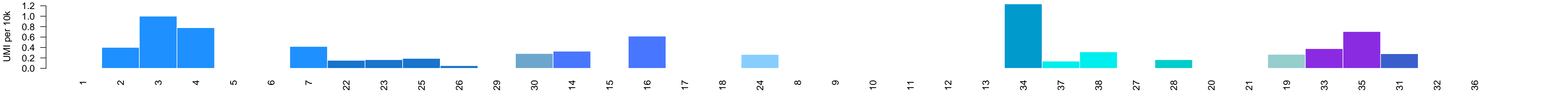


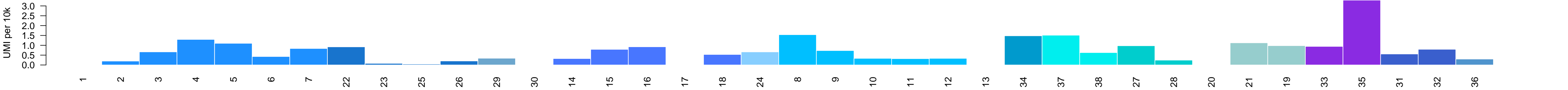
Acetyltransf_1.HG1.1:like:AANAT/GNPNAT1/NAA40/NAA50/NAA60/NAA80/NAT8/NAT8B/NAT8L/SAT1/SAT2/SATL1
Hhon_g01422.t1



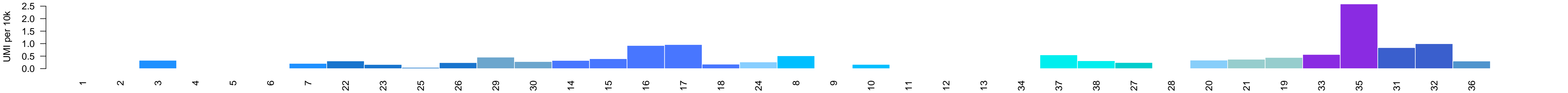
Acetyltransf_1.HG1.16:GNPNAT1
Hhon_g03106.t1



Acetyltransf_1.HG1.8:NAA20
Hhon_g07286.t1



Acetyltransf_1.HG2.0:NA
Hhon_g10162.t1



Acetyltransf_1.HG2.0:NA
Hhon_g11808.t1

UMI per 10k

2.0
1.5
1.0
0.5
0.0

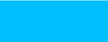
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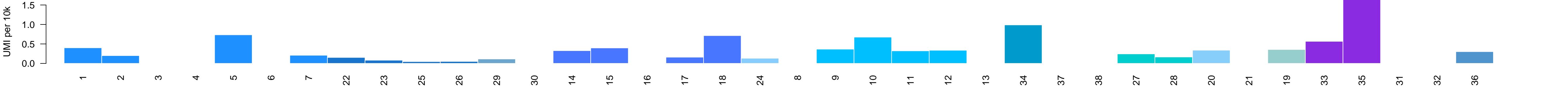


19

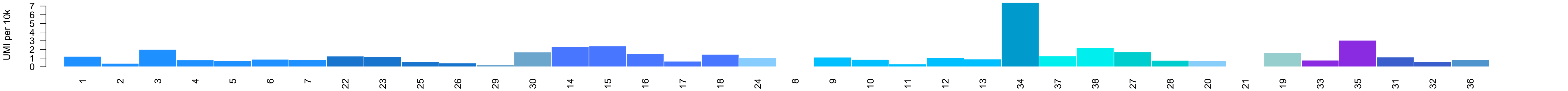


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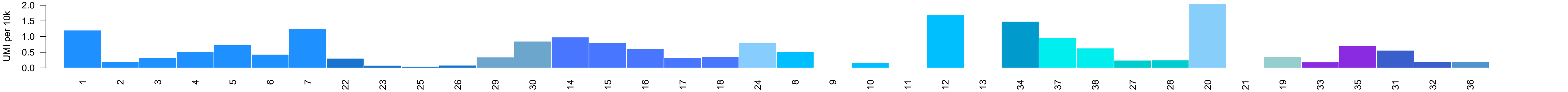
BAH.HG1.3:PBRM1;Bromodomain.HG1.29:PBRM1
Hhon_g09288.t1



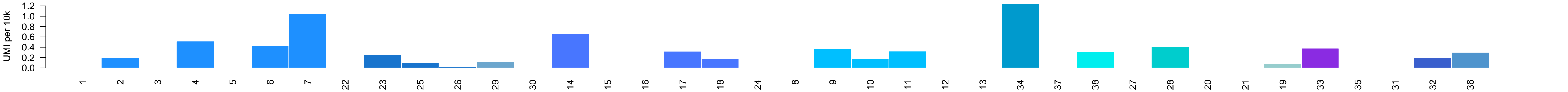
BAH.HG1.6:RERE
Hhon_g01820.t1



BIR.HG1.5:BIRC6
Hhon_g04612.t1

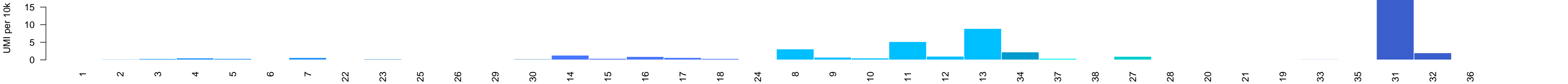


BIR.HG1.6:BIRC5
Hhon_g02999.t1

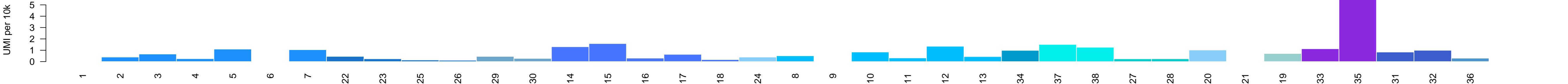


Bromodomain.HG1.17:BRD7/BRD9

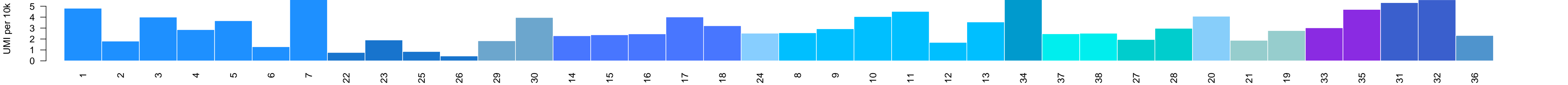
Hhon_g04486.t1



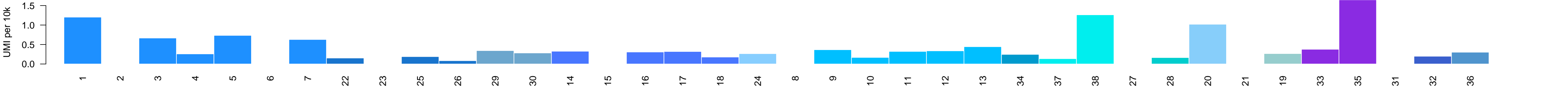
Bromodomain.HG1.18:BRD1/BRPF1/BRPF3;PWWP.HG1.12:BRD1/BRPF1/BRPF3
Hhon_g02173.t1



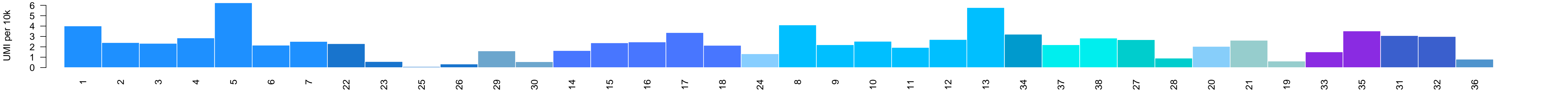
Bromodomain.HG1.20:BRD2/BRD3/BRD4/BRDT
Hhon_g03776.t1



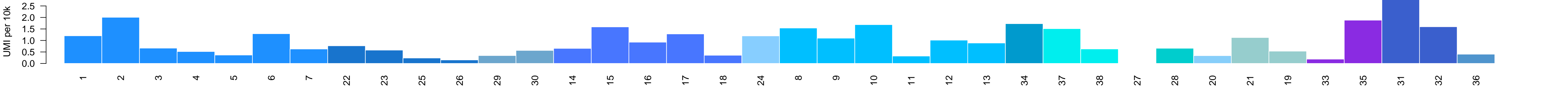
Bromodomain.HG1.21:BRWD1/BRWD3/PHIP
Hhon_g08078.t1



Bromodomain.HG1.25:CREBBP/EP300;KAT11.HG1.2:CREBBP/EP300
Hhon_g05460.t1



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



Bromodomain.HG1.4:BAZ1A
Hhon_g07271.t1

UMI per 10k

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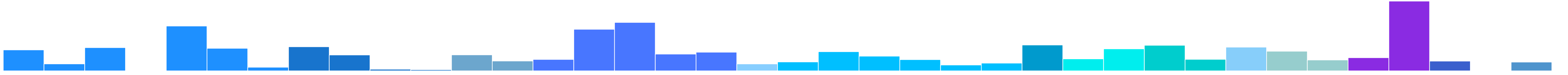
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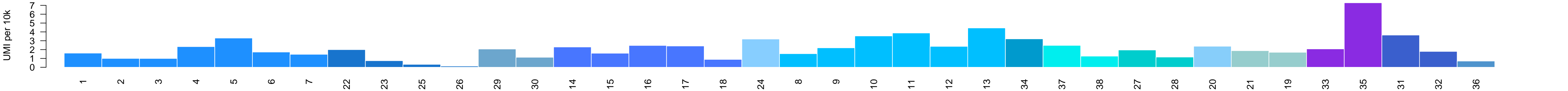
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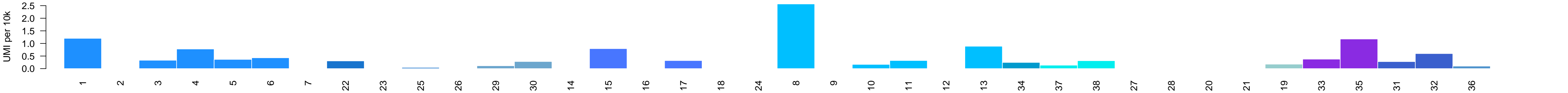
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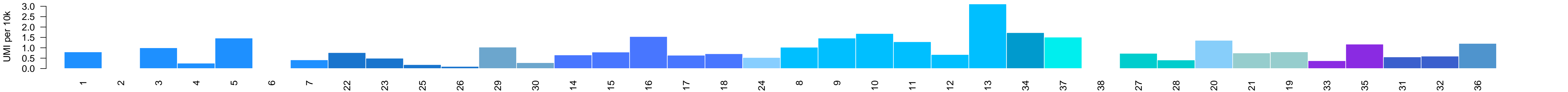
Bromodomain.HG1.5:BAZ2A/BAZ2B
Hhon_g04767.t1



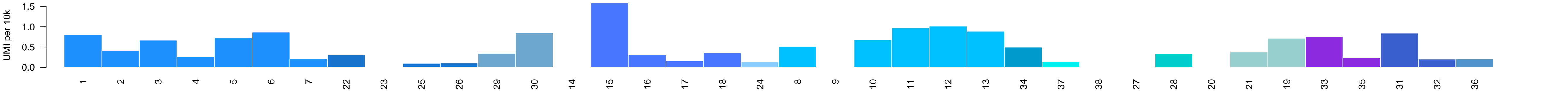
Bromodomain.HG1.9:like:BRD2/BRD3/BRD4/BRDT
Hhon_g09299.t1



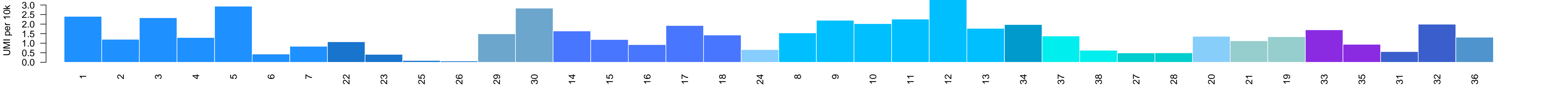
Chromo.HG1.13:CHD6/CHD7/CHD8/CHD9;SNF2_N.HG1.28:CHD6/CHD7/CHD8/CHD9
Hhon_g07827.t1



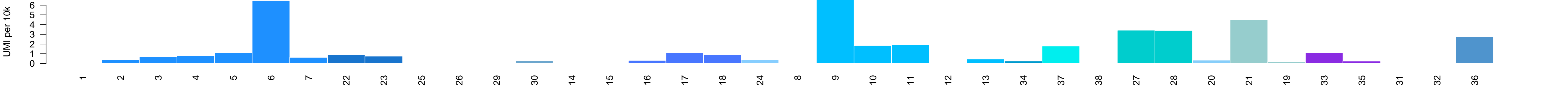
Chromo.HG1.6:SUV39H1/SUV39H2
Hhon_g09127.t1



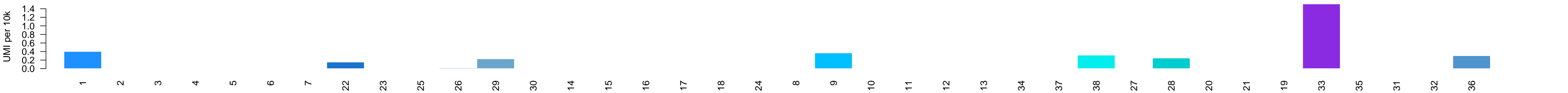
CupinJmjC.HG1.24:KDM7A/PHF2/PHF8
Hhon_g06125.t1



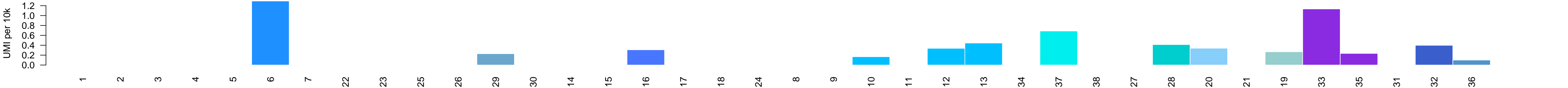
CupinJmjC.HG1.25:JMJD6
Hhon_g07323.t1



CupinJmjC.HG1.39:like:JMJD7/JMJD7-PLA2G4B
Hhon_g03153.t1



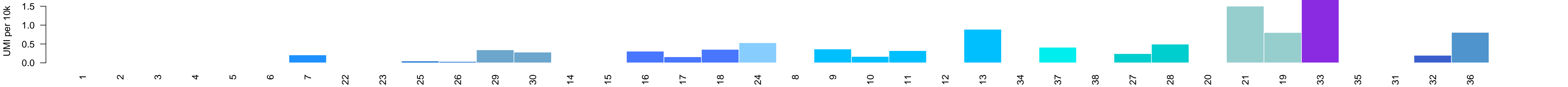
CupinJmjC.HG1.46:like:JMJD7/JMJD7-PLA2G4B
Hhon_g01871.t1



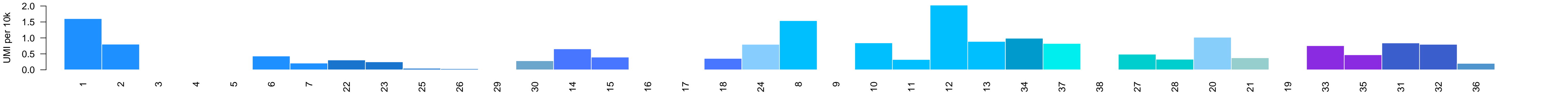
CupinJmjC.HG1.46:like:JMJD7/JMJD7-PLA2G4B
Hhon_g09522.t1



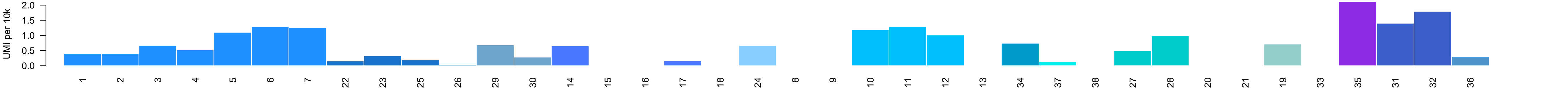
CupinJmjC.HG1.8:like:JMJD4/JMJD6/JMJD8
Hhon_g02577.t1



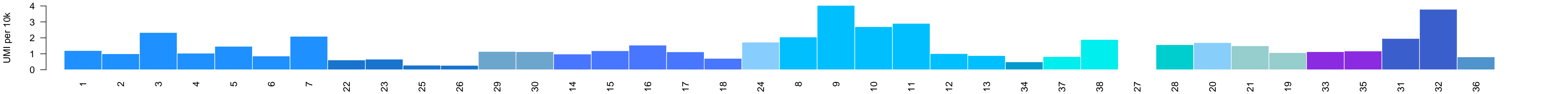
CupinJmjC.HG2.0:JARID2
Hhon_g08230.t1



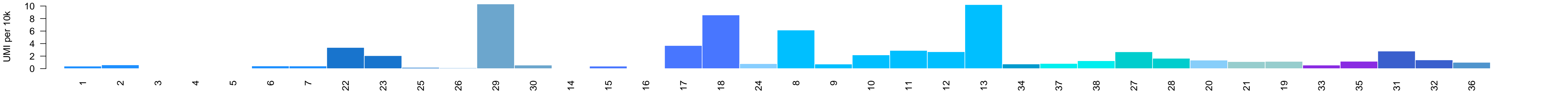
DOT1.HG1.3:DOT1L
Hhon_g00168.t1



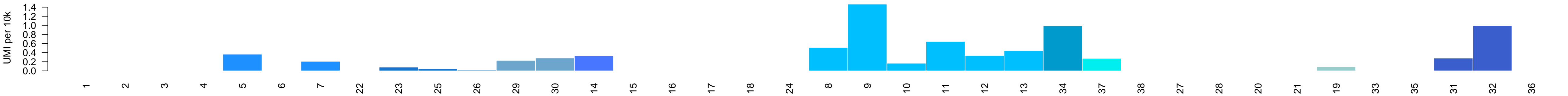
Hist_deacetyl.HG1.1:HDAC11
Hhon_g04359.t1



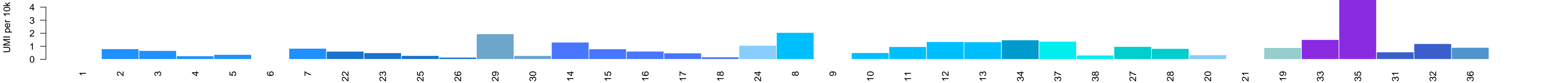
Hist_deacetyl.HG1.4:HDAC4/HDAC5/HDAC7/HDAC9
Hhon_g06280.t1



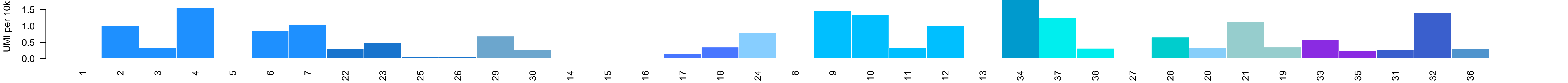
Hist_deacetyl.HG1.6:like:HDAC1/HDAC2/HDAC3
Hhon_g06894.t1



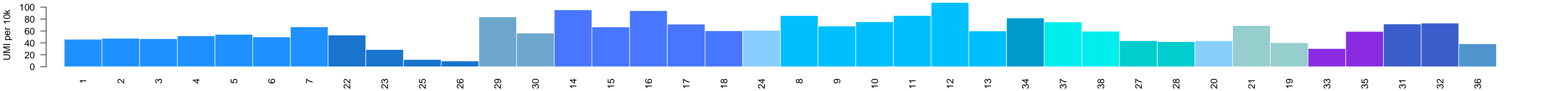
Hist_deacetyl.HG1.7:HDAC1/HDAC2
Hhon_g00226.t1



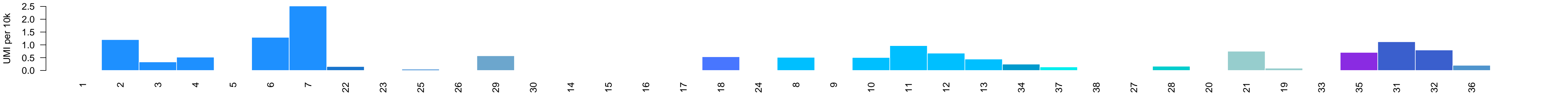
Histone.HG2.1:CENPA
Hhon_g09811.t1



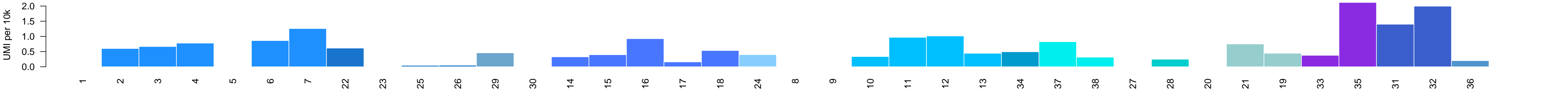
Histone.HG2.13:like:H3-3A/H3-3B/H3-5/H3Y1/H3Y2
Hhon_g06924.t1



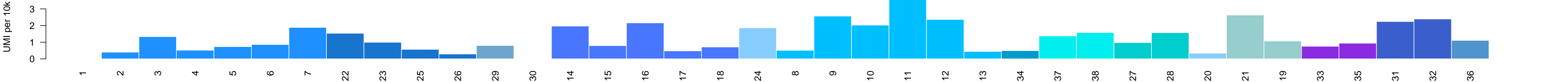
Histone.HG3.0:DRAP1
Hhon_g07246.t1



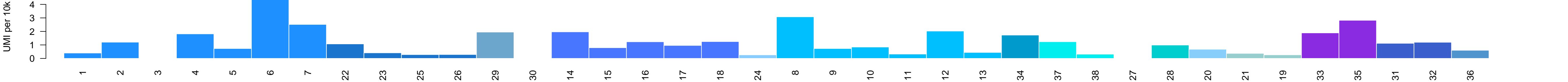
Histone.HG3.4:CHRA1
Hhon_g07697.t1



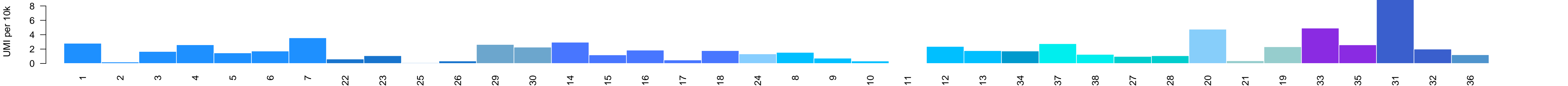
Histone.HG4.1:NFYB
Hhon_g07946.t1



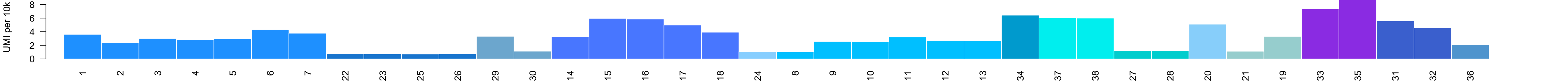
Histone.HG4.2:DR1
Hhon_g02746.t1



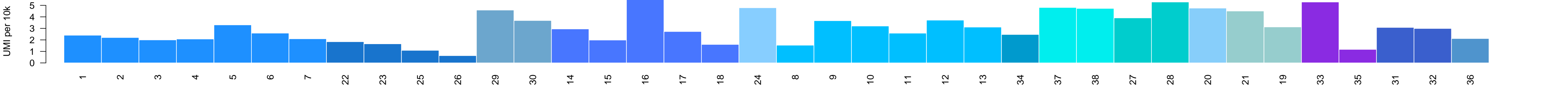
Histone.Unclassified
Hhon_g00184.t1



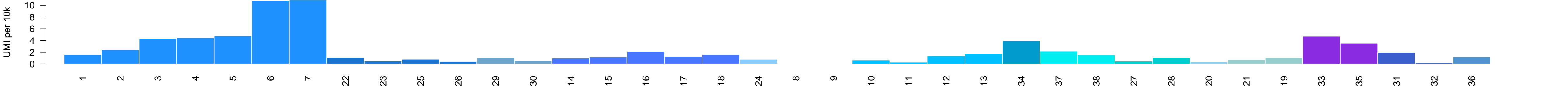
Histone.Unclassified
Hhon_g02201.t1



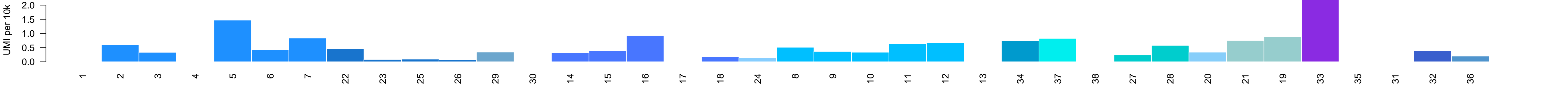
Histone.Unclassified
Hhon_g02498.t1



Histone.Unclassified
Hhon_g04967.t1

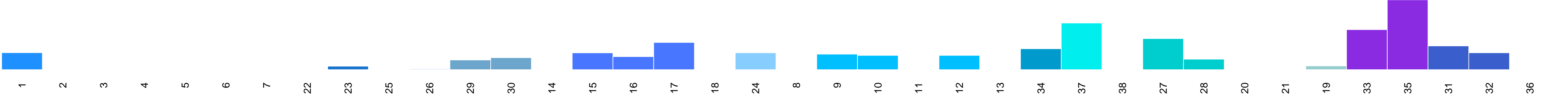
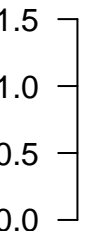


Histone.Unclassified
Hhon_g08198.t1

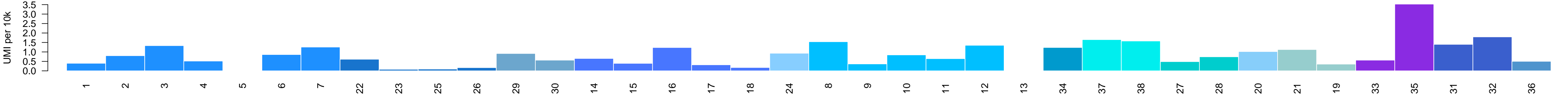


ING.HG1.1:ING3
Hhon_g06127.t1

UMI per 10k

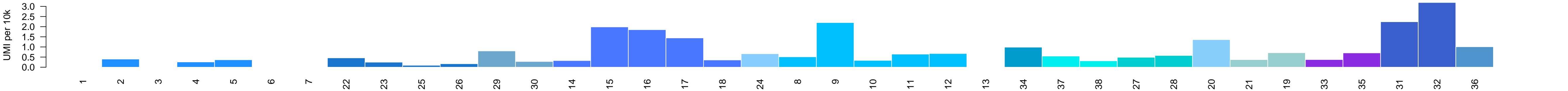


ING.HG1.3:ING1/ING2
Hhon_g00590.t1

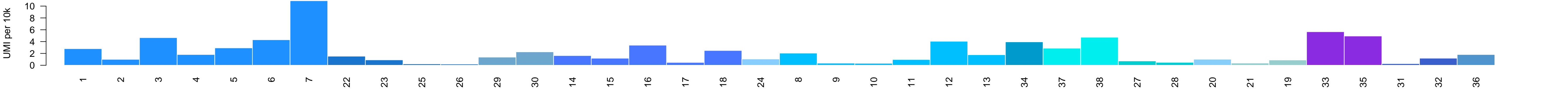


LinkerHistone.HG1.1:like:H1-0/H1-1/H1-2/H1-3/H1-4/H1-5/H1-6/H1-8/H1-10/HP1BP3/SHPRH;MOZ_SAS.HG1.6:KAT6A/KAT6B

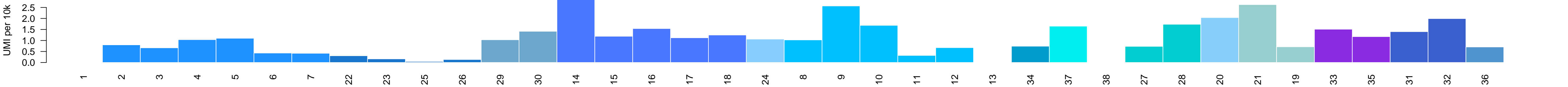
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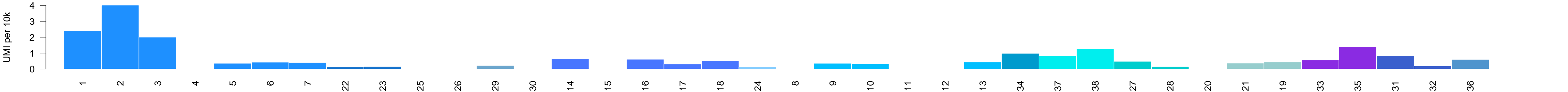
LinkerHistone.HG1.12:like:H1-0/H1-10/SHPRH
Hhon_g00465.t1



MOZ_SAS.HG1.1:like:KAT5/KAT6A/KAT6B/KAT7/KAT8
Hhon_g08105.t1

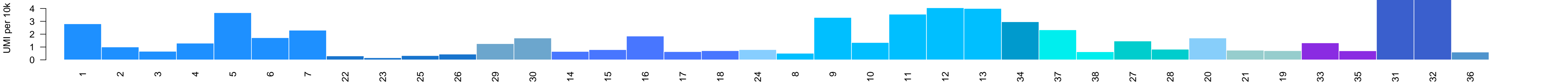


MOZ_SAS.HG1.5:KAT8
Hhon_g03304.t1



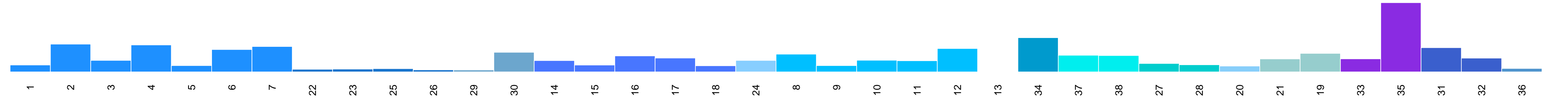
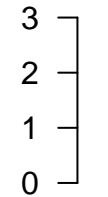
NAP.HG1.0:SET/SETSIP/TSPY1/TSPY2/TSPY3/TSPY4/TSPY8/TSPY9P/TSPY10/TSPYL1/TSPYL2/TSPYL4/TSPYL5/TSPYL6

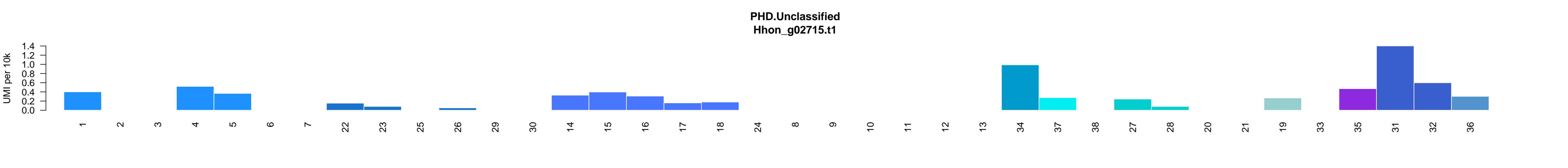
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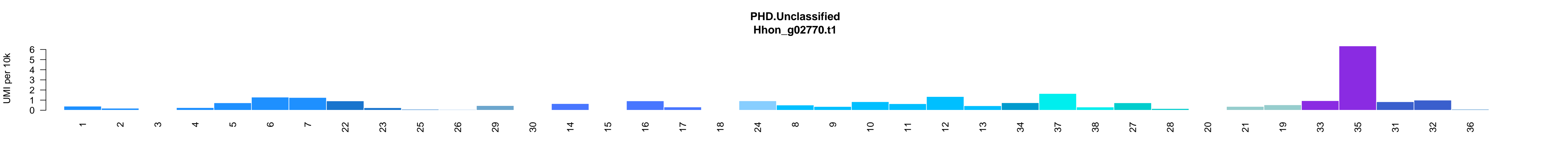


PHD.Unclassified
Hhon_g02669.t1

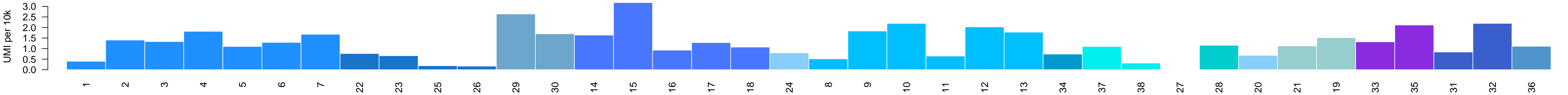
UMI per 10k

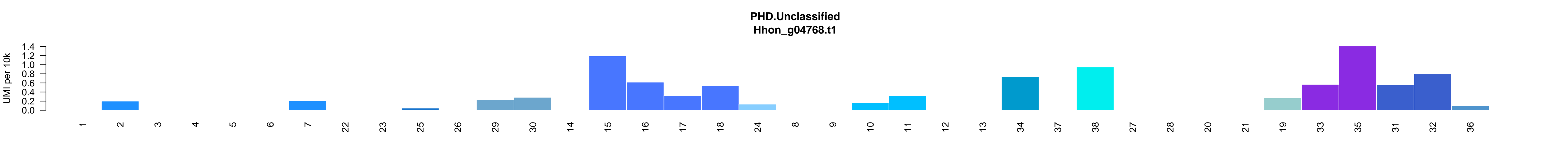


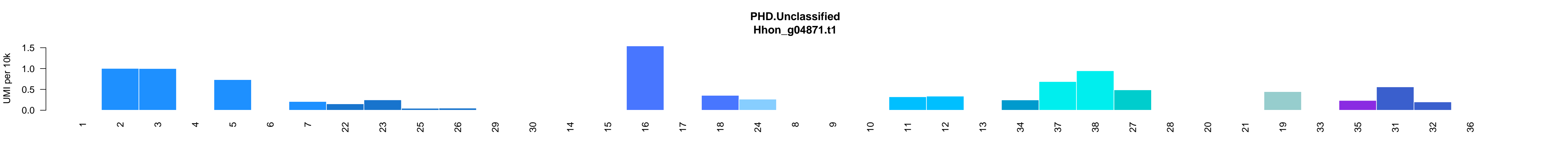


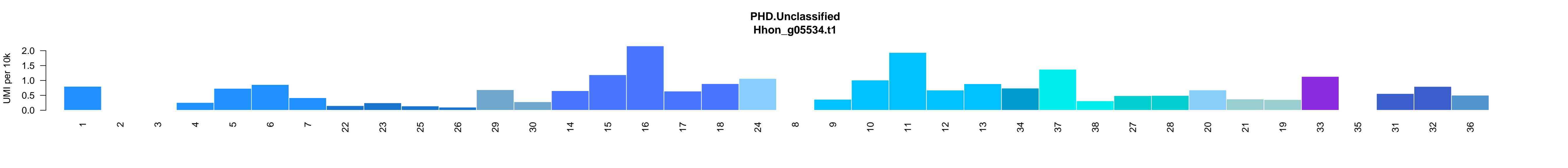


PHD.Unclassified
Hhon_g03693.t1

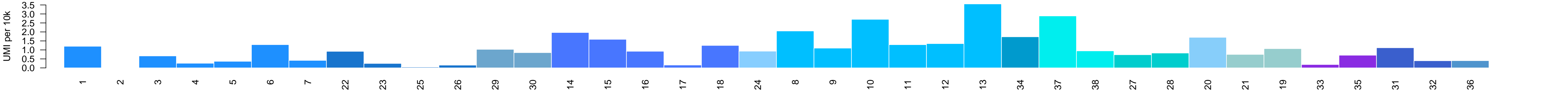




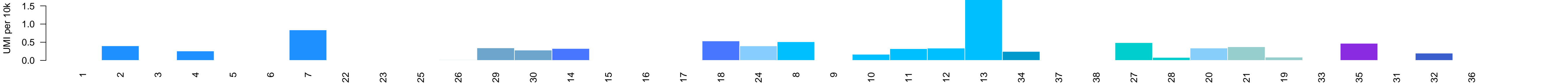




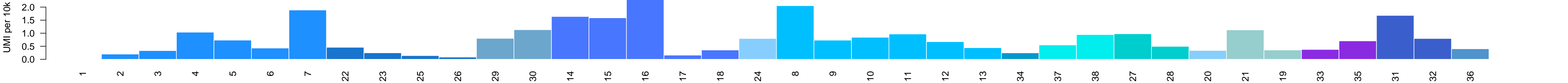
PHD.Unclassified
Hhon_g07626.t1



PHD.Unclassified
Hhon_g09435.t1



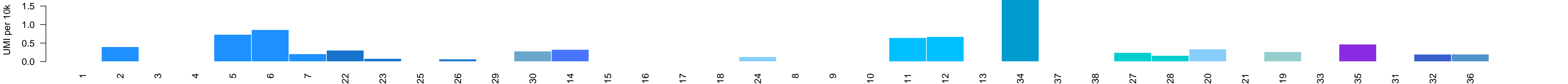
PWWP.HG1.17:HDGFL1/HDGFL2/HDGFL3/PSIP1
Hhon_g08117.t1



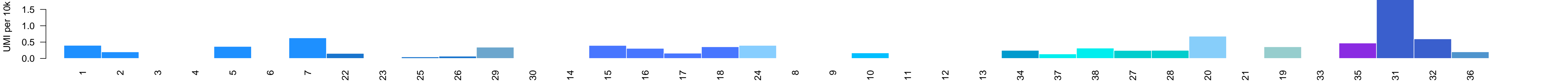
SET.HG1.21:EZH1/EZH2
Hhon_g04096.t1



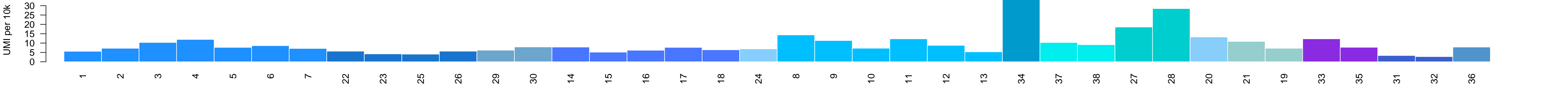
SET.HG1.34:SETD2
Hhon_g00607.t1



SET.HG1.45:SETDB1/SETDB2
Hhon_g05556.t1

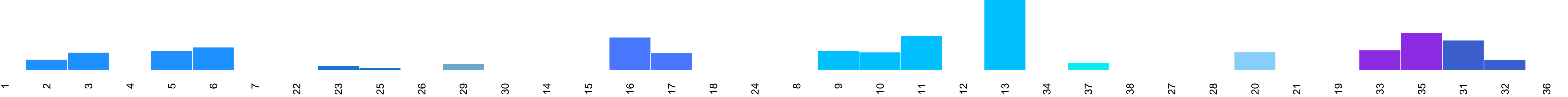
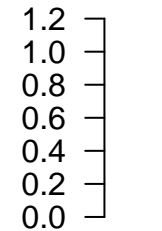


SNF2_N.HG1.0:ERCC6L
Hhon_g00800.t1

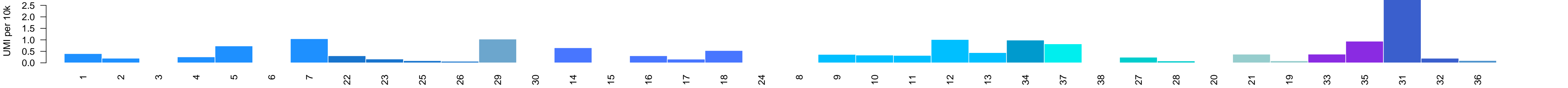


SNF2_N.HG1.1:ERCC6
Hhon_g01503.t1

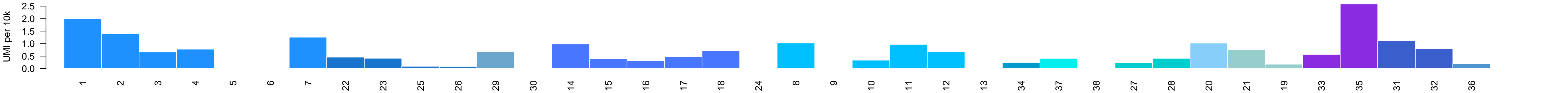
UMI per 10k



SNF2_N.HG1.15:INO80
Hhon_g07285.t1



SNF2_N.HG1.16:AC106886.5/EP400/SRCAP
Hhon_g00751.t1



SNF2_N.HG1.23:like:SMARCA5
Hhon_g05458.t1

UMI per 10k

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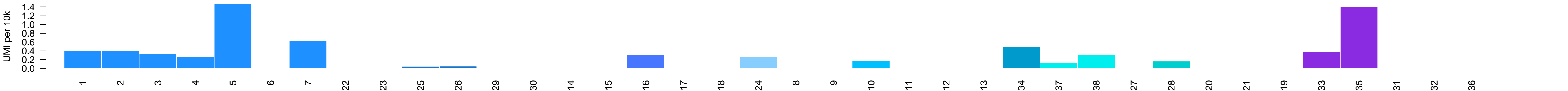
36



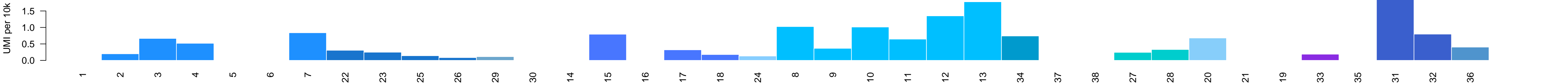
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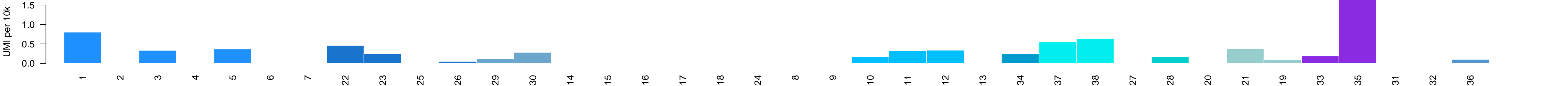
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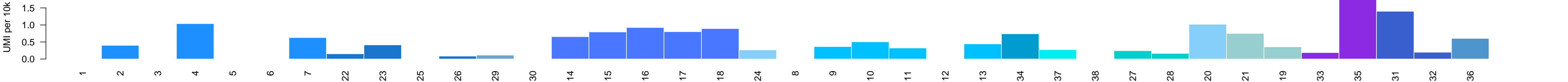
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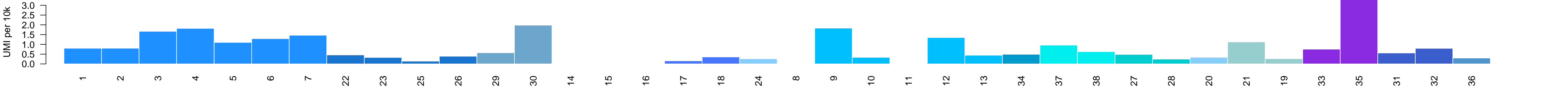
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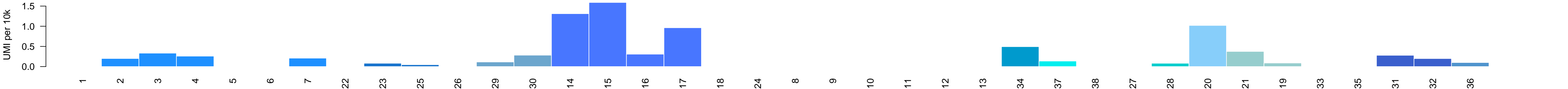
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TUDOR.Unclassified
Hhon_g02353.t1



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



YEATS.HG1.1:YEATS4
Hhon_g01885.t1

