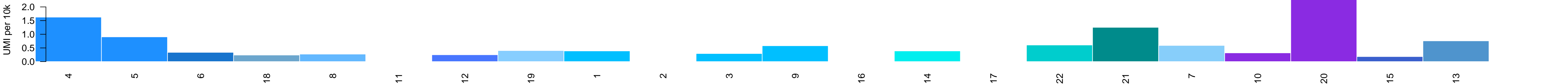
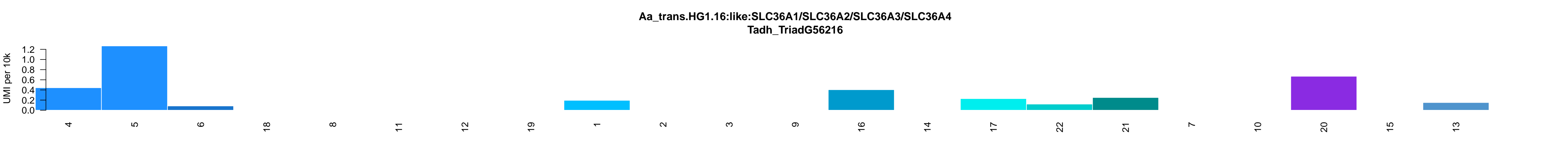


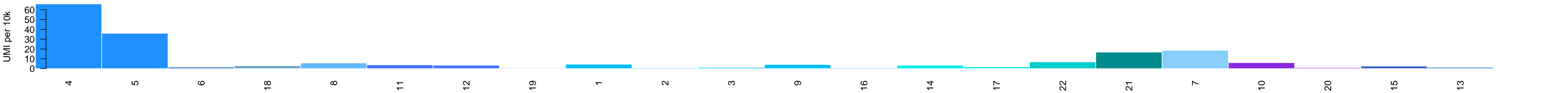
Aa_trans.HG1.12:SLC36A1/SLC36A2/SLC36A3/SLC36A4

Tadh_TriadG56468





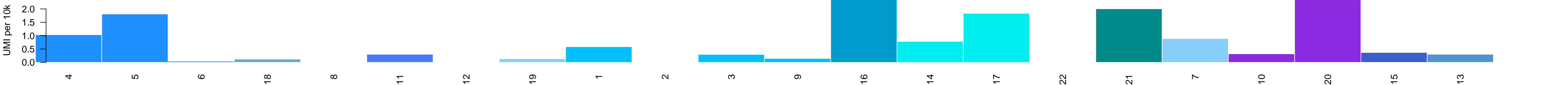
Aa_trans.HG1.2:SLC38A9
Tadh_TriadG29059



Aa_trans.HG1.2:SLC38A9
Tadh_TriadG29062



Aa_trans.HG1.2:SLC38A9
Tadh_TriadG29287



Aa_trans.HG1.4:SLC38A11
Tadh_TriadG22089

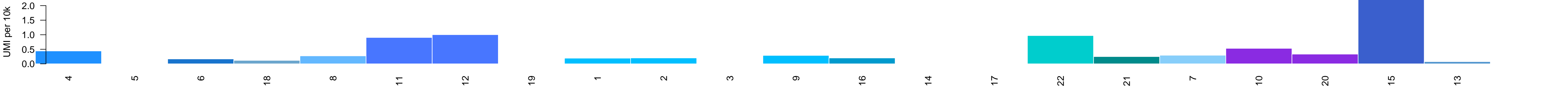
UMI per 10k

1.5
1.0
0.5
0.0

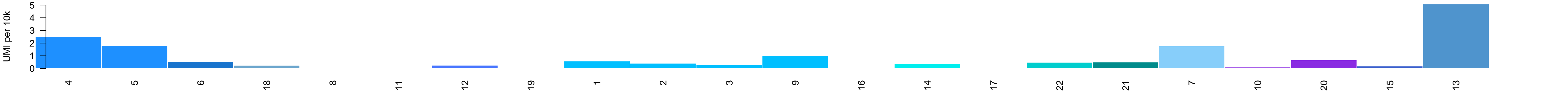
4 5 6 18 8 11 12 19 1 2 3 9 16 14 17 22 21 7 10 20 15 13



Aa_trans.HG1.4:SLC38A11
Tadh_TriadG22445



Aa_trans.HG1.4:SLC38A11
Tadh_TriadG23041



ANF_receptor.HG1.18:like:GRM1/GRM2/GRM3/GRM4/GRM5/GRM6/GRM7/GRM8
Tadh_TriadG11098

UMI per 10k

1.5
1.0
0.5
0.0



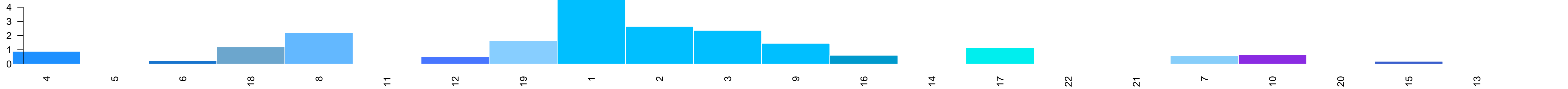
ANF_receptor.HG1.18:like:GRM1/GRM2/GRM3/GRM4/GRM5/GRM6/GRM7/GRM8
Tadh_TriadG25549





ANF_receptor.HG1.37:like:CASR/GPRC6A/TAS1R1/TAS1R2/TAS1R3
Tadh_wf_g964

UMI per 10k



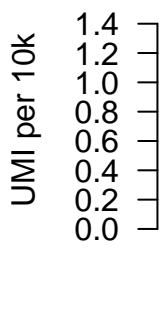
ANF_receptor.HG1.48:like:CASR/GPRC6A/TAS1R1/TAS1R2/TAS1R3
Tadh_TriadG25328



ANF_receptor.HG1.51:like:CASR/GPRC6A/TAS1R1/TAS1R2/TAS1R3

Tadh_TriadG29555

UMI per 10k



ANF_receptor.HG1.52:like:CASR/GPRC6A/TAS1R1/TAS1R2/TAS1R3
Tadh_wf_g3864



ANF_receptor.HG1.61:like:CASR/GPRC6A/TAS1R1/TAS1R2/TAS1R3
Tadh_TriadG29561

UMI per 10k

1.5
1.0
0.5
0.0

4 5 6 18 8 11 12 19 1 2 3 9 16 14 17 22 21 7 10 20 15 13



ANF_receptor.HG1.63:like:CASR/GPRC6A/TAS1R1/TAS1R2/TAS1R3
Tadh_TriadG29631



ANF_receptor.HG1.64:like:CASR/GPRC6A/TAS1R1/TAS1R2/TAS1R3
Tadh_TriadG59554

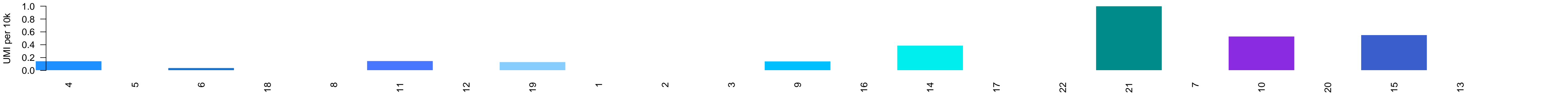
UMI per 10k

1.2
1.0
0.8
0.6
0.4
0.2
0.0

4 5 6 18 8 11 12 19 1 2 3 9 16 14 17 22 21 7 10 20 15 13

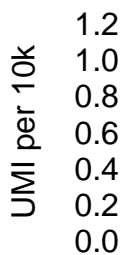


ANF_receptor.HG2.10:like:GABBR1/GABBR2
Tadh_TriadG26726

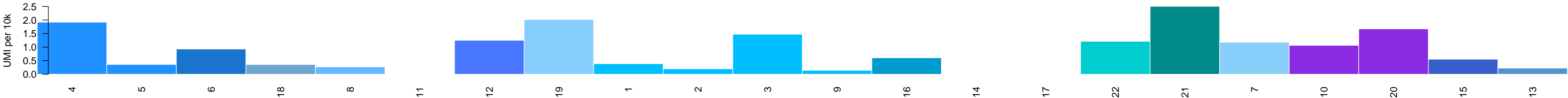


ANF_receptor.HG2.10:like:GABBR1/GABBR2
Tadh_TriadG52983

UMI per 10k



ANF_receptor.HG2.10:like:GABBR1/GABBR2
Tadh_TriadG57193



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



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

UMI per 10k

3.0
2.5
2.0
1.5
1.0
0.5
0.0

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

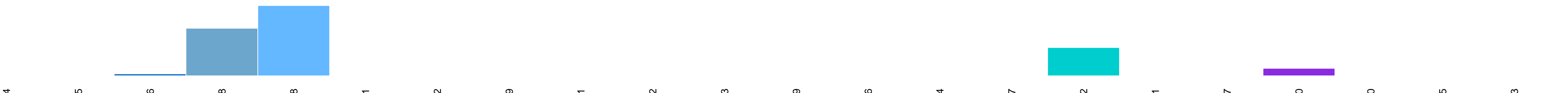
7

10

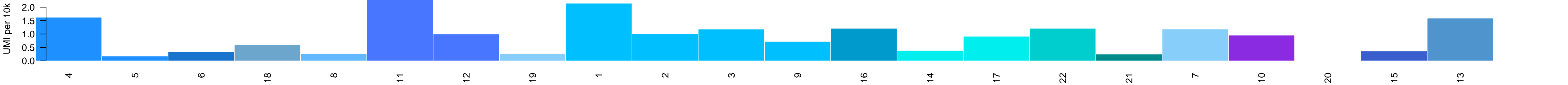
20

15

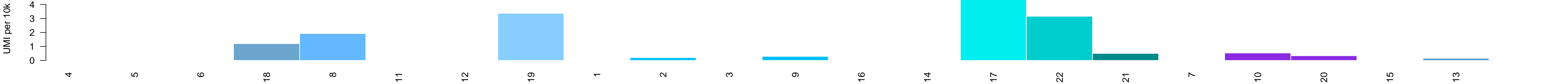
13



ANF_receptor.HG2.14:like:GABBR1/GABBR2
Tadh_TriadG61926



ANF_receptor.HG2.16:like:GABBR1/GABBR2
Tadh_TriadG28568



ANF_receptor.HG2.17:like:GABBR1/GABBR2
Tadh_TriadG28937

UMI per 10k

6
5
4
3
2
1
0

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

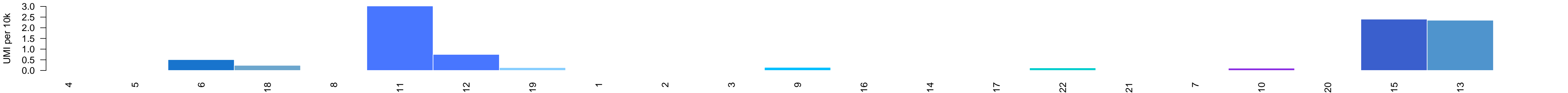
13

4

5

6

ANF_receptor.HG2.18:like:GABBR1/GABBR2
Tadh_TriadG59025



ANF_receptor.HG2.19:like:GABBR1/GABBR2
Tadh_TriadG59024

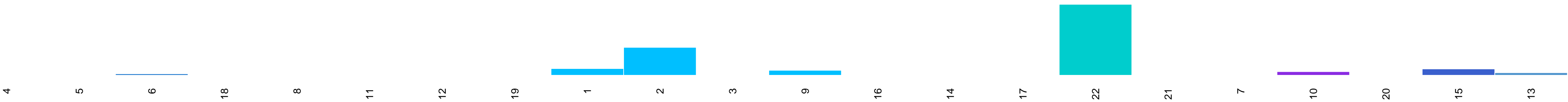


ANF_receptor.HG2.23:like:GABBR1/GABBR2
Tadh_TriadG54955

UMI per 10k

2.0
1.5
1.0
0.5
0.0

4 5 6 18 8 11 12 19 1 2 3 9 16 14 17 22 21 7 10 20 15 13



ANF_receptor.HG2.24:like:GABBR1/GABBR2
Tadh_TriadG54957



ANF_receptor.HG2.24:like:GABBR1/GABBR2
Tadh_wf_g11837

UMI per 10k

1.2
1.0
0.8
0.6
0.4
0.2
0.0

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

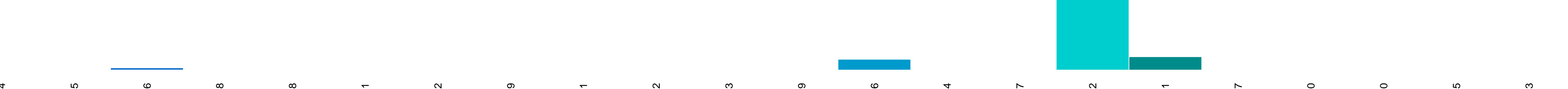
7

10

20

15

13



ANF_receptor.HG2.3:like:GABBR1/GABBR2
Tadh_TriadG30267

UMI per 10k

30
20
10
0

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

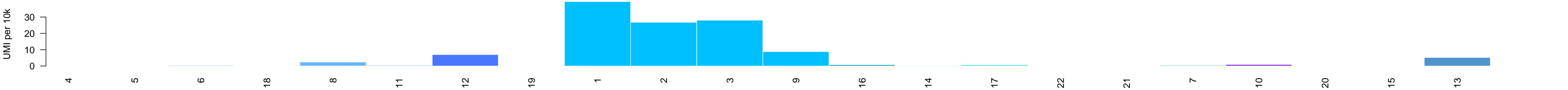
7

10

20

15

13



ANF_receptor.HG2.3:like:GABBR1/GABBR2
Tadh_TriadG55385

UMI per 10k

2.0
1.5
1.0
0.5
0.0

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

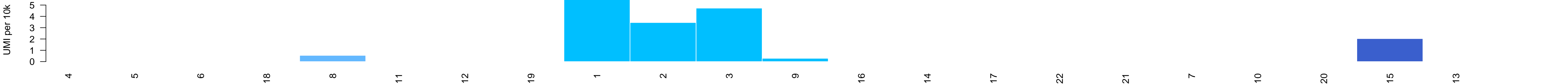
20

15

13



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



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

UMI per 10k

20
15
10
5
0

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

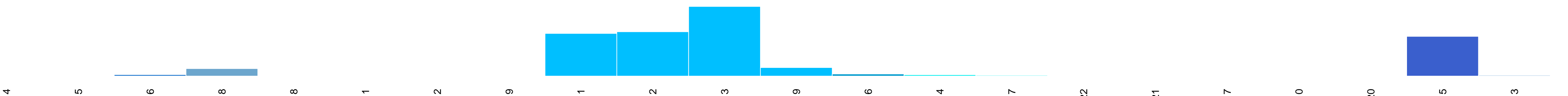
7

10

20

15

13



ANF_receptor.HG2.3:like:GABBR1/GABBR2
Tadh_TriadG60050

UMI per 10k

10
8
6
4
2
0

4 5 6 18 8 11 12 19 1 2 3 9 16 14 17 22 21 7 10 20 15 13



ANF_receptor.HG2.31:like:GABBR1/GABBR2
Tadh_TriadG58939

UMI per 10k

3
2
1
0

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

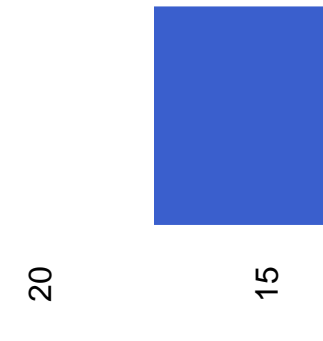
7

10

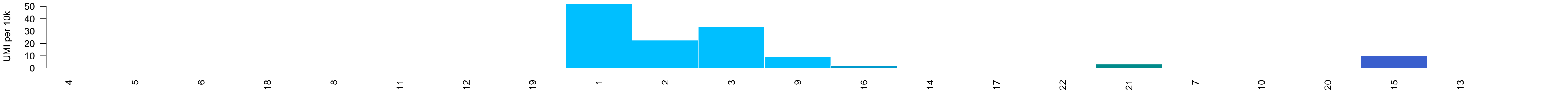
20

15

13



ANF_receptor.HG2.32:like:GABBR1/GABBR2
Tadh_TriadG58938



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



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

UMI per 10k

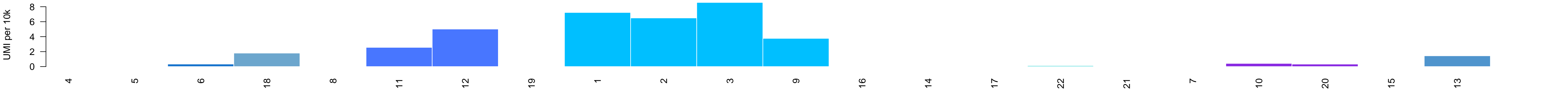
35
30
25
20
15
10
5
0



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



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



ANF_receptor.HG2.4:like:GABBR1/GABBR2
Tadh_TriadG55429

UMI per 10k

4
3
2
1
0

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

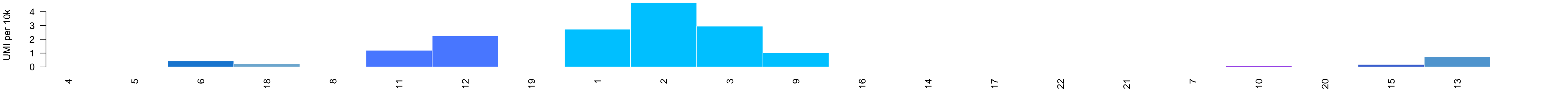
7

10

20

15

13



ANF_receptor.HG2.5:like:GABBR1/GABBR2
Tadh_TriadG52576

UMI per 10k

1.0
0.8
0.6
0.4
0.2
0.0

4 5 6 18 8 11 12 19 1 2 3 9 16 14 17 22 21 7 10 20 15 13



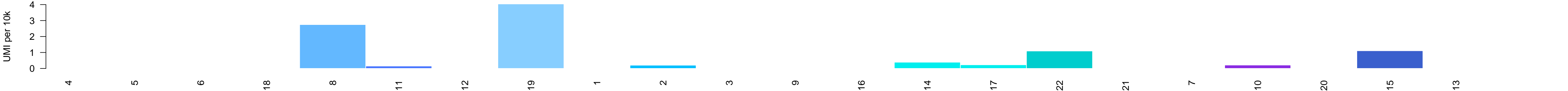
ANF_receptor.HG2.5:like:GABBR1/GABBR2
Tadh_TriadG52577



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



ANF_receptor.HG2.5:like:GABBR1/GABBR2
Tadh_TriadG52579



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

UMI per 10k

2.5
2.0
1.5
1.0
0.5
0.0

4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

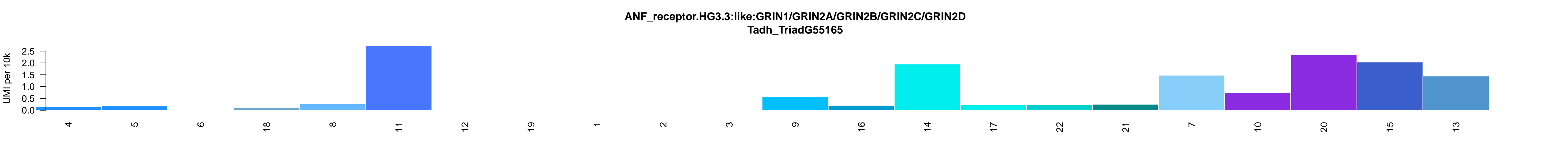
10

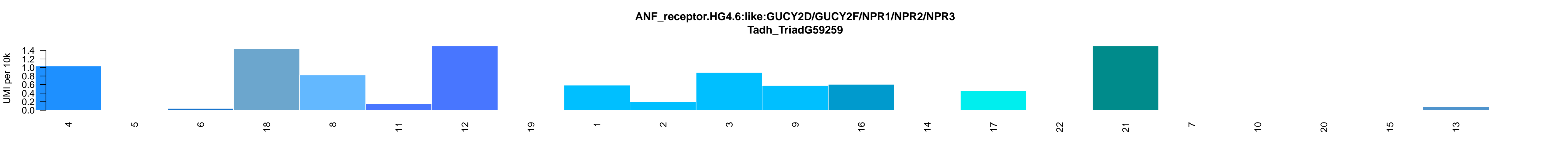
20

15

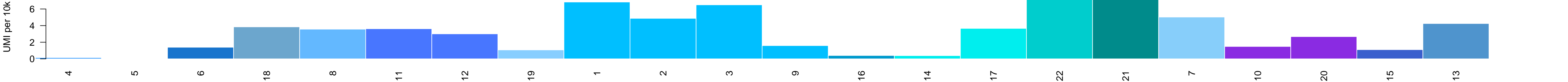
13



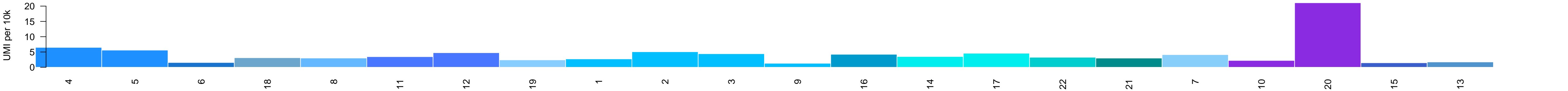




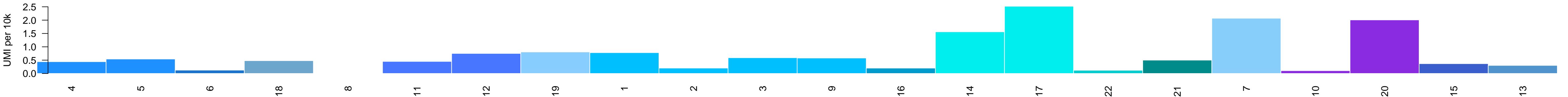
ANF_receptor.HG4.9:like:GUCY2D/GUCY2F/NPR1/NPR2/NPR3
Tadh_TriadG30655



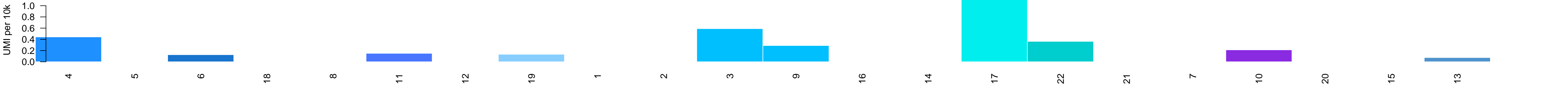
CaMKII.HG1.1:CAMK2A/CAMK2B/CAMK2D/CAMK2G
Tadh_TriadG50756



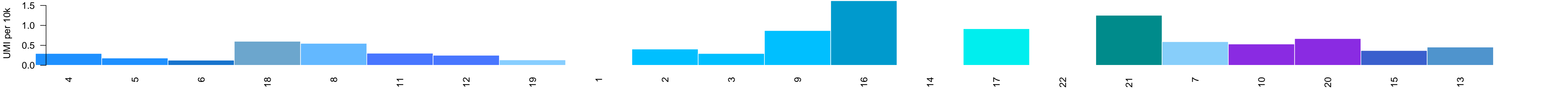
Caps.HG1.1:UNC13A/UNC13B/UNC13C;Synaptotagmin.HG1.66:UNC13A/UNC13C;Unc13.HG1.1:UNC13A/UNC13B/UNC13C
Tadh_TriadG22095



Citron.HG1.6:VPS39
Tadh_TriadG61902

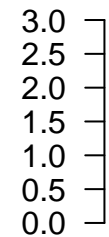


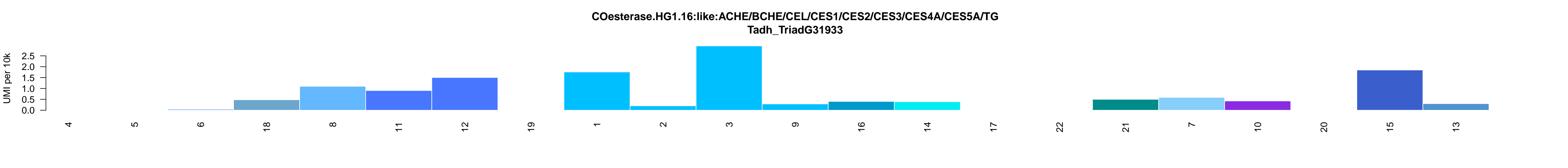
Citron.HG2.0:MAP4K1/MAP4K2/MAP4K3/MAP4K5
Tadh_TriadG51772



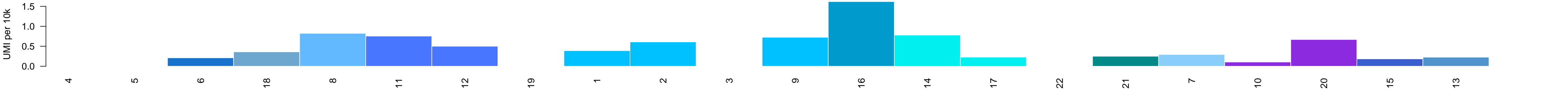
COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG31923

UMI per 10k

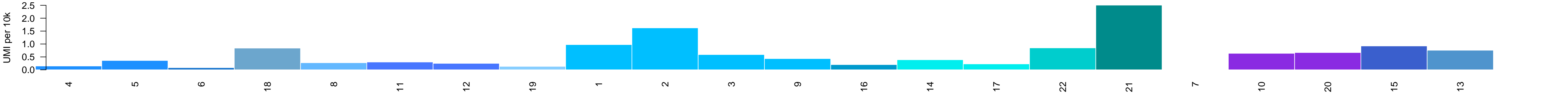




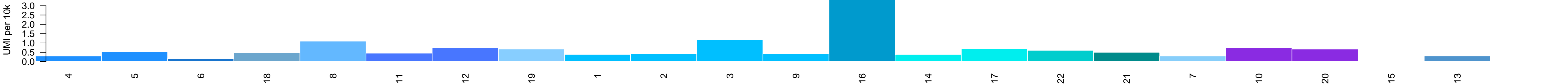
COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Tadh_TriadG61120



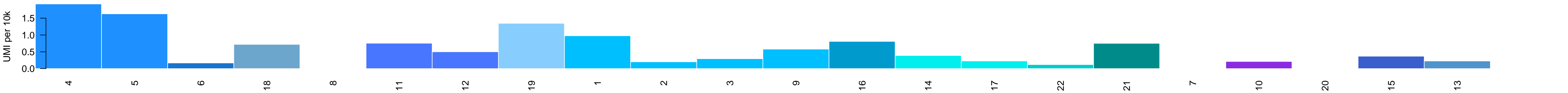
FERM_M.HG1.1:like:MYO10/PLEKHH1/PLEKHH2/PLEKHH3
Tadh_TriadG59851



FERM_M.HG1.18:TLN1/TLN2
Tadh_TriadG19795

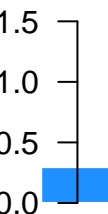


FERM_M.HG1.24:FRMD1/FRMD6/FRMPD2/PTPN13
Tadh_TriadG18577

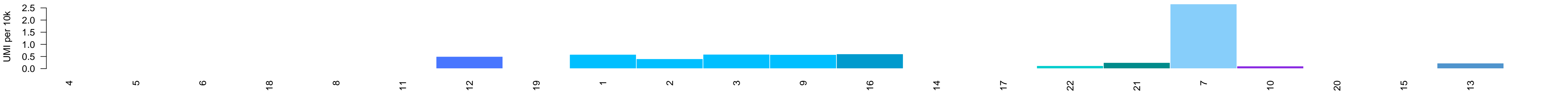


FERM_M.HG1.26:FERMT1/FERMT2/FERMT3
Tadh_TriadG33394

UMI per 10k

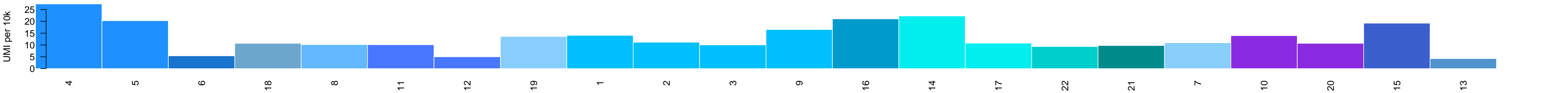


FERM_M.HG1.3:PLEKHH1/PLEKHH2
Tadh_TriadG33842

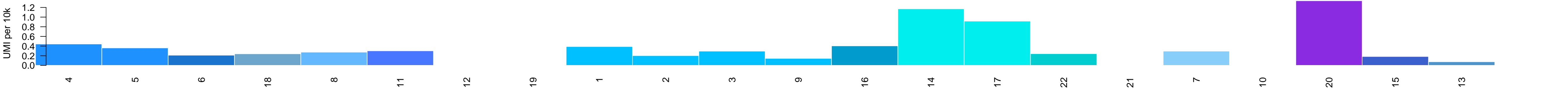


FERM_M.HG1.32:EPB41/EPB41L1/EPB41L2/EPB41L3/EPB41L4A/EPB41L4B/EPB41L5/FARP1/FARP2/FRMD3/FRMD5/FRMD7/PTPN3/PTPN4

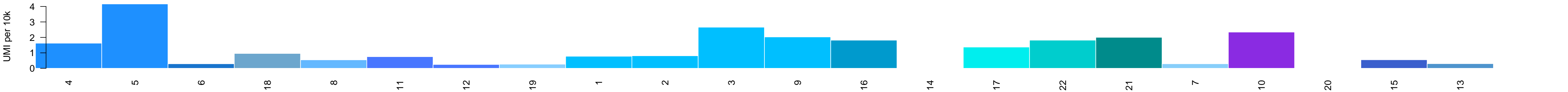
Tadh_TriadG64069



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



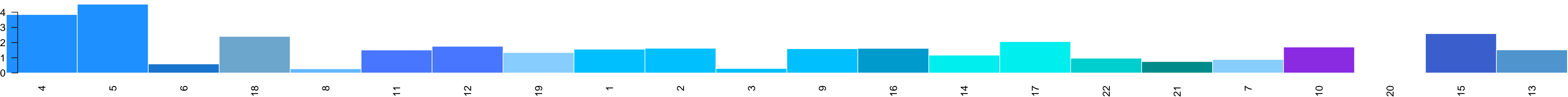
FERM_M.HG1.33:like:EZR/MSN/NF2/RDX
Tadh_TriadG29123



FERM_M.HG1.40:EZR/MSN/RDX

Tadh_TriadG25177

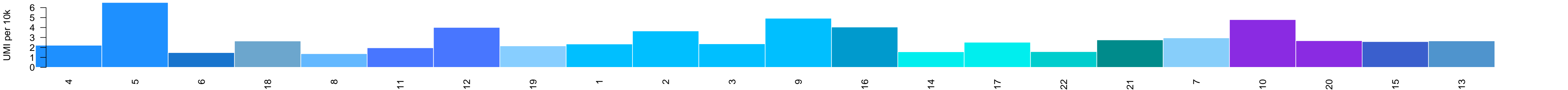
UMI per 10k



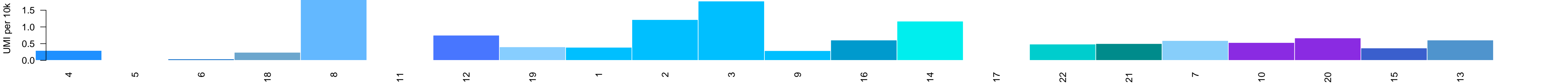
Guanylate_kin.HG1.12:LRGUK
Tadh_TriadG52391



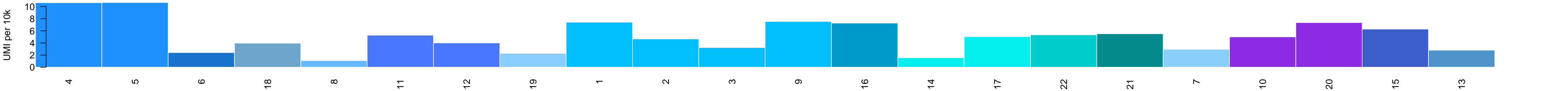
Guanylate_kin.HG1.14:DLG1/DLG2/DLG3/DLG4
Tadh_TriadG63520



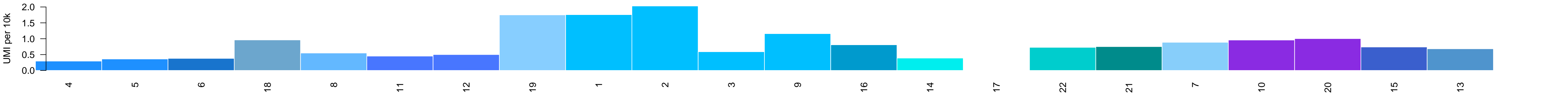
Guanylate_kin.HG1.16:GUK1
Tadh_TriadG50000



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



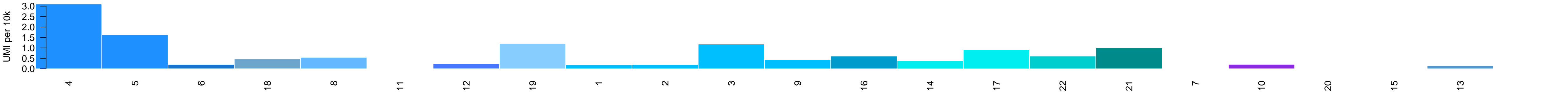
Guanylate_kin.HG2.0:CACNB1/CACNB2/CACNB3/CACNB4
Tadh_TriadG22195



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

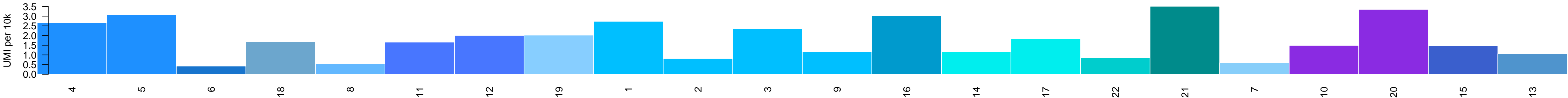


Homer.HG1.2:HOMER1/HOMER2/HOMER3
Tadh_TriadG22944

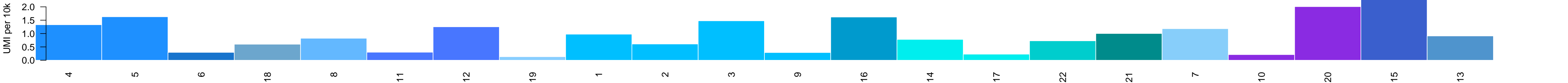


MARVEL.HG1.13:SYNPR/SYP/SYPL1/SYPL2

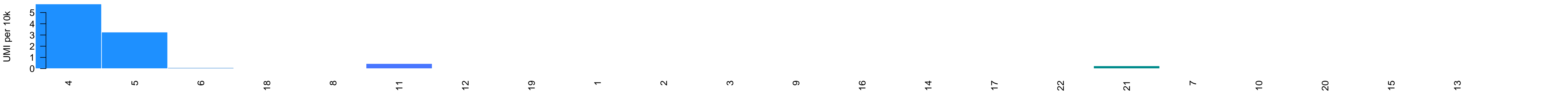
Tadh_TriadG52445

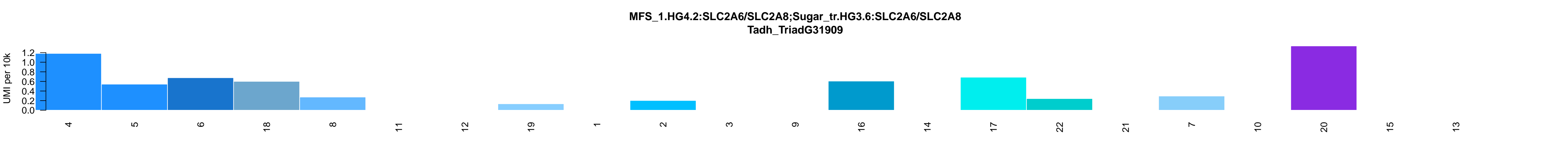


MFS_1.HG10.1:SLC49A3
Tadh_TriadG32990

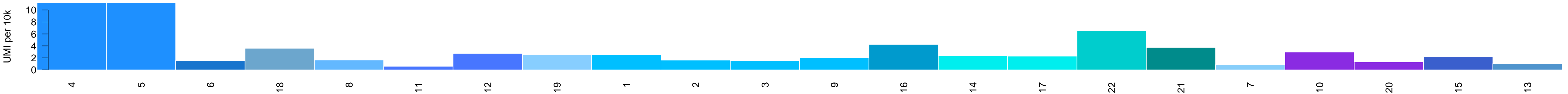


MFS_1.HG12.2:like:MFSD6
Tadh_TriadG60583





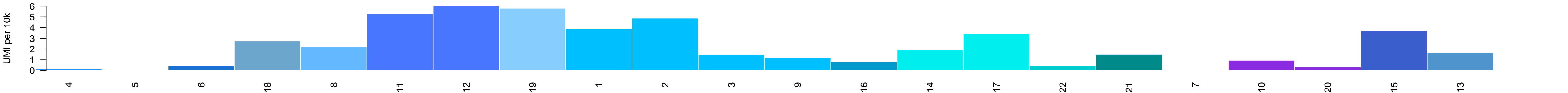
MFS_1.HG4.3:like:SLC2A10/SLC2A12;Sugar_tr.HG3.4:like:SLC2A10/SLC2A12/SLC2A13
Tadh_TriadG64243



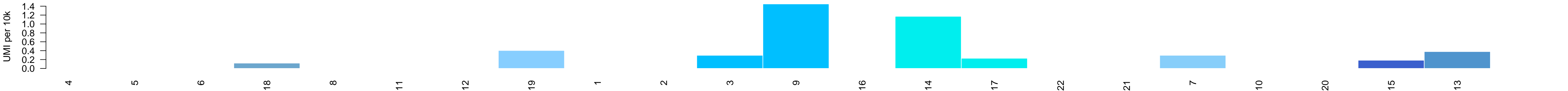
MFS_1.HG5.21:SLC17A6/SLC17A7/SLC17A8
Tadh_TriadG36003



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



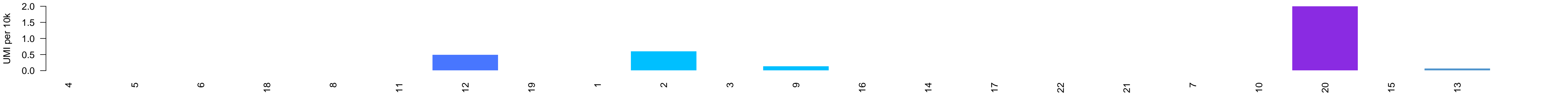
MFS_1.HG6.1:like:SLCO1A2/SLCO1B1/SLCO1B3/SLCO1B3–SLCO1B7/SLCO1C1/SLCO2A1/SLCO2B1/SLCO3A1/SLCO4A1/SLCO4C1/SLCO5A1/SLCO6A1
Tadh_TriadG1002



MFS_1.HG6.1:like:SLCO1A2/SLCO1B1/SLCO1B3/SLCO1B3–SLCO1B7/SLCO1C1/SLCO2A1/SLCO2B1/SLCO3A1/SLCO4A1/SLCO4C1/SLCO5A1/SLCO6A1
Tadh_TriadG11683



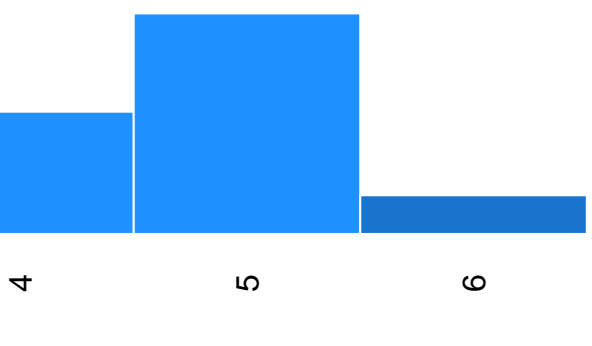
MFS_1.HG6.1:like:SLCO1A2/SLCO1B1/SLCO1B3/SLCO1B3–SLCO1B7/SLCO1C1/SLCO2A1/SLCO2B1/SLCO3A1/SLCO4A1/SLCO4C1/SLCO5A1/SLCO6A1
Tadh_TriadG11804



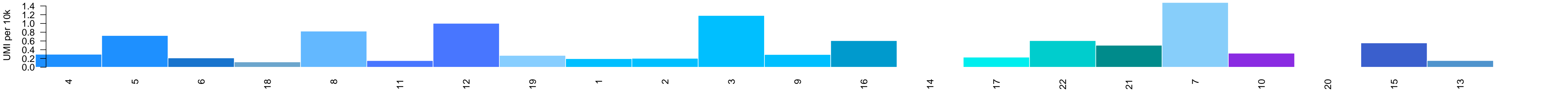
MFS_1.HG7.0:SLC18A1/SLC18A2/SLC18A3/SLC18B1
Tadh_TriadG58957

UMI per 10k

5
4
3
2
1
0



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



MFS_1.HG8.11:SLC22A18
Tadh_TriadG58186

UMI per 10k

1.5
1.0
0.5
0.0



4

5

6

18

8

11

12

19

1

2

3

9

16

14

17

22

21

7

10

20

15

13

MFS_1.HG8.5:MFSD9;Sugar_tr.HG2.5:MFSD9
Tadh_TriadG25848

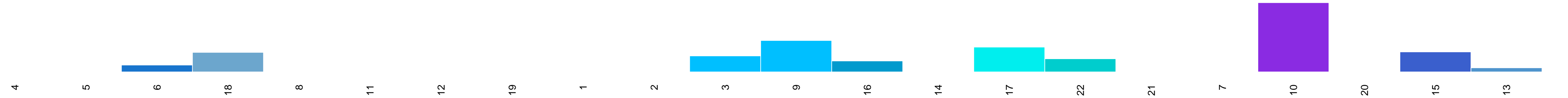
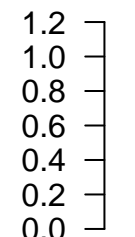


NCam.HG1.3:NCAM1/NCAM2
Tadh_TriadG59528

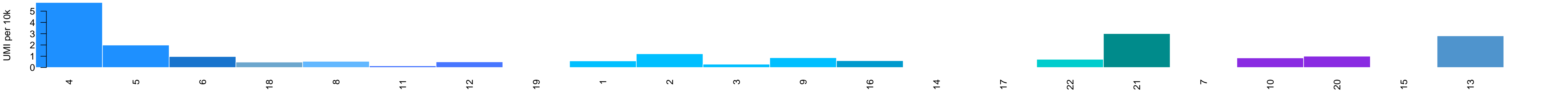


NO_synthase.HG1.0:NOS1/NOS2/NOS3
Tadh_TriadG19964

UMI per 10k



PDZ.HG2.2:like:PARD6A/PARD6B/PARD6G
Tadh_TriadG58026

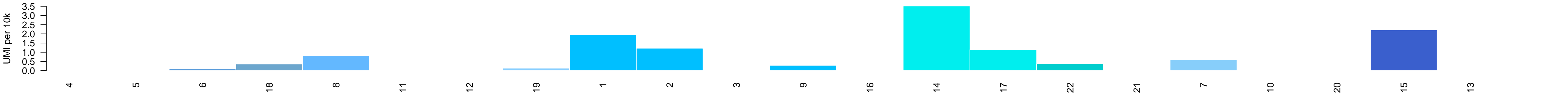


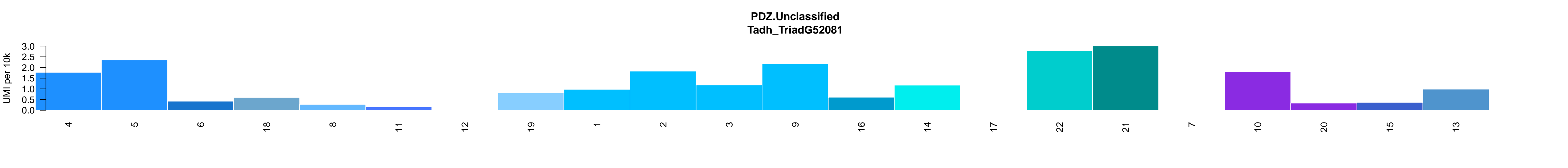
PDZ.HG4.9:like:HTRA1/HTRA2/HTRA3/HTRA4
Tadh_TriadG58565

UMI per 10k

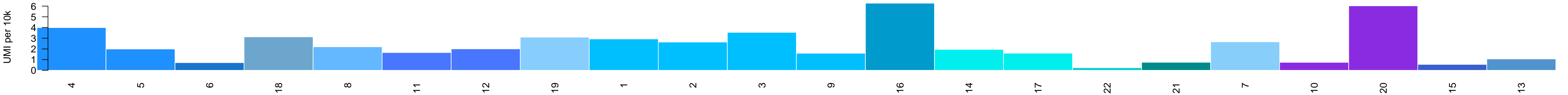


PDZ.Unclassified
Tadh_TriadG28602

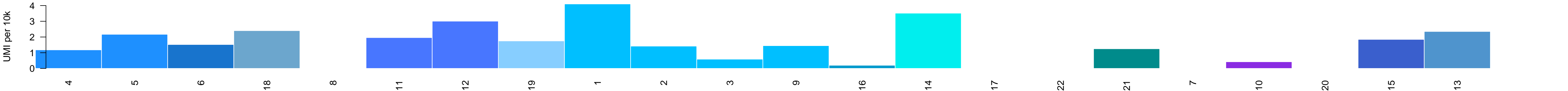


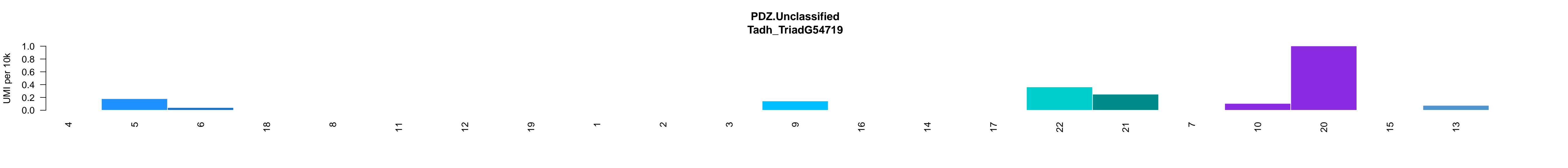


PDZ.Unclassified
Tadh_TriadG53974

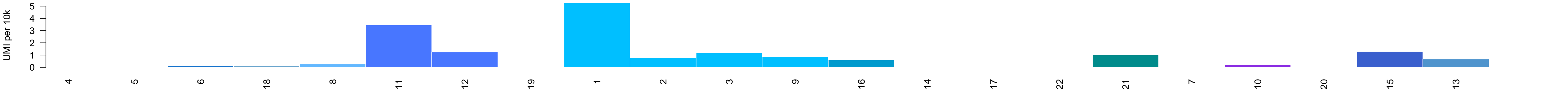


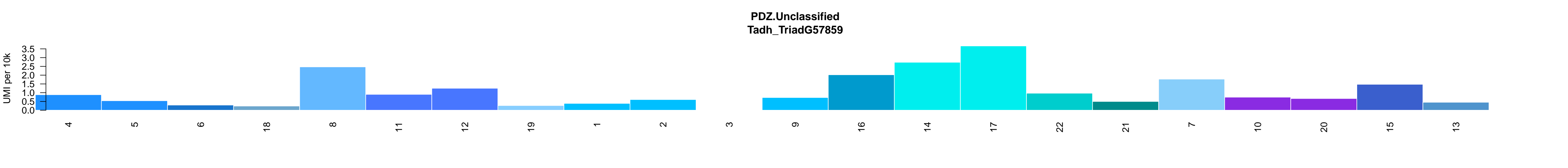
PDZ.Unclassified
Tadh_TriadG54453



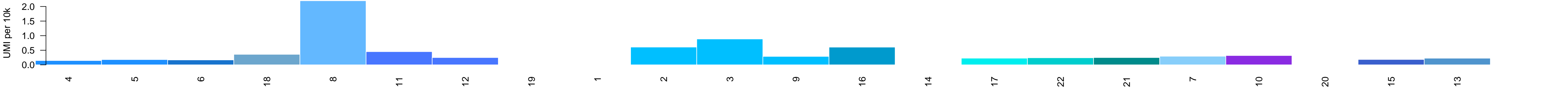


PDZ.Unclassified
Tadh_TriadG55409

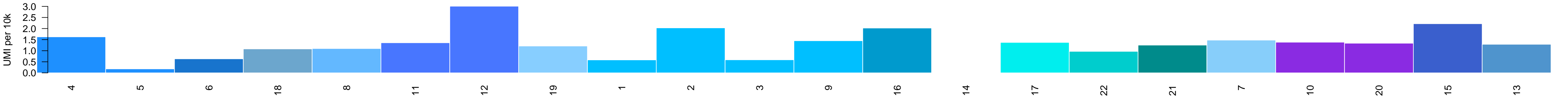




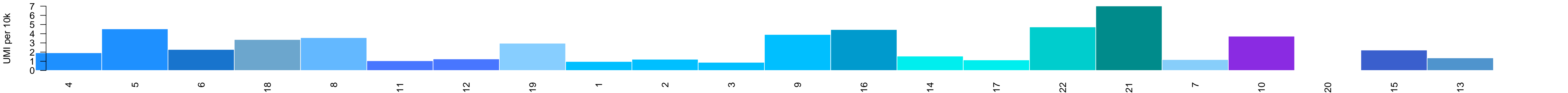
PDZ.Unclassified
Tadh_TriadG58260



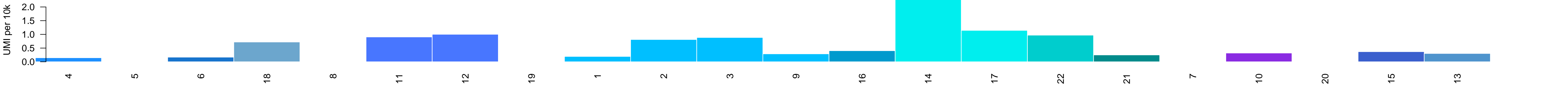
PDZ.Unclassified
Tadh_TriadG59148



PDZ.Unclassified
Tadh_TriadG61629

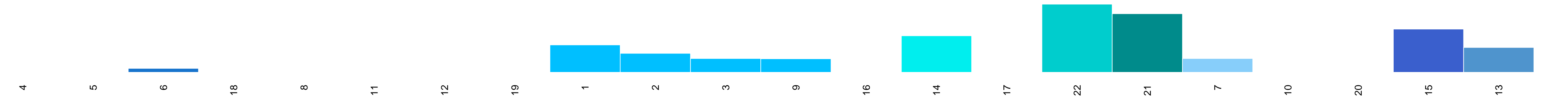
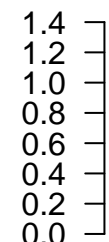


PDZ.Unclassified
Tadh_TriadG62108

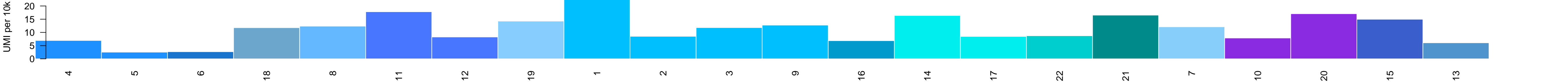


PDZ.Unclassified
Tadh_TriadG62142

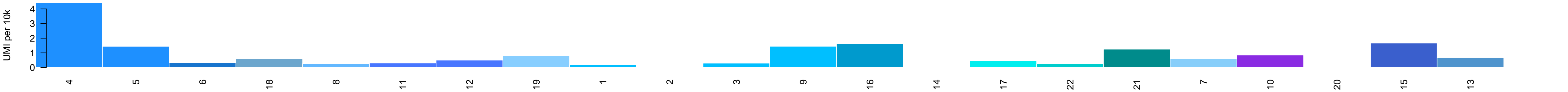
UMI per 10k



PDZ.Unclassified
Tadh_TriadG64065



PDZ.Unclassified
Tadh_wf_g4045



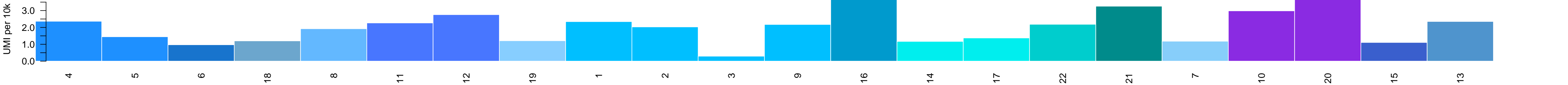
PID.HG1.0:RABGAP1
Tadh_TriadG21193



PMCA.HG1.2:ATP2C1/ATP2C2
Tadh_TriadG19369



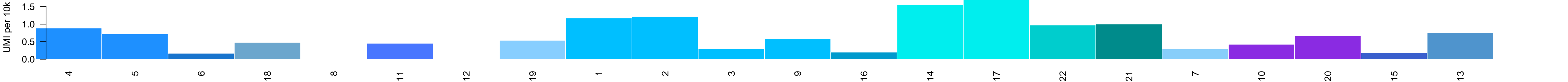
PMCA.HG1.5:AC010616.1/ATP1A1/ATP1A2/ATP1A3/ATP1A4/ATP4A/ATP12A
Tadh_TriadG63163



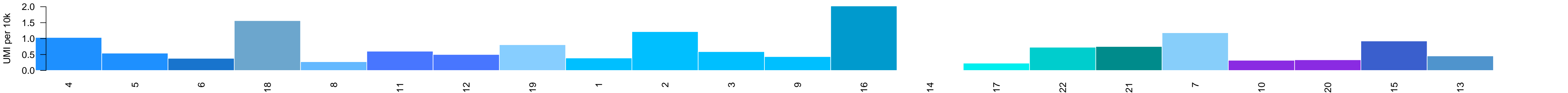
Pyridoxal_deC.HG1.4:CSAD/GAD1/GAD2/GADL1
Tadh_TriadG60611



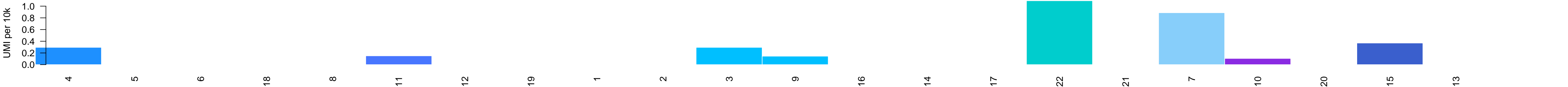
RasGAP.HG1.2:NF1
Tadh_TriadG59057



RasGAP.HG1.4:IQGAP1/IQGAP2
Tadh_TriadG37304

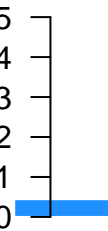


RasGAP.HG1.5:DAB2IP/RASAL2/RASAL3/SYNGAP1;SynGAP.HG1.2:like:DAB2IP/RASAL2/SYNGAP1
Tadh_wf_g1897



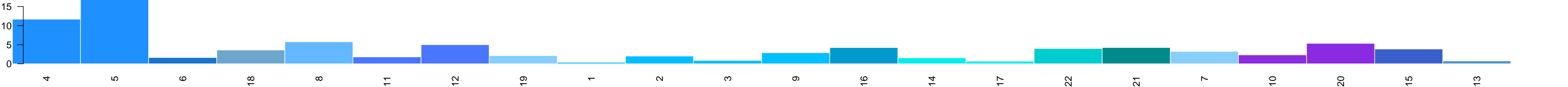
RasGAP.HG1.7:RASA1
Tadh_wf_g3134

UMI per 10k



SDF.HG1.1:SLC1A1/SLC1A2/SLC1A3/SLC1A4/SLC1A5/SLC1A6/SLC1A7
Tadh_TriadG11741

UMI per 10k



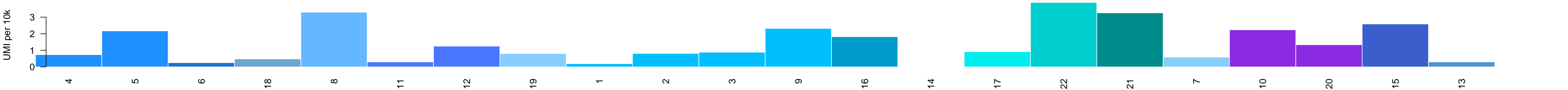
SNF.HG1.1:like:AC136616.1/SLC6A1/SLC6A2/SLC6A3/SLC6A4/SLC6A5/SLC6A6/SLC6A7/SLC6A8/SLC6A9/SLC6A11/SLC6A12/SLC6A13/SLC6A14/SLC6A15/SLC6A16/SLC6A17/SLC6A18/SLC6A19/SLC6A20

Tadh_TriadG33521



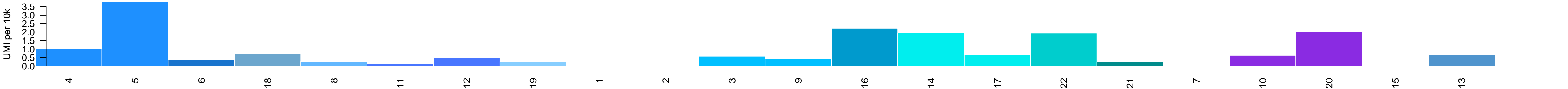
SNF.HG1.2:like:AC136616.1/SLC6A1/SLC6A2/SLC6A3/SLC6A4/SLC6A5/SLC6A6/SLC6A7/SLC6A8/SLC6A9/SLC6A11/SLC6A12/SLC6A13/SLC6A14/SLC6A15/SLC6A16/SLC6A17/SLC6A18/SLC6A19/SLC6A20

Tadh_TriadG27813



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_TriadG29421



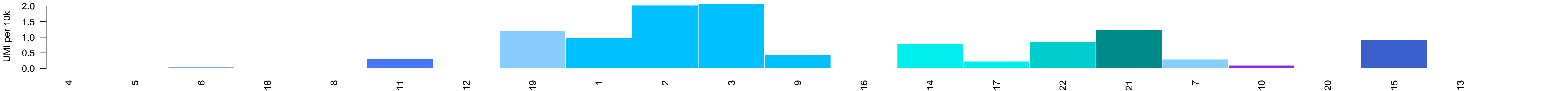
SNF.HG1.2:like:AC136616.1/SLC6A1/SLC6A2/SLC6A3/SLC6A4/SLC6A5/SLC6A6/SLC6A7/SLC6A8/SLC6A9/SLC6A11/SLC6A12/SLC6A13/SLC6A14/SLC6A15/SLC6A16/SLC6A17/SLC6A18/SLC6A19/SLC6A20

Tadh_TriadG33157



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

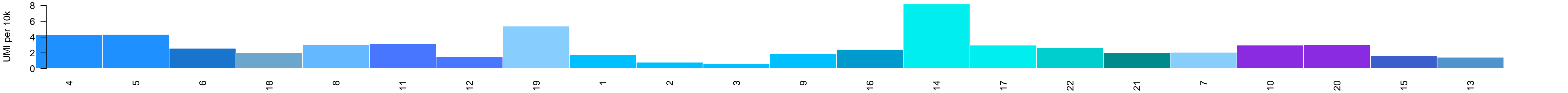
Tadh_TriadG59303



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_TriadG61885



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

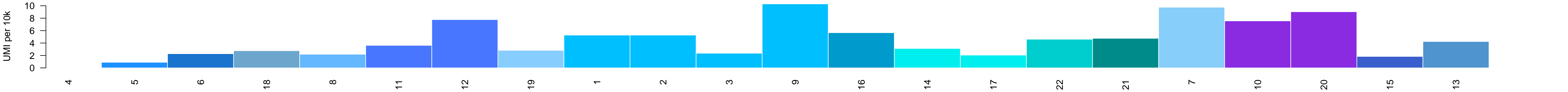


SNF.HG1.30:like:SLC6A5/SLC6A7/SLC6A9/SLC6A14
Tadh_TriadG50950



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

Tadh_TriadG38332



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

UMI per 10k

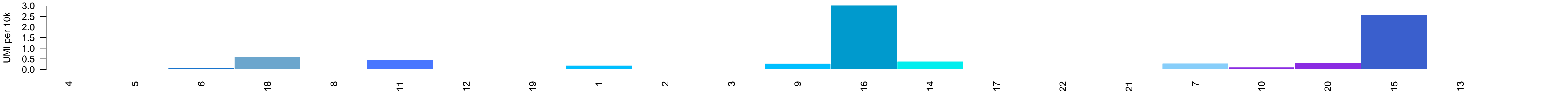


SNF.HG1.46:like:SLC6A5/SLC6A7/SLC6A9/SLC6A14
Tadh_TriadG54650



SNF.HG1.5:like:AC136616.1/SLC6A1/SLC6A2/SLC6A3/SLC6A4/SLC6A5/SLC6A6/SLC6A7/SLC6A8/SLC6A9/SLC6A11/SLC6A12/SLC6A13/SLC6A14/SLC6A15/SLC6A16/SLC6A17/SLC6A18/SLC6A19/SLC6A20

Tadh_TriadG33106

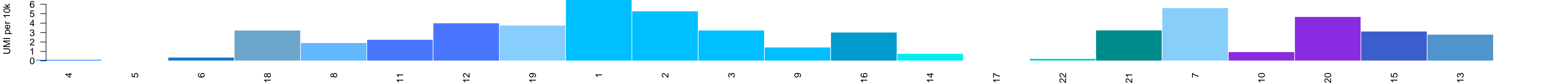


SNF.HG1.6:like:AC136616.1/SLC6A1/SLC6A2/SLC6A3/SLC6A4/SLC6A5/SLC6A6/SLC6A7/SLC6A8/SLC6A9/SLC6A11/SLC6A12/SLC6A13/SLC6A14/SLC6A15/SLC6A16/SLC6A17/SLC6A18/SLC6A19/SLC6A20

Tadh_TriadG33097



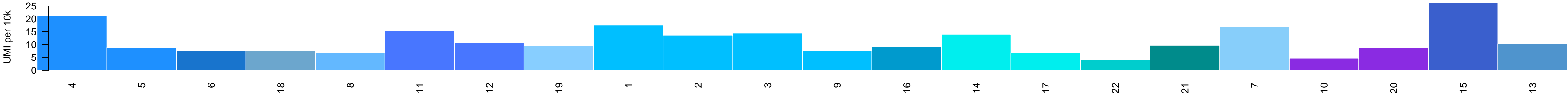
Synaphin.HG2.0:NA
Tadh_wf_g9757



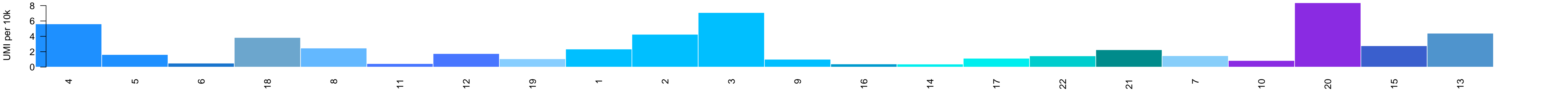
Synaptotagmin.HG1.0:SYT7
Tadh_TriadG32942



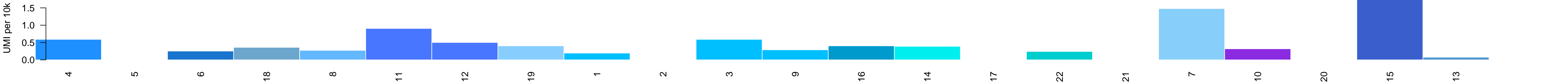
Synaptotagmin.HG1.2:SYT1/SYT2/SYT5/SYT8
Tadh_TriadG51110



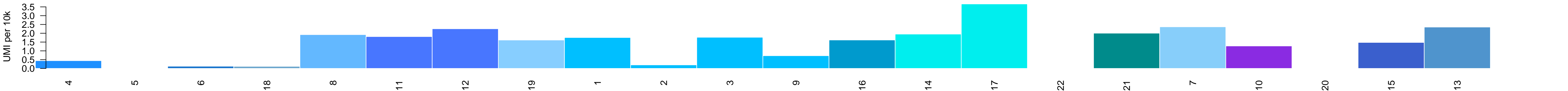
Synaptotagmin.HG1.32:BAIAP3/UNC13D
Tadh_TriadG60504



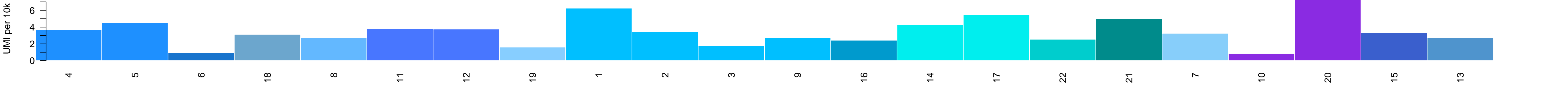
Synaptotagmin.HG1.42:MCTP1
Tadh_TriadG60419



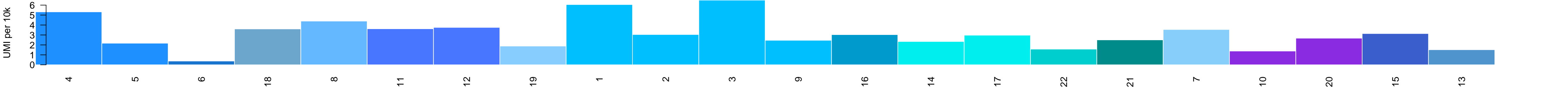
Synaptotagmin.HG1.45:like:SYTL1/SYTL2/SYTL3/SYTL4/SYTL5
Tadh_TriadG56437



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



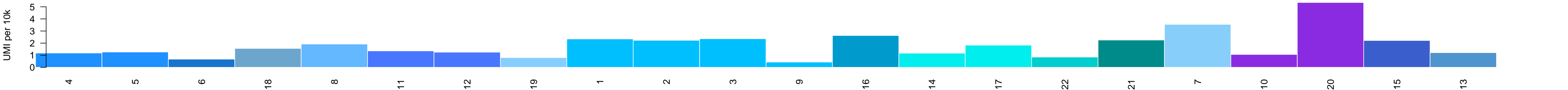
Syntaxin.HG1.2:STX7/STX12
Tadh_TriadG27557



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



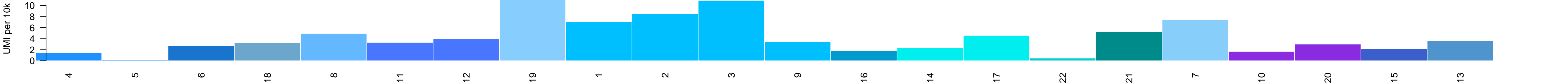
Syntaxin.HG1.4:STX1A/STX1B/STX2/STX3/STX4
Tadh_TriadG28648



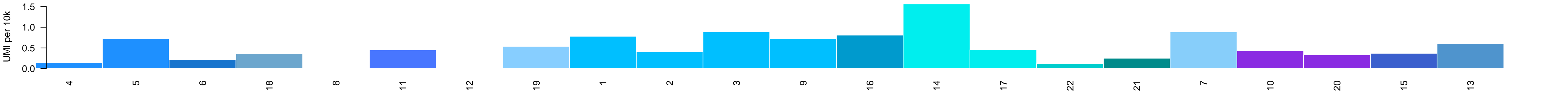
Y_phosphatase.HG1.26:like:PTPN9
Tadh_TriadG54170



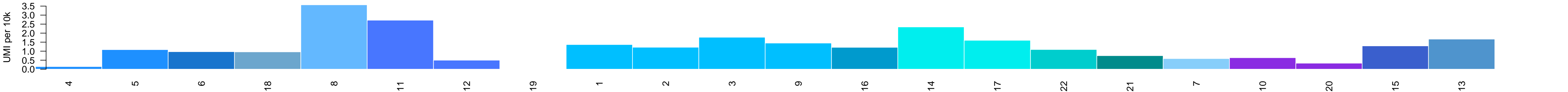
Y_phosphatase.HG1.29:PTPRN/PTPRN2
Tadh_TriadG55048



Y_phosphatase.HG1.52:PTPRG/PTPRZ1
Tadh_TriadG7563



Y_phosphatase.HG2.6:PTPDC1
Tadh_TriadG56940



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

UMI per 10k

2.5
2.0
1.5
1.0
0.5
0.0

