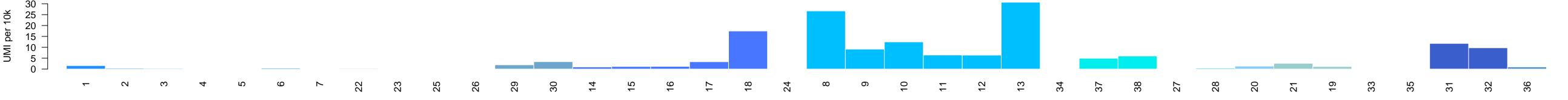
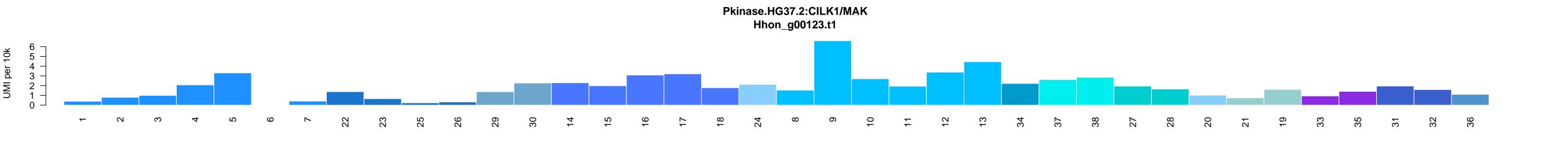
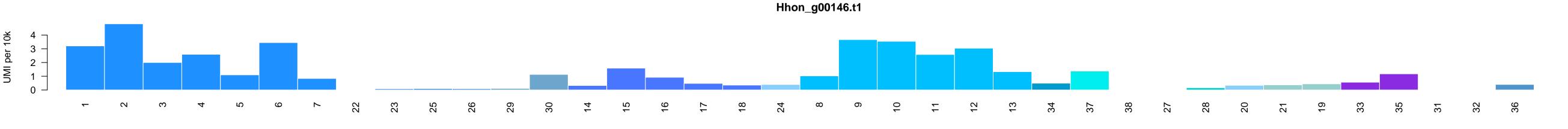
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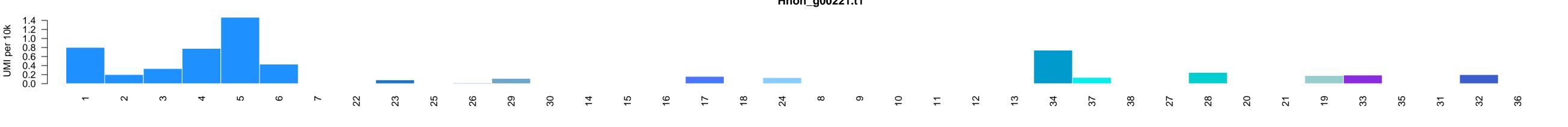


HG5.21:MOS _g00146.t1

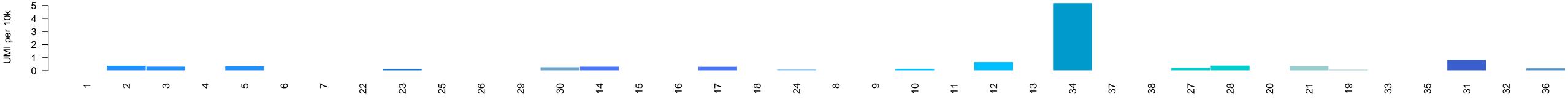


Pkinase.HG2.34:NIM1K Hhon_g00182.t1 per 10k

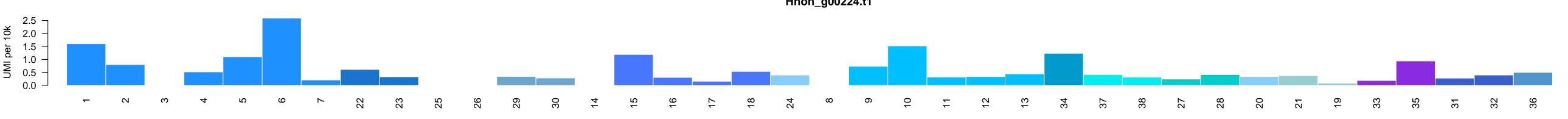
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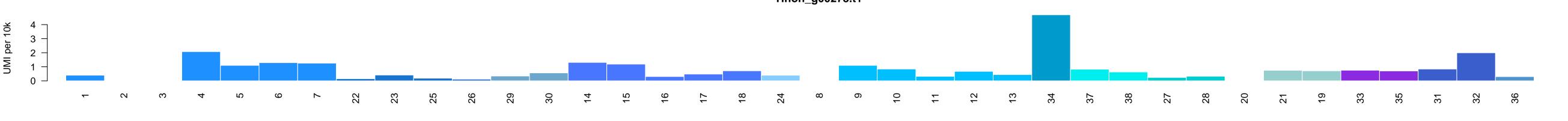
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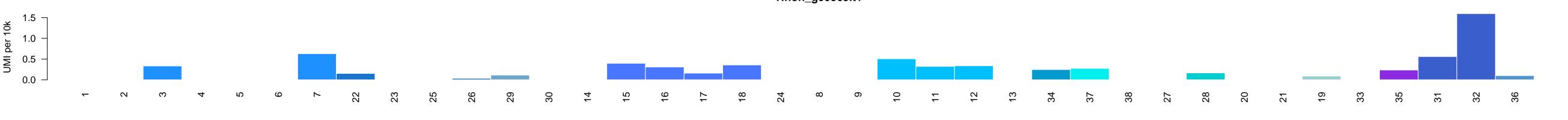
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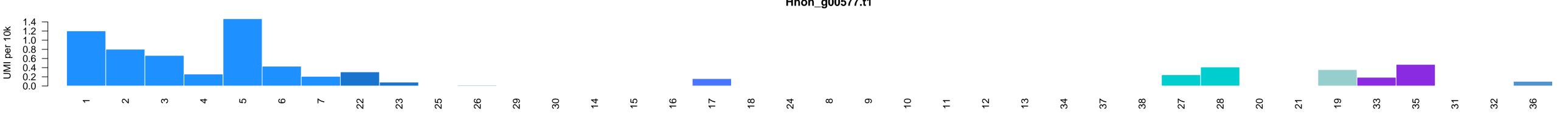
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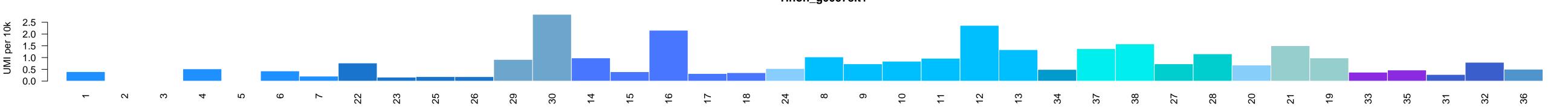
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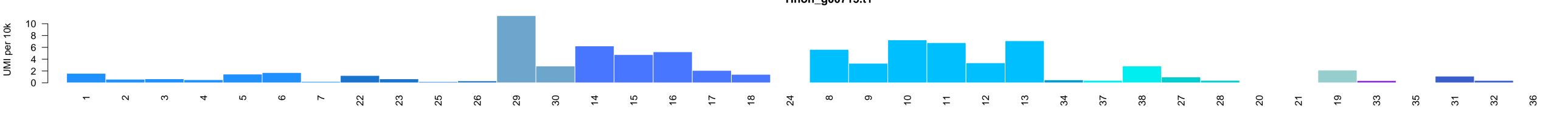
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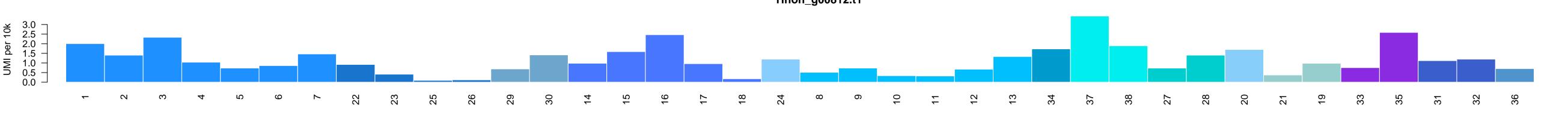
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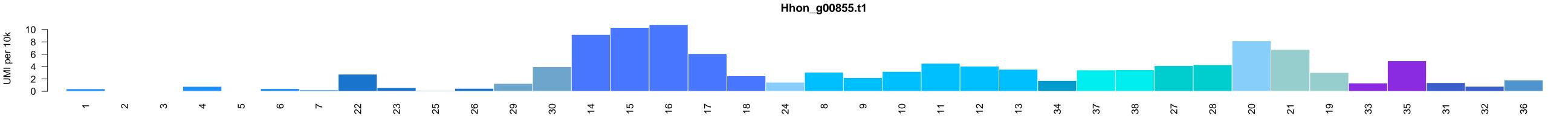
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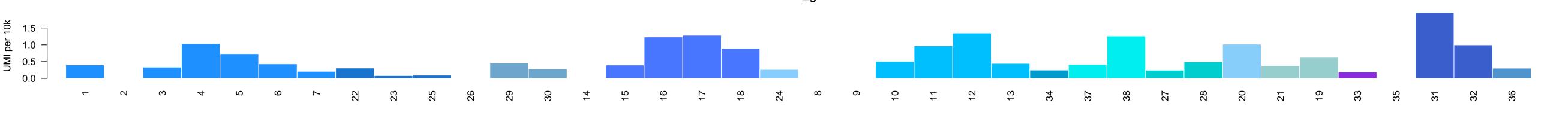
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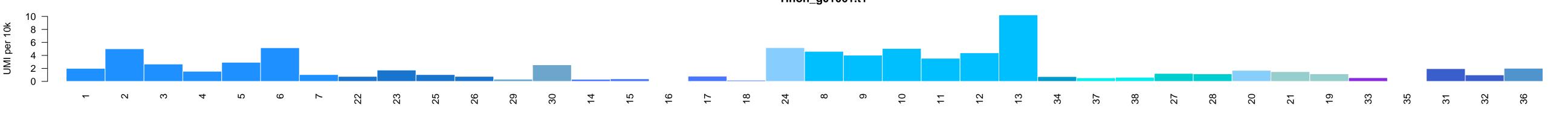
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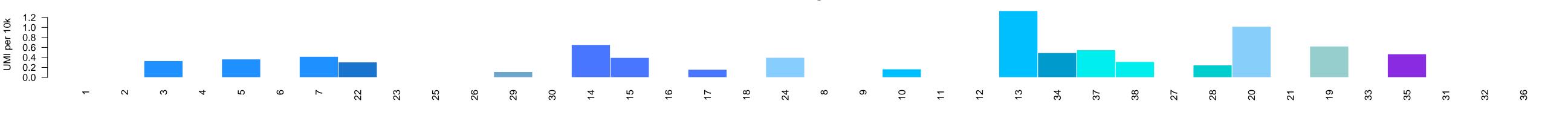
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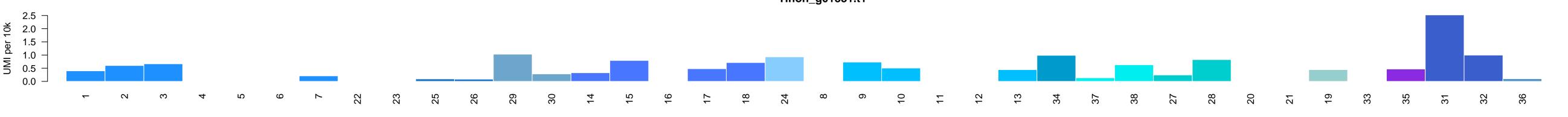
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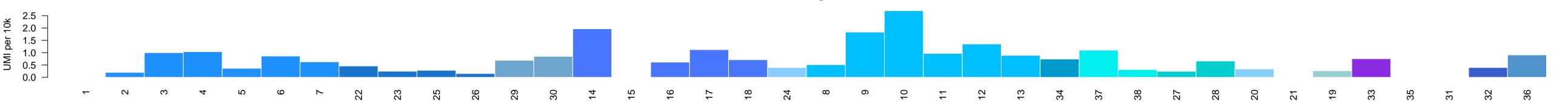
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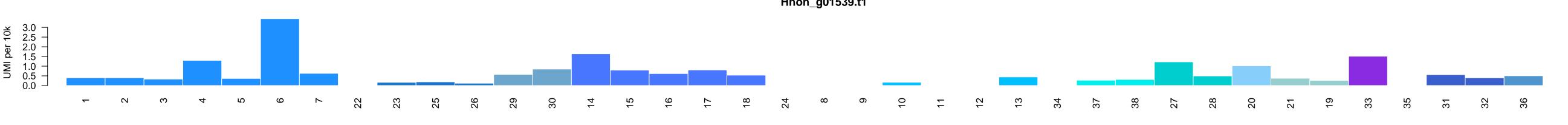
Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
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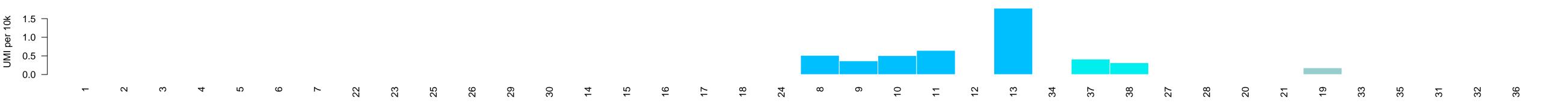
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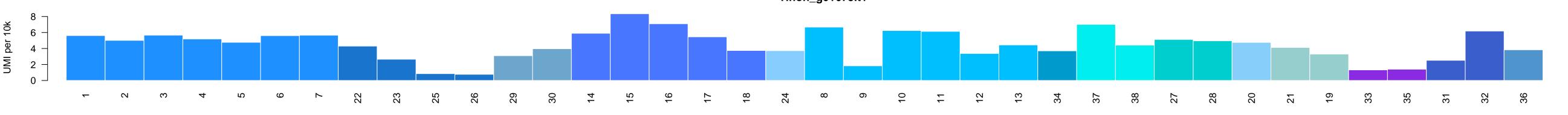
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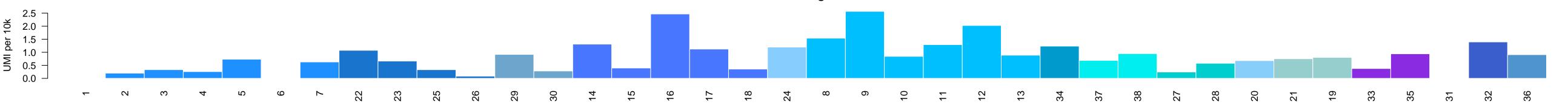
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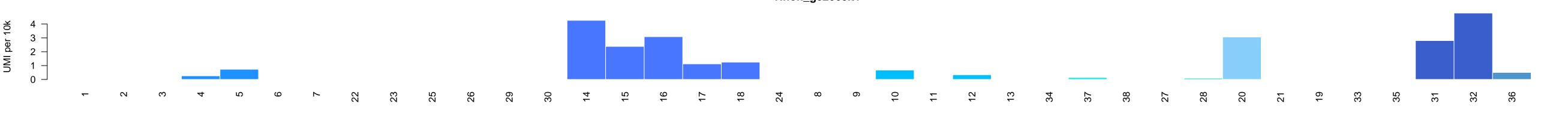
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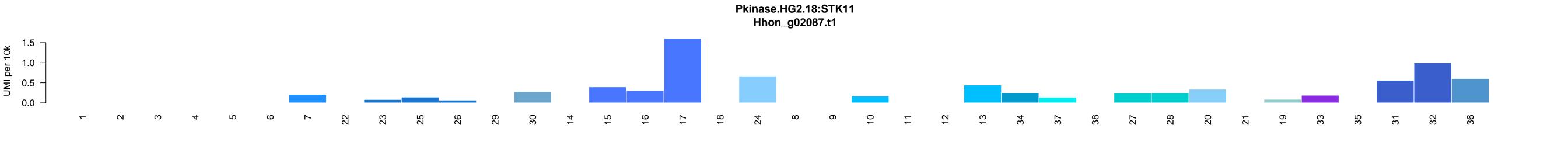


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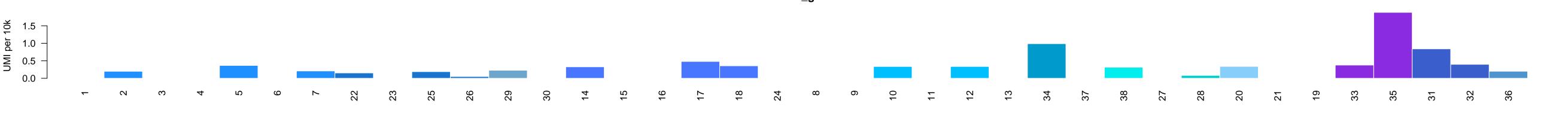


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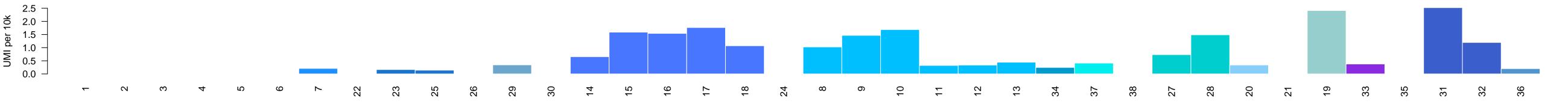




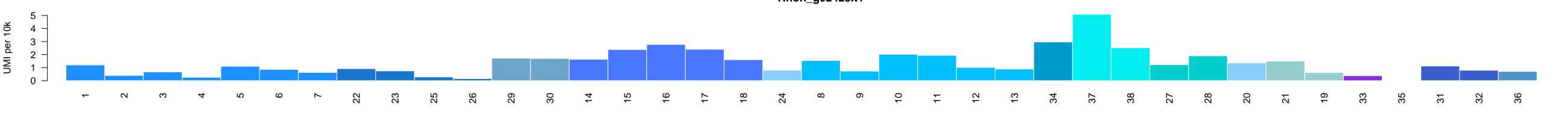
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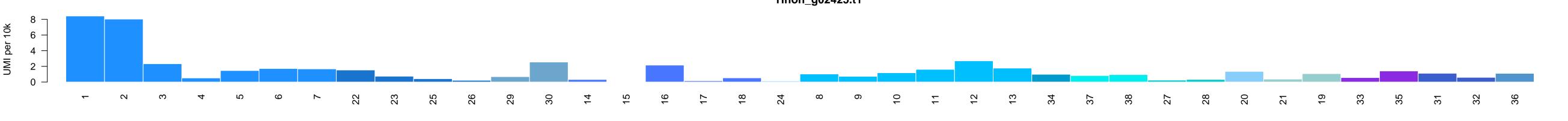
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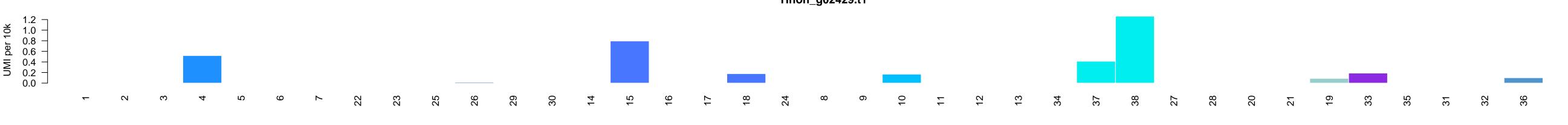
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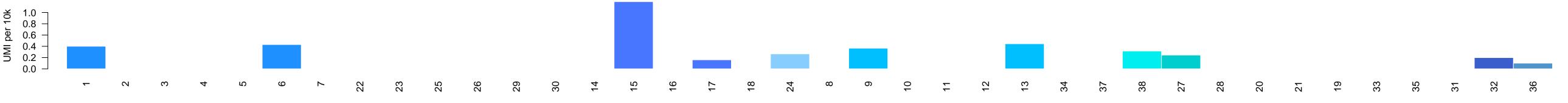
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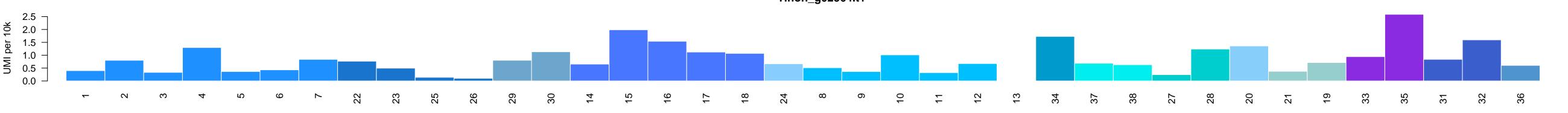
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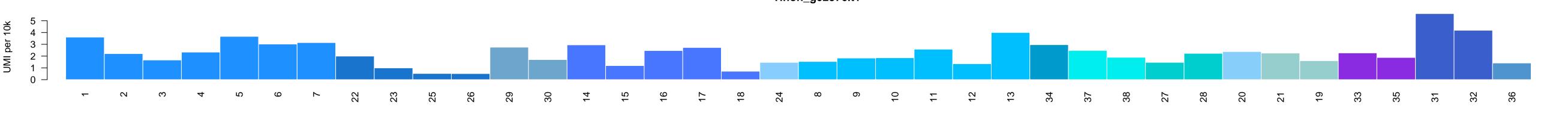
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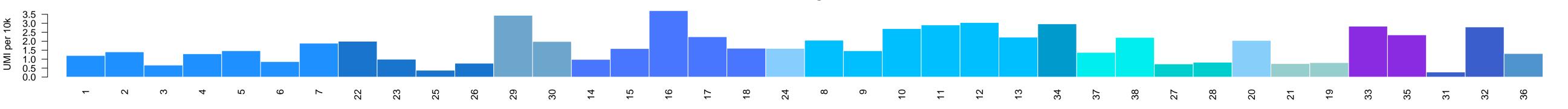
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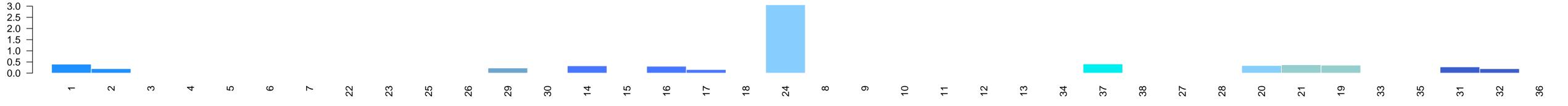
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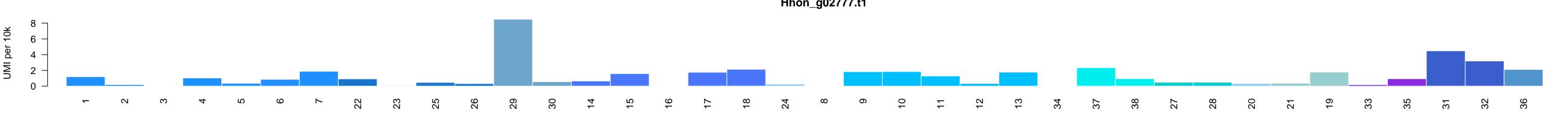
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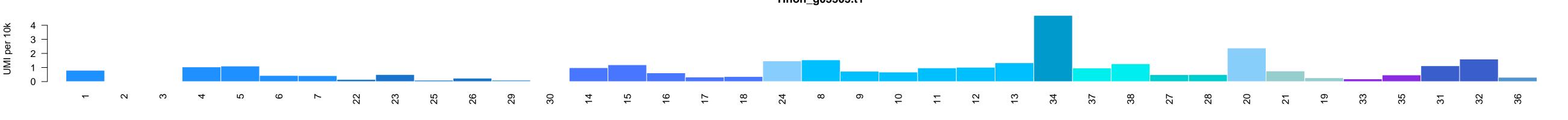
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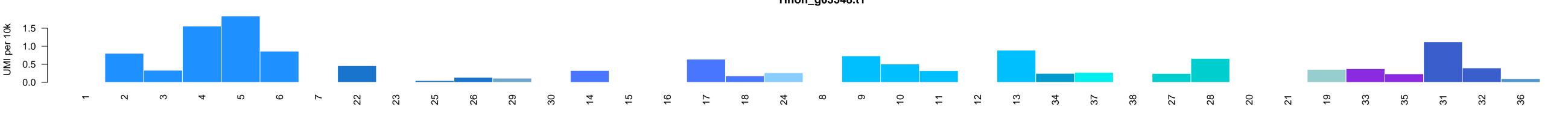
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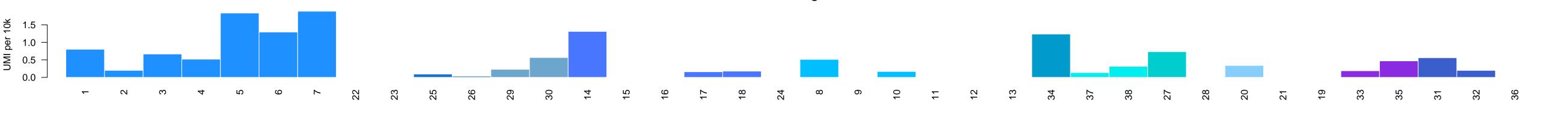
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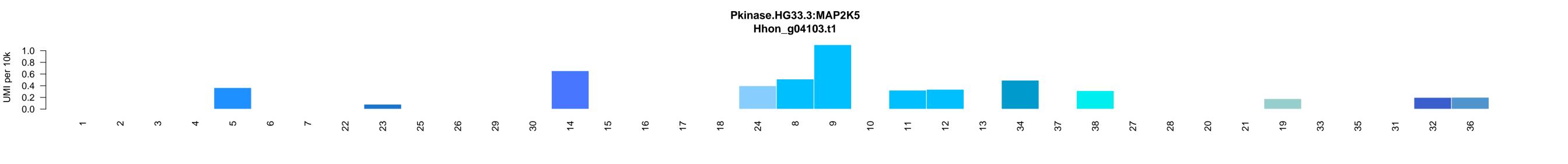
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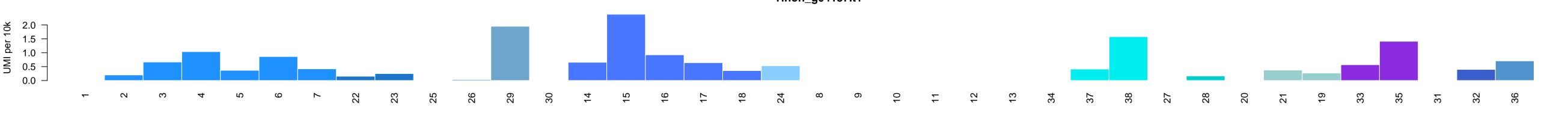
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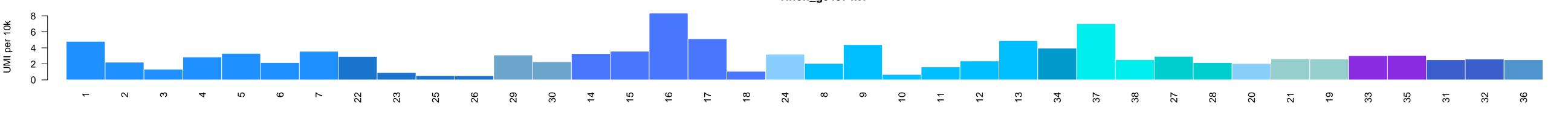
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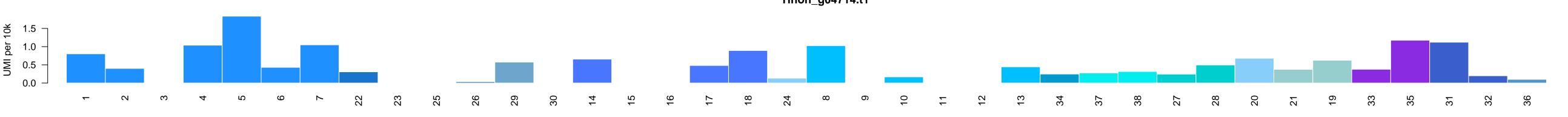
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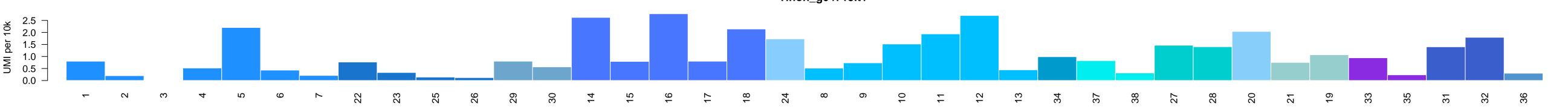
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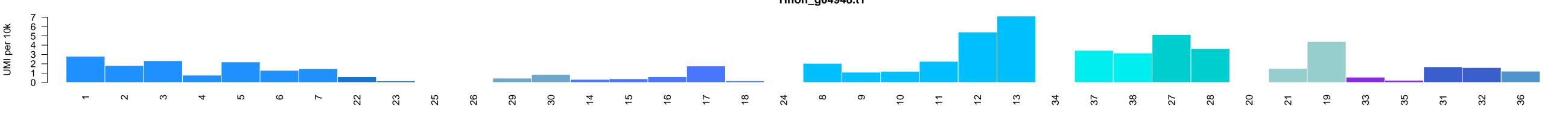
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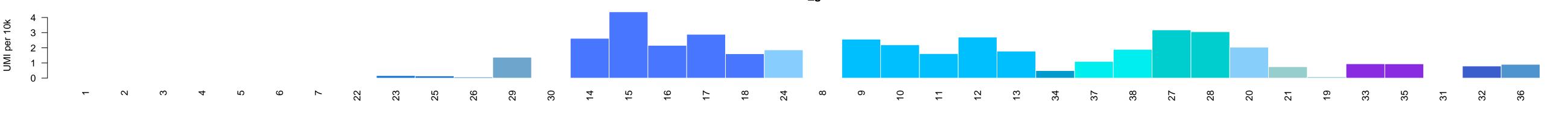
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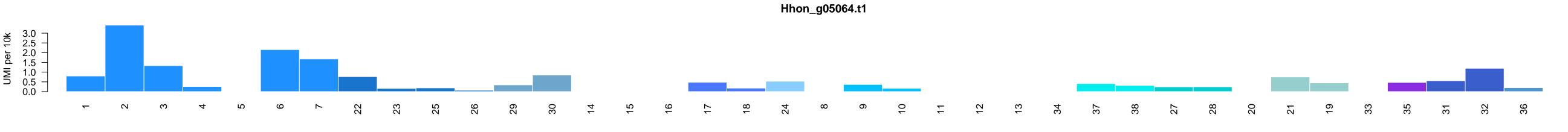
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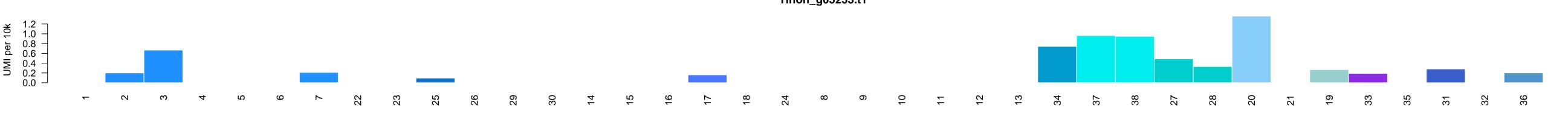
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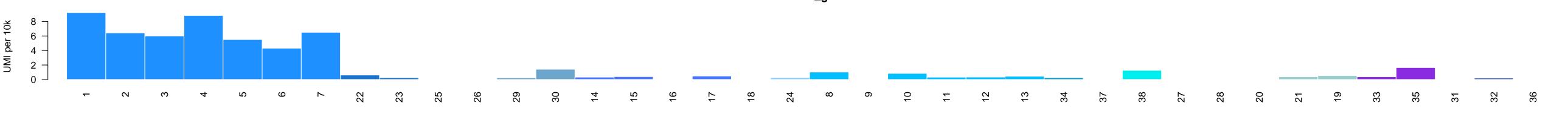
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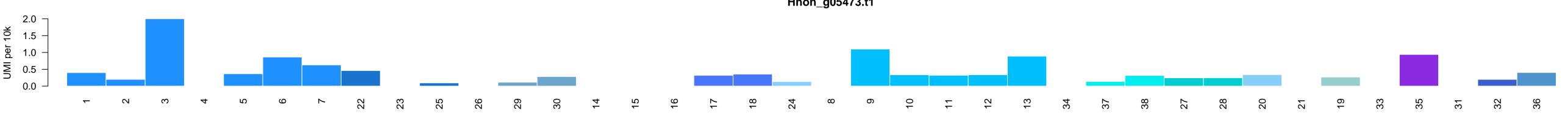
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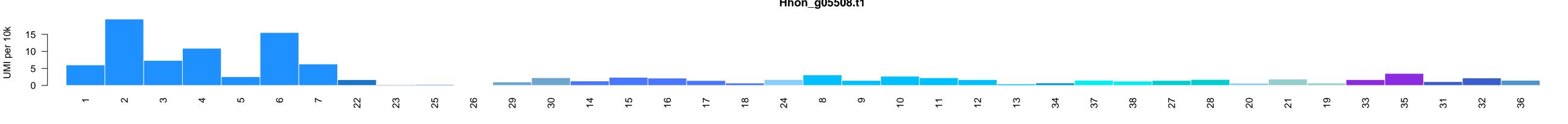
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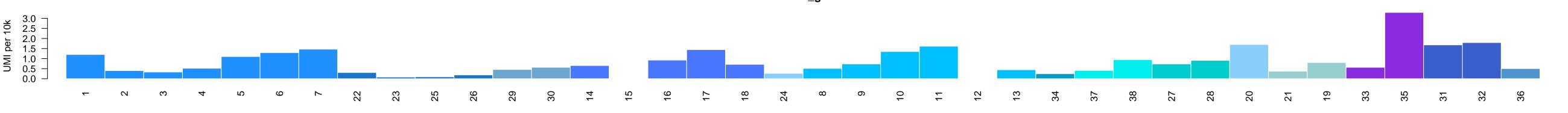
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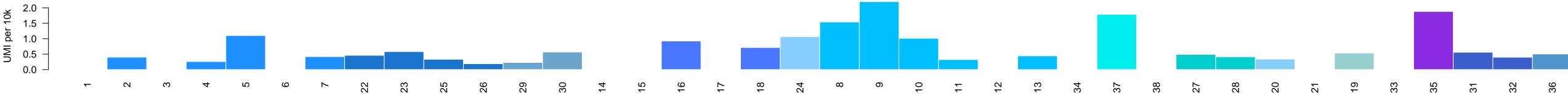
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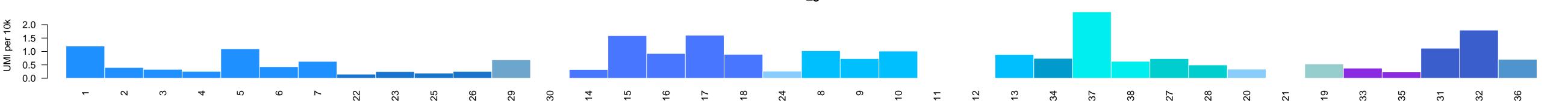
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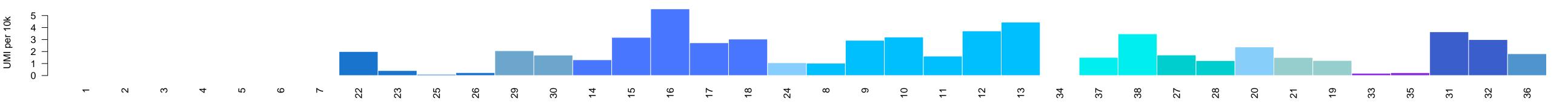
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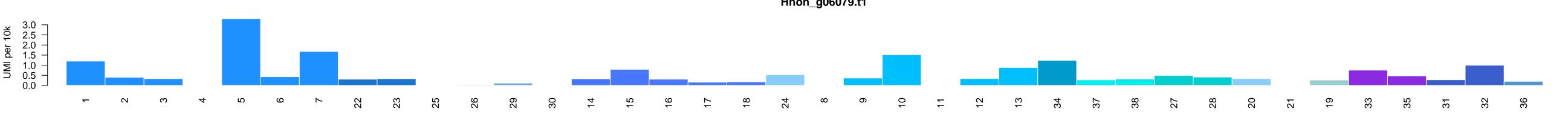
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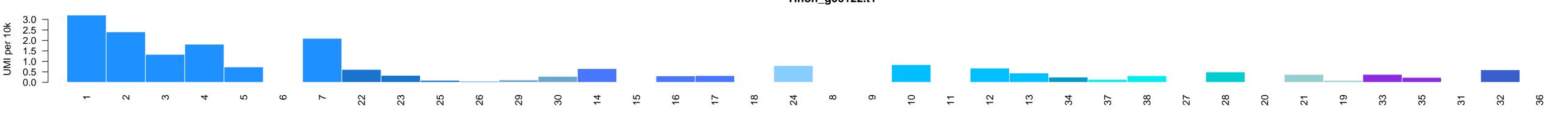
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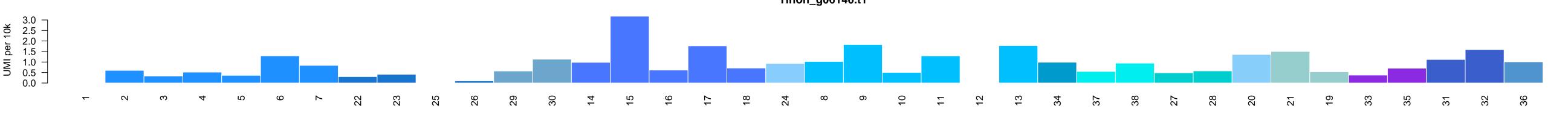
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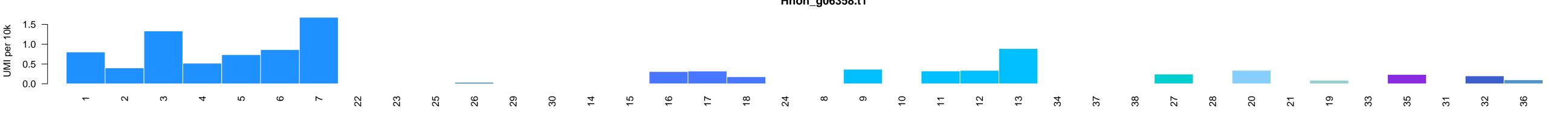
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Pkinase.HG4.18:CDK16/CDK17/CDK18 Hhon_g06140.t1

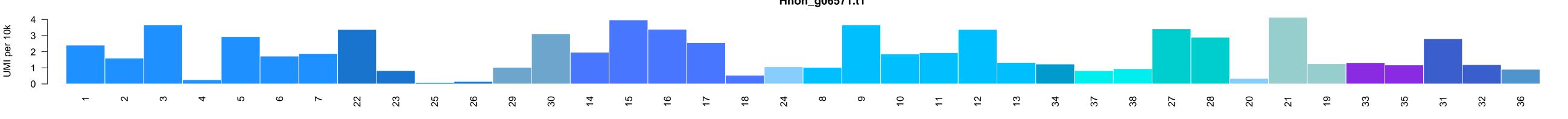


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Pkinase.HG12.1:AKT1/AKT2/AKT3 Hhon_g06571.t1

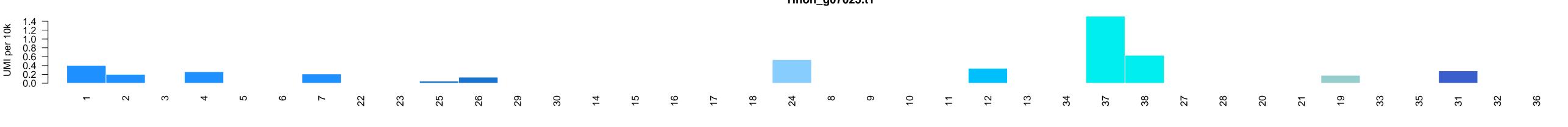


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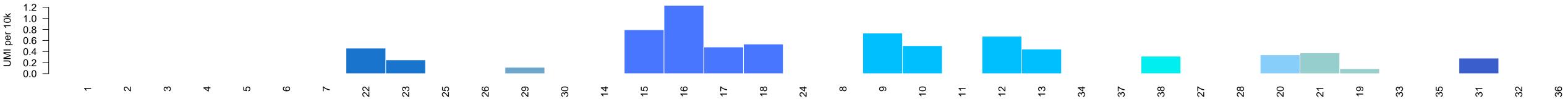
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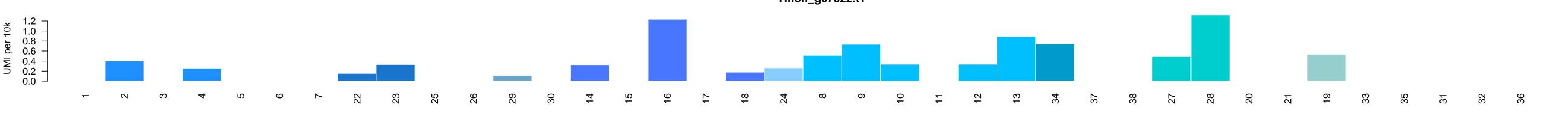
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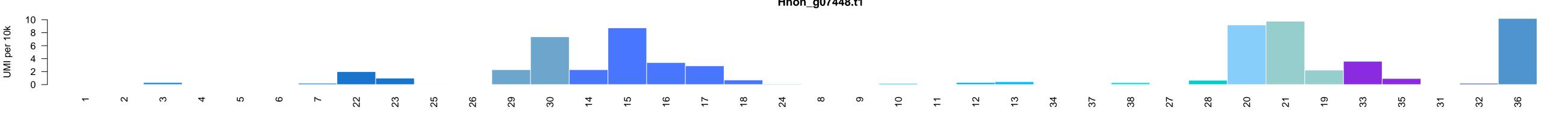
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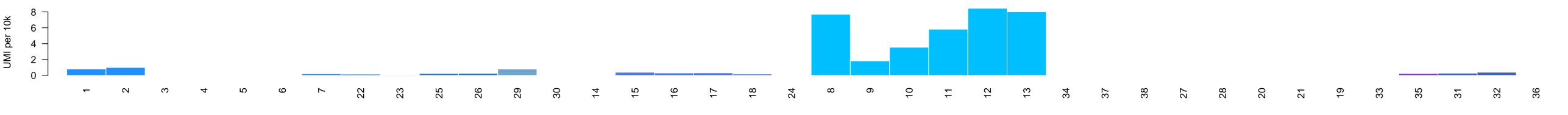
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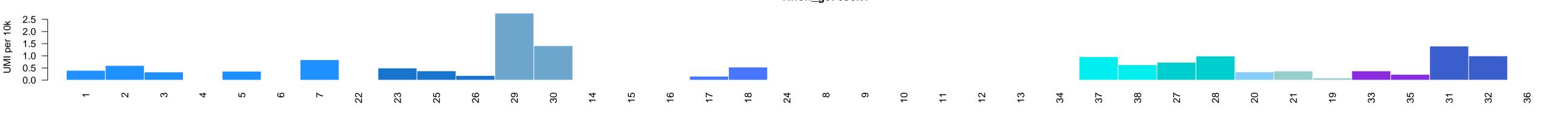
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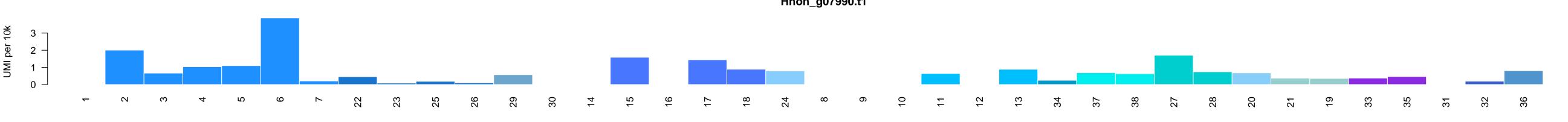
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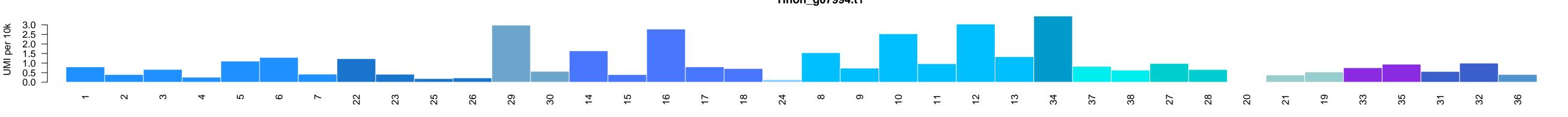
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Pkinase.HG2.37:SIK1/SIK1B/SIK2/SIK3 Hhon_g07990.t1

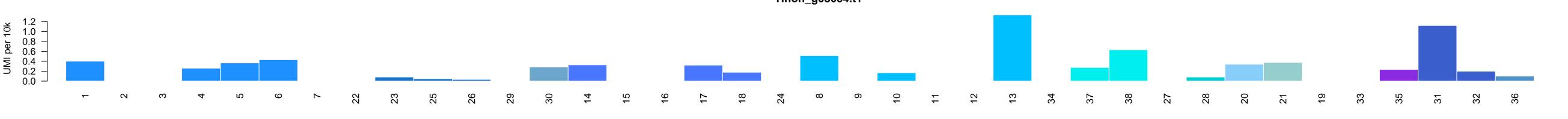


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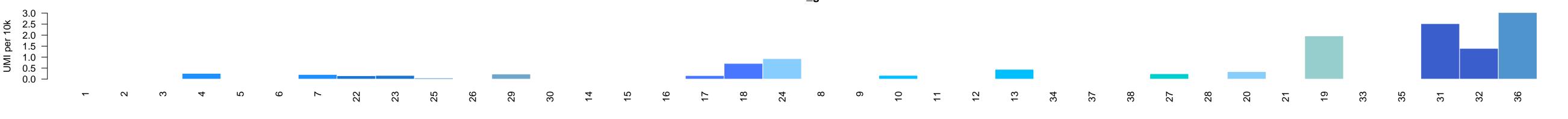


Pkinase.HG37.1:MOK Hhon_g08033.t1 1.5 The second of the second o

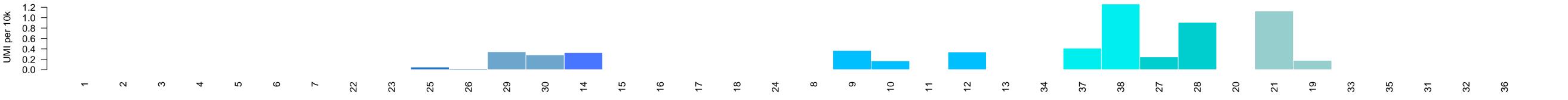
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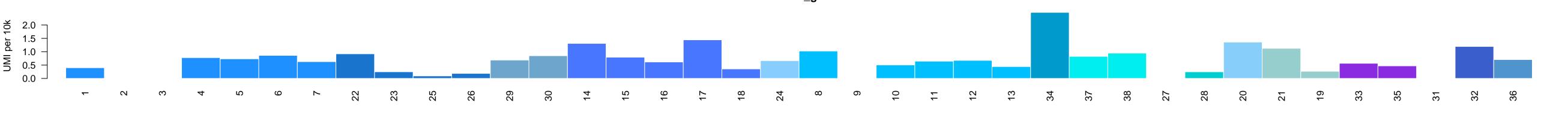
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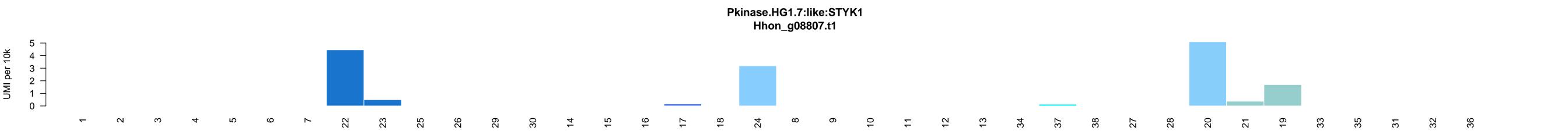
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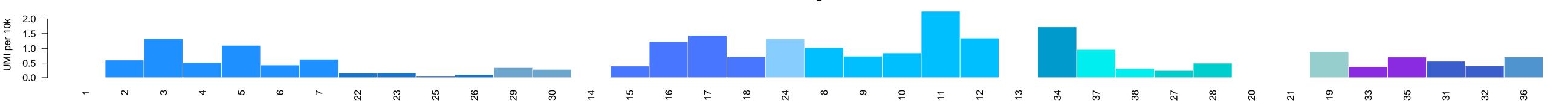
Pkinase.HG45.1:WNK1/WNK2/WNK3/WNK4 Hhon_g08521.t1



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Pkinase.HG17.8:MAPK8/MAPK9/MAPK10 Hhon_g09098.t1

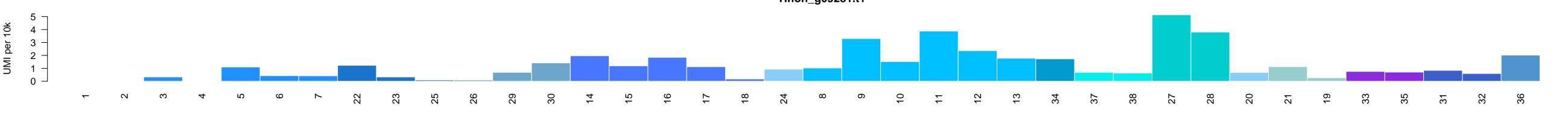


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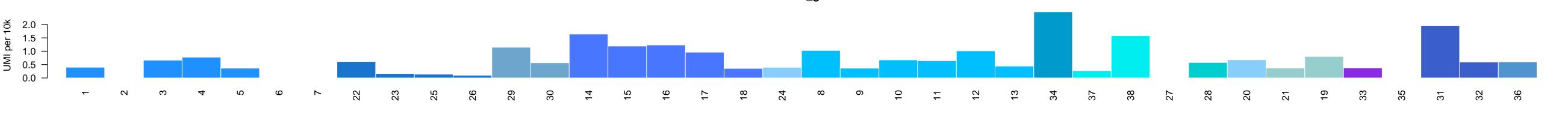
Pkinase.HG2.24:BRSK1/BRSK2 Hhon_g09281.t1



Pkinase.HG34.12:like:IRAK1/IRAK2/IRAK3/IRAK4 Hhon_g09836.t1

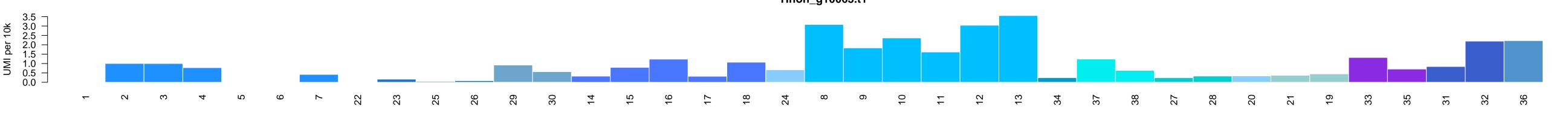


Pkinase.HG39.2:CDKL2/CDKL3 Hhon_g09904.t1

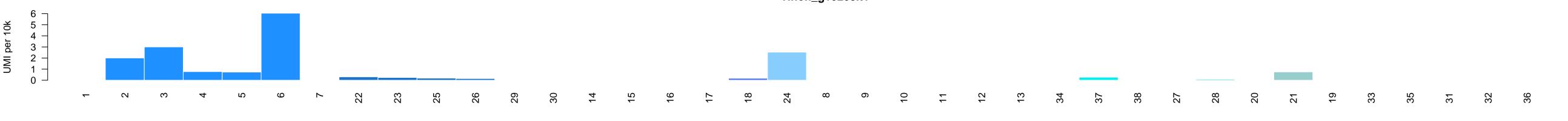


Pkinase.HG15.10:DYRK2/DYRK3 Hhon_g09946.t1 1.5 The second of the second o 30 15 17 18 18 18 $\overline{}$

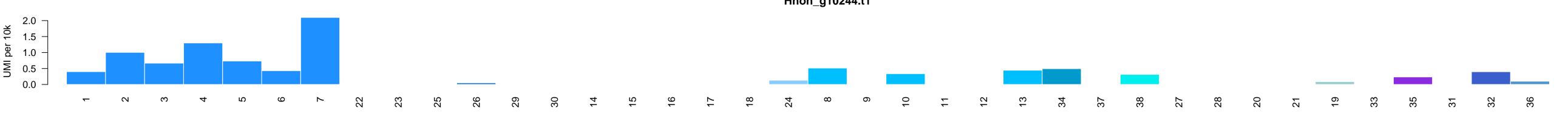
Pkinase.HG53.1:SRPK1/SRPK2/SRPK3 Hhon_g10063.t1



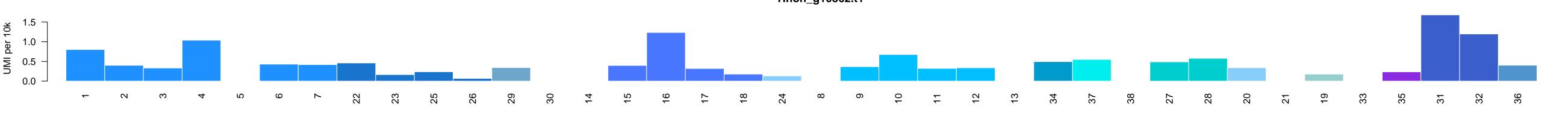
Pkinase.HG34.12:like:IRAK1/IRAK2/IRAK3/IRAK4 Hhon_g10208.t1



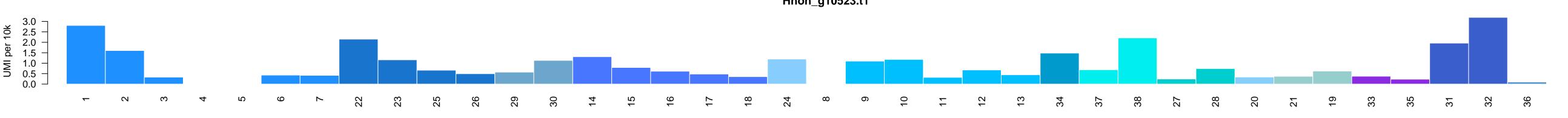
Pkinase.HG38.0:STK16 Hhon_g10244.t1



Pkinase.HG5.7:MAP3K7 Hhon_g10362.t1



Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3
Hhon_g10523.t1



Pkinase.HG12.10:AL355916.3/PRKCE/PRKCH Hhon_g10573.t1

