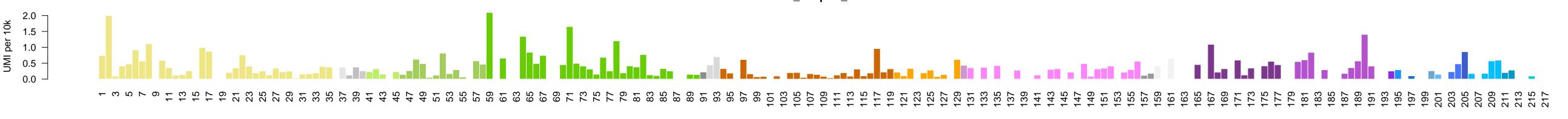
BAH.HG1.5:BAHD1 TrH2_TrispH2_003138



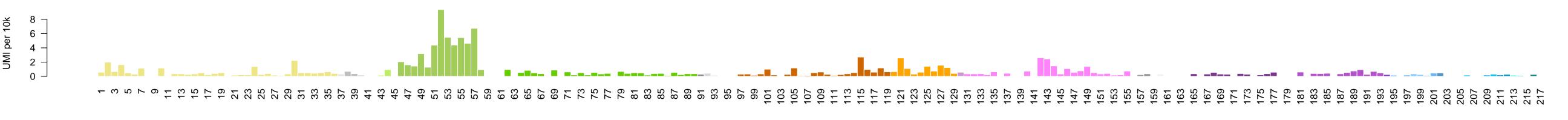
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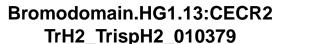


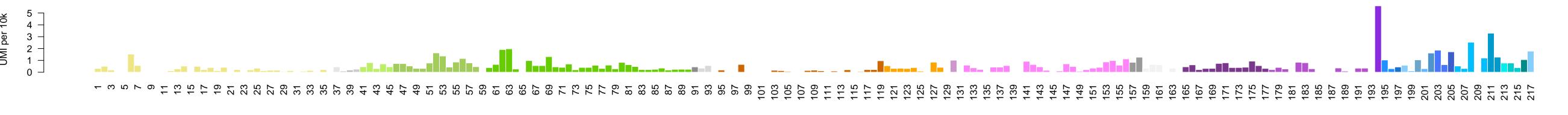
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TrH2_TrispH2_003261



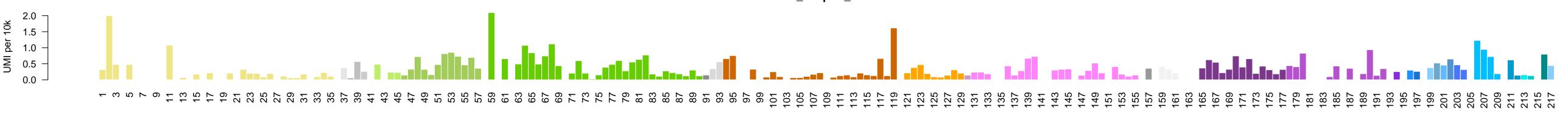
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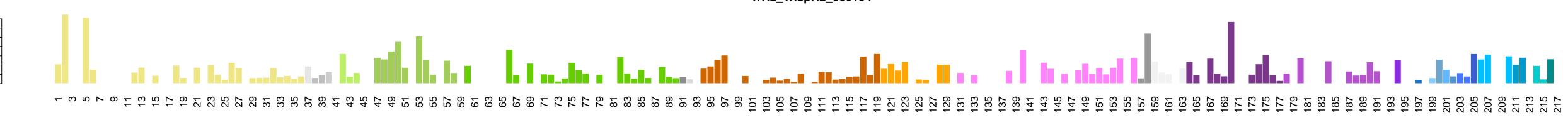




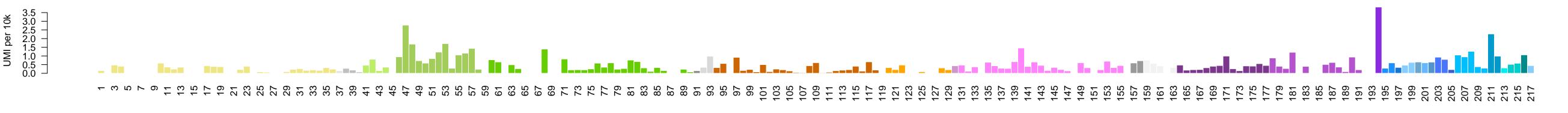
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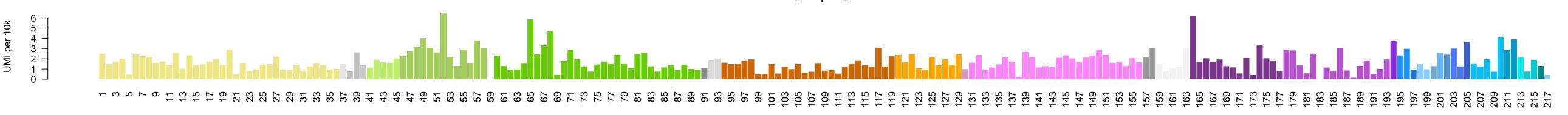
Bromodomain.HG1.17:BRD7/BRD9 TrH2 TrispH2 000194



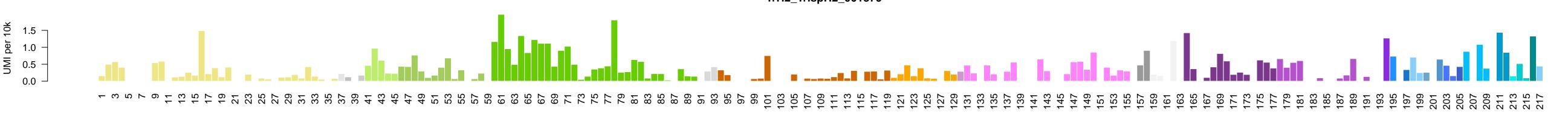
Bromodomain.HG1.18:BRD1/BRPF1/BRPF3;PWWP.HG1.12:BRD1/BRPF1/BRPF3 TrH2_TrispH2_004877



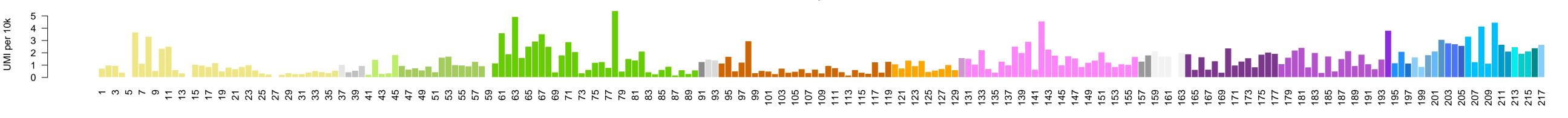
Bromodomain.HG1.20:BRD2/BRD3/BRD4/BRDT TrH2_TrispH2_003484



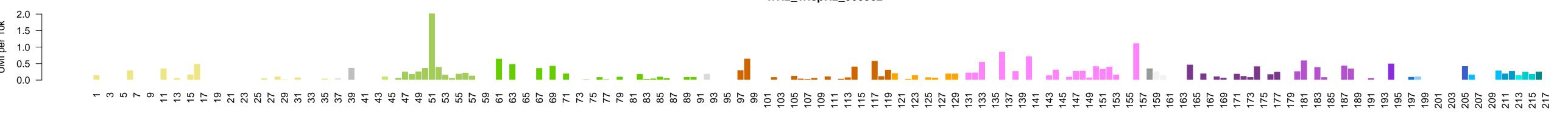
Bromodomain.HG1.21:BRWD1/BRWD3/PHIP TrH2_TrispH2_001879



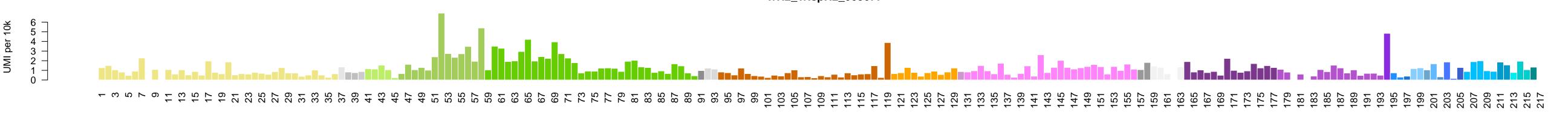
Bromodomain.HG1.25:CREBBP/EP300;KAT11.HG1.2:CREBBP/EP300 TrH2_TrispH2_009021



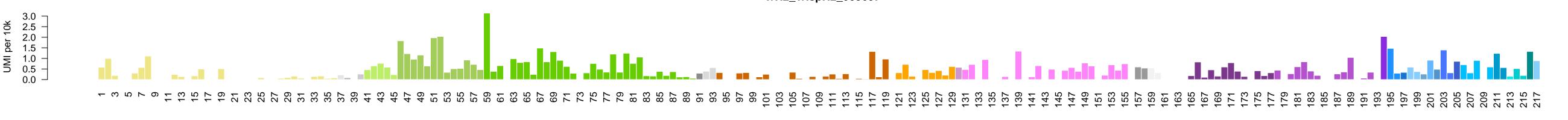
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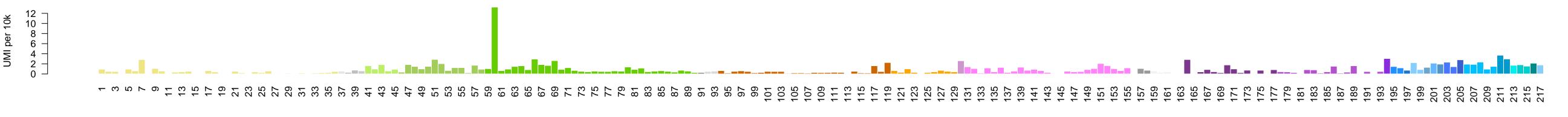
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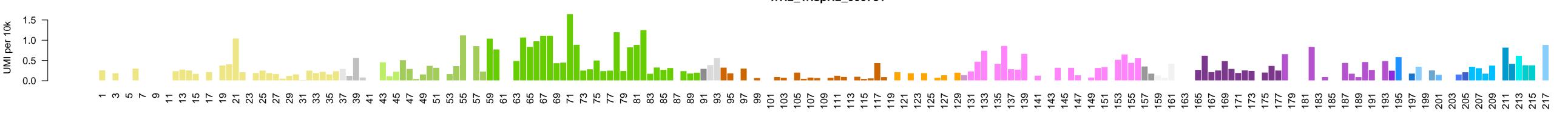
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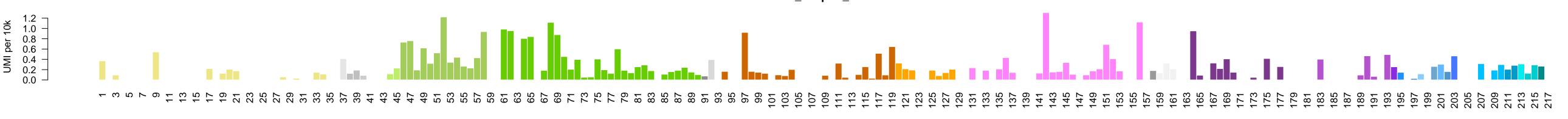
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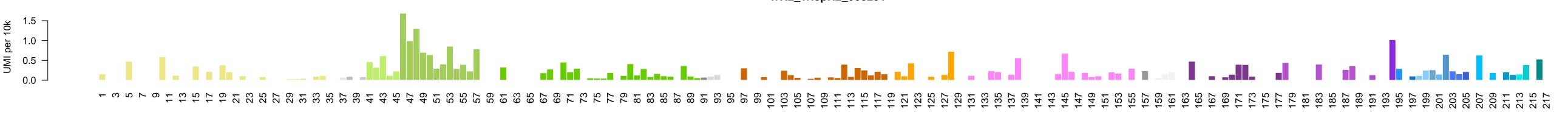
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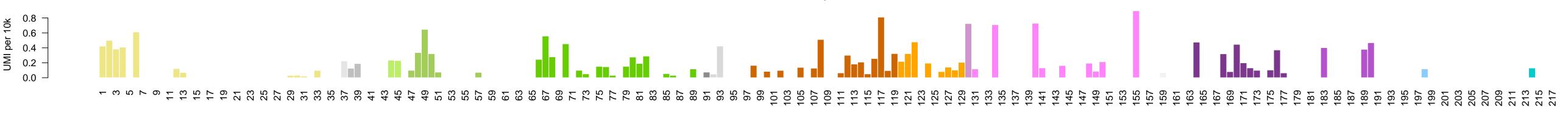
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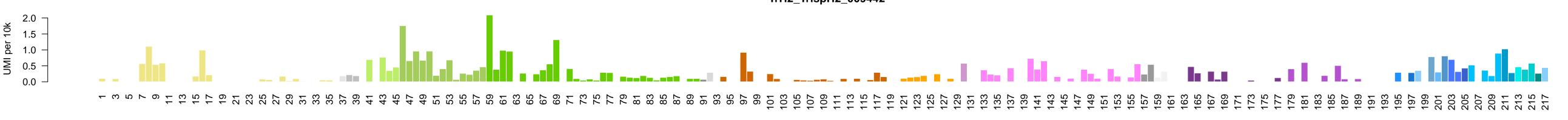
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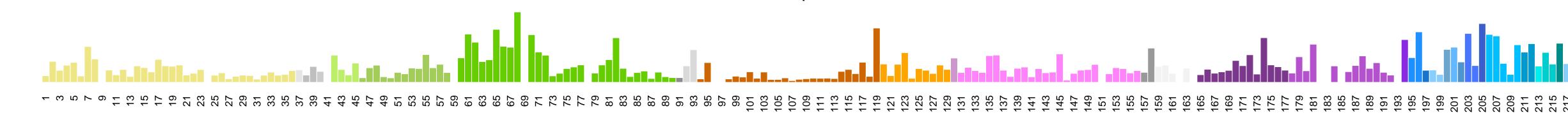
CAF1A.HG1.0:CHAF1A TrH2_TrispH2_010973



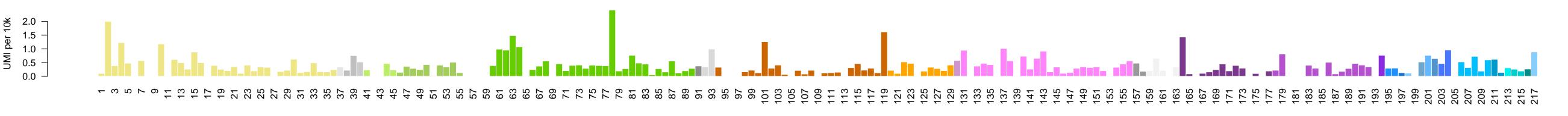
Chromo.HG1.12:AC006064.6/CHD3/CHD4/CHD5;SNF2_N.HG1.26:AC006064.6/CHD3/CHD4/CHD5
TrH2_TrispH2_009442



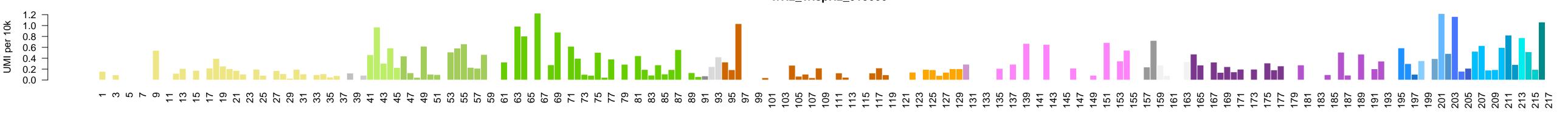
Chromo.HG1.13:CHD6/CHD7/CHD8/CHD9;SNF2_N.HG1.28:CHD6/CHD7/CHD8/CHD9 TrH2_TrispH2_000692



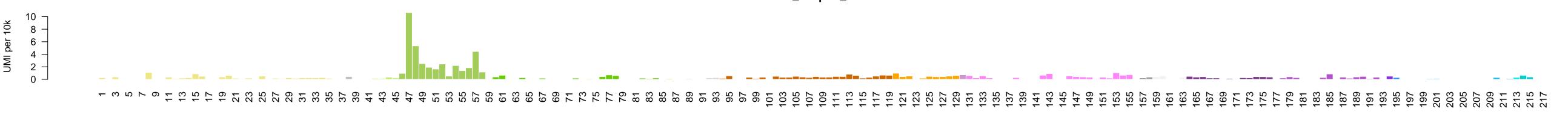
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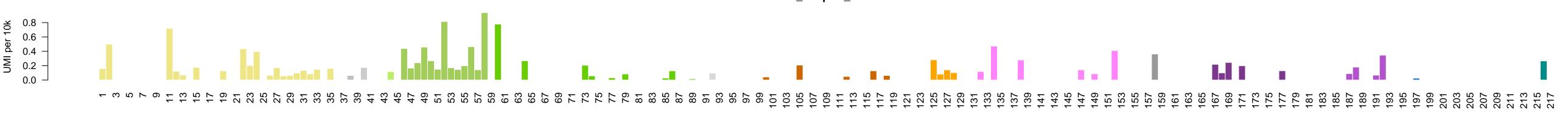
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TrH2_TrispH2_010666

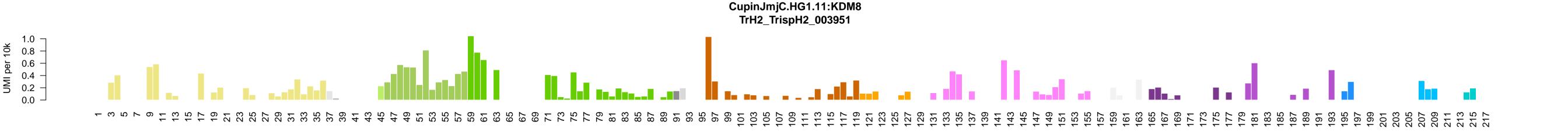


Chromo.HG1.6:SUV39H1/SUV39H2 TrH2_TrispH2_010891

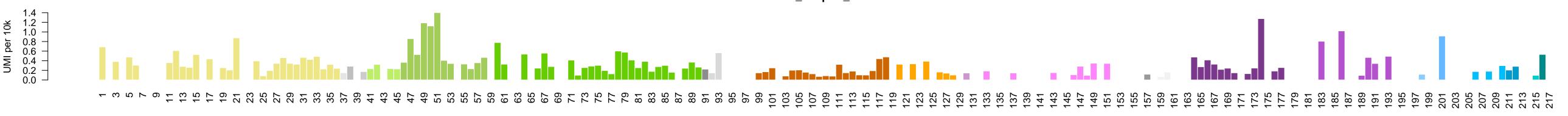


CupinJmjC.HG1.1:HSPBAP1 TrH2_TrispH2_011284

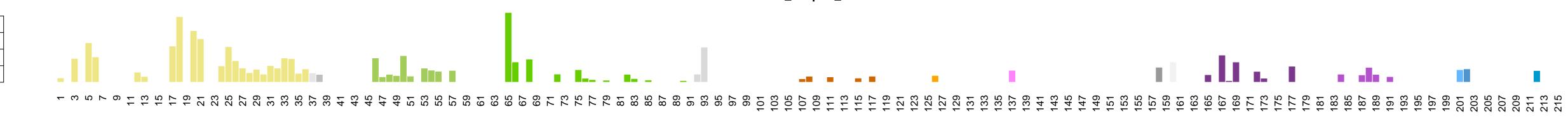




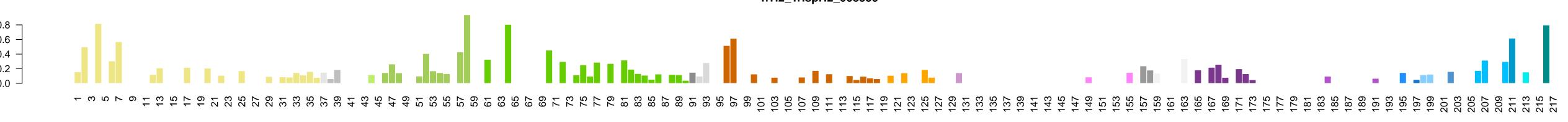
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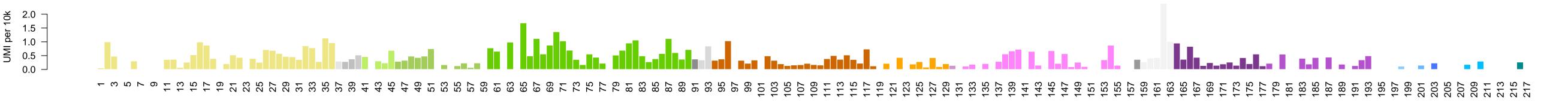
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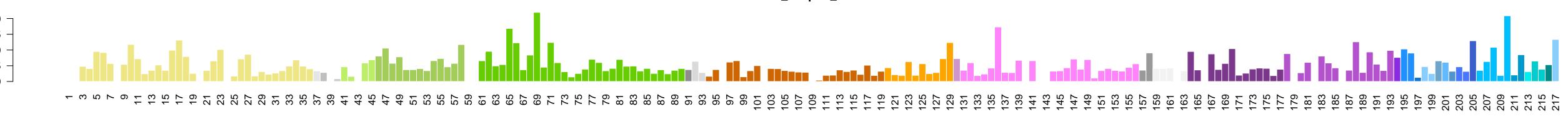
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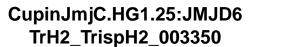


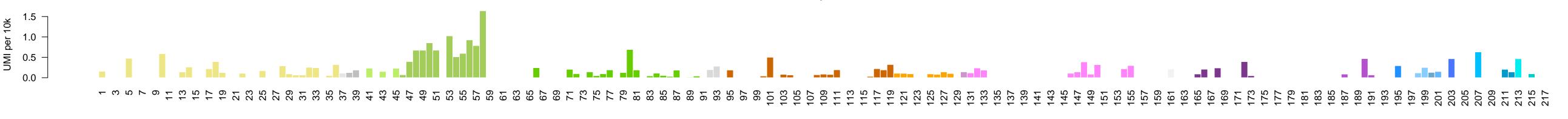
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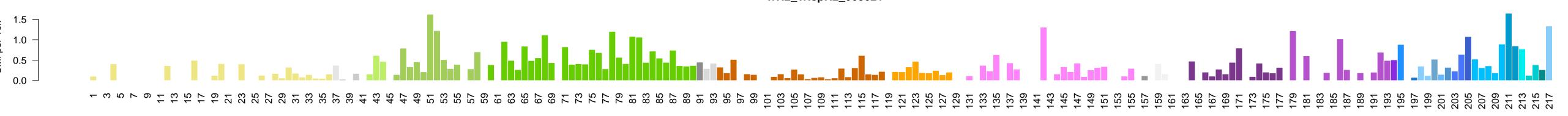
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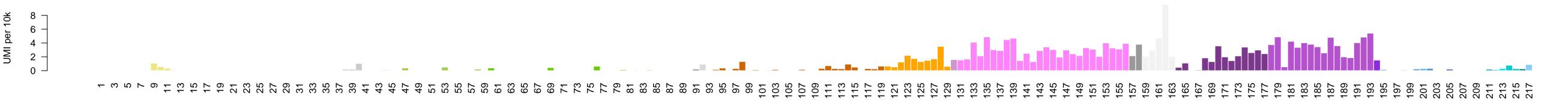




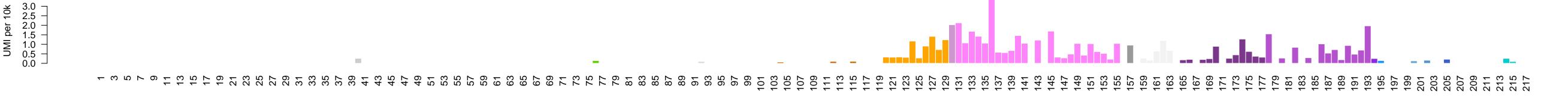
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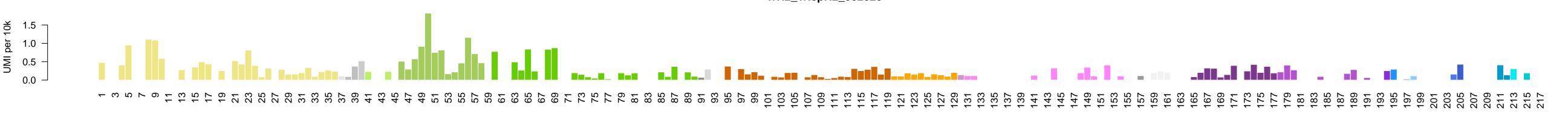
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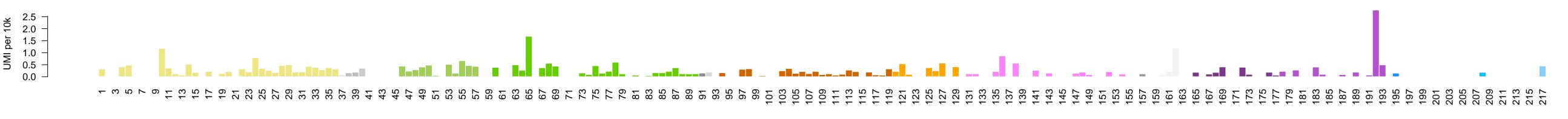
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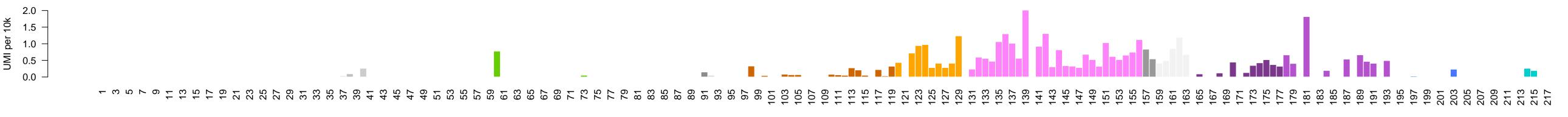
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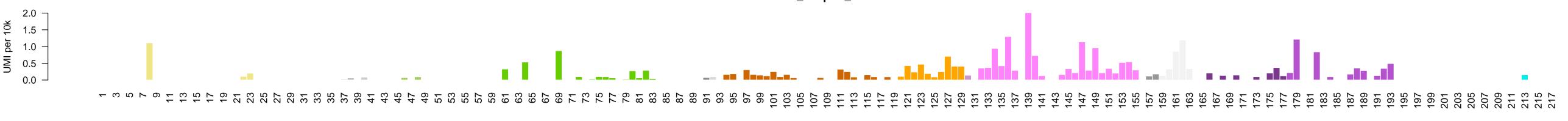
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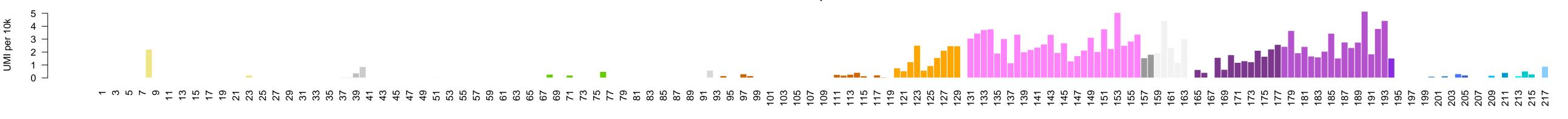
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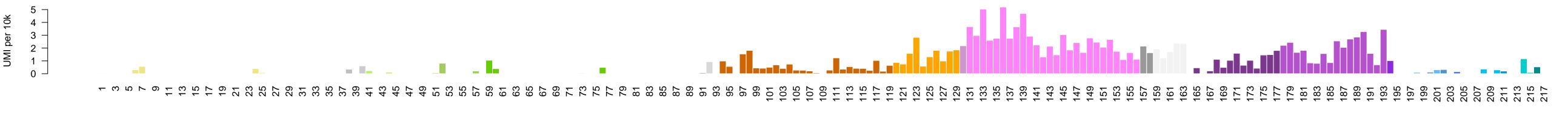
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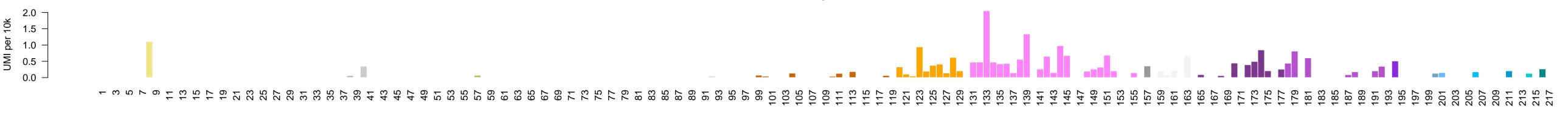
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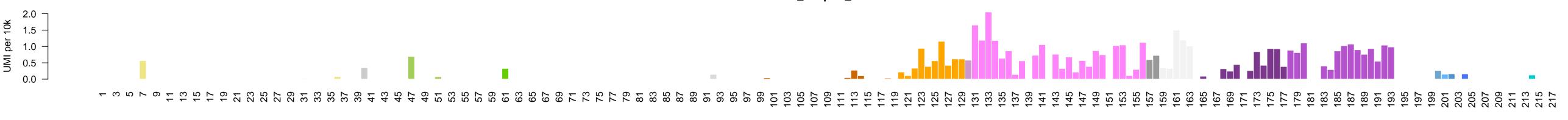
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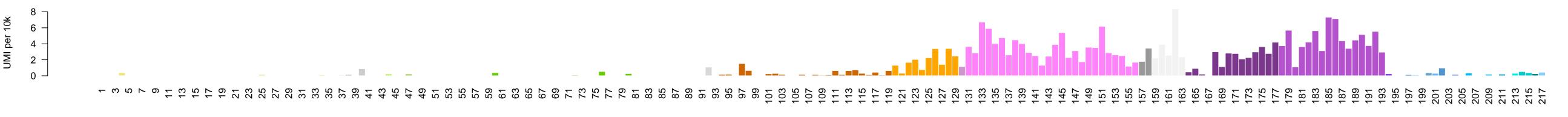
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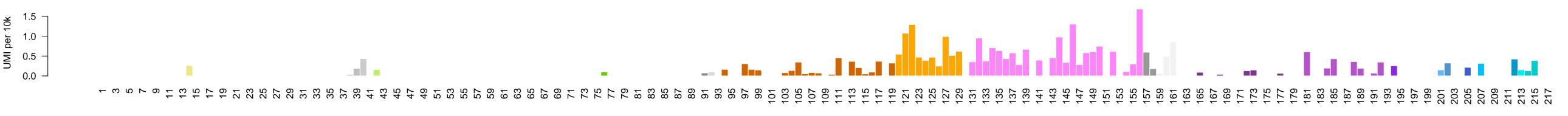
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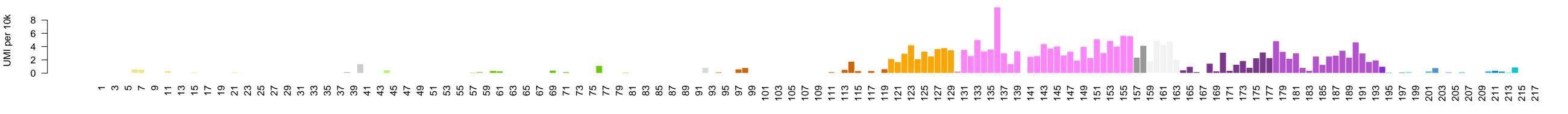
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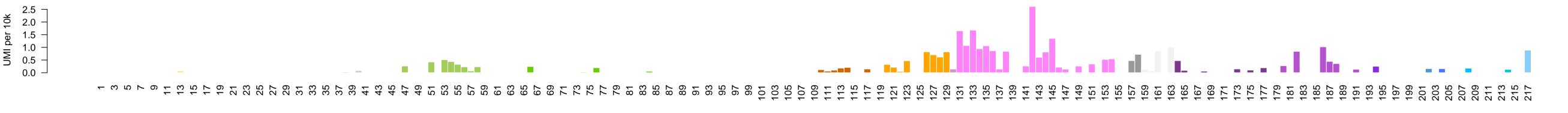
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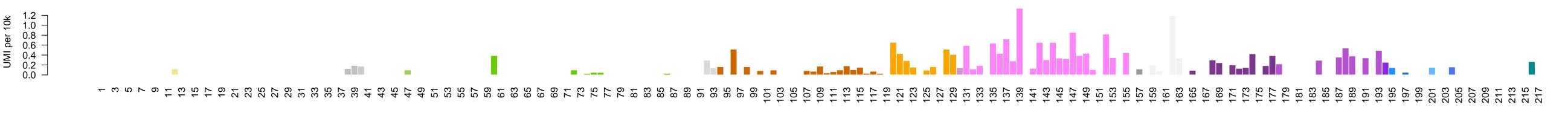
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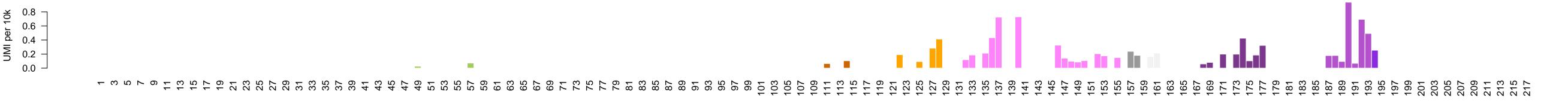
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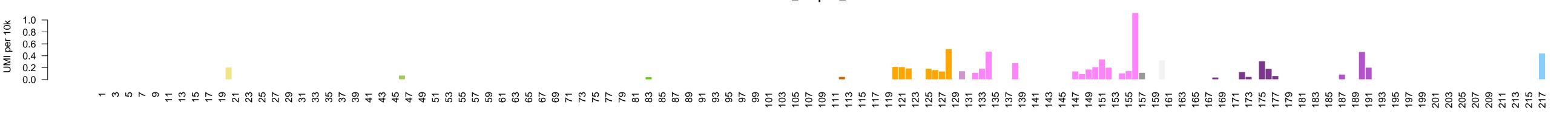
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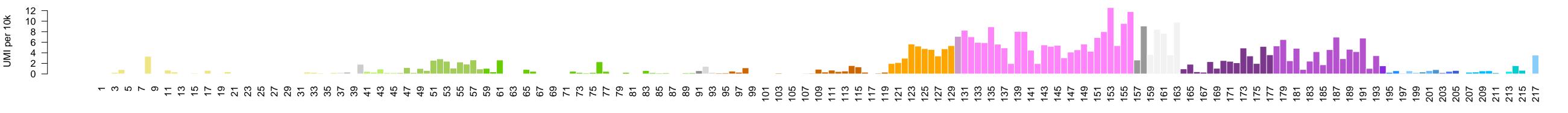
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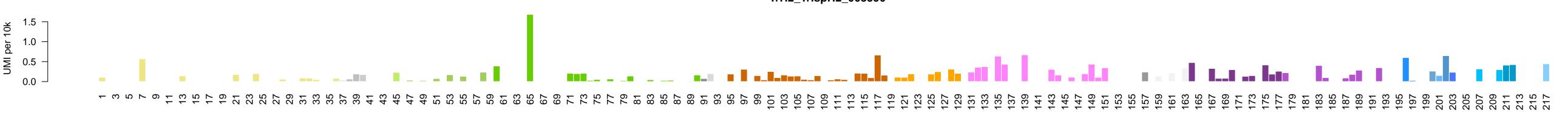
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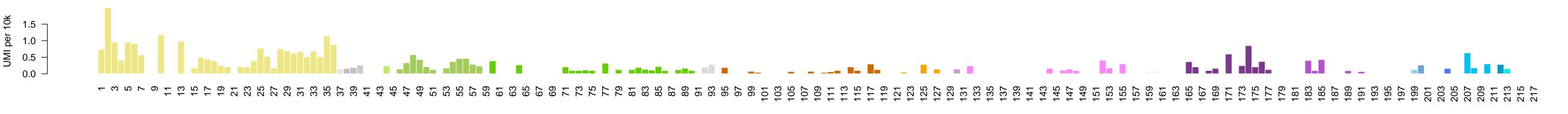
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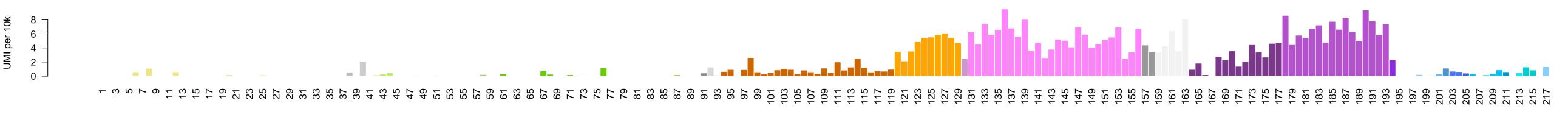
CupinJmjC.HG1.6:JMJD1C/KDM3A/KDM3B TrH2 TrispH2 003590



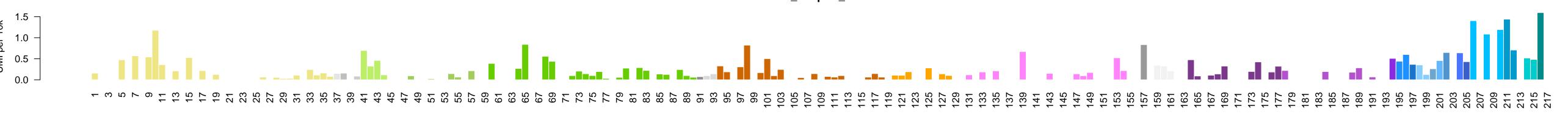
CupinJmjC.HG1.7:like:KDM2A/KDM2B/KDM7A/PHF2/PHF8 TrH2_TrispH2_003357



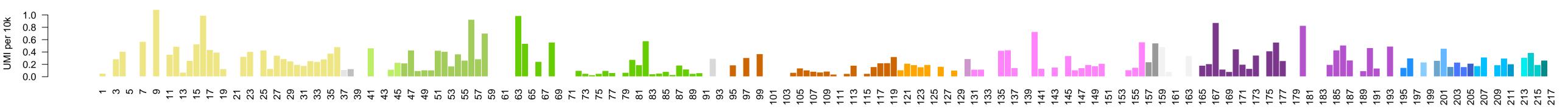
CupinJmjC.HG1.8:like:JMJD4/JMJD6/JMJD8 TrH2_TrispH2_001540



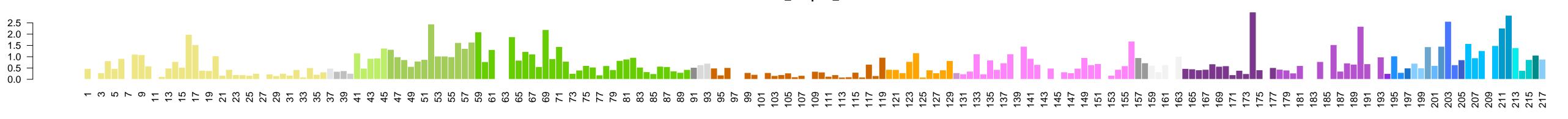
CupinJmjC.HG2.0:JARID2 TrH2_TrispH2_001569



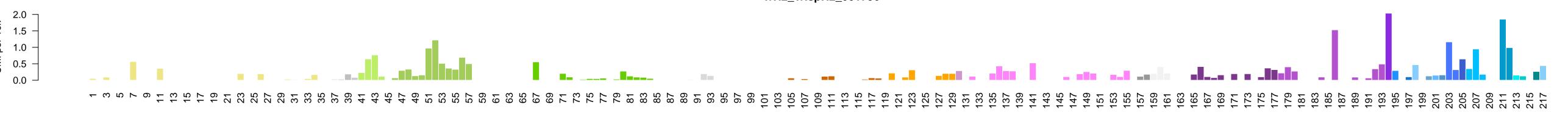
CupinJmjC.HG2.2:AL451062.2/KDM4A/KDM4B/KDM4C/KDM4D/KDM4E/KDM4F TrH2_TrispH2_001378



CupinJmjC.HG2.3:KDM5A/KDM5B/KDM5C/KDM5D TrH2_TrispH2_004411

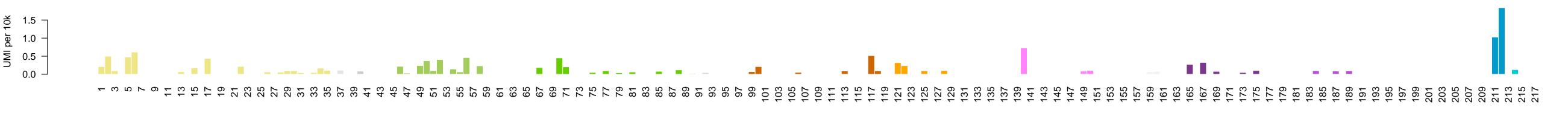


CupinJmjC.Unclassified TrH2_TrispH2_001736

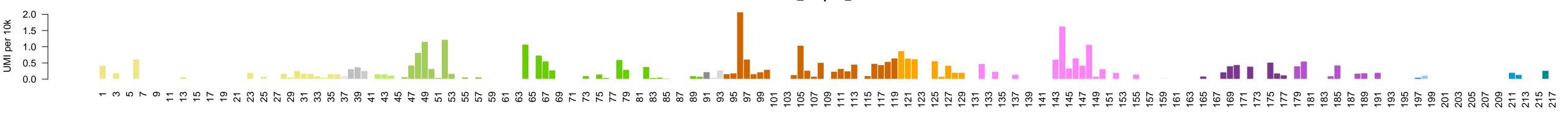




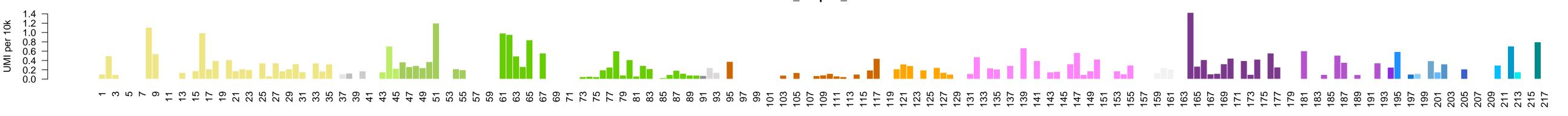
GNAT_acetyltr_2.HG1.0:NAT10 TrH2_TrispH2_000898



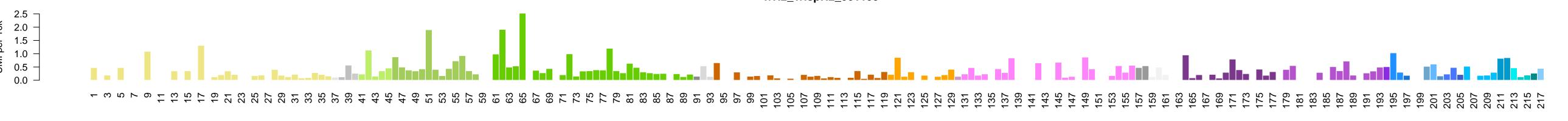
Hat1_N.HG1.0:HAT1;MOZ_SAS.HG1.0:like:KAT5/KAT6A/KAT6B/KAT7/KAT8 TrH2_TrispH2_002269



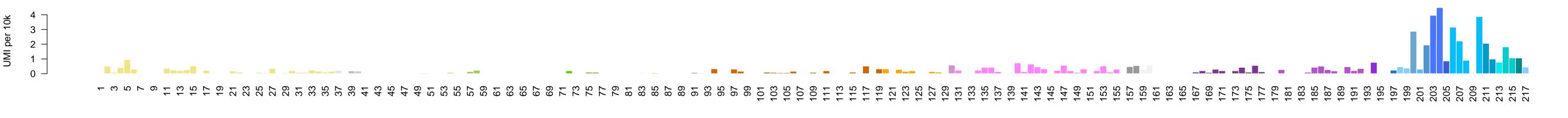
Hira.HG1.0:HIRA TrH2 TrispH2 008630



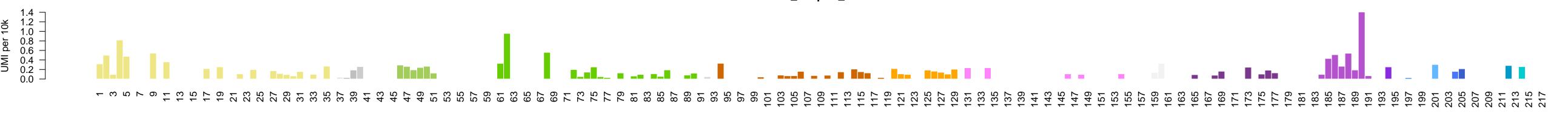
Hist_deacetyl.HG1.1:HDAC11 TrH2 TrispH2 001159



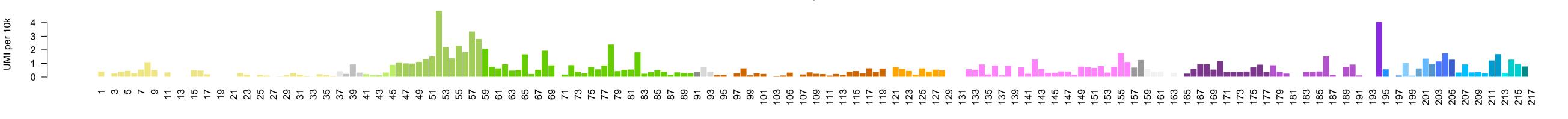
Hist_deacetyl.HG1.4:HDAC4/HDAC5/HDAC7/HDAC9
TrH2_TrispH2_005725

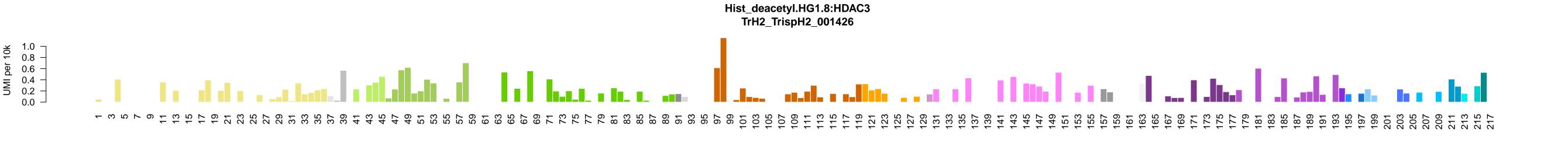


Hist_deacetyl.HG1.5:AL133500.1/HDAC8 TrH2_TrispH2_005857

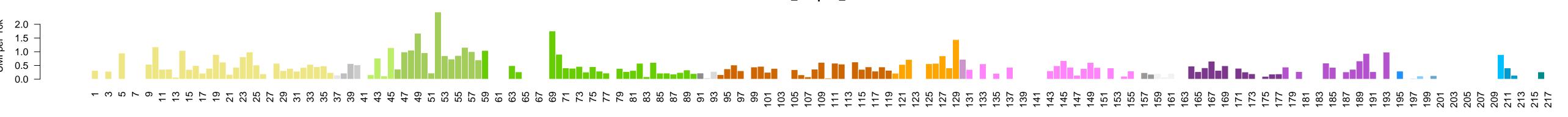


Hist_deacetyl.HG1.7:HDAC1/HDAC2 TrH2_TrispH2_006473

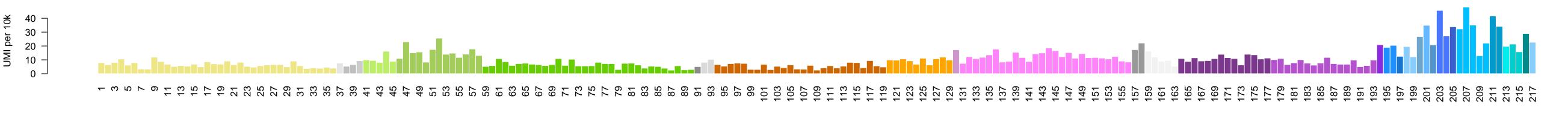




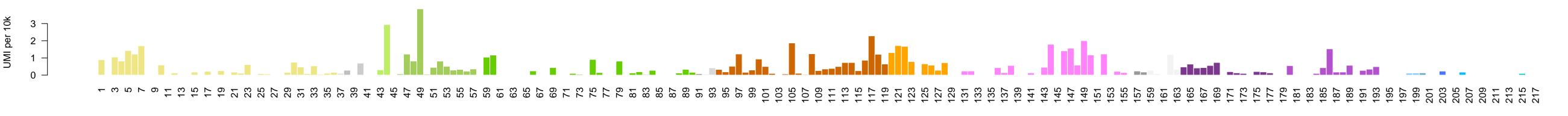
Histone.HG2.1:CENPA TrH2 TrispH2 009346

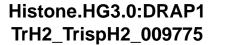


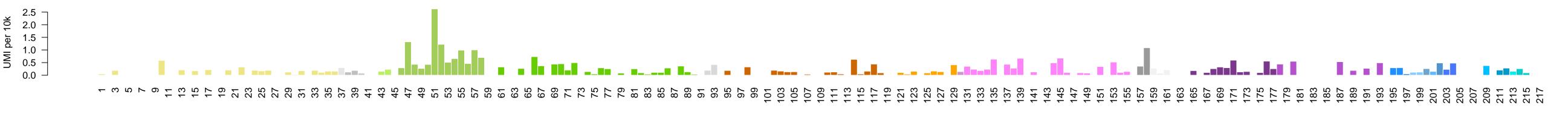
Histone.HG2.13:like:H3-3A/H3-3B/H3-5/H3Y1/H3Y2 TrH2_TrispH2_004552



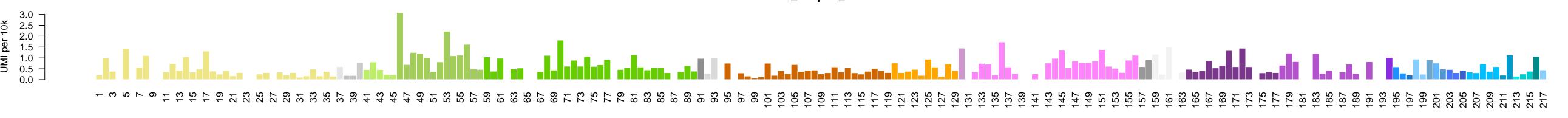
Histone.HG2.16:H3C13/H3C14 TrH2_TrispH2_005838



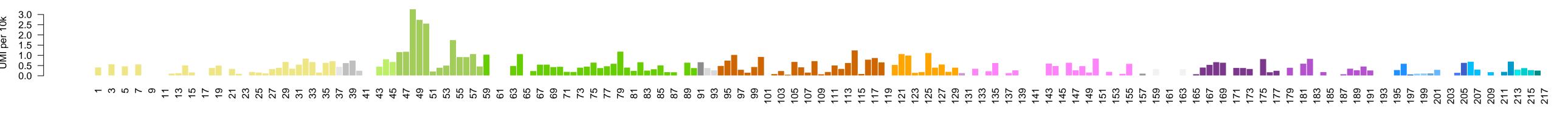




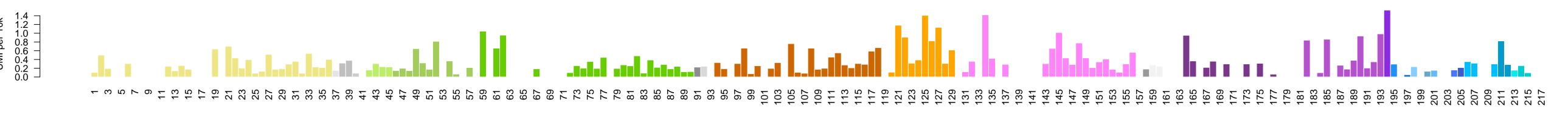
Histone.HG3.1:NFYC
TrH2 TrispH2 004524



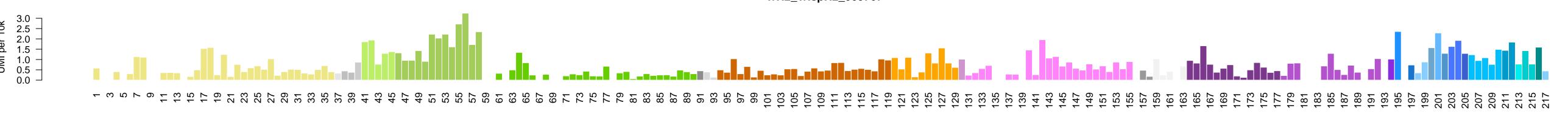
Histone.HG3.3:POLE4
TrH2_TrispH2_005489



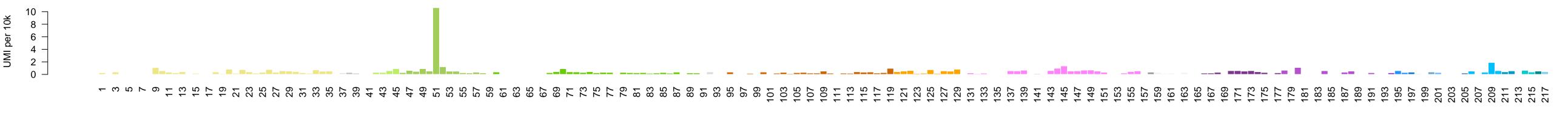
Histone.HG3.4:CHRAC1 TrH2 TrispH2 003697



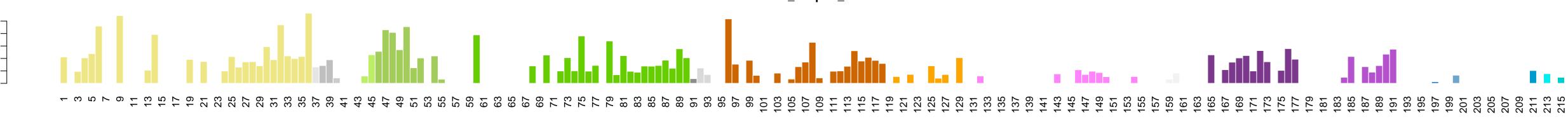
Histone.HG4.1:NFYB TrH2 TrispH2 005767

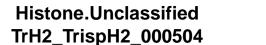


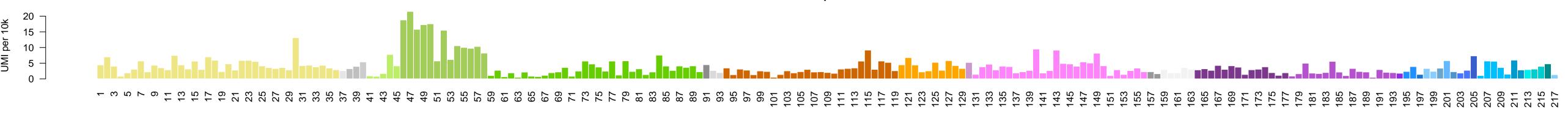
Histone.HG4.2:DR1
TrH2_TrispH2_002518

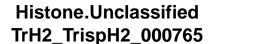


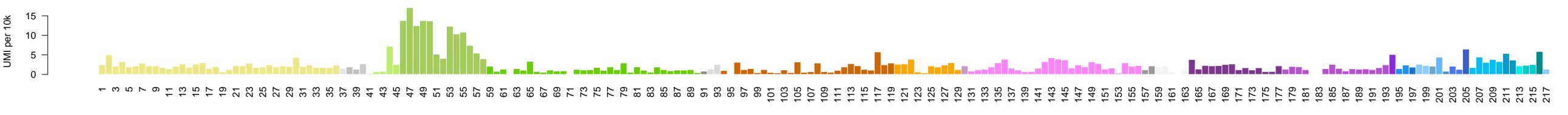
Histone.HG5.1:CENPS/CENPS-CORT TrH2_TrispH2_005420



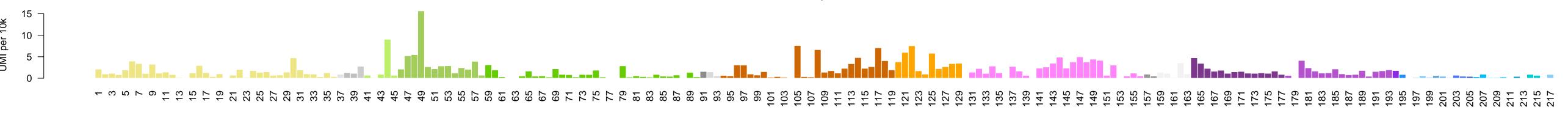




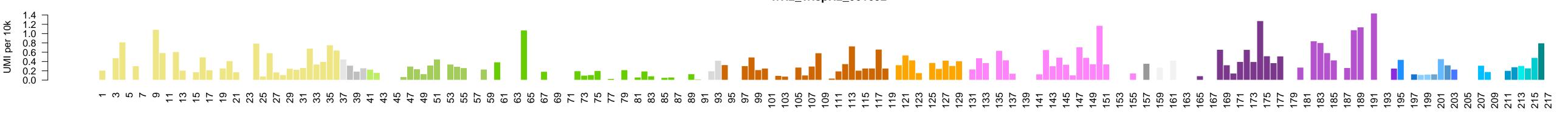




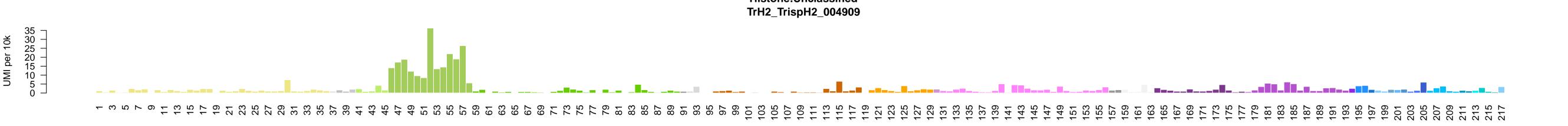




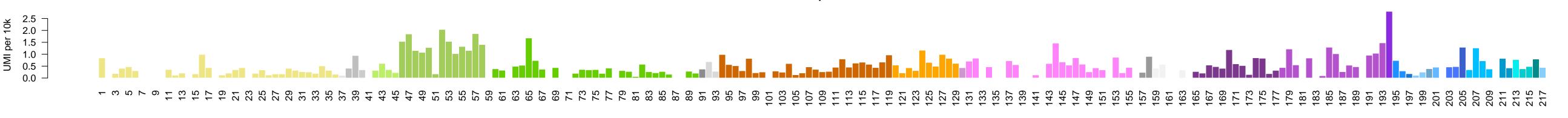
Histone.Unclassified TrH2 TrispH2 001652



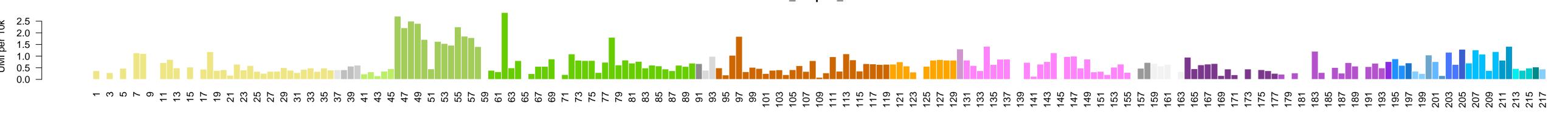
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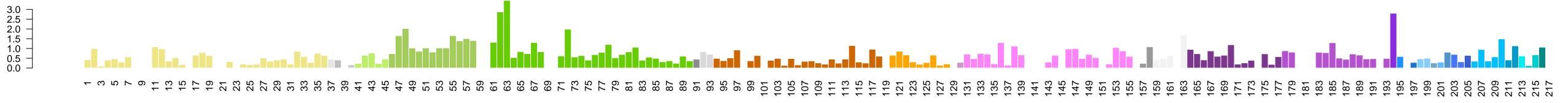
ING.HG1.1:ING3
TrH2 TrispH2 000072



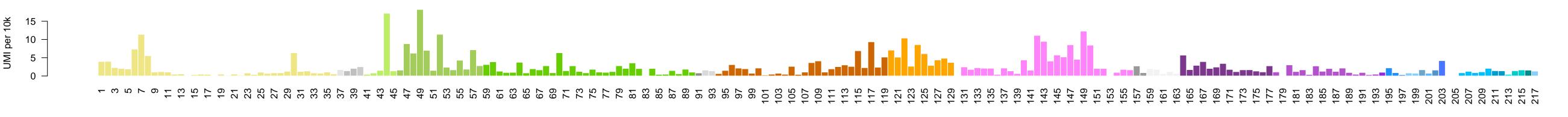
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TrH2_TrispH2_004617



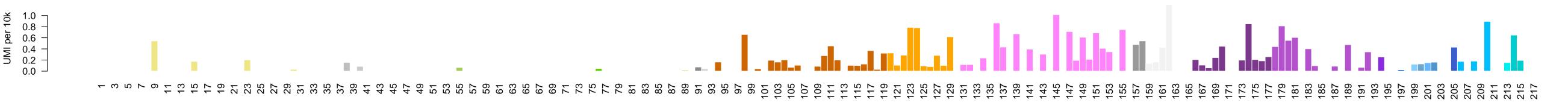
ING.HG1.3:ING1/ING2
TrH2_TrispH2_010373



LinkerHistone.HG1.12:like:H1-0/H1-10/SHPRH TrH2_TrispH2_002729

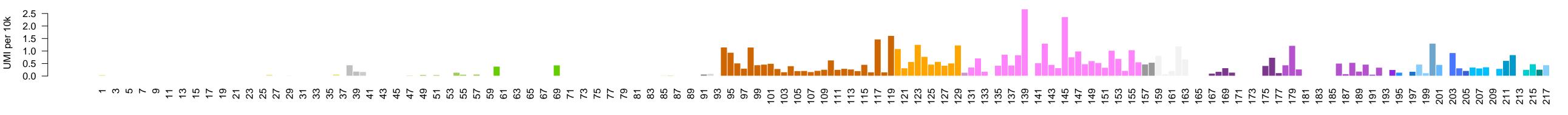


MBT.HG1.1:like:PHF20L1 TrH2_TrispH2_001486

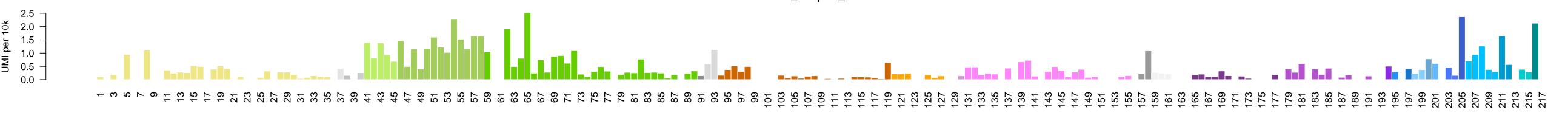


MBT.HG1.2:like:AL031681.2/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2

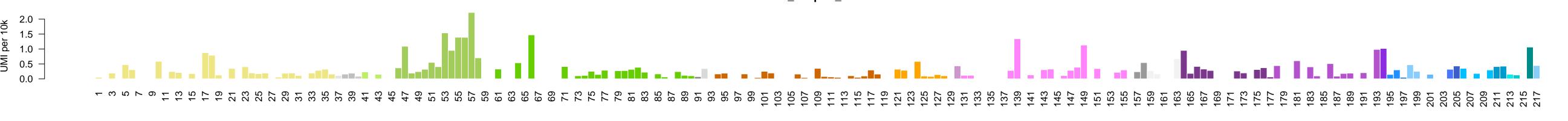
TrH2_TrispH2_004509



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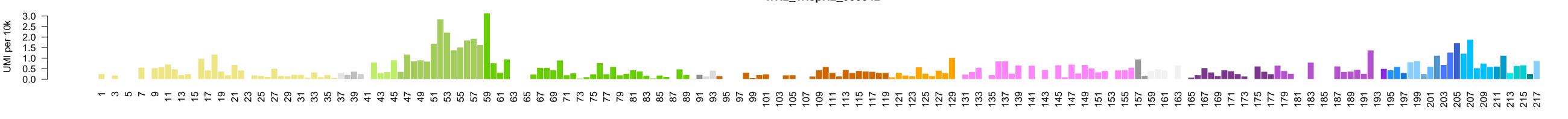


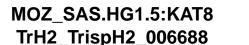
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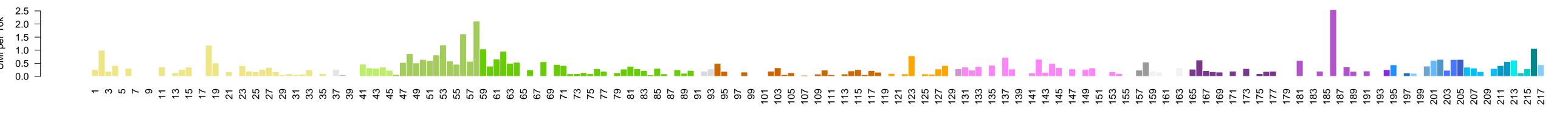


MOZ_SAS.HG1.2:like:KAT5/KAT6A/KAT6B/KAT7/KAT8

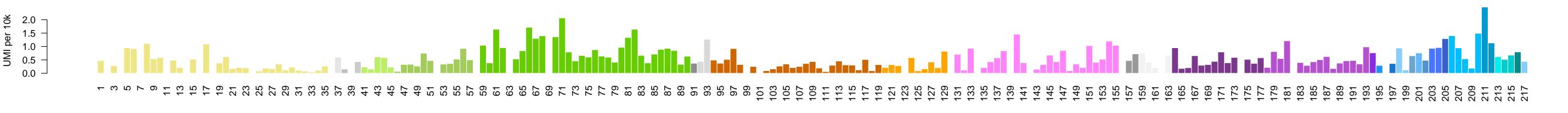
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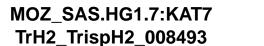


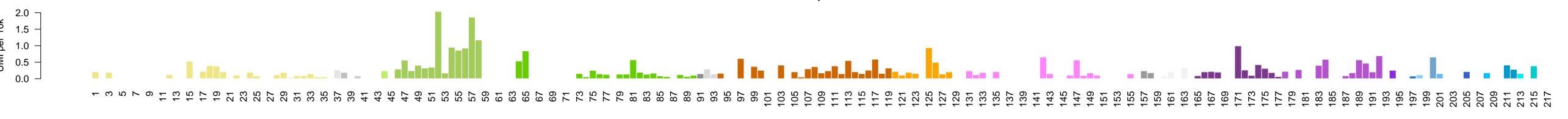




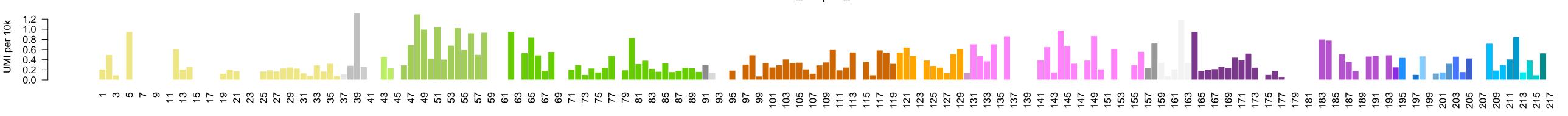
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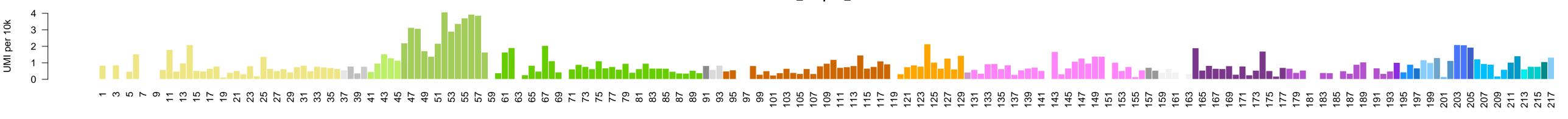




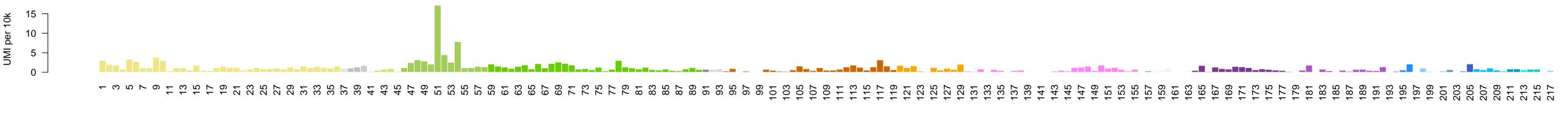
MRG.HG1.1:MORF4L1/MORF4L2 TrH2_TrispH2_008230



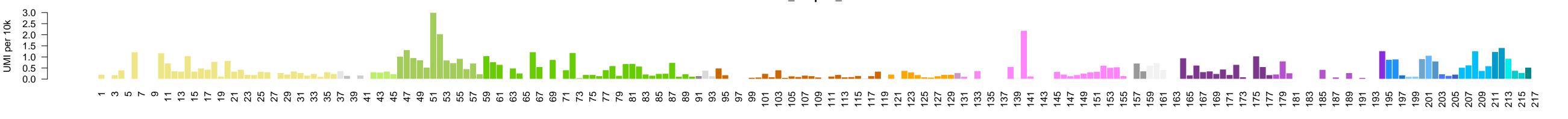
NAP.HG1.0:SET/SETSIP/TSPY1/TSPY2/TSPY3/TSPY4/TSPY8/TSPY9P/TSPY10/TSPYL1/TSPYL2/TSPYL4/TSPYL5/TSPYL6 TrH2_TrispH2_003034



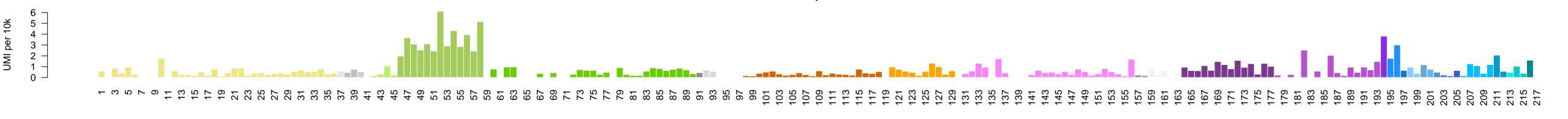
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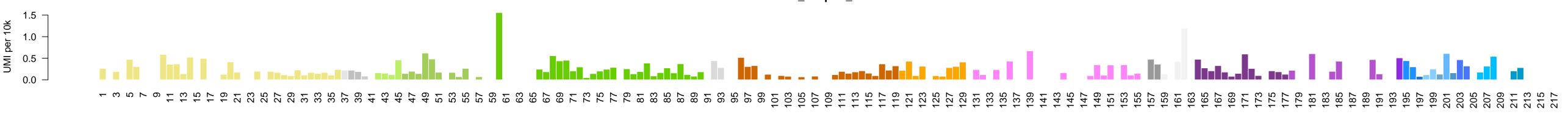
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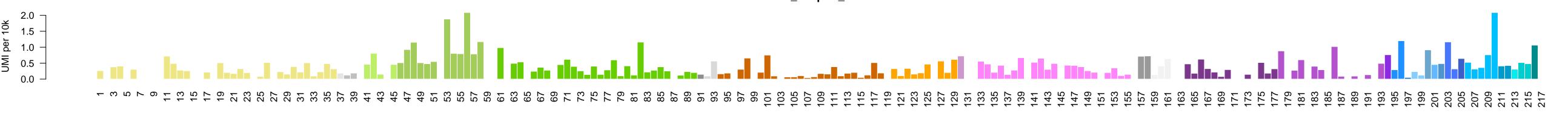
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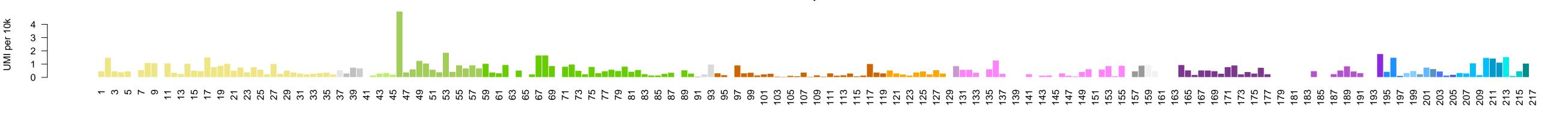
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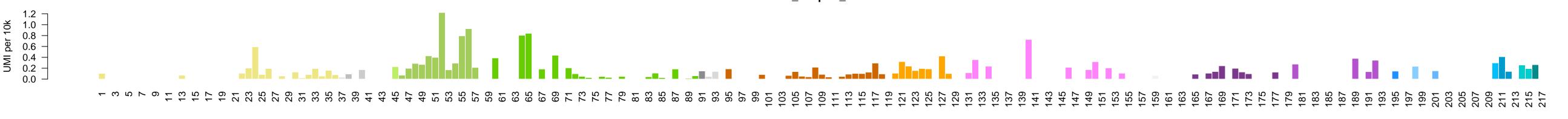
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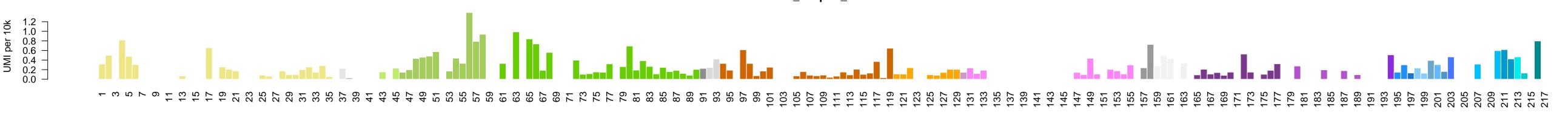
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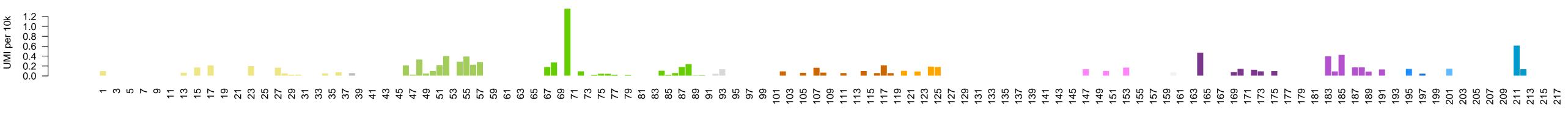
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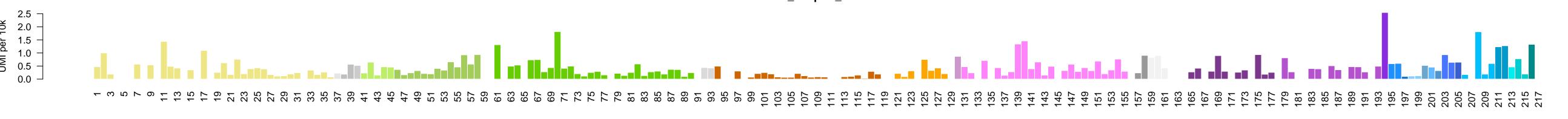
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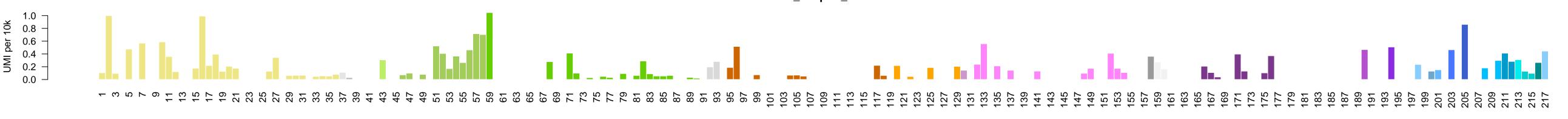




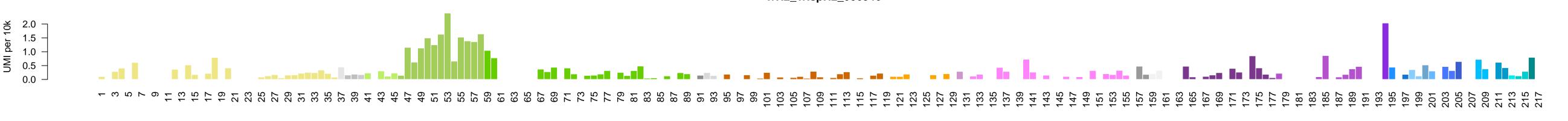
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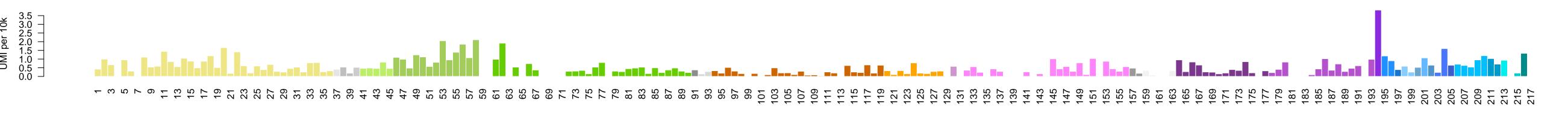
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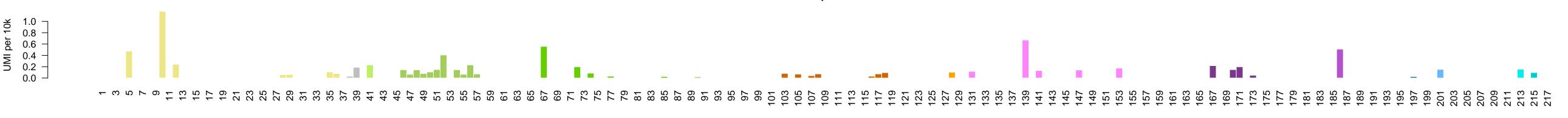
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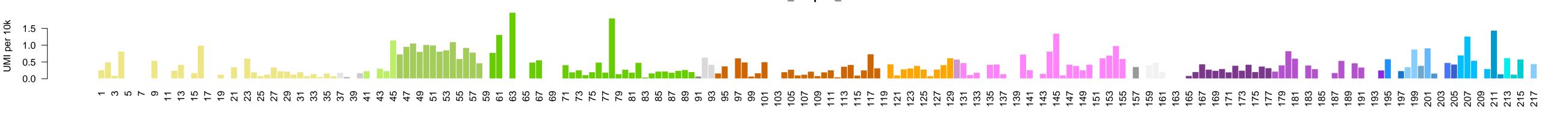
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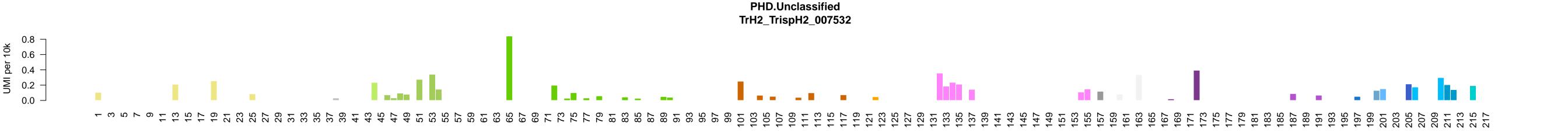


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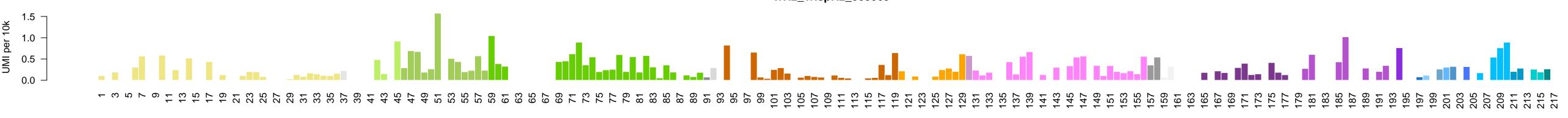


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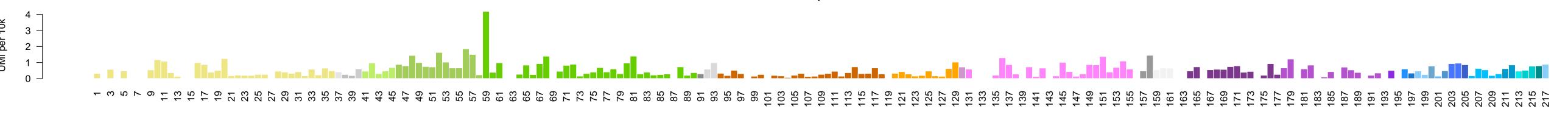




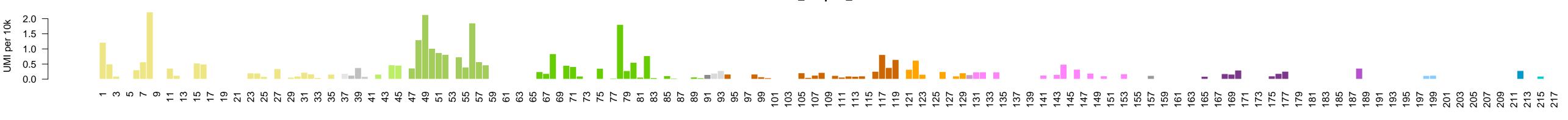
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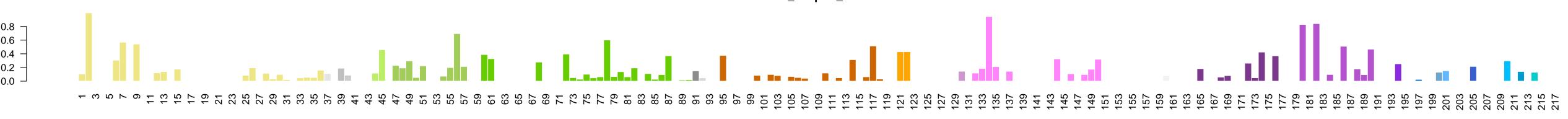
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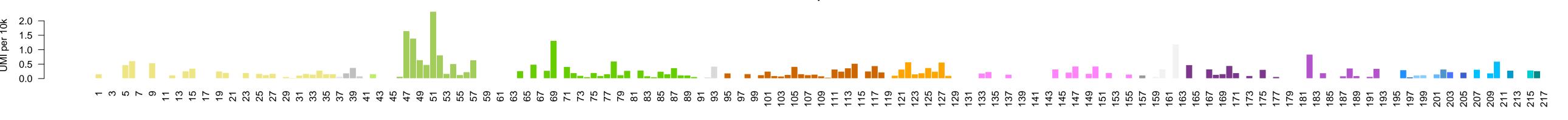
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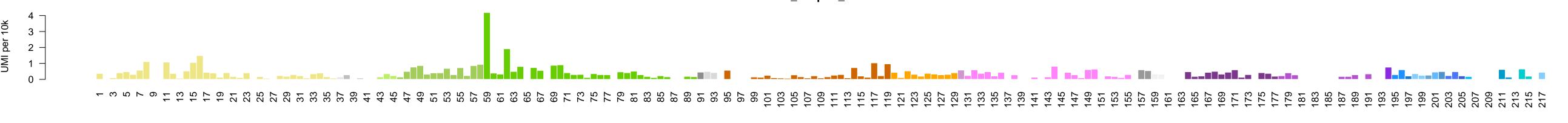
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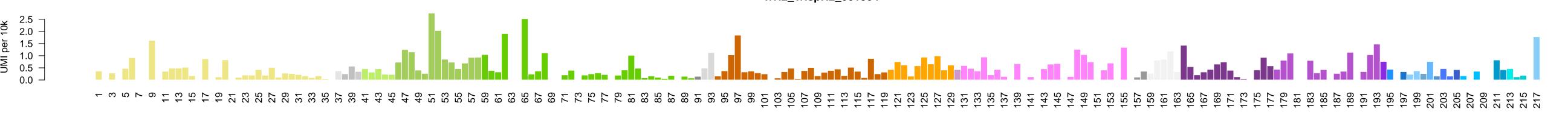
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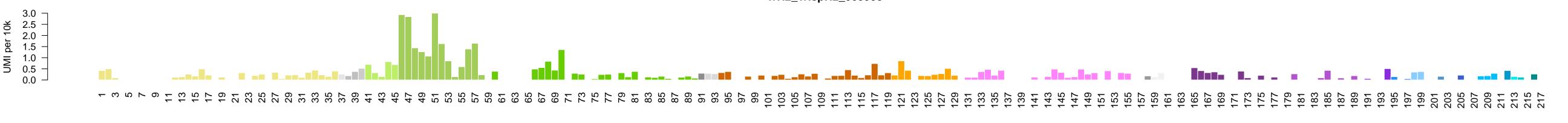
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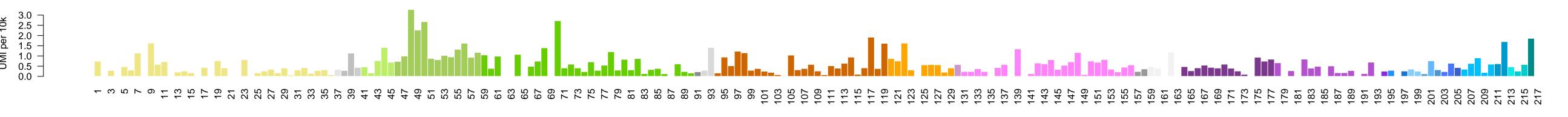
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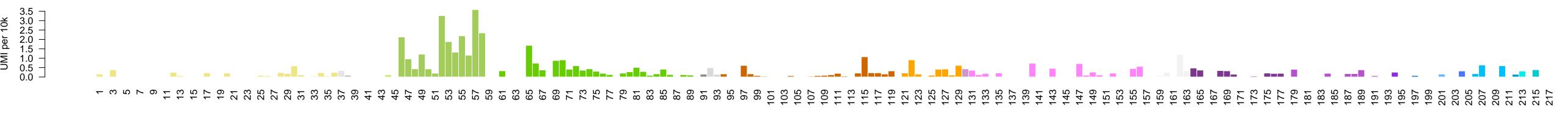
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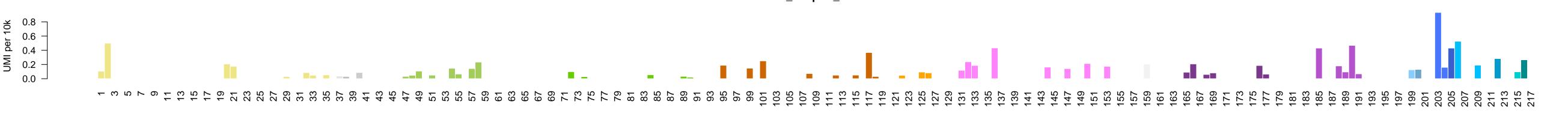
Rtt106.HG2.0:SUPT16H;SPT16.HG1.0:SUPT16H TrH2_TrispH2_000175



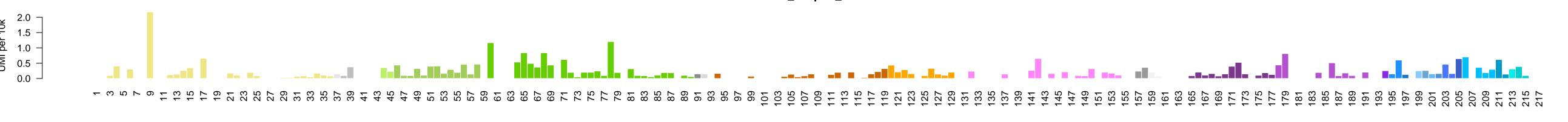
SET.HG1.0:KMT5B/KMT5C TrH2_TrispH2_011149



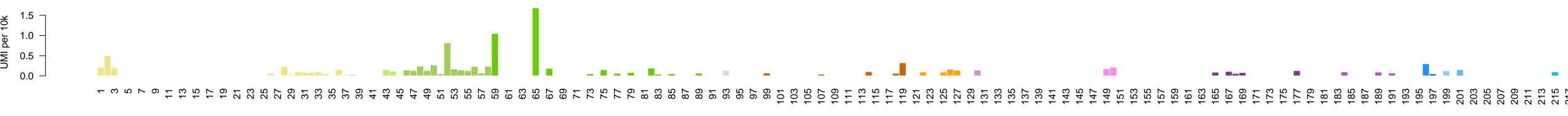
SET.HG1.21:EZH1/EZH2 TrH2 TrispH2 007277



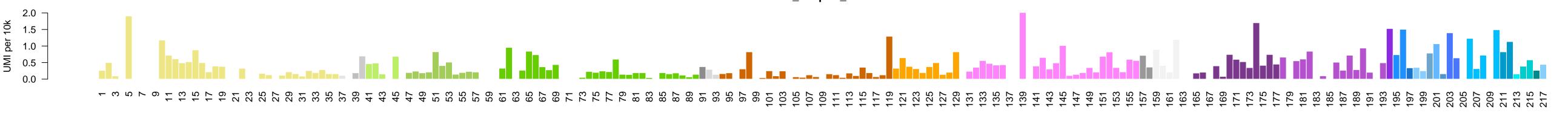
SET.HG1.25:SETD1A/SETD1B TrH2 TrispH2 011464



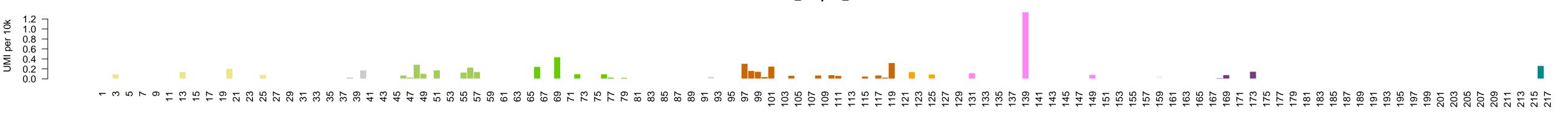
SET.HG1.30:SMYD4 TrH2_TrispH2_009620



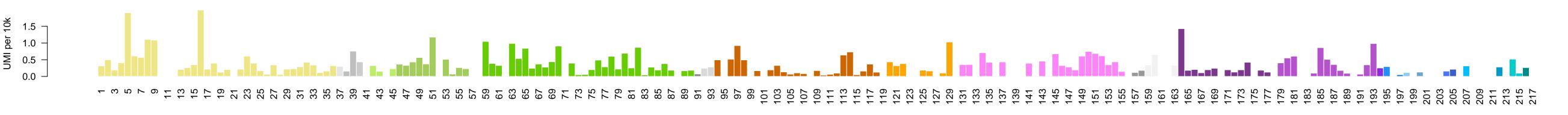
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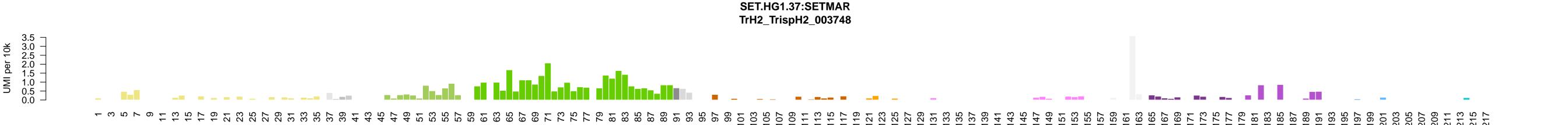


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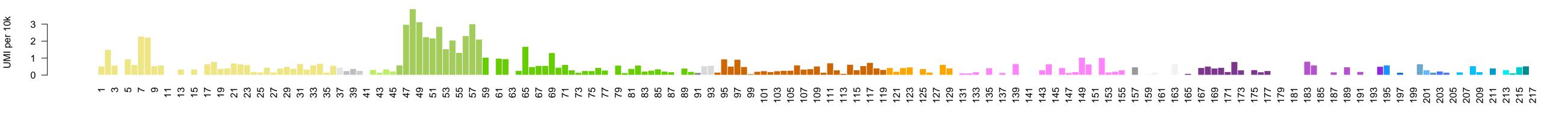


SET.HG1.34:SETD2 TrH2_TrispH2_009709

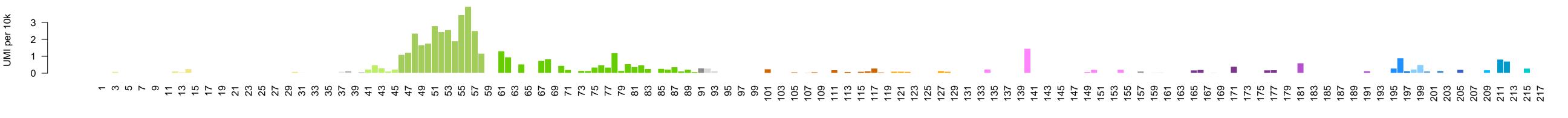




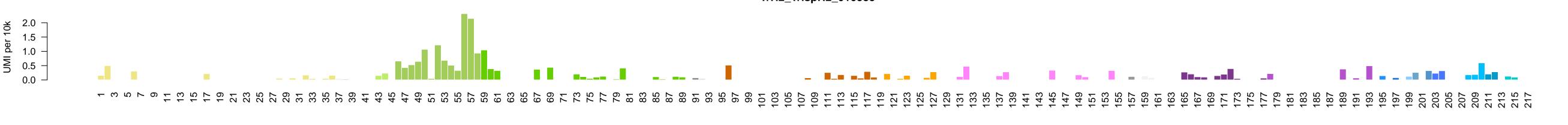
SET.HG1.44:EHMT1/EHMT2 TrH2_TrispH2_003966



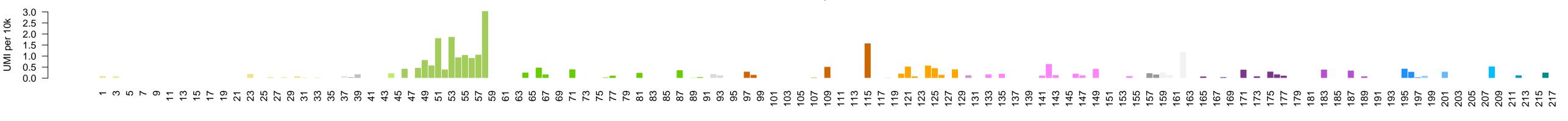
SET.HG1.45:SETDB1/SETDB2 TrH2_TrispH2_010383



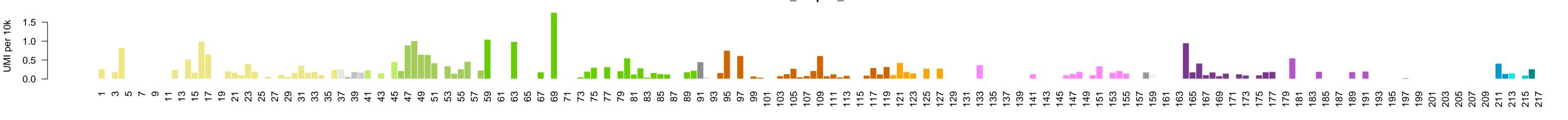
SET.HG1.45:SETDB1/SETDB2 TrH2_TrispH2_010385



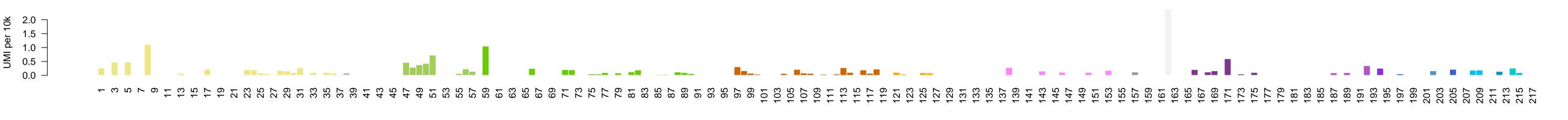
SET.HG1.49:KMT5A TrH2_TrispH2_005943



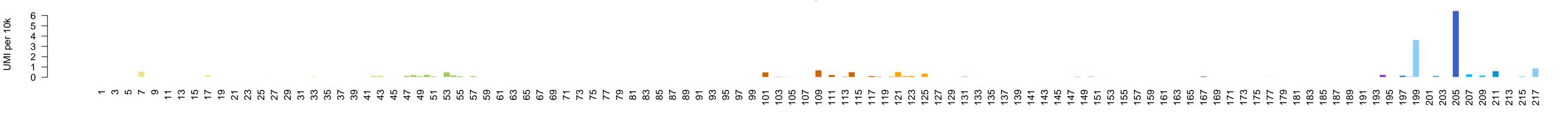
SET.HG1.52:SMYD5 TrH2 TrispH2 007115



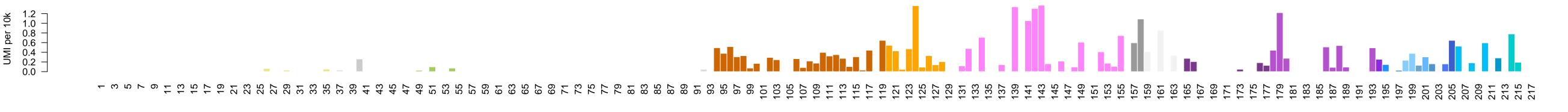
SET.HG1.6:SETD7 TrH2_TrispH2_005474



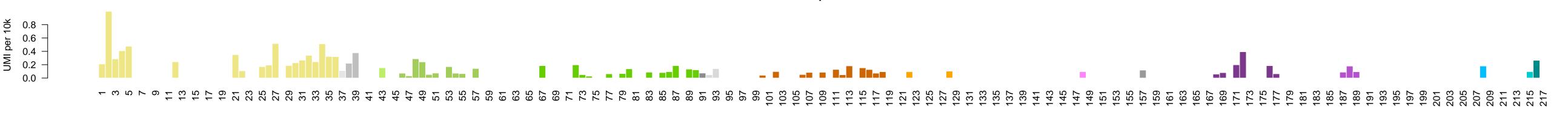
SET.HG2.2:PRDM1/PRDM6/PRDM14 TrH2 TrispH2 002621



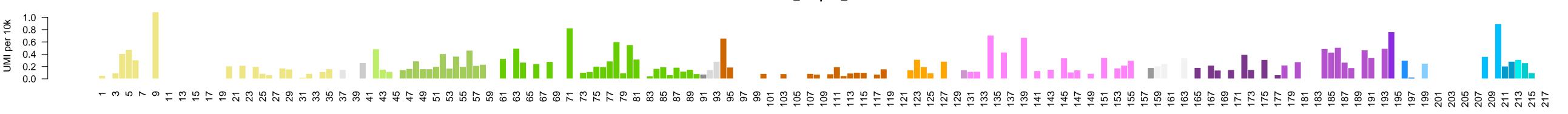
SIR2.HG1.12:like:SIRT3 TrH2_TrispH2_005203



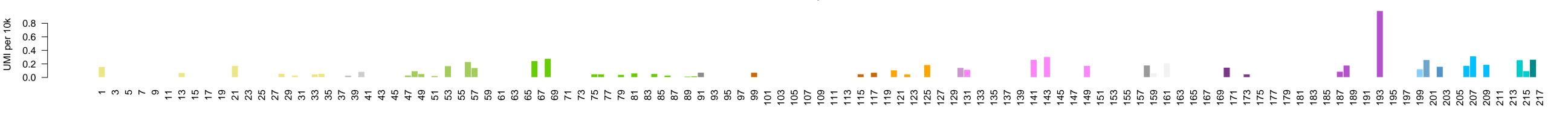
SIR2.HG1.13:SIRT3 TrH2_TrispH2_005204



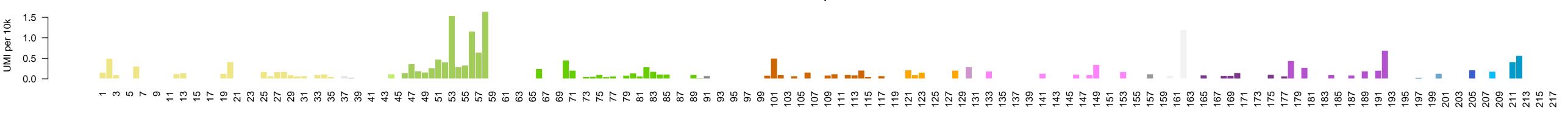
SIR2.HG1.16:SIRT2 TrH2_TrispH2_003287



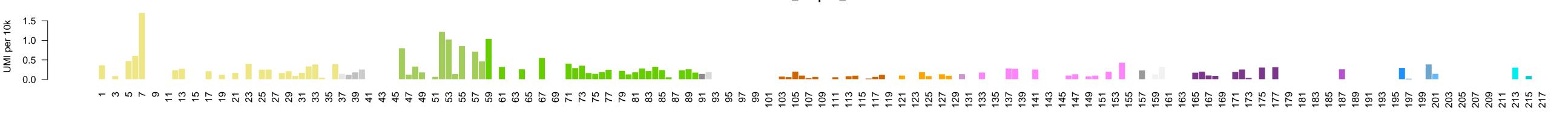
SIR2.HG1.2:SIRT7 TrH2_TrispH2_002249



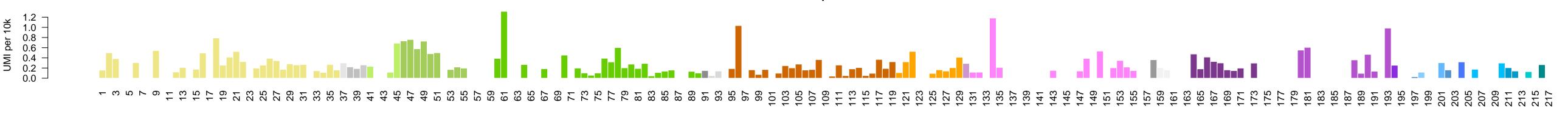
SIR2.HG1.3:SIRT4 TrH2_TrispH2_006893



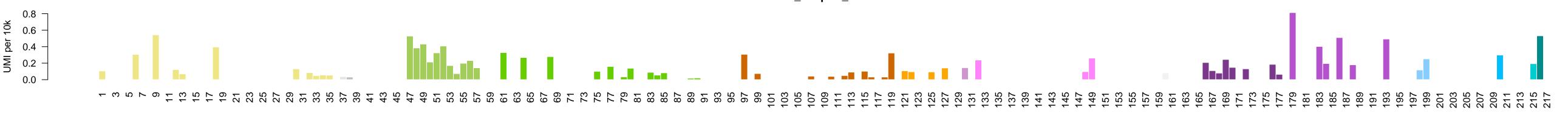
SIR2.HG1.4:like:SIRT1/SIRT2/SIRT3 TrH2_TrispH2_007426



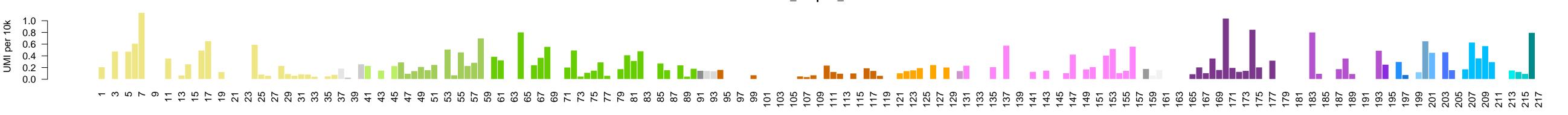
SIR2.HG1.6:SIRT5 TrH2_TrispH2_003851



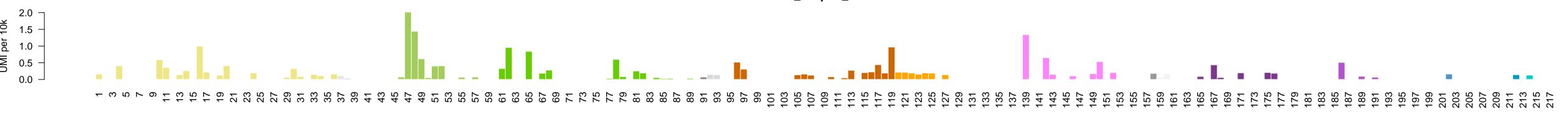
SNF2_N.HG1.0:ERCC6L TrH2_TrispH2_001971



SNF2_N.HG1.1:ERCC6 TrH2_TrispH2_002335

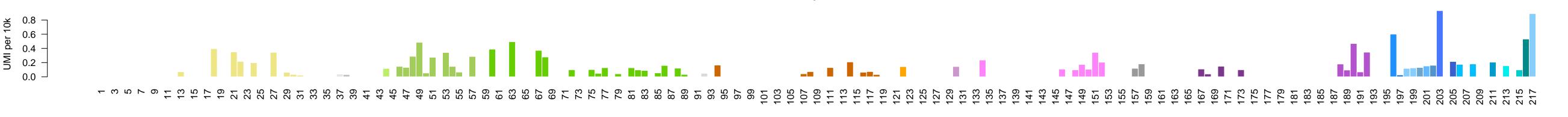


SNF2_N.HG1.10:RAD54B TrH2_TrispH2_002880

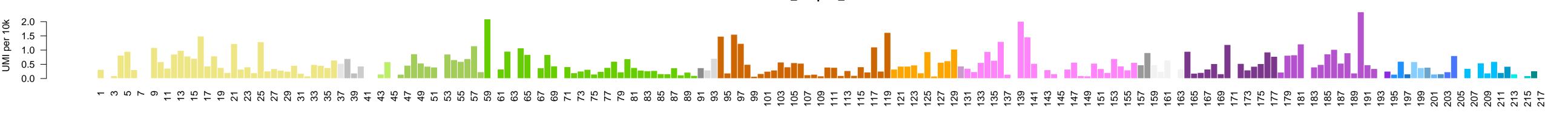


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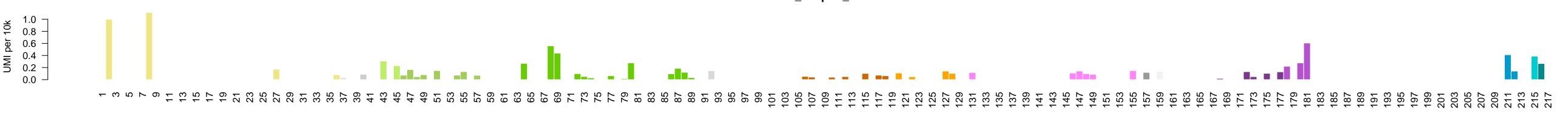
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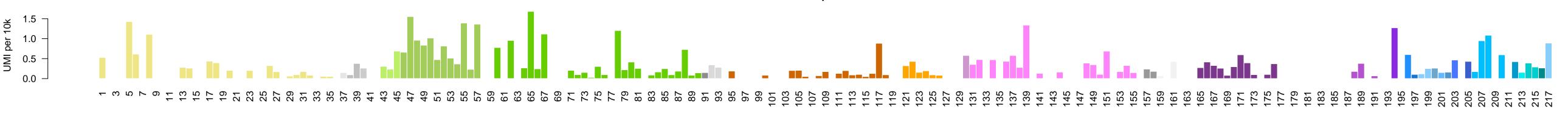
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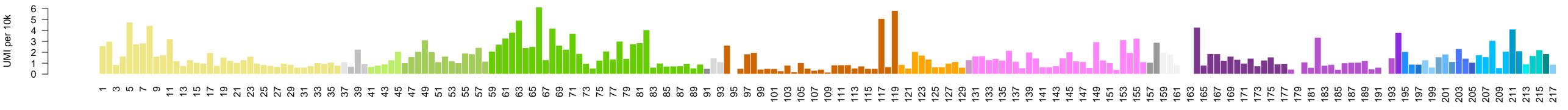
SNF2_N.HG1.13:SHPRH TrH2_TrispH2_002706



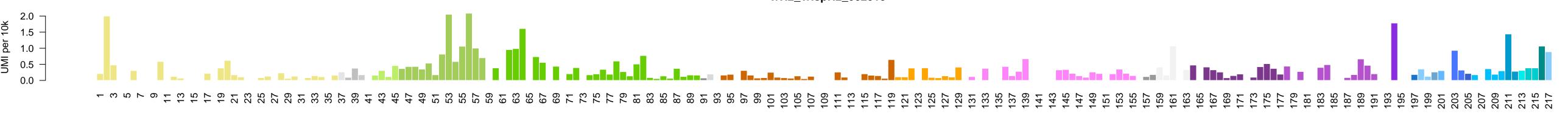
SNF2_N.HG1.15:INO80 TrH2_TrispH2_005586



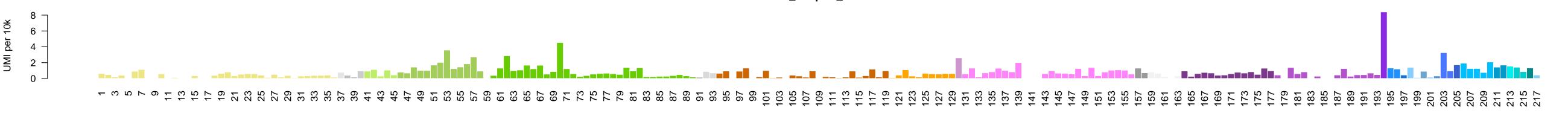
SNF2_N.HG1.16:AC106886.5/EP400/SRCAP TrH2_TrispH2_002018

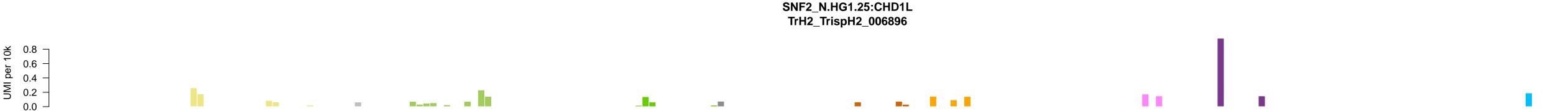


SNF2_N.HG1.2:BTAF1 TrH2_TrispH2_002319

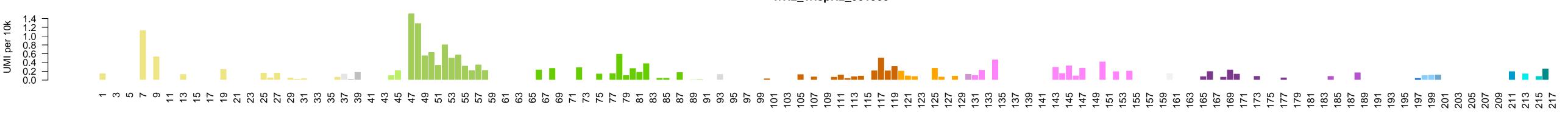


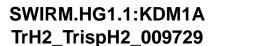
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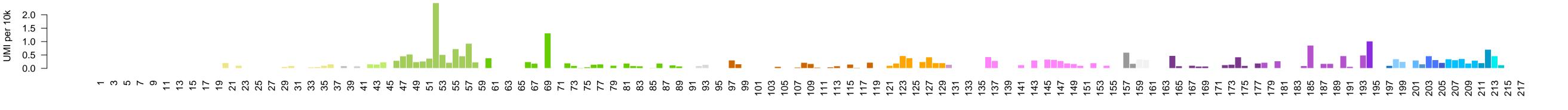




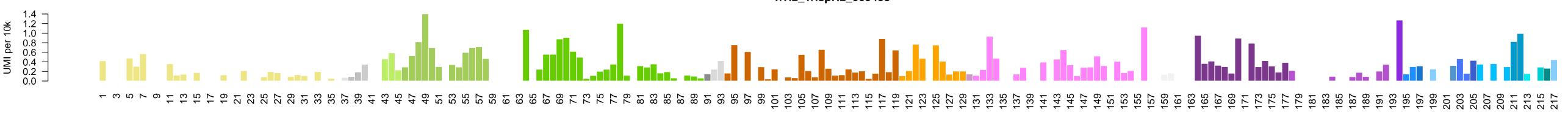
SNF2_N.HG1.7:SMARCAD1 TrH2_TrispH2_001905

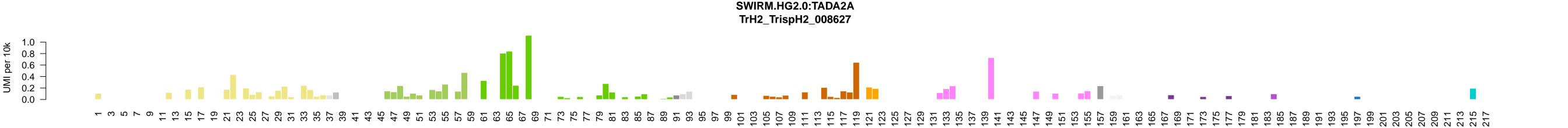


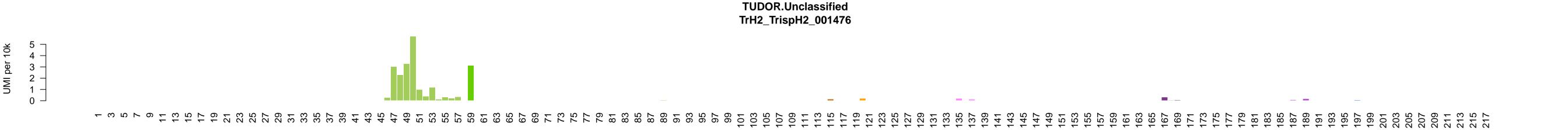


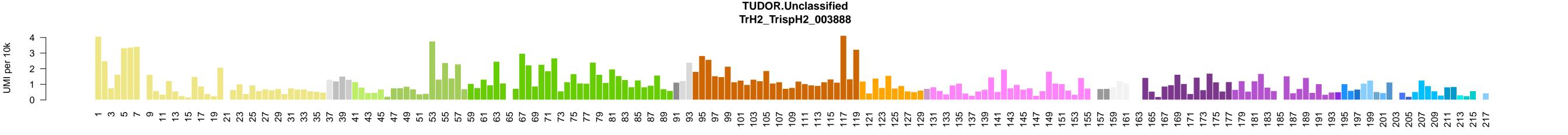


SWIRM.HG1.3:SMARCC1/SMARCC2 TrH2_TrispH2_009433

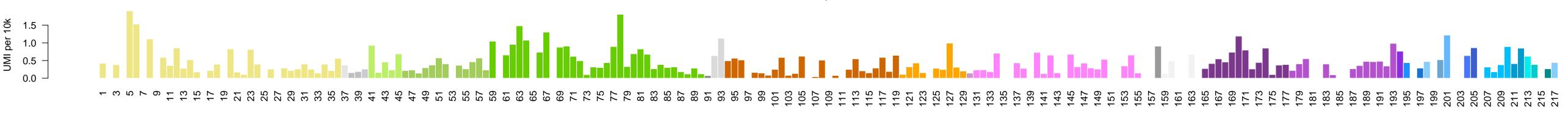




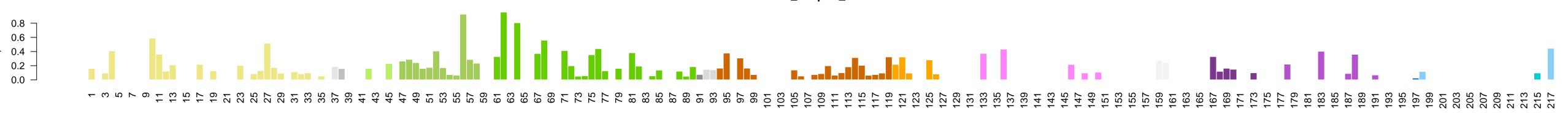




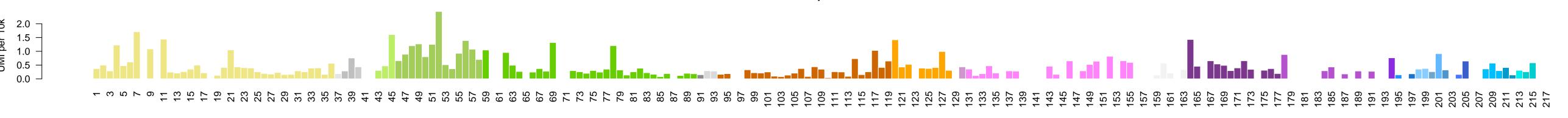
TUDOR.Unclassified
TrH2 TrispH2 006558



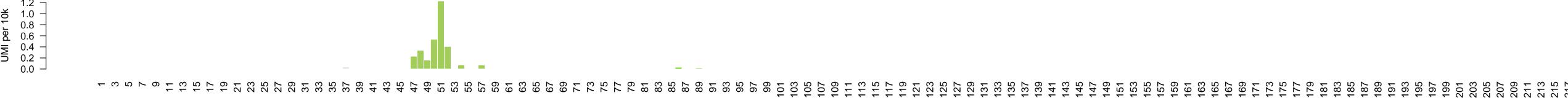
TUDOR.Unclassified TrH2 TrispH2 007007



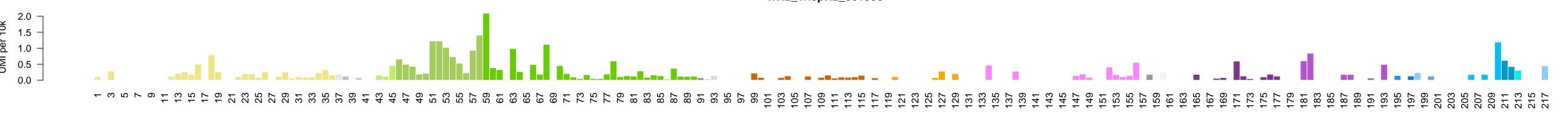
TUDOR.Unclassified TrH2 TrispH2 008567



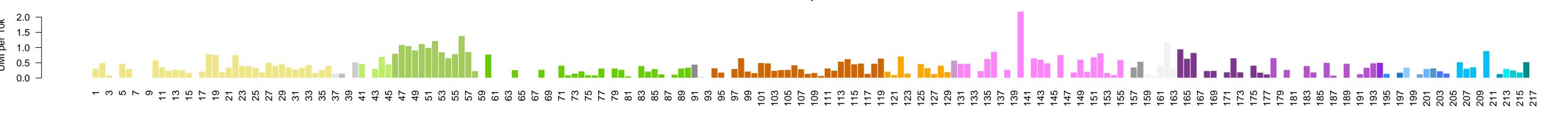




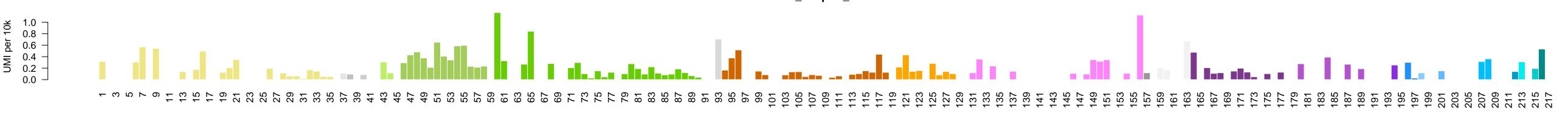
YEATS.HG1.0:MLLT1/MLLT3 TrH2_TrispH2_001306



YEATS.HG1.1:YEATS4
TrH2_TrispH2_007327



YEATS.HG1.2:YEATS2 TrH2 TrispH2 002142



TrH2_TrispH2_009035

zf-CCHH.HG1.2:like:APLF

zf-CCHH.HG1.8:like:APLF TrH2 TrispH2 004577

