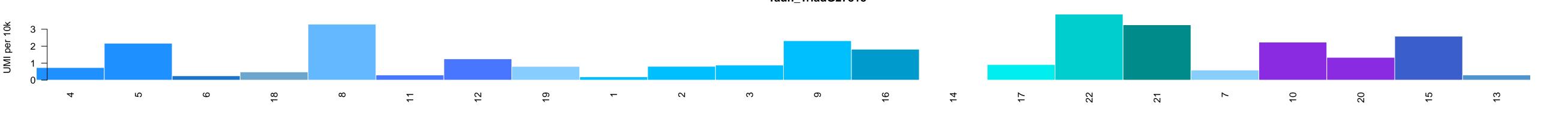
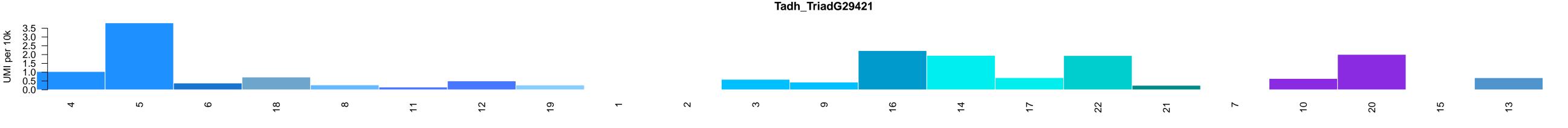
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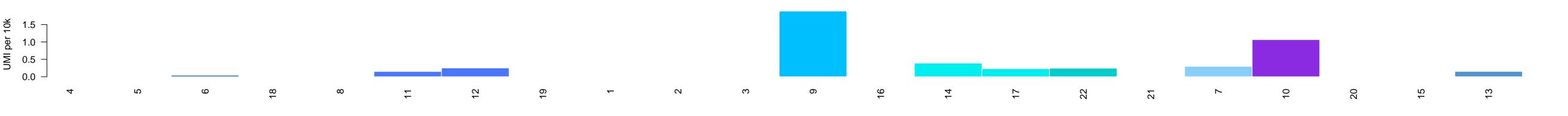


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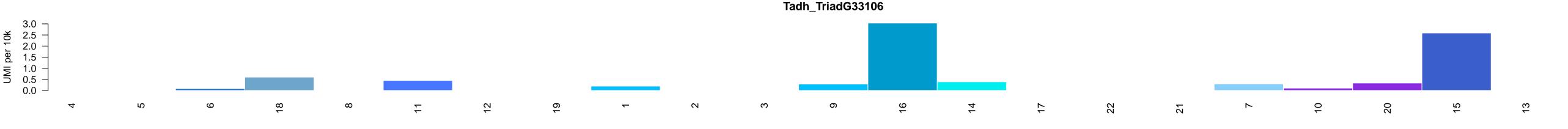


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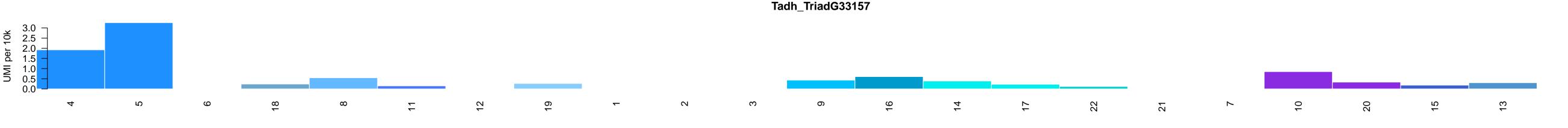
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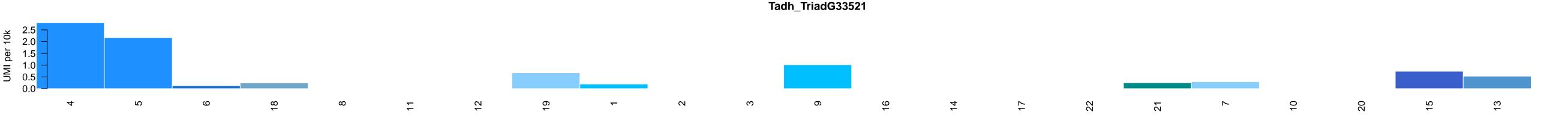
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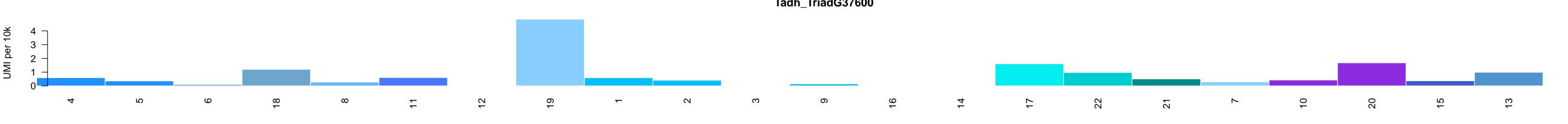
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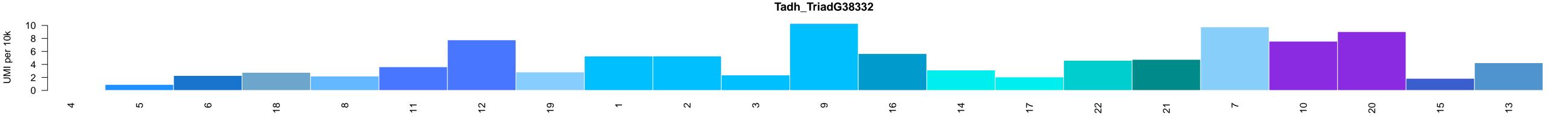
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SNF.HG1.4:SLC6A15/SLC6A16/SLC6A17/SLC6A18/SLC6A19/SLC6A20 Tadh_TriadG37600



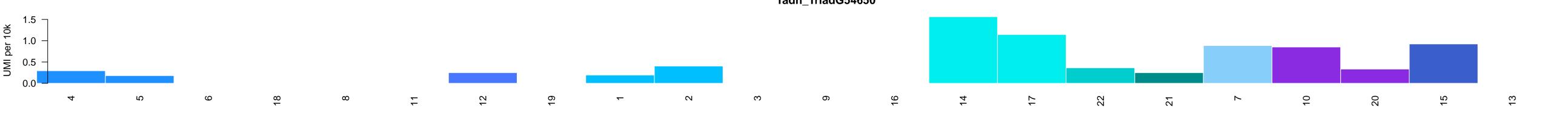
SNF.HG1.31:like:SLC6A5/SLC6A7/SLC6A9/SLC6A14 Tadh_TriadG38332

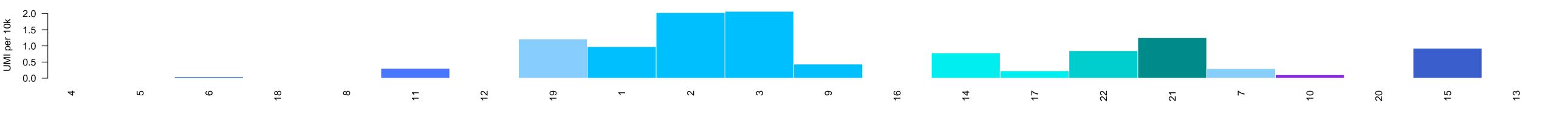


SNF.HG1.30:like:SLC6A5/SLC6A7/SLC6A9/SLC6A14 Tadh_TriadG50950

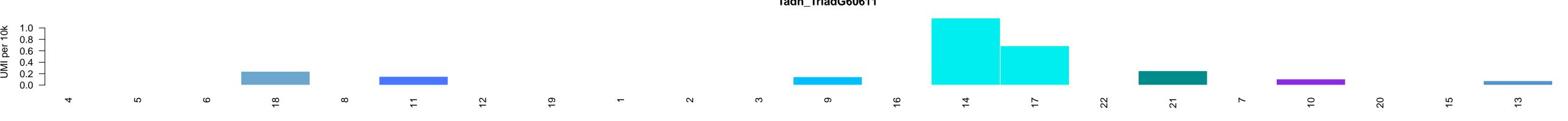


SNF.HG1.46:like:SLC6A5/SLC6A7/SLC6A9/SLC6A14 Tadh_TriadG54650



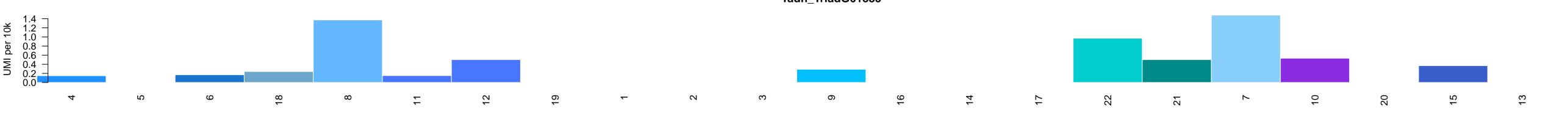


Pyridoxal_deC.HG1.4:CSAD/GAD1/GAD2/GADL1
Tadh_TriadG60611

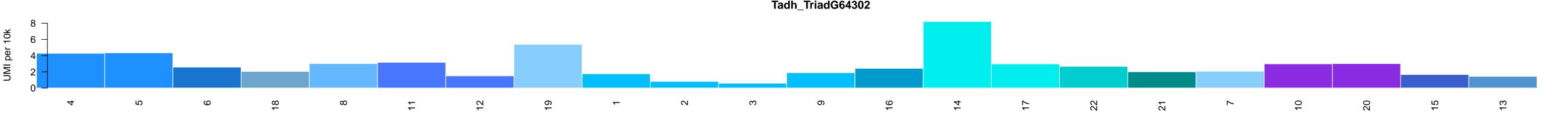


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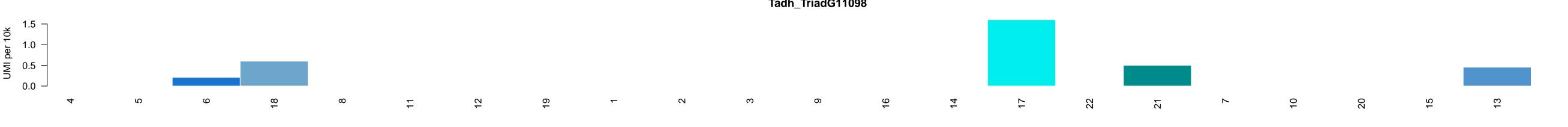
Tadh_TriadG61885

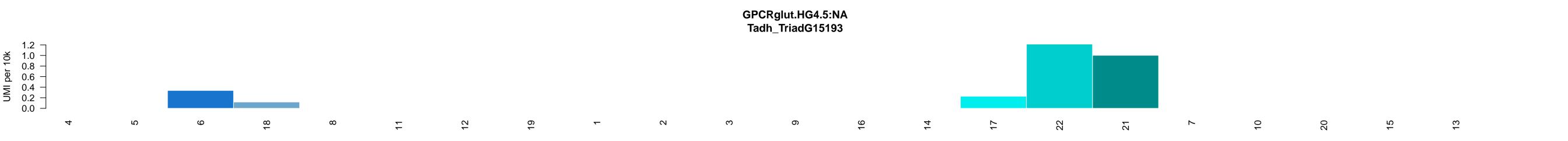


SNF.HG1.25:like:SLC6A5/SLC6A7/SLC6A9/SLC6A14 Tadh_TriadG64302



GPCRglut.HG2.4:like:GRM1/GRM2/GRM3/GRM4/GRM5/GRM6/GRM7/GRM8 Tadh_TriadG11098





GPCRglut.HG1.35:like:GABBR1/GABBR2
Tadh_TriadG16716



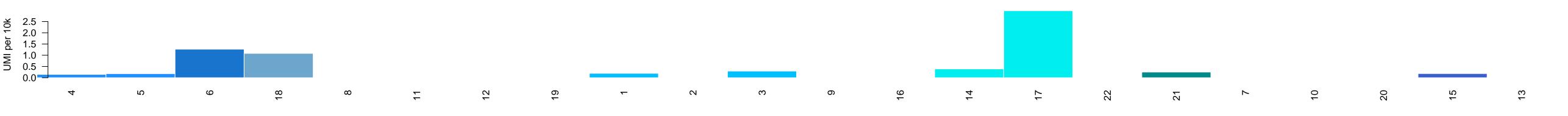
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Tadh_TriadG25328

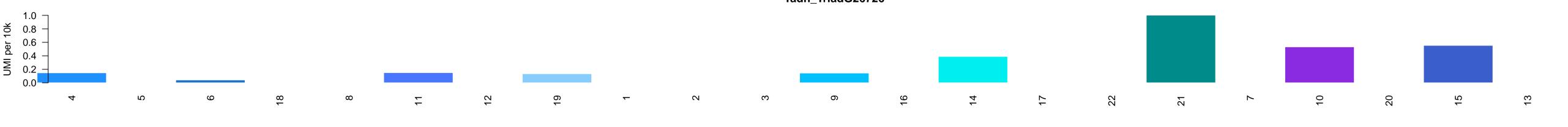


GPCRglut.HG2.4:like:GRM1/GRM2/GRM3/GRM4/GRM5/GRM6/GRM7/GRM8

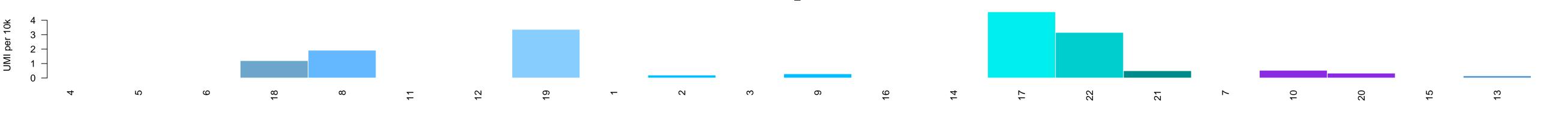
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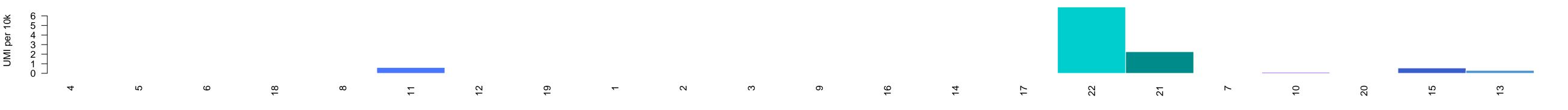
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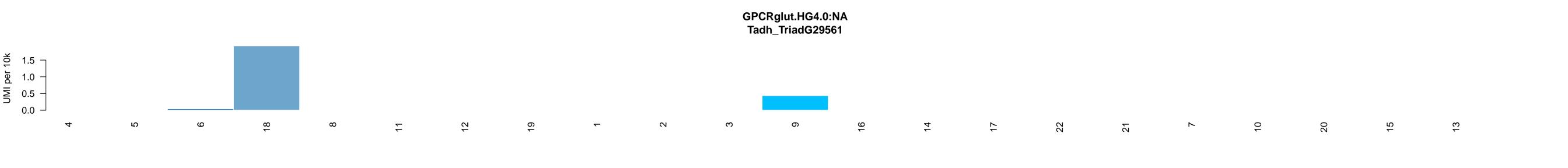
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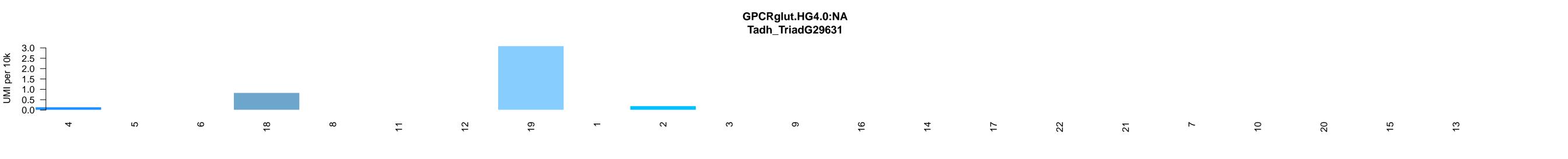
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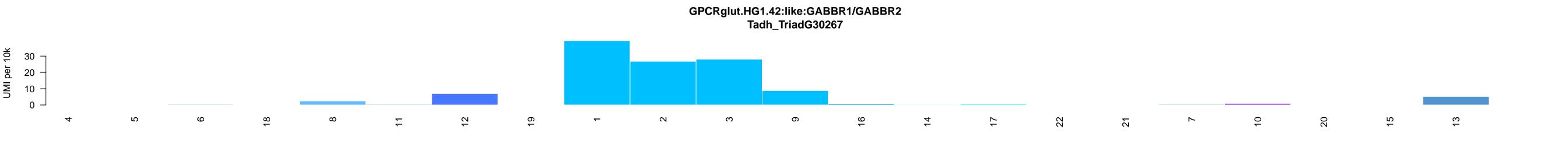












GPCRglut.HG1.29:like:GABBR1/GABBR2 Tadh_TriadG30374

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GPCRglut.HG1.11:like:GABBR1/GABBR2/GPR156 Tadh_TriadG52576



GPCRglut.HG1.9:like:GABBR1/GABBR2/GPR156 Tadh_TriadG52577

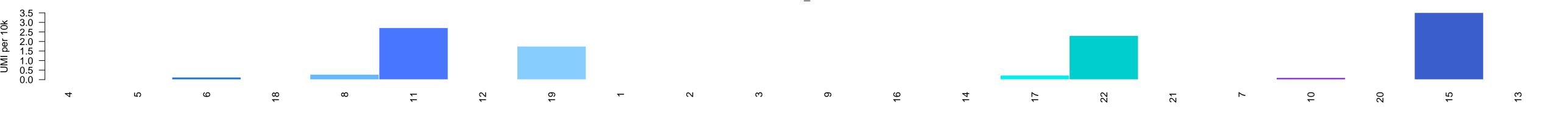


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GPCRglut.HG1.5:like:GABBR1/GABBR2/GPR156 Tadh_TriadG52578



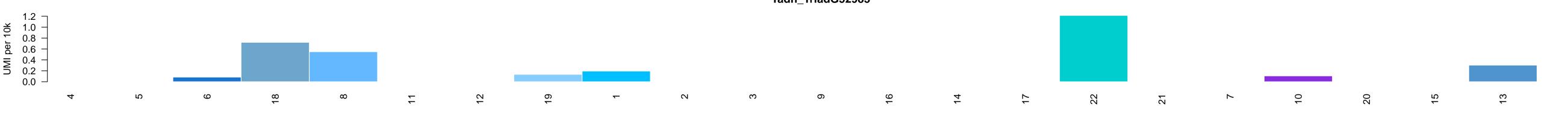




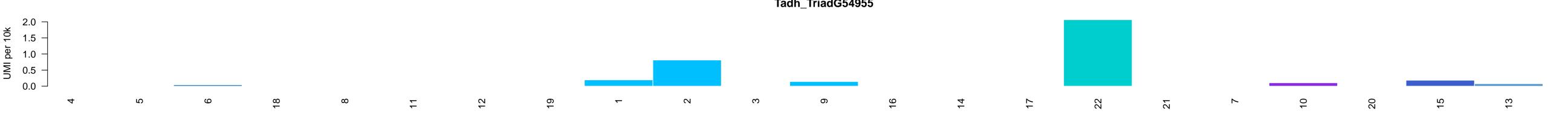
GPCRglut.HG1.7:like:GABBR1/GABBR2/GPR156 Tadh_TriadG52581



GPCRglut.HG1.6:GABBR1/GABBR2 Tadh_TriadG52983



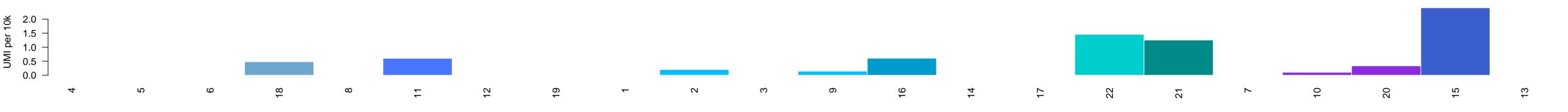
GPCRglut.HG1.30:like:GABBR1/GABBR2 Tadh_TriadG54955



GPCRglut.HG1.30:like:GABBR1/GABBR2 Tadh_TriadG54957



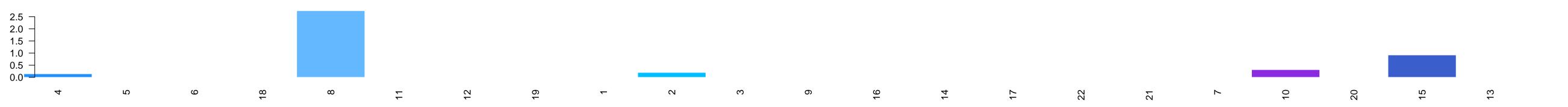
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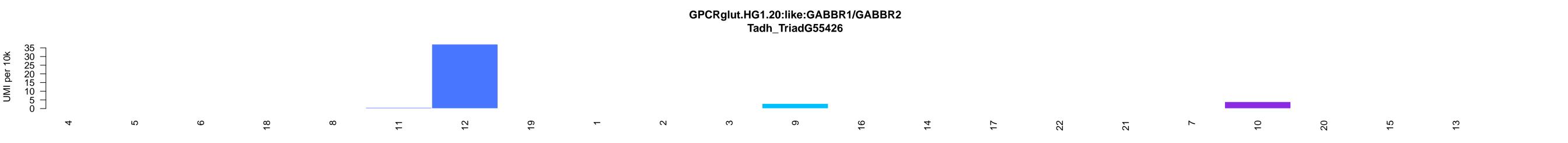


GPCRglut.HG1.45:like:GABBR1/GABBR2 Tadh_TriadG55386



GPCRglut.HG1.20:like:GABBR1/GABBR2 Tadh_TriadG55425





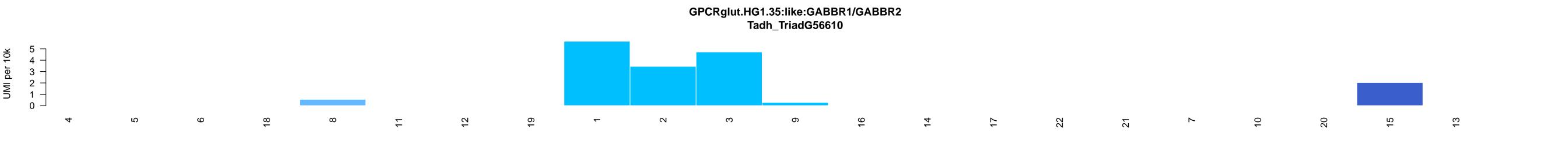
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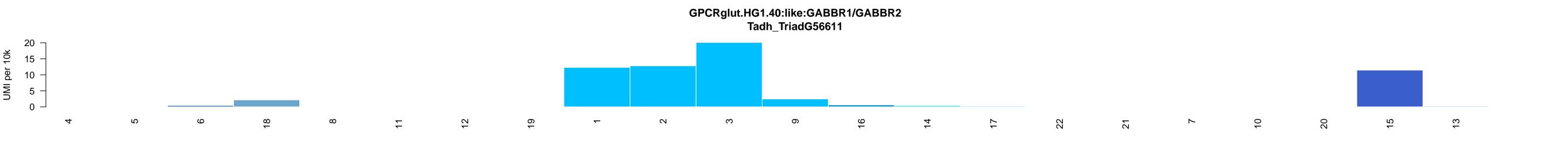
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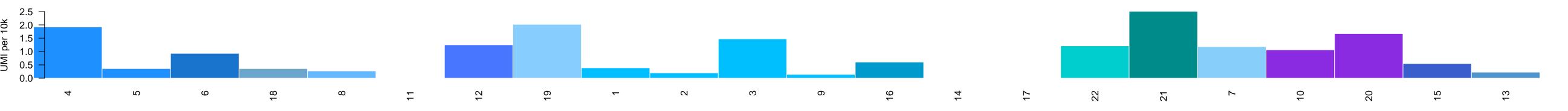
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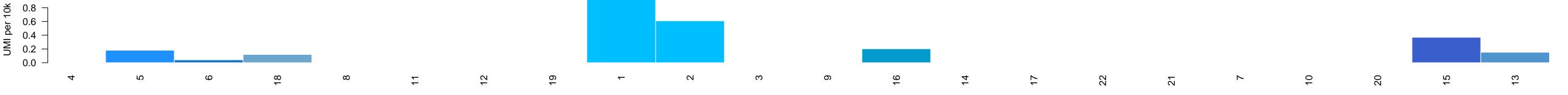




GPCRglut.HG1.10:like:GABBR1/GABBR2 Tadh_TriadG57193



GPCRglut.HG1.30:like:GABBR1/GABBR2 Tadh_TriadG58937



GPCRglut.HG1.30:like:GABBR1/GABBR2
Tadh_TriadG58938

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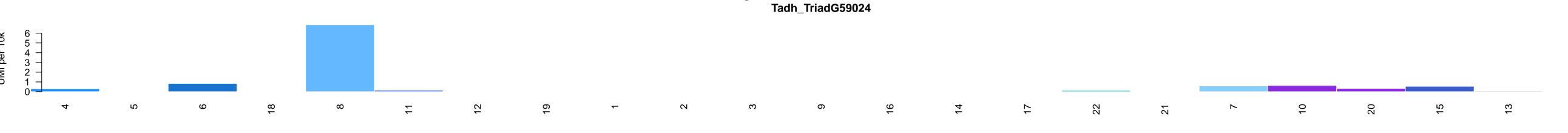
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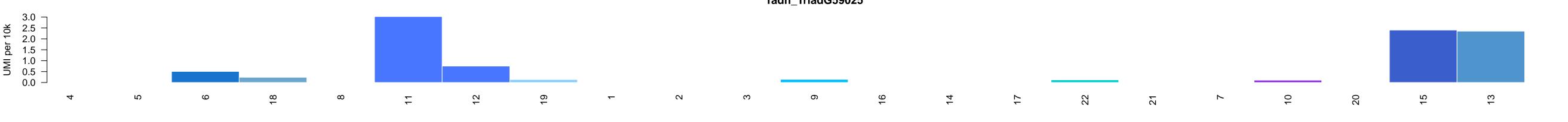
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GPCRglut.HG1.30:like:GABBR1/GABBR2 Tadh_TriadG59024







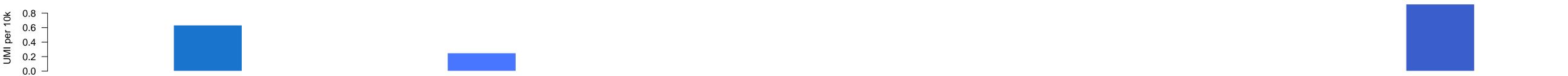


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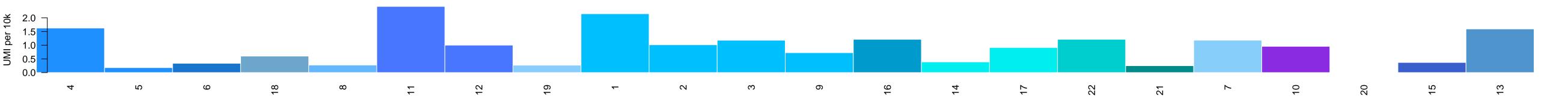


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GPCRglut.HG1.2:GPR156 Tadh_TriadG61926







GPCRglut.HG4.14:NA Tadh_wf_g3864

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