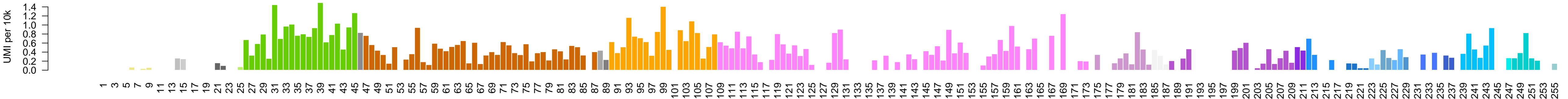
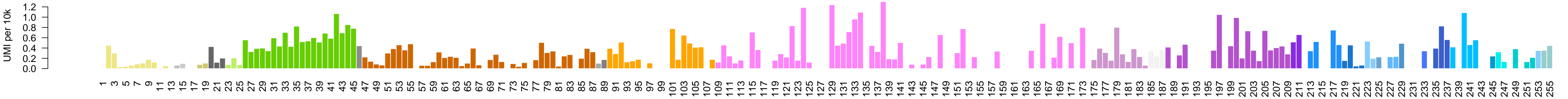


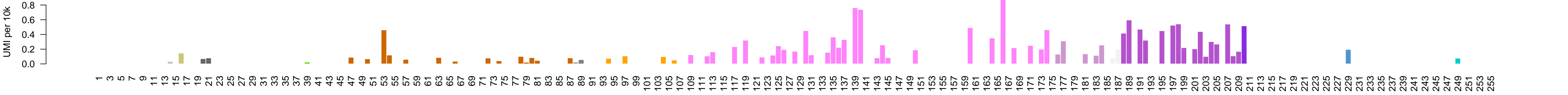
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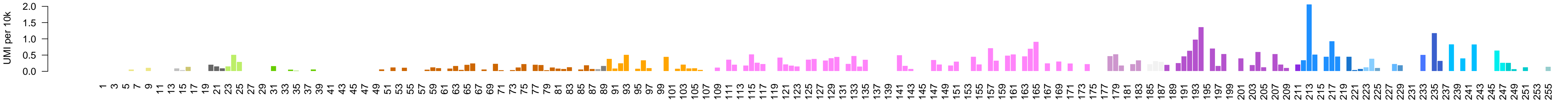
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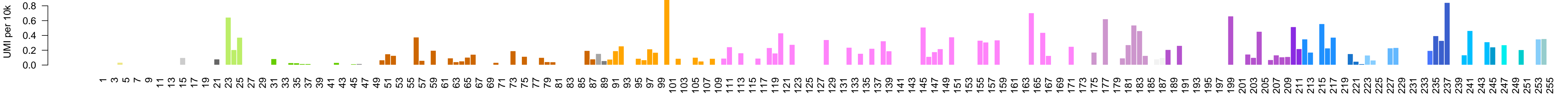


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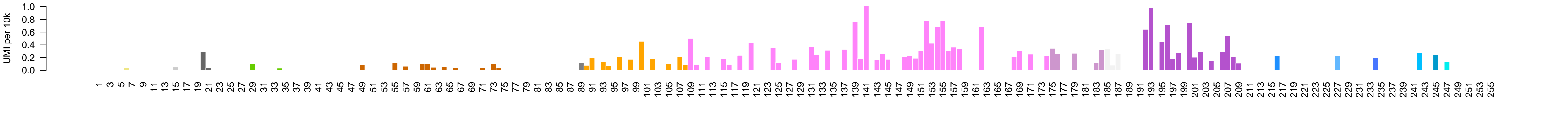


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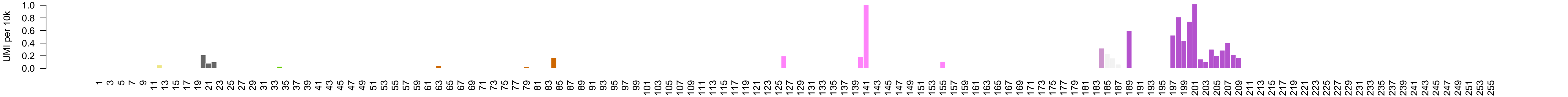




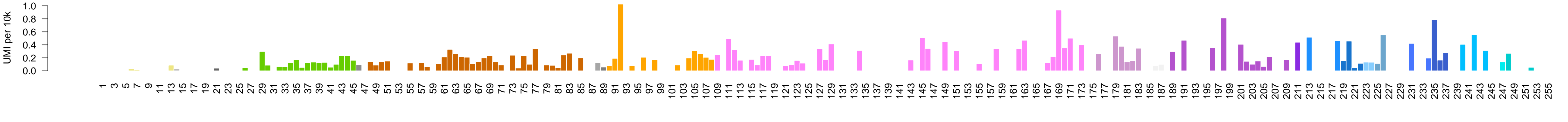
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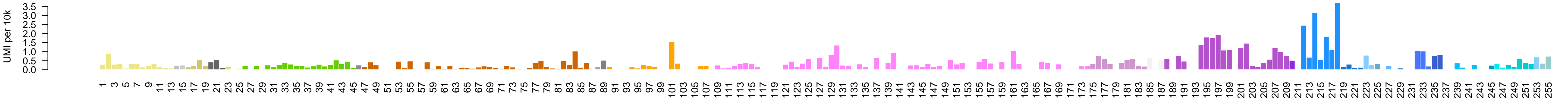
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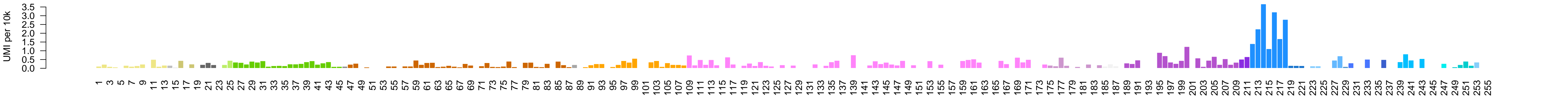
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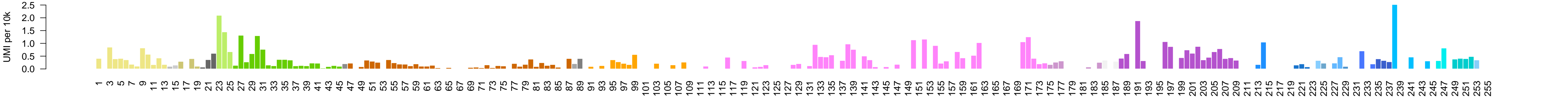
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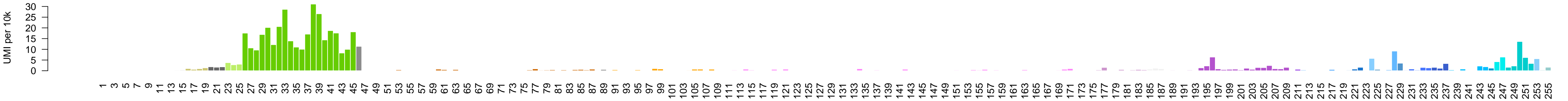
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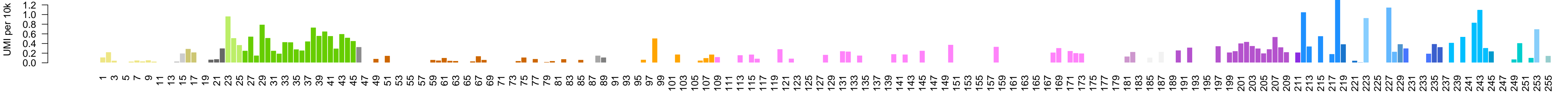
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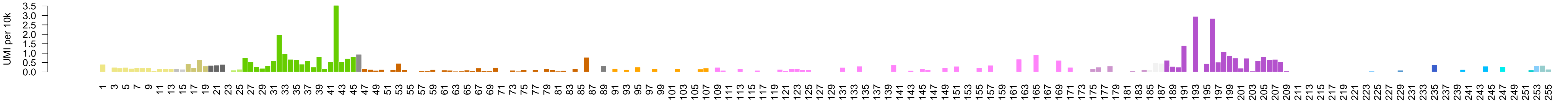
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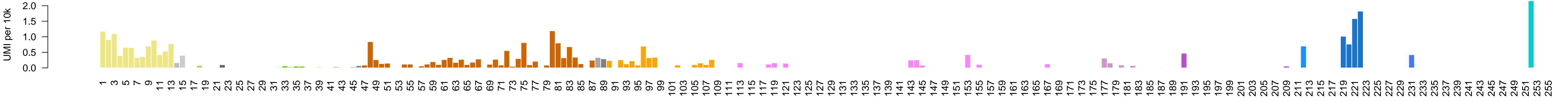
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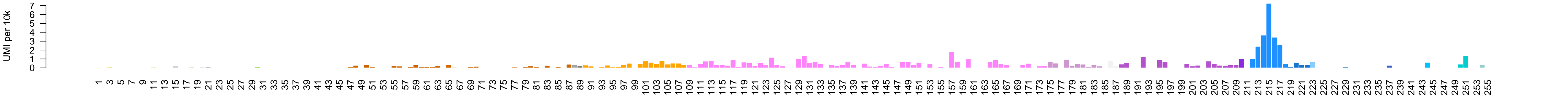
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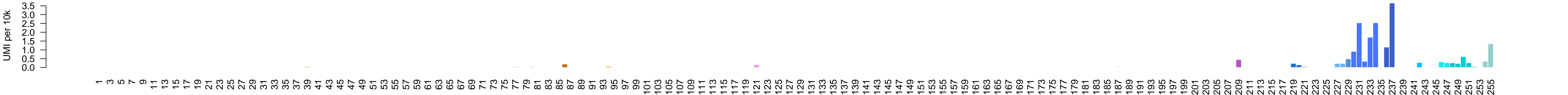
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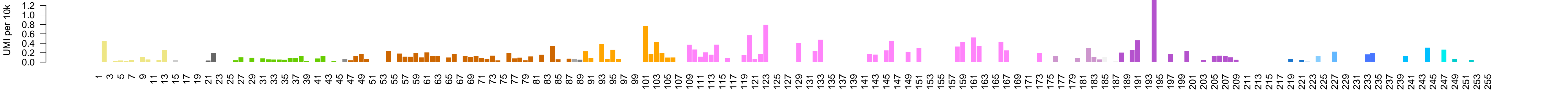
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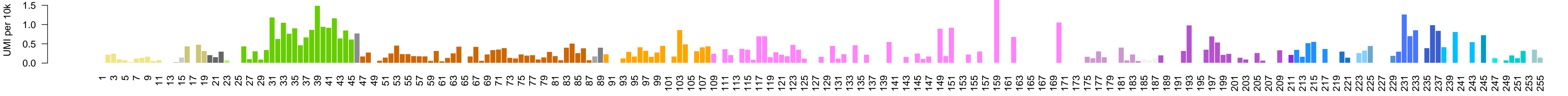
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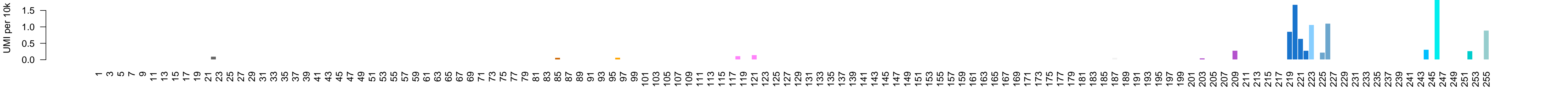


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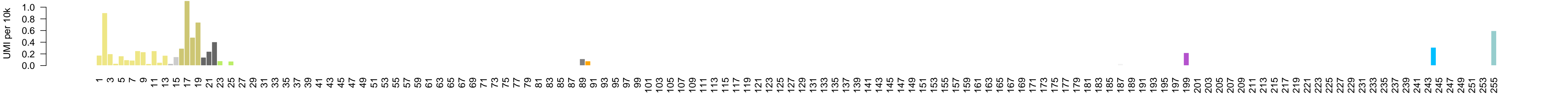


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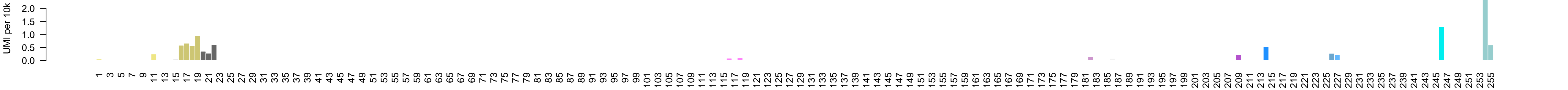
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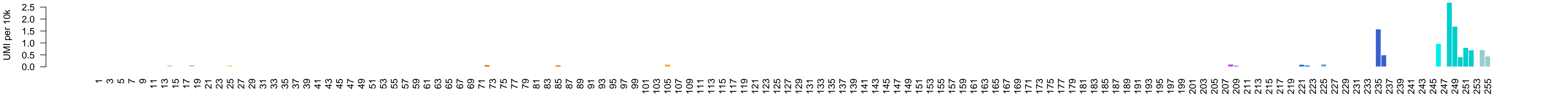
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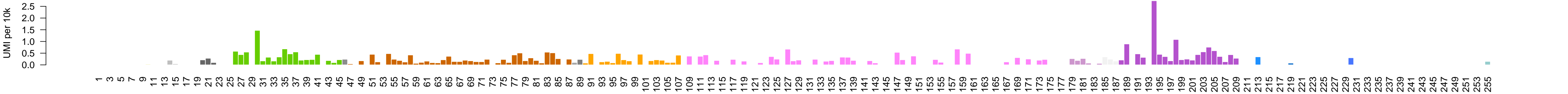
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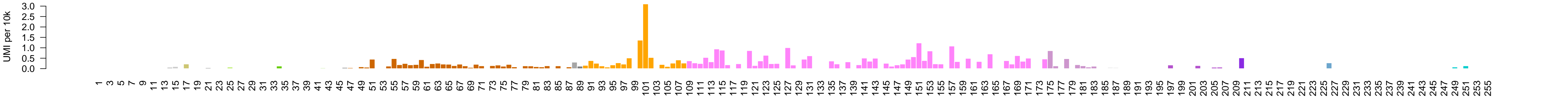
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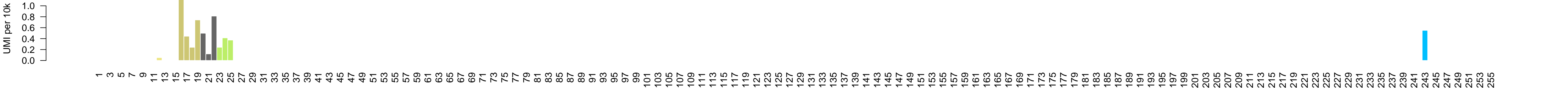
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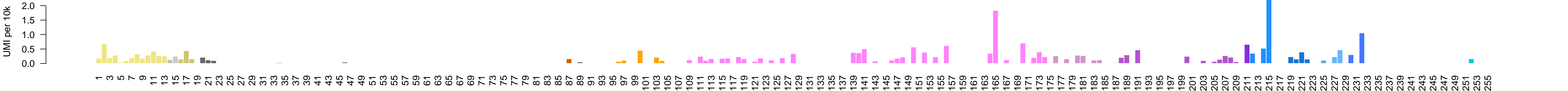
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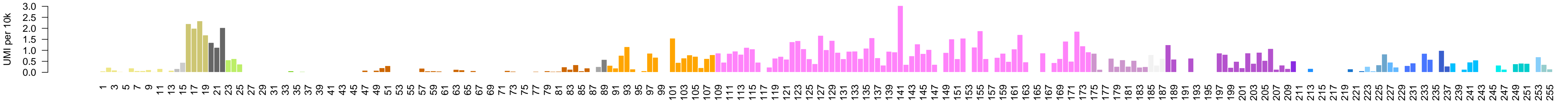
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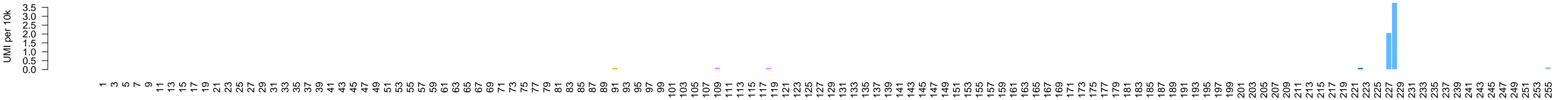
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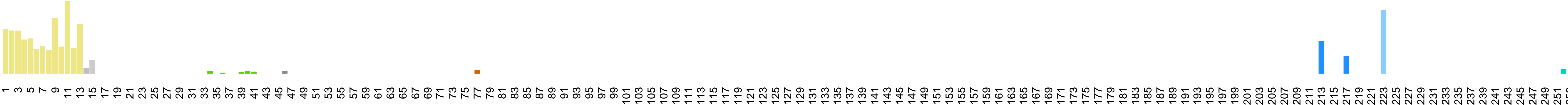
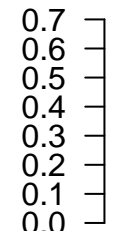


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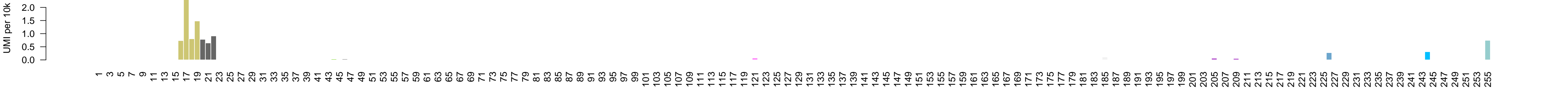


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UMI per 10k

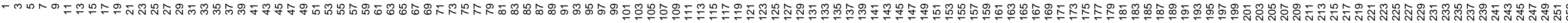
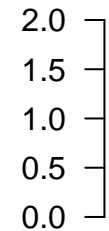


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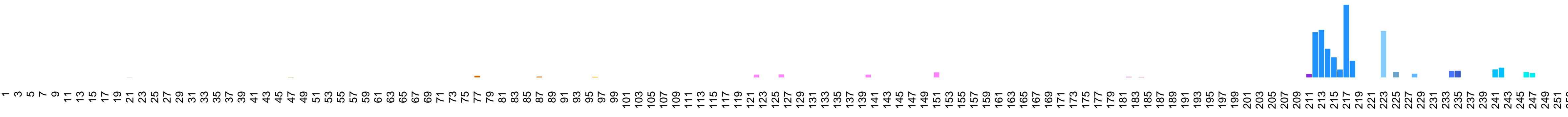
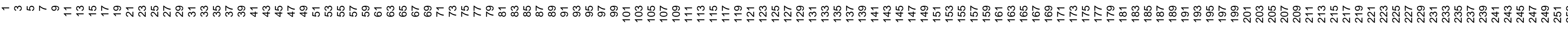
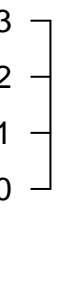
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UMI per 10k

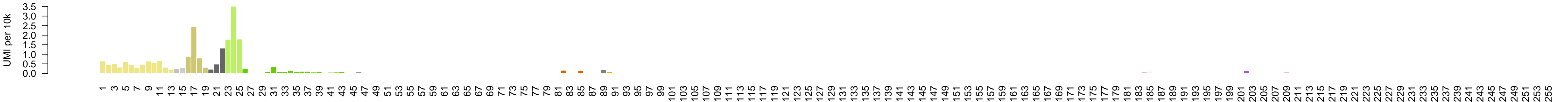


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UMI per 10k



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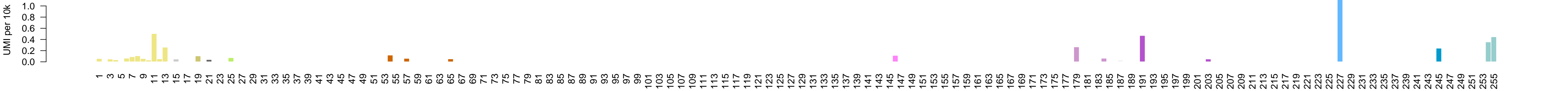
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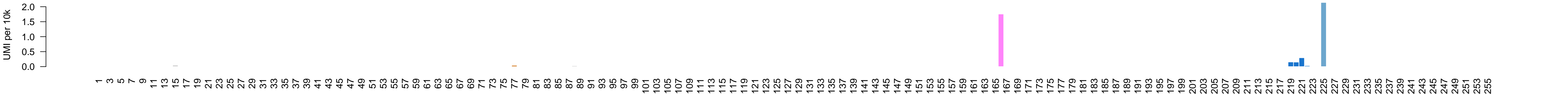
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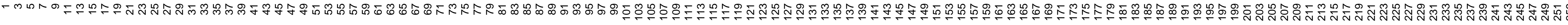
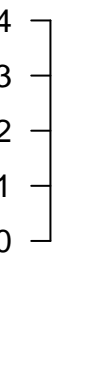


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Hhon_g03029.t1



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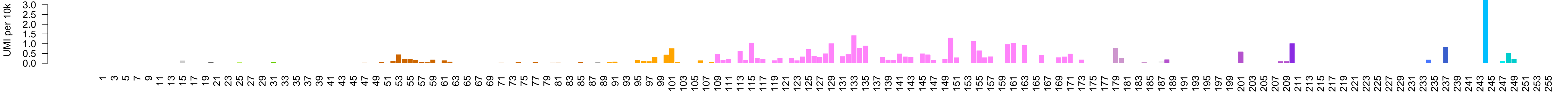
UMI per 10k



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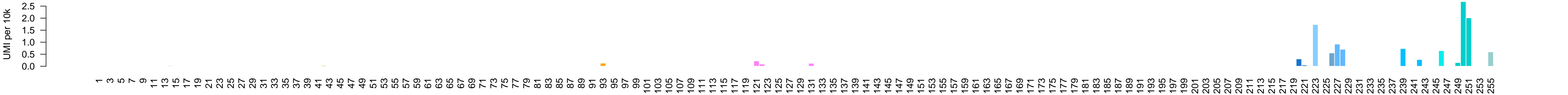
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Hhon_g04628.t1



ANF_receptor.HG2.10:like:GABBR1/GABBR2
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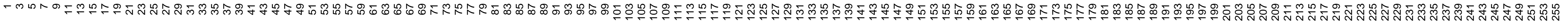
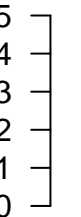


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Hhon_g05486.t1



ANF_receptor.HG2.10:like:GABBR1/GABBR2
Hhon_g11350.t1

UMI per 10k



ANF_receptor.HG2.11:like:GABBR1/GABBR2
Hhon_g07928.t1

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 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189 191 193 195 197 199 201 203 205 207 209 211 213 215 217 219 221 223 225 227 229 231 233 235 237 239 241 243 245 247 249 251 253 255

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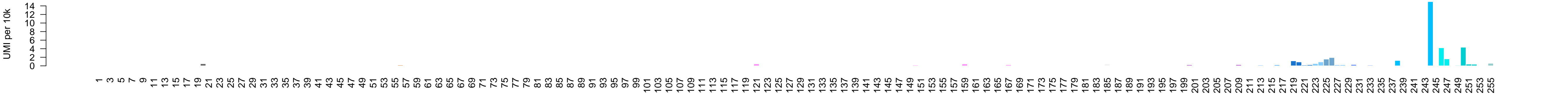
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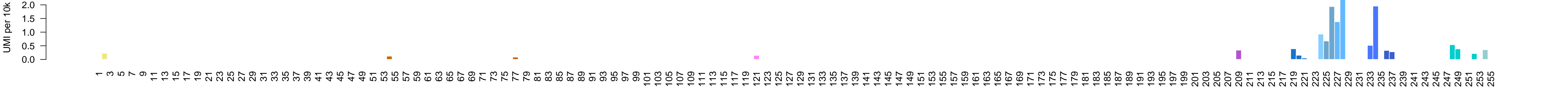
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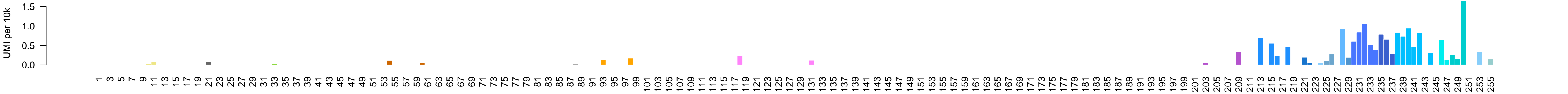
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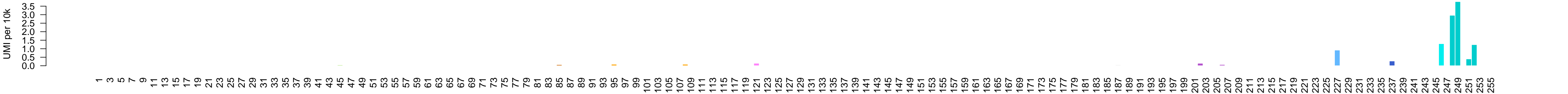
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Hhon_g09726.t1



ANF_receptor.HG2.14:like:GABBR1/GABBR2
Hhon_g05112.t1



ANF_receptor.HG2.16:like:GABBR1/GABBR2
Hhon_g02705.t1



ANF_receptor.HG2.17:like:GABBR1/GABBR2
Hhon_g02704.t1

UMI per 10k

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UMI per 10k

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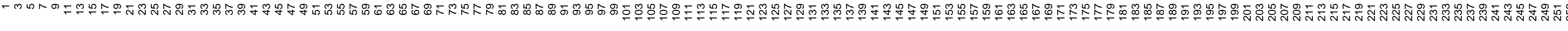
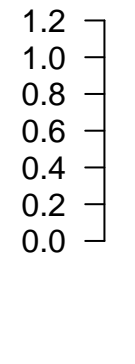
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ANF_receptor.HG2.19:like:GABBR1/GABBR2

Hhon_g02707.t1

UMI per 10k



ANF_receptor.HG2.20:like:GABBR1/GABBR2
Hhon_g07966.t1

UMI per 10k

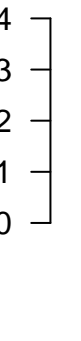
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1 3 5 7 9 11 13 15 17 19 21 23 25 27 29 31 33 35 37 39 41 43 45 47 49 51 53 55 57 59 61 63 65 67 69 71 73 75 77 79 81 83 85 87 89 91 93 95 97 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189 191 193 195 197 199 201 203 205 207 209 211 213 215 217 219 221 223 225 227 229 231 233 235 237 239 241 243 245 247 249 251 253 255



ANF_receptor.HG2.21:like:GABBR1/GABBR2
Hhon_g01249.t1

UMI per 10k



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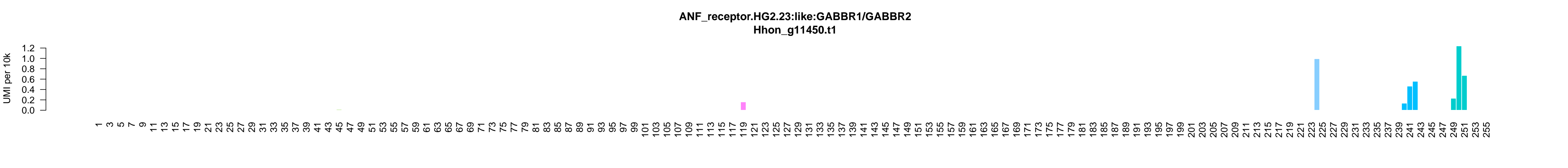
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Hhon_g01250.t1

UMI per 10k

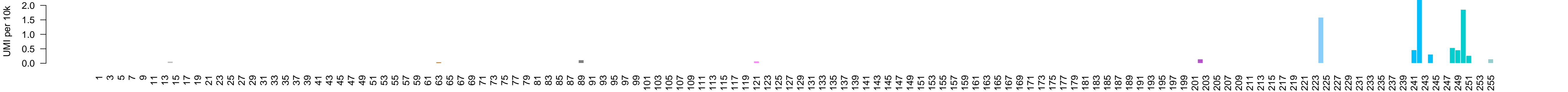
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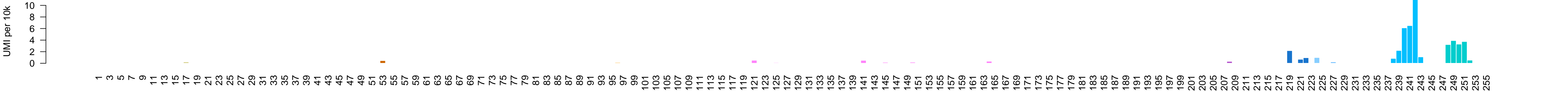




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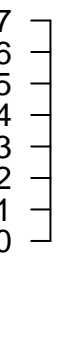


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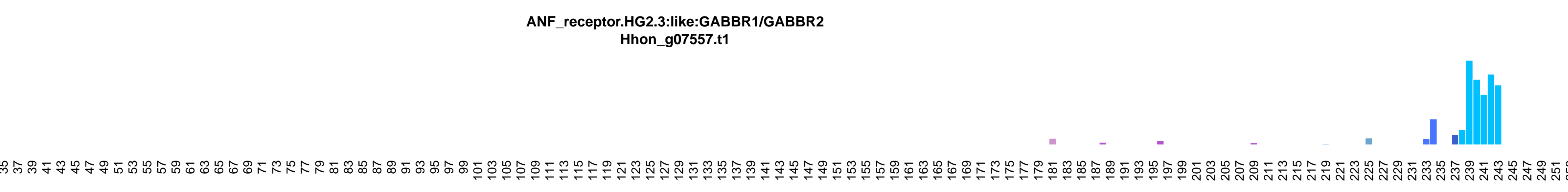


ANF_receptor.HG2.3:like:GABBR1/GABBR2
Hhon_g07557.t1

UMI per 10k



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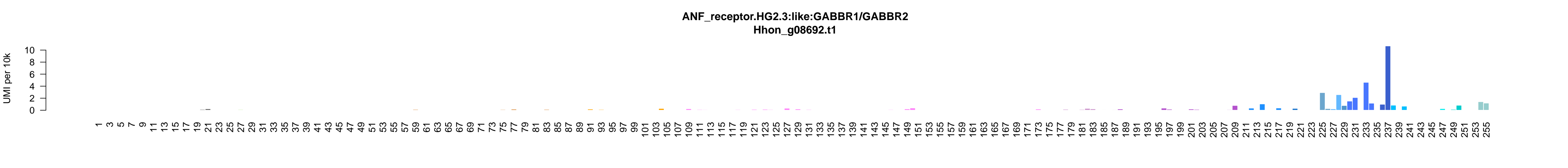


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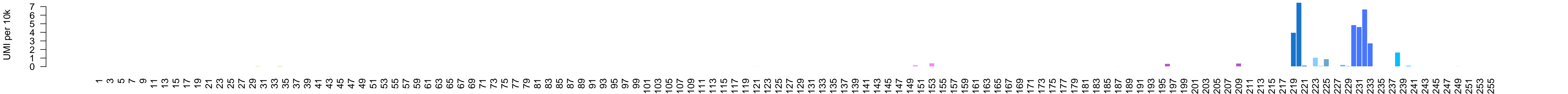
UMI per 10k

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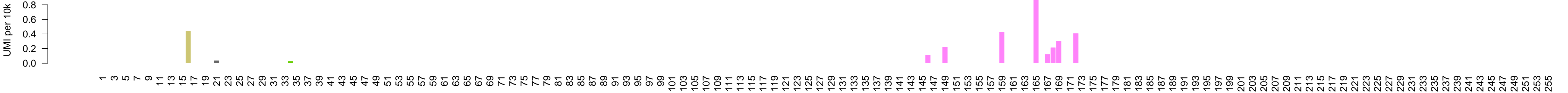
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ANF_receptor.HG2.3:like:GABBR1/GABBR2
Hhon_g08729.t1



ANF_receptor.HG2.3:like:GABBR1/GABBR2
Hhon_g10005.t1



ANF_receptor.HG2.3:like:GABBR1/GABBR2
Hhon_g10188.t1

UMI per 10k

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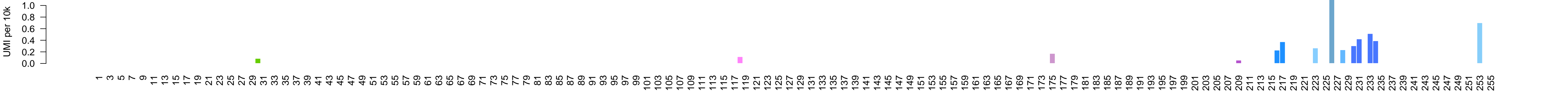
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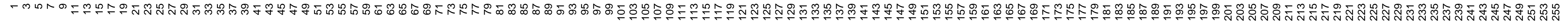
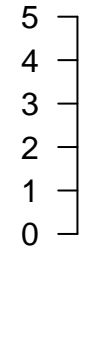
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Hhon_g10508.t1



ANF_receptor.HG2.3:like:GABBR1/GABBR2
Hhon_g10739.t1

UMI per 10k



ANF_receptor.HG2.3:like:GABBR1/GABBR2
Hhon_g10740.t1

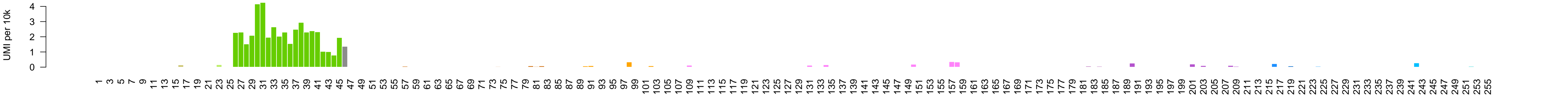
UMI per 10k

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ANF_receptor.HG2.30:GABBR2
Hhon_g06259.t1



ANF_receptor.HG2.31:like:GABBR1/GABBR2
Hhon_g07965.t1

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 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189 191 193 195 197 199 201 203 205 207 209 211 213 215 217 219 221 223 225 227 229 231 233 235 237 239 241 243 245 247 249 251 253 255

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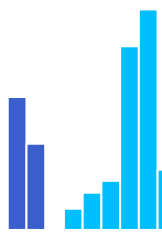
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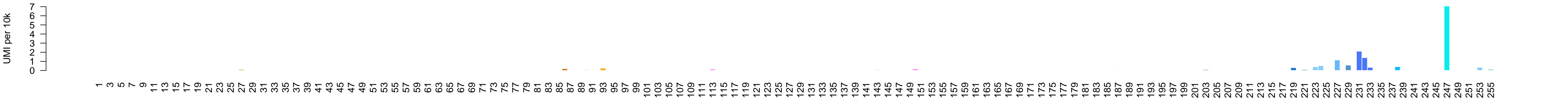
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Hhon_g07967.t1



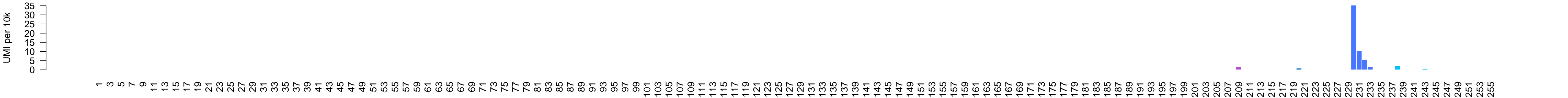
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Hhon_g09350.t1



ANF_receptor.HG2.4:like:GABBR1/GABBR2
Hhon_g09351.t1



ANF_receptor.HG2.4:like:GABBR1/GABBR2
Hhon_g09352.t1

UMI per 10k

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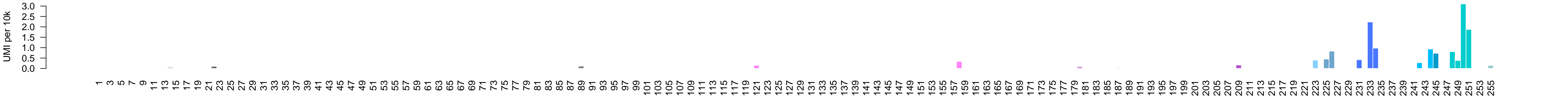
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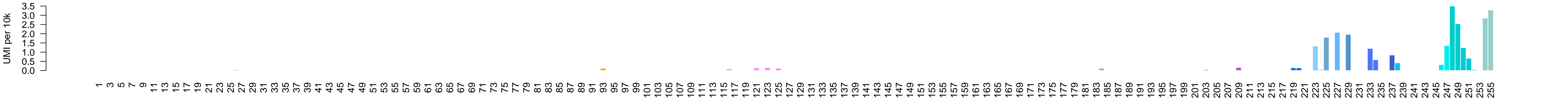
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Hhon_g01977.t1



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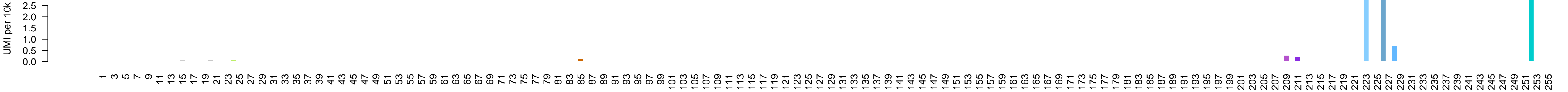
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ANF_receptor.HG2.5:like:GABBR1/GABBR2
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ANF_receptor.HG2.5:like:GABBR1/GABBR2
Hhon_g05298.t1

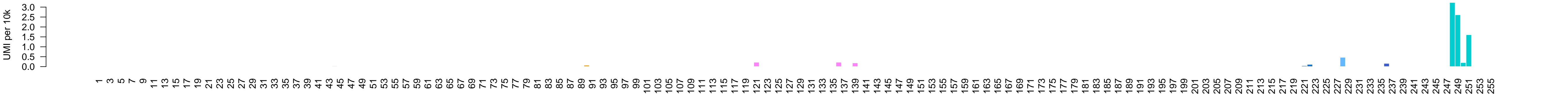
UMI per 10k

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ANF_receptor.HG2.5:like:GABBR1/GABBR2
Hhon_g05728.t1



ANF_receptor.HG2.5:like:GABBR1/GABBR2
Hhon_g08349.t1

UMI per 10k

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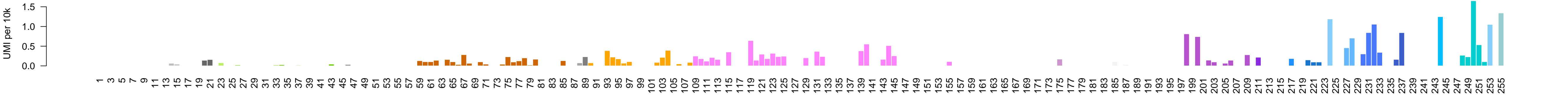
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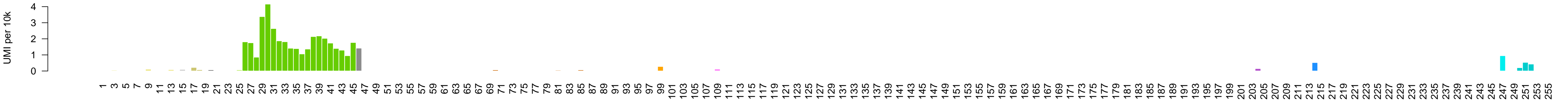
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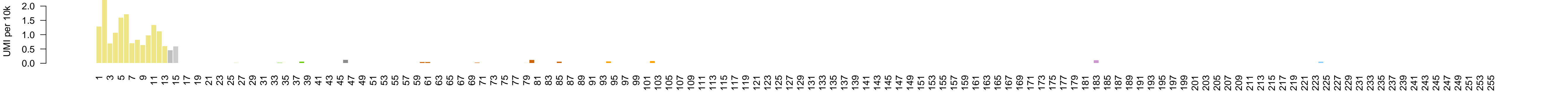
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ANF_receptor.HG2.6:like:GABBR1/GABBR2
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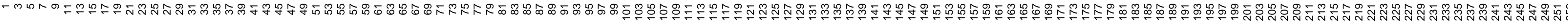
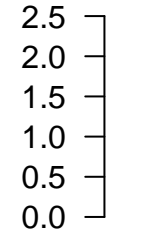
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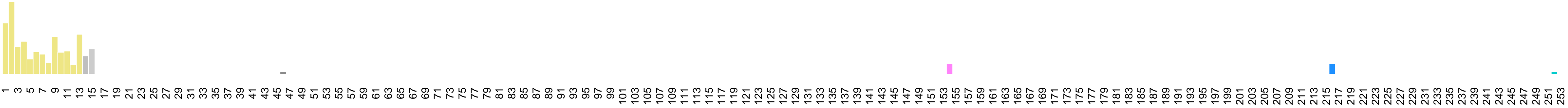
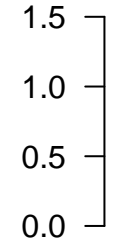
UMI per 10k



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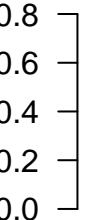
UMI per 10k



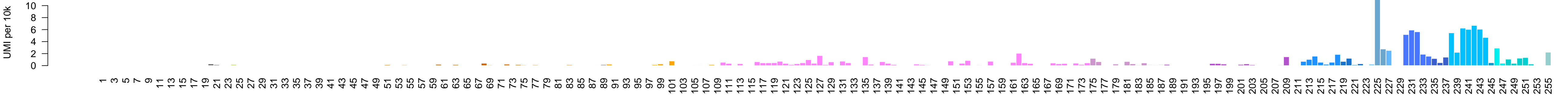
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Hhon_g02972.t1

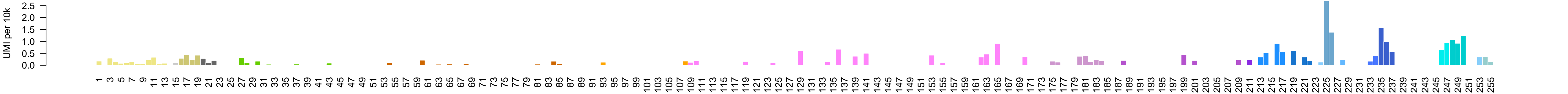
UMI per 10k



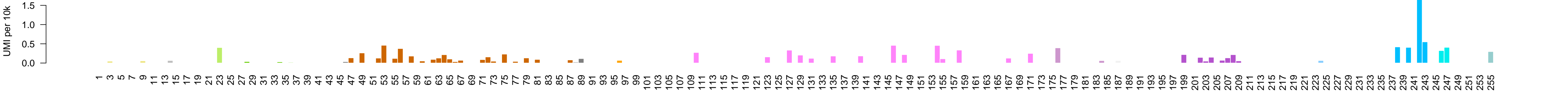
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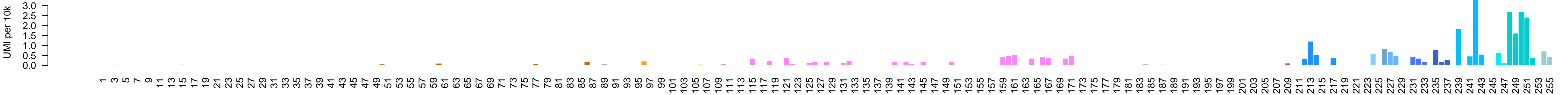
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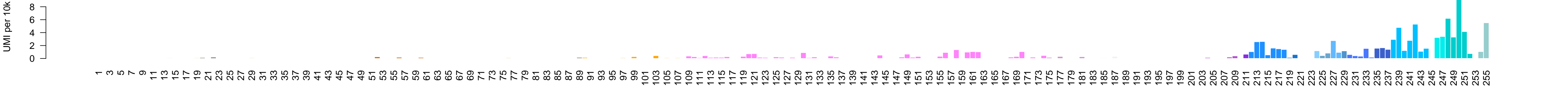
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Hhon_g01654.t1



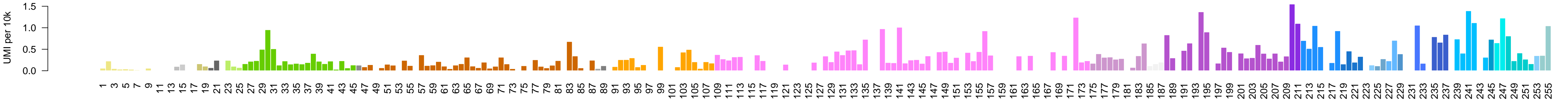
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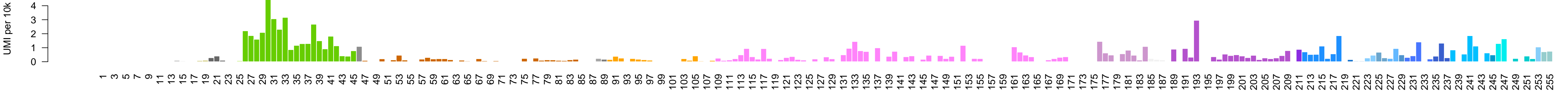
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Hhon_g04948.t1



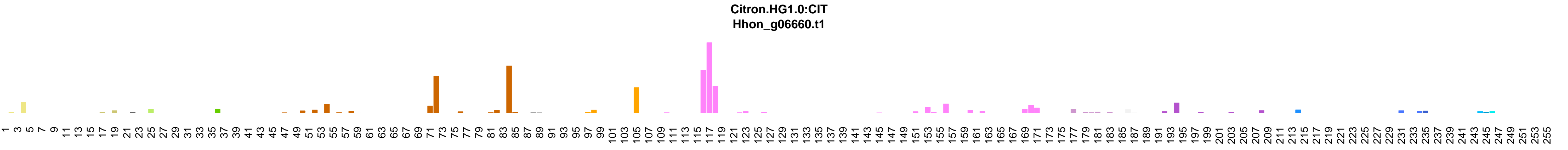
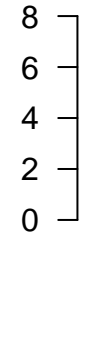
Caps.HG1.0:CADPS/CADPS2
Hhon_g02164.t1



Caps.HG1.1:UNC13A/UNC13B/UNC13C;Synaptotagmin.HG1.66:UNC13A/UNC13C;Unc13.HG1.1:UNC13A/UNC13B/UNC13C
Hhon_g00633.t1

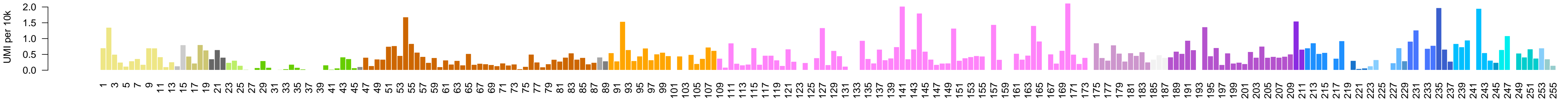


UMI per 10k

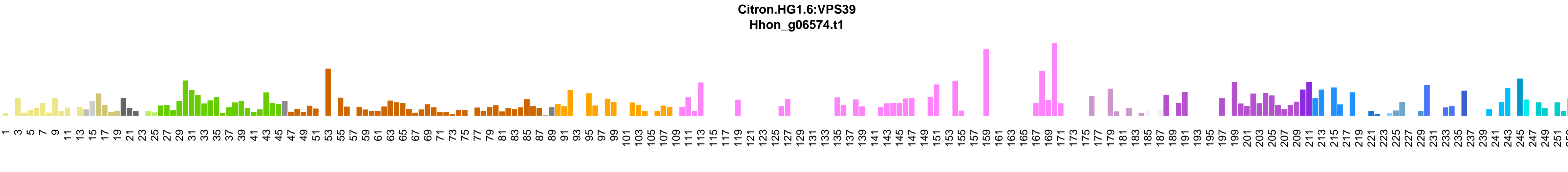
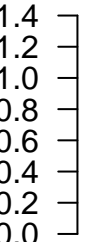


Citron.HG1.0:CIT
Hhon_g06660.t1

Citron.HG1.1: CDC42BPA/CDC42BPB/CDC42BPG
Hhon_g06582.t1

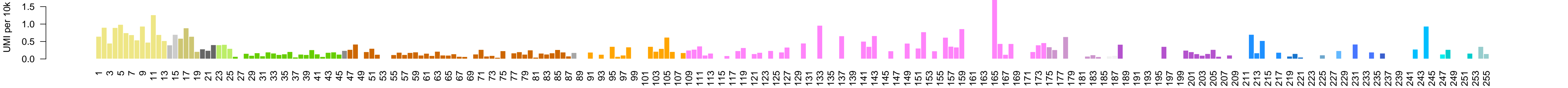


UMI per 10k



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

Hhon_g09097.t1



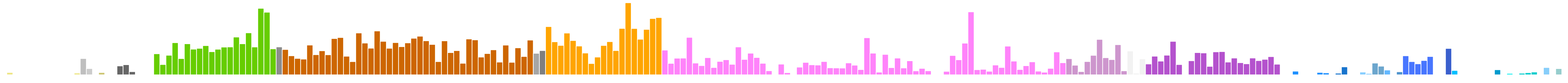
COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG

Hhon_g01143.t1

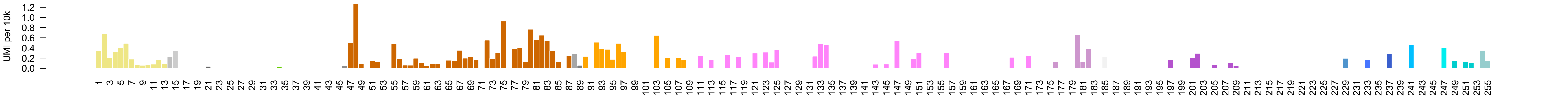
UMI per 10k

6
5
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 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189 191 193 195 197 199 201 203 205 207 209 211 213 215 217 219 221 223 225 227 229 231 233 235 237 239 241 243 245 247 249 251 253 255

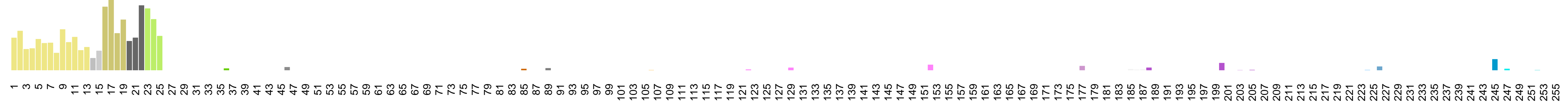
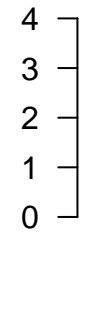


COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Hhon_g01144.t1

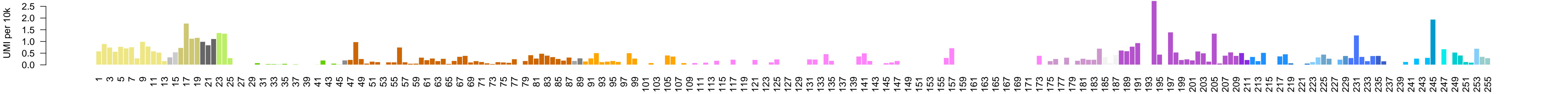


COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Hhon_g01145.t1

UMI per 10k



COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Hhon_g01146.t1



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

Hhon_g01147.t1

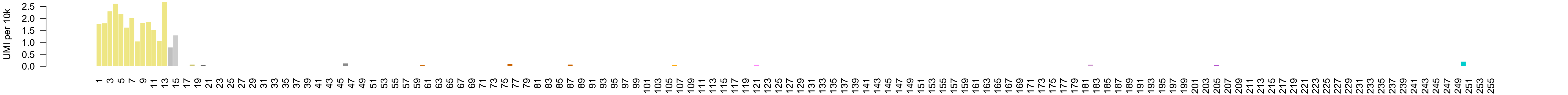


COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Hhon_g01148.t1



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

Hhon_g01149.t1

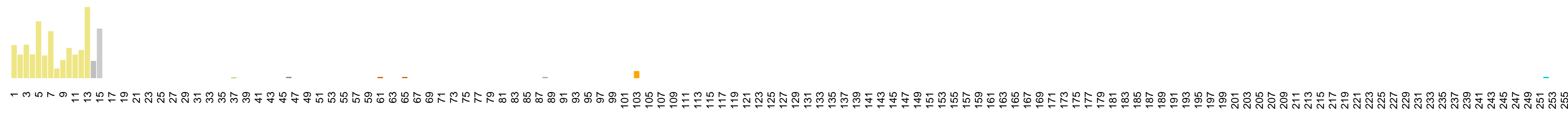


COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG

Hhon_g01150.t1

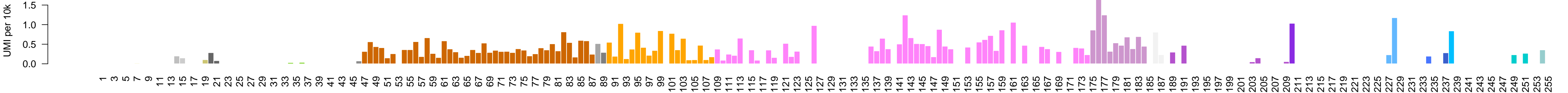
UMI per 10k

2.0
1.5
1.0
0.5
0.0



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

Hhon_g01151.t1



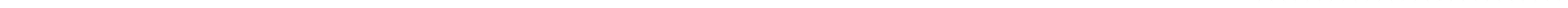
COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG

Hhon_g01153.t1

UMI per 10k

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 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189 191 193 195 197 199 201 203 205 207 209 211 213 215 217 219 221 223 225 227 229 231 233 235 237 239 241 243 245 247 249 251 253 255



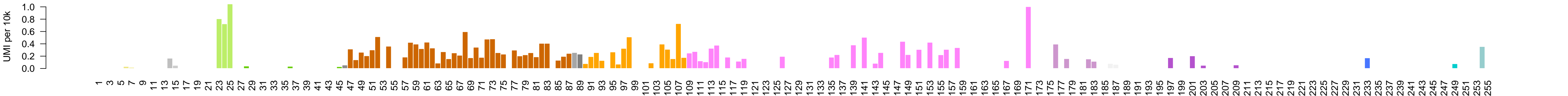
COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG

Hhon_g01156.t1



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

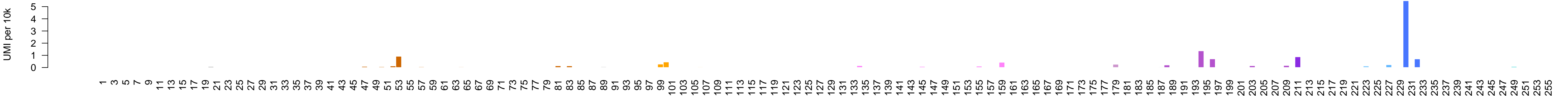
Hhon_g01157.t1



COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Hhon_g01159.t1



COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Hhon_g01160.t1



COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG
Hhon_g01161.t1



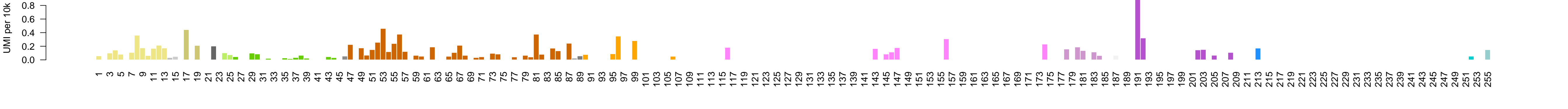
COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG

Hhon_g01162.t1



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

Hhon_g03041.t1



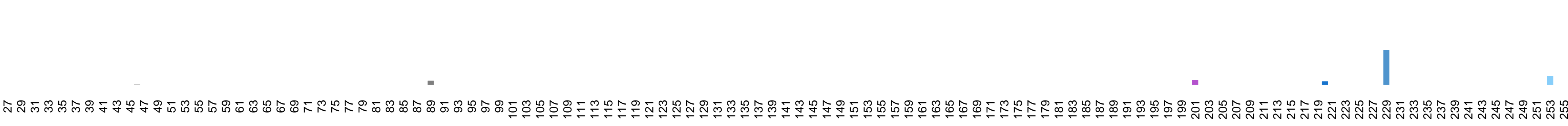
COesterase.HG1.16:like:ACHE/BCHE/CEL/CES1/CES2/CES3/CES4A/CES5A/TG

Hhon_g09790.t1

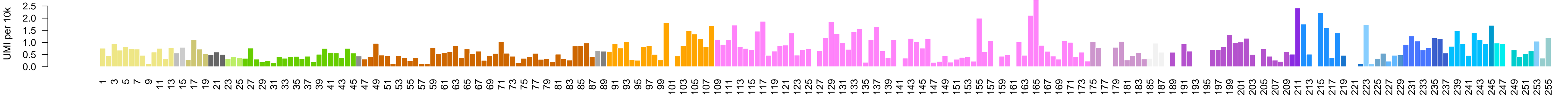
UMI per 10k

2.5
2.0
1.5
1.0
0.5
0.0

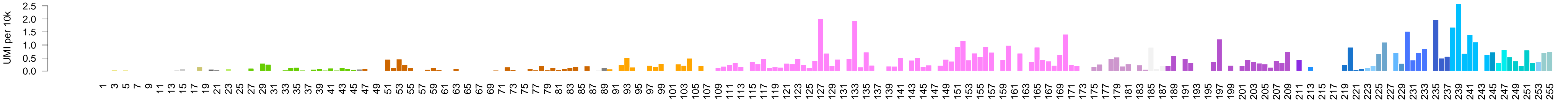
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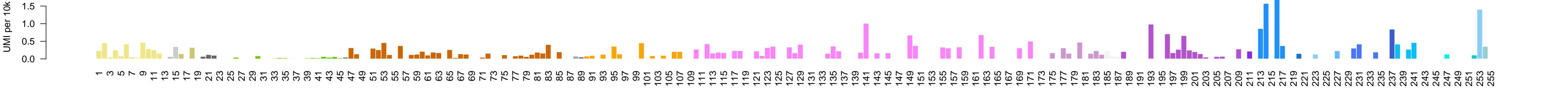
Cript.HG1.0:CRIP
Hhon_g05823.t1



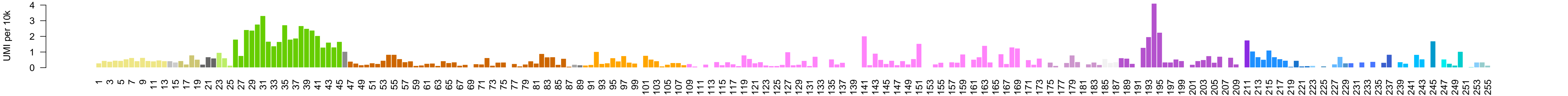
FERM_M.HG1.1:like:MYO10/PLEKHH1/PLEKHH2/PLEKHH3
Hhon_g09383.t1



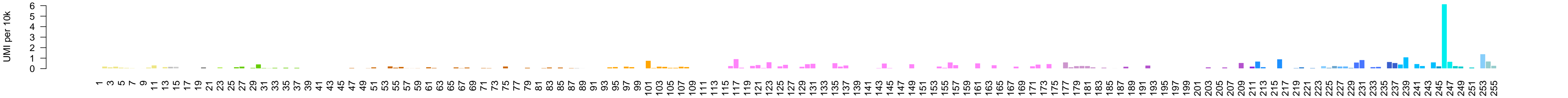
FERM_M.HG1.14:like:MYO7A/MYO7B
Hhon_g04928.t1



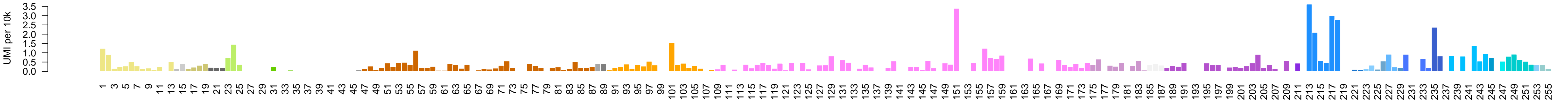
FERM_M.HG1.18:TLN1/TLN2
Hhon_g04057.t1



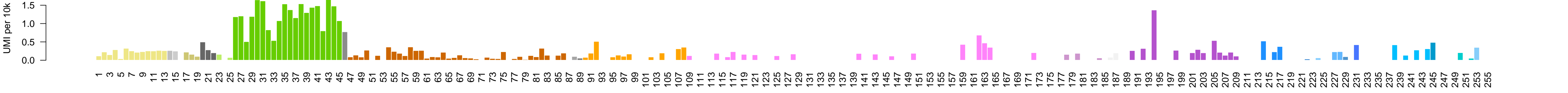
FERM_M.HG1.22:PTPN14/PTPN21;Y_phosphatase.HG1.37:PTPN14/PTPN21
Hhon_g10875.t1



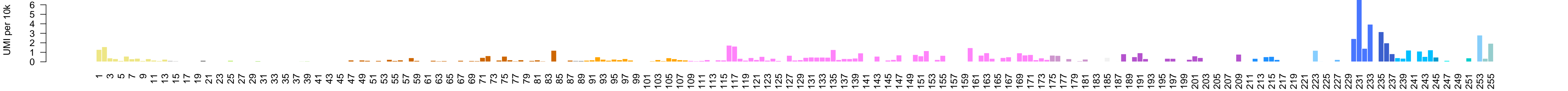
FERM_M.HG1.24:FRMD1/FRMD6/FRMPD2/PTPN13
Hhon_g03085.t1



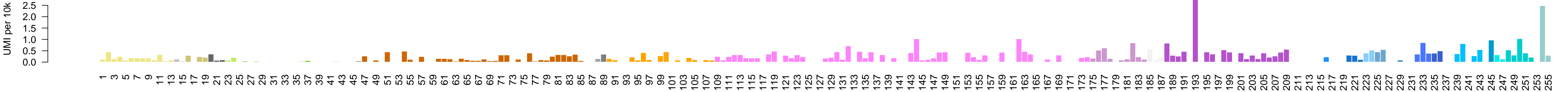
FERM_M.HG1.26:FERMT1/FERMT2/FERMT3
Hhon_g03113.t1



FERM_M.HG1.3:PLEKHH1/PLEKHH2
Hhon_g09776.t1

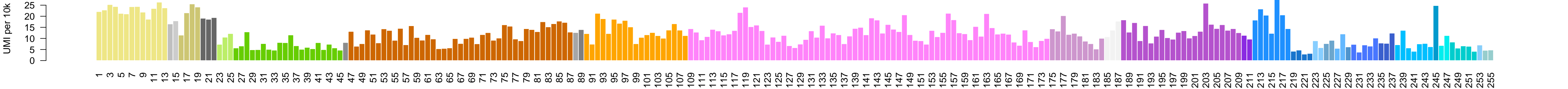


FERM_M.HG1.30:FRMD4A/FRMD4B
Hhon_g05657.t1



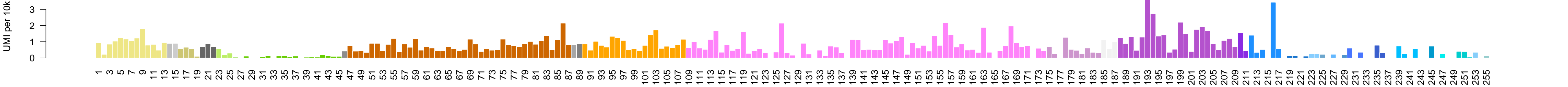
FERM_M.HG1.32:EPB41/EPB41L1/EPB41L2/EPB41L3/EPB41L4A/EPB41L4B/EPB41L5/FARP1/FARP2/FRMD3/FRMD5/FRMD7/PTPN3/PTPN4

Hhon_g00260.t1



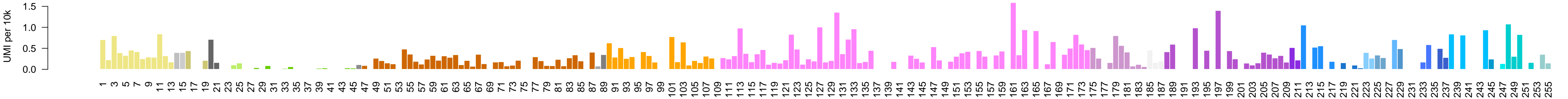
FERM_M.HG1.32:EPB41/EPB41L1/EPB41L2/EPB41L3/EPB41L4A/EPB41L4B/EPB41L5/FARP1/FARP2/FRMD3/FRMD5/FRMD7/PTPN3/PTPN4

Hhon_g06296.t1

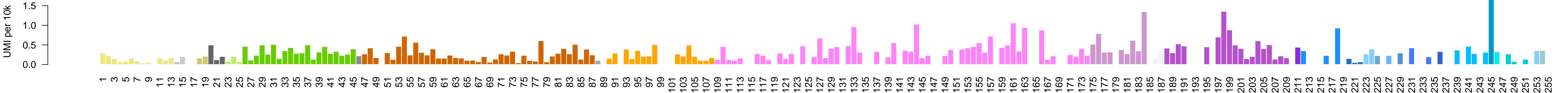


FERM_M.HG1.32:EPB41/EPB41L1/EPB41L2/EPB41L3/EPB41L4A/EPB41L4B/EPB41L5/FARP1/FARP2/FRMD3/FRMD5/FRMD7/PTPN3/PTPN4

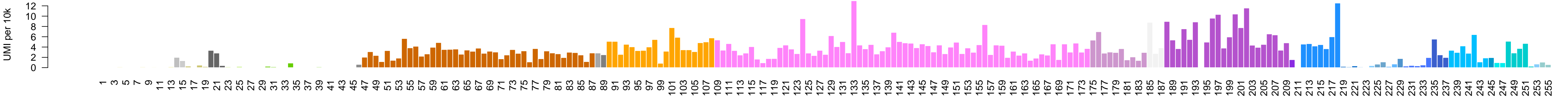
Hhon_g08340.t1



FERM_M.HG1.32:EPB41/EPB41L1/EPB41L2/EPB41L3/EPB41L4A/EPB41L4B/EPB41L5/FARP1/FARP2/FRMD3/FRMD5/FRMD7/PTPN3/PTPN4;Y_phosphatase.HG1.38:PTPN3/PTPN4
Hhon_g06298.t1



FERM_M.HG1.33:like:EZR/MSN/NF2/RDX
Hhon_g00802.t1

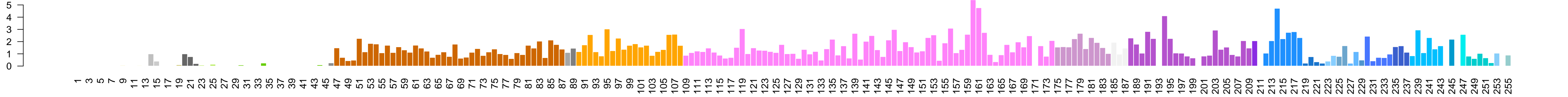


FERM_M.HG1.4:MYO10/PLEKHH3
Hhon_g02820.t1

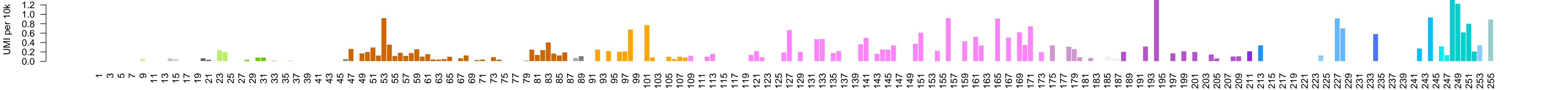


FERM_M.HG1.40:EZR/MSN/RDX
Hhon_g05421.t1

UMI per 10k

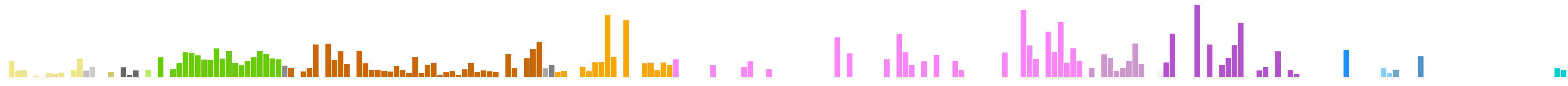
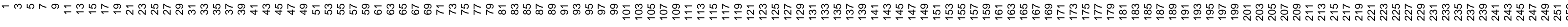
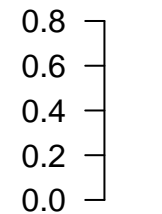


FERM_M.HG1.7:like:FERMT1/FERMT2/FERMT3/TLN1/TLN2
Hhon_g08184.t1

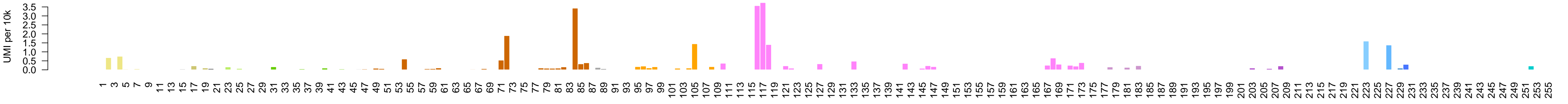


FERM_M.HG2.1:FRMPD1/FRMPD3/FRMPD4
Hhon_g09737.t1

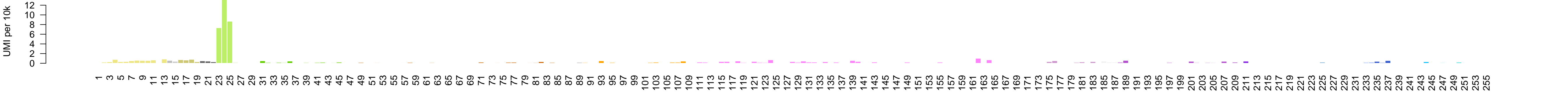
UMI per 10k



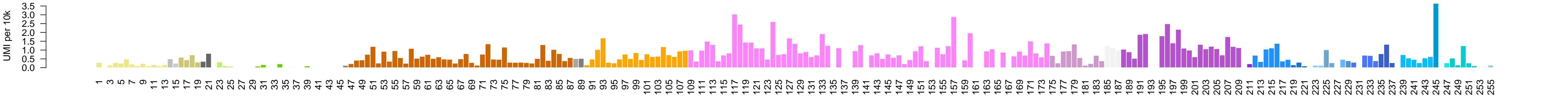
GKAP.HG1.0:DLGAP5
Hhon_g03573.t1



GKAP.HG1.1:DLGAP1/DLGAP2/DLGAP3/DLGAP4
Hhon_g07014.t1

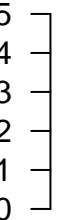


Guanylate_kin.HG1.0:MAGI1/MAGI2/MAGI3
Hhon_g05213.t1

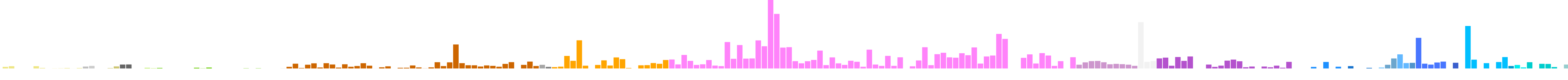


Guanylate_kin.HG1.12:LRGUK
Hhon_g06532.t1

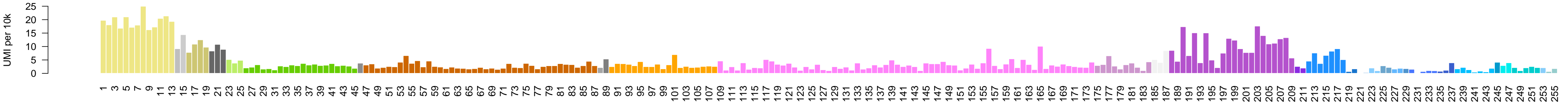
UMI per 10k



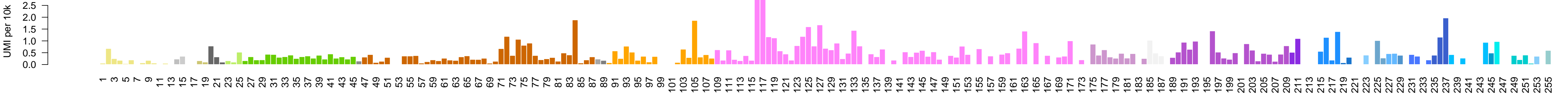
1 3 5 7 9 11 13 15 17 19 21 23 25 27 29 31 33 35 37 39 41 43 45 47 49 51 53 55 57 59 61 63 65 67 69 71 73 75 77 79 81 83 85 87 89 91 93 95 97 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189 191 193 195 197 199 201 203 205 207 209 211 213 215 217 219 221 223 225 227 229 231 233 235 237 239 241 243 245 247 249 251 253 255



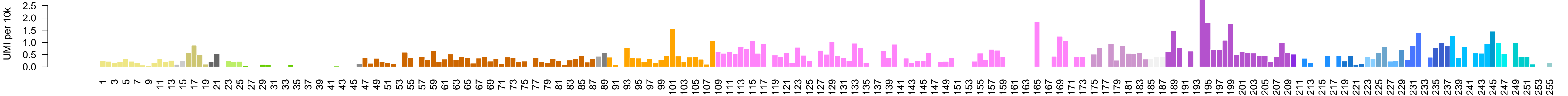
Guanylate_kin.HG1.14:DLG1/DLG2/DLG3/DLG4
Hhon_g03162.t1



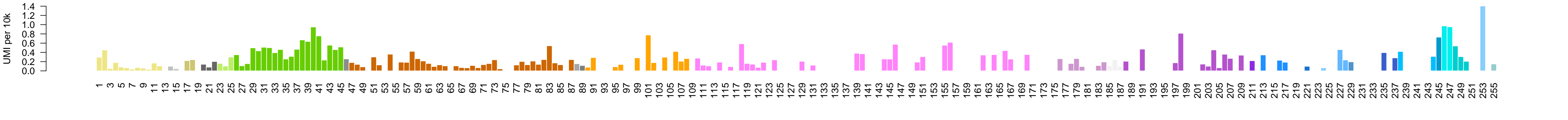
Guanylate_kin.HG1.16:GUK1
Hhon_g02406.t1



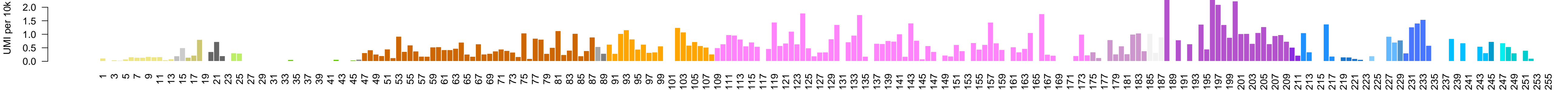
Guanylate_kin.HG1.2:like:AL358113.1/CASK/DLG1/DLG2/DLG3/DLG4/DLG5/GUK1/LRGUK/MPP1/MPP2/MPP3/MPP4/MPP5/MPP6/MPP7/TJP1/TJP2/TJP3;PDZ.HG2.11:MPP5
Hhon_g09855.t1



Guanylate_kin.HG1.20:CASK/MPP1;PDZ.HG2.8:CASK
Hhon_g05233.t1

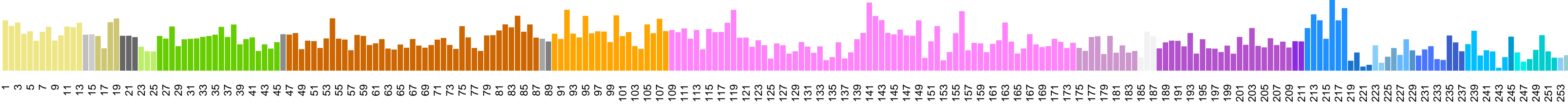
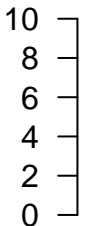


Guanylate_kin.HG1.3:MPP3/MPP4/MPP7;PDZ.HG2.10:like:MPP5
Hhon_g02443.t1

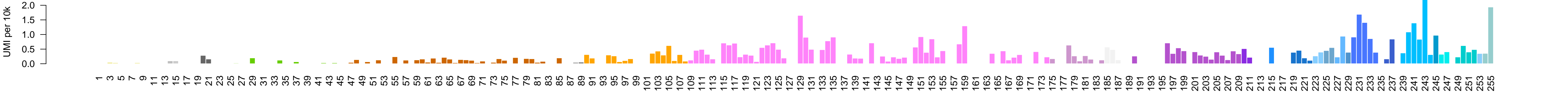


Guanylate_kin.HG1.9:like:CASK/MPP1/MPP2/MPP6;PDZ.HG2.5:MPP2/MPP6
Hhon_g02440.t1

UMI per 10k

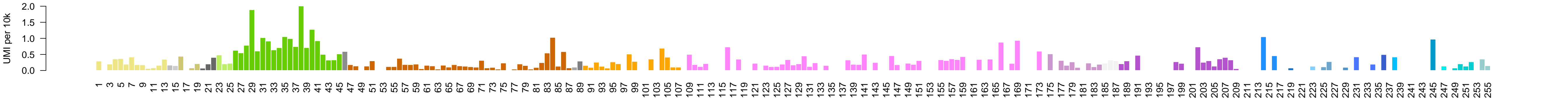


Guanylate_kin.HG2.0:CACNB1/CACNB2/CACNB3/CACNB4
Hhon_g00617.t1

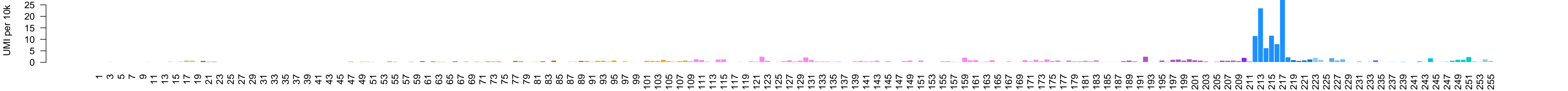


Homer.HG1.1:ENAH/EVL/SPRED1/SPRED2/SPRED3/VASP

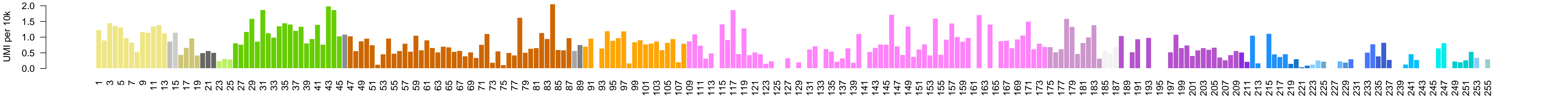
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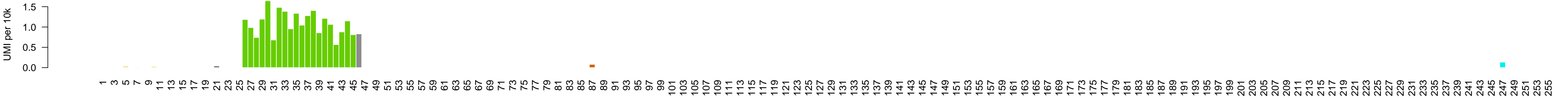
Homer.HG1.2:HOMER1/HOMER2/HOMER3
Hhon_g05897.t1



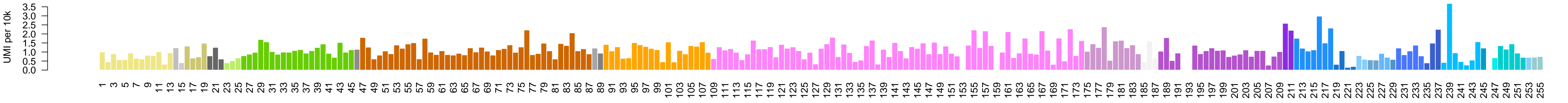
Homer.HG1.9:WAS/WASL
Hhon_g08647.t1



Homer.HG1.9:WAS/WASL
Hhon_g08649.t1



MARVEL.HG1.13:SYNPR/SYP/SYPL1/SYPL2
Hhon_g06148.t1



MARVEL.HG1.7:like:OCLN/SYNPR/SYP/SYPL1/SYPL2

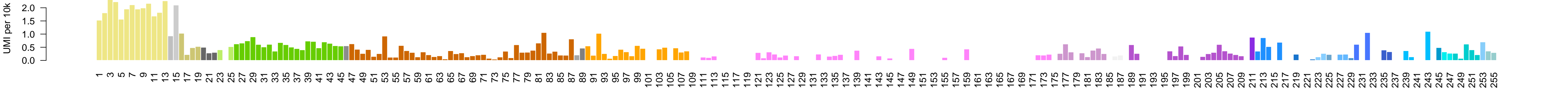
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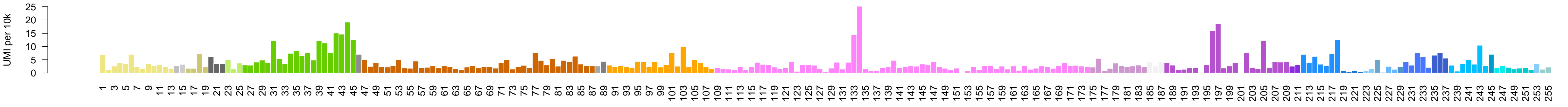
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MARVEL.HG2.0:SYNGR1/SYNGR2/SYNGR4
Hhon_g01440.t1



MARVEL.HG2.0:SYNGR1/SYNGR2/SYNGR4
Hhon_g01441.t1



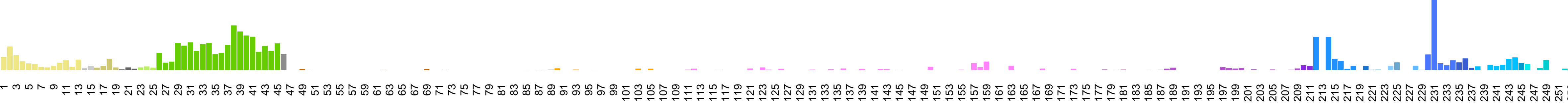
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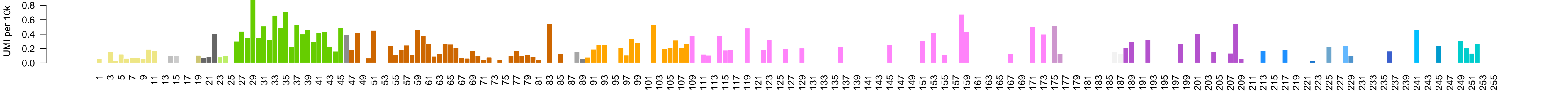
UMI per 10k

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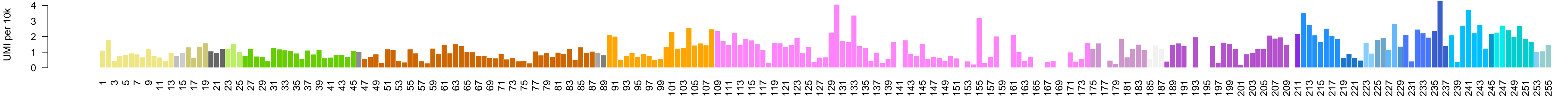
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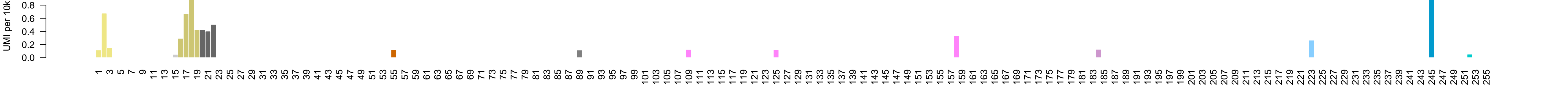
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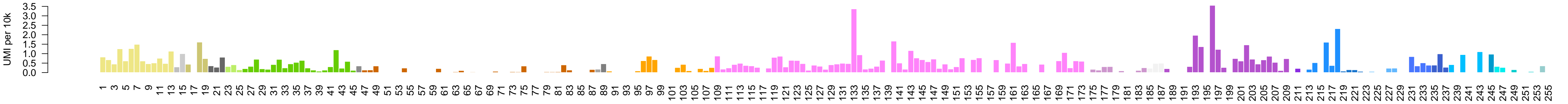
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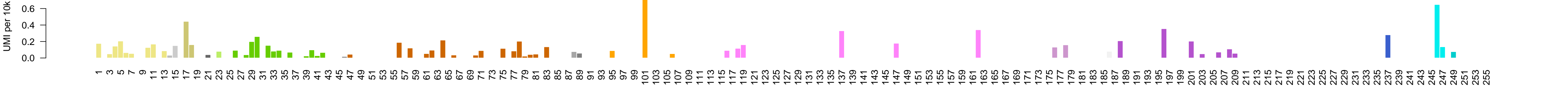
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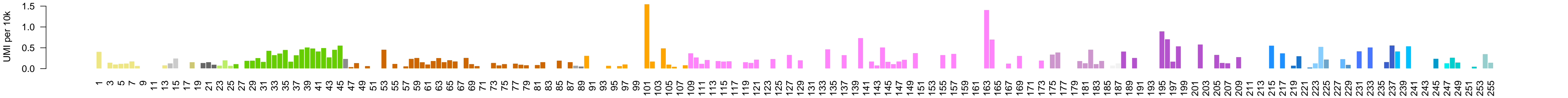
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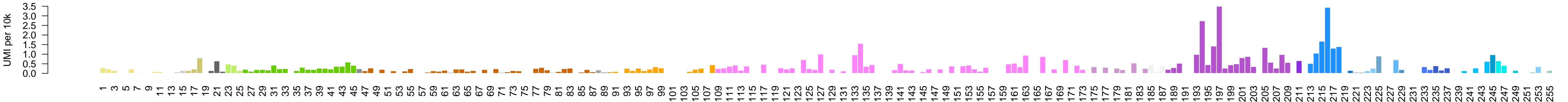
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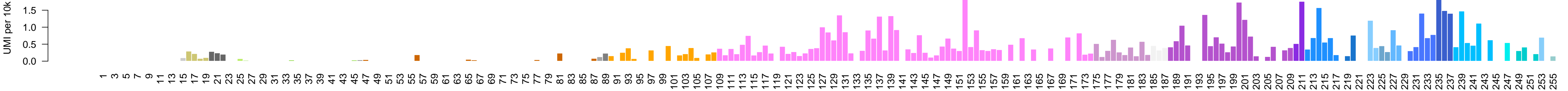
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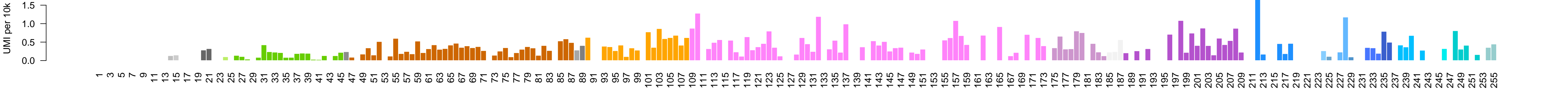
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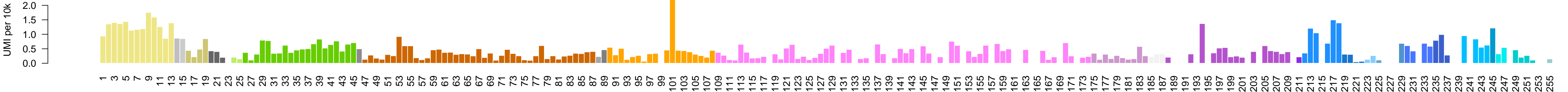
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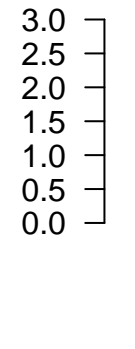


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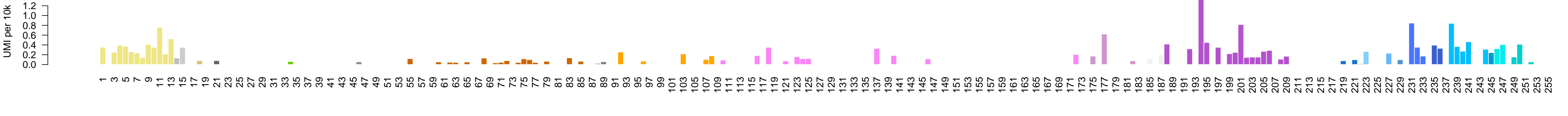
UMI per 10k



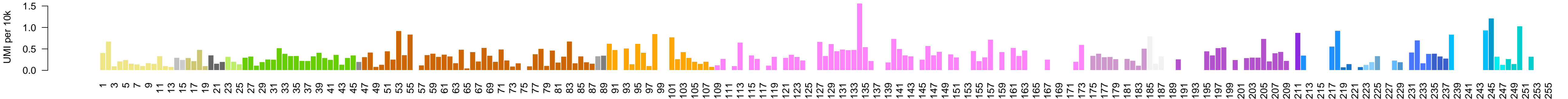
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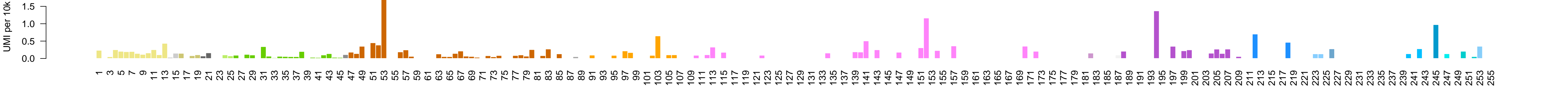
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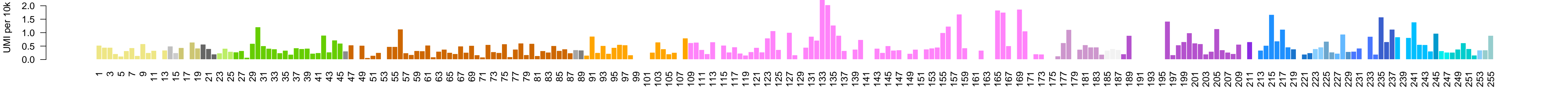
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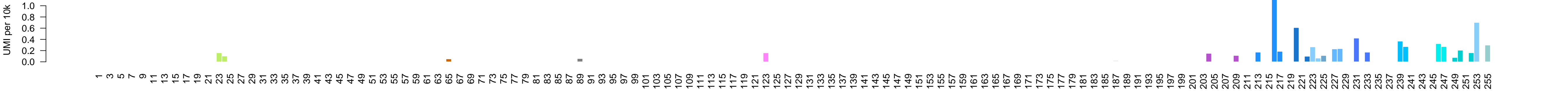
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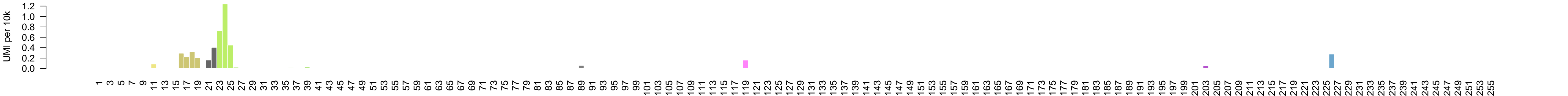
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MFS_1.HG15.1:SLC45A1/SLC45A2/SLC45A3
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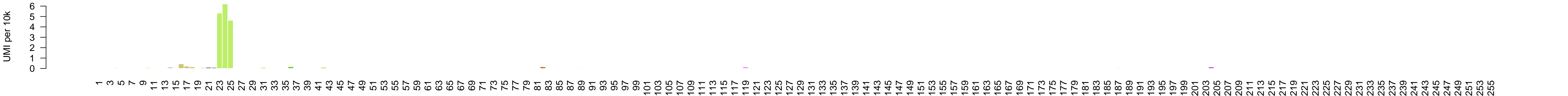
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MFS_1.HG15.1:SLC45A1/SLC45A2/SLC45A3
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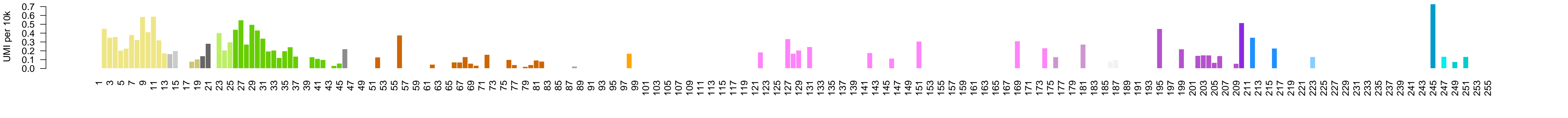
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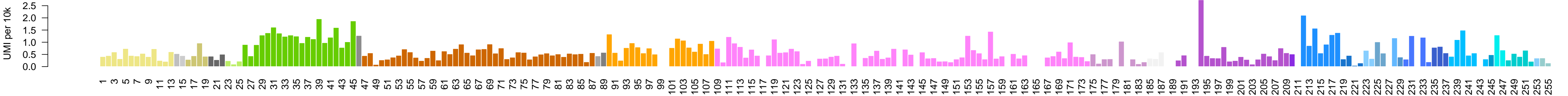
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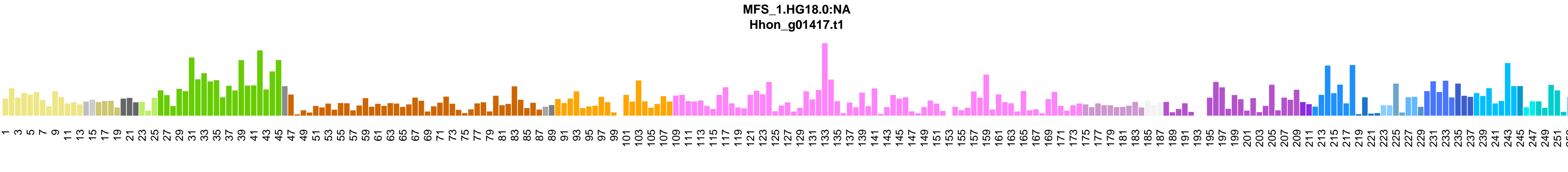


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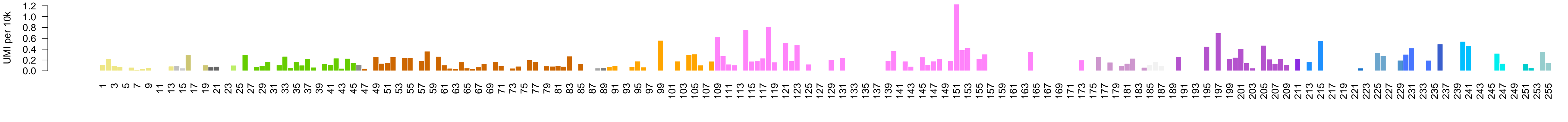


UMI per 10k

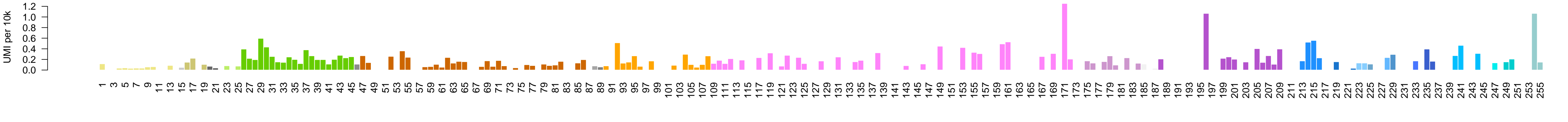
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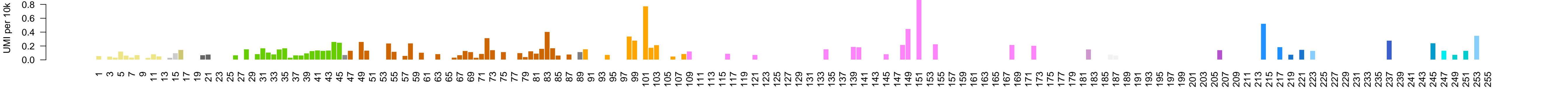
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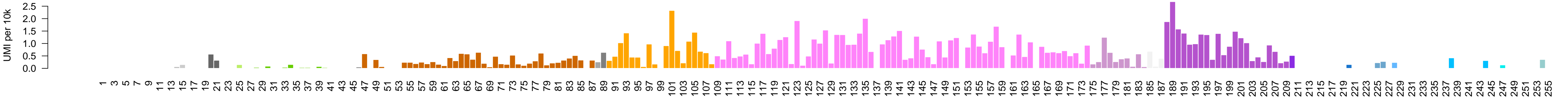
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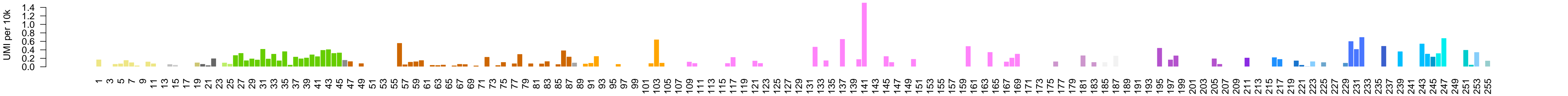
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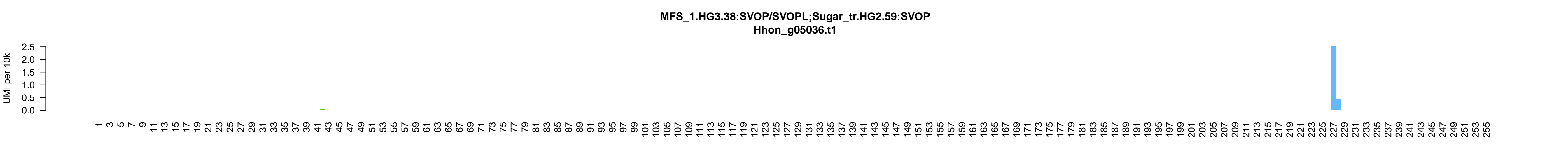


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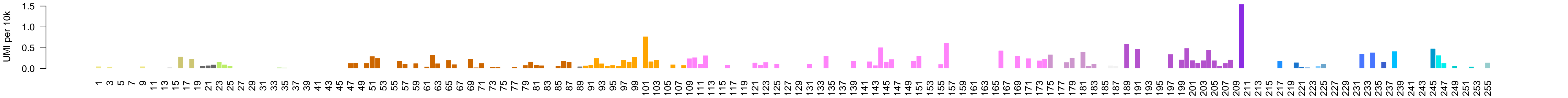


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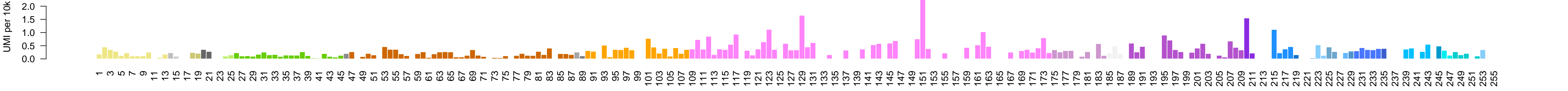




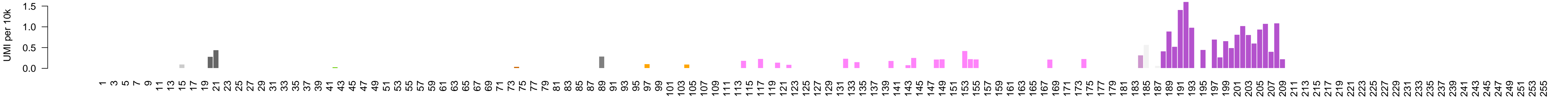
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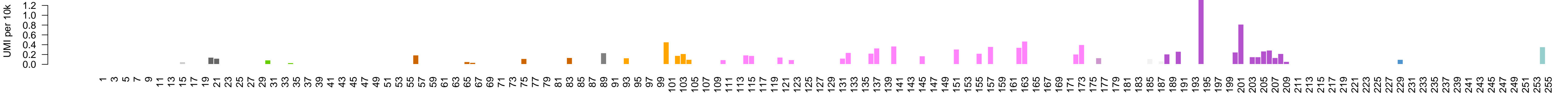
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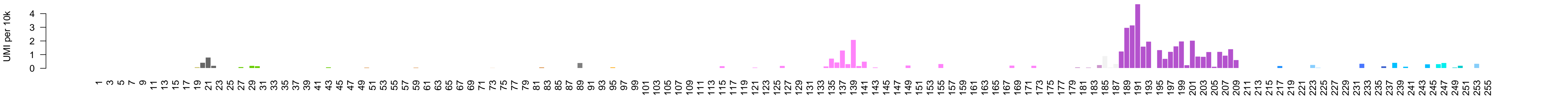
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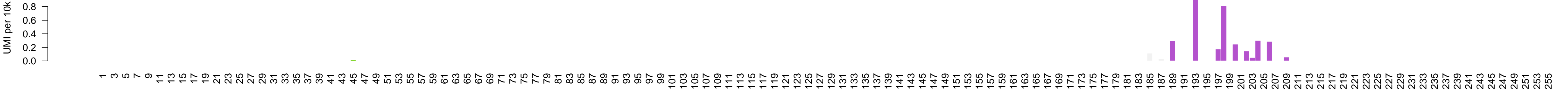
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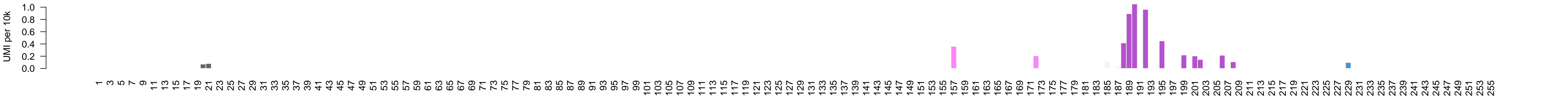
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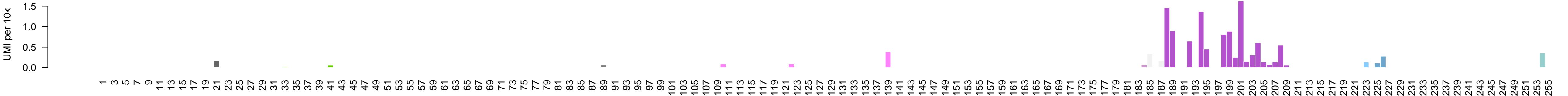
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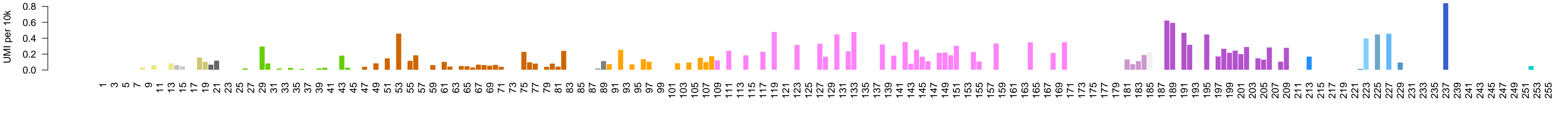
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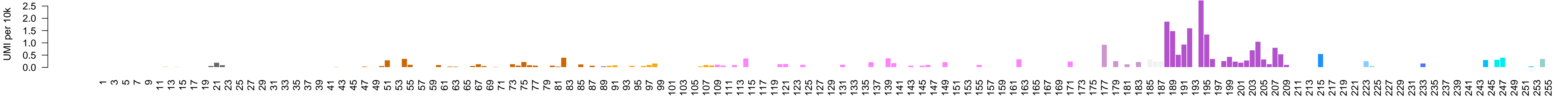
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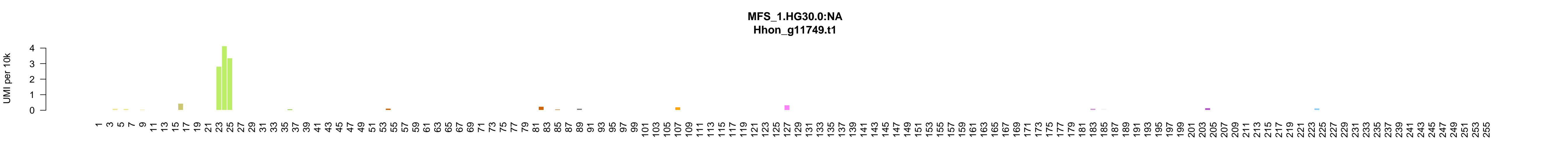


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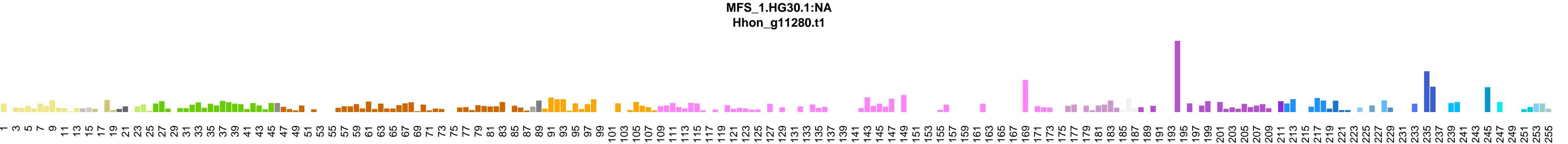
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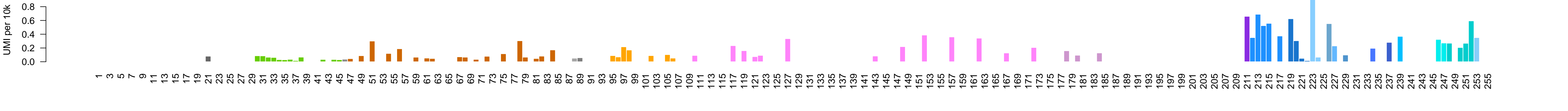


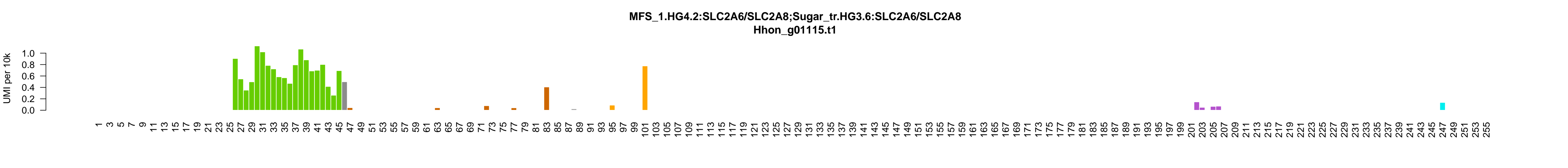
UMI per 10k

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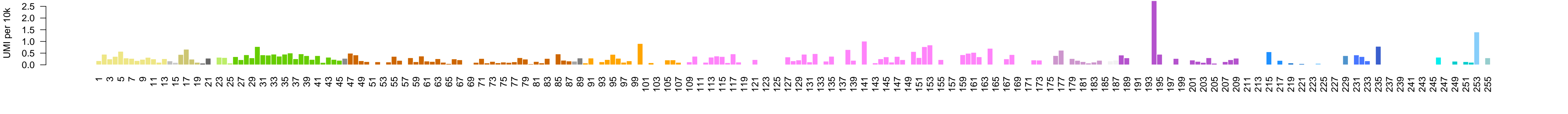


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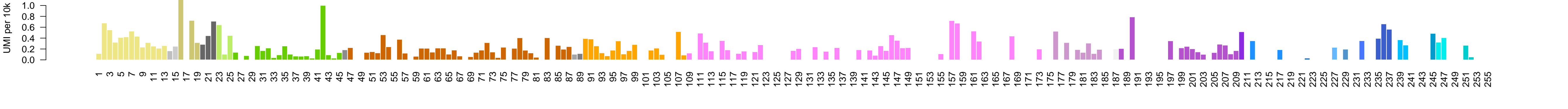




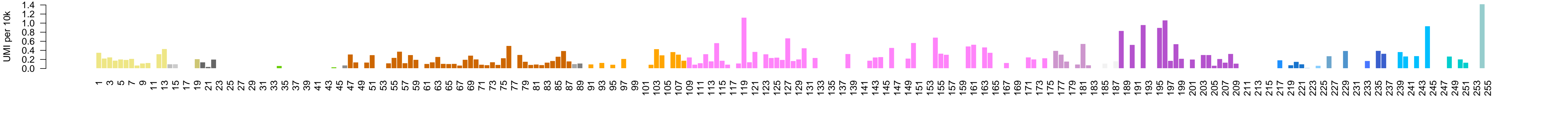
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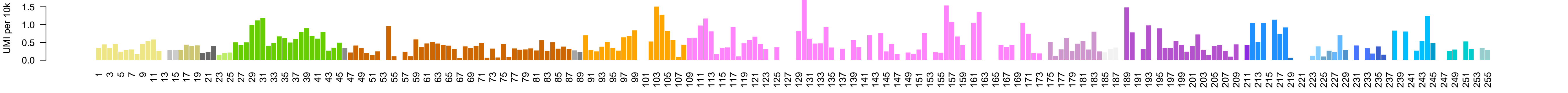
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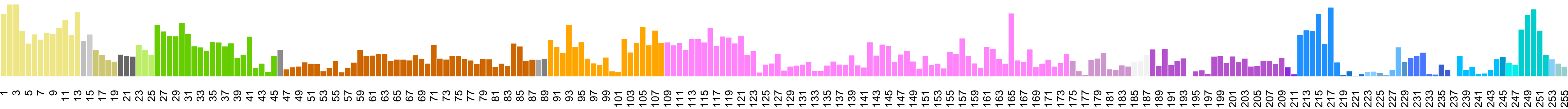
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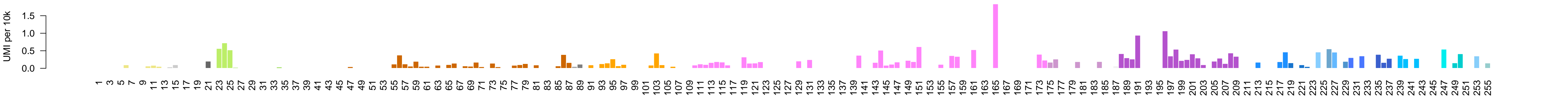
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UMI per 10k

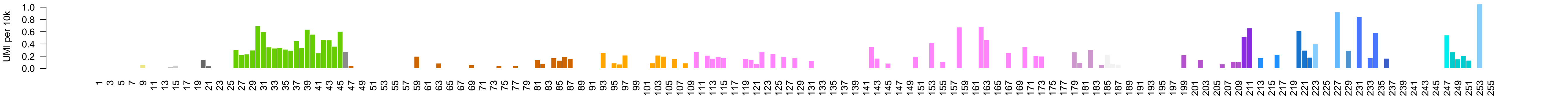
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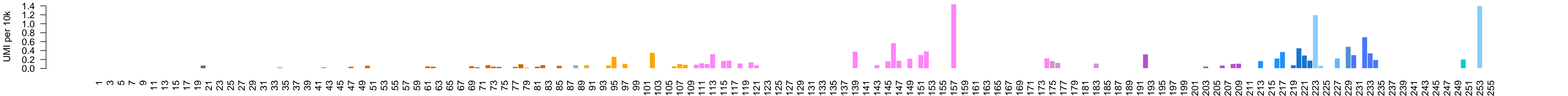
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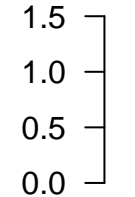


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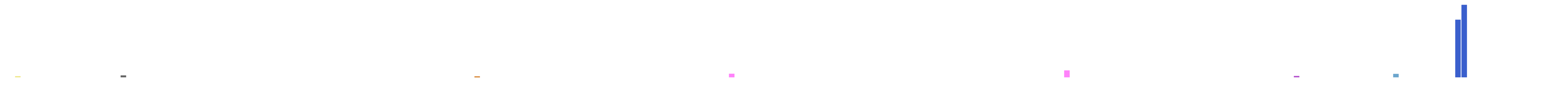


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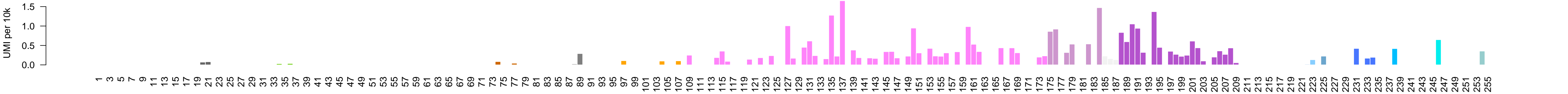
UMI per 10k



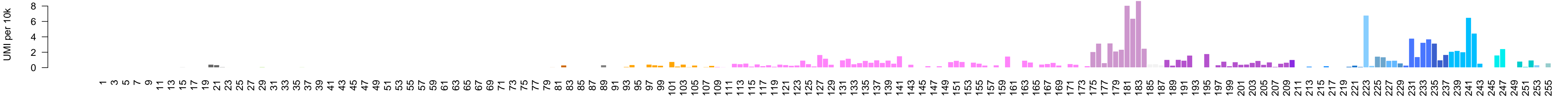
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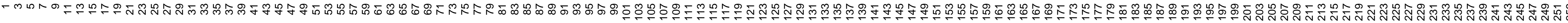
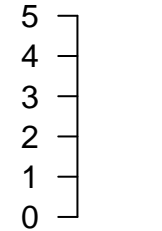
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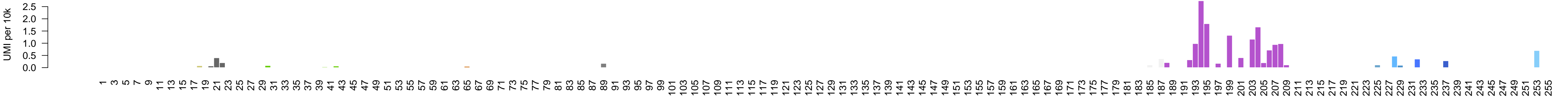
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UMI per 10k

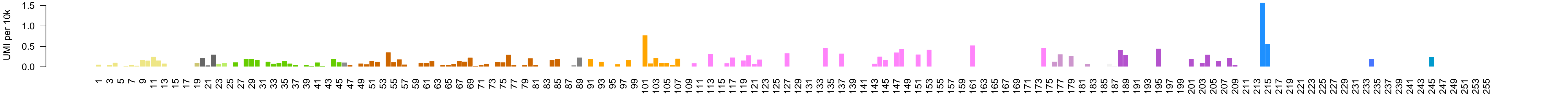


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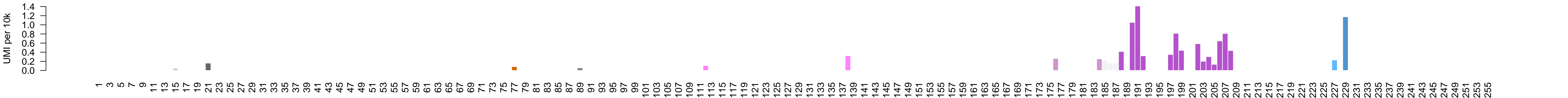


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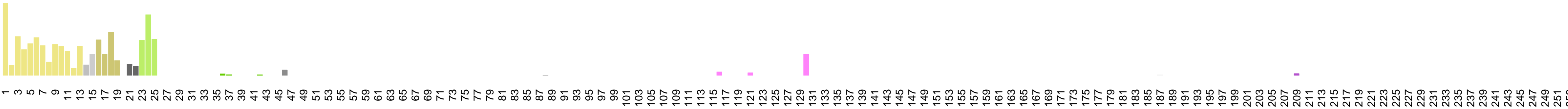
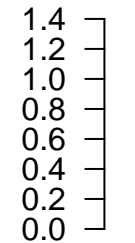
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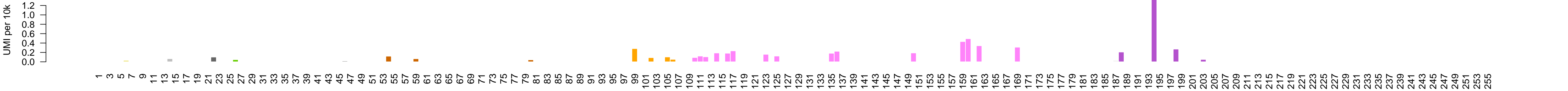
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UMI per 10k



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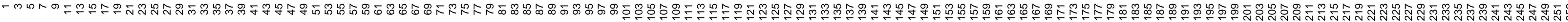
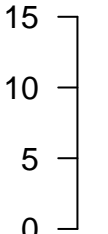
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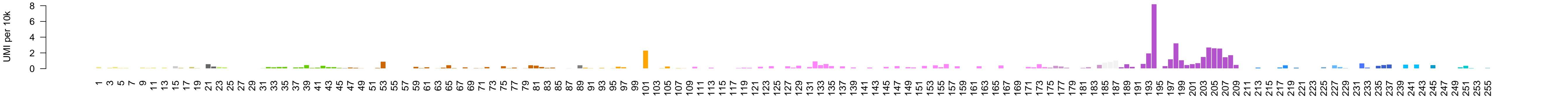
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UMI per 10k



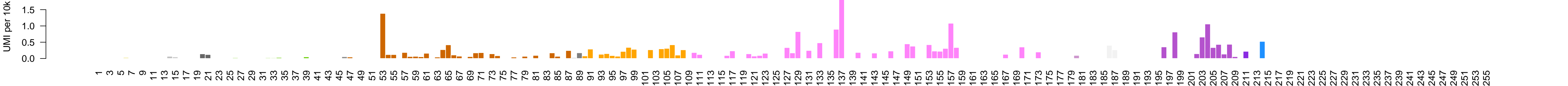
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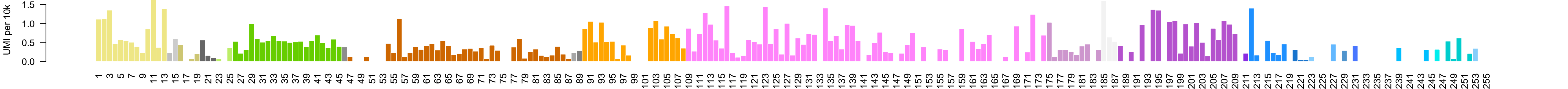
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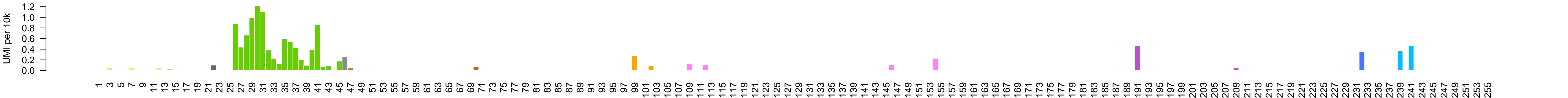
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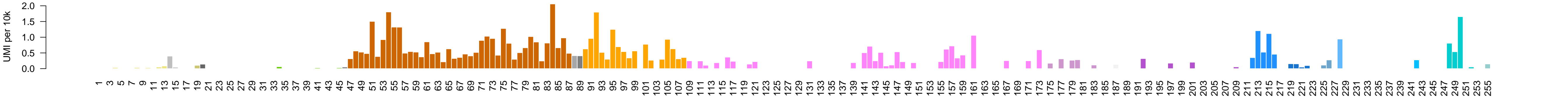
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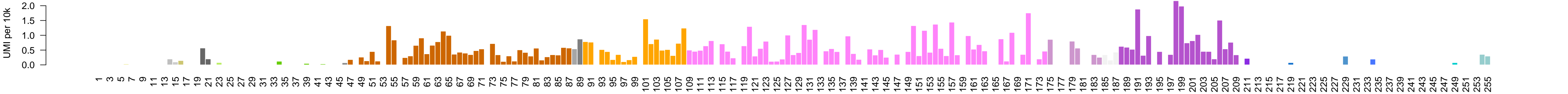
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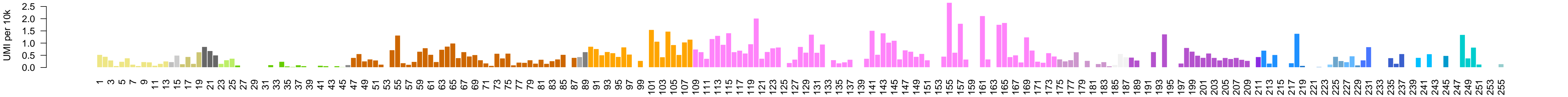
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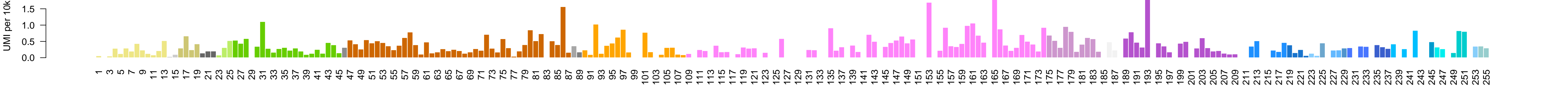
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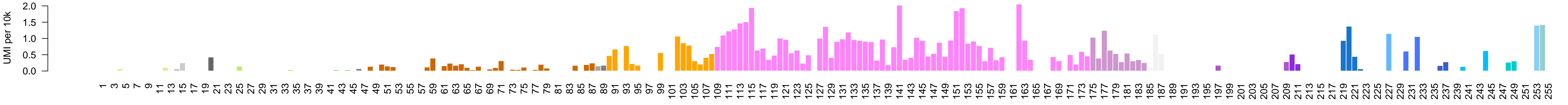


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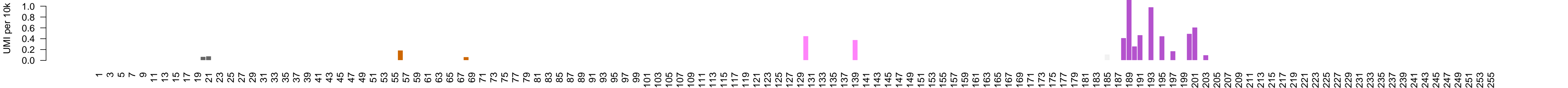
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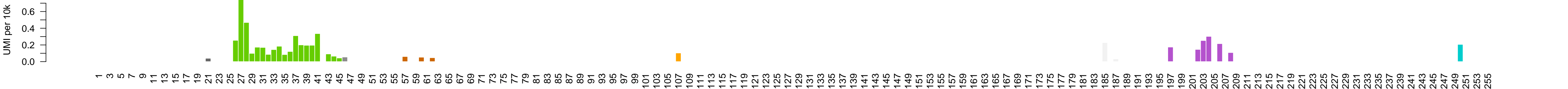
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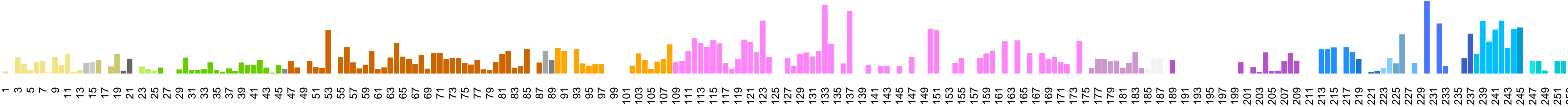
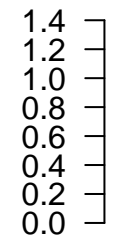


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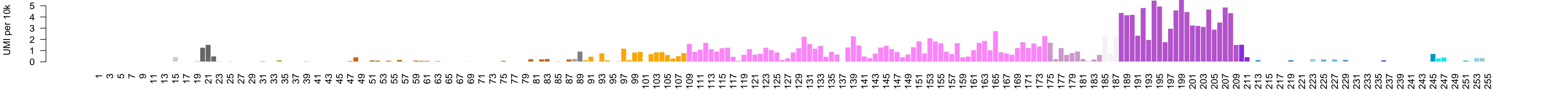


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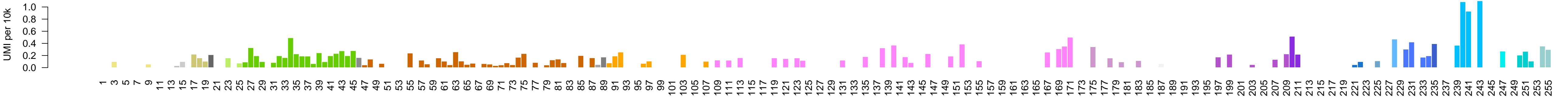
UMI per 10k



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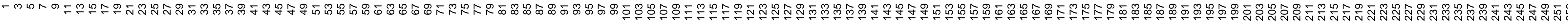
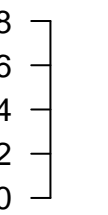


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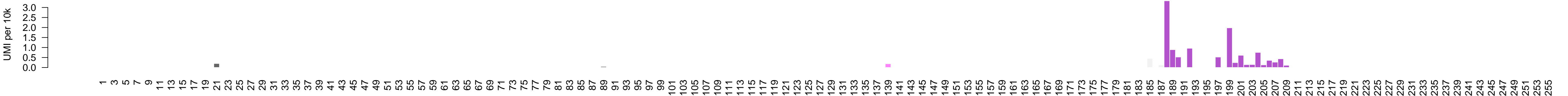


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UMI per 10k

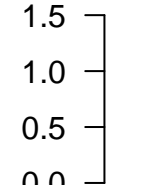


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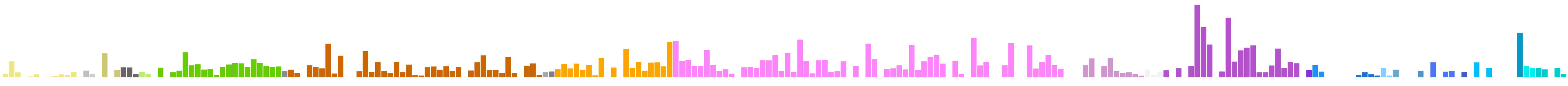


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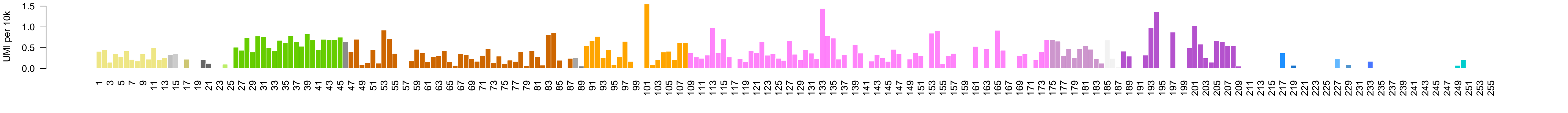
UMI per 10k



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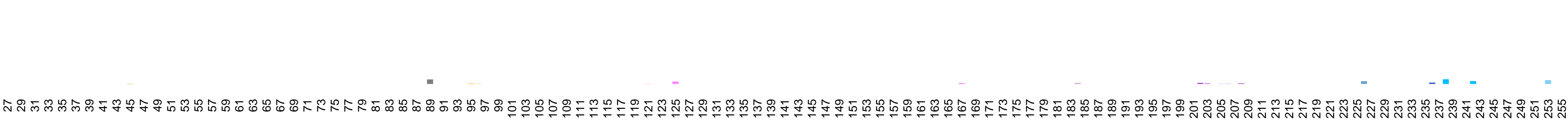


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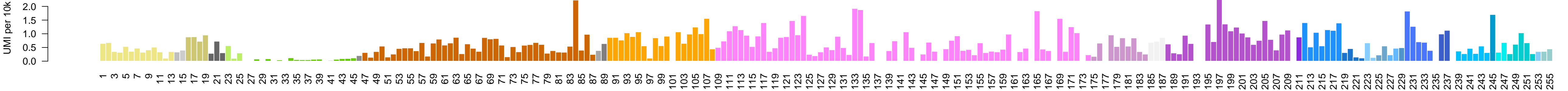
UMI per 10k

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1
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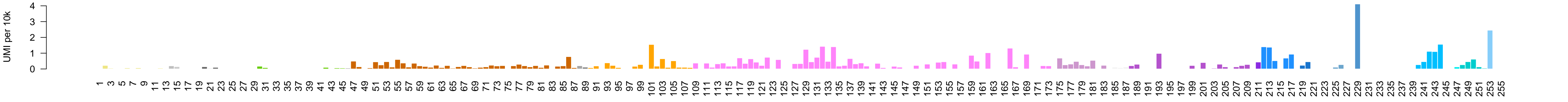
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PDZ.HG2.1:PARD6A/PARD6B/PARD6G
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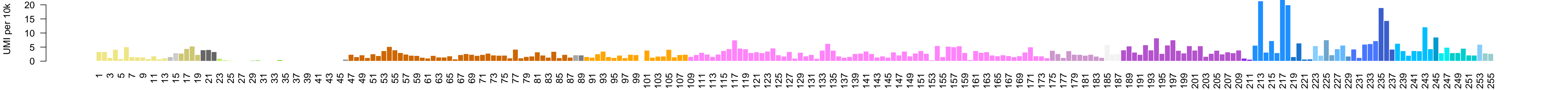


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PDZ.HG3.14:SHROOM2/SHROOM3/SHROOM4/SYNPO2/SYNPO2L

Hhon_g03284.t1



UMI per 10k

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0.0

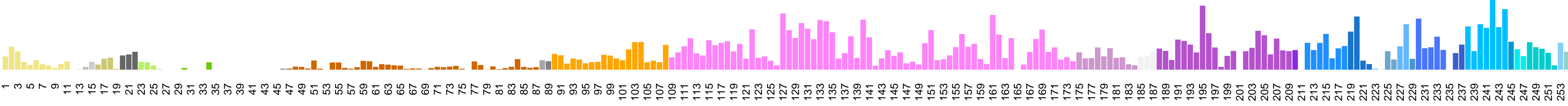
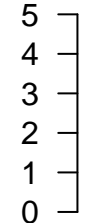
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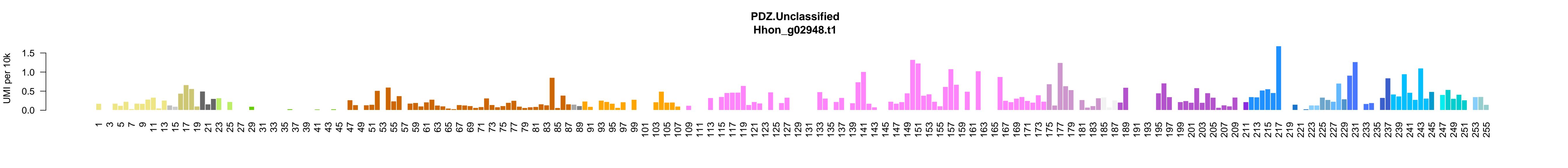
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PDZ.Unclassified
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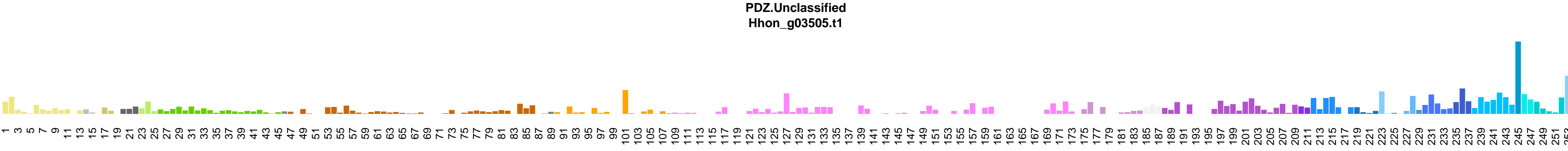
UMI per 10k





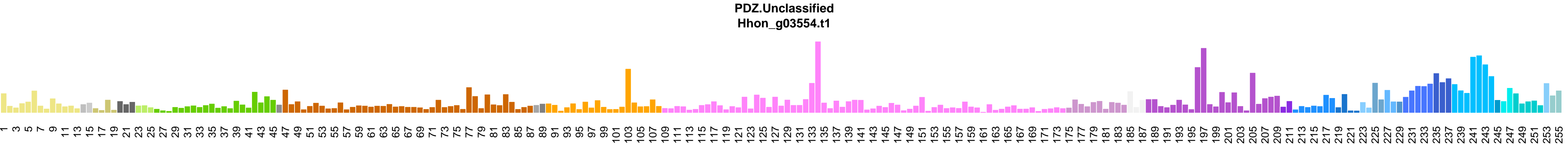
UMI per 10k

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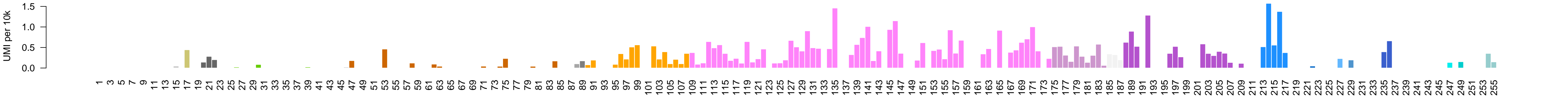


UMI per 10k

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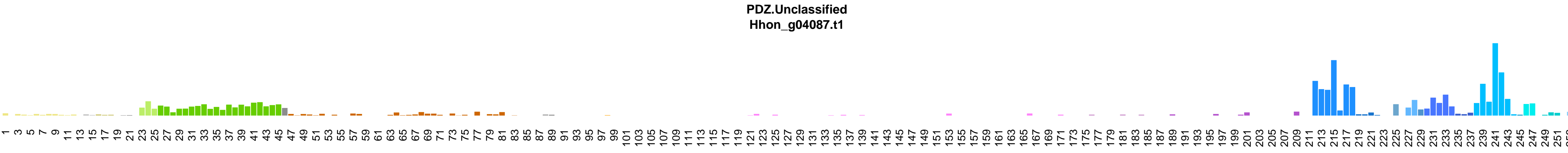


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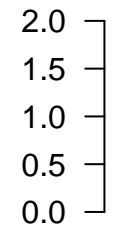


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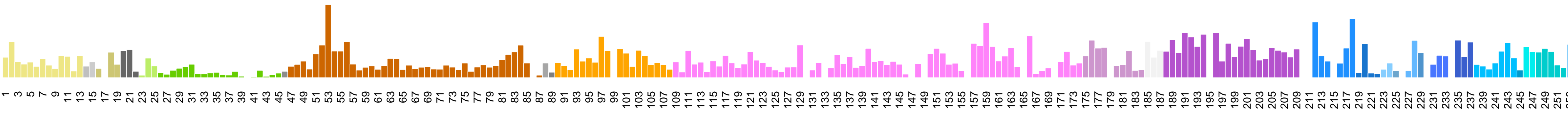
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UMI per 10k

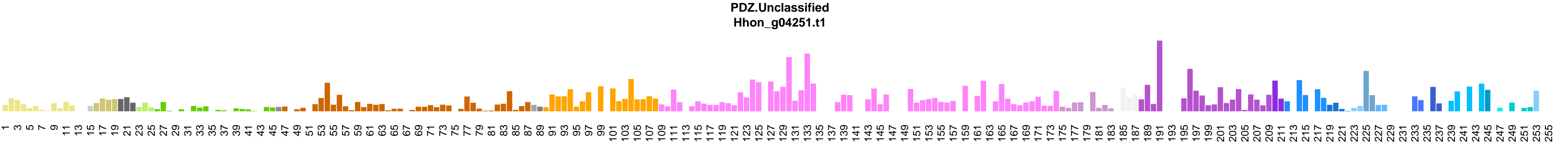


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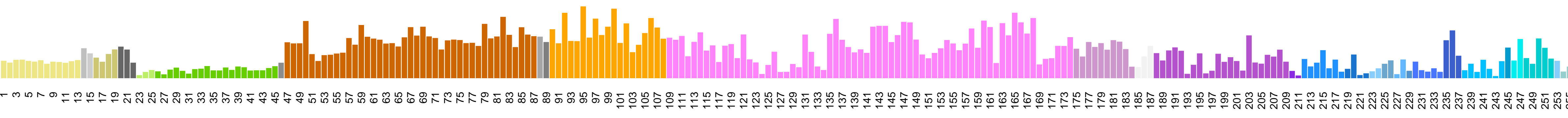
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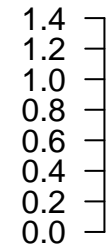


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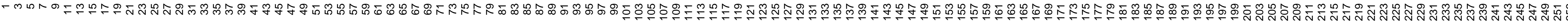
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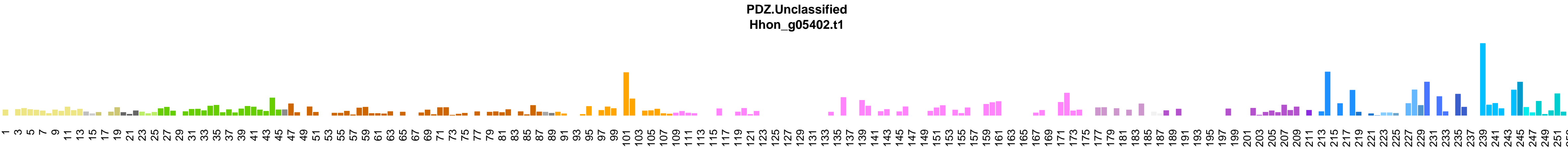
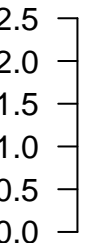
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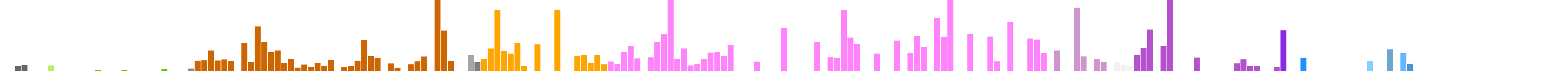


UMI per 10k

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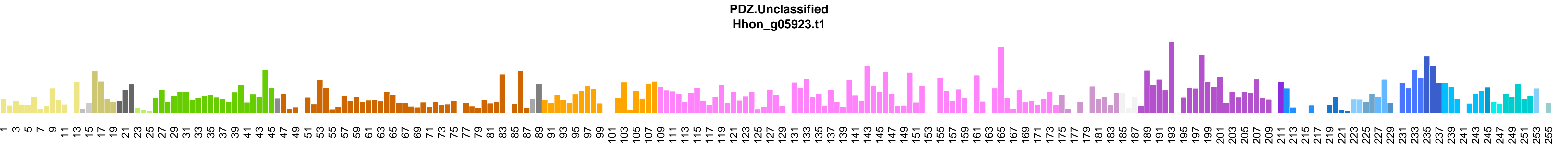
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UMI per 10k

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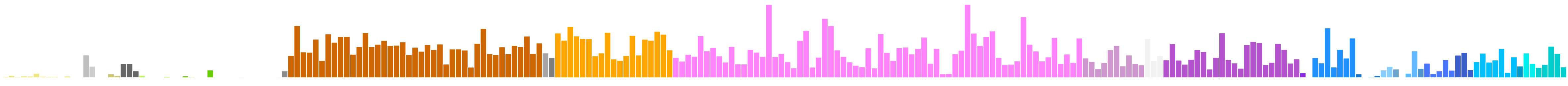


UMI per 10k

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PDZ.Unclassified
Hhon_g06231.t1

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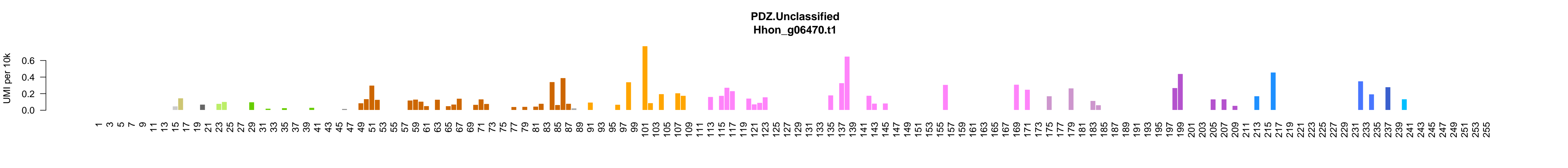


UMI per 10k

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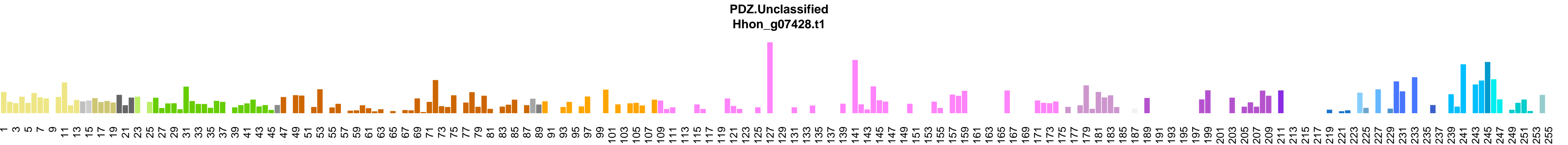
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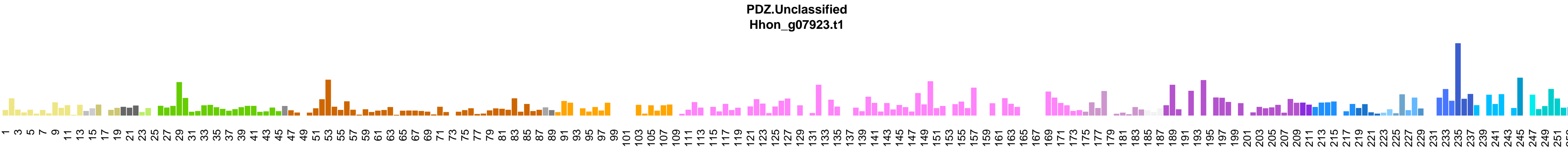
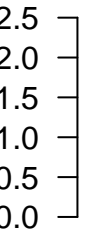


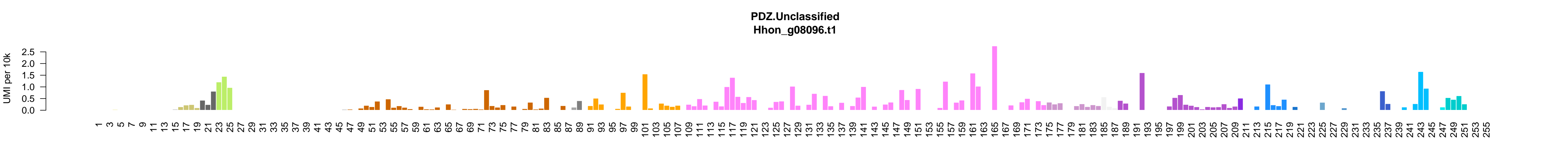
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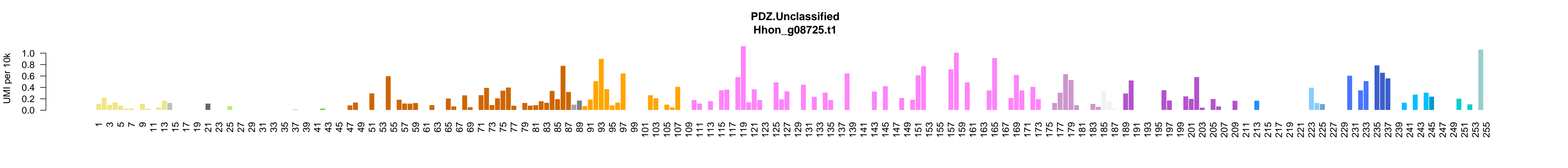
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0.2
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UMI per 10k





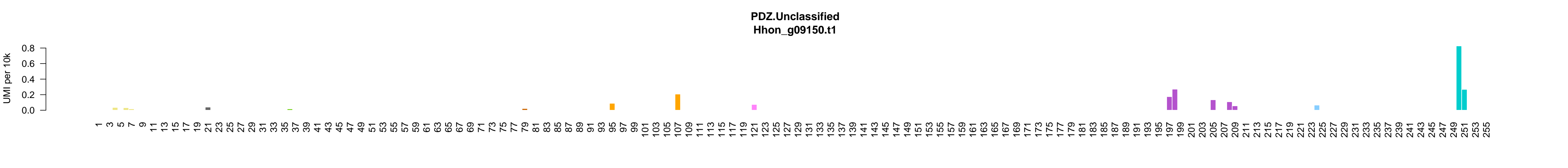


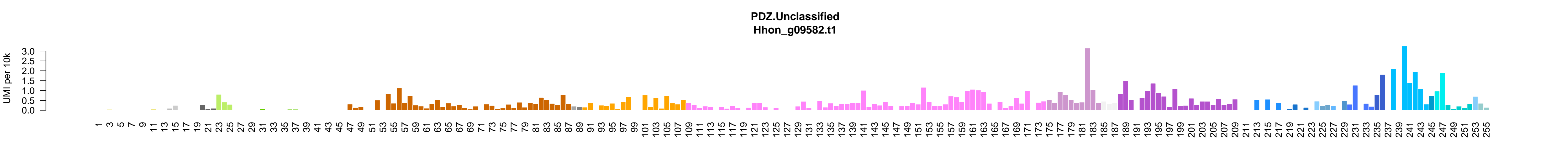
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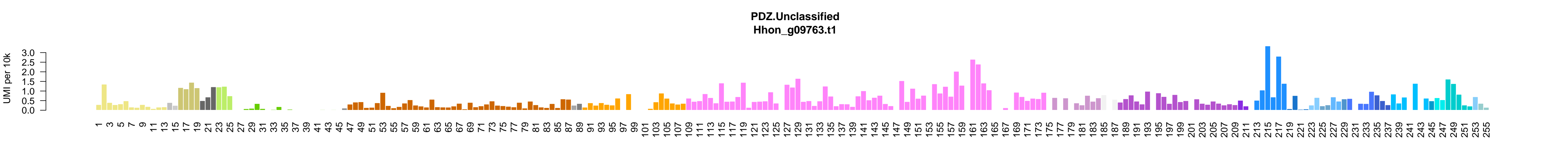
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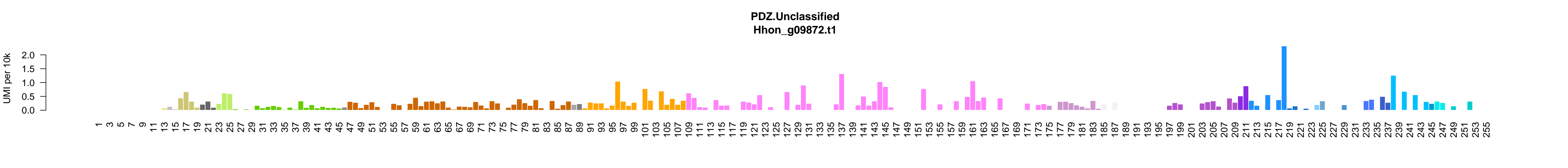
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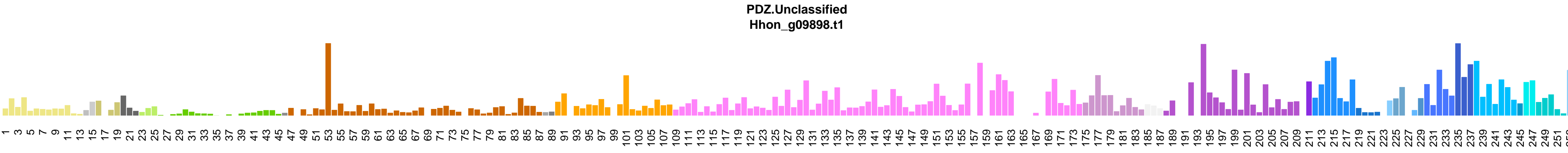
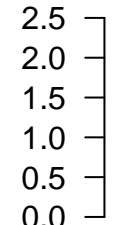


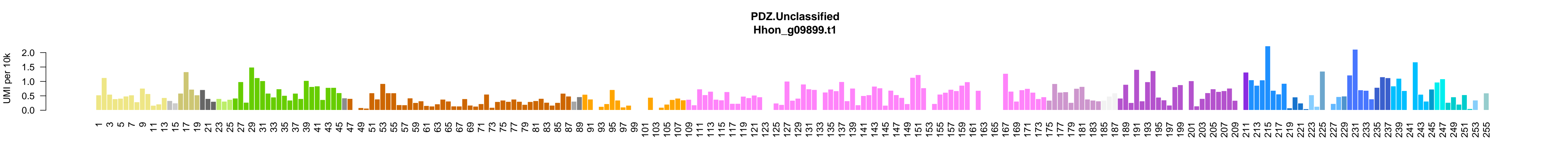






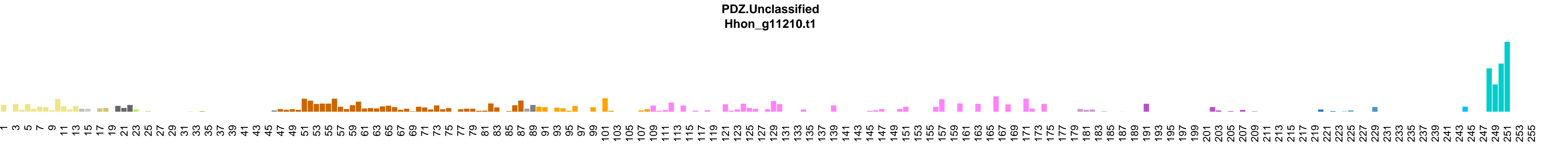
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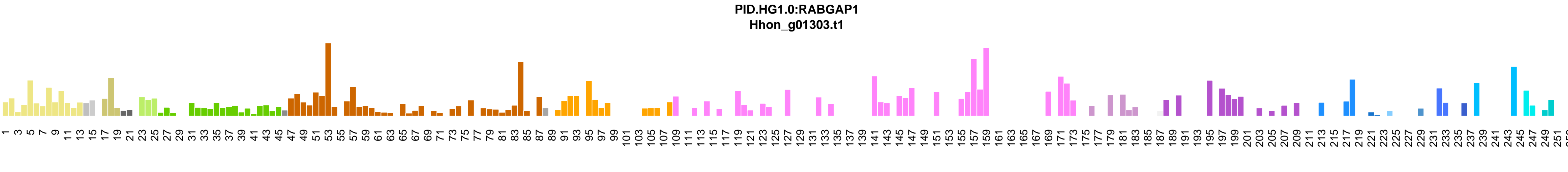


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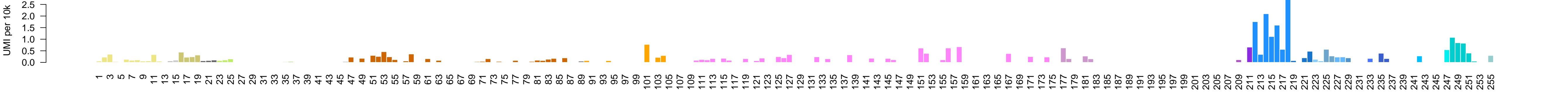
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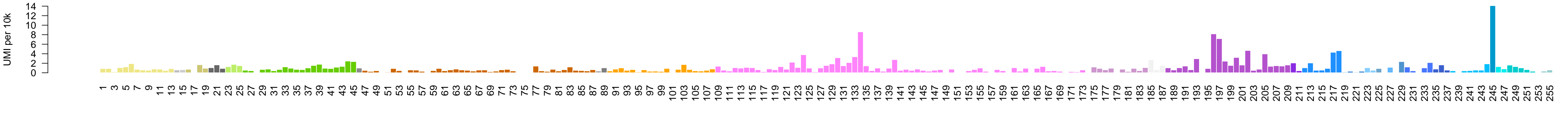
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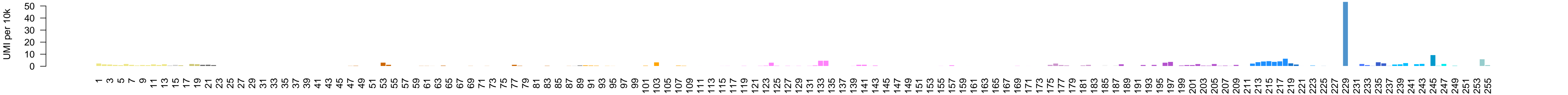
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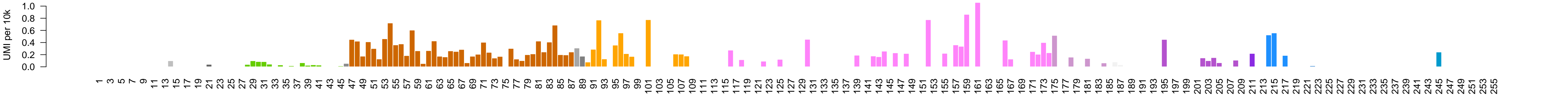
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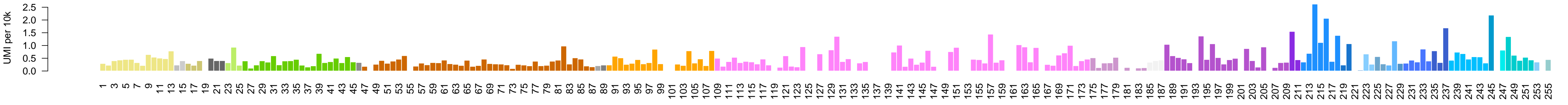
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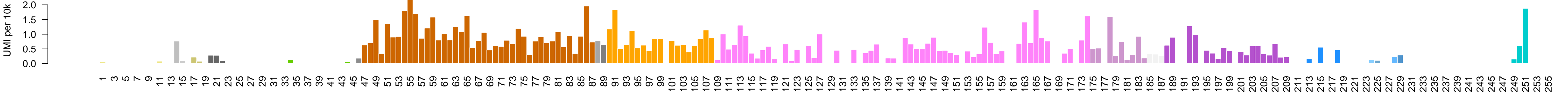
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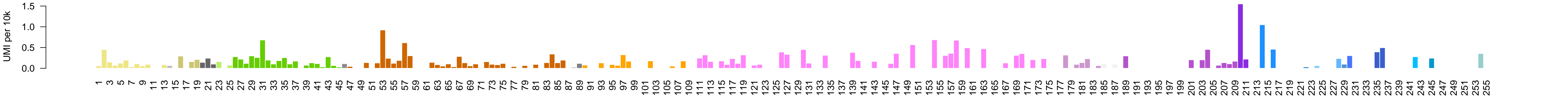


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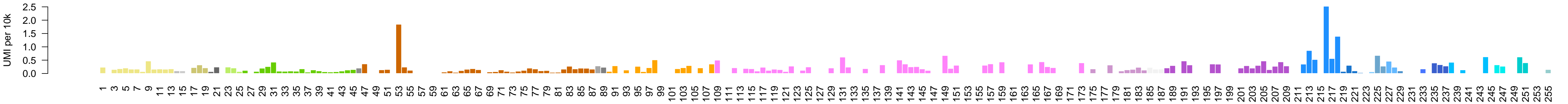


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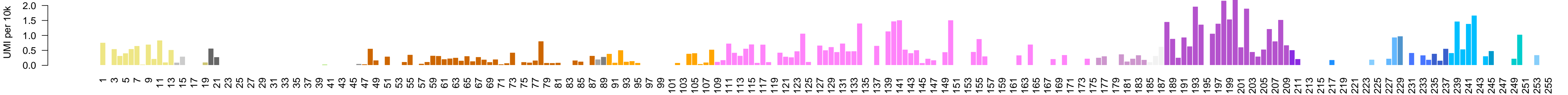
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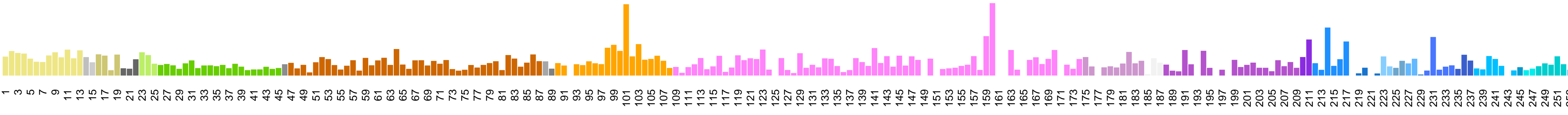
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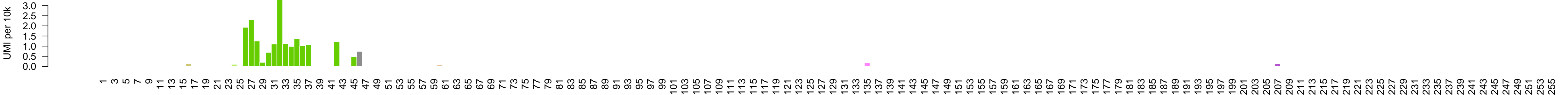
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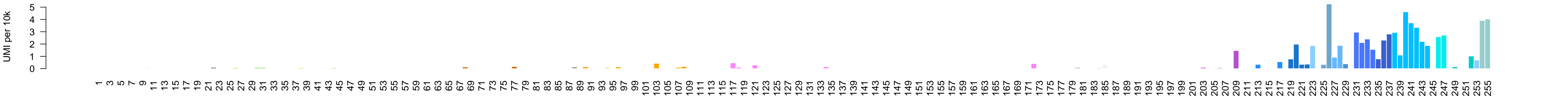
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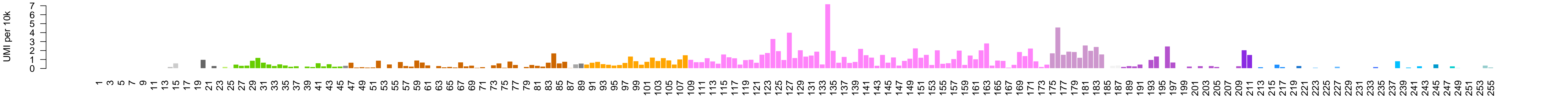
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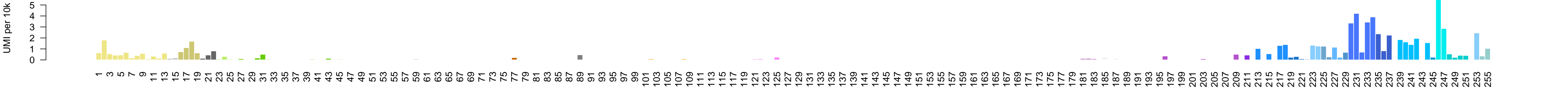
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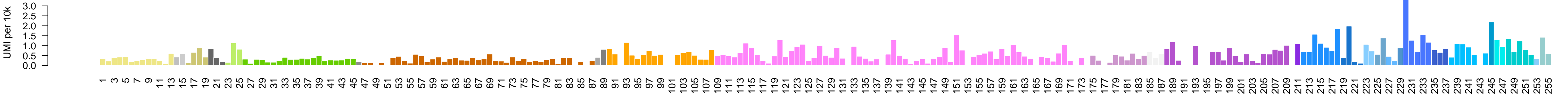
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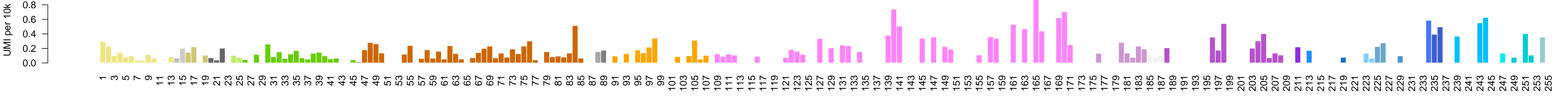


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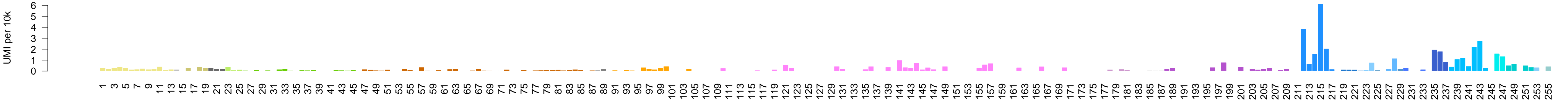


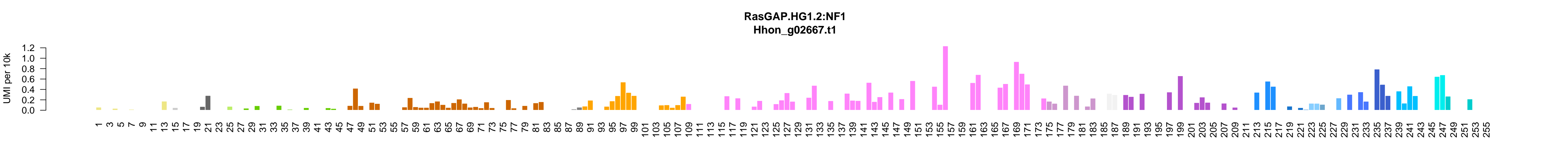
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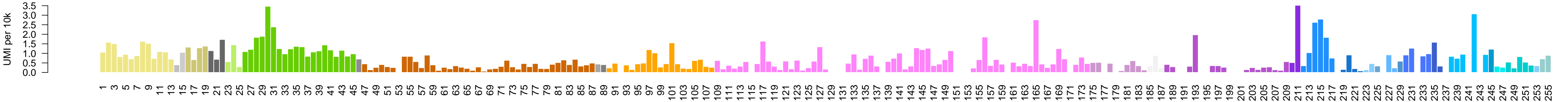


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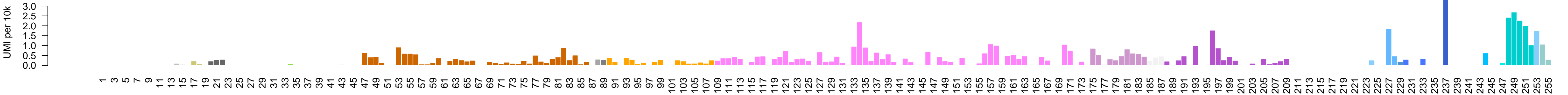




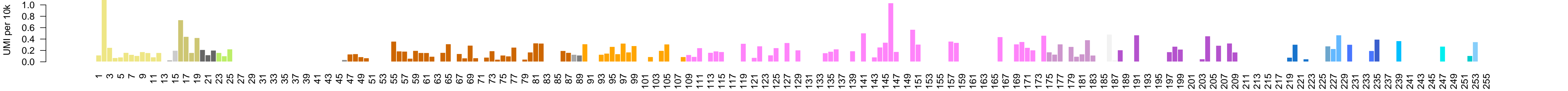
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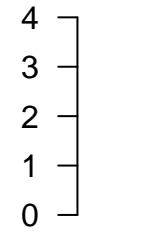


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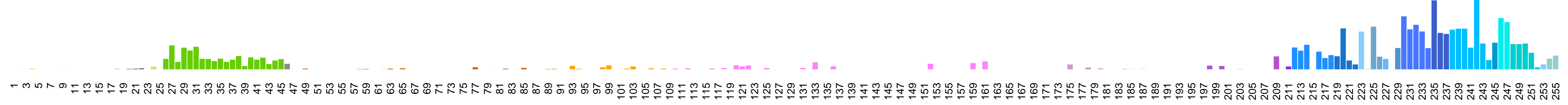


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UMI per 10k

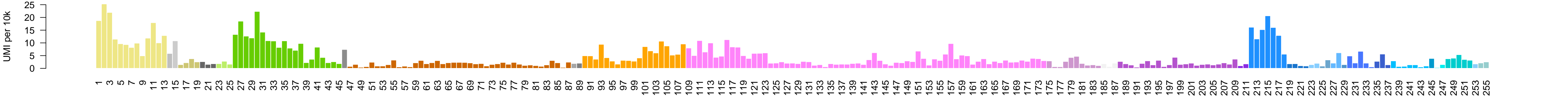


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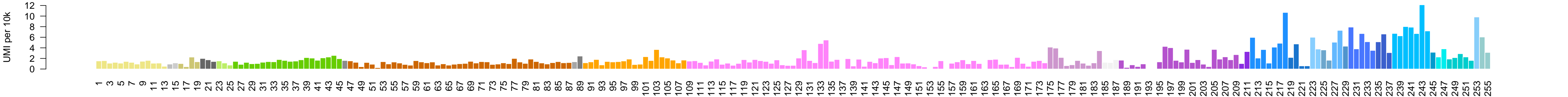


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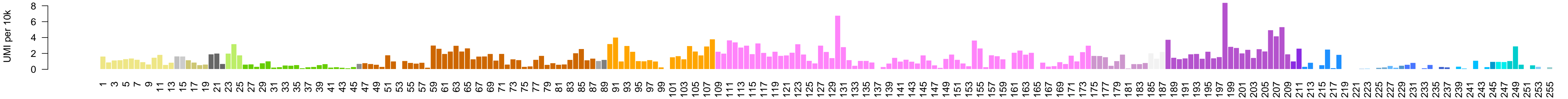


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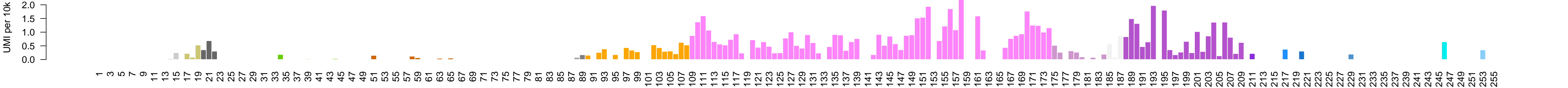
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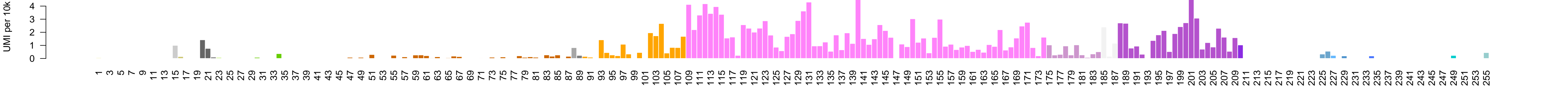
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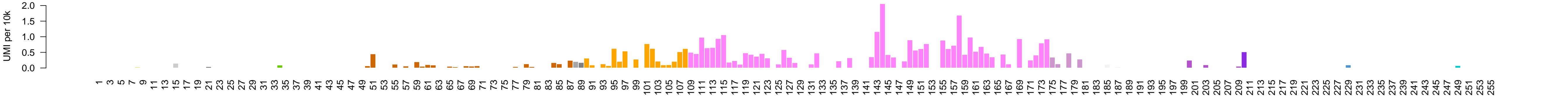
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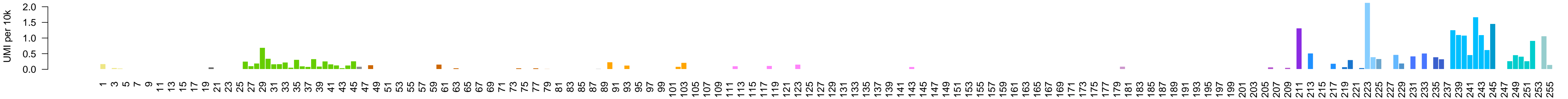
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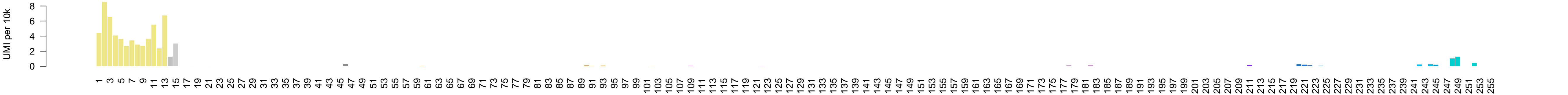
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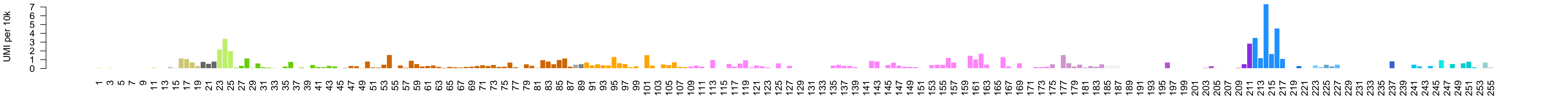
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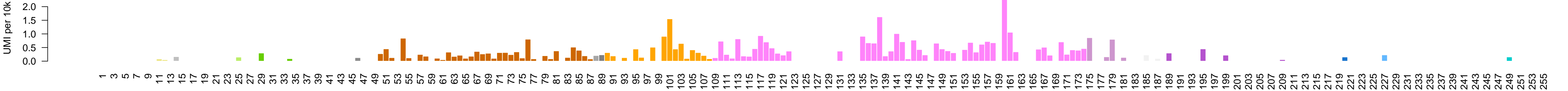
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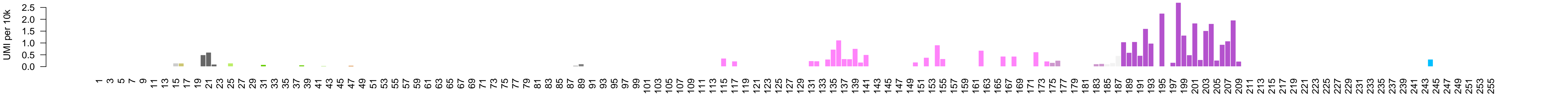
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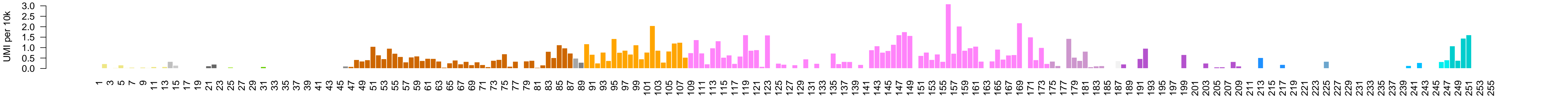
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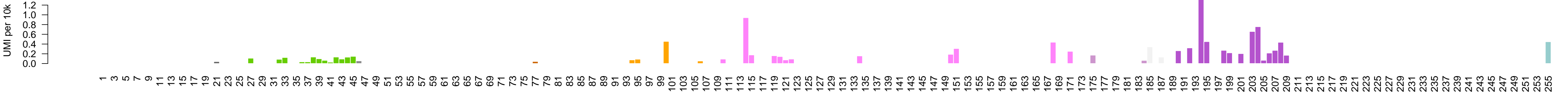
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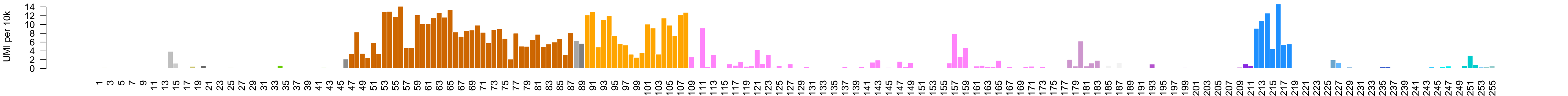


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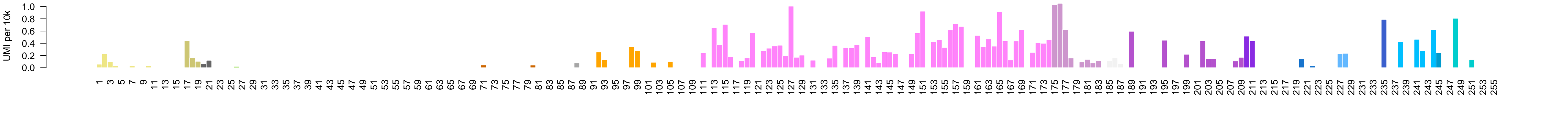
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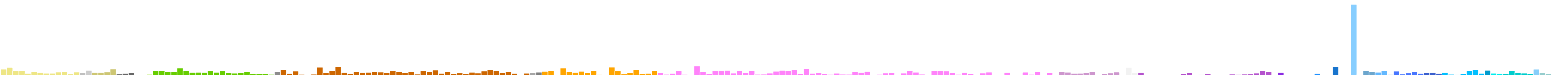
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UMI per 10k

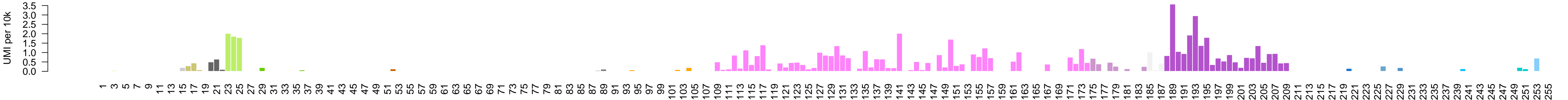
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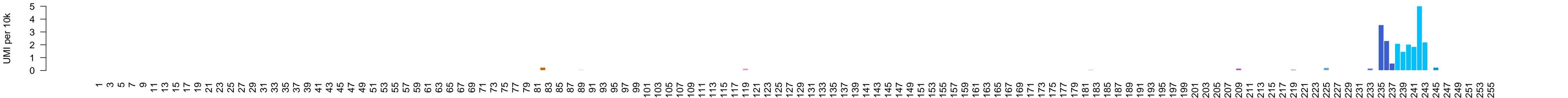
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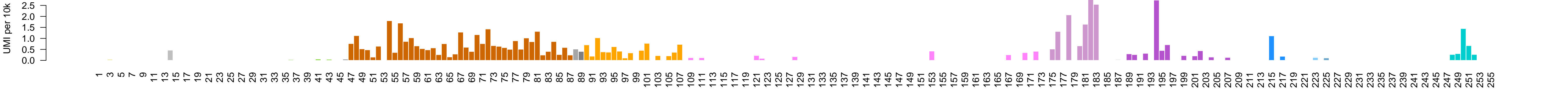
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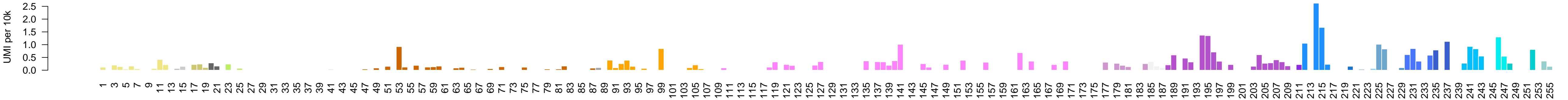
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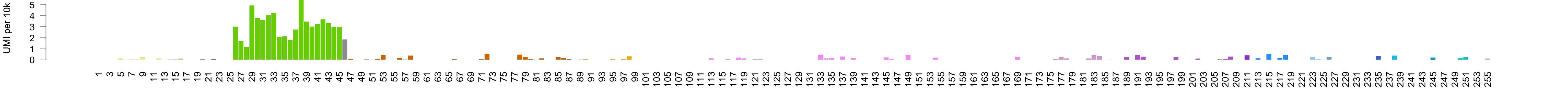
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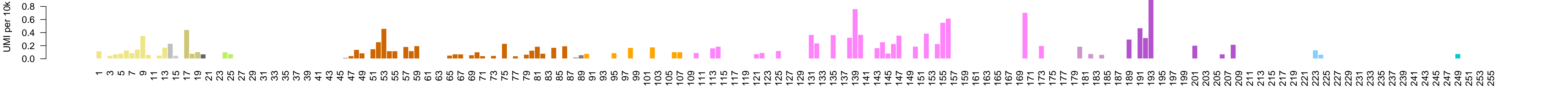


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Hhon_g01176.t1



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Hhon_g04889.t1



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

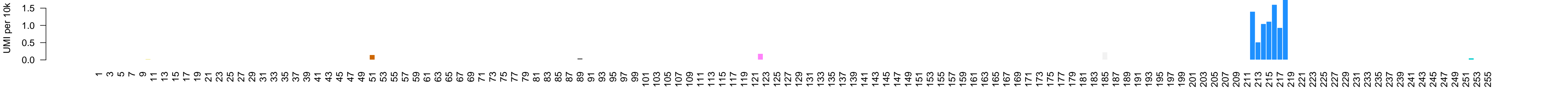
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UMI per 10k

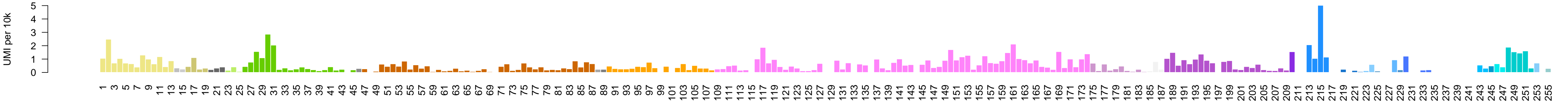
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1 3 5 7 9 11 13 15 17 19 21 23 25 27 29 31 33 35 37 39 41 43 45 47 49 51 53 55 57 59 61 63 65 67 69 71 73 75 77 79 81 83 85 87 89 91 93 95 97 99 101 103 105 107 109 111 113 115 117 119 121 123 125 127 129 131 133 135 137 139 141 143 145 147 149 151 153 155 157 159 161 163 165 167 169 171 173 175 177 179 181 183 185 187 189 191 193 195 197 199 201 203 205 207 209 211 213 215 217 219 221 223 225 227 229 231 233 235 237 239 241 243 245 247 249 251 253 255

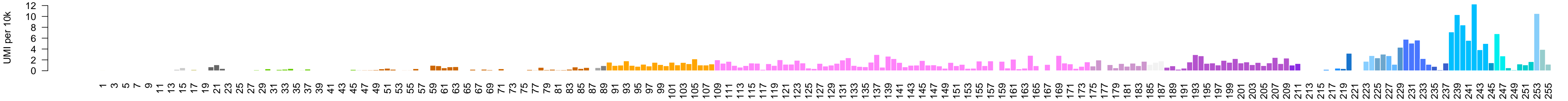
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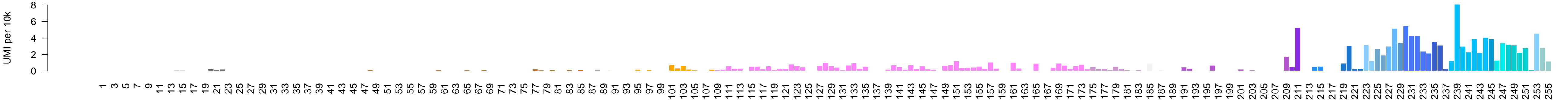
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Hhon_g08441.t1



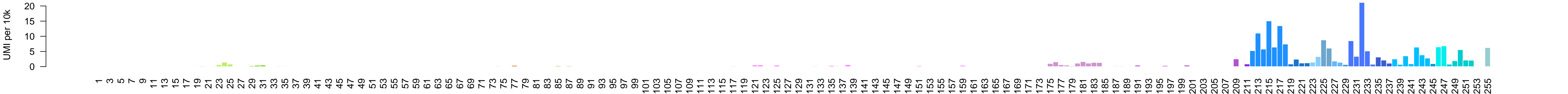
Synaphin.HG2.0:NA
Hhon_g07161.t1



Synaptotagmin.HG1.0:SYT7
Hhon_g03364.t1

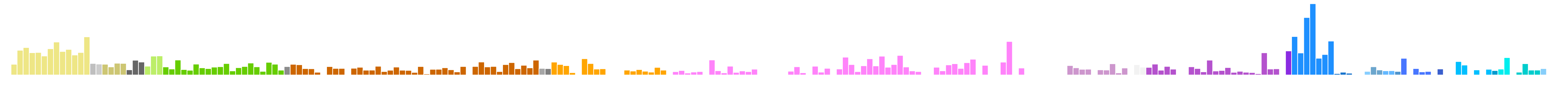
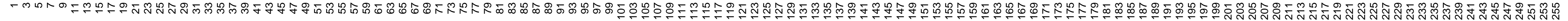
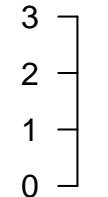


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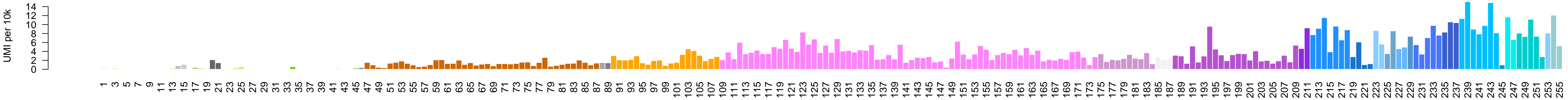


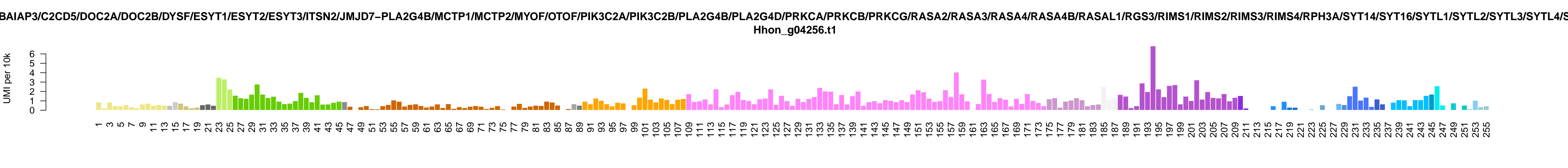
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Hhon_g05915.t1

UMI per 10k



Synaptotagmin.HG1.2:SYT1/SYT2/SYT5/SYT8
Hhon_g03362.t1

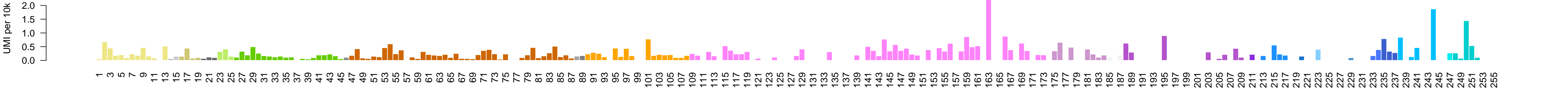




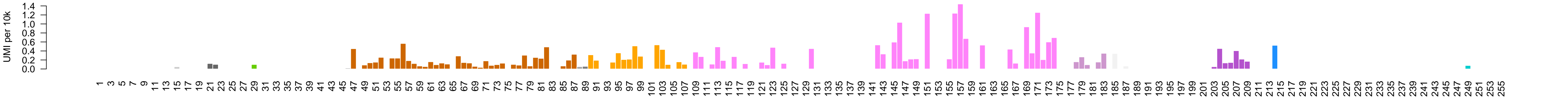
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Hhon_g07466.t1



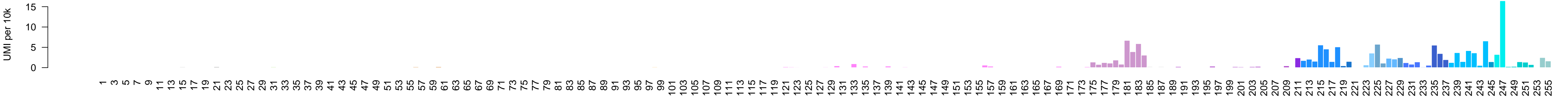
Synaptotagmin.HG1.30:like:BAIAP3/UNC13D
Hhon_g08515.t1



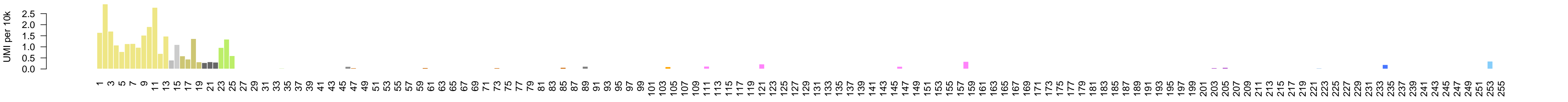
Synaptotagmin.HG1.32:BAIAP3/UNC13D
Hhon_g03866.t1



Synaptotagmin.HG1.32:BAIAP3/UNC13D
Hhon_g03867.t1



Synaptotagmin.HG1.34:like:DYSF/MYOF
Hhon_g02267.t1

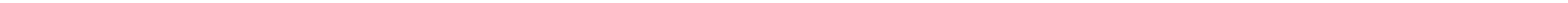


Synaptotagmin.HG1.35:DYSF/MYOF
Hhon_g02268.t1

UMI per 10k

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3
2
1
0

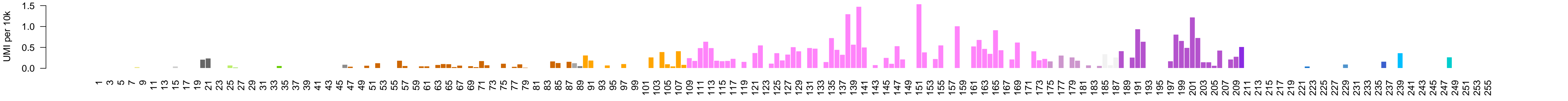
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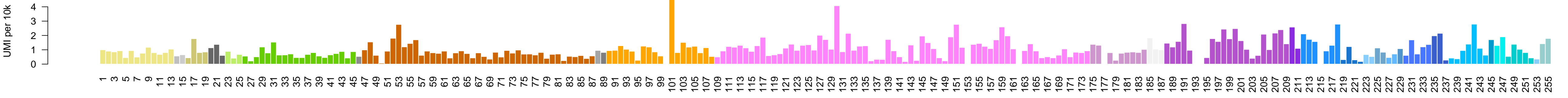
Synaptotagmin.HG1.49:ESYT1/ESYT2/ESYT3
Hhon_g05559.t1



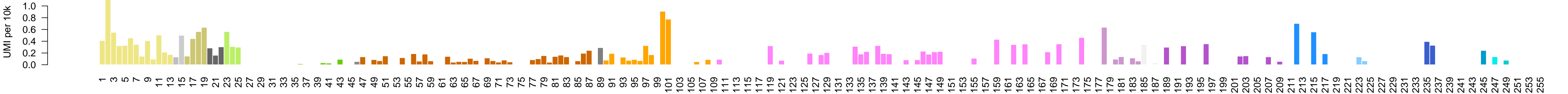
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Hhon_g01247.t1



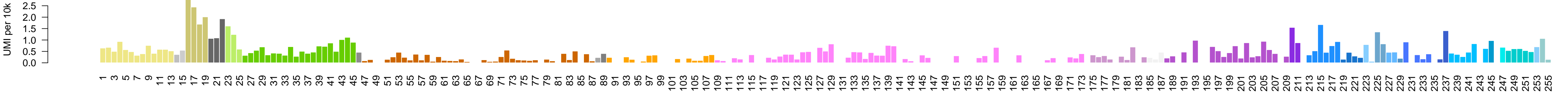
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Hhon_g02376.t1



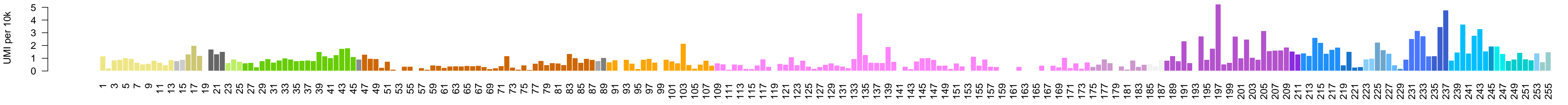
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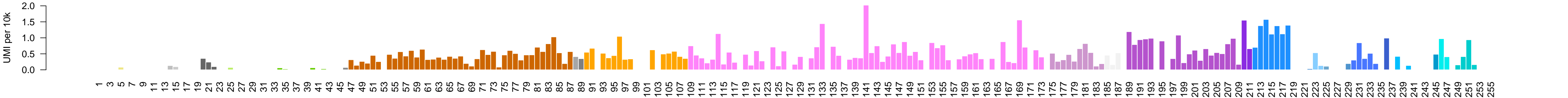
Synaptotagmin.HG1.62:PRKCA/PRKCB/PRKCG
Hhon_g04801.t1



Syntaxin.HG1.2:STX7/STX12
Hhon_g00160.t1

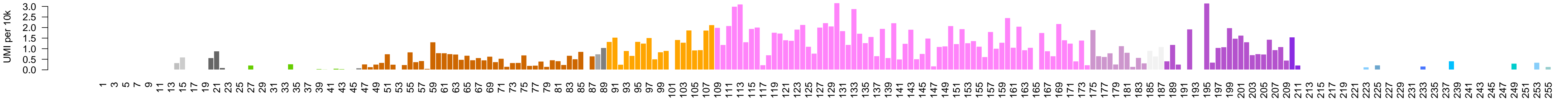


Syntaxin.HG1.4:STX1A/STX1B/STX2/STX3/STX4
Hhon_g06927.t1



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

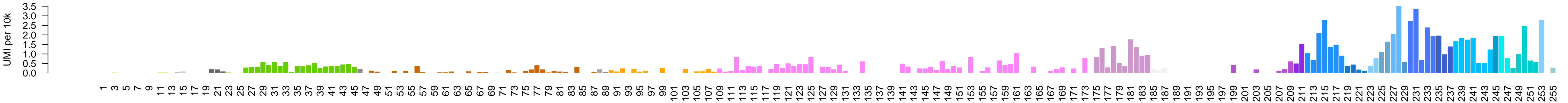
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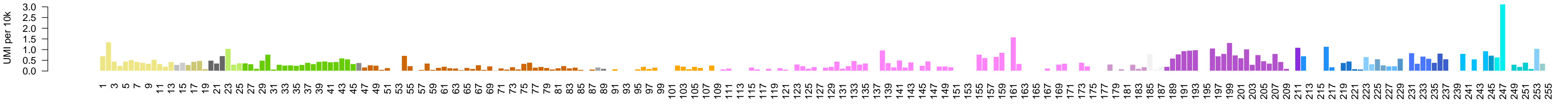
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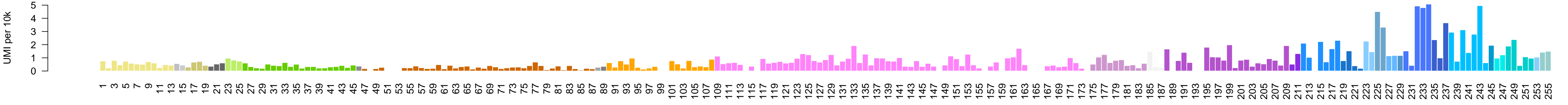
Syntaxin.HG1.4:STX1A/STX1B/STX2/STX3/STX4
Hhon_g07820.t1



Y_phosphatase.HG1.13:PTPN6/PTPN11
Hhon_g10340.t1



Y_phosphatase.HG1.14:PTPN5/PTPN7/PTPRR
Hhon_g08135.t1



Y_phosphatase.HG1.16:like:PTPRK/PTPRM/PTPRT/PTPRU
Hhon_g02885.t1

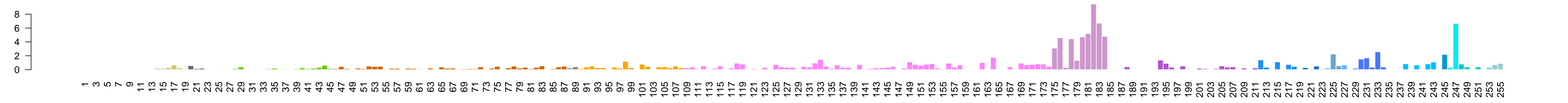


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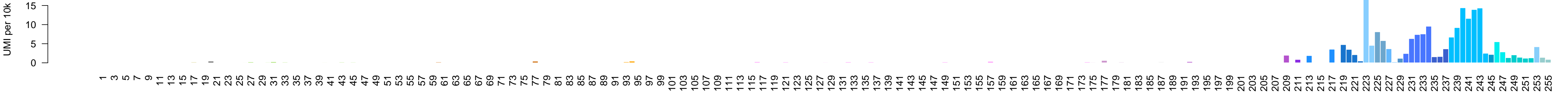
UMI per 10k

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6
4
2
0

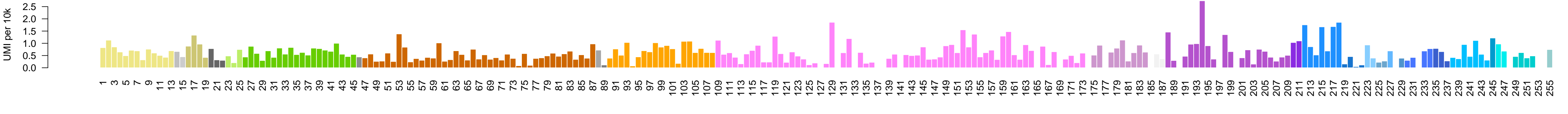
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Y_phosphatase.HG1.29:PTPRN/PTPRN2
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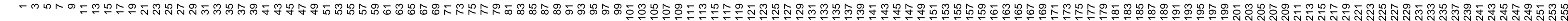
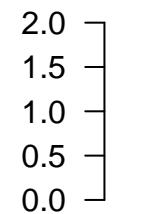


Y_phosphatase.HG1.30:PTPN1/PTPN2
Hhon_g06178.t1

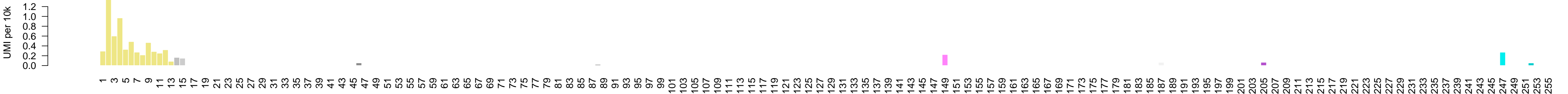


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Hhon_g06297.t1

UMI per 10k

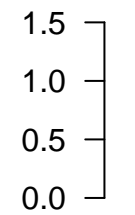


Y_phosphatase.HG1.41:like:PTPN5/PTPN7/PTPRR
Hhon_g03898.t1

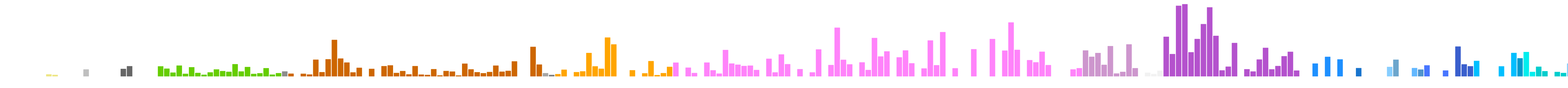


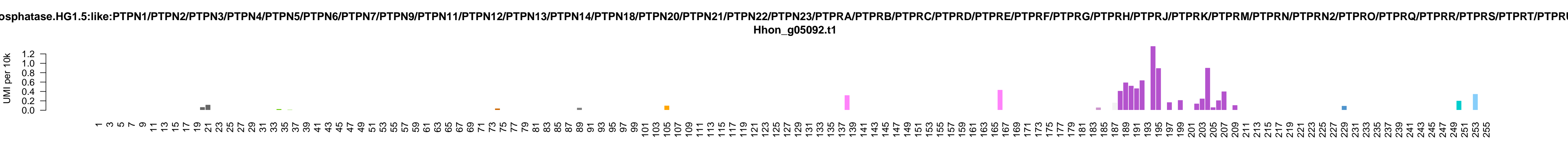
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UMI per 10k

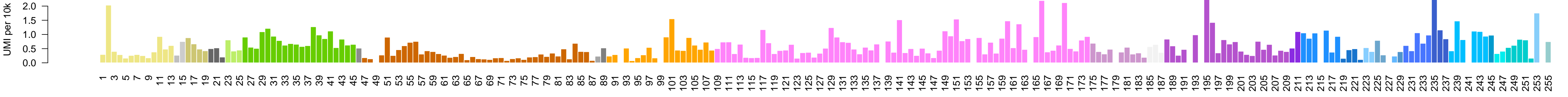


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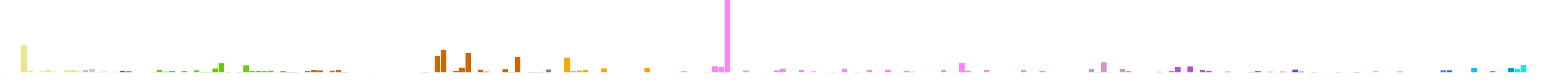
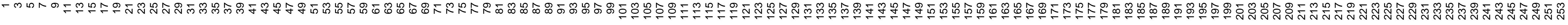
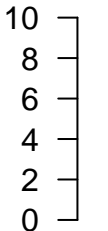


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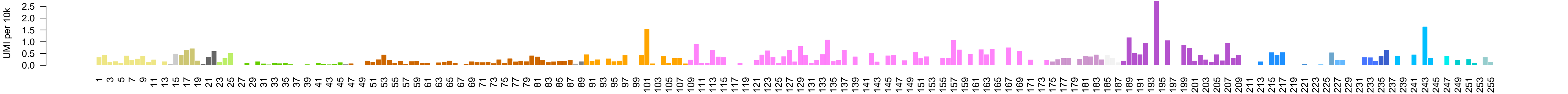
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UMI per 10k



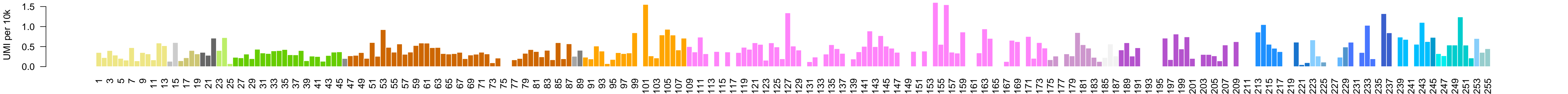
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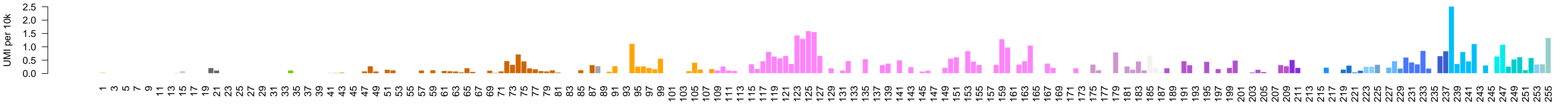


Y_phosphatase.HG2.5:like: CDC14A/CDC14B/CDC14C/DUSP23/PTP4A1/PTP4A2/PTP4A3/PTPDC1

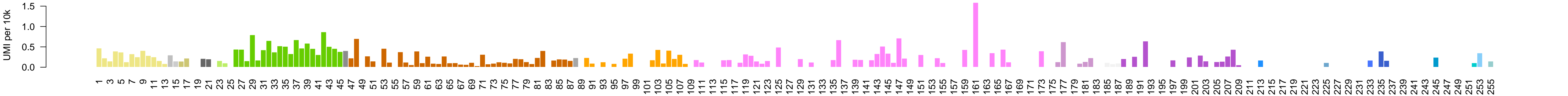
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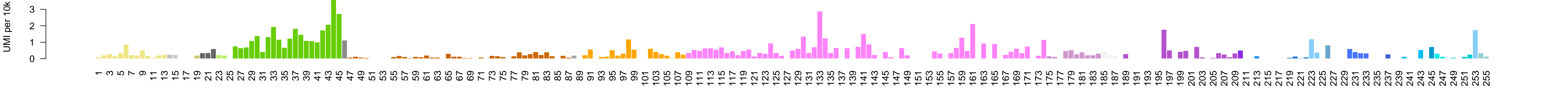
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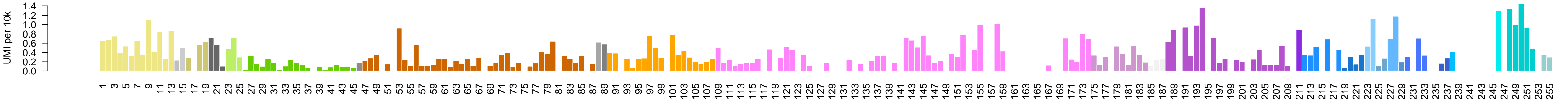
Y_phosphatase.HG3.1:like:DUSP1/DUSP4/DUSP5
Hhon_g07988.t1



Y_phosphatase.HG3.4:like:DUSP1/DUSP4/DUSP5
Hhon_g05020.t1



Y_phosphatase.HG3.4:like:DUSP1/DUSP4/DUSP5
Hhon_g07846.t1



Y_phosphatase.HG6.0:NA
Hhon_g06487.t1

