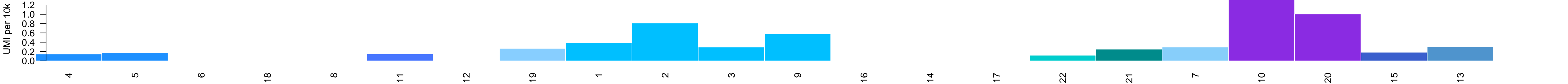
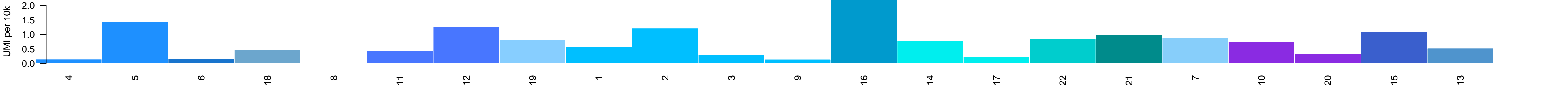


CYP3A7_CYP3A51P_readthrough | p450/
Tadh_TriadG13325



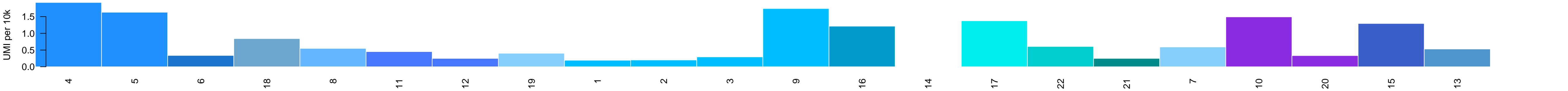
retinol_dehydrogenase_13 | adh_short/
Tadh_TriadG19476



retinol_dehydrogenase_13 | adh_short/
Tadh_TriadG19714



hydroxysteroid_17_beta_dehydrogenase_10 | adh_short/
Tadh_TriadG22420



cytochrome_P450_family_51_subfamily_A_member_1 | p450/
Tadh_TriadG22633

UMI per 10k

10
8
6
4
2
0

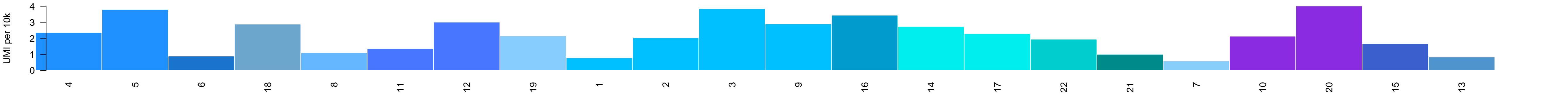
4 5 6 18 8 11 12 19 1 2 3 9 16 14 17 22 21 7 10 20 15 13



cytochrome_P450_family_3_subfamily_A_member_4 | p450/
Tadh_TriadG23953



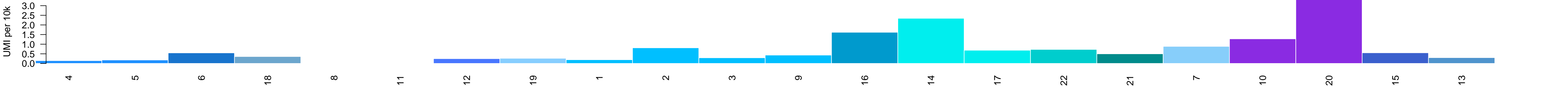
ubiquinol_cytochrome_c_reductase_Rieske_iron_sulfur_polypeptide_1 | UCR_TM/Rieske/
Tadh_TriadG24746



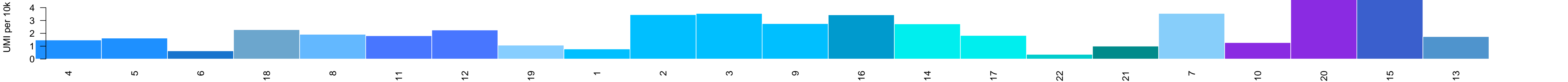
cytochrome_P450_family_3_subfamily_A_member_5 | p450/
Tadh_TriadG25372



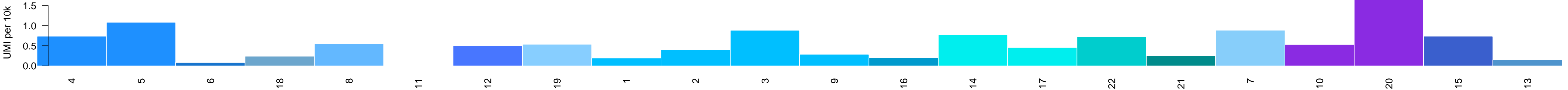
sigma_non_opioid_intracellular_receptor_1 | ERG2_Sigma1R/
Tadh_TriadG28090



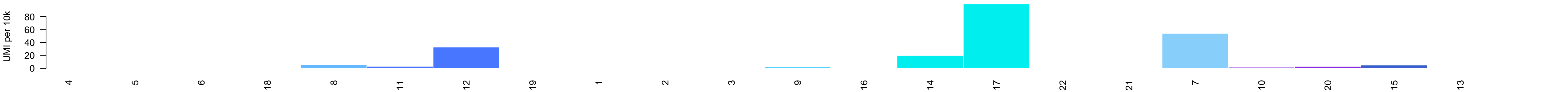
retinol_dehydrogenase_12 | adh_short/
Tadh_TriadG28989



hydroxysteroid_17_beta_dehydrogenase_2 | adh_short/
Tadh_TriadG31297



cytochrome_P450_family_24_subfamily_A_member_1 | p450/
Tadh_TriadG34090



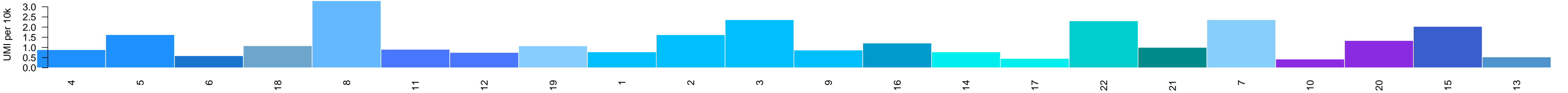
retinol_dehydrogenase_16 | adh_short/
Tadh_TriadG35982



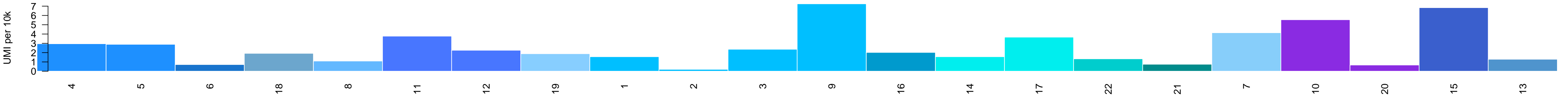
fatty_acid_desaturase_3 | Cyt-b5/FA_desaturase/
Tadh_TriadG36979



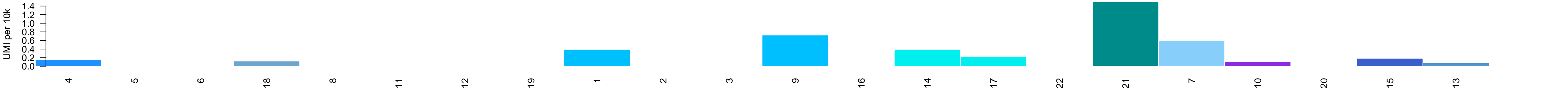
fatty_acid_2_hydroxylase | Cyt-b5/FA_hydroxylase/
Tadh_TriadG37592



15_hydroxyprostaglandin_dehydrogenase | adh_short/
Tadh_TriadG37607



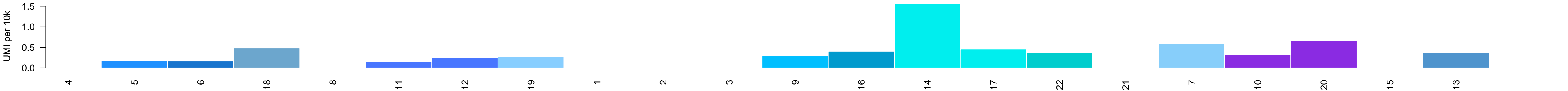
retinol_dehydrogenase_13 | adh_short/
Tadh_TriadG49836



stearoyl_CoA_desaturase | FA_desaturase/
Tadh_TriadG51148



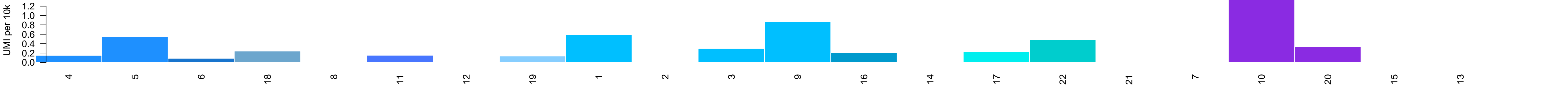
NAD_P_dependent_steroid_dehydrogenase_like | 3Beta_HSD/
Tadh_TriadG52229



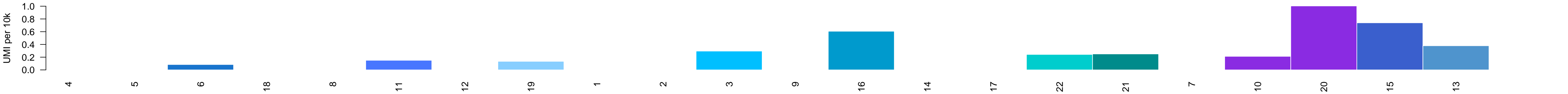
retinol_dehydrogenase_12 | adh_short/
Tadh_TriadG54391



retinol_dehydrogenase_16 | adh_short/
Tadh_TriadG54855



cytochrome_P450_family_3_subfamily_A_member_4 | p450/p450/
Tadh_TriadG55168



cytochrome_P450_family_3_subfamily_A_member_5 | p450/
Tadh_TriadG56203

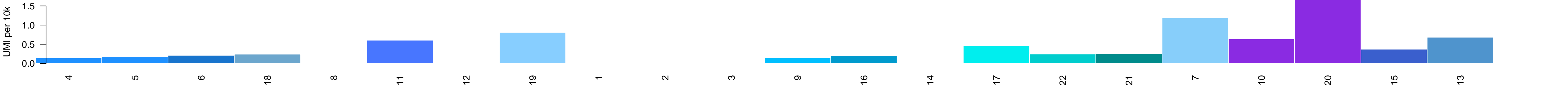
UMI per 10k

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



apoptosis_inducing_factor_mitochondria_associated_3 | Rieske/Pyr_redox_2/Reductase_C/
Tadh_TriadG59728



cytochrome_P450_family_26_subfamily_B_member_1 | p450/
Tadh_TriadG60776

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

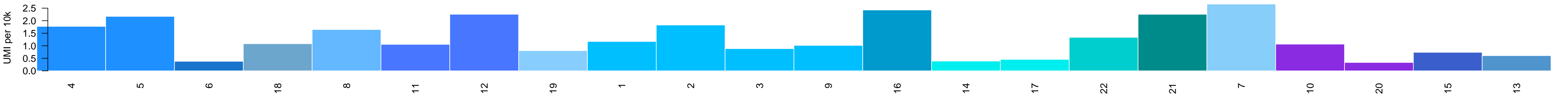
20

15

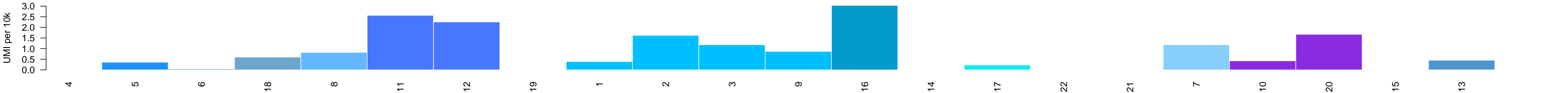
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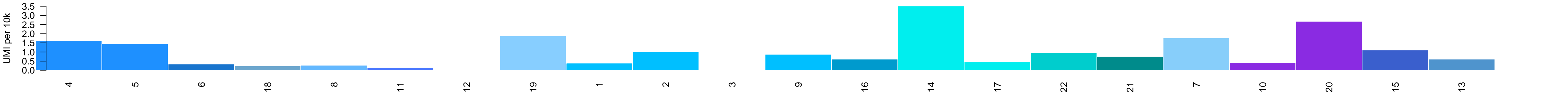
short_chain_dehydrogenase_reductase_family_16C_member_5 | adh_short/
Tadh_TriadG62171



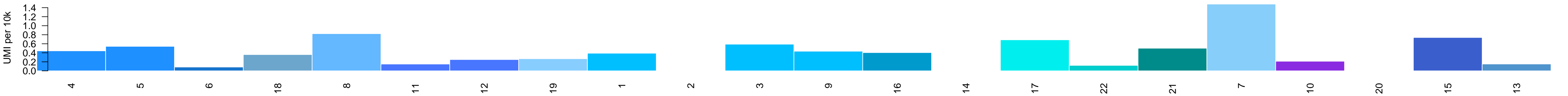
cytochrome_P450_family_3_subfamily_A_member_4 | p450/
Tadh_TriadG62362

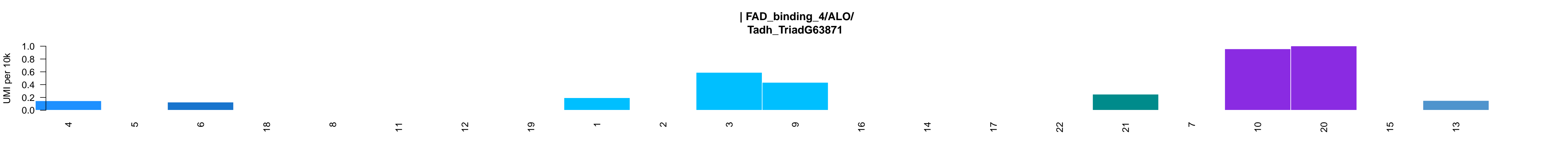


stearoyl_CoA_desaturase | FA_desaturase/
Tadh_TriadG63195



squalene_epoxidase | SE/
Tadh_TriadG63342





Rieske_Fe_S_domain_containing | Rieske/
Tadh_wf_g866

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

