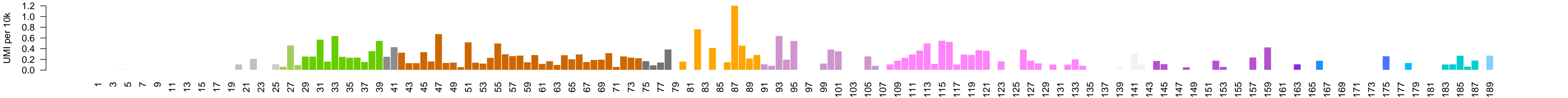
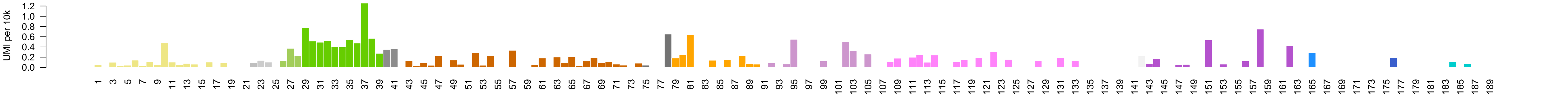


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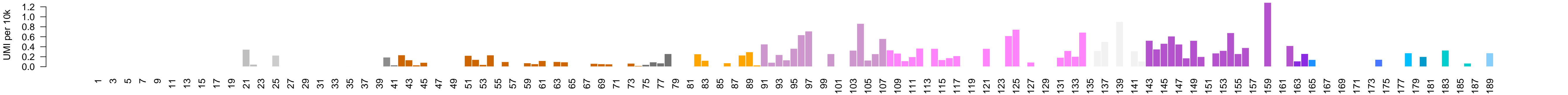


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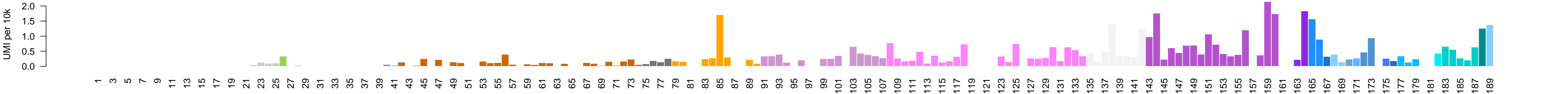


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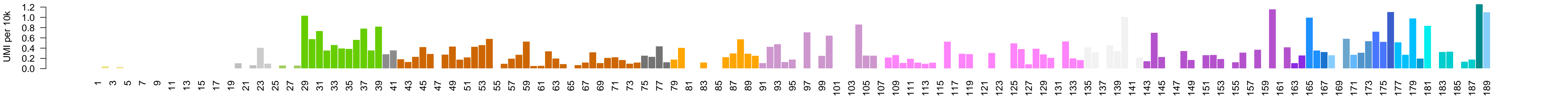
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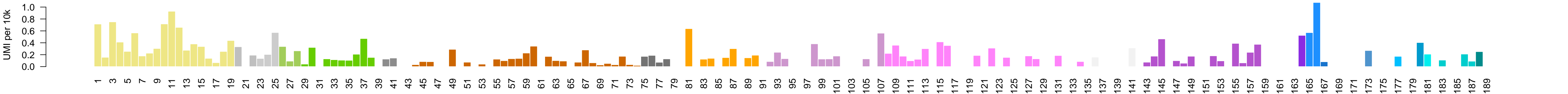
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Tadh_TriadG56468



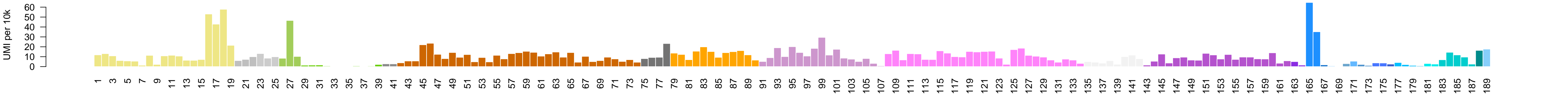
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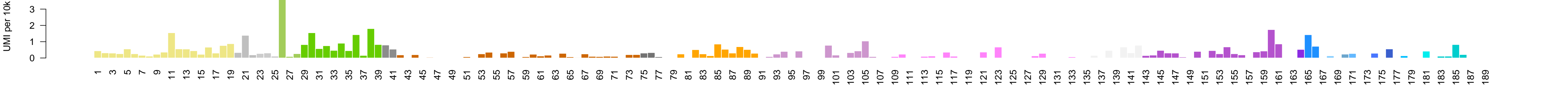
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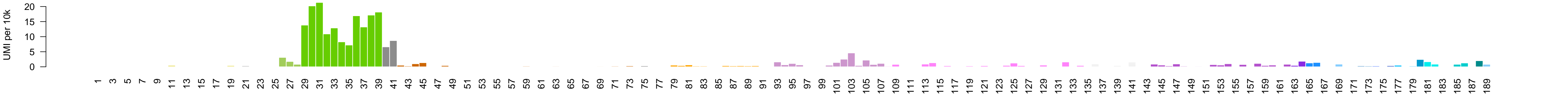
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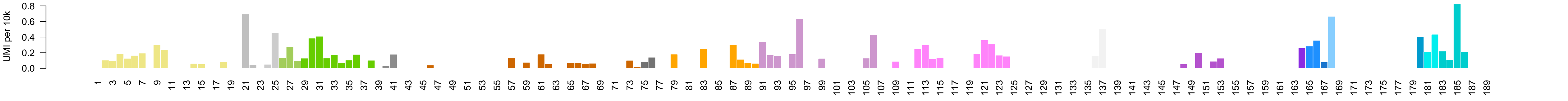
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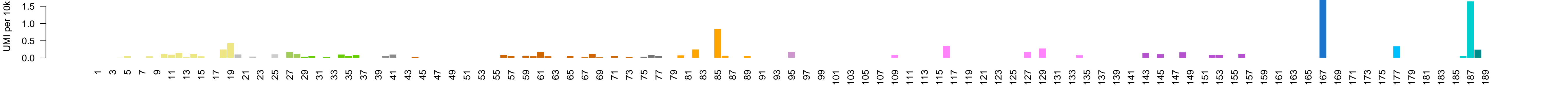
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Tadh_TriadG29287



Aa_trans.HG1.2:SLC38A9
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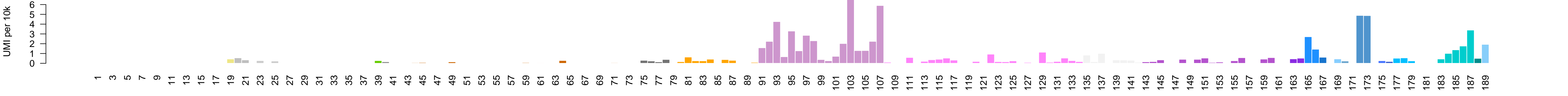
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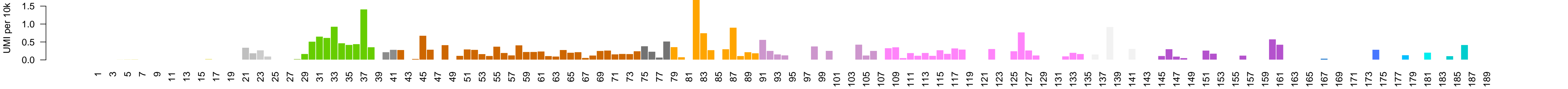
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Tadh_TriadG22445



Aa_trans.HG1.4:SLC38A11
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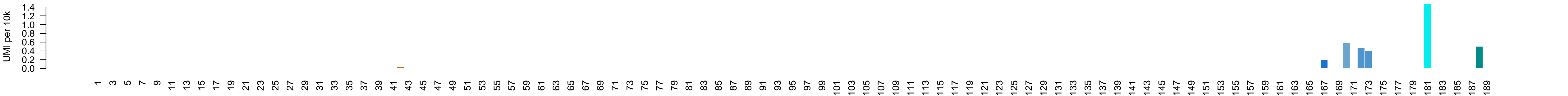


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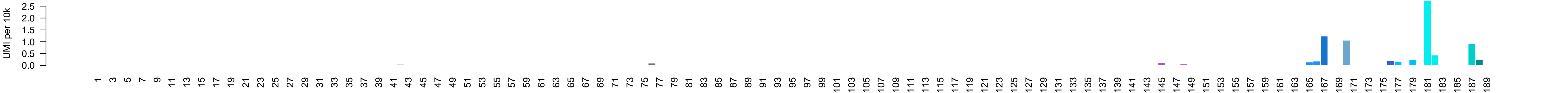
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Tadh_TriadG11098

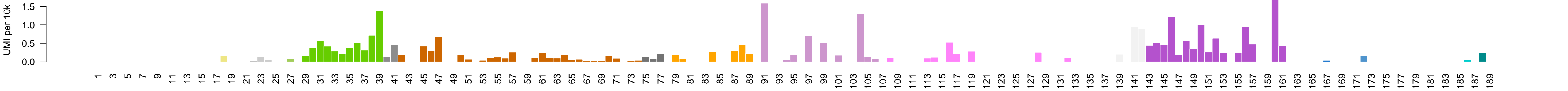


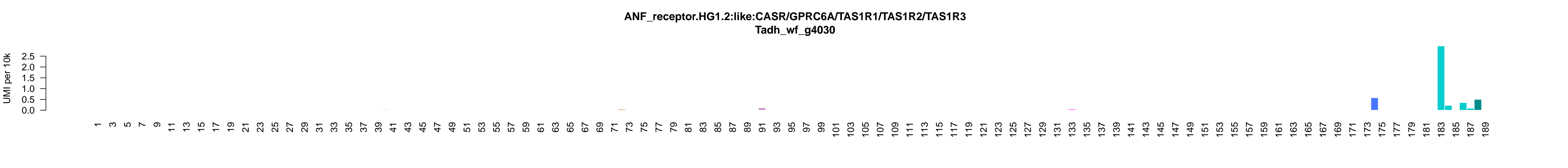
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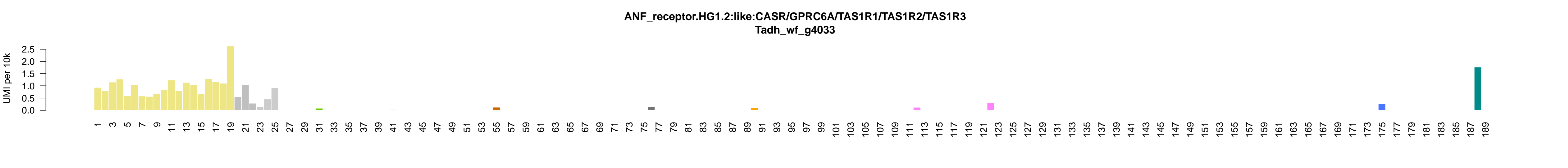
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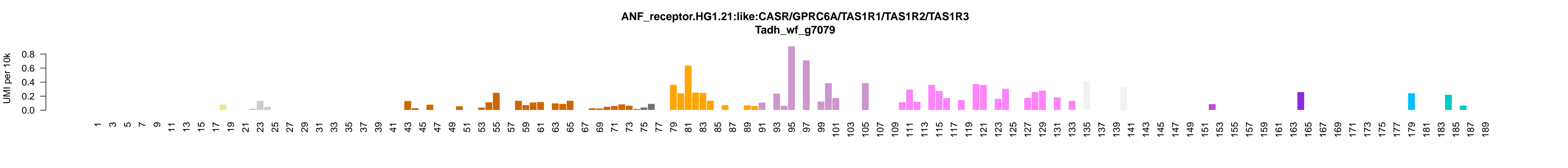


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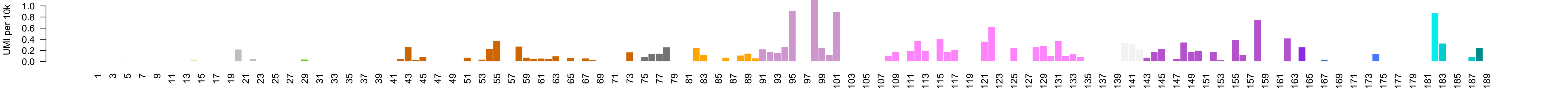


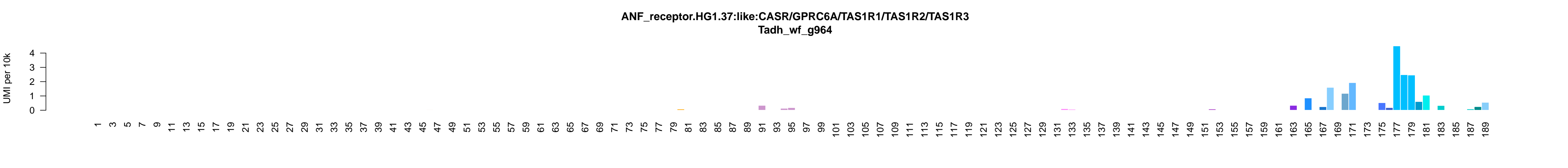




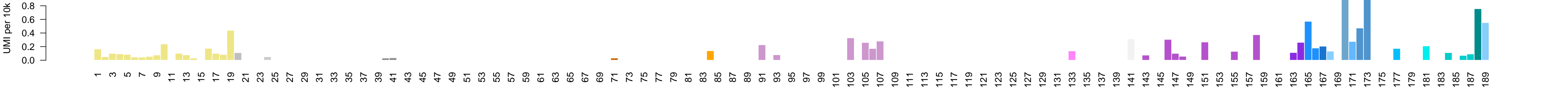


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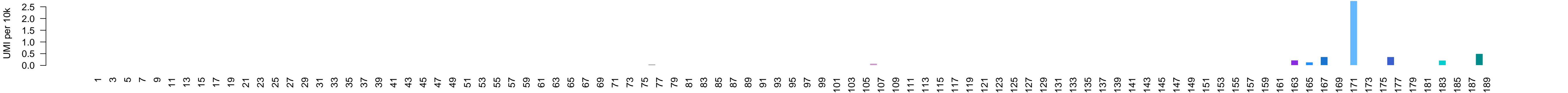




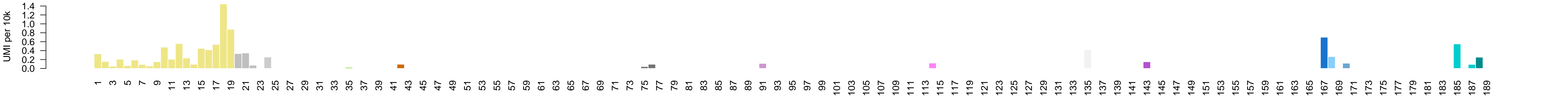
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Tadh_wf_g965

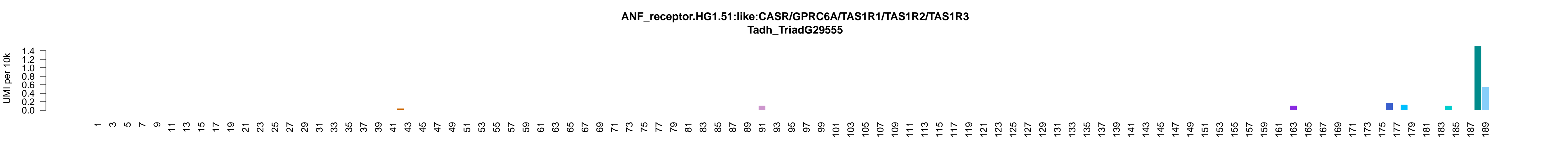


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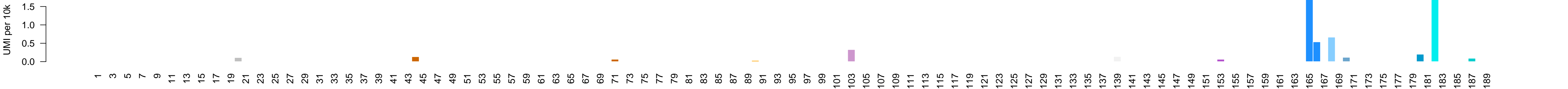


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Tadh_TriadG11320

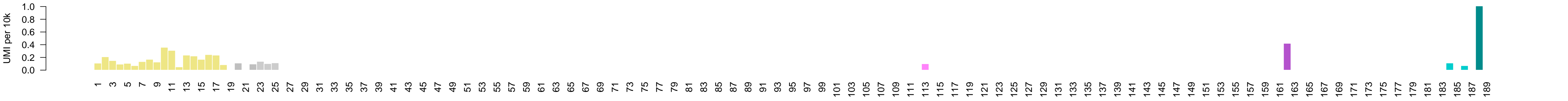




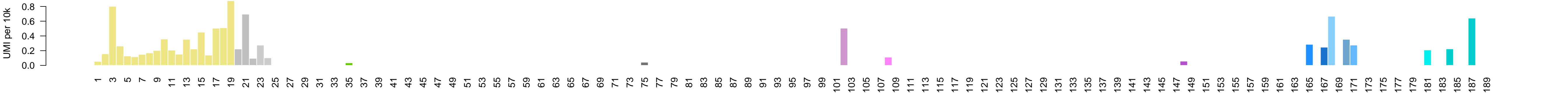
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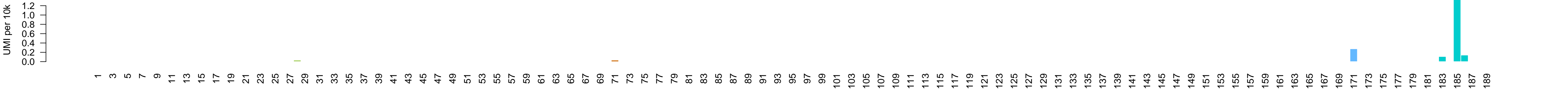
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Tadh_TriadG59566



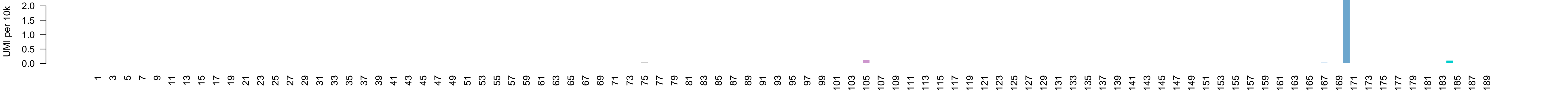
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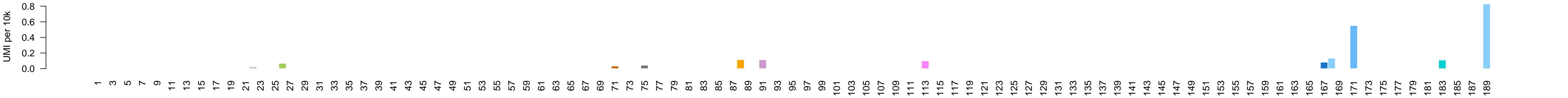
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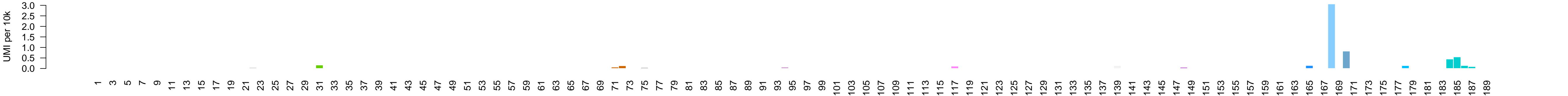
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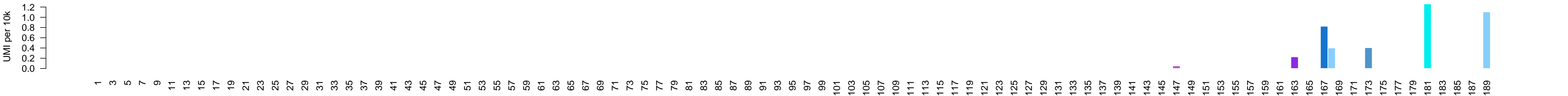
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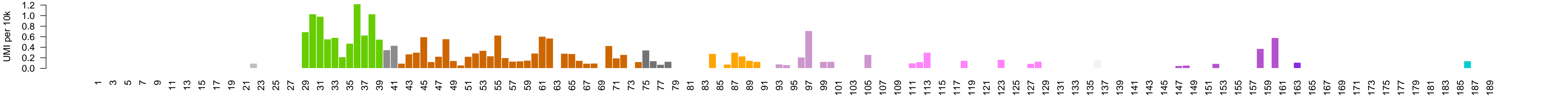
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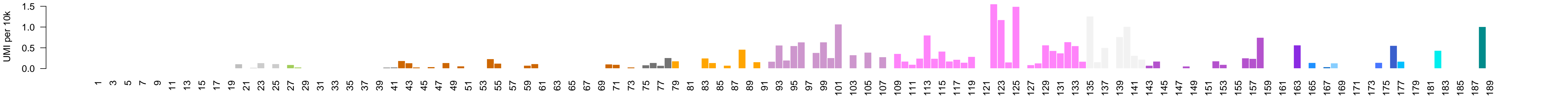
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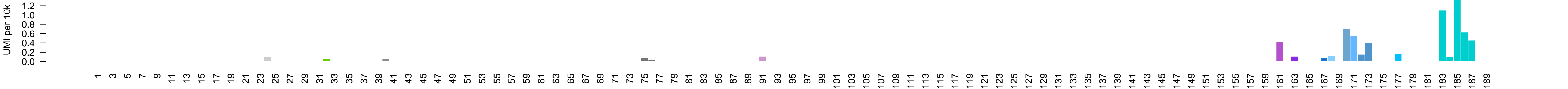
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Tadh_wf_g10200



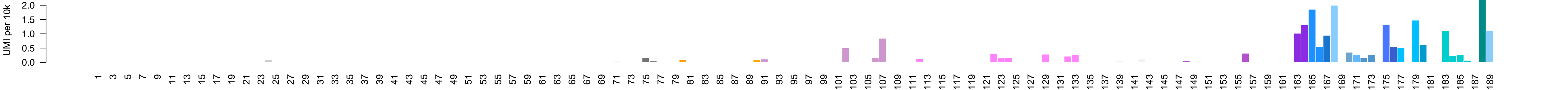
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Tadh_TriadG26726



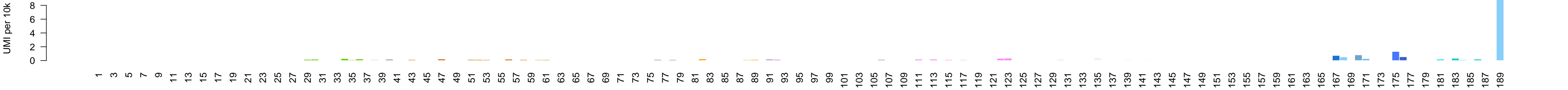
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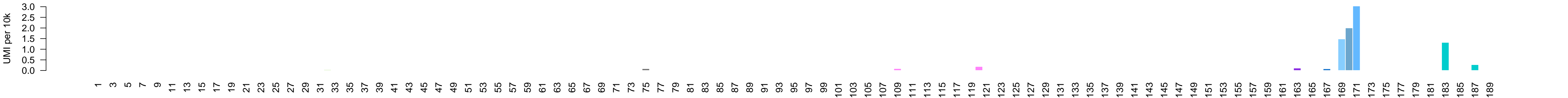
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Tadh_TriadG57193



ANF_receptor.HG2.11:like:GABBR1/GABBR2
Tadh_TriadG58979



ANF_receptor.HG2.13:like:GABBR1/GABBR2
Tadh_TriadG64091



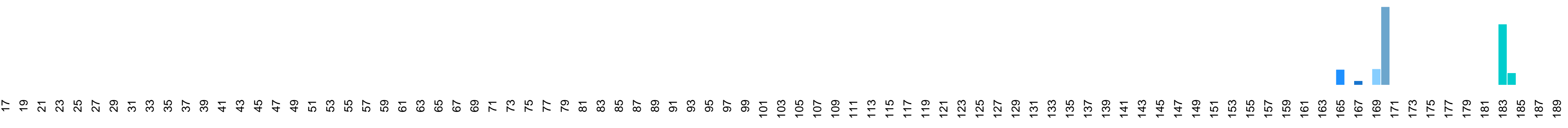
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Tadh_wf_g7102

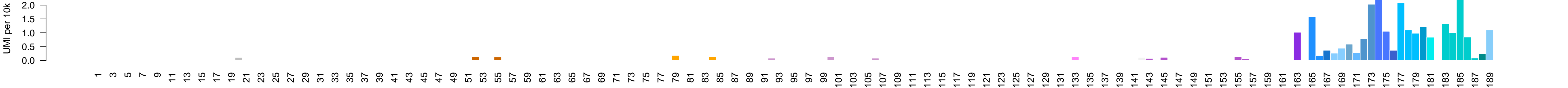
UMI per 10k

0.7
0.6
0.5
0.4
0.3
0.2
0.1
0.0

1 3 5 7 9 11 13 15 17 19 21 23 25 27 29 31 33 35 37 39 41 43 45 47 49 51 53 55 57 59 61 63 65 67 69 71 73 75 77 79 81 83 85 87 89 91 93 95 97 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189



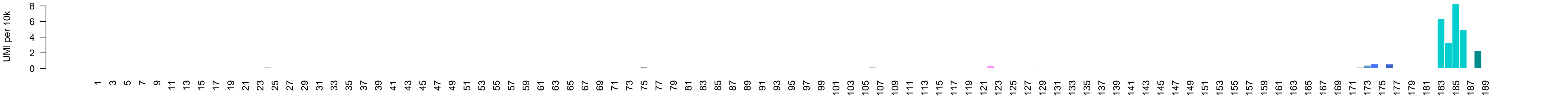
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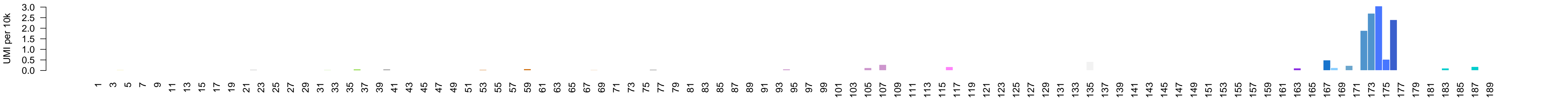
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Tadh_TriadG28568



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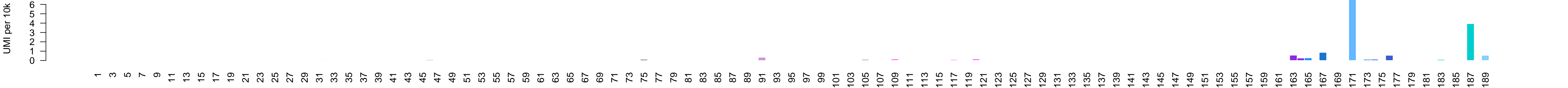


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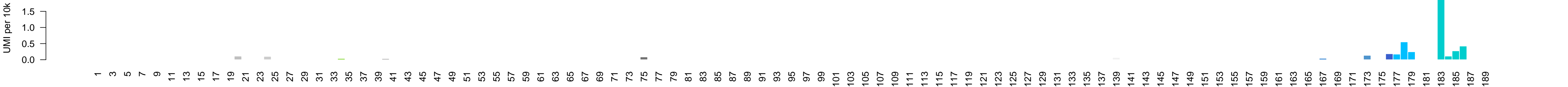


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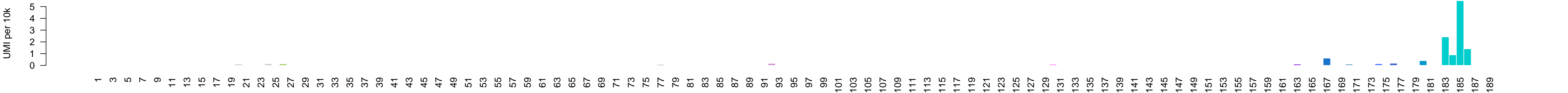
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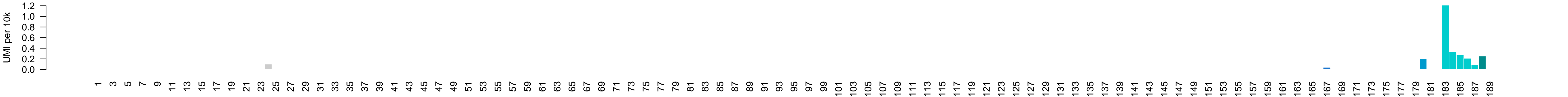
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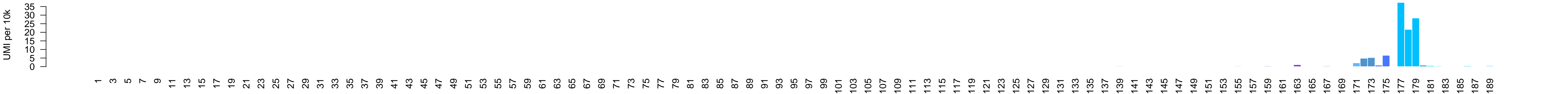
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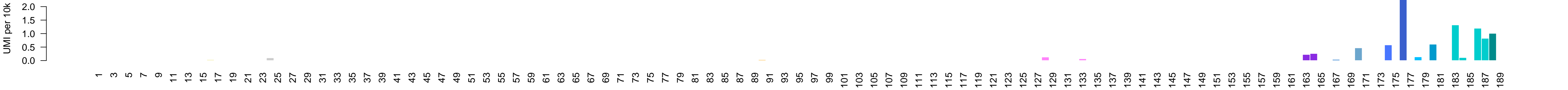
ANF_receptor.HG2.24:like:GABBR1/GABBR2
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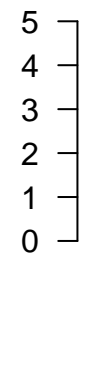


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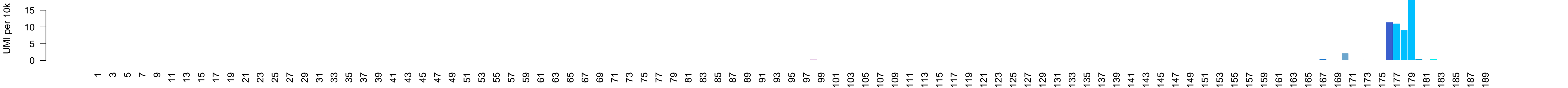


ANF_receptor.HG2.3:like:GABBR1/GABBR2
Tadh_TriadG56610

UMI per 10k



ANF_receptor.HG2.3:like:GABBR1/GABBR2
Tadh_TriadG56611



ANF_receptor.HG2.3:like:GABBR1/GABBR2

Tadh_TriadG60050

UMI per 10k

10
8
6
4
2
0

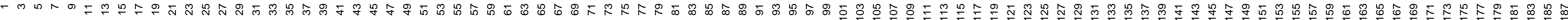
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0

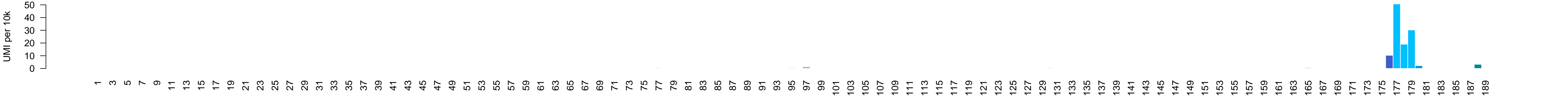
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UMI per 10k

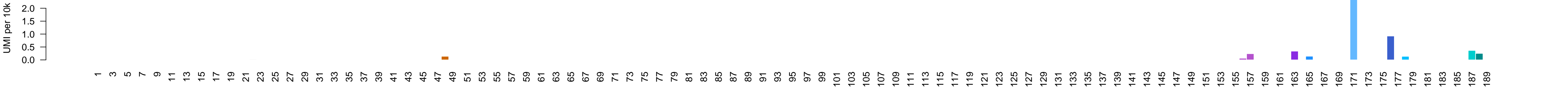
3
2
1
0



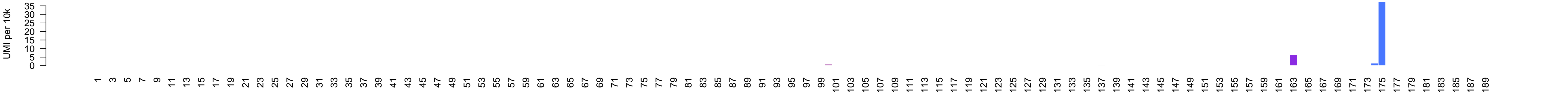
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Tadh_TriadG58938



ANF_receptor.HG2.4:like:GABBR1/GABBR2
Tadh_TriadG55425



ANF_receptor.HG2.4:like:GABBR1/GABBR2
Tadh_TriadG55426



ANF_receptor.HG2.4:like:GABBR1/GABBR2
Tadh_TriadG55427

UMI per 10k

4
3
2
1
0

1 3 5 7 9 11 13 15 17 19 21 23 25 27 29 31 33 35 37 39 41 43 45 47 49 51 53 55 57 59 61 63 65 67 69 71 73 75 77 79 81 83 85 87 89 91 93 95 97 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189

0

1

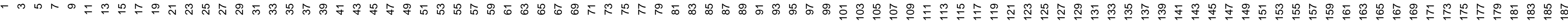
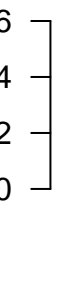
2

3

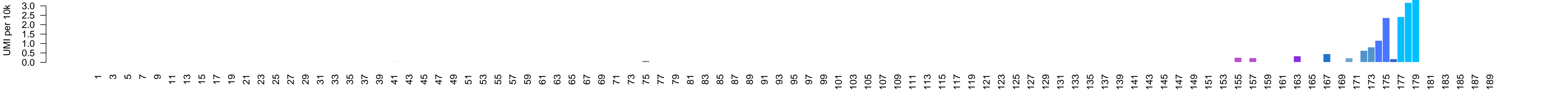
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ANF_receptor.HG2.4:like:GABBR1/GABBR2
Tadh_TriadG55428

UMI per 10k



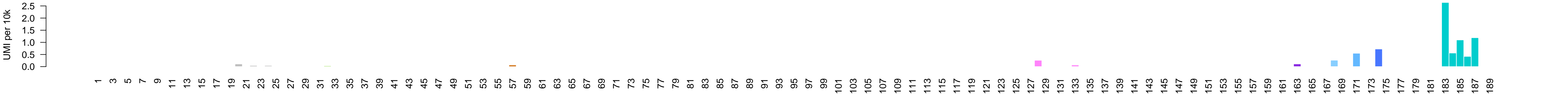
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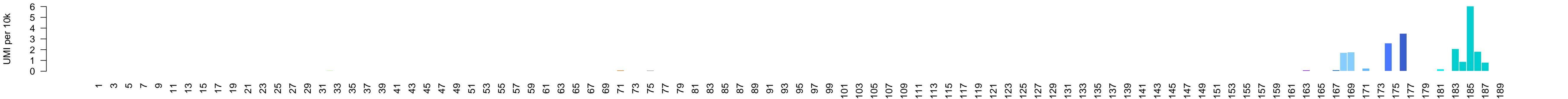
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Tadh_TriadG52576



ANF_receptor.HG2.5:like:GABBR1/GABBR2
Tadh_TriadG52577



ANF_receptor.HG2.5:like:GABBR1/GABBR2
Tadh_TriadG52578

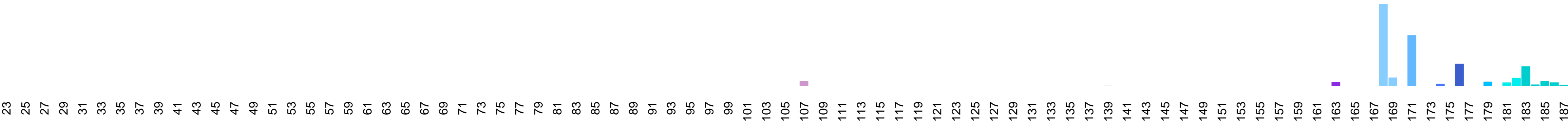


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Tadh_TriadG52579

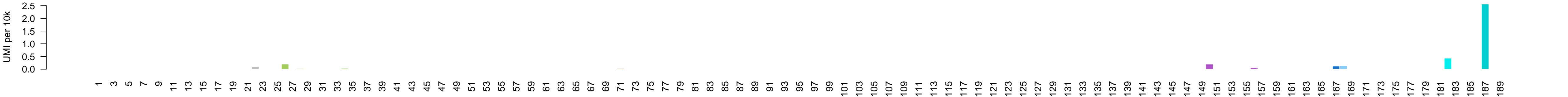
UMI per 10k

3
2
1
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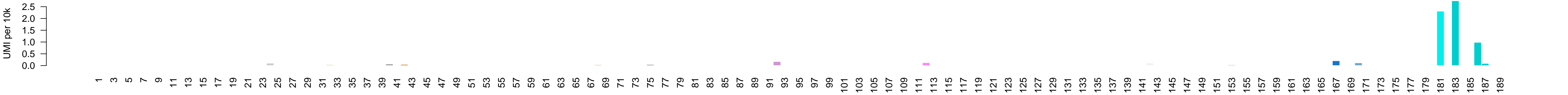
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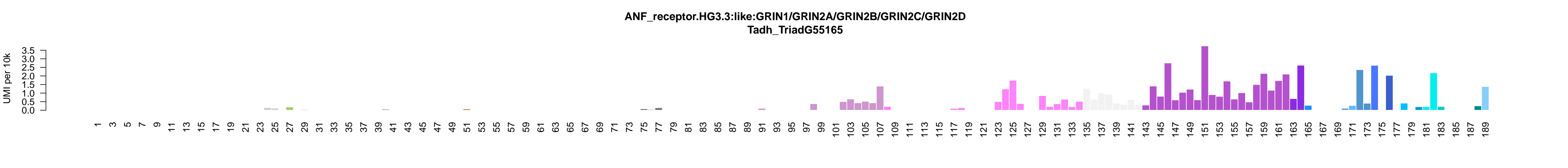


ANF_receptor.HG2.5:like:GABBR1/GABBR2
Tadh_TriadG52582

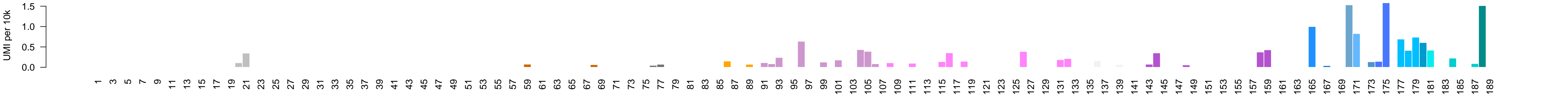


ANF_receptor.HG2.5:like:GABBR1/GABBR2
Tadh_TriadG55500

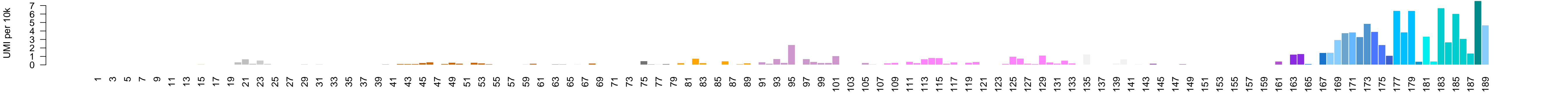




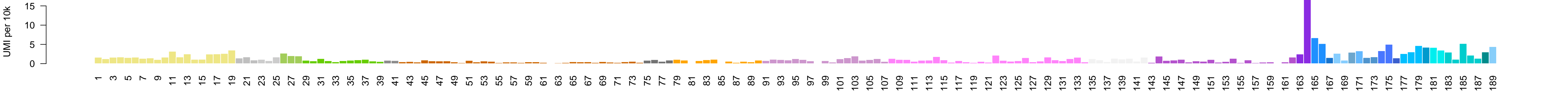
ANF_receptor.HG4.6:like:GUCY2D/GUCY2F/NPR1/NPR2/NPR3
Tadh_TriadG59259



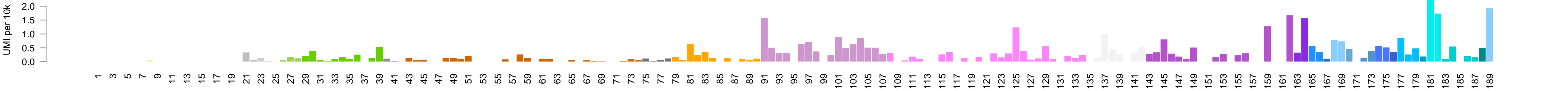
ANF_receptor.HG4.9:like:GUCY2D/GUCY2F/NPR1/NPR2/NPR3
Tadh_TriadG30655



CaMKII.HG1.1:CAMK2A/CAMK2B/CAMK2D/CAMK2G
Tadh_TriadG50756



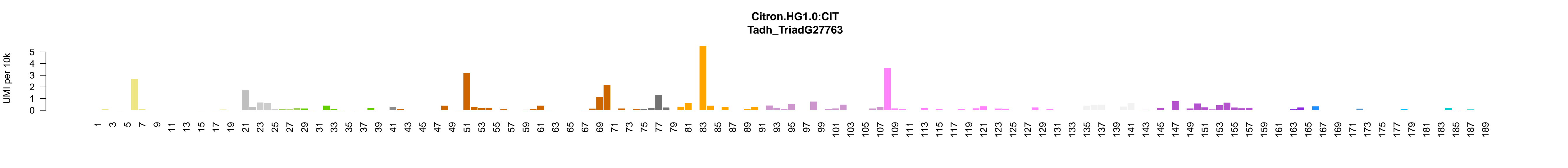
Caps.HG1.1:UNC13A/UNC13B/UNC13C;Synaptotagmin.HG1.66:UNC13A/UNC13C;Unc13.HG1.1:UNC13A/UNC13B/UNC13C
Tadh_TriadG22095



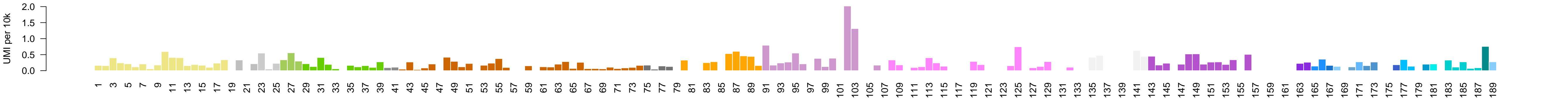
UMI per 10k

Citron.HG1.0:CIT
Tadh_TriadG27763

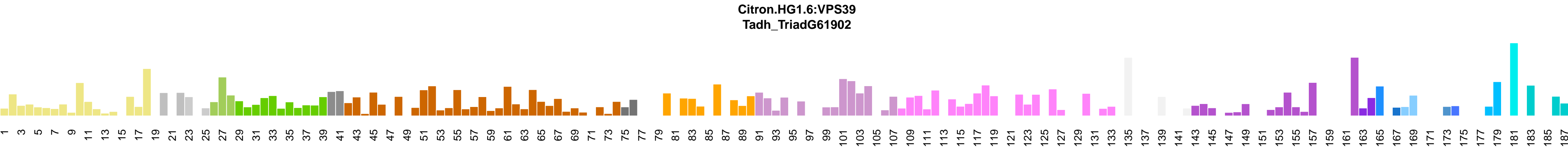
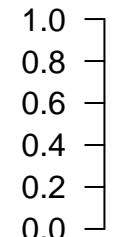
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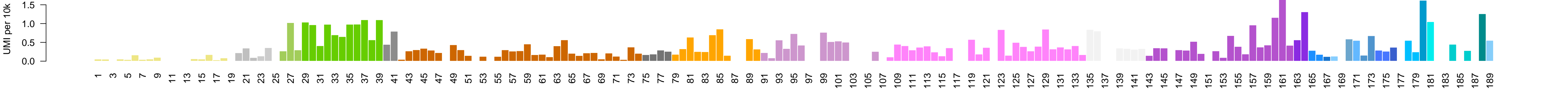
Citron.HG1.3:MAP4K4/MINK1/TNIK
Tadh_TriadG52207



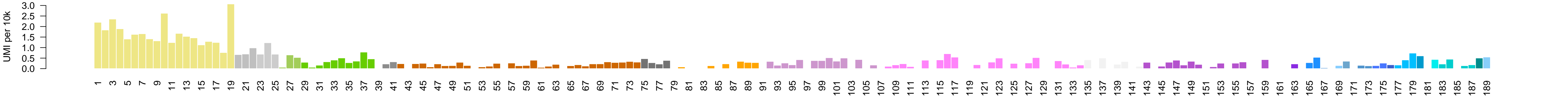
UMI per 10k



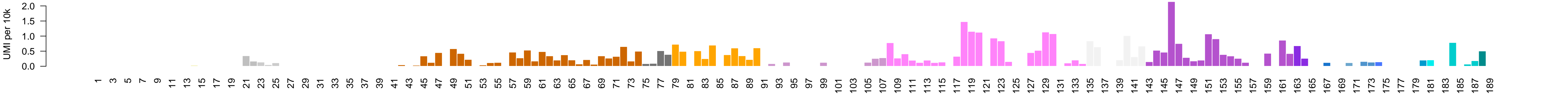
Citron.HG2.0:MAP4K1/MAP4K2/MAP4K3/MAP4K5
Tadh_TriadG51772



COesterase.HG1.1:like:AADAC/AADACL2/ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/NLGN1/NLGN2/NLGN3/NLGN4X/NLGN4Y/TG
Tadh_TriadG59917



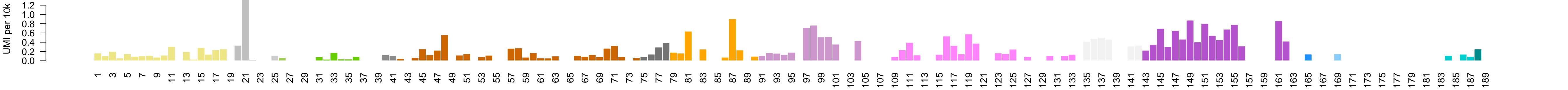
COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG1836



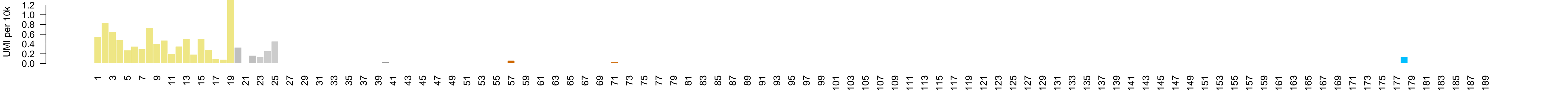
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Tadh_TriadG1914



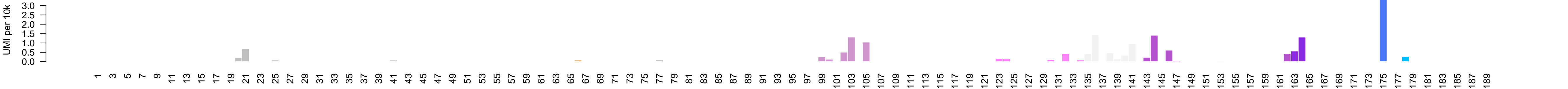
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Tadh_TriadG19242



COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG2392



COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG31923

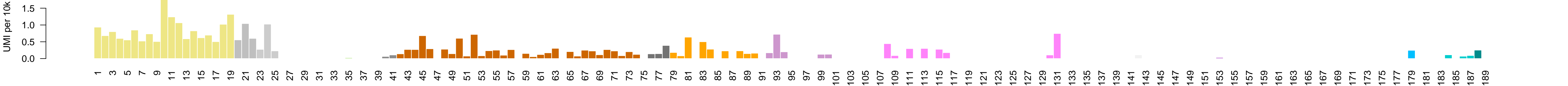


COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG31933

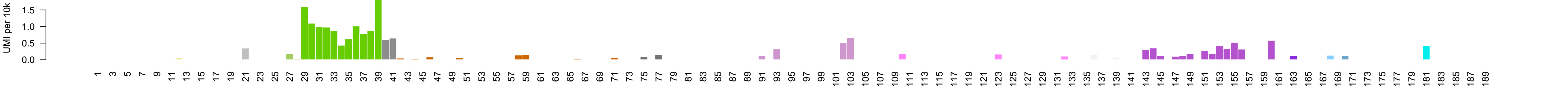


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Tadh_TriadG31975

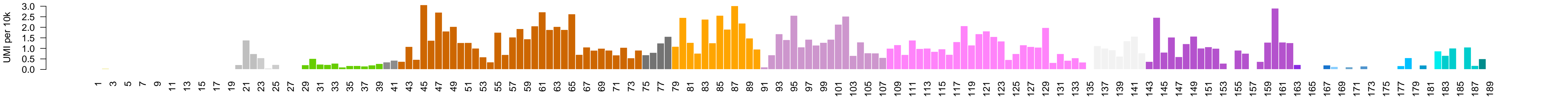


COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG32031

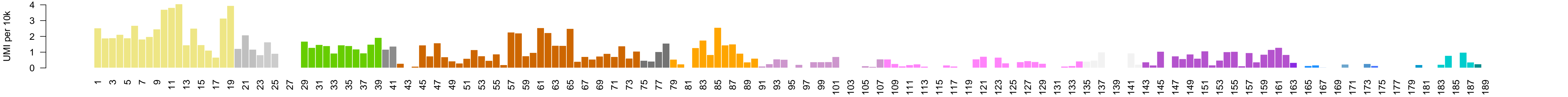


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Tadh_TriadG32071

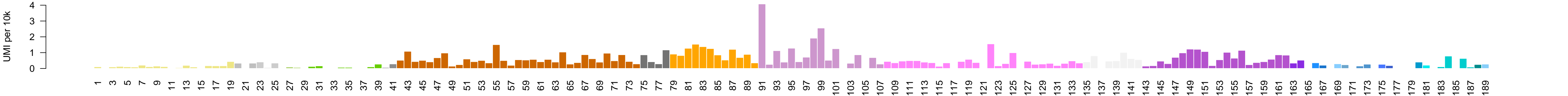


COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG32073



COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG

Tadh_TriadG32085

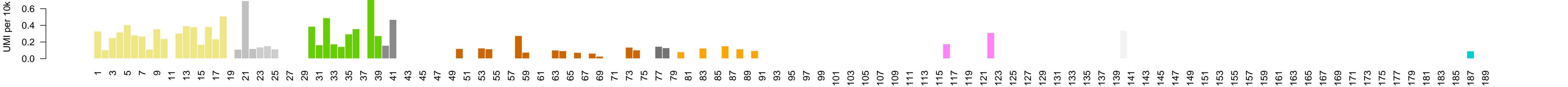


COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG

Tadh_TriadG32115

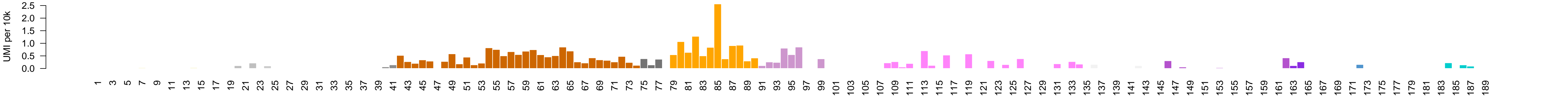


COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
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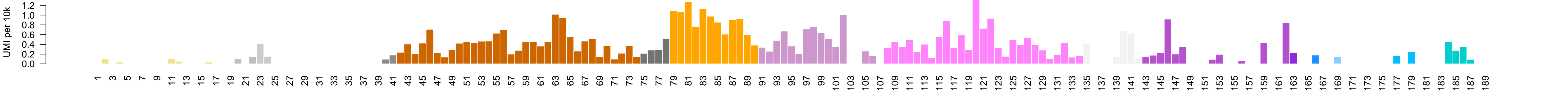


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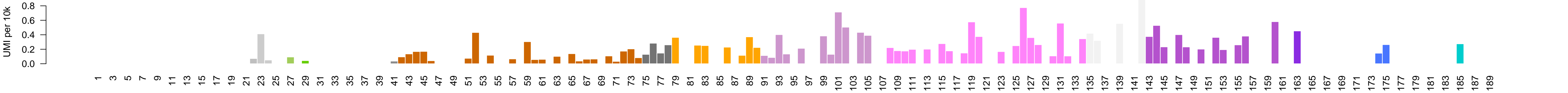
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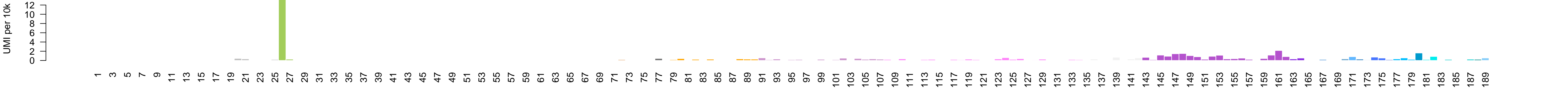
COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG61106



COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG61115



COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG61120



COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
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UMI per 10k

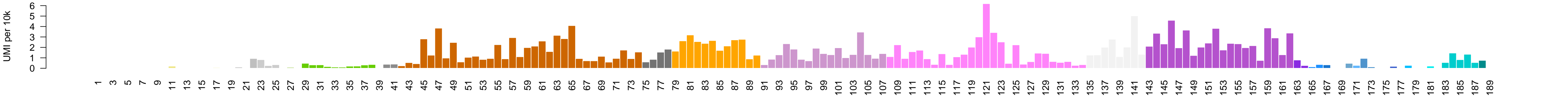
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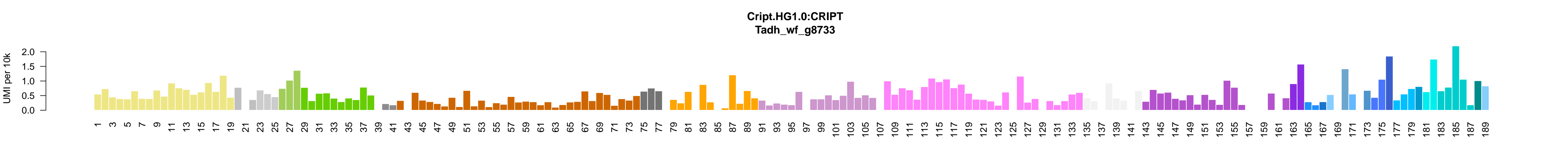
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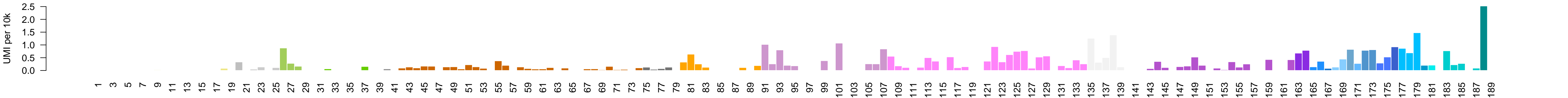
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Tadh_TriadG64430

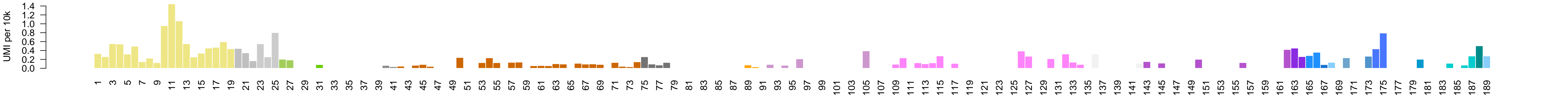




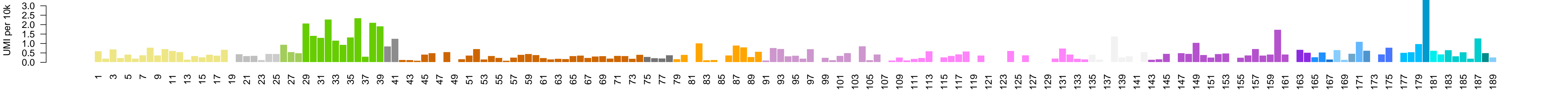
FERM_M.HG1.1:like:MYO10/PLEKHH1/PLEKHH2/PLEKHH3
Tadh_TriadG59851



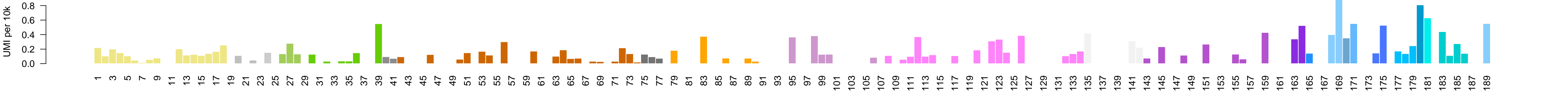
FERM_M.HG1.14:like:MYO7A/MYO7B
Tadh_TriadG61967



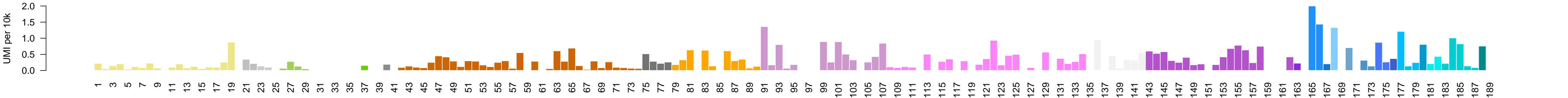
FERM_M.HG1.18:TLN1/TLN2
Tadh_TriadG19795



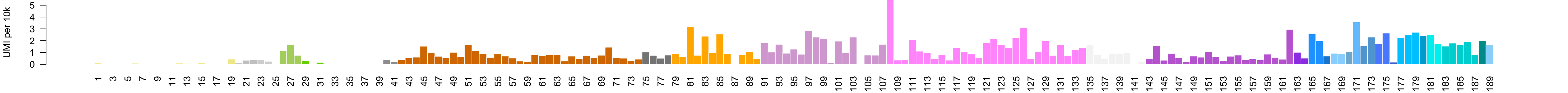
FERM_M.HG1.22:PTPN14/PTPN21;Y_phosphatase.HG1.37:PTPN14/PTPN21
Tadh_TriadG61918



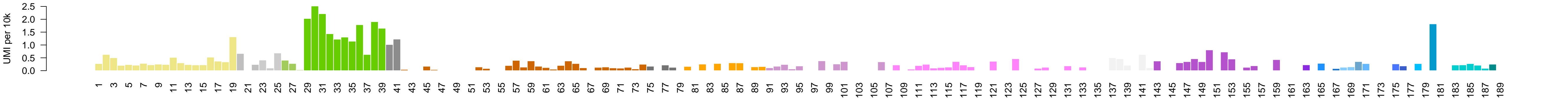
FERM_M.HG1.24:FRMD1/FRMD6/FRMPD2/PTPN13
Tadh_TriadG18577



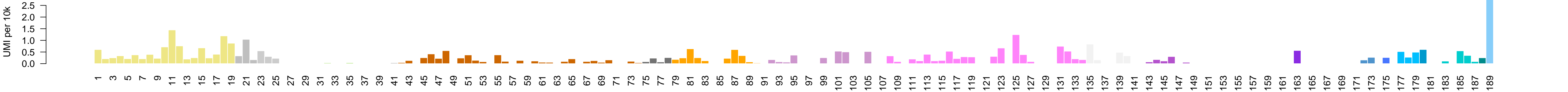
FERM_M.HG1.24:FRMD1/FRMD6/FRMPD2/PTPN13;Y_phosphatase.HG1.21:PTPN13/PTPN20
Tadh_TriadG64201



FERM_M.HG1.26:FERMT1/FERMT2/FERMT3
Tadh_TriadG33394

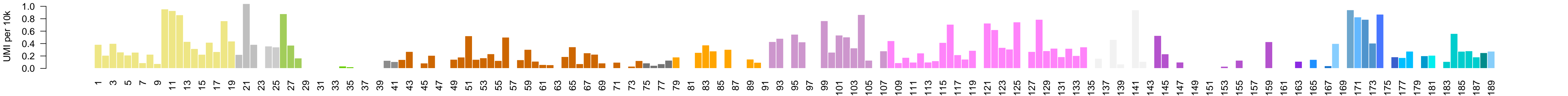


FERM_M.HG1.3:PLEKHH1/PLEKHH2
Tadh_TriadG33842



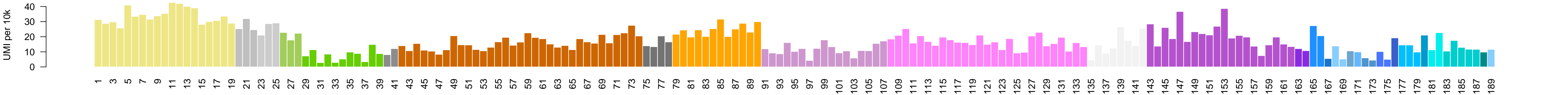
FERM_M.HG1.32:EPB41/EPB41L1/EPB41L2/EPB41L3/EPB41L4A/EPB41L4B/EPB41L5/FARP1/FARP2/FRMD3/FRMD5/FRMD7/PTPN3/PTPN4

Tadh_TriadG23231



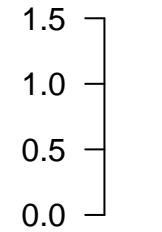
FERM_M.HG1.32:EPB41/EPB41L1/EPB41L2/EPB41L3/EPB41L4A/EPB41L4B/EPB41L5/FARP1/FARP2/FRMD3/FRMD5/FRMD7/PTPN3/PTPN4

Tadh_TriadG64069

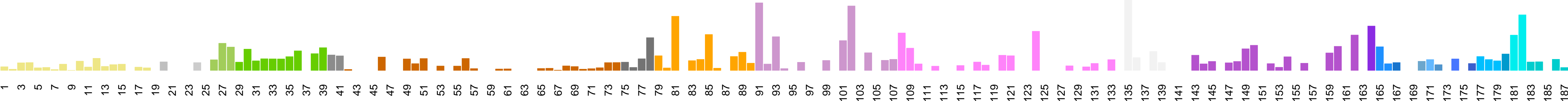


FERM_M.HG1.32:EPB41/EPB41L1/EPB41L2/EPB41L3/EPB41L4A/EPB41L4B/EPB41L5/FARP1/FARP2/FRMD3/FRMD5/FRMD7/PTPN3/PTPN4;Y_phosphatase.HG1.38:PTPN3/PTPN4
Tadh_TriadG63802

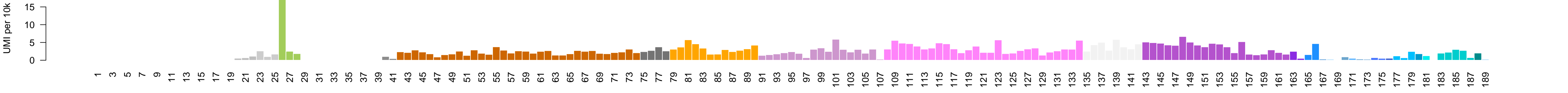
UMI per 10k



1 3 5 7 9 11 13 15 17 19 21 23 25 27 29 31 33 35 37 39 41 43 45 47 49 51 53 55 57 59 61 63 65 67 69 71 73 75 77 79 81 83 85 87 89 91 93 95 97 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189



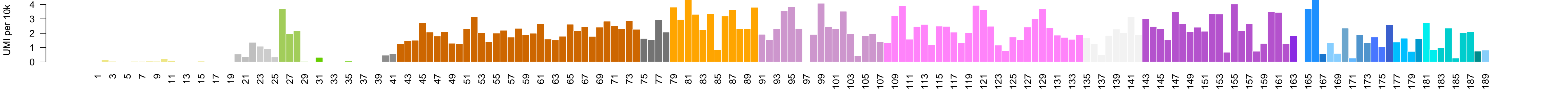
FERM_M.HG1.33:like:EZR/MSN/NF2/RDX
Tadh_TriadG29123



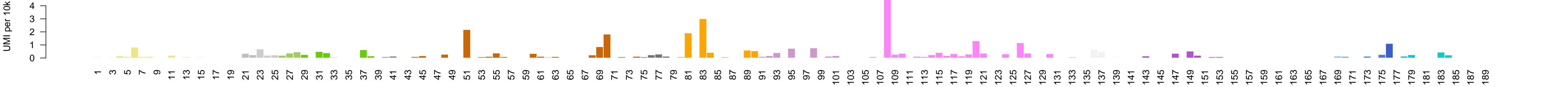
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Tadh_TriadG54062



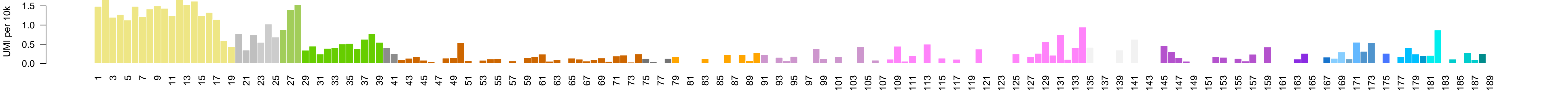
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Tadh_TriadG25177



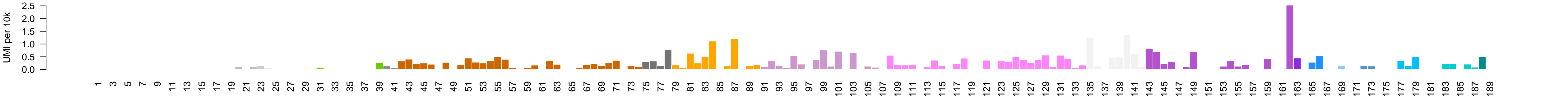
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Tadh_TriadG57926



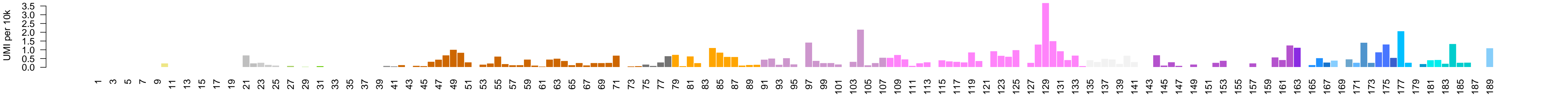
GKAP.HG1.1:DLGAP1/DLGAP2/DLGAP3/DLGAP4
Tadh_TriadG60595



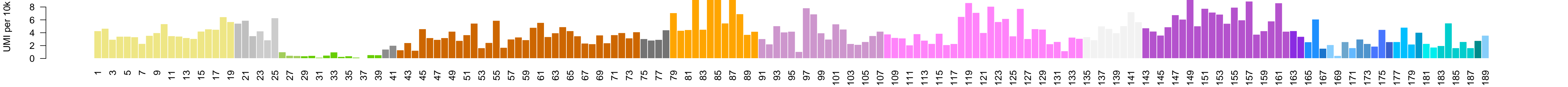
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Tadh_TriadG60291



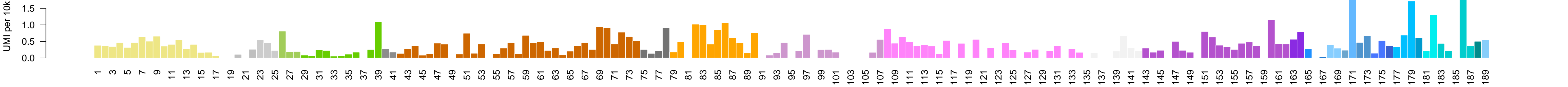
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Tadh_TriadG52391



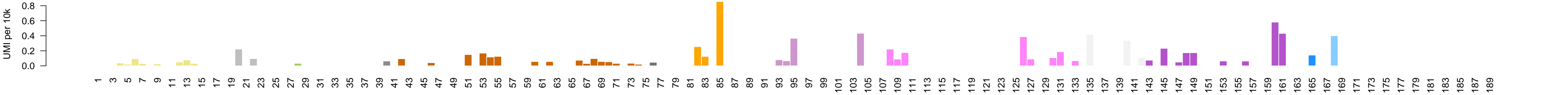
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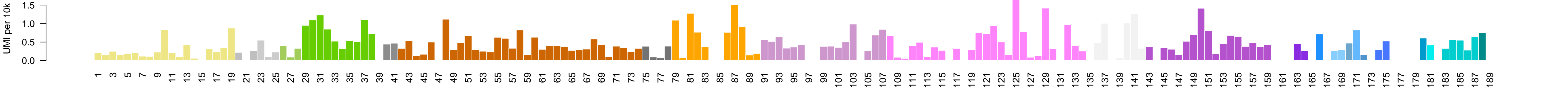
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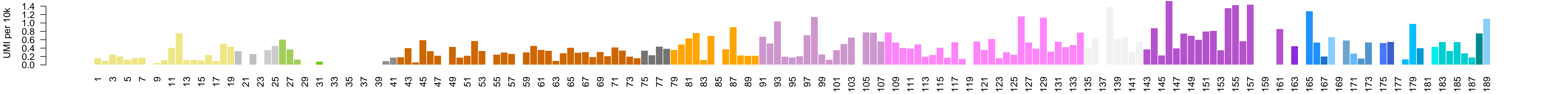
Guanylate_kin.HG1.2:like:AL358113.1/CASK/DLG1/DLG2/DLG3/DLG4/DLG5/GUK1/LRGUK/MPP1/MPP2/MPP3/MPP4/MPP5/MPP6/MPP7/TJP1/TJP2/TJP3;PDZ.HG2.11:MPP5
Tadh_TriadG52734



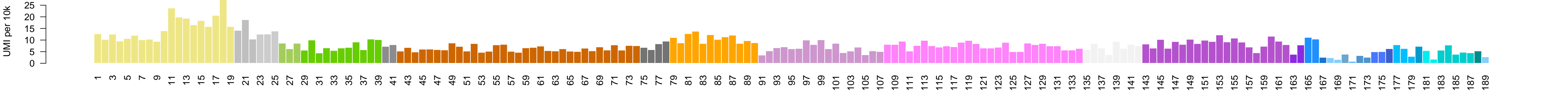
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Tadh_TriadG55243



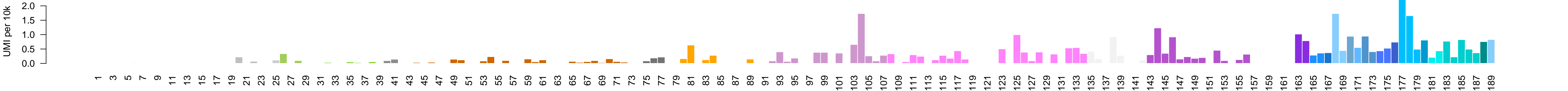
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Tadh_TriadG54413



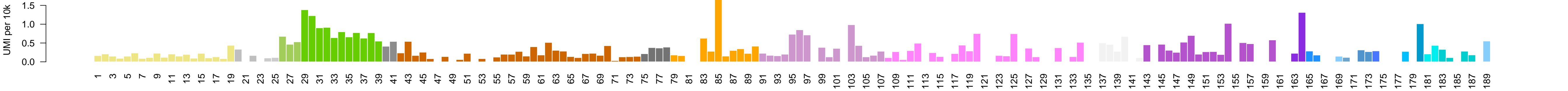
Guanylate_kin.HG1.9:like:CASK/MPP1/MPP2/MPP6;PDZ.HG2.5:MPP2/MPP6
Tadh_TriadG63748



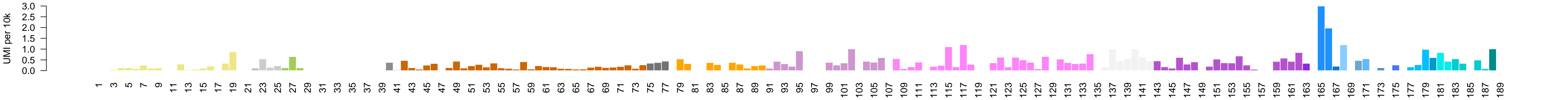
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Tadh_TriadG22195



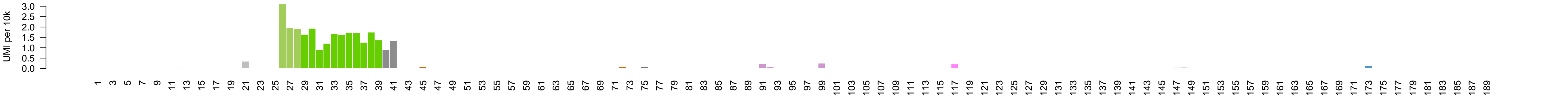
Homer.HG1.1:ENAH/EVL/SPRED1/SPRED2/SPRED3/VASP
Tadh_TriadG57130



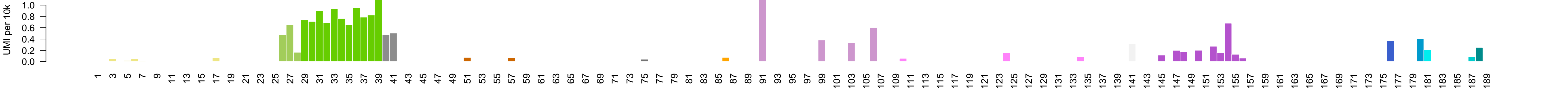
Homer.HG1.2:HOMER1/HOMER2/HOMER3
Tadh_TriadG22944



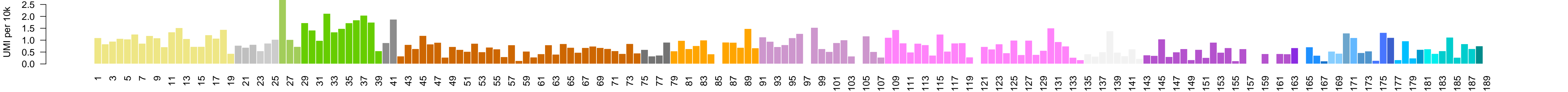
Homer.HG1.6:like:WAS/WASL
Tadh_TriadG54495



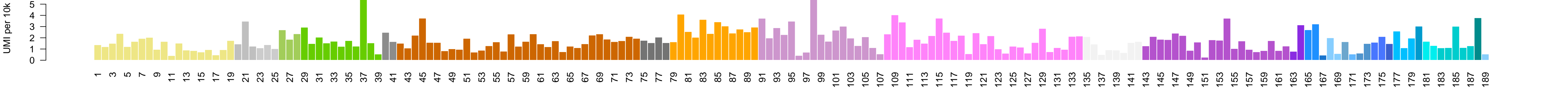
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Tadh_TriadG54960



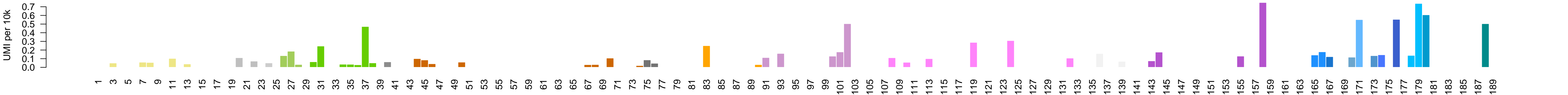
Homer.HG1.9:WAS/WASL
Tadh_TriadG54962



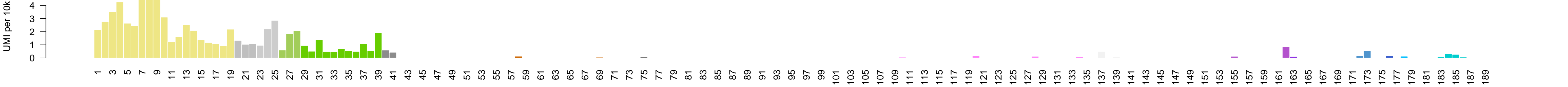
MARVEL.HG1.13:SYNPR/SYP/SYPL1/SYPL2
Tadh_TriadG52445



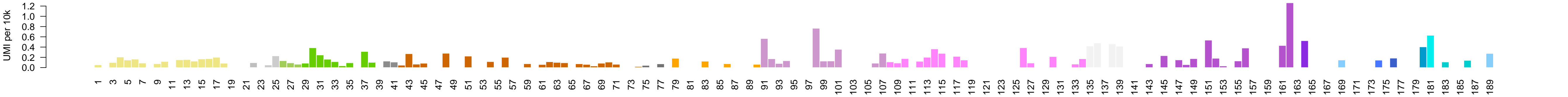
MARVEL.HG1.13:SYNPR/SYP/SYPL1/SYPL2
Tadh_TriadG62635



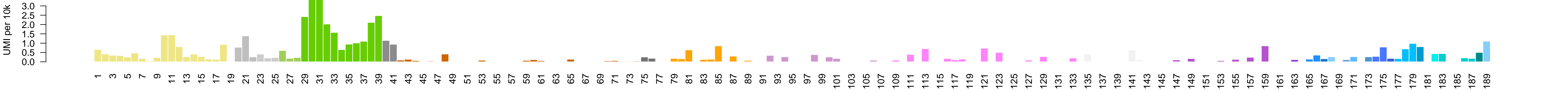
MARVEL.HG1.8:like:OCLN/SYNPR/SYP/SYPL1/SYPL2
Tadh_TriadG63863



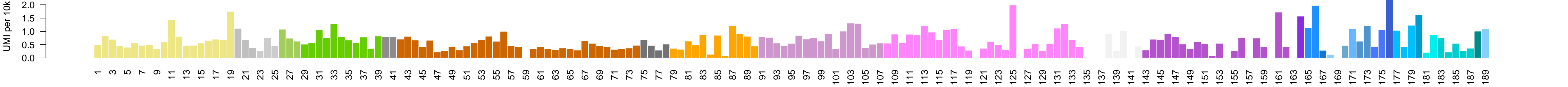
MARVEL.HG2.0:SYNGR1/SYNGR2/SYNGR4
Tadh_TriadG56946



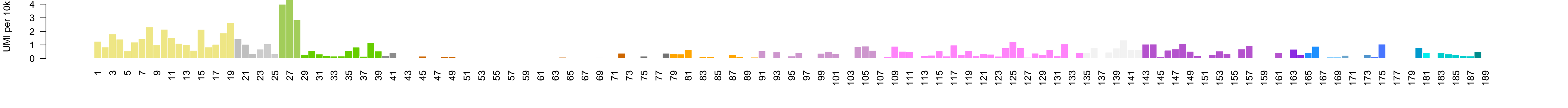
MFS_1.HG1.16:SLC16A2/SLC16A10
Tadh_TriadG19777



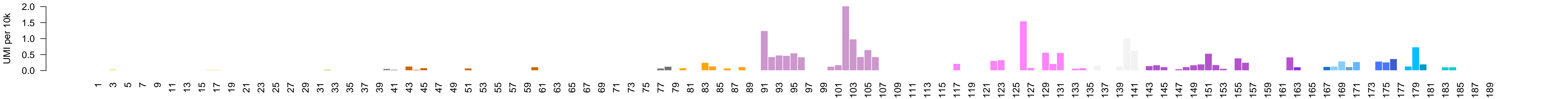
MFS_1.HG10.1:SLC49A3
Tadh_TriadG32990



MFS_1.HG10.3:like:SLC49A3
Tadh_TriadG60867

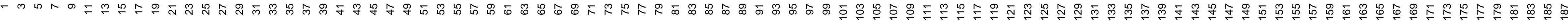
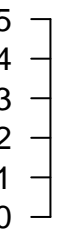


MFS_1.HG11.7:SLC37A1/SLC37A2
Tadh_TriadG60236

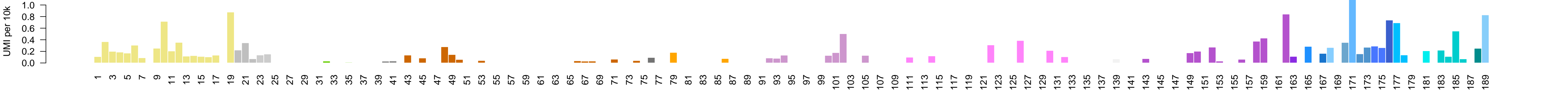


MFS_1.HG12.2:like:MFSD6
Tadh_TriadG60583

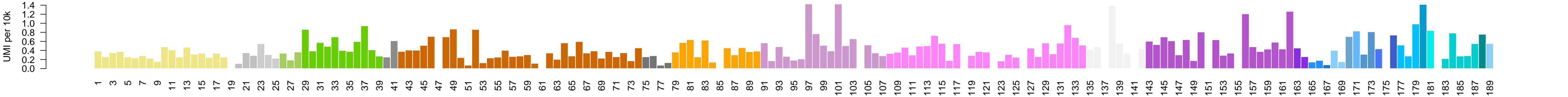
UMI per 10k



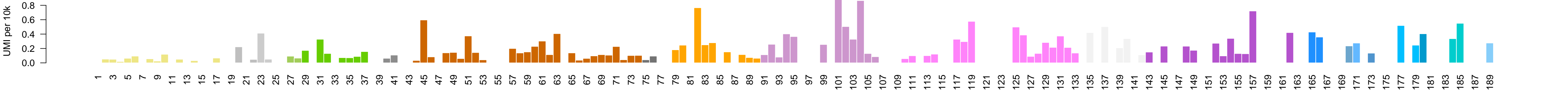
MFS_1.HG12.3:MFSD6
Tadh_TriadG1695



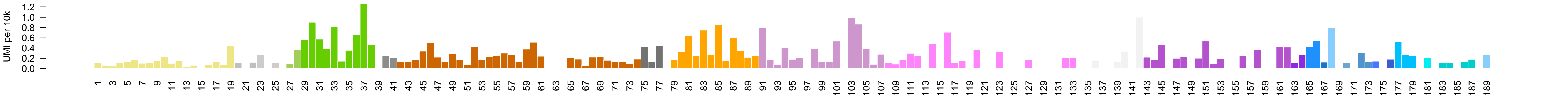
MFS_1.HG14.0:SPNS1/SPNS2/SPNS3
Tadh_TriadG63211



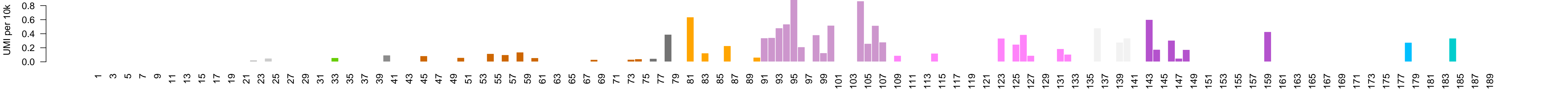
FS_1.HG19.2:MFSD12
Tadh_TriadG30948



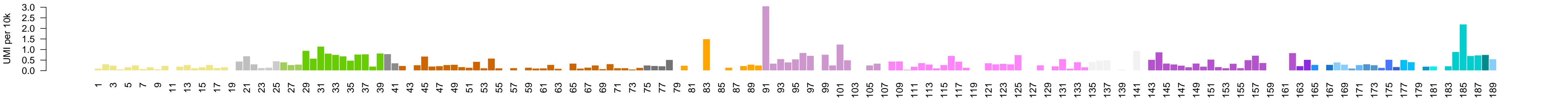
MFS_1.HG21.1:SLC15A3/SLC15A4
Tadh_TriadG57267



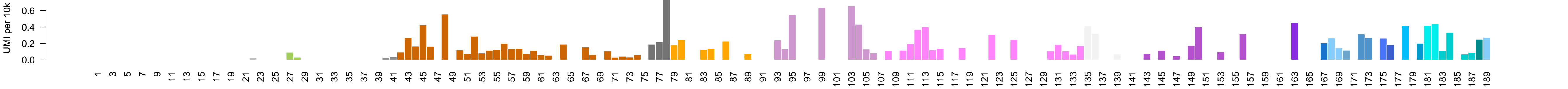
MFS_1.HG3.29:like:SV2A/SV2B/SV2C;
Sugar_tr.HG2.28:like:SV2A/SV2B/SV2C
Tadh_TriadG2343



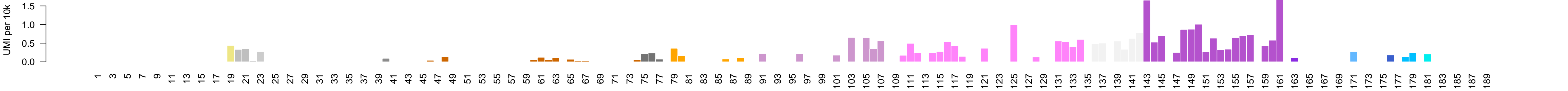
MFS_1.HG3.3:SV2A/SV2B/SV2C;Sugar_tr.HG2.26:like:SV2A/SV2B/SV2C
Tadh_TriadG36521



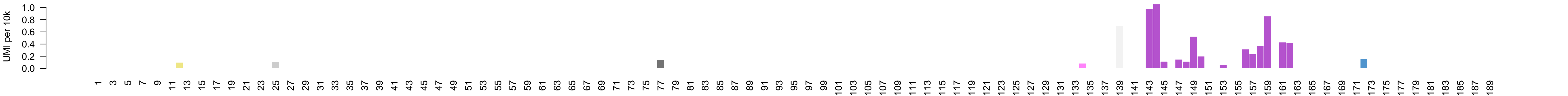
MFS_1.HG3.38:SVOP/SVOPL;Sugar_tr.HG2.59:SVOP
Tadh_TriadG58334



MFS_1.HG3.38:SVOP/SVOPL;Sugar_tr.HG2.59:SVOP
Tadh_TriadG58337



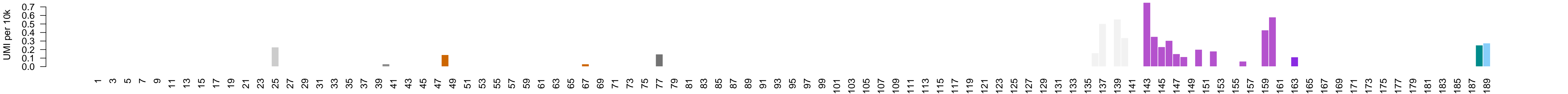
MFS_1.HG3.39:like:SV2A/SV2B/SV2C;
Tadh_TriadG31400
Sugar_tr.HG2.56:like:SV2A/SV2B/SV2C



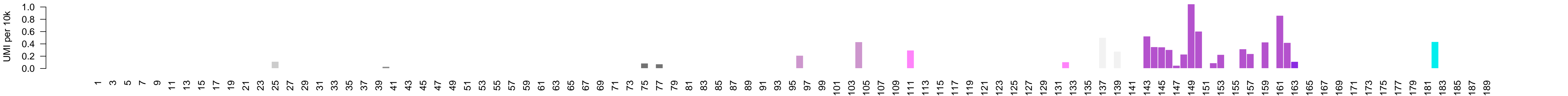
MFS_1.HG3.39:like:SV2A/SV2B/SV2C;
Tadh_TriadG60673



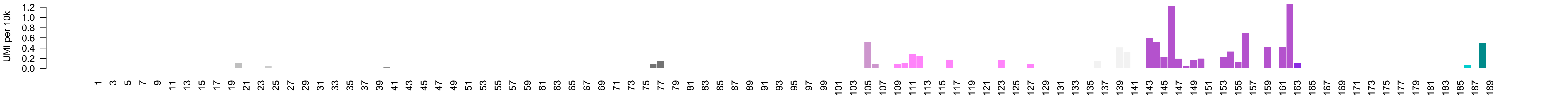
MFS_1.HG3.42:like:SV2A/SV2B/SV2C;
Sugar_tr.HG2.49:like:SV2A/SV2B/SV2C
Tadh_TriadG31425



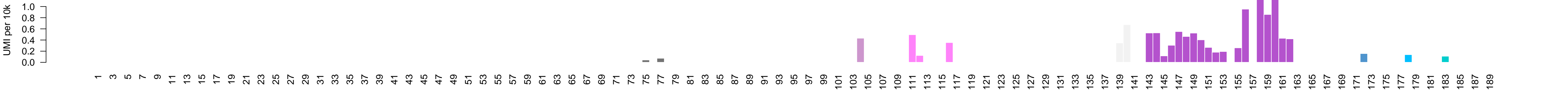
MFS_1.HG3.44:like:SV2A/SV2B/SV2C;
Tadh_TriadG60671



MFS_1.HG3.53:like:SV2A/SV2B/SV2C;
Sugar_tr.HG2.63:like:SV2A/SV2B/SV2C
Tadh_TriadG60690



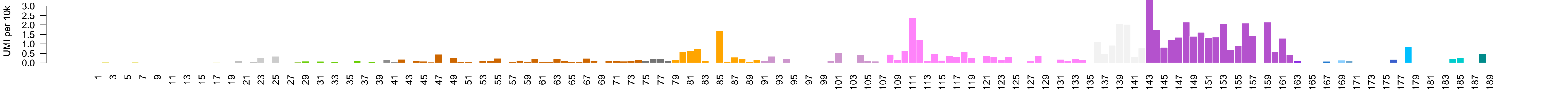
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Tadh_TriadG60691



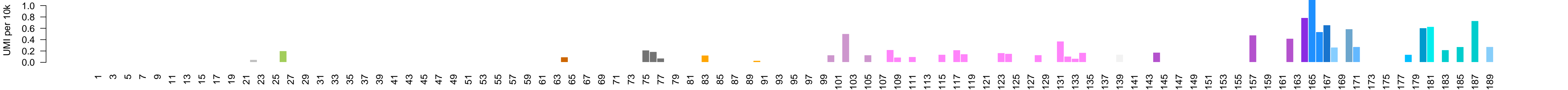
MFS_1.HG3.53:like:SV2A/SV2B/SV2C;
Sugar_tr.HG2.63:like:SV2A/SV2B/SV2C
Tadh_TriadG60692



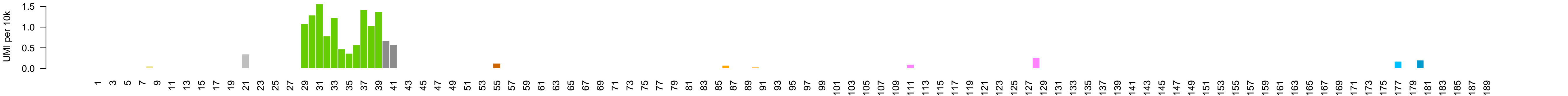
MFS_1.HG3.54:like:SV2A/SV2B/SV2C;
Sugar_tr.HG2.64:like:SV2A/SV2B/SV2C
Tadh_TriadG596

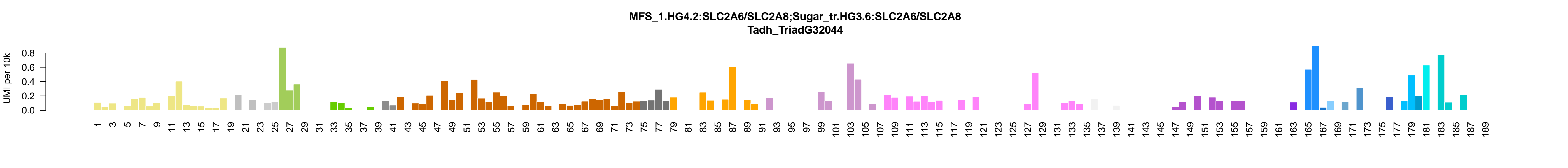


MFS_1.HG4.2:SLC2A6/SLC2A8;Sugar_tr.HG3.6:SLC2A6/SLC2A8
Tadh_TriadG31909

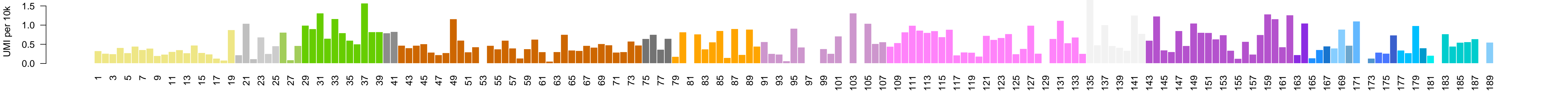


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Tadh_TriadG32030

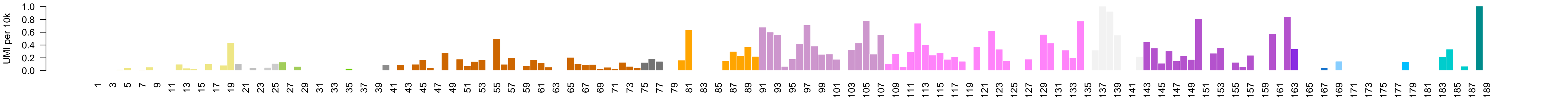




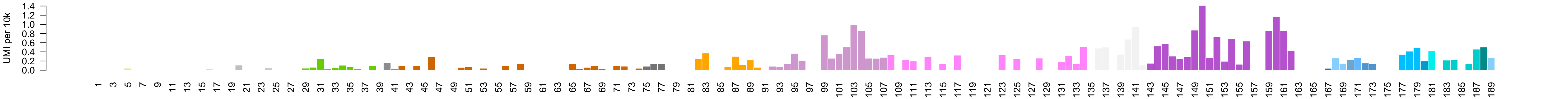
MFS_1.HG4.2:SLC2A6/SLC2A8;Sugar_tr.HG3.6:SLC2A6/SLC2A8
Tadh_TriadG38377



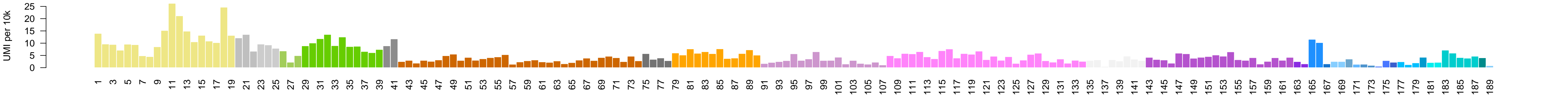
MFS_1.HG4.2:SLC2A6/SLC2A8;Sugar_tr.HG3.6:SLC2A6/SLC2A8
Tadh_TriadG58298



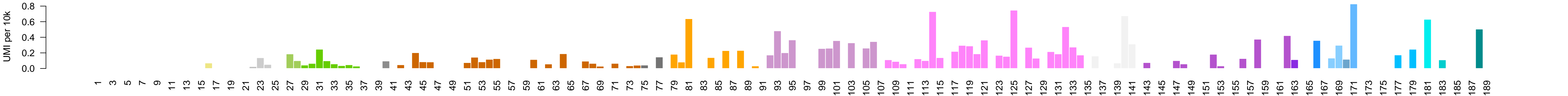
MFS_1.HG4.2:SLC2A6/SLC2A8;Sugar_tr.HG3.6:SLC2A6/SLC2A8
Tadh_TriadG61088



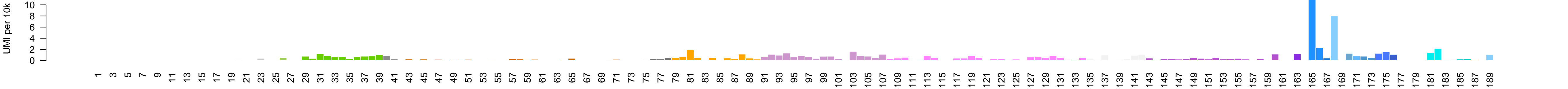
MFS_1.HG4.3:like:SLC2A10/SLC2A12;Sugar_tr.HG3.4:like:SLC2A10/SLC2A12/SLC2A13
Tadh_TriadG64243



MFS_1.HG4.4:SLC2A10/SLC2A12;Sugar_tr.HG3.3:SLC2A10/SLC2A12
Tadh_TriadG58068



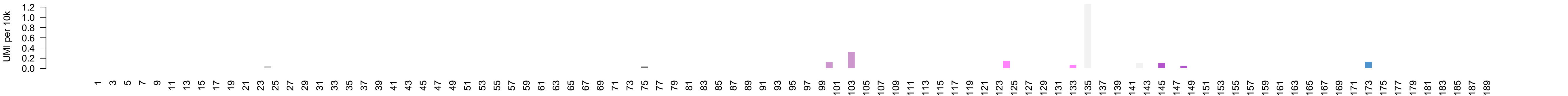
MFS_1.HG5.21:SLC17A6/SLC17A7/SLC17A8
Tadh_TriadG36003



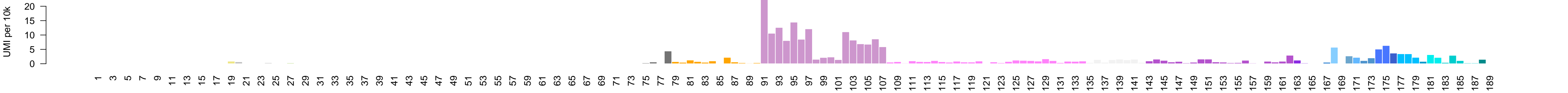
MFS_1.HG5.22:like:SLC17A6/SLC17A7/SLC17A8
Tadh_TriadG61197



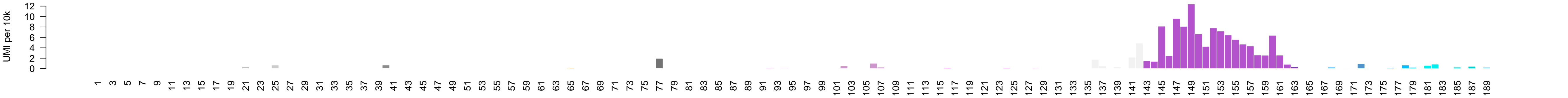
MFS_1.HG5.8:SLC17A9
Tadh_TriadG18643



MFS_1.HG5.8:SLC17A9;Sugar_tr.HG5.5:NA
Tadh_TriadG36875

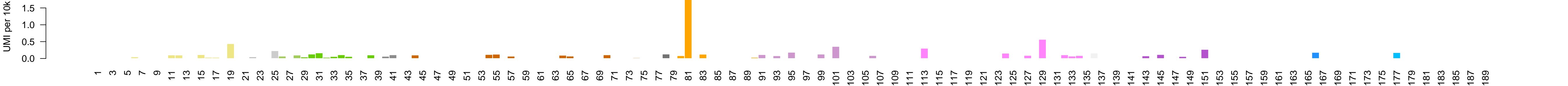


MFS_1.HG6.1:like:SLCO1A2/SLCO1B1/SLCO1B3/SLCO1B3–SLCO1B7/SLCO1C1/SLCO2A1/SLCO2B1/SLCO3A1/SLCO4A1/SLCO4C1/SLCO5A1/SLCO6A1
Tadh_TriadG1002



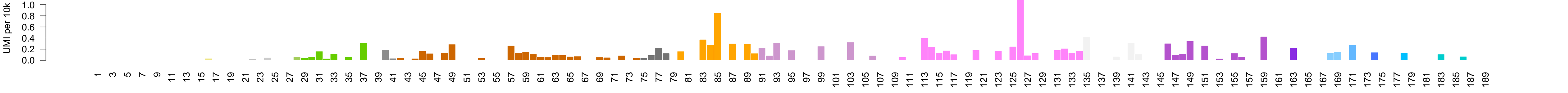
MFS_1.HG6.1:like:SLCO1A2/SLCO1B1/SLCO1B3/SLCO1B3–SLCO1B7/SLCO1C1/SLCO2A1/SLCO2B1/SLCO3A1/SLCO4A1/SLCO4C1/SLCO5A1/SLCO6A1

Tadh_TriadG1023



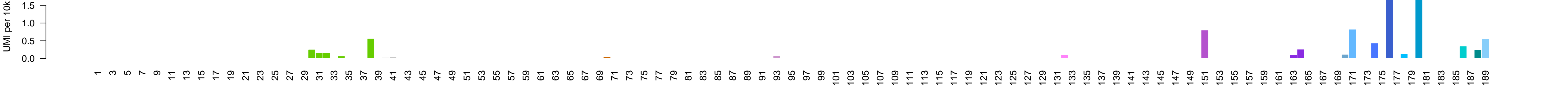
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Tadh_TriadG11569



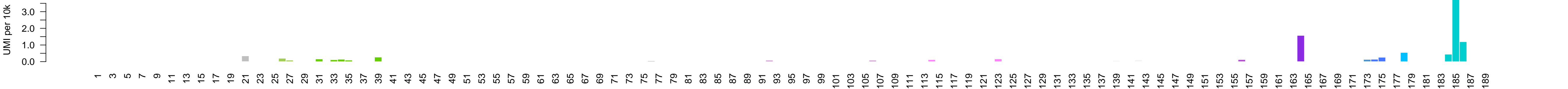
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Tadh_TriadG11683



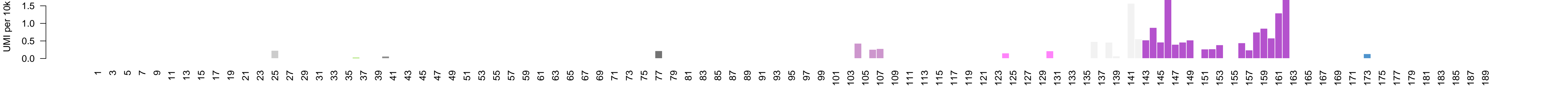
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Tadh_TriadG11804



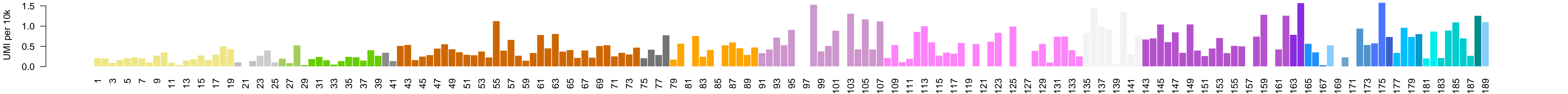
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Tadh_TriadG18955

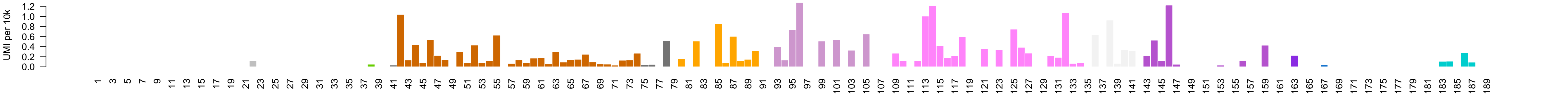


MFS_1.HG6.1:like:SLCO1A2/SLCO1B1/SLCO1B3/SLCO1B3–SLCO1B7/SLCO1C1/SLCO2A1/SLCO2B1/SLCO3A1/SLCO4A1/SLCO4C1/SLCO5A1/SLCO6A1

Tadh_TriadG31000

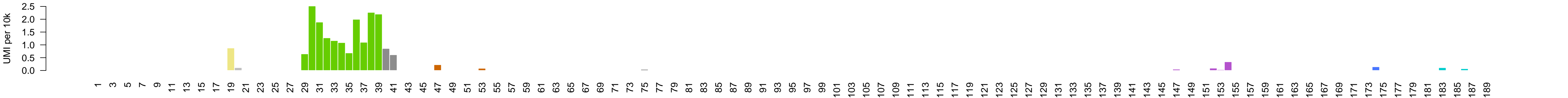


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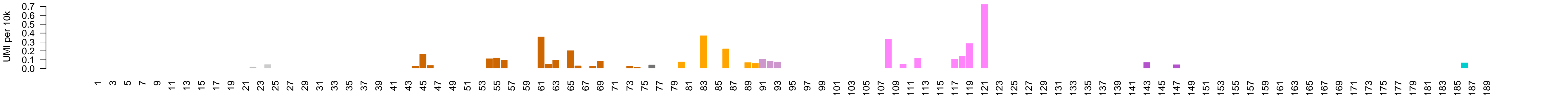


MFS_1.HG6.1:like:SLCO1A2/SLCO1B1/SLCO1B3/SLCO1B3–SLCO1B7/SLCO1C1/SLCO2A1/SLCO2B1/SLCO3A1/SLCO4A1/SLCO4C1/SLCO5A1/SLCO6A1

Tadh_TriadG33189

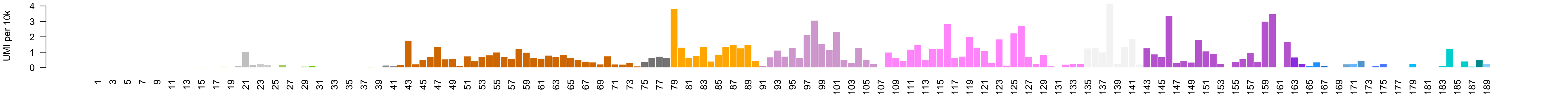


MFS_1.HG6.1:like:SLCO1A2/SLCO1B1/SLCO1B3/SLCO1B3–SLCO1B7/SLCO1C1/SLCO2A1/SLCO2B1/SLCO3A1/SLCO4A1/SLCO4C1/SLCO5A1/SLCO6A1
Tadh_TriadG58340

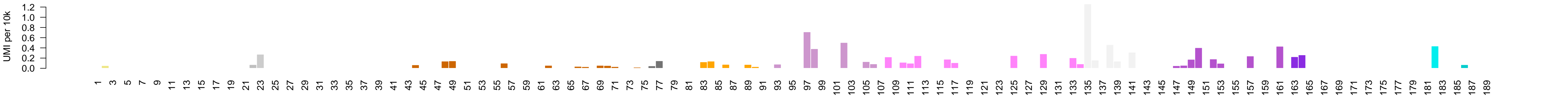


MFS_1.HG6.1:like:SLCO1A2/SLCO1B1/SLCO1B3/SLCO1B3–SLCO1B7/SLCO1C1/SLCO2A1/SLCO2B1/SLCO3A1/SLCO4A1/SLCO4C1/SLCO5A1/SLCO6A1

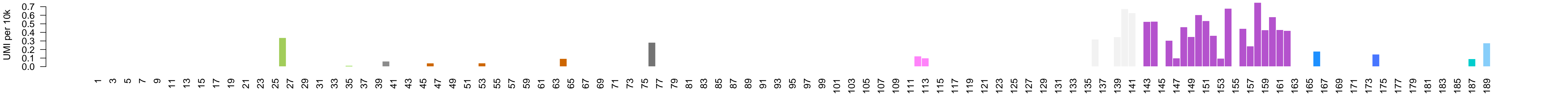
Tadh_TriadG968



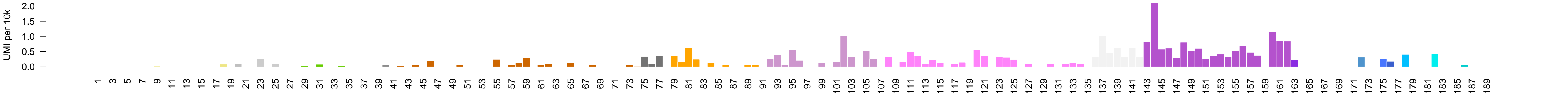
MFS_1.HG7.0:SLC18A1/SLC18A2/SLC18A3/SLC18B1
Tadh_TriadG51495



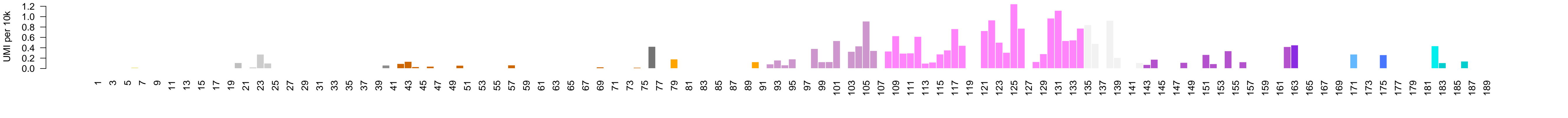
MFS_1.HG7.0:SLC18A1/SLC18A2/SLC18A3/SLC18B1
Tadh_TriadG58799



MFS_1.HG7.0:SLC18A1/SLC18A2/SLC18A3/SLC18B1
Tadh_TriadG58954

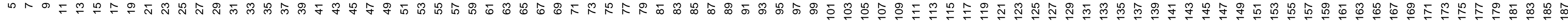
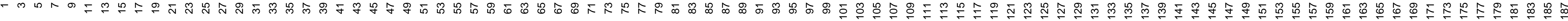
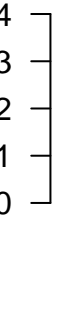


MFS_1.HG7.0:SLC18A1/SLC18A2/SLC18A3/SLC18B1
Tadh_TriadG58955

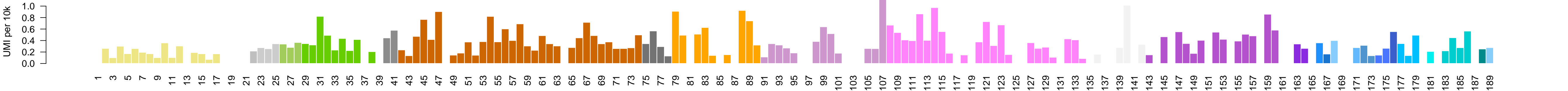


MFS_1.HG7.0:SLC18A1/SLC18A2/SLC18A3/SLC18B1
Tadh_TriadG58957

UMI per 10k

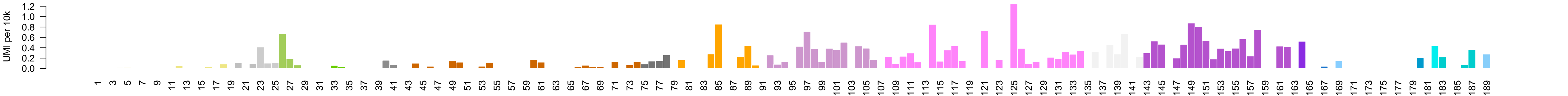


MFS_1.HG7.0:SLC18A1/SLC18A2/SLC18A3/SLC18B1
Tadh_TriadG58958

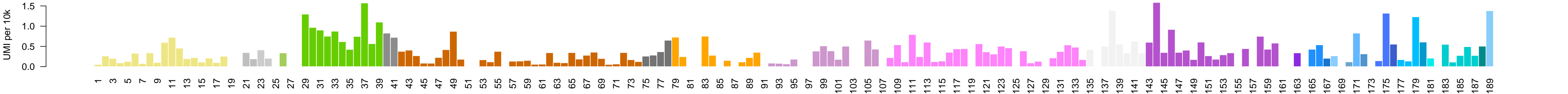


MFS_1.HG7.0:SLC18A1/SLC18A2/SLC18A3/SLC18B1

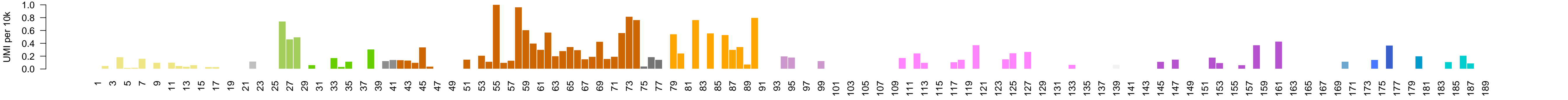
Tadh_TriadG58959



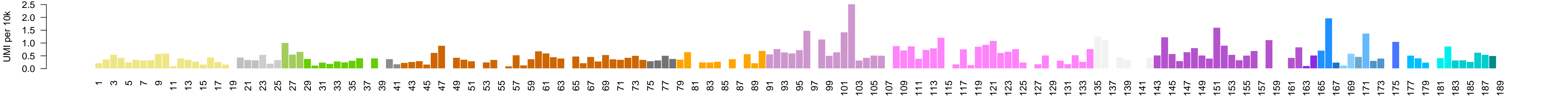
MFS_1.HG7.0:SLC18A1/SLC18A2/SLC18A3/SLC18B1
Tadh_TriadG64145



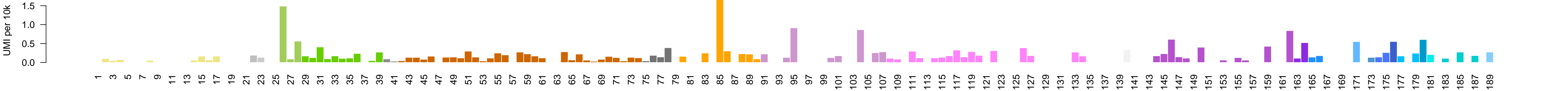
MFS_1.HG8.11:SLC22A18
Tadh_TriadG27771



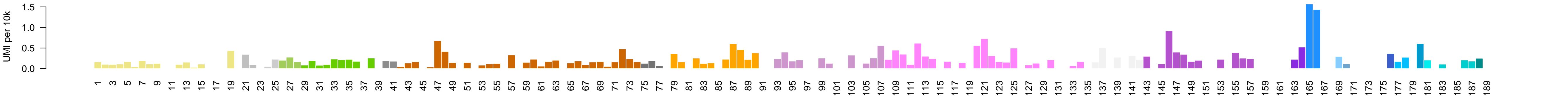
MFS_1.HG8.11:SLC22A18
Tadh_TriadG58186



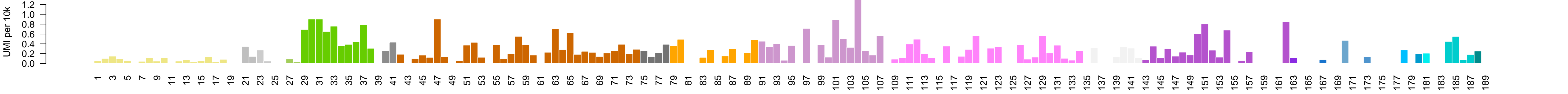
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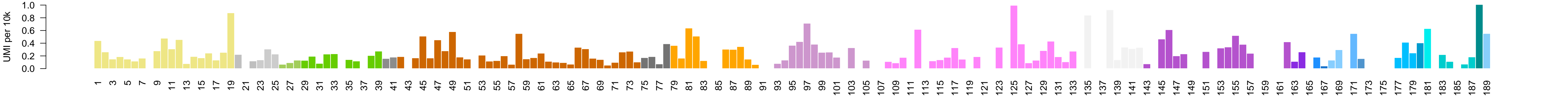
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Tadh_TriadG25848



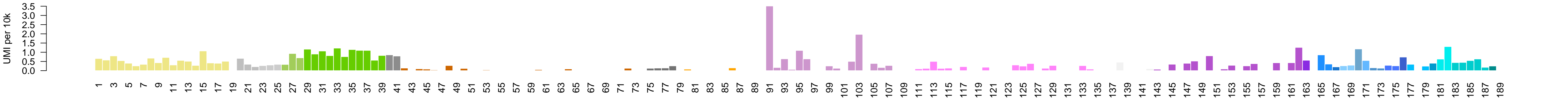
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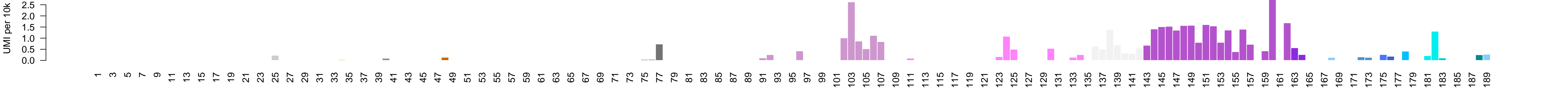
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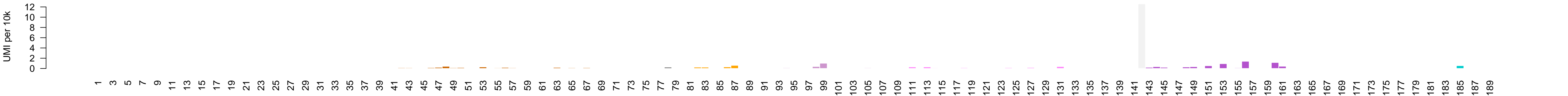
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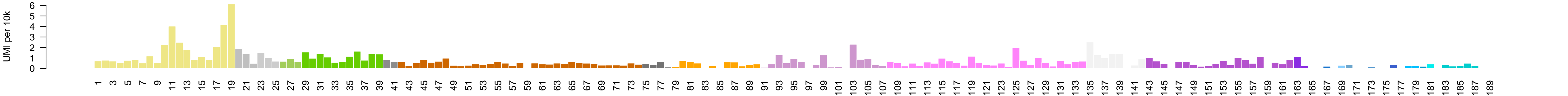
NCam.HG1.3:NCAM1/NCAM2
Tadh_TriadG59528



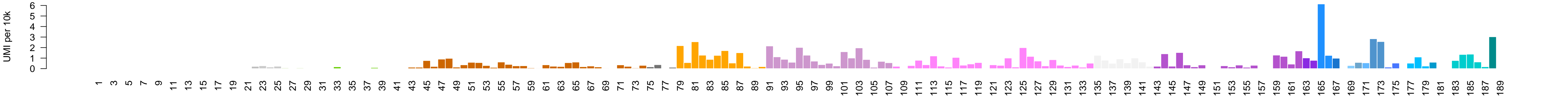
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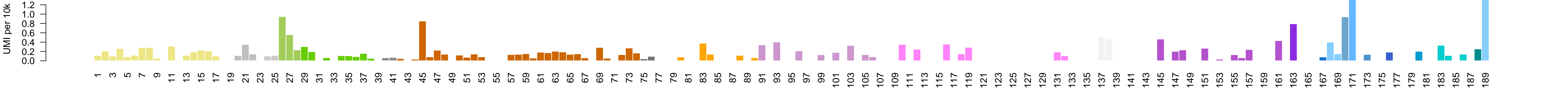
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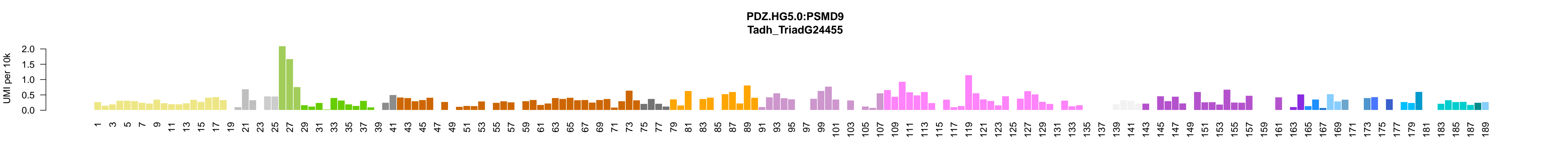


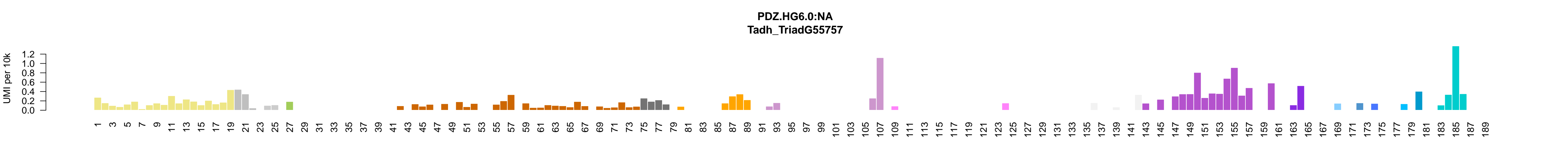
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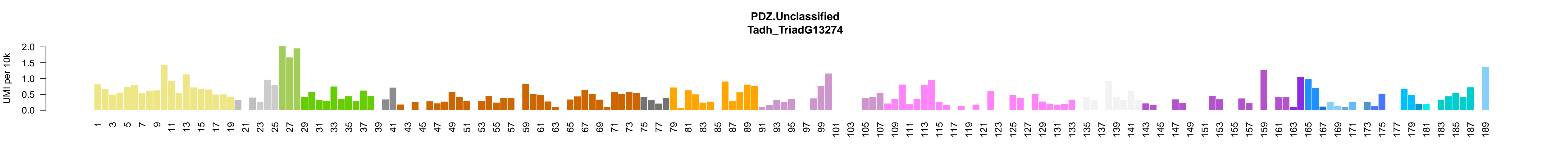


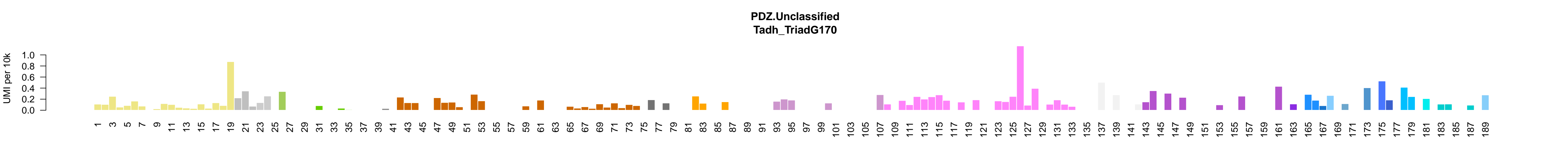
PDZ.HG4.9:like:HTRA1/HTRA2/HTRA3/HTRA4
Tadh_TriadG58565

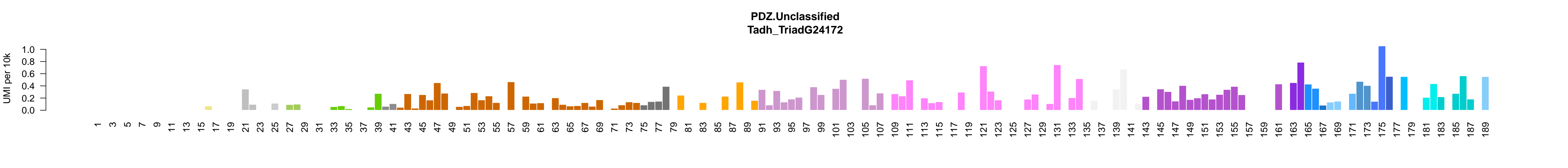


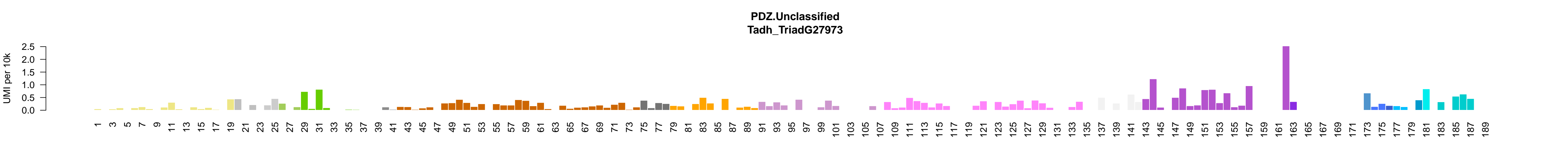










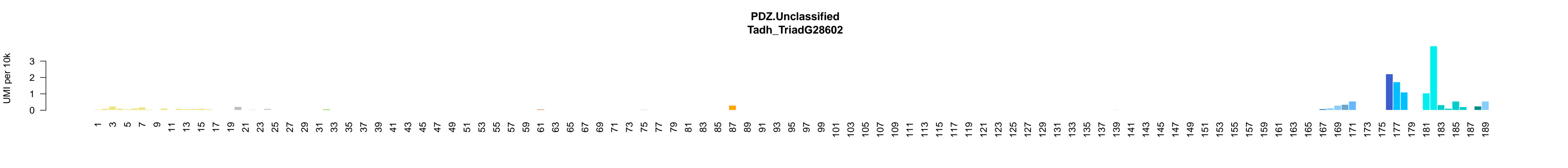


UMI per 10k

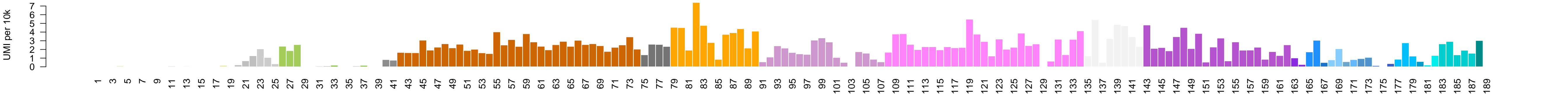
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1 3 5 7 9 11 13 15 17 19 21 23 25 27 29 31 33 35 37 39 41 43 45 47 49 51 53 55 57 59 61 63 65 67 69 71 73 75 77 79 81 83 85 87 89 91 93 95 97 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189

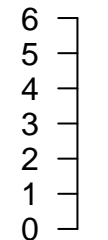
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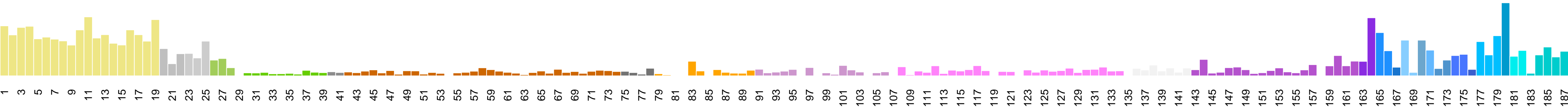
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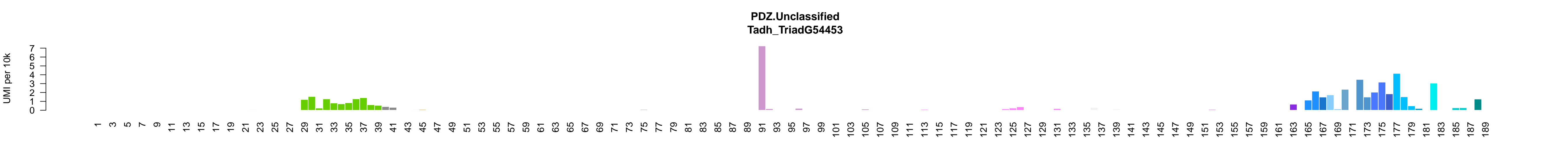


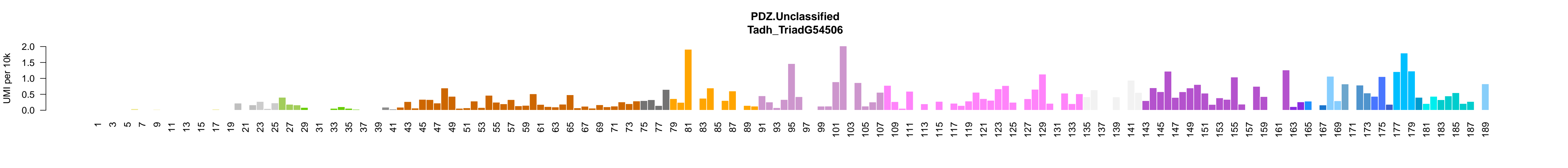
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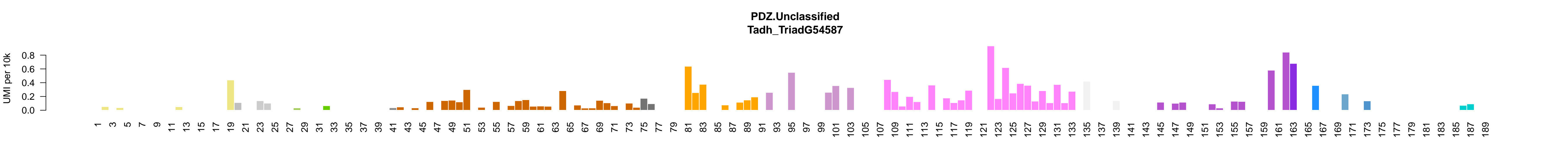


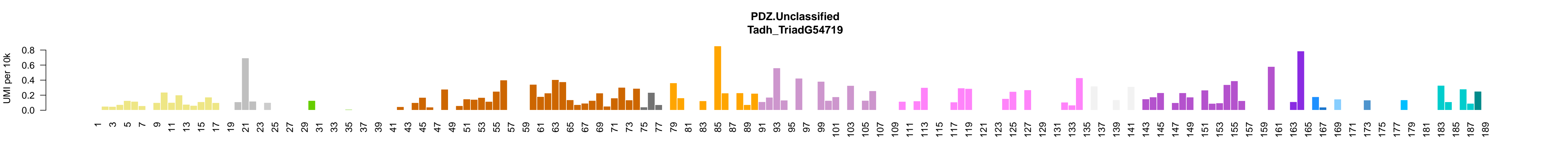
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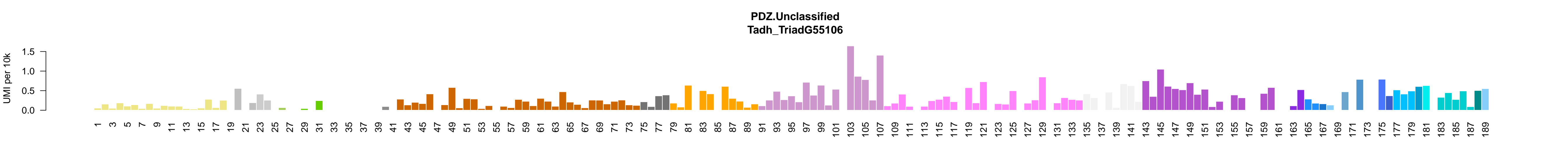




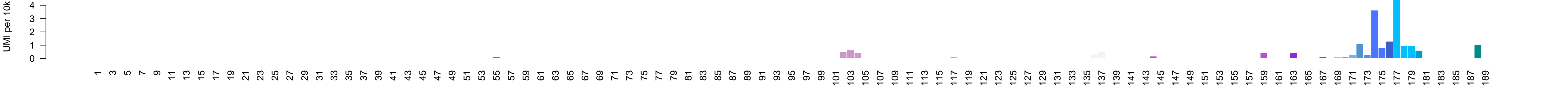


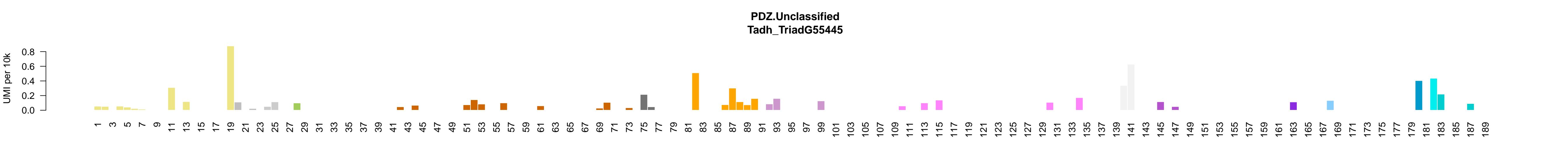


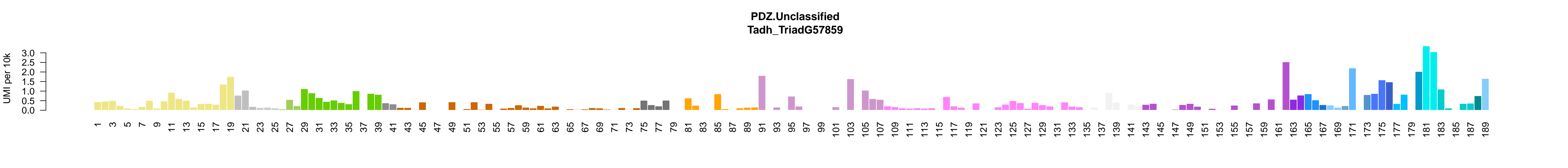


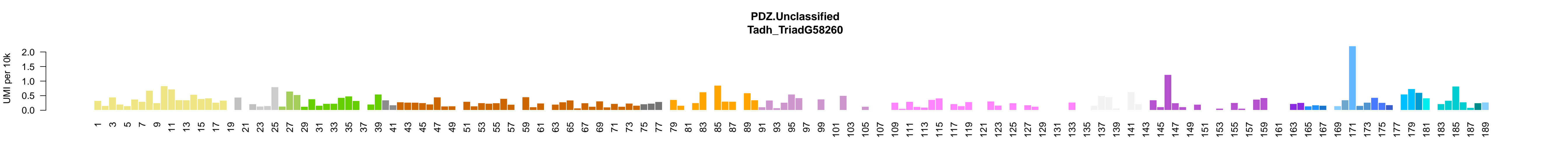


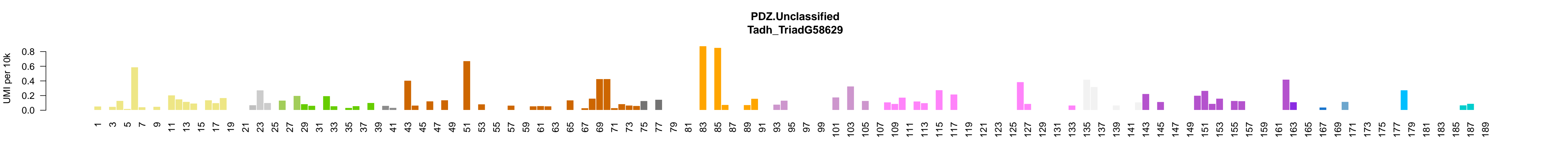
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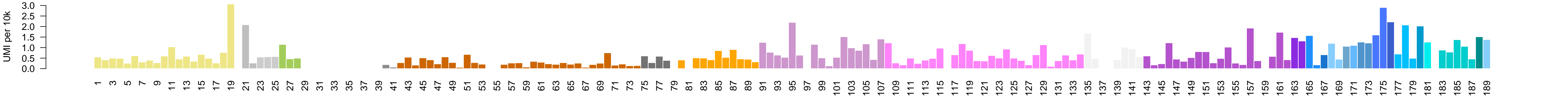


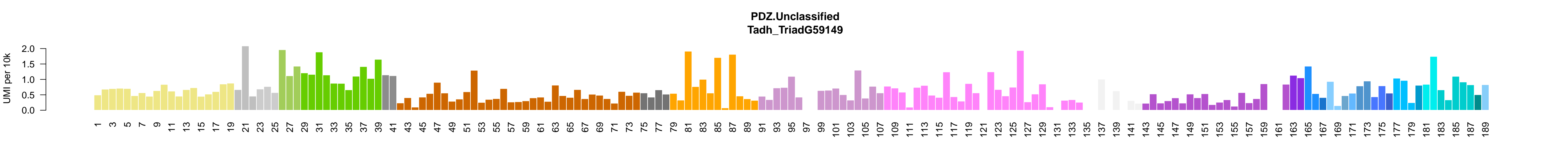






PDZ.Unclassified
Tadh_TriadG59148

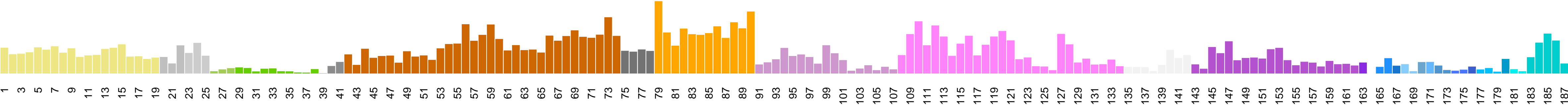




UMI per 10k

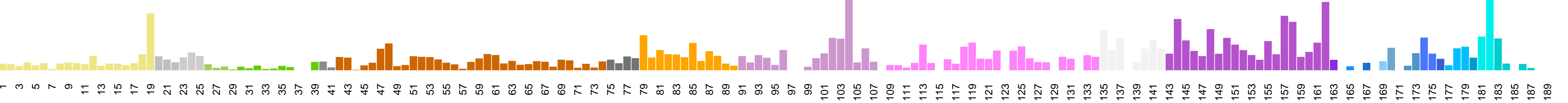
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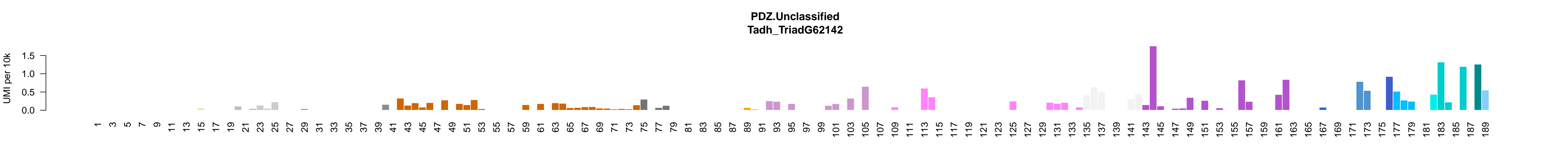
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UMI per 10k

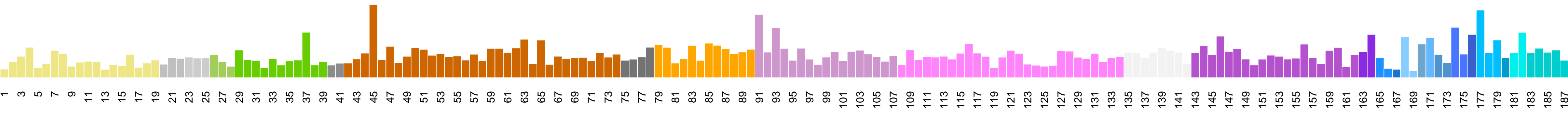
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0.5
0.0

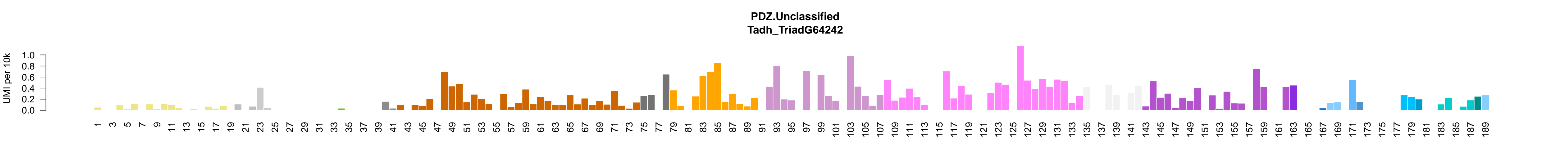




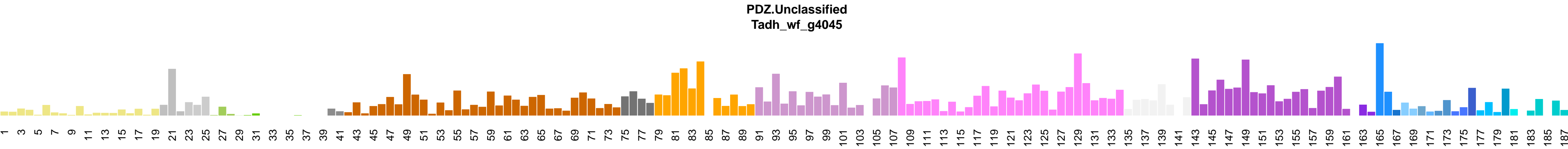
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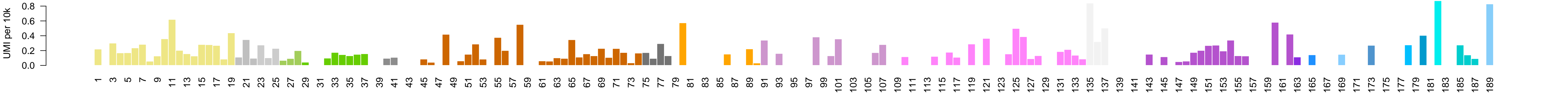
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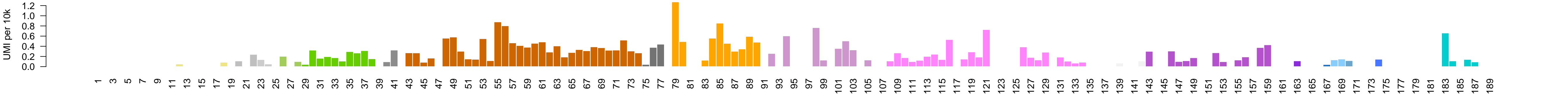


UMI per 10k

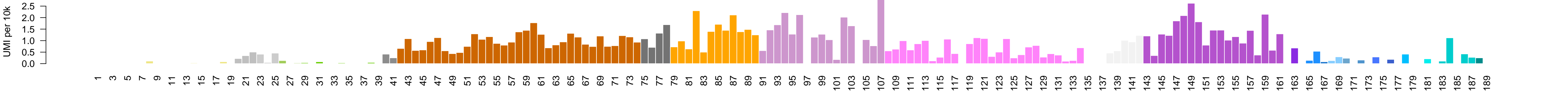




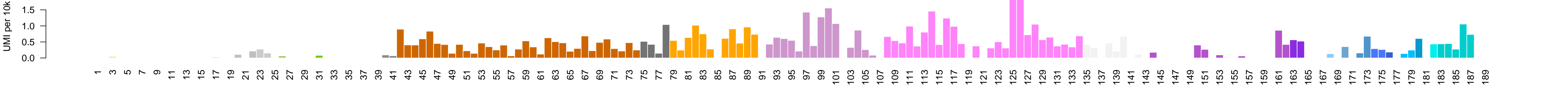
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Tadh_TriadG54315



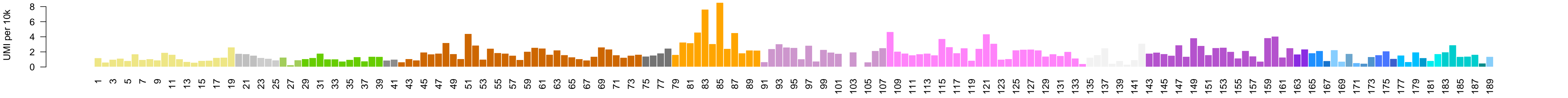
PID.HG1.21:DAB1/DAB2
Tadh_TriadG58883



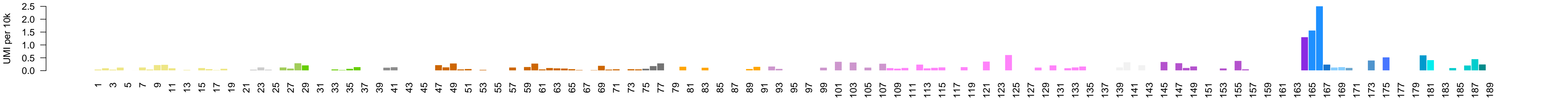
PID.HG1.8:like:LDLRAP1/PID1
Tadh_TriadG57901



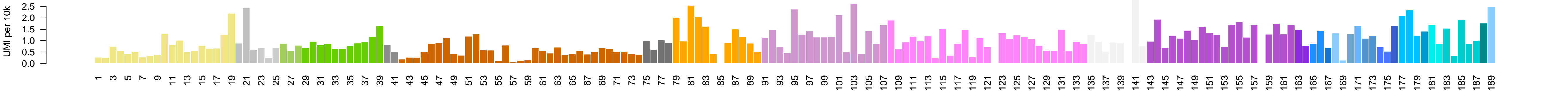
PMCA.HG1.1:ATP2A1/ATP2A2/ATP2A3
Tadh_TriadG28963



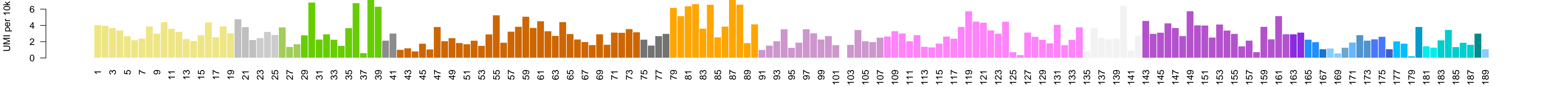
PMCA.HG1.2:ATP2C1/ATP2C2
Tadh_TriadG19369



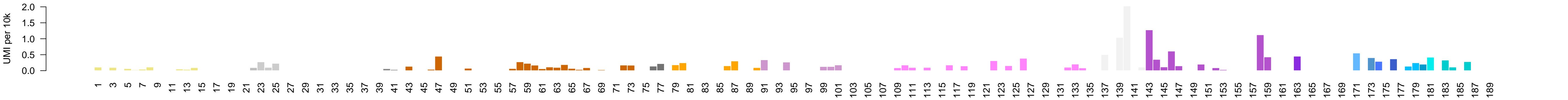
PMCA.HG1.3:ATP2B1/ATP2B2/ATP2B3/ATP2B4
Tadh_TriadG25320



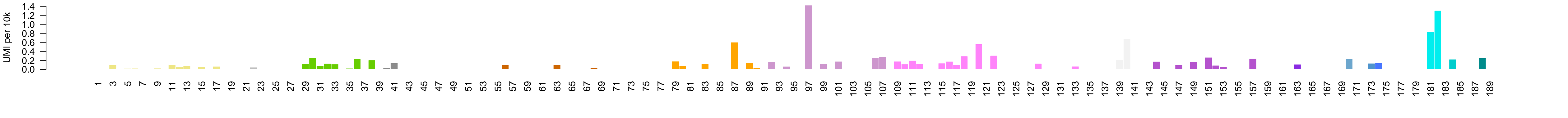
PMCA.HG1.5:AC010616.1/ATP1A1/ATP1A2/ATP1A3/ATP1A4/ATP4A/ATP12A
Tadh_TriadG63163



Pyridoxal_deC.HG1.4:CSAD/GAD1/GAD2/GADL1
Tadh_TriadG50906

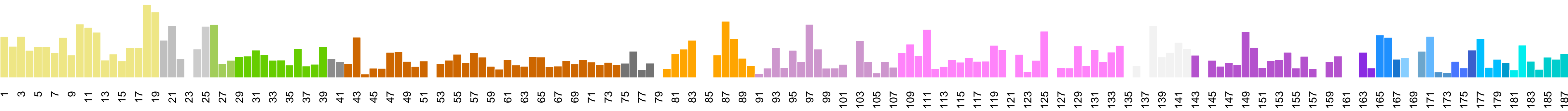
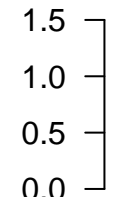


Pyridoxal_deC.HG1.4:CSAD/GAD1/GAD2/GADL1
Tadh_TriadG60611



Pyridoxal_deC.HG1.7:SGPL1
Tadh_TriadG28705

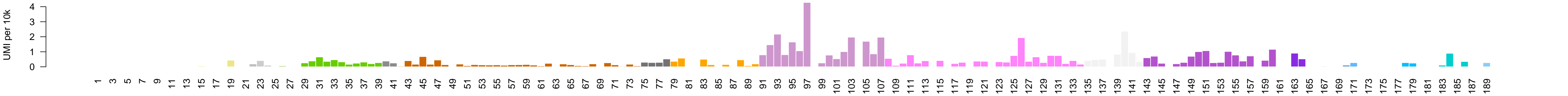
UMI per 10k



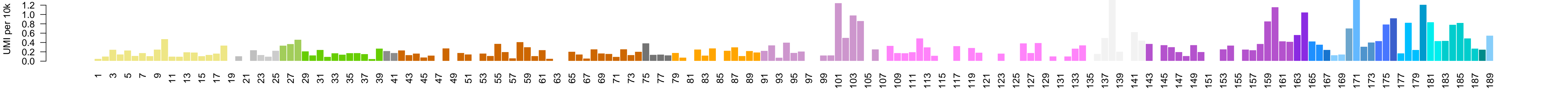
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Tadh_TriadG56436



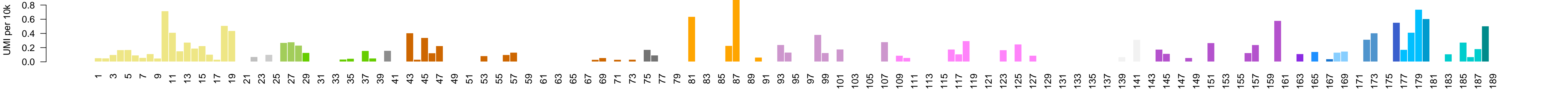
RabphilinRIM.HG1.3:RIMS1/RIMS2
Tadh_TriadG58069



RabphilinRIM.HG1.4:RPH3A/RPH3AL;Synaptotagmin.HG1.40:DOC2A/DOC2B/RPH3A
Tadh_TriadG59690



RasGAP.HG1.1:like:IQGAP1/IQGAP2/IQGAP3
Tadh_TriadG55233



UMI per 10k

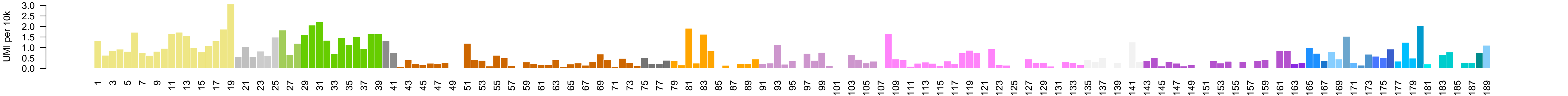
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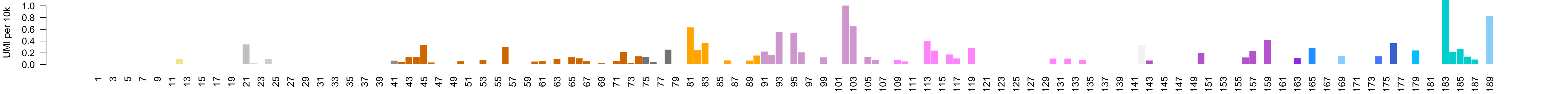
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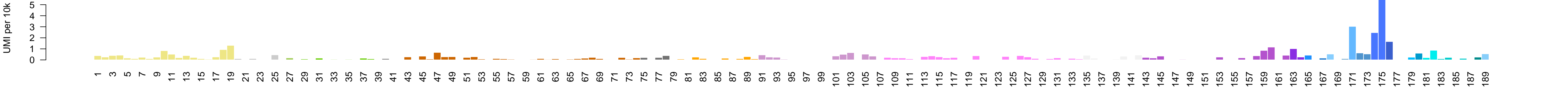
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Tadh_TriadG37304



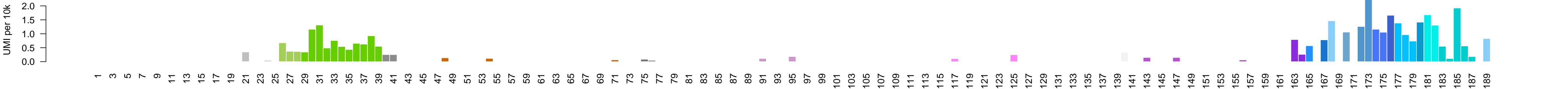
RasGAP.HG1.5:DAB2IP/RASAL2/RASAL3/SYNGAP1;SynGAP.HG1.2:like:DAB2IP/RASAL2/SYNGAP1
Tadh_wf_g1897



RasGAP.HG1.7:RASA1
Tadh_wf_g3134

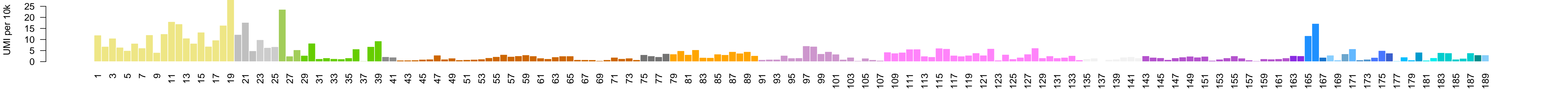


RasGAP.HG1.9:RASA2/RASA3
Tadh_TriadG63821

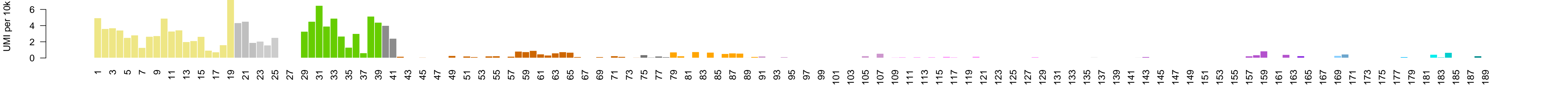


SDF.HG1.1:SLC1A1/SLC1A2/SLC1A3/SLC1A4/SLC1A5/SLC1A6/SLC1A7

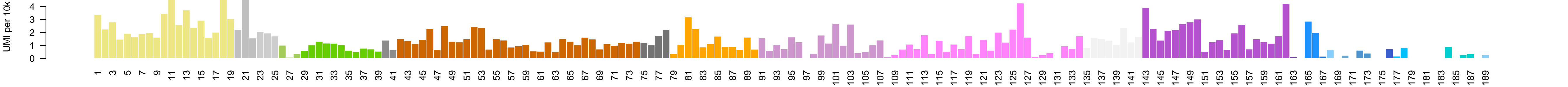
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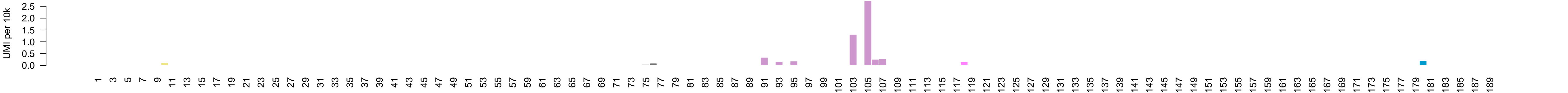
SDF.HG1.1:SLC1A1/SLC1A2/SLC1A3/SLC1A4/SLC1A5/SLC1A6/SLC1A7
Tadh_TriadG50710



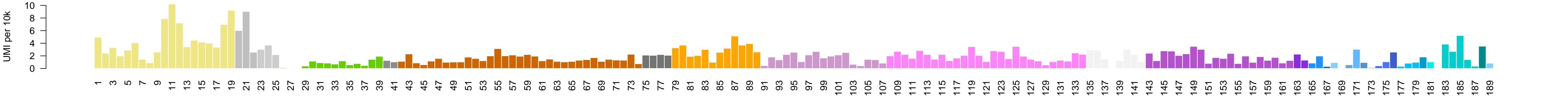
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Tadh_TriadG33521



SNF.HG1.17:like:SLC6A5/SLC6A7/SLC6A9/SLC6A14
Tadh_TriadG31455



SNF.HG1.2:like:AC136616.1/SLC6A1/SLC6A2/SLC6A3/SLC6A4/SLC6A5/SLC6A6/SLC6A7/SLC6A8/SLC6A9/SLC6A11/SLC6A12/SLC6A13/SLC6A14/SLC6A15/SLC6A16/SLC6A17/SLC6A18/SLC6A19/SLC6A20
Tadh_TriadG27813

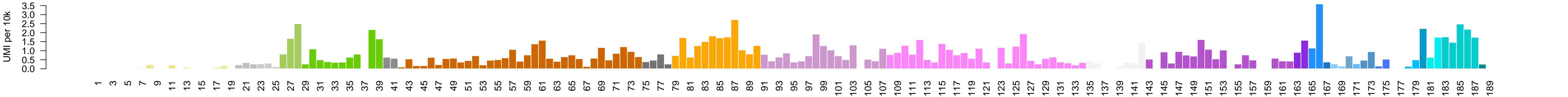


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Tadh_TriadG29105

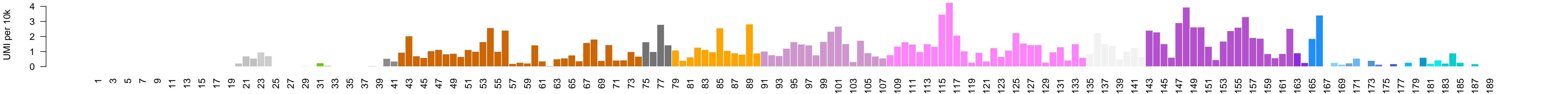


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Tadh_TriadG29421



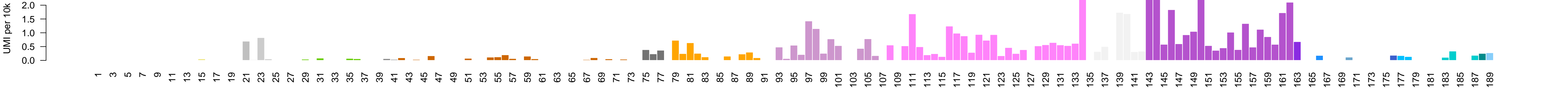
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Tadh_TriadG33157

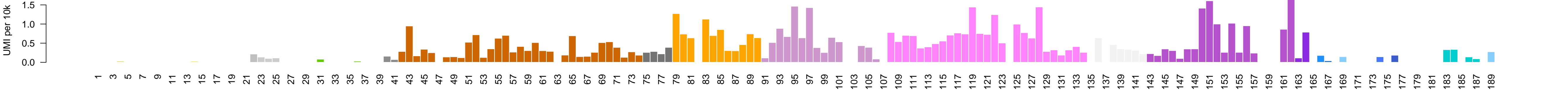


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Tadh_TriadG34381

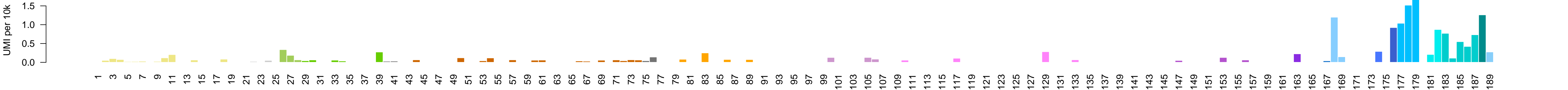


SNF.HG1.2:like:AC136616.1/SLC6A1/SLC6A2/SLC6A3/SLC6A4/SLC6A5/SLC6A6/SLC6A7/SLC6A8/SLC6A9/SLC6A11/SLC6A12/SLC6A13/SLC6A14/SLC6A15/SLC6A16/SLC6A17/SLC6A18/SLC6A19/SLC6A20
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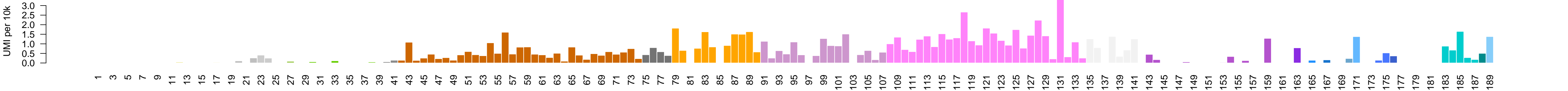
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Tadh_TriadG59303

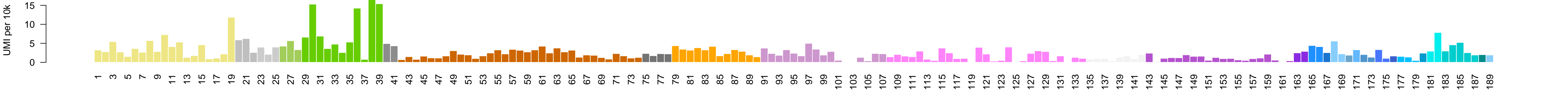


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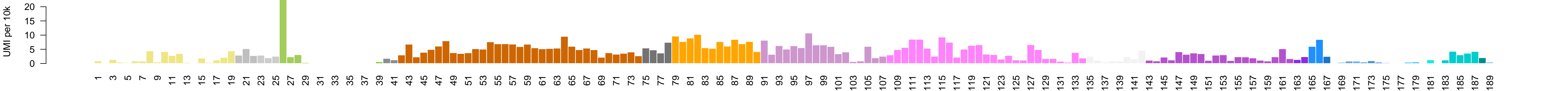
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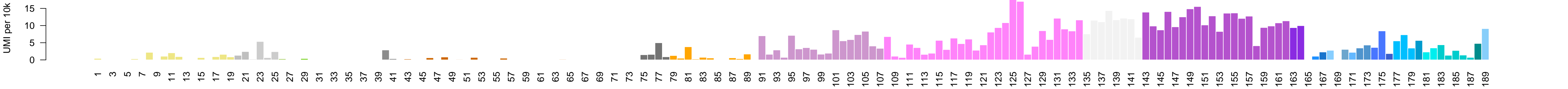
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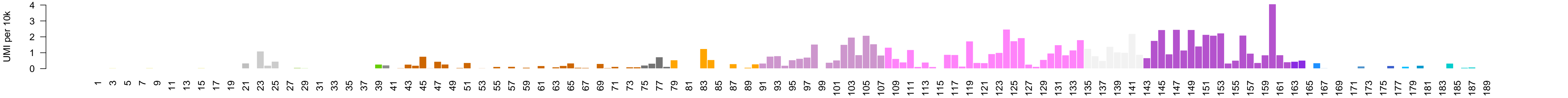
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Tadh_TriadG50950



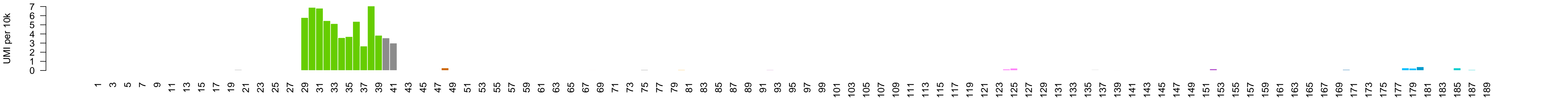
SNF.HG1.31:like:SLC6A5/SLC6A7/SLC6A9/SLC6A14
Tadh_TriadG38332



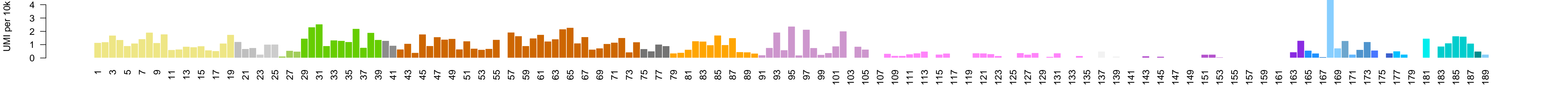
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Tadh_TriadG23982



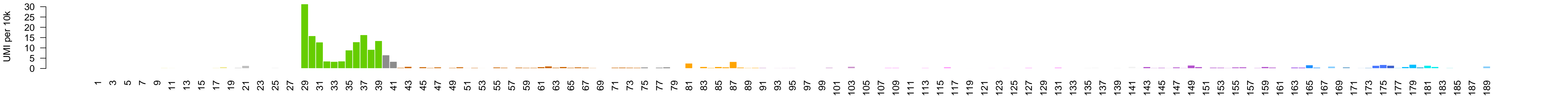
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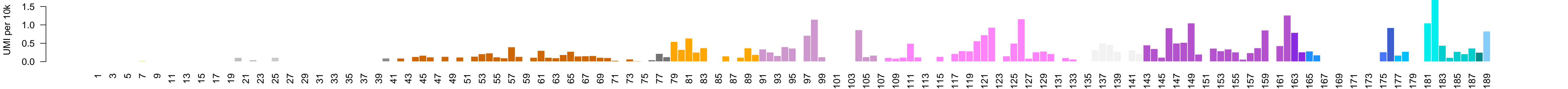
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Tadh_TriadG37600



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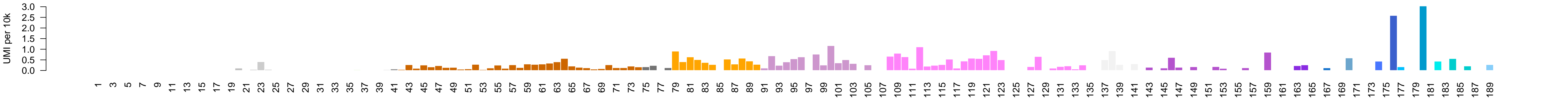


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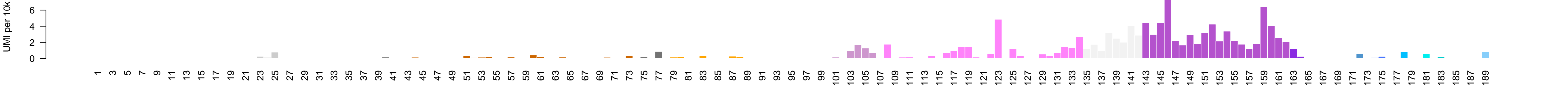
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Tadh_TriadG33106

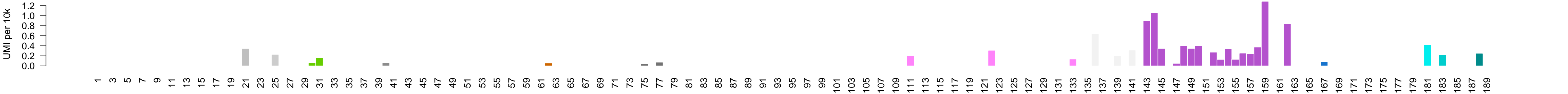


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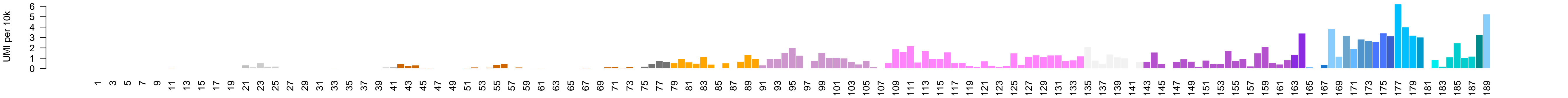
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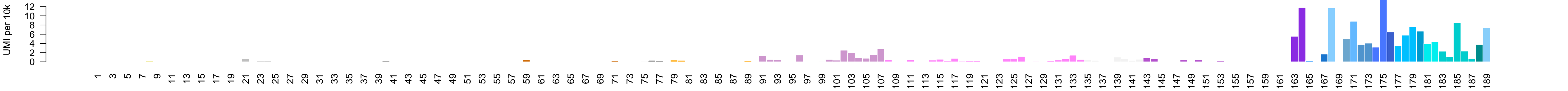
Sugar_tr.HG2.59:SVOP
Tadh_TriadG62146



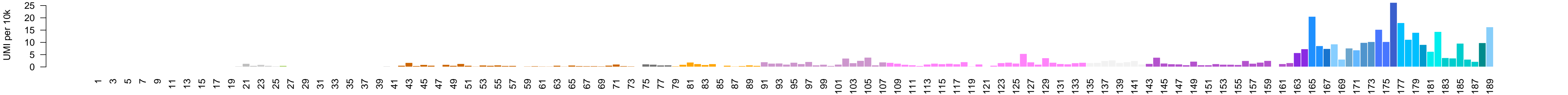
Synaphin.HG2.0:NA
Tadh_wf_g9757



Synaptotagmin.HG1.0:SYT7
Tadh_TriadG32942



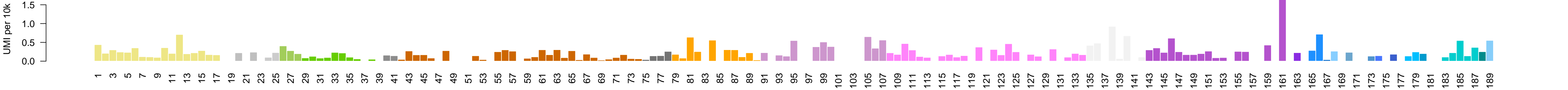
Synaptotagmin.HG1.2:SYT1/SYT2/SYT5/SYT8
Tadh_TriadG51110



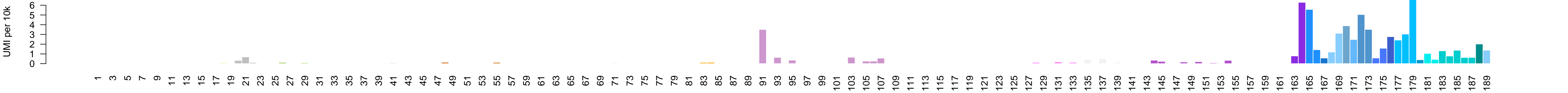
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Tadh_TriadG57029



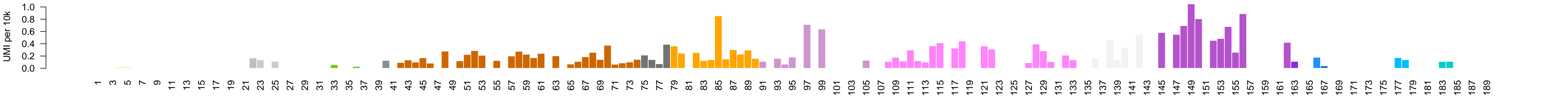
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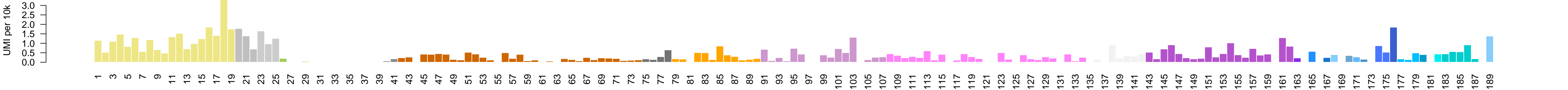
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Tadh_TriadG60504



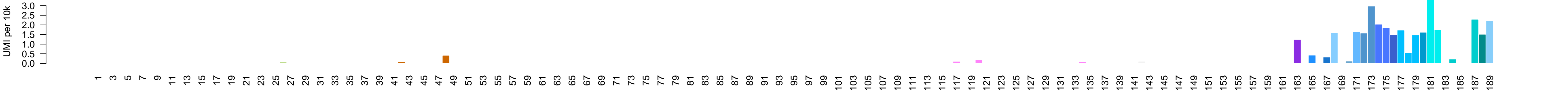
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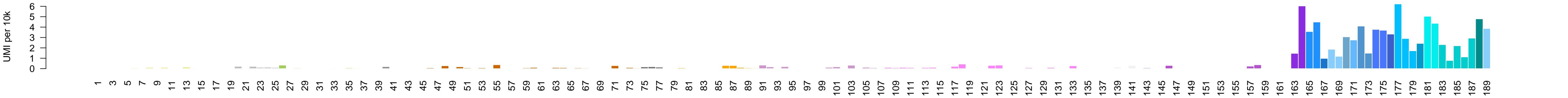
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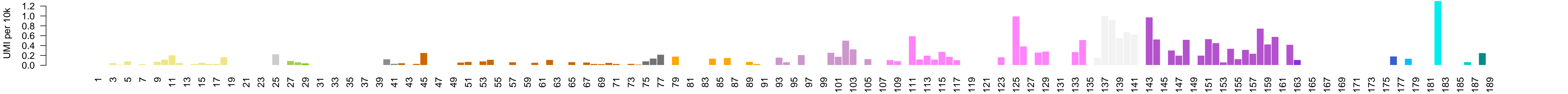
Synaptotagmin.HG1.45:like:SYTL1/SYTL2/SYTL3/SYTL4/SYTL5
Tadh_TriadG56437



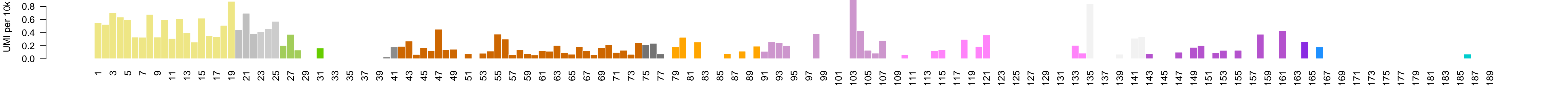
Synaptotagmin.HG1.49:ESYT1/ESYT2/ESYT3
Tadh_TriadG61664



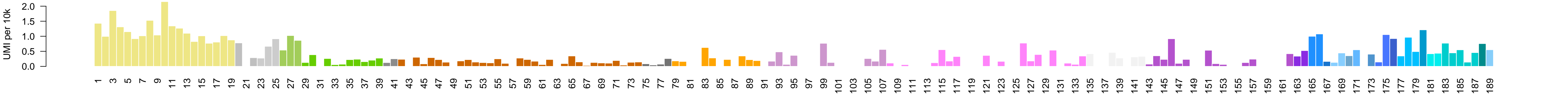
Synaptotagmin.HG1.5:like:FO681492.1/SYT12/SYT13/SYT15
Tadh_TriadG53128



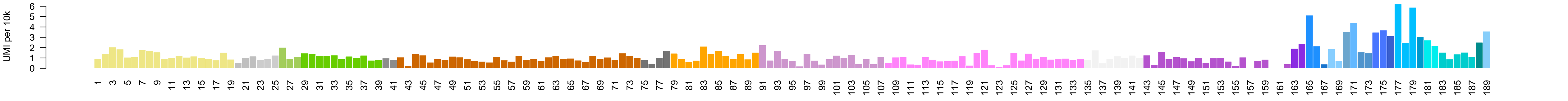
Synaptotagmin.HG1.60:like:ITSN2/UNC13A/UNC13C
Tadh_TriadG51955



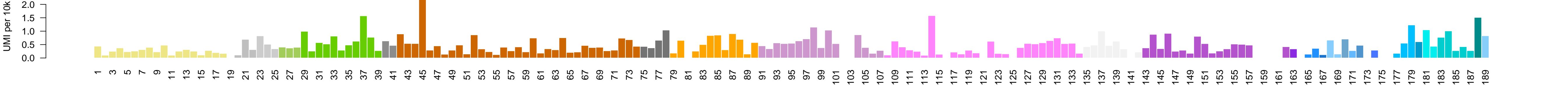
Syntaxin.HG1.1:STX16/STX16-NPEPL1
Tadh_TriadG52504



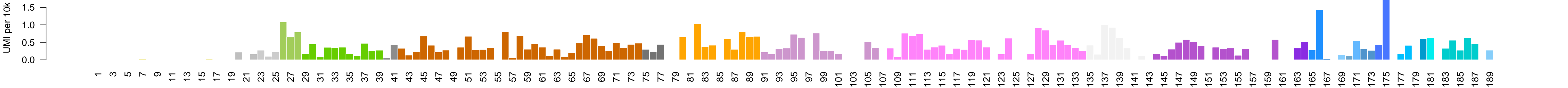
Syntaxin.HG1.2:STX7/STX12
Tadh_TriadG27557



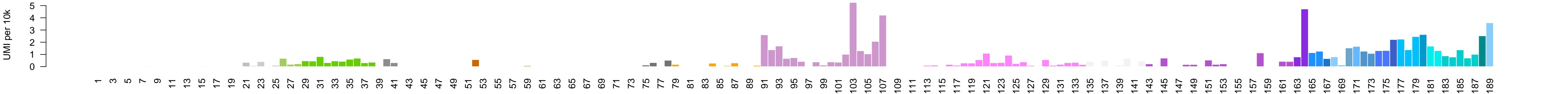
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Tadh_TriadG58273



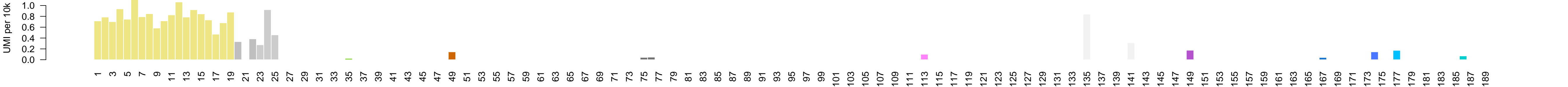
Syntaxin.HG1.4:STX1A/STX1B/STX2/STX3/STX4
Tadh_TriadG26532



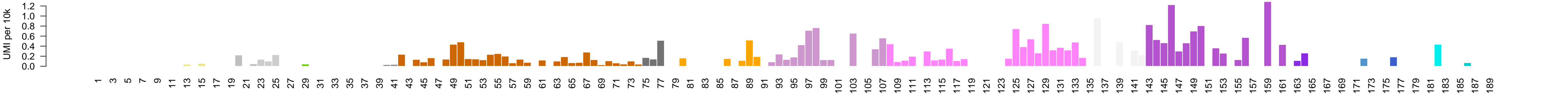
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Tadh_TriadG28648



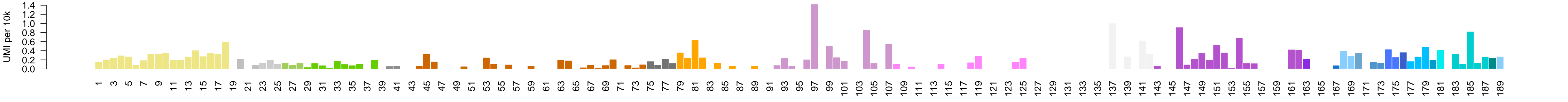
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Tadh_TriadG28962



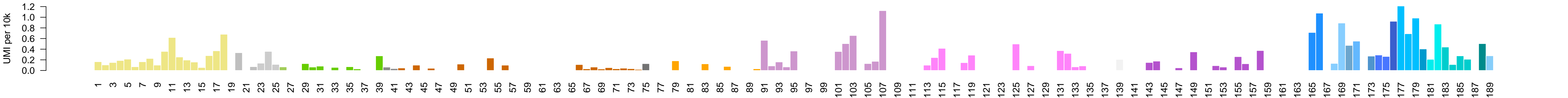
Syntaxin.HG1.4:STX1A/STX1B/STX2/STX3/STX4
Tadh_TriadG59128



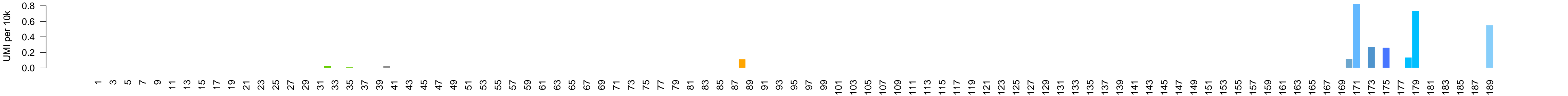
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Tadh_wf_g6284



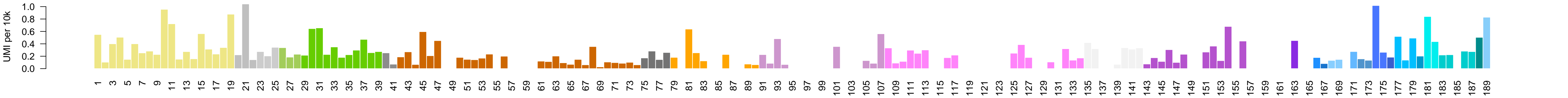
Y_phosphatase.HG1.14:PTPN5/PTPN7/PTPRR
Tadh_TriadG51484



Y_phosphatase.HG1.16:like:PTPRK/PTPRM/PTPRT/PTPRU
Tadh_wf_g1014

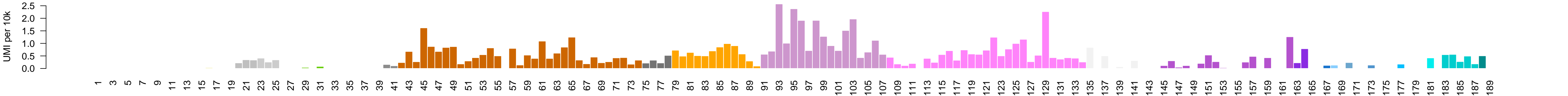


Y_phosphatase.HG1.20:PTPN23
Tadh_TriadG60390

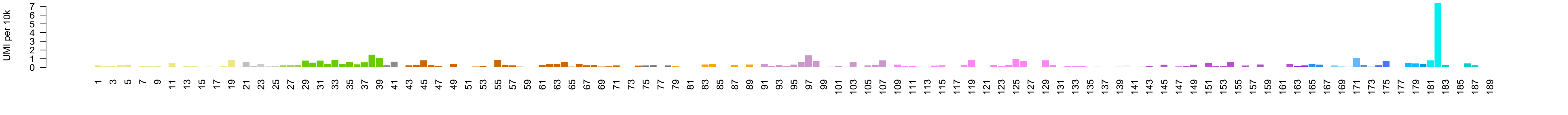


Y_phosphatase.HG1.24:like:PTPRA/PTPRC/PTPRD/PTPRE/PTPRF/PTPRG/PTPRS/PTPRZ1

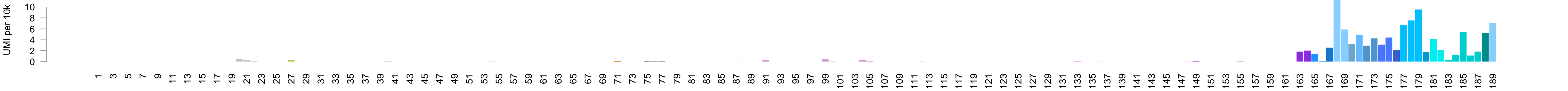
Tadh_TriadG27501



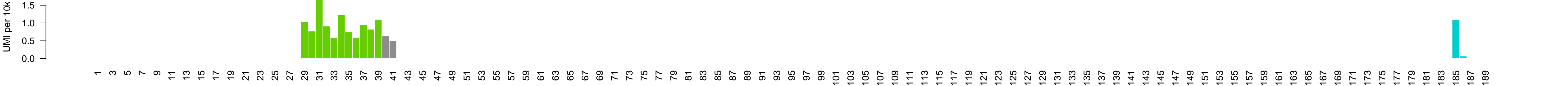
Y_phosphatase.HG1.26:like:PTPN9
Tadh_TriadG54170



Y_phosphatase.HG1.29:PTPRN/PTPRN2
Tadh_TriadG55048



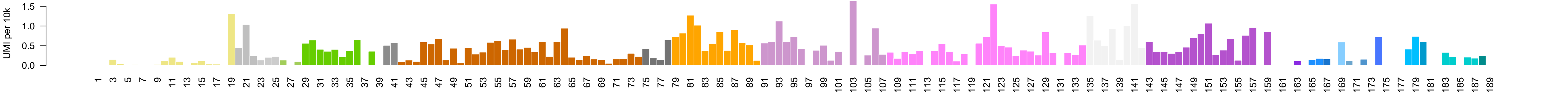
Y_phosphatase.HG1.38:PTPN3/PTPN4
Tadh_TriadG55016

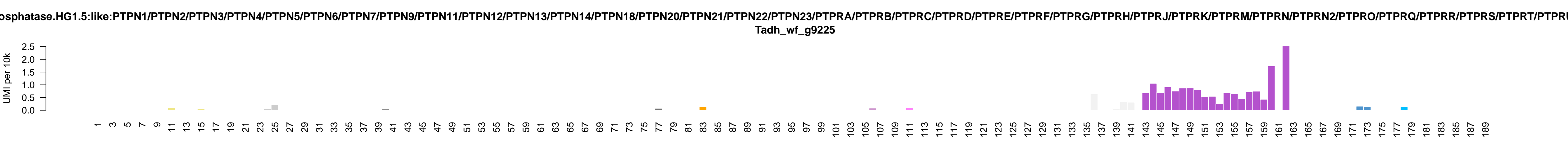


Y_phosphatase.HG1.41:like:PTPN5/PTPN7/PTPRR
Tadh_TriadG60483

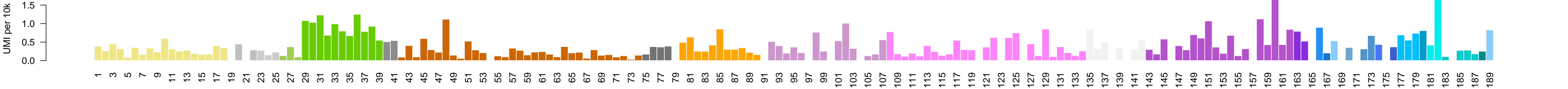


Y_phosphatase.HG1.45:like:PTPRA/PTPRC/PTPRD/PTPRE/PTPRF/PTPRG/PTPRS/PTPRZ1
Tadh_TriadG27423

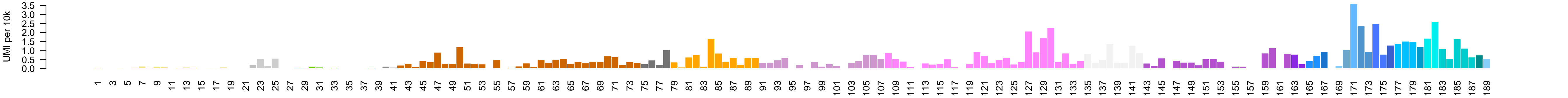




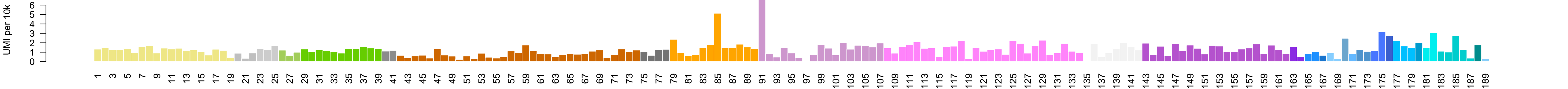
Y_phosphatase.HG1.52:PTPRG/PTPRZ1
Tadh_TriadG7563



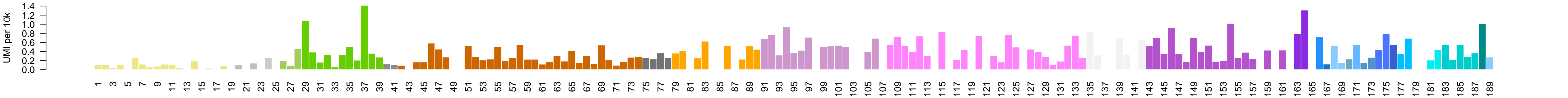
Y_phosphatase.HG2.6:PTPDC1
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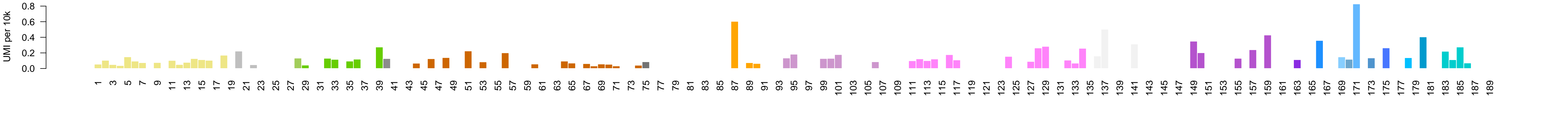
Y_phosphatase.HG2.8:PTP4A1/PTP4A2/PTP4A3
Tadh_TriadG27879



Y_phosphatase.HG3.4:like:DUSP1/DUSP4/DUSP5
Tadh_TriadG60150



Y_phosphatase.HG3.4:like:DUSP1/DUSP4/DUSP5
Tadh_TriadG9723



Y_phosphatase.HG6.0:NA
Tadh_TriadG64052

