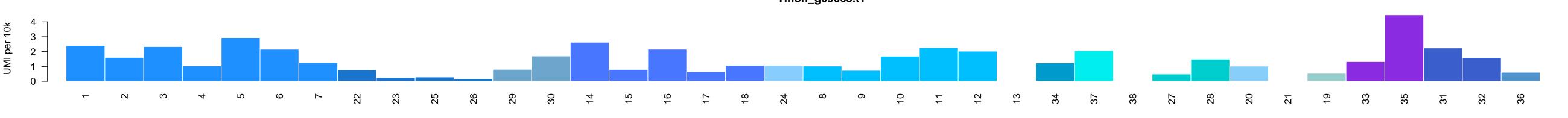
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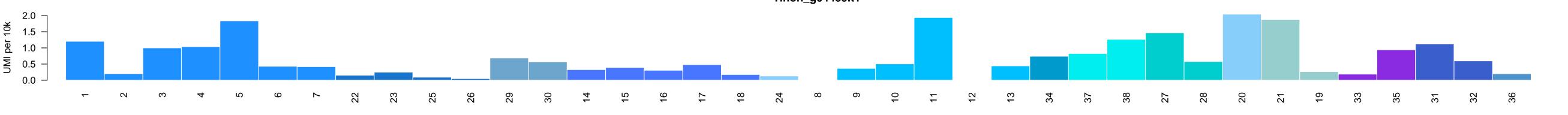
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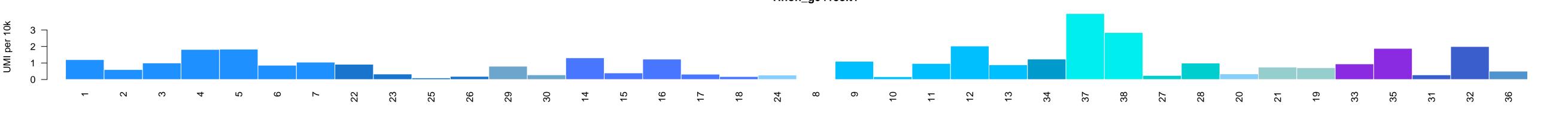
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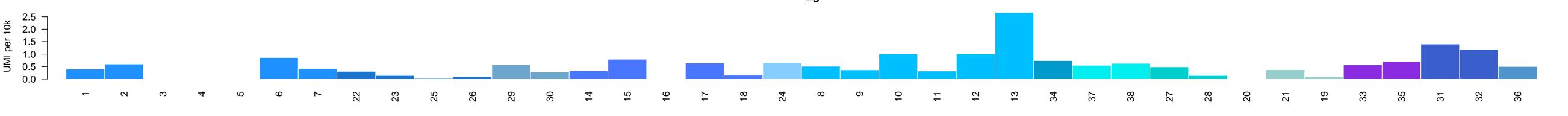
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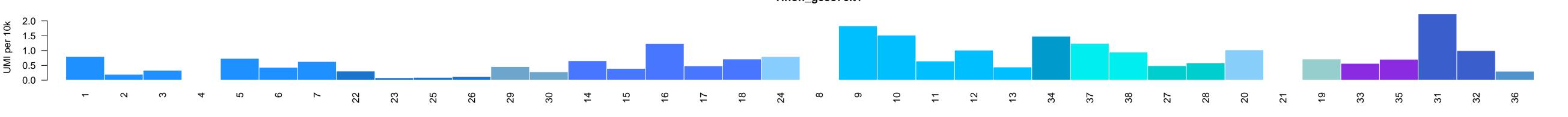
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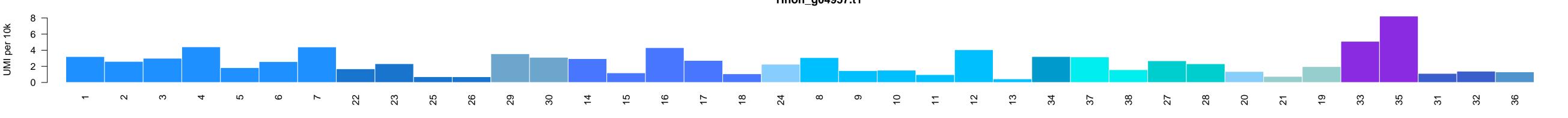


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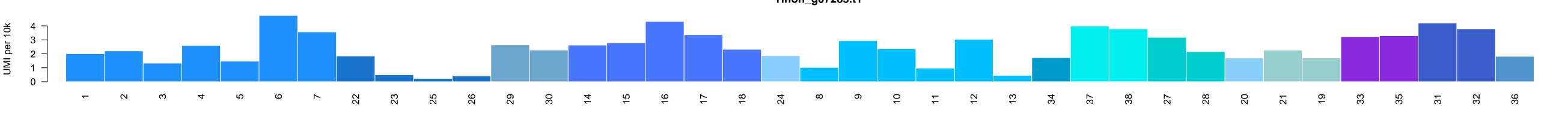


DEAD.HG3.14:DHX15 Hhon\_g03332.t1 3.5 - 3.0 - 2.5 - 2.0 - 1.5 - 1.5 - 3.5 - 

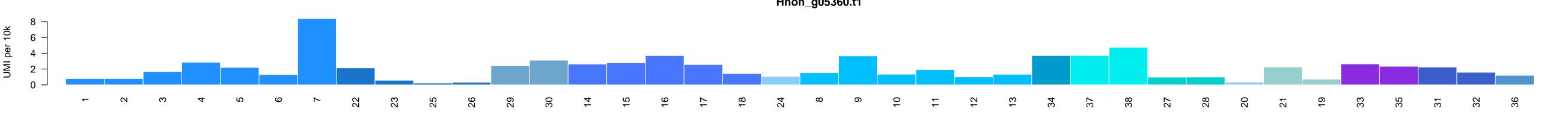
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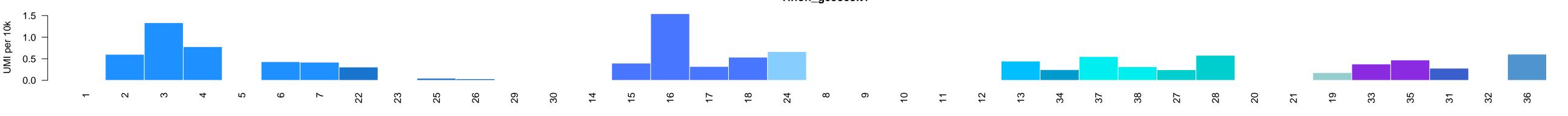
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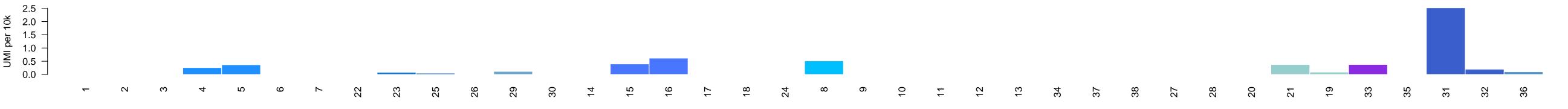
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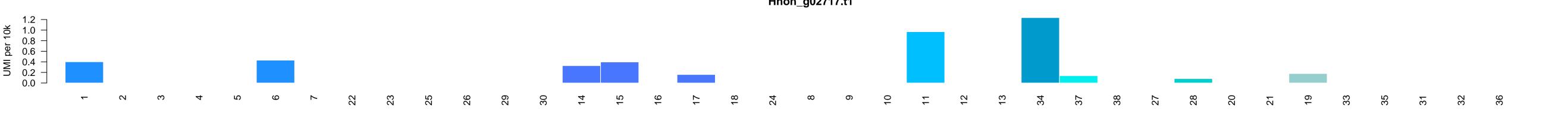
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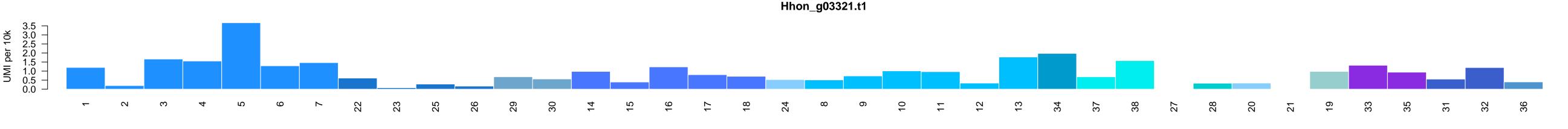
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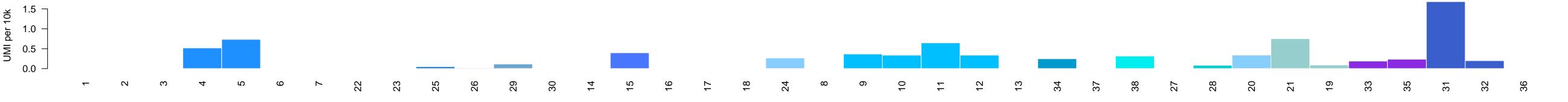
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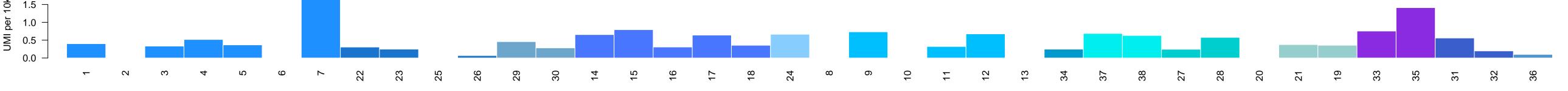
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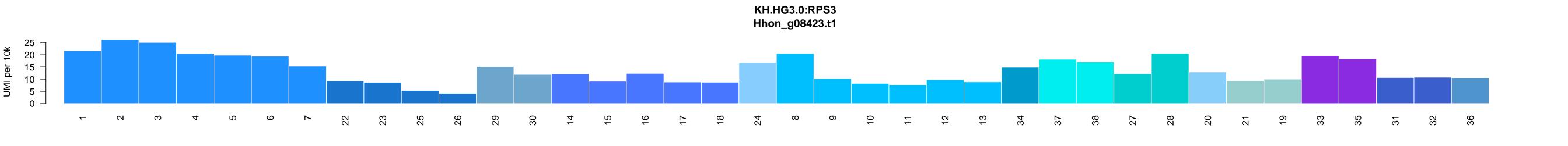


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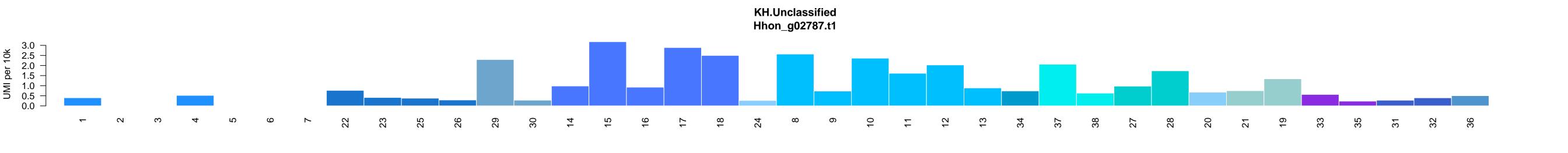
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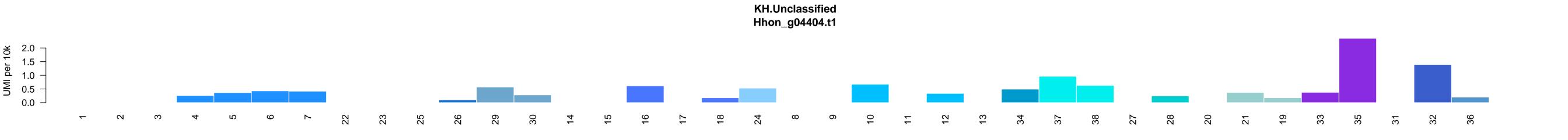




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Hhon\_g01752.t1 



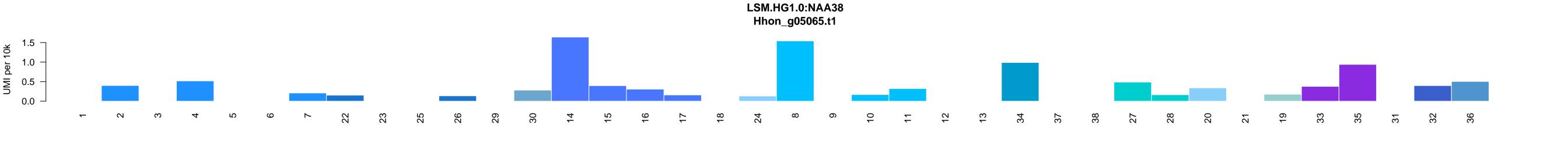


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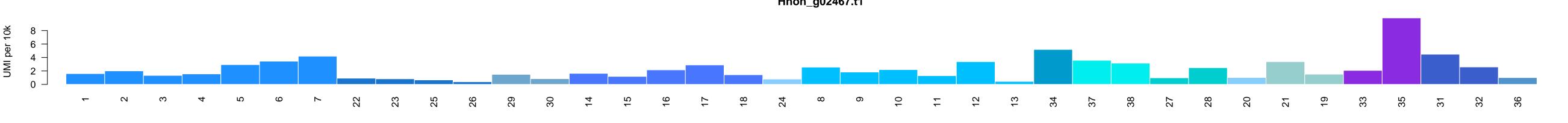
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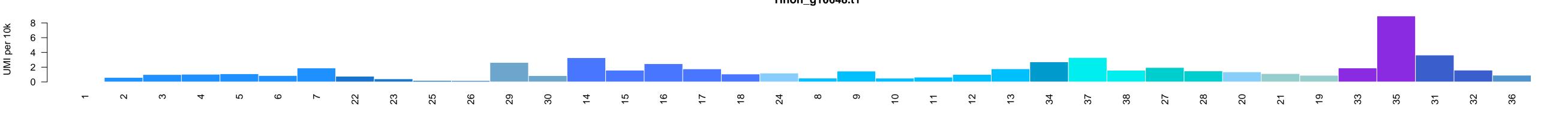
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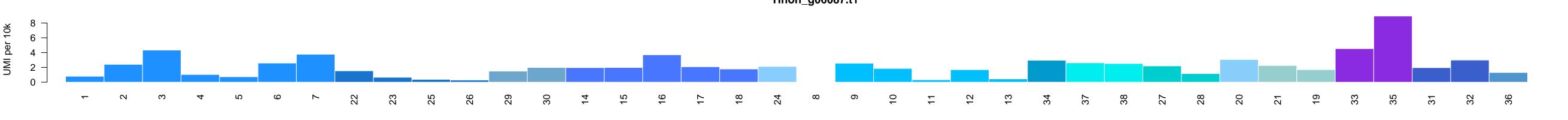
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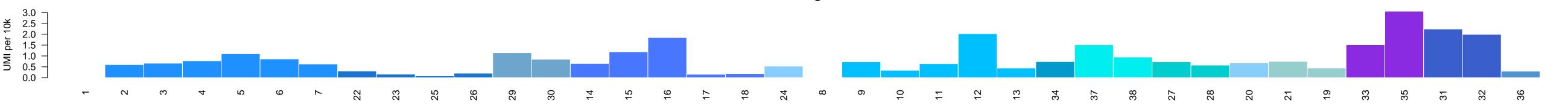
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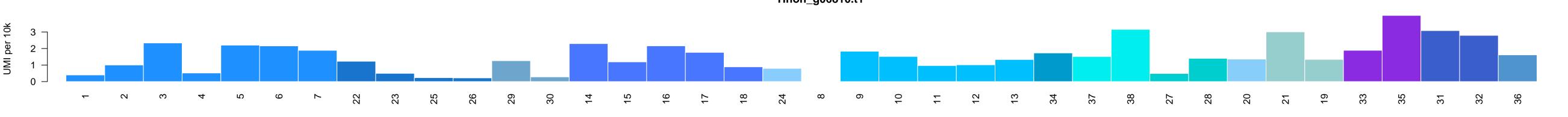
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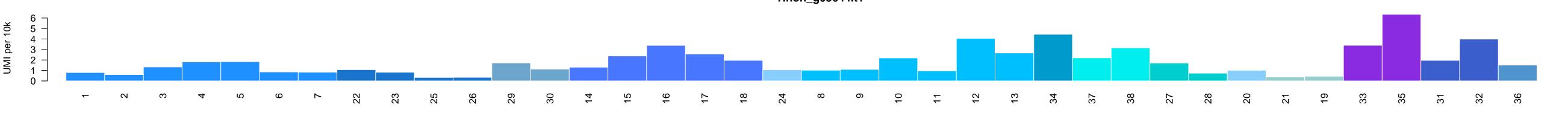
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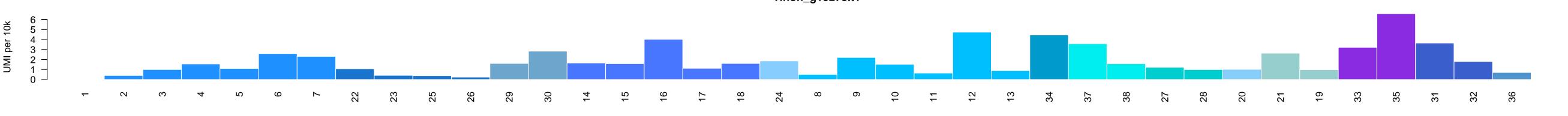
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LSM.HG1.14:AP000356.5/SNRPD3 Hhon\_g03644.t1



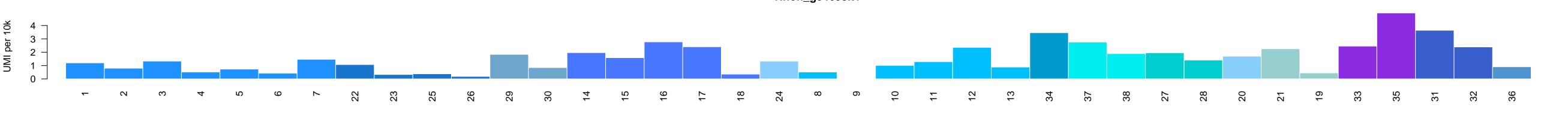
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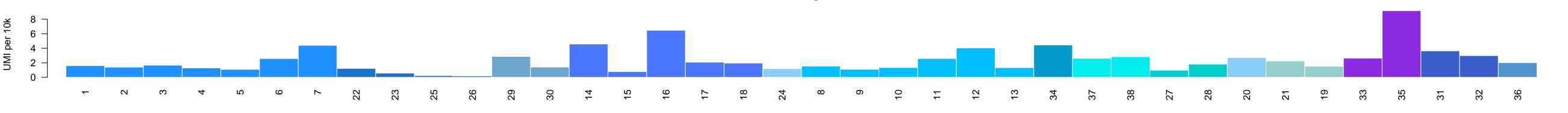
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Hhon\_g02424.t1 

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LSM.HG1.4:SNRPG Hhon\_g04415.t1

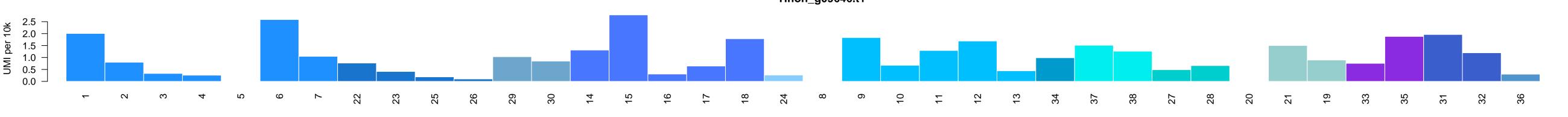


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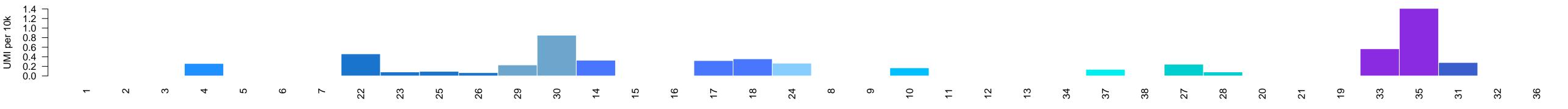
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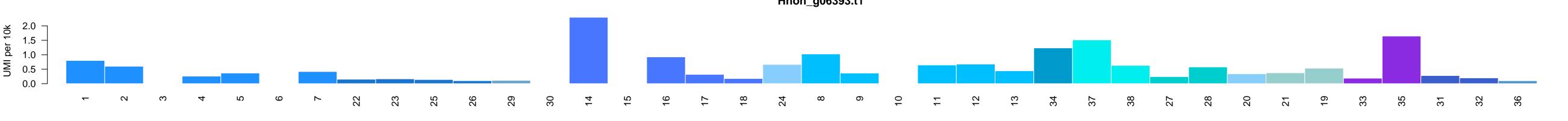


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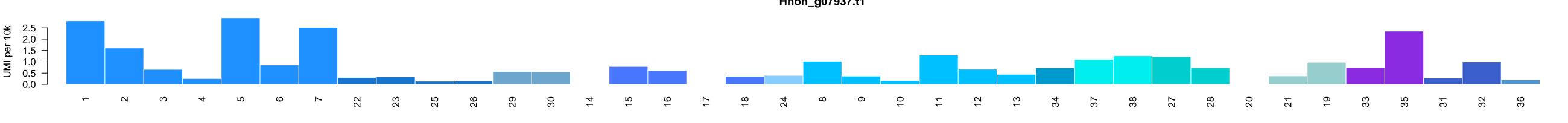
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PUA.HG3.1:MCTS1/MCTS2P Hhon\_g06393.t1

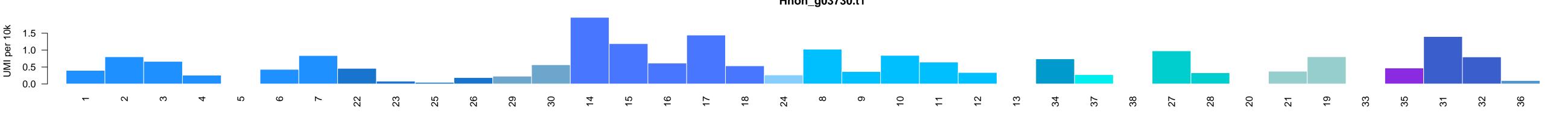


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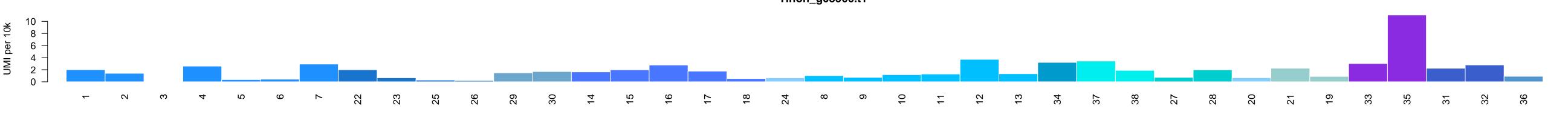


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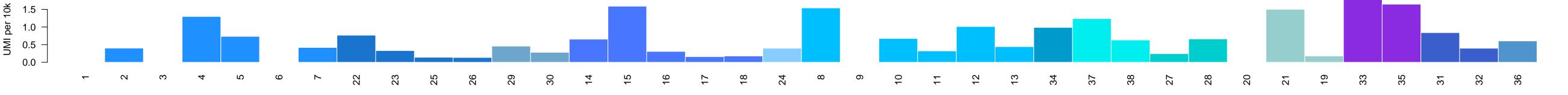
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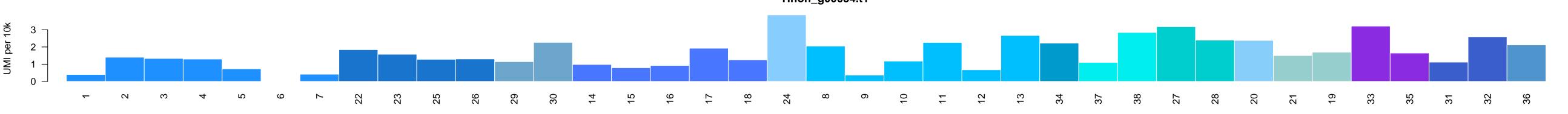
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Hhon\_g08566.t1



RRM.HG1.19:SF3B4
Hhon\_g03429.t1

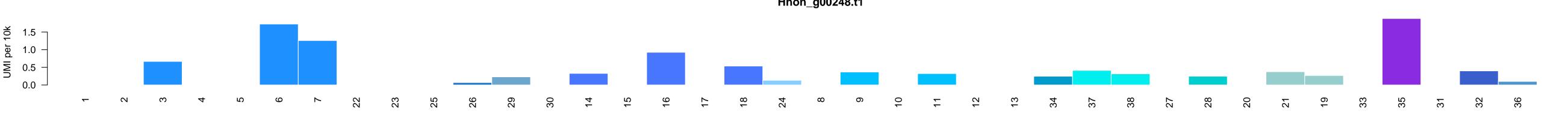


RRM.HG1.24:PABPC1/PABPC1L/PABPC1L2A/PABPC1L2B/PABPC3/PABPC4/PABPC4L/PABPC5
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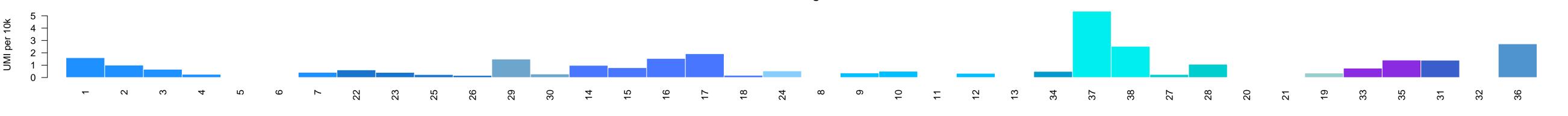


RRM.HG1.26:TIA1/TIAL1 Hhon\_g02252.t1 文 3 기 

RRM.HG1.27:TRNAU1AP Hhon\_g00248.t1

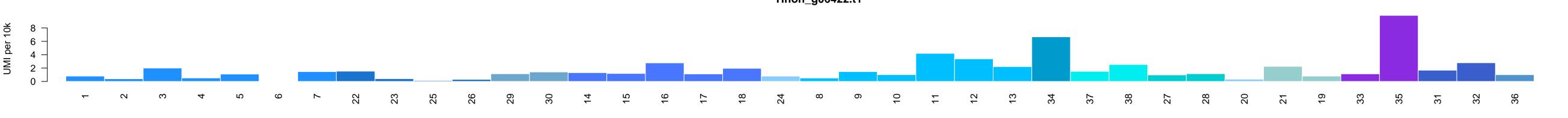


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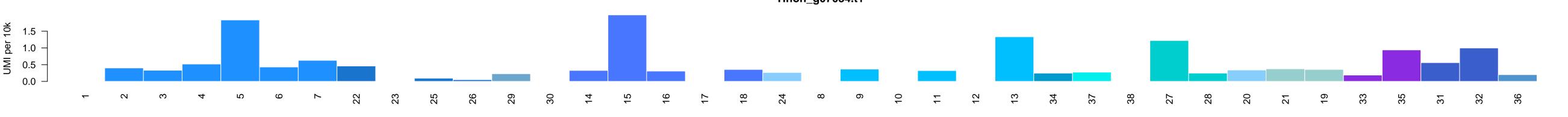


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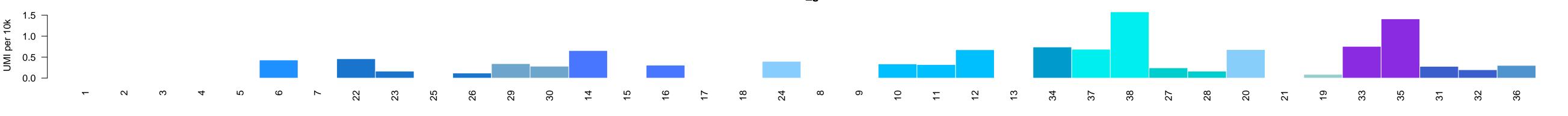
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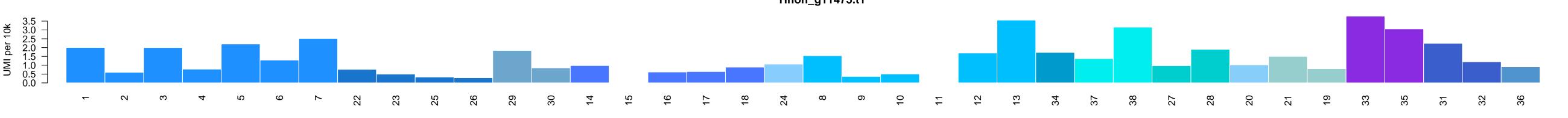
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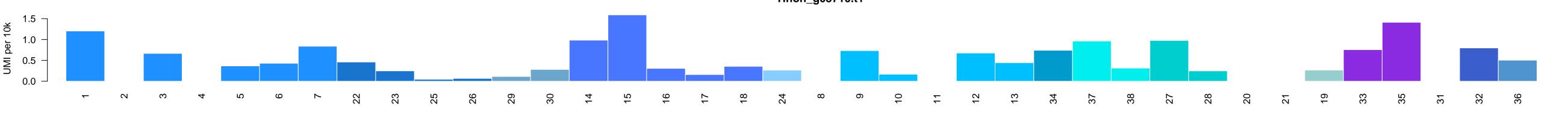
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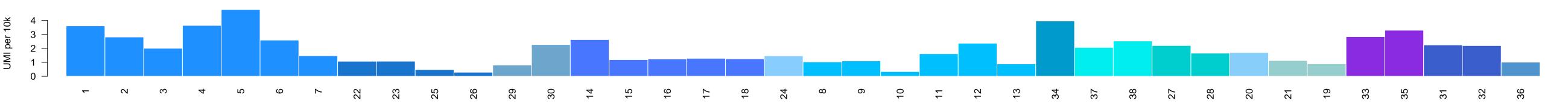
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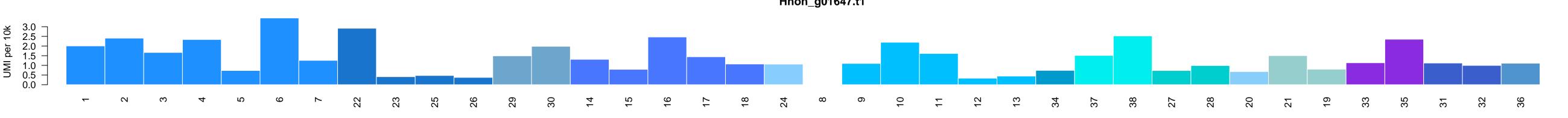
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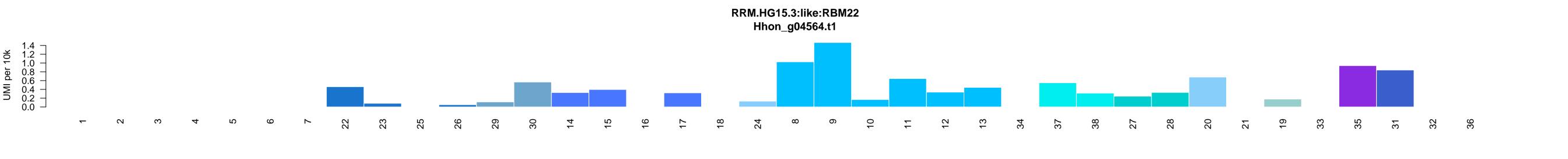
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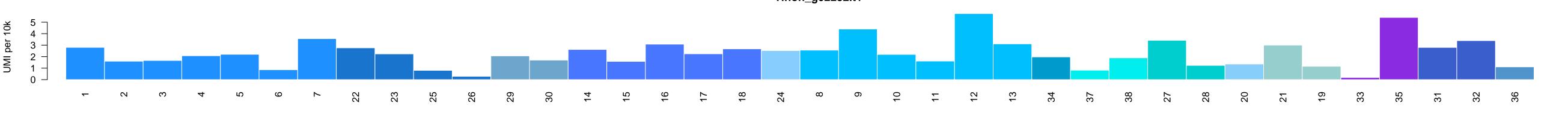
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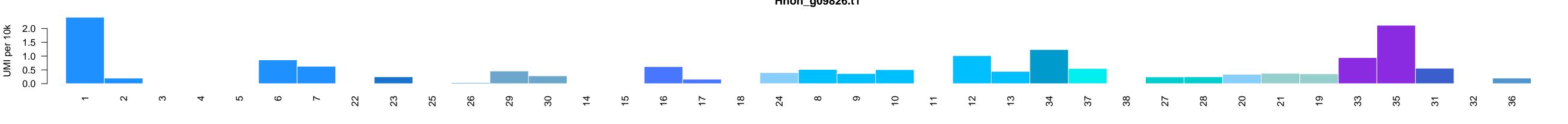
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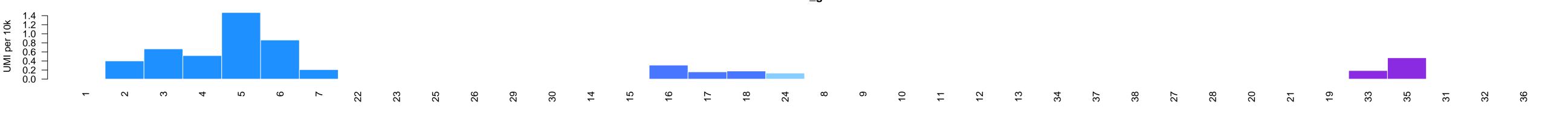


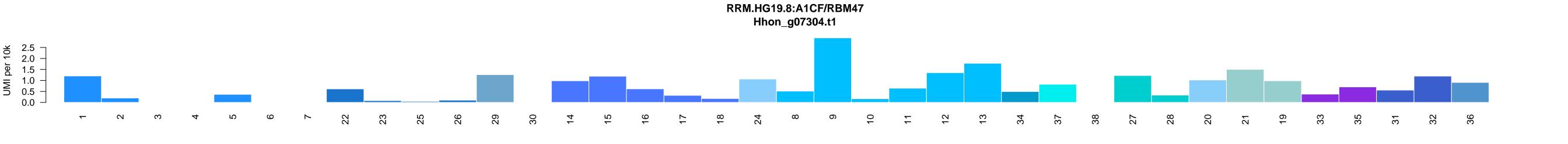
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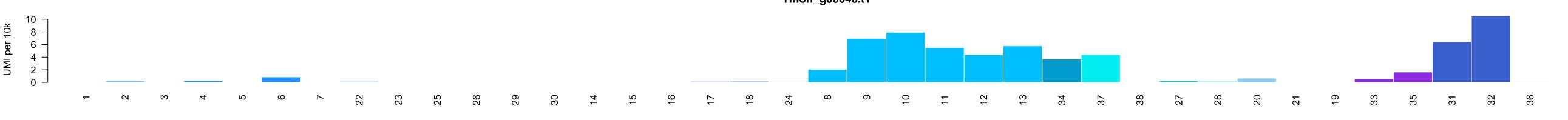
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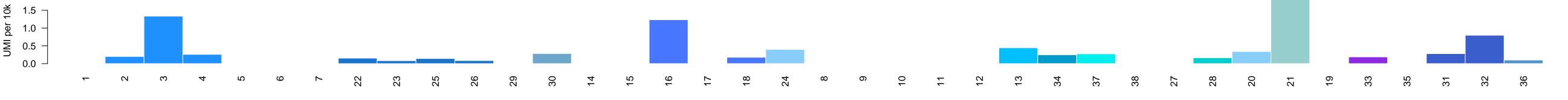




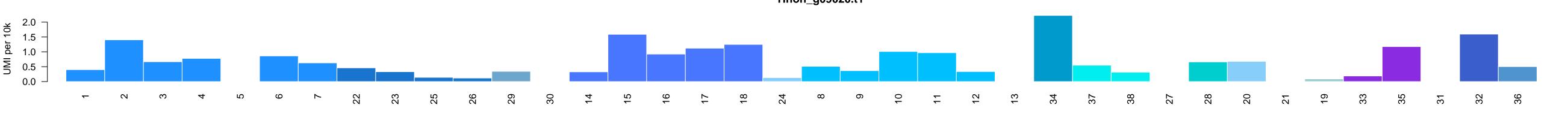
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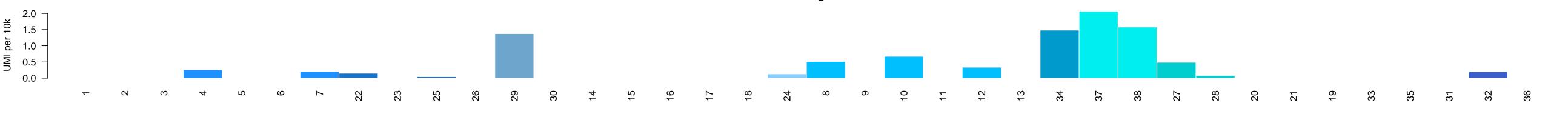
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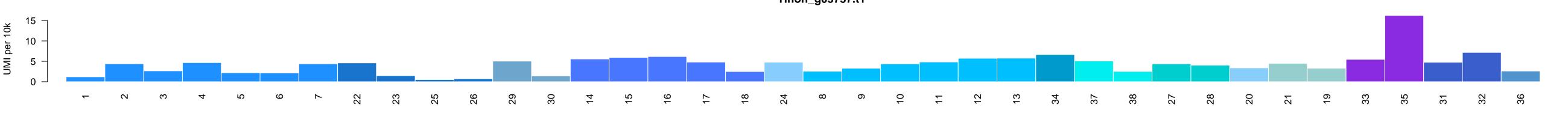
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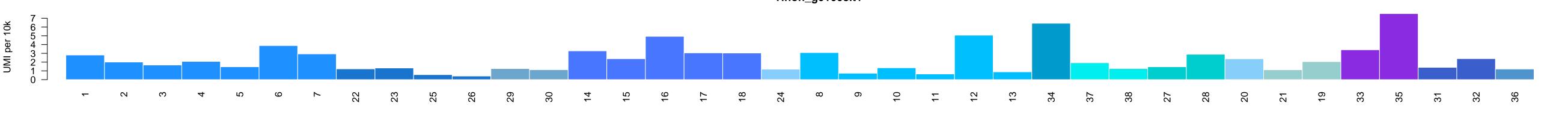
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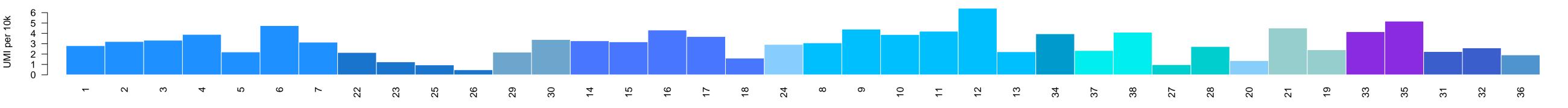
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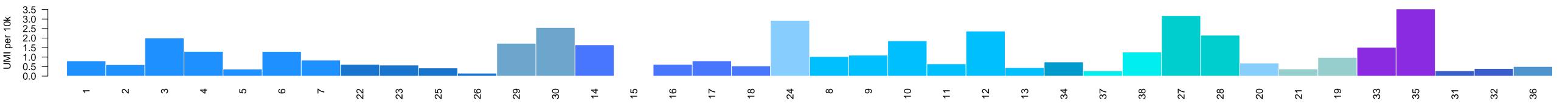
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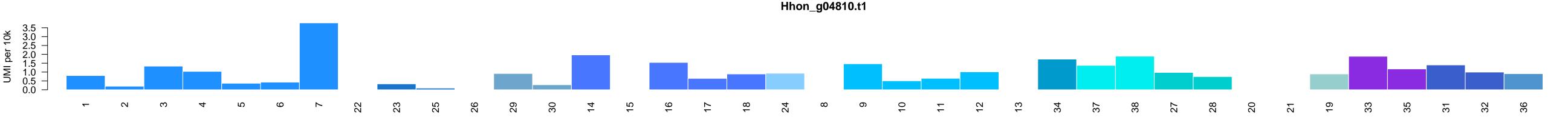
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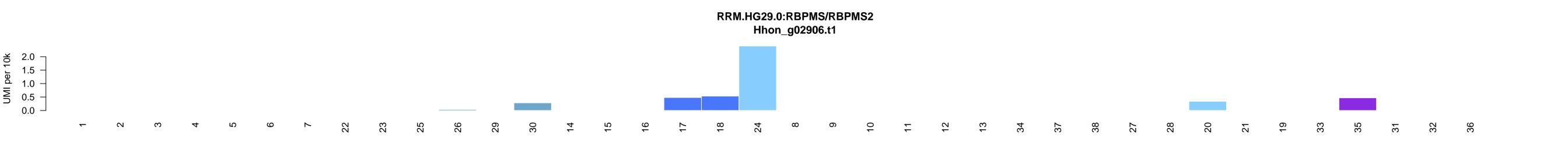
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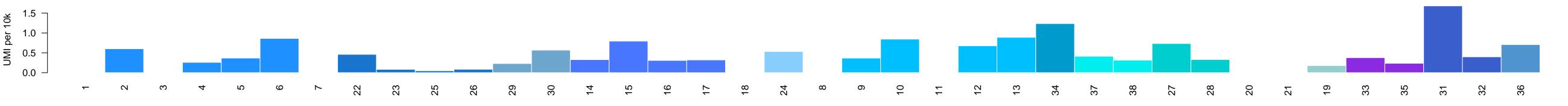
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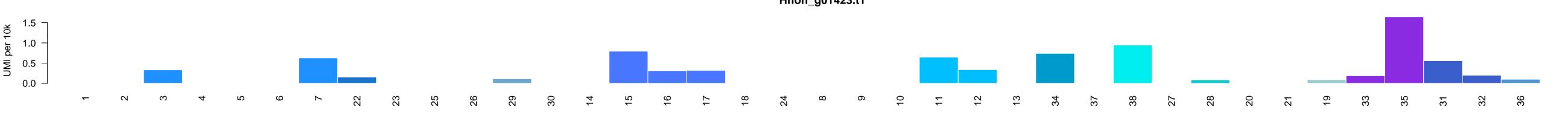
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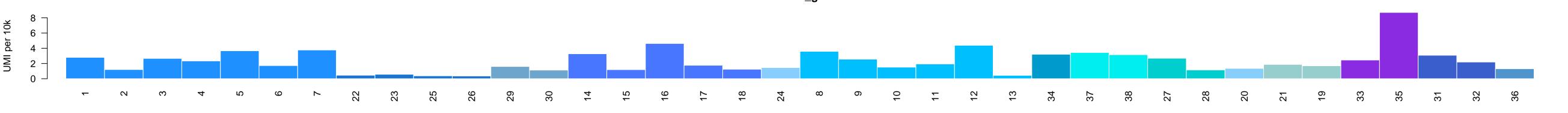


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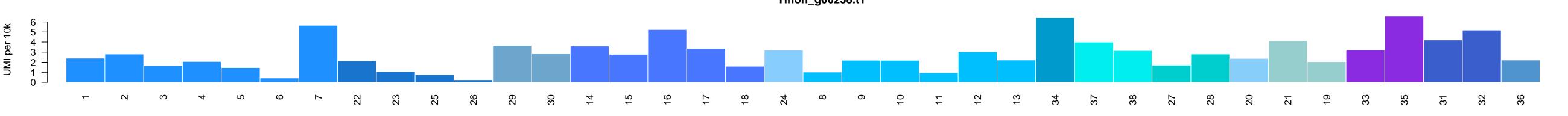


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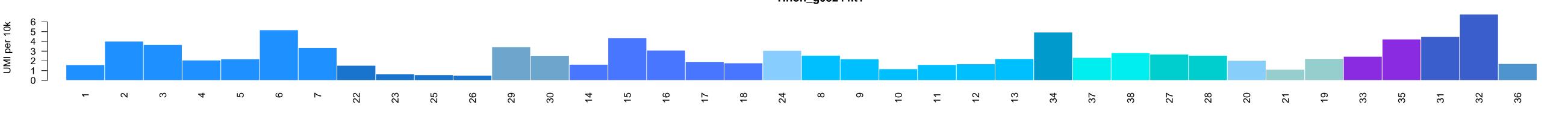
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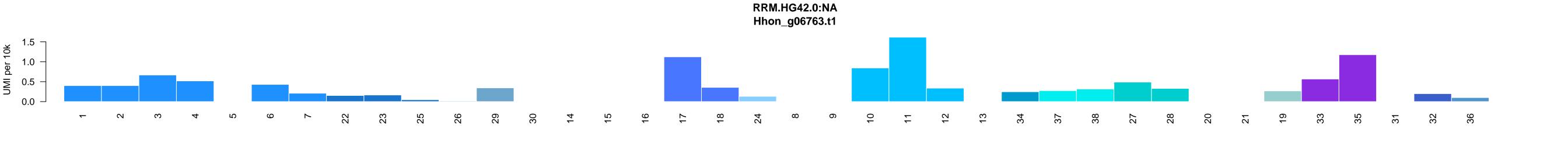


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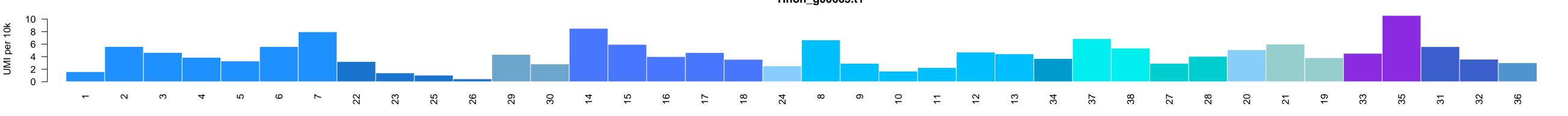


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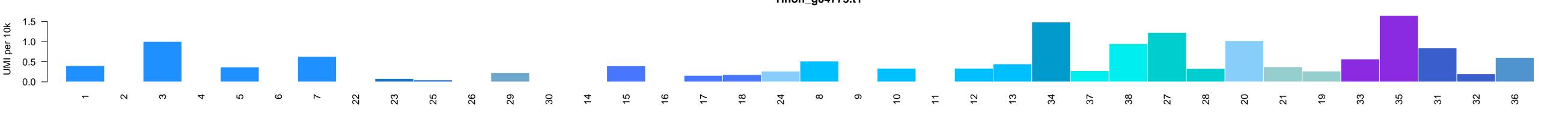




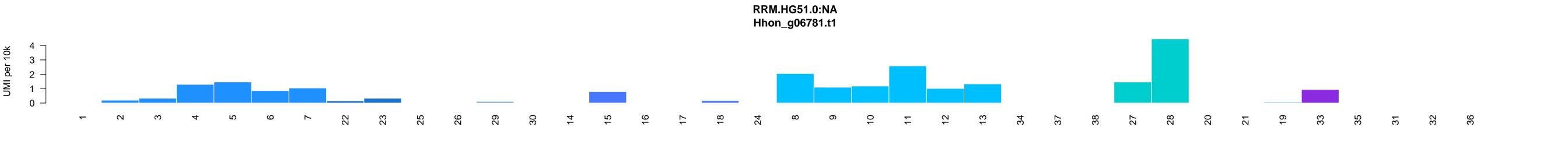
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RRM.HG5.11:U2AF2 Hhon\_g04775.t1

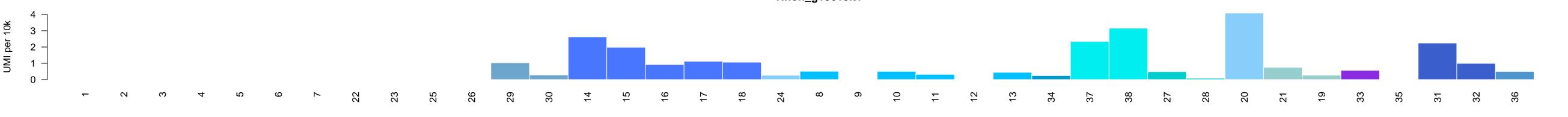


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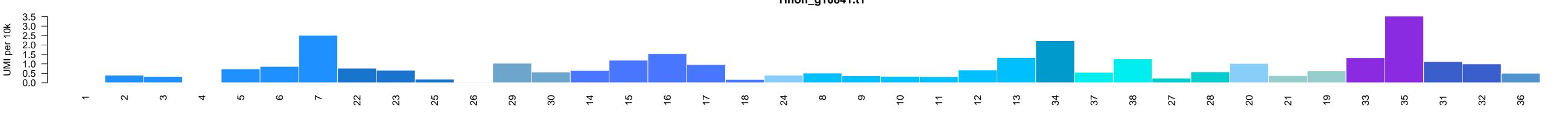


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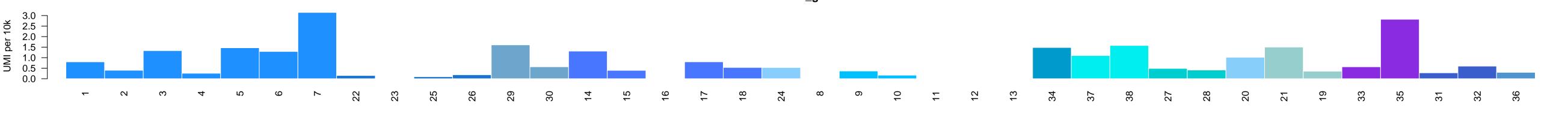
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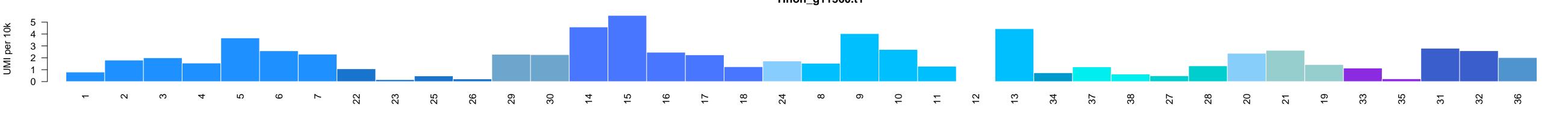
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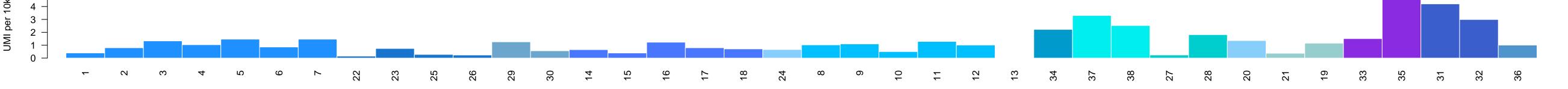


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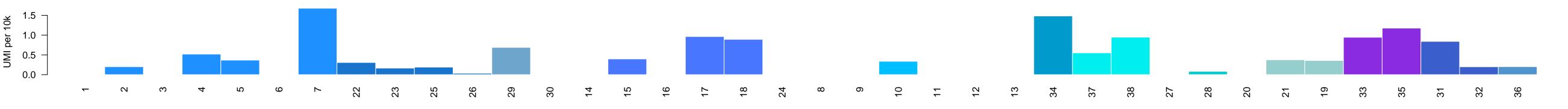




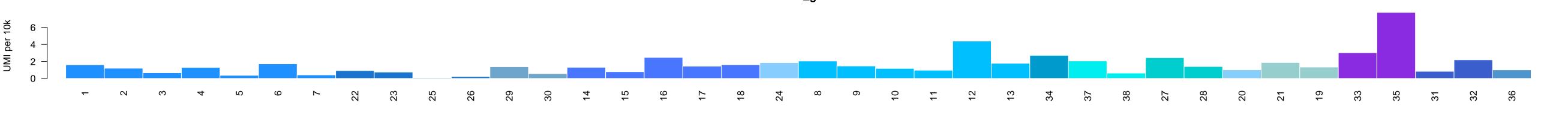
RRM.HG8.0:RNPS1
Hhon\_g04811.t1

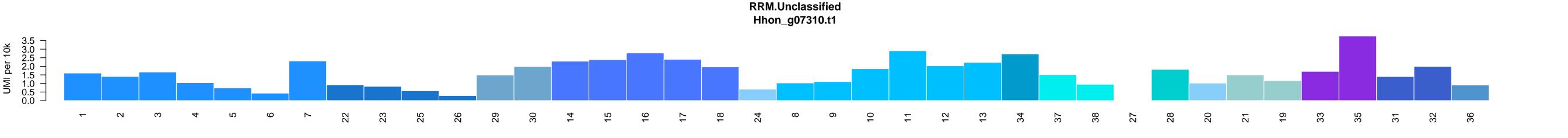


RRM.HG8.8:SRSF2/SRSF8 Hhon\_g01419.t1



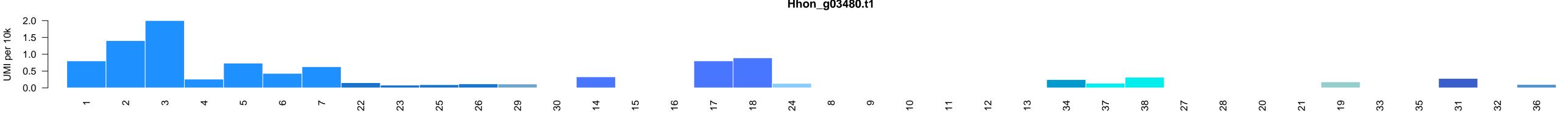
HG9.9:like:SRSF3/SRSF7 Hhon\_g09736.t1





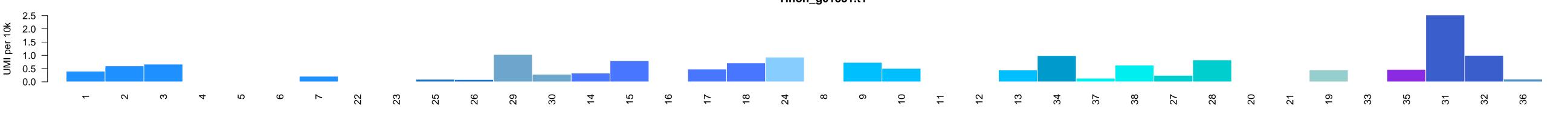
Hhon\_g04631.t1 3.0 7 2.5 -2.0 -1.5 -1.0 -

SAM.HG1.17:CASKIN1/CASKIN2 Hhon\_g03480.t1

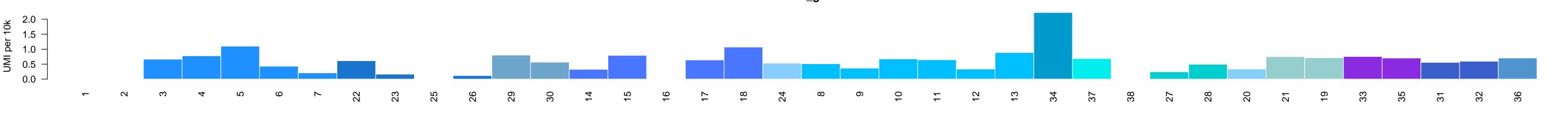


SAM.HG1.6:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6

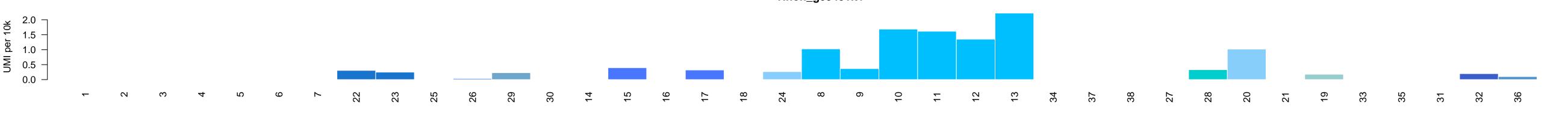
Hhon\_g01331.t1



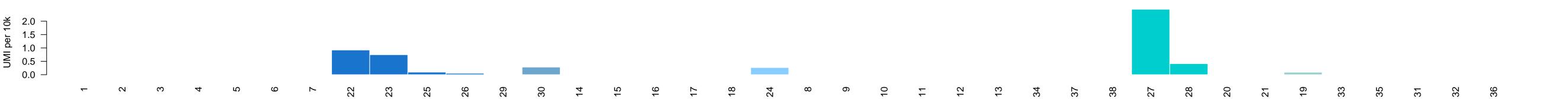
SAM.HG13.1:SAMD4A/SAMD4B Hhon\_g03982.t1



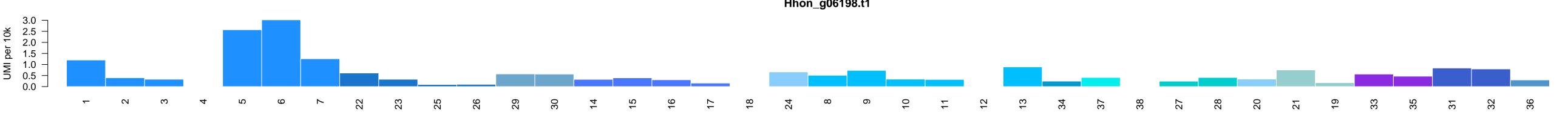
SAM.HG2.16:like:SAMD10/SAMD12 Hhon\_g03431.t1

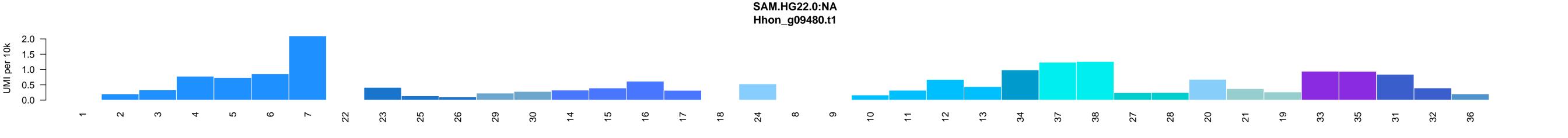


## SAM.HG2.22:like:SAMD10/SAMD12 Hhon\_g00609.t1

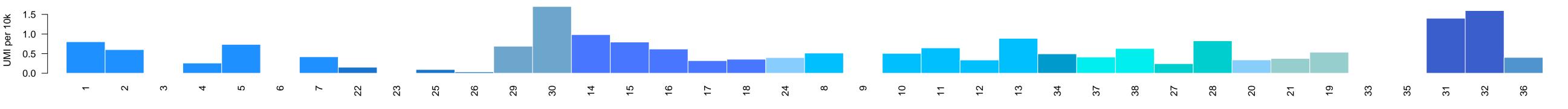


SAM.HG2.7:DGKD/DGKH Hhon\_g06198.t1

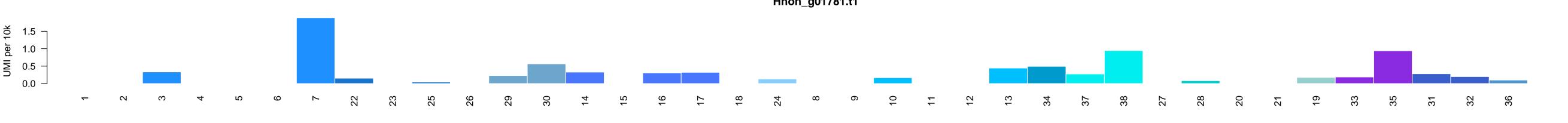


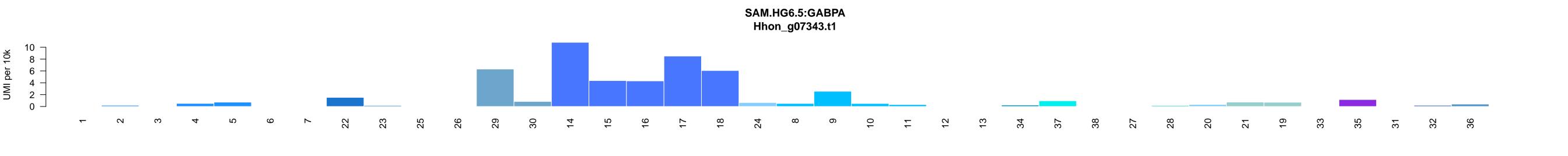


SAM.HG3.0:LRSAM1 Hhon\_g09700.t1



SAM.HG4.6:SAMD7/SAMD11 Hhon\_g01781.t1

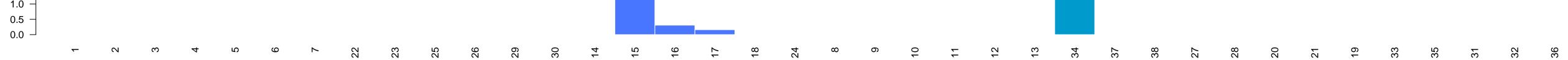




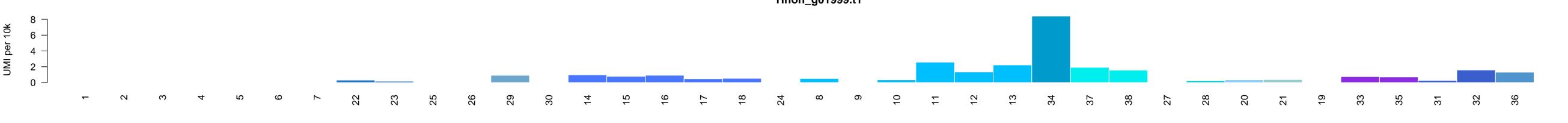
SAM.HG6.6:ETS1/ETS2
Hhon\_g07347.t1



SAM.HG7.0:PPFIBP2
Hhon\_g03133.t1



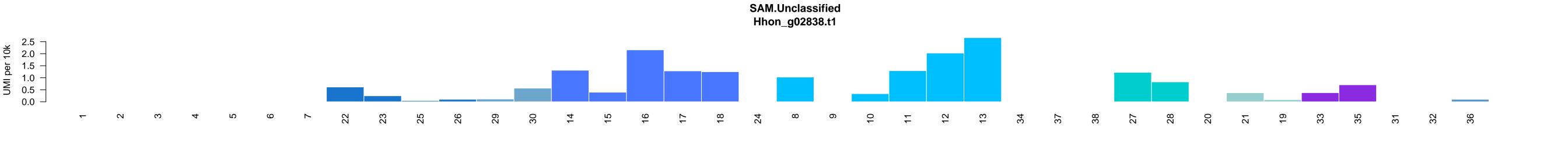
SAM.HG7.3:PPFIA1/PPFIA2/PPFIA3/PPFIA4 Hhon\_g01999.t1



SAM.HG8.2:SAMD5
Hhon\_g05434.t1

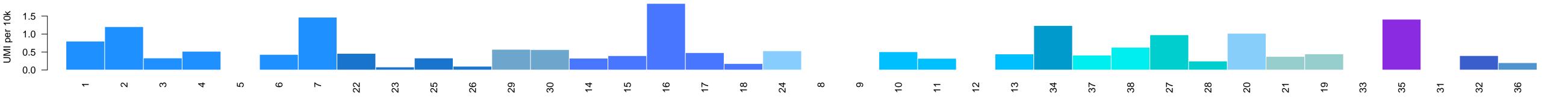
출 1.5 기



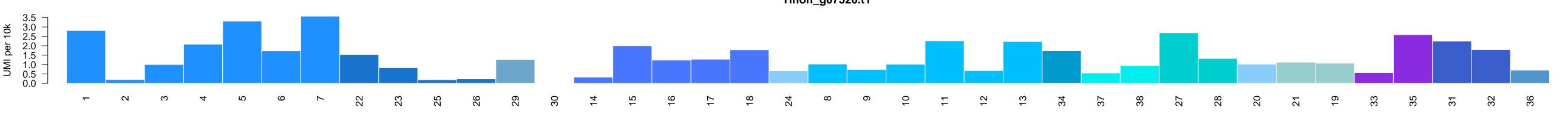


Hhon\_g06904.t1 

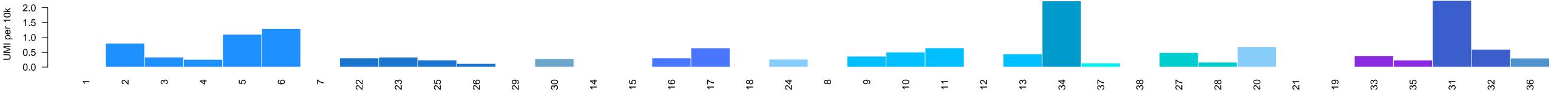
SURP.HG1.4:SF3A1 Hhon\_g03694.t1



YTH.HG1.3:YTHDC1 Hhon\_g07520.t1



zf-CCCH.HG1.5:ZC3H15 Hhon\_g10263.t1



zf-CCCH.HG2.0:like:AC091959.1/HELZ/RBM27/RC3H1/RC3H2/RNF113A/RNF113B/UNK/UNKL/ZC3H7A/ZC3H7B/ZC3H13/ZFP36/ZFP36L1/ZFP36L2/ZMAT5
Hhon\_g04302.t1



zf-nanos.HG1.1:NANOS1/NANOS2/NANOS3 Hhon\_g01355.t1