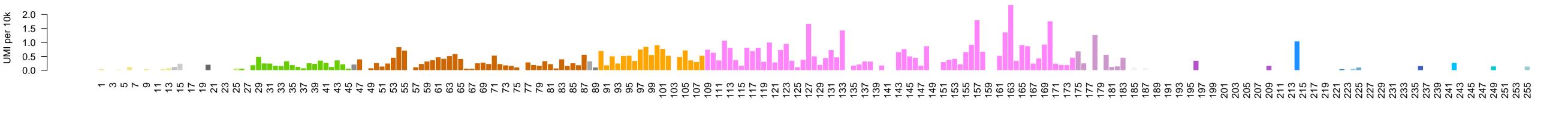
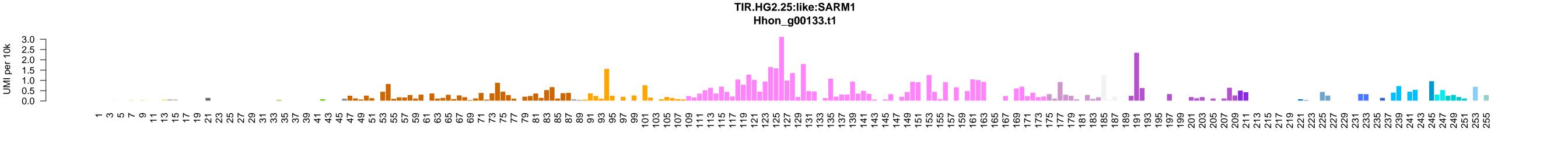
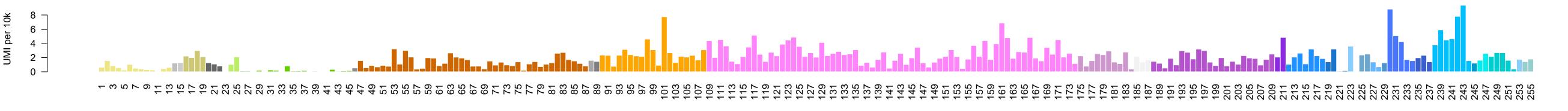
TGF_beta.HG1.4:BMP3/GDF10 Hhon_g00066.t1



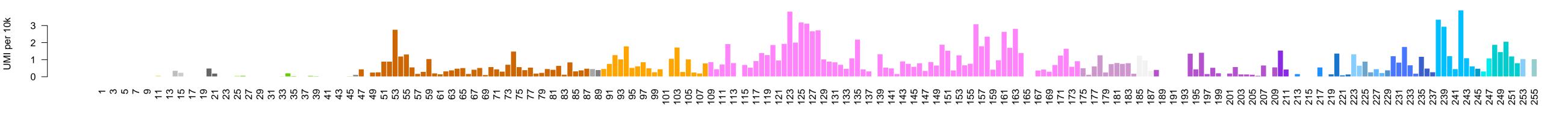
RGS.HG1.13:RGS22 Hhon_g00113.t1



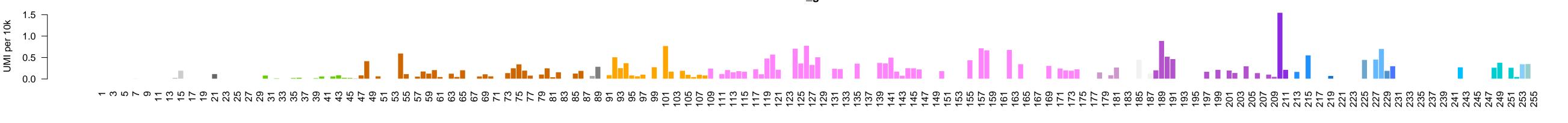
Ldl_recept_b.Unclassified Hhon_g00195.t1



FGF.HG1.2:FGF8/FGF17/FGF18 Hhon_g00258.t1

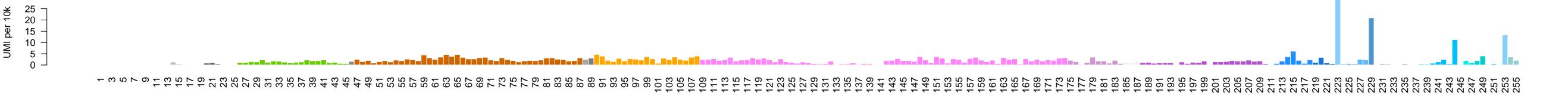


R.HG2.10:like:SARM1 Hhon_g00266.t1

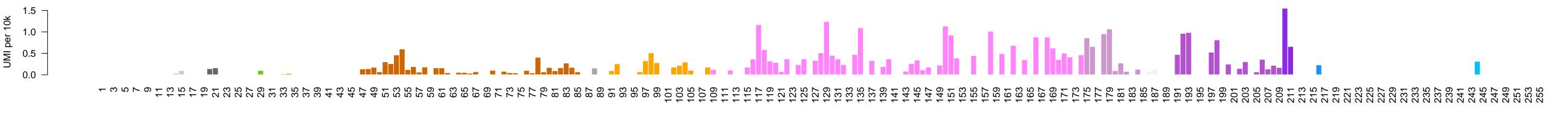


RGS.HG1.18:like:RGS1/RGS2/RGS3/RGS4/RGS5/RGS8/RGS13/RGS16/RGS17/RGS18/RGS19/RGS20/RGS21

Hhon_g01310.t1



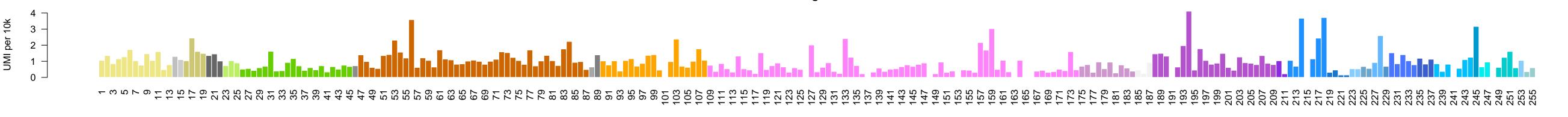
Notch.HG1.1:NOTCH1/NOTCH2/NOTCH3/NOTCH4
Hhon_g01551.t1



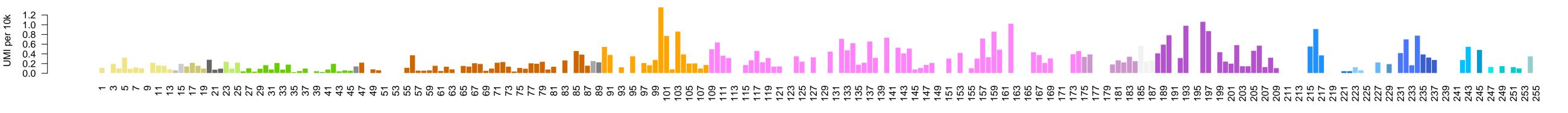
TIR.HG1.19:like:MYD88 Hhon_g01741.t1



Patched.HG1.3:HMGCR Hhon_g02462.t1



R.HG2.33:like:SARM1 Hhon_g02603.t1

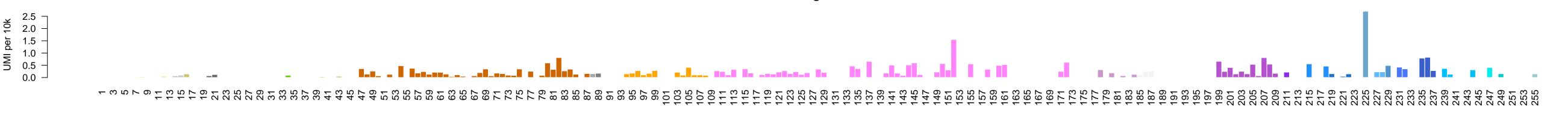


TGF_beta.HG1.32:like:INHBA/INHBB/INHBC/INHBE

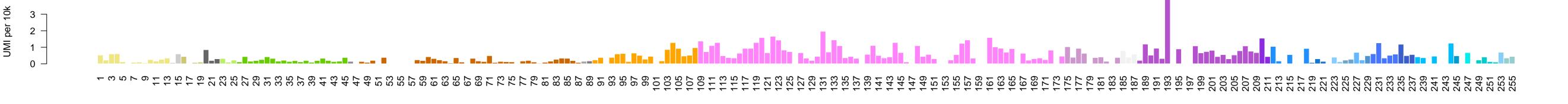
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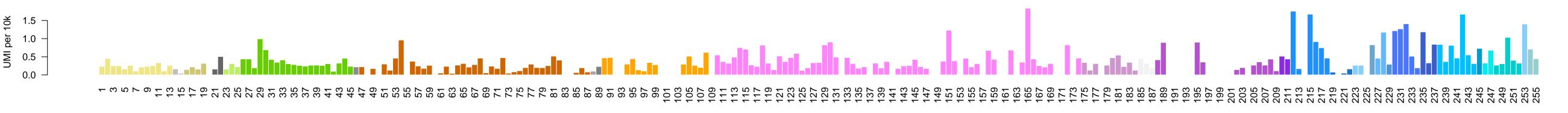




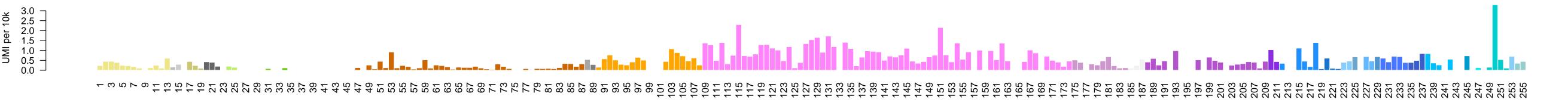
MH2.HG1.0:SMAD6/SMAD7 Hhon_g04010.t1



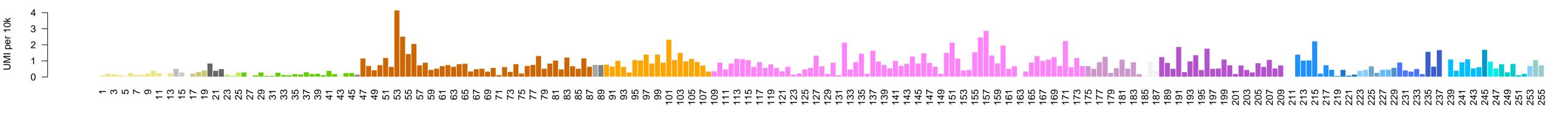
Patched.HG1.1:like:NPC1/NPC1L1/PTCH1/PTCH2/PTCHD1/PTCHD3/PTCHD4
Hhon_g04014.t1



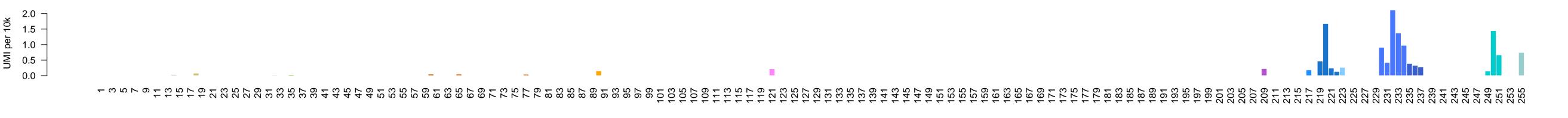
Frizzled.HG1.12:FZD3/FZD6 Hhon_g04136.t1



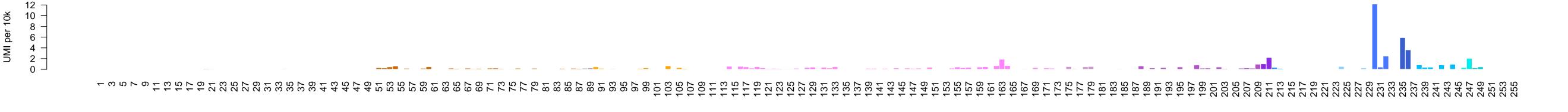
Ldl_recept_b.Unclassified Hhon_g04149.t1



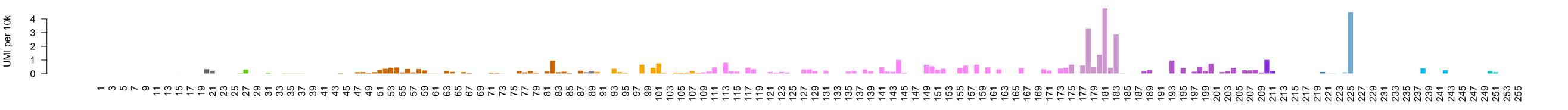
Frizzled.HG1.43:like:MUSK/ROR1/ROR2 Hhon_g04472.t1



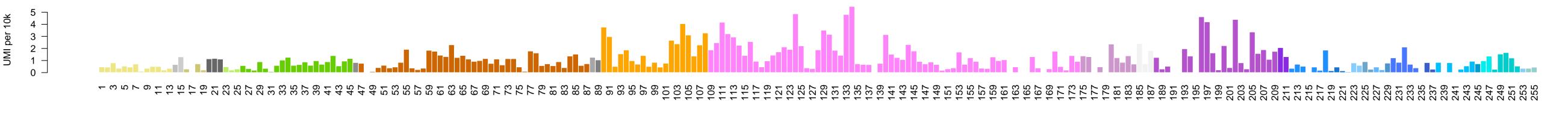
LAG1-DNAbind.HG1.0:RBPJ/RBPJL Hhon_g05183.t1



Ldl_recept_b.Unclassified Hhon_g05269.t1

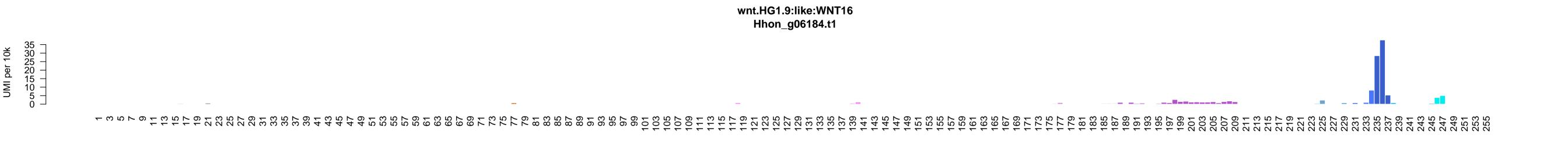


MH2.HG1.13:like:SMAD2/SMAD3 Hhon_g05417.t1

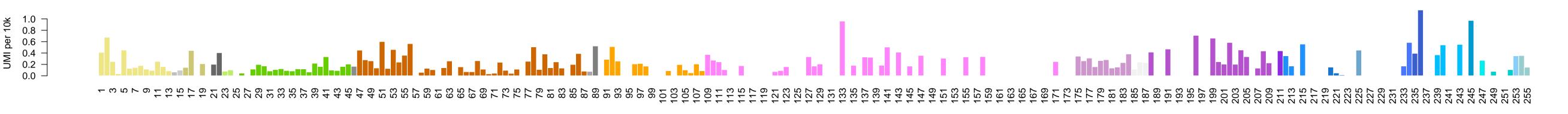








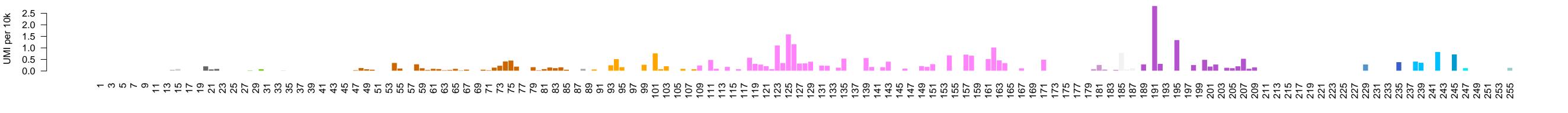
Patched.HG1.4:SCAP
Hhon_g06299.t1



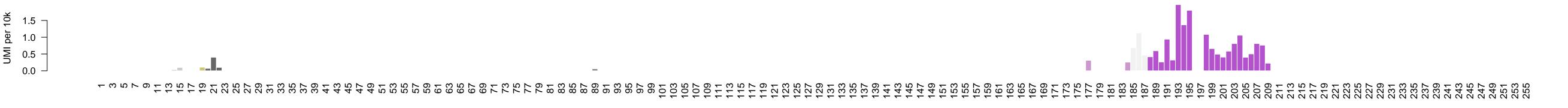
MH2.HG1.14:SMAD2/SMAD3 Hhon_g06533.t1



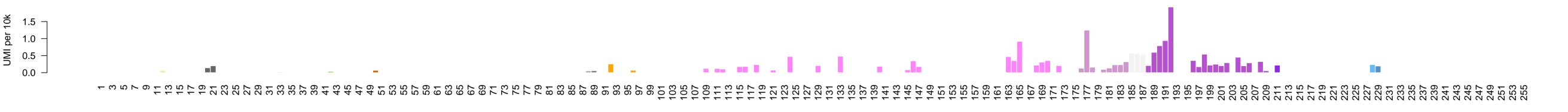
RGS.HG1.10:like:RGSL1 Hhon_g06572.t1



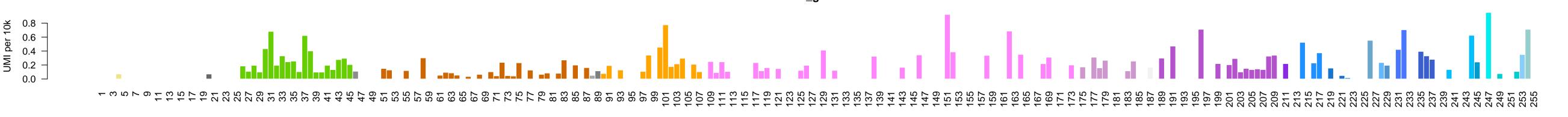
TGF_beta.HG1.15:BMP5/BMP6/BMP7/BMP8A/BMP8B Hhon_g06684.t1



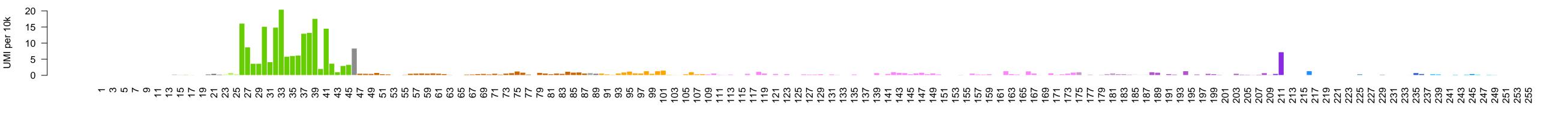
TGF_beta.HG1.15:BMP5/BMP6/BMP7/BMP8A/BMP8B Hhon_g06685.t1

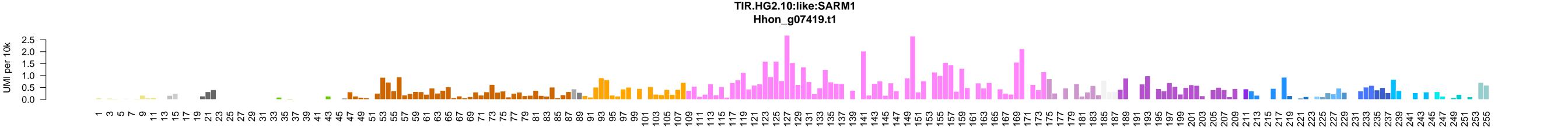


Patched.HG1.2:DISP3 Hhon_g06932.t1

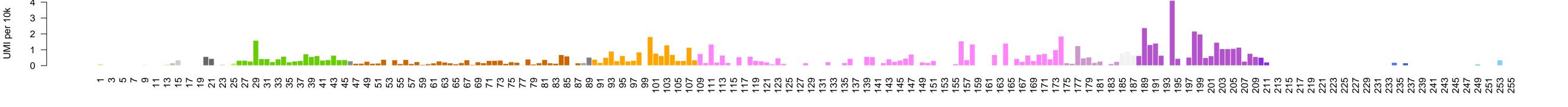


R.HG2.15:like:SARM1 Hhon_g07040.t1

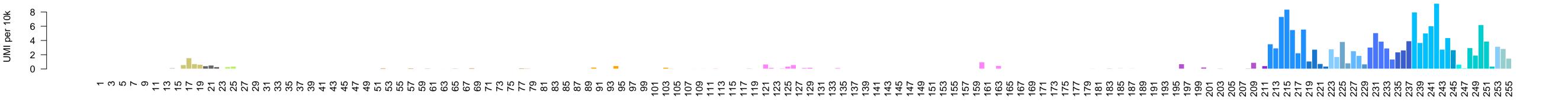




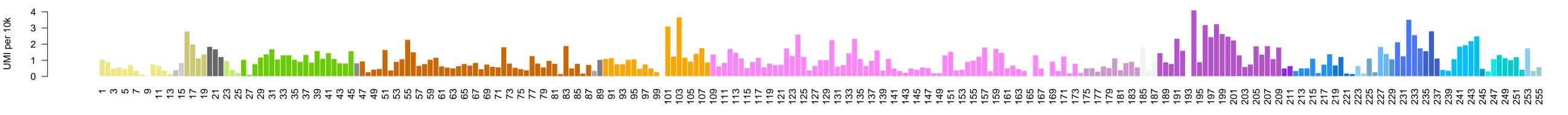
TGF_beta.HG1.13:BMP2/BMP4 Hhon_g07488.t1



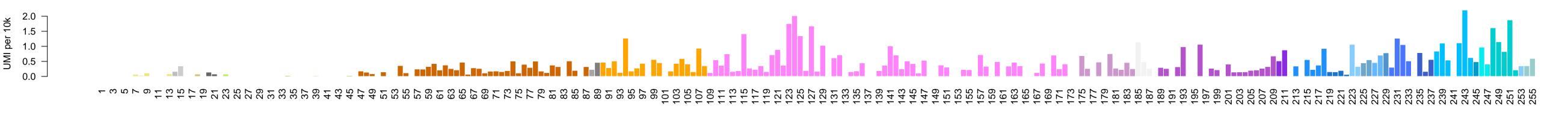
RGS.HG1.21:RGS17/RGS19/RGS20 Hhon_g07606.t1



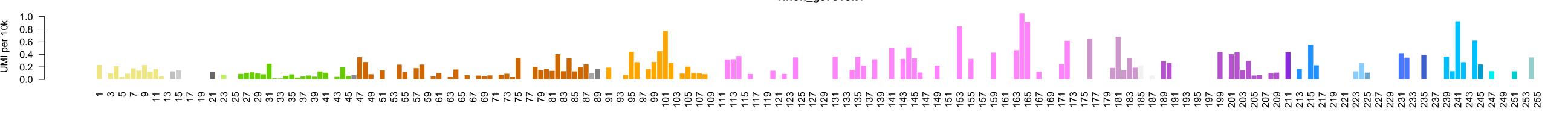
Patched.HG1.13:NPC1/NPC1L1 Hhon_g07714.t1



TIR.HG2.33:like:SARM1 Hhon_g07813.t1



TIR.HG2.19:like:SARM1 Hhon_g07918.t1



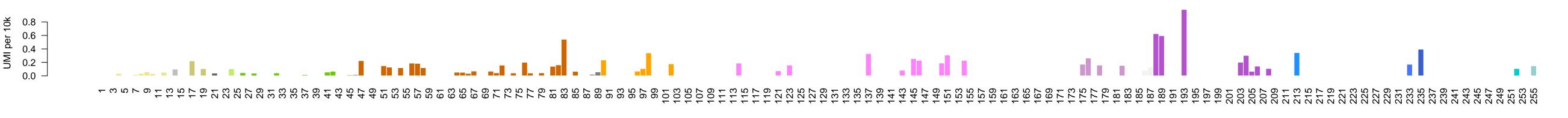
TIR.HG2.14:like:SARM1 Hhon_g08043.t1

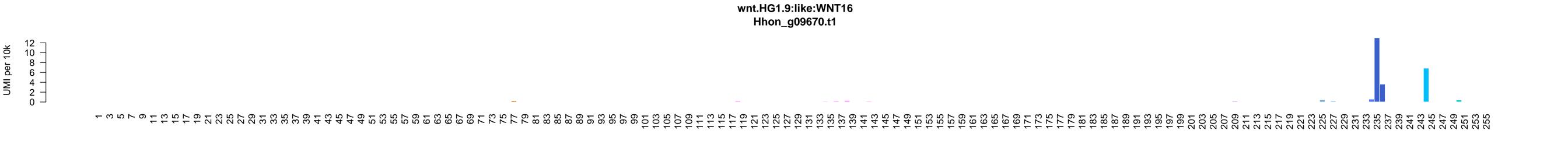


Hhon_g08109.t1

MH2.HG1.1:SMAD4

R.HG2.35:like:SARM1 Hhon_g08301.t1

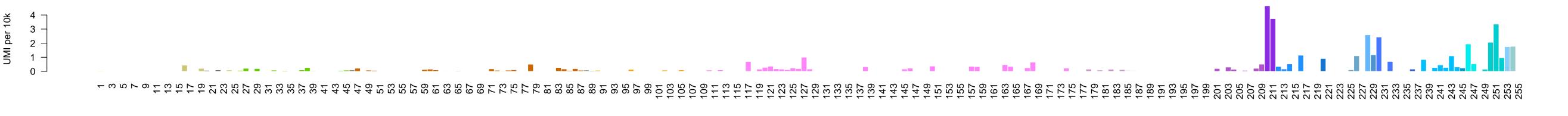




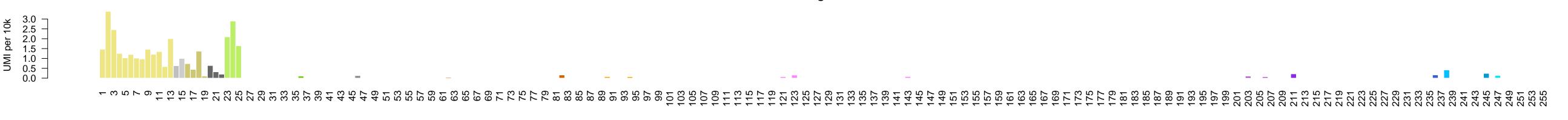
wnt.HG1.21:WNT7A/WNT7B Hhon_g10513.t1



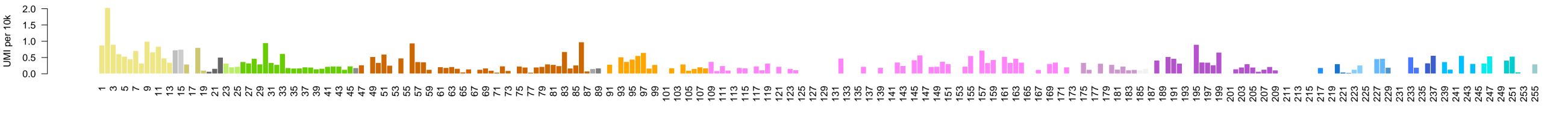
DSL.HG1.4:DLL1/DLL4/JAG1/JAG2 Hhon_g10574.t1



RGS.HG1.11:RGS6/RGS7/RGS9/RGS11 Hhon_g10853.t1







MH2.HG1.14:SMAD2/SMAD3 Hhon_g11830.t1



R.HG2.19:like:SARM1 Hhon_g11841.t1

