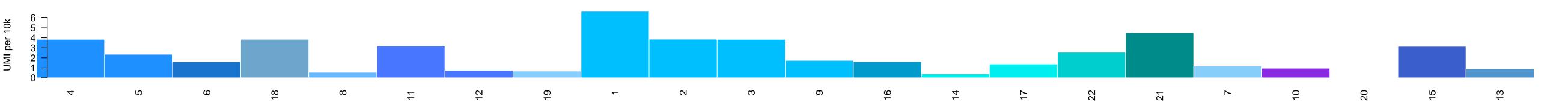
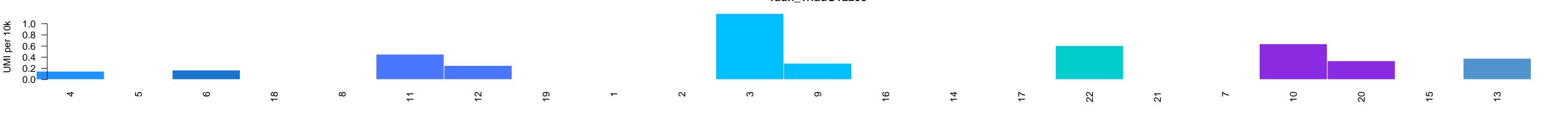
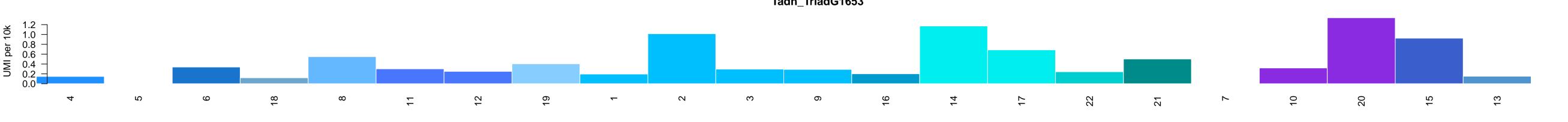
Pkinase.HG2.34:NIM1K Tadh_TriadG11807



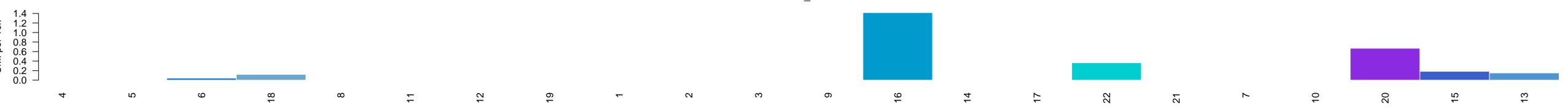
Pkinase.HG8.16:AATK/IGF1R/INSR/INSRR/LMTK2/LMTK3 Tadh_TriadG12208



Pkinase.HG5.11:MAP3K12/MAP3K13 Tadh_TriadG1653

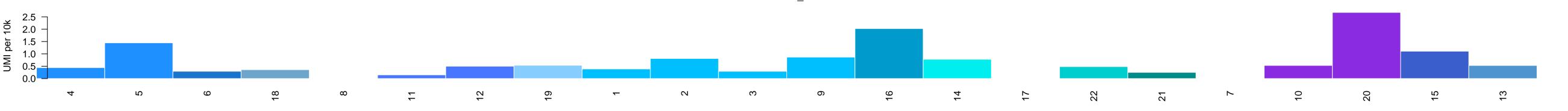


Pkinase.HG22.0:TTBK1/TTBK2 Tadh_TriadG19397

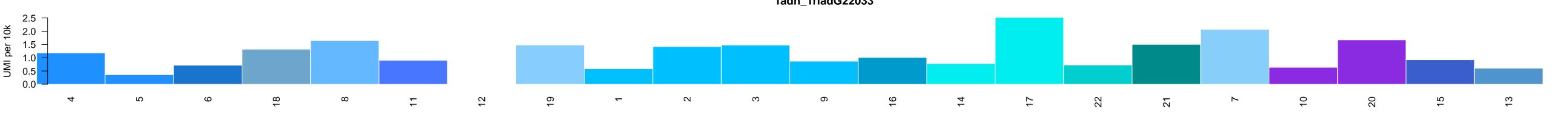


Pkinase.HG39.5:CDKL1/CDKL4

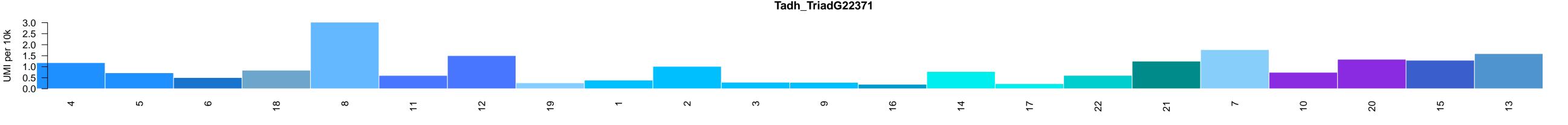
Pkinase.HG33.7:MAP2K4 Tadh_TriadG21381



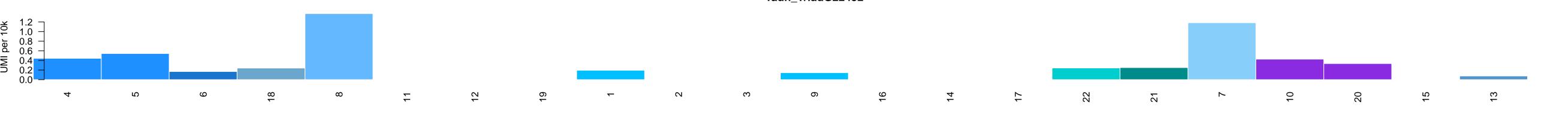
Pkinase.HG28.16:like:ACVR1/ACVR1B/ACVR1C/ACVRL1/BMPR1A/BMPR1B/TGFBR1
Tadh_TriadG22033



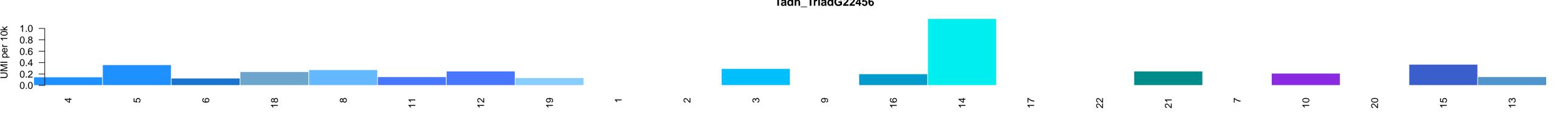
Pkinase.HG22.11:CSNK1G1/CSNK1G2/CSNK1G3 Tadh_TriadG22371



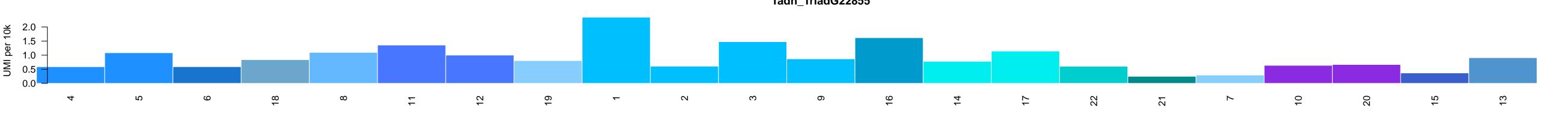
Pkinase.HG28.18:ACVR1/ACVRL1/BMPR1A/BMPR1B Tadh_TriadG22452



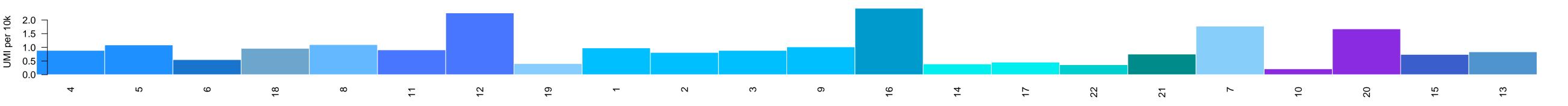
Pkinase.HG19.1:MAP3K2/MAP3K3 Tadh_TriadG22456



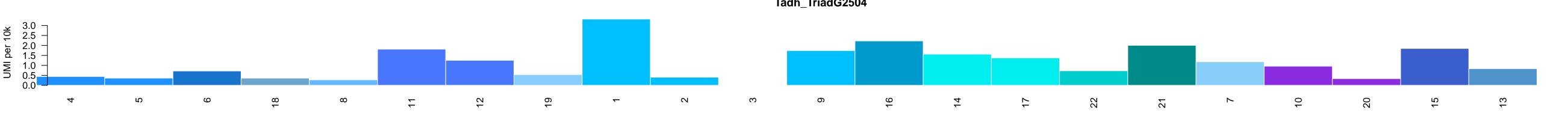
Pkinase.HG8.16:AATK/IGF1R/INSR/INSRR/LMTK2/LMTK3 Tadh_TriadG22855



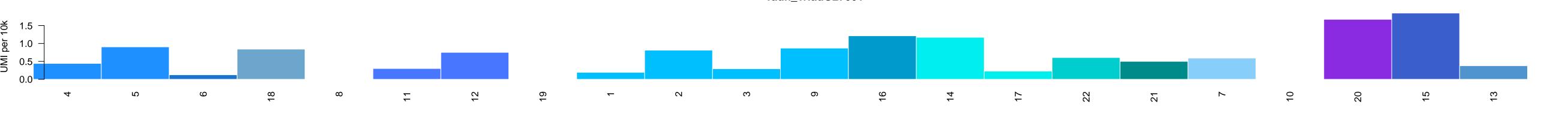
Pkinase.HG2.37:SIK1/SIK1B/SIK2/SIK3
Tadh_TriadG23373



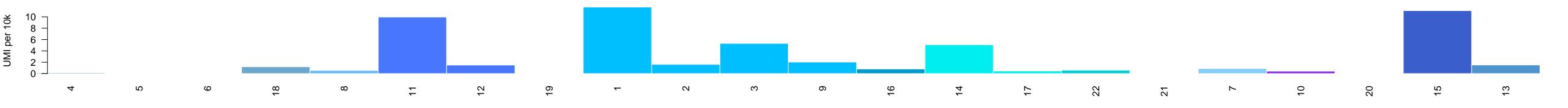
Pkinase.HG15.10:DYRK2/DYRK3 Tadh_TriadG2504



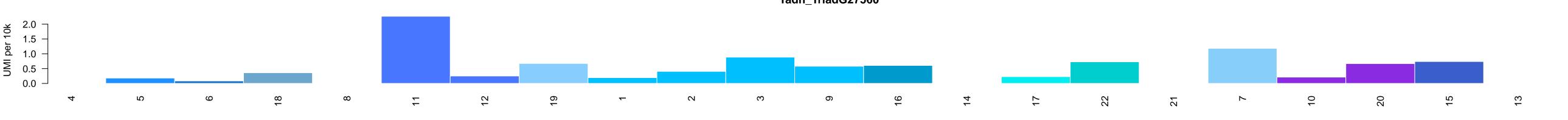
Pkinase.HG2.39:MARK1/MARK2/MARK3/MARK4 Tadh_TriadG27001



Pkinase.HG7.12:like:NEK6/NEK7 Tadh_TriadG27553



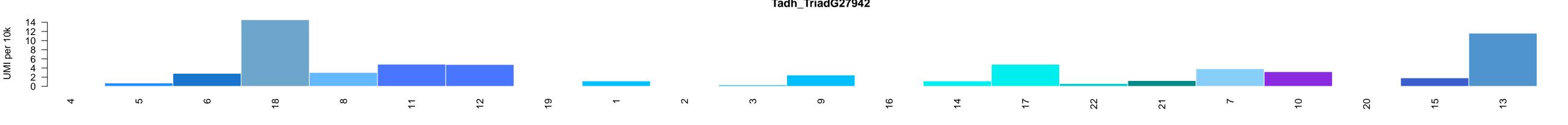
Pkinase.HG28.18:ACVR1/ACVRL1/BMPR1A/BMPR1B Tadh_TriadG27560



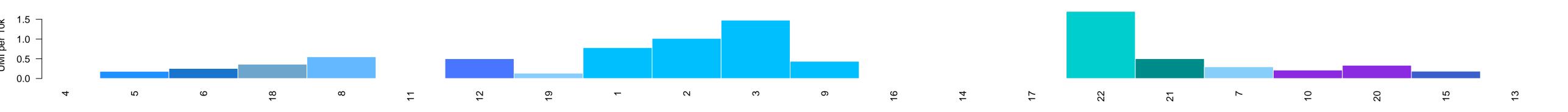
Pkinase.HG37.2:CILK1/MAK Tadh_TriadG27686

Pkinase.HG2.3:PASK Tadh_TriadG27874 3 2

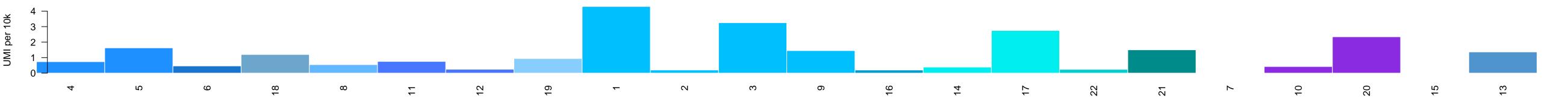
Pkinase.HG7.11:like:NEK6/NEK7 Tadh_TriadG27942



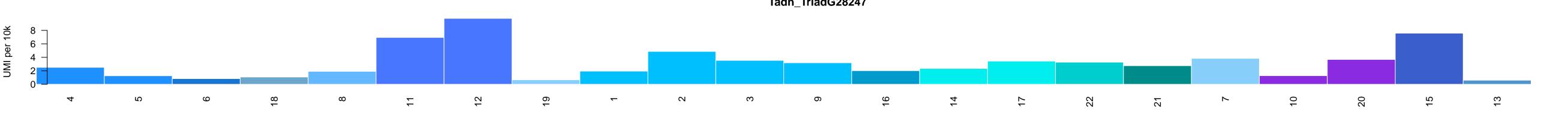
Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 Tadh_TriadG27989



Pkinase.HG5.21:MOS Tadh_TriadG28028

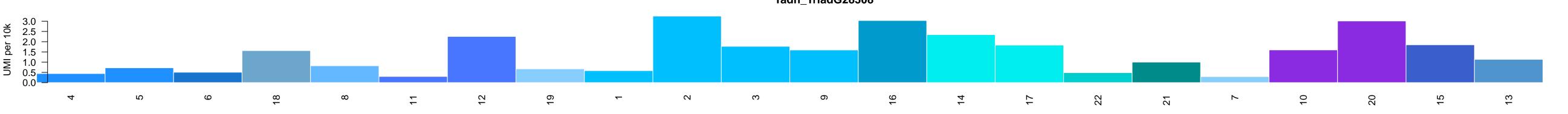


Pkinase.HG14.3:STK32A/STK32B/STK32C Tadh_TriadG28247

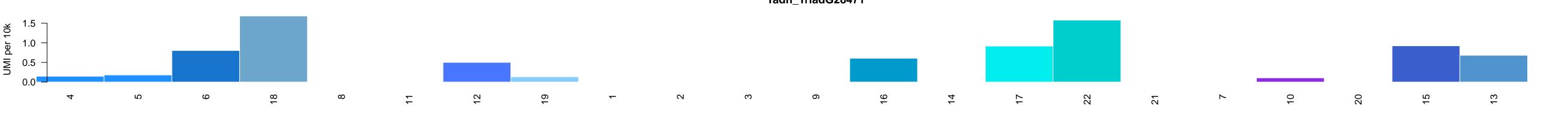


Pkinase.HG40.4:like:GRK1/GRK2/GRK3/GRK4/GRK5/GRK6/GRK7;RGS.HG2.0:GRK1/GRK4/GRK5/GRK6/GRK7

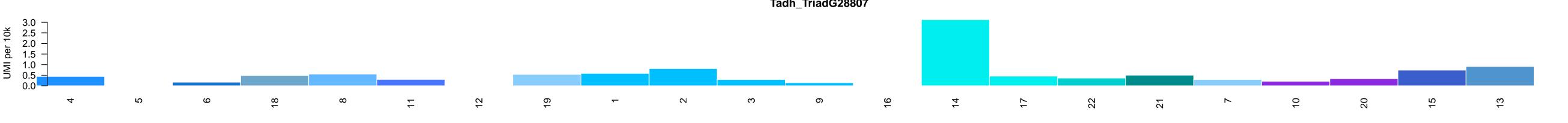
Tadh_TriadG28308



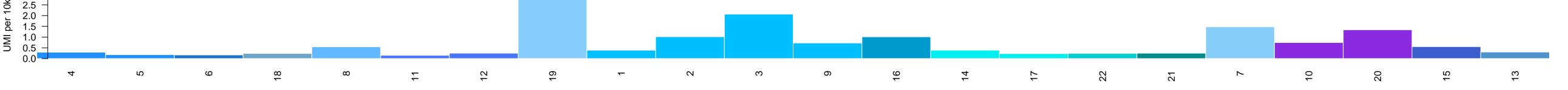
Pkinase.HG13.15:GUCY2D/GUCY2F Tadh_TriadG28471



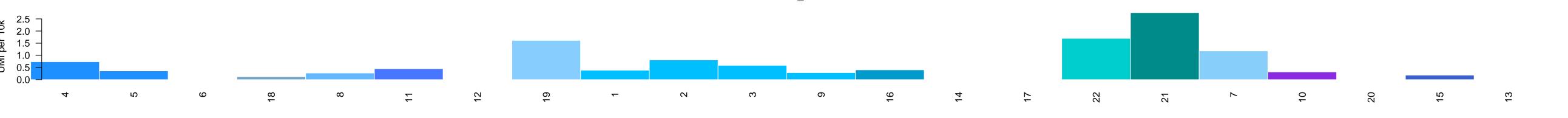
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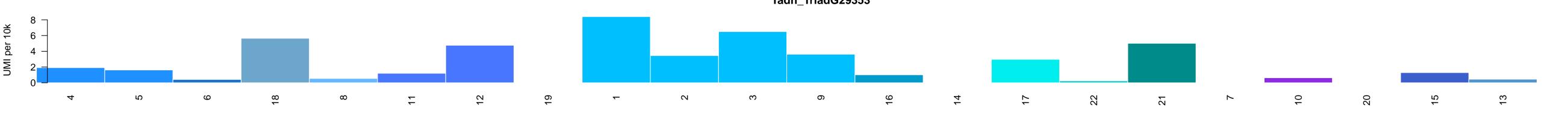
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Pkinase.HG2.23:CAMKK1/CAMKK2 Tadh_TriadG28940

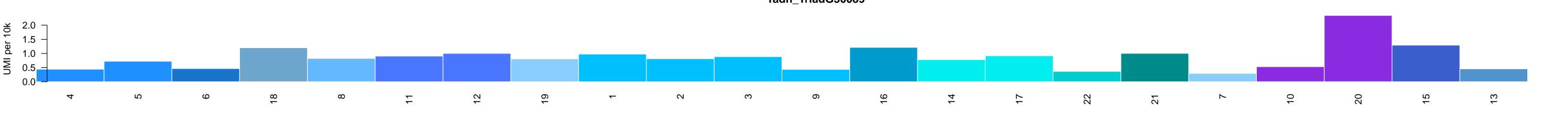


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Pkinase.HG12.10:AL355916.3/PRKCE/PRKCH Tadh_TriadG30067

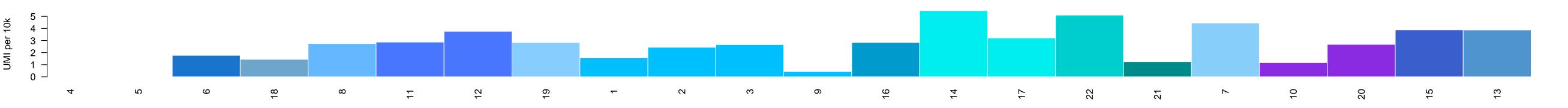
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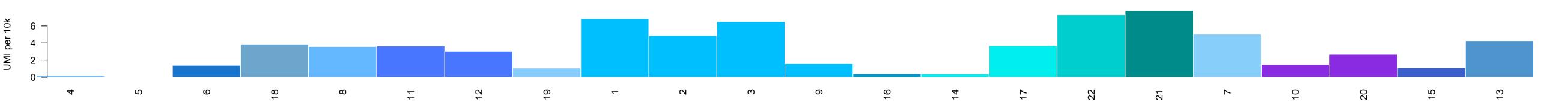
Pkinase.HG10.6:TAOK1/TAOK2/TAOK3 Tadh_TriadG30095

Pkinase.HG17.8:MAPK8/MAPK9/MAPK10 Tadh_TriadG30150

Pkinase.HG29.15:like:PRKG1 Tadh_TriadG30166



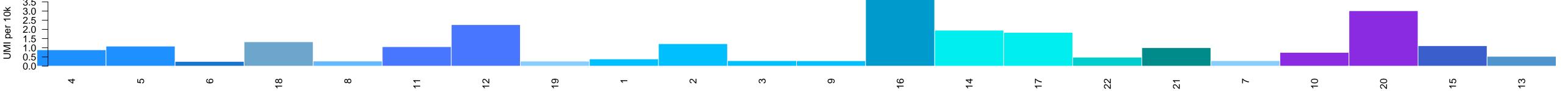
Pkinase.HG13.5:like:GUCY2C/GUCY2D/GUCY2F/NPR1/NPR2 Tadh_TriadG30655



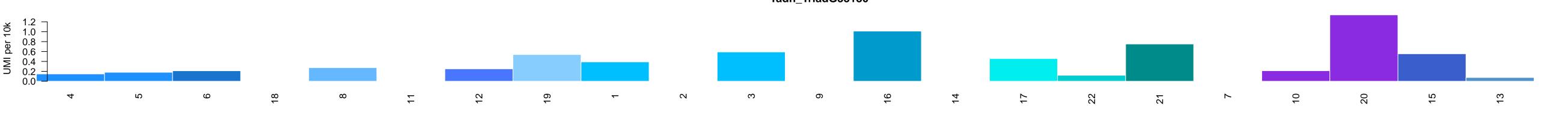
Pkinase.HG28.3:ACVR2A/ACVR2B Tadh_TriadG3190

Pkinase.HG10.8:STK3/STK4 Tadh_TriadG32096 2

Pkinase.HG15.4:HIPK1/HIPK2/HIPK3/HIPK4
Tadh_TriadG32819



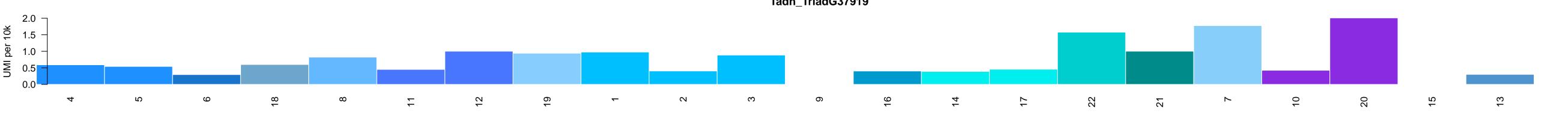
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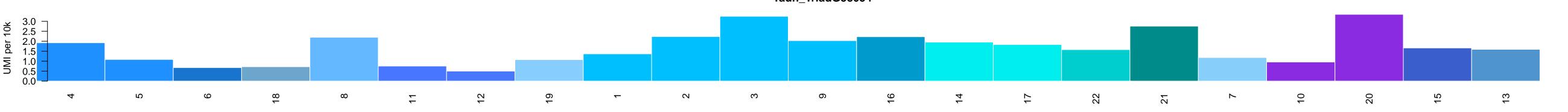
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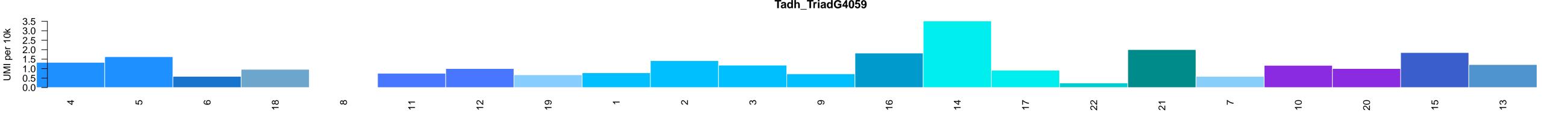
Pkinase.HG2.25:PRKAA1/PRKAA2 Tadh_TriadG37919



Pkinase.HG17.3:NLK Tadh_TriadG38054



Pkinase.HG45.1:WNK1/WNK2/WNK3/WNK4 Tadh_TriadG4059



Pkinase.HG2.32:HUNK
Tadh_TriadG4614

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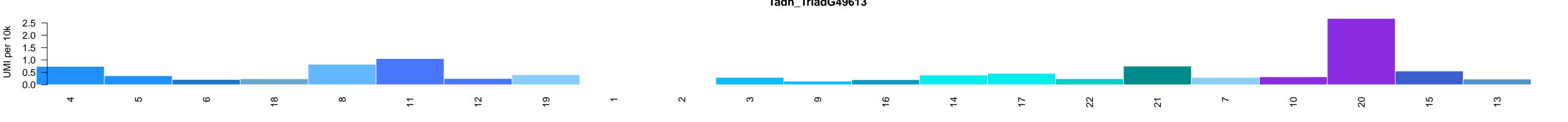
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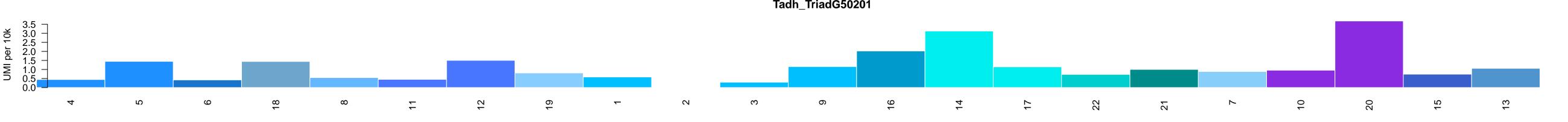
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Pkinase.HG10.10:STK24/STK25/STK26 Tadh_TriadG49722

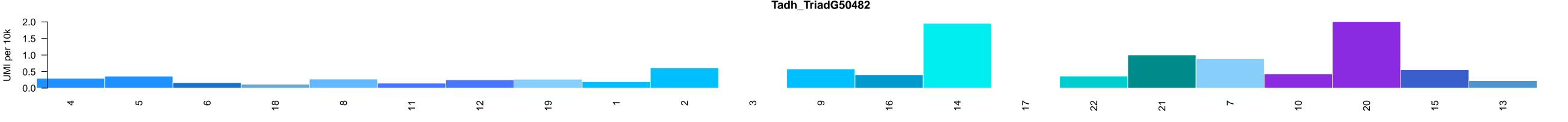


Pkinase.HG4.17:CDK14/CDK15 Tadh_TriadG50028 2

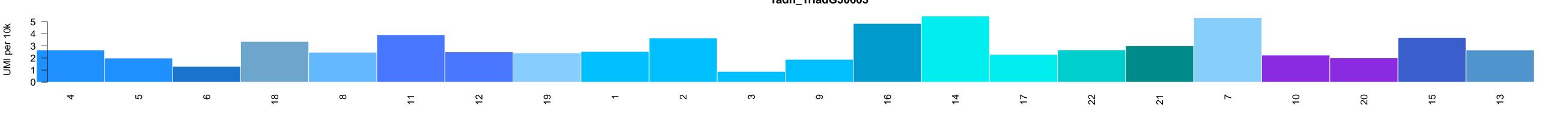
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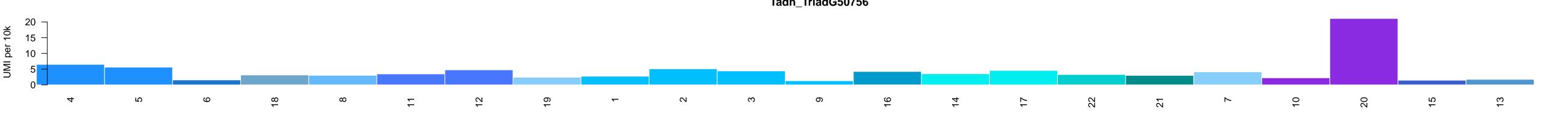
Pkinase.HG3.33:like:BLK/FGR/FYN/HCK/LCK/LYN/SRC/YES1 Tadh_TriadG50482



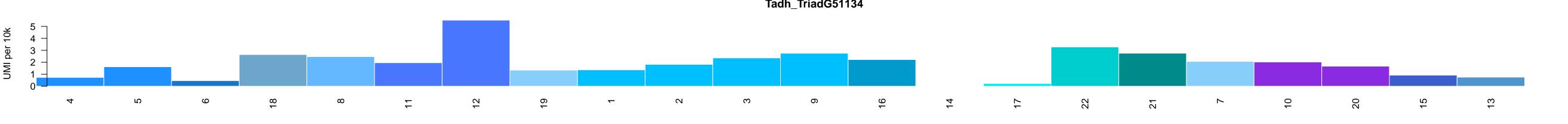
Pkinase.HG22.8:CSNK1A1/CSNK1A1L Tadh_TriadG50603



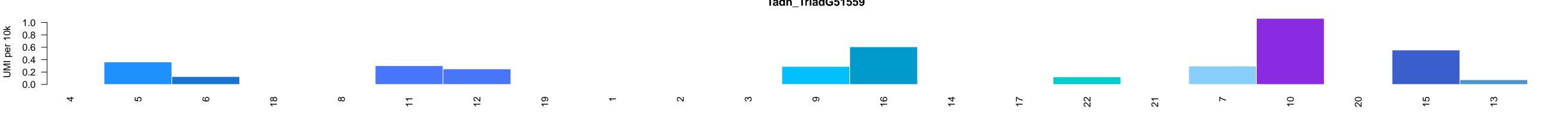
Pkinase.HG6.22:CAMK2A/CAMK2B/CAMK2D/CAMK2G Tadh_TriadG50756



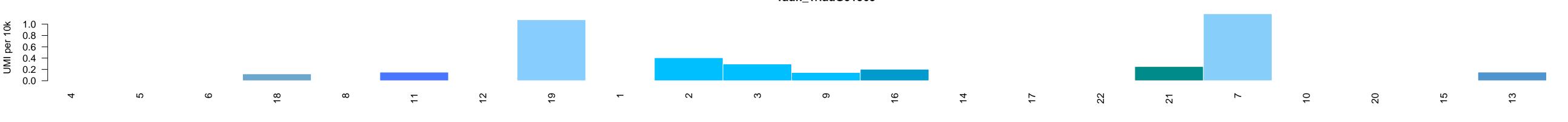
Pkinase.HG12.1:AKT1/AKT2/AKT3 Tadh_TriadG51134



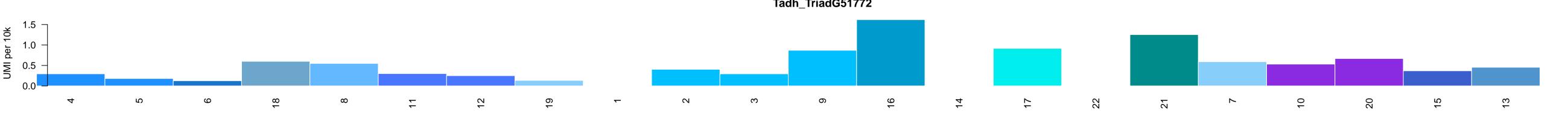
Pkinase.HG34.12:like:IRAK1/IRAK2/IRAK3/IRAK4 Tadh_TriadG51559



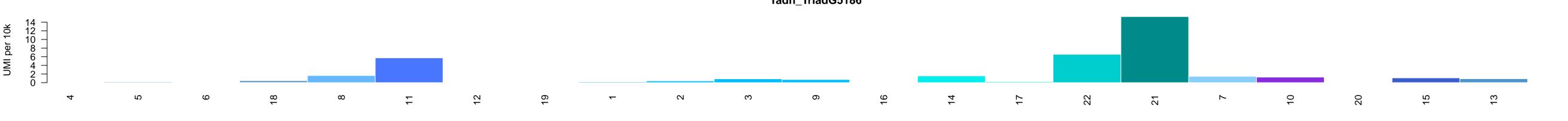
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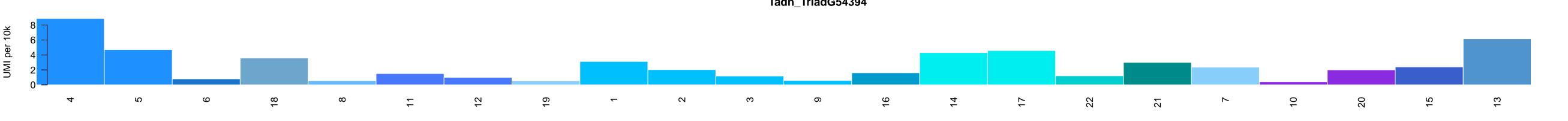
Pkinase.HG10.9:MAP4K1/MAP4K2/MAP4K3/MAP4K5 Tadh_TriadG51772



Pkinase.HG6.0:like:CHEK2/PRKD1/PRKD2/PRKD3/STK33 Tadh_TriadG5186



Pkinase.HG23.0:KALRN/TRIO Tadh_TriadG54394



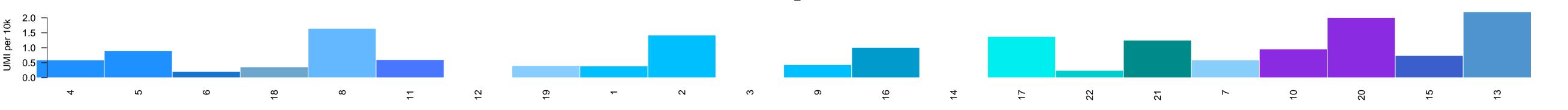
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Pkinase.HG31.14:like:DSTYK Tadh_TriadG55120

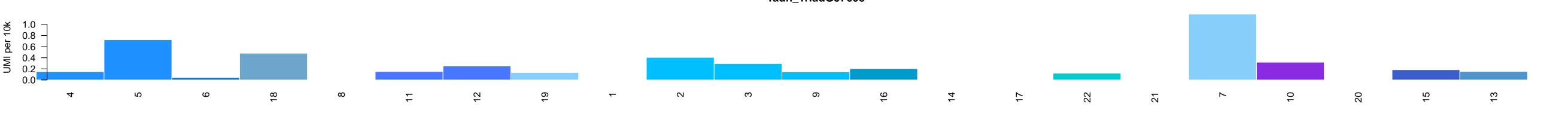




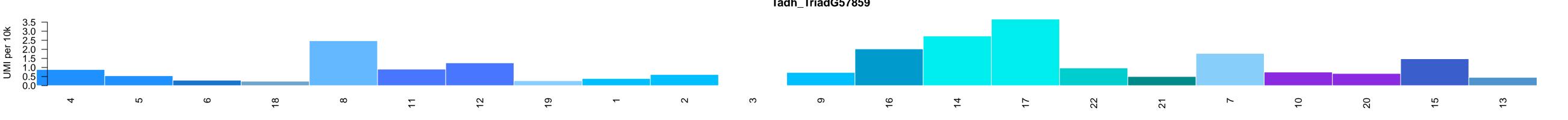
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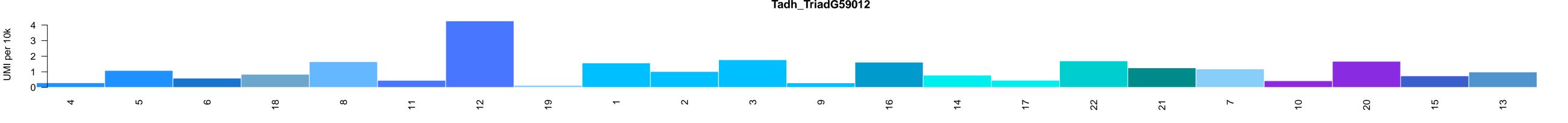
Pkinase.HG12.6:PRKCI/PRKCZ Tadh_TriadG57608



Pkinase.HG16.2:MAST1/MAST2/MAST3/MAST4 Tadh_TriadG57859



Pkinase.HG40.10:GRK2/GRK3;RGS.HG2.2:GRK2/GRK3 Tadh_TriadG59012



Pkinase.HG13.1:like:GUCY2C/GUCY2D/GUCY2F/NPR1/NPR2
Tadh_TriadG59364



Pkinase.HG6.7:CAMK4 Tadh_TriadG60371

