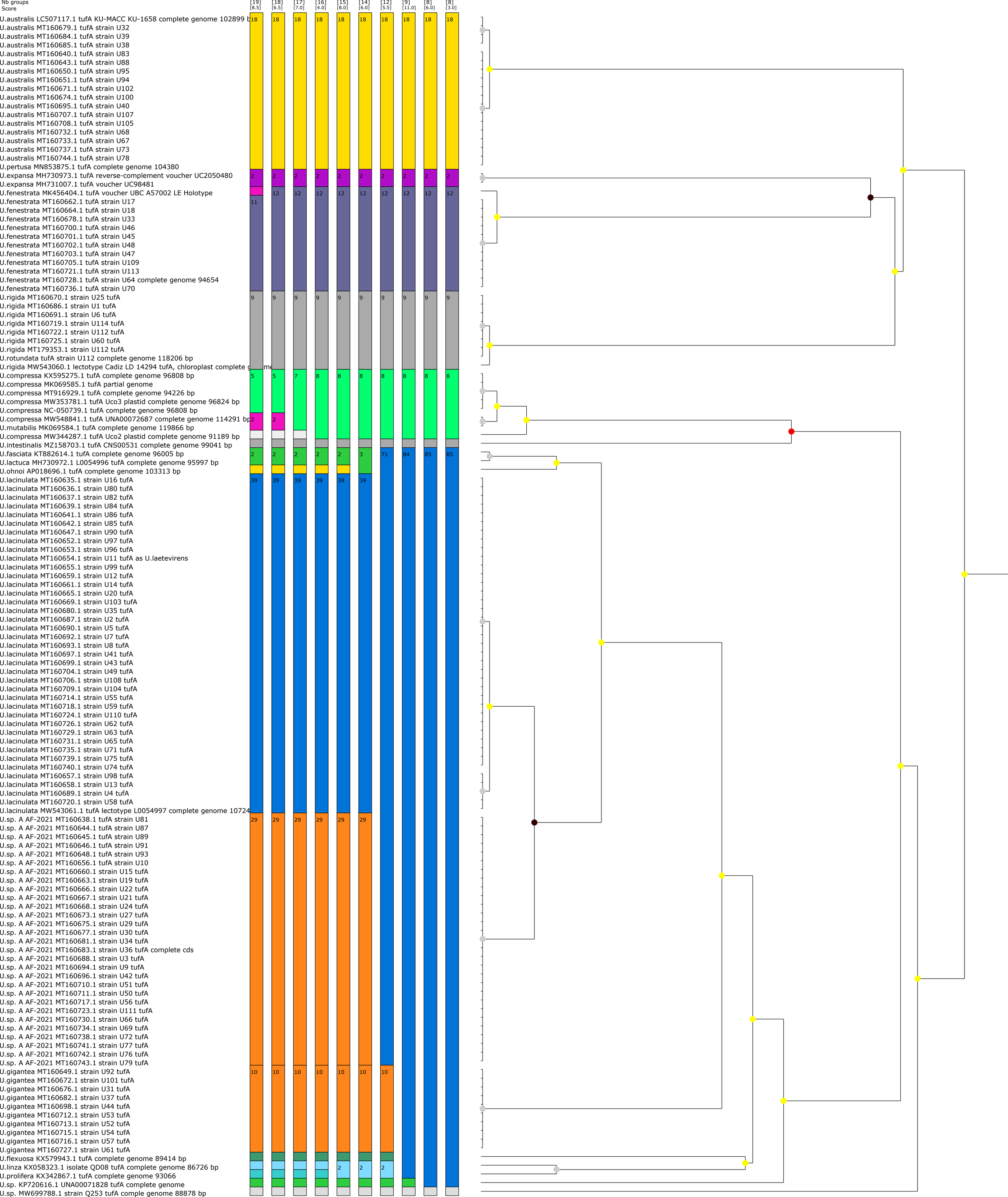
Nb groups
Score
[19] [8.5]
[18] [6.5]
[17] [7.0]
[16] [4.0]
[15] [8.0]
[14] [6.0]
[12] [5.5]
[9] [11.0]
[8] [6.0]
[8] [3.0]
U.australis LC507117.1 tufA KU-MACC KU-1658 complete genome 102899 b
U.australis MT160679.1 tufA strain U32
U.australis MT160684.1 tufA strain U39
U.australis MT160685.1 tufA strain U38
U.australis MT160640.1 tufA strain U83
U.australis MT160643.1 tufA strain U88
U.australis MT160650.1 tufA strain U95
U.australis MT160651.1 tufA strain U94
U.australis MT160671.1 tufA strain U102
U.australis MT160674.1 tufA strain U100
U.australis MT160695.1 tufA strain U40
U.australis MT160707.1 tufA strain U107
U.australis MT160708.1 tufA strain U105
U.australis MT160732.1 tufA strain U68
U.australis MT160733.1 tufA strain U67
U.australis MT160737.1 tufA strain U73
U.australis MT160744.1 tufA strain U78
U.pertusa MN853875.1 tufA complete genome 104380
U.expansa MH730973.1 tufA reverse-complement voucher UC2050480
U.expansa MH731007.1 tufA voucher UC98481
U.fenestrata MK456404.1 tufA voucher UBC A57002 LE Holotype
U.fenestrata MT160662.1 tufA strain U17
U.fenestrata MT160664.1 tufA strain U18
U.fenestrata MT160678.1 tufA strain U33
U.fenestrata MT160700.1 tufA strain U46
U.fenestrata MT160701.1 tufA strain U45
U.fenestrata MT160702.1 tufA strain U48
U.fenestrata MT160703.1 tufA strain U47
U.fenestrata MT160705.1 tufA strain U109
U.fenestrata MT160721.1 tufA strain U113
U.fenestrata MT160728.1 tufA strain U64 complete genome 94654
U.fenestrata MT160736.1 tufA strain U70
U.rigida MT160670.1 strain U25 tufA
U.rigida MT160686.1 strain U1 tufA
U.rigida MT160691.1 strain U6 tufA
U.rigida MT160719.1 strain U114 tufA
U.rigida MT160722.1 strain U112 tufA
U.rigida MT160725.1 strain U60 tufA
U.rigida MT179353.1 strain U112 tufA
U.rotundata tufA strain U112 complete genome 118206 bp
U.rigida MW543060.1 lectotype Cadiz LD 14294 tufA, chloroplast complete g me
U.compressa KX595275.1 tufA complete genome 96808 bp
U.compressa MK069585.1 tufA partial genome
U.compressa MT916929.1 tufA complete genome 94226 bp
U.compressa MW353781.1 tufA Uco3 plastid complete genome 96824 bp
U.compressa NC-050739.1 tufA complete genome 96808 bp
U.compressa MW548841.1 tufA UNA00072687 complete genome 114291 bp
U.mutabilis MK069584.1 tufA complete genome 119866 bp
U.compressa MW344287.1 tufA Uco2 plastid complete genome 91189 bp
U.intestinalis MZ158703.1 tufA CNS00531 complete genome 99041 bp
U.fasciata KT882614.1 tufA complete genome 96005 bp
U.lactuca MH730972.1 L0054996 tufA complete genome 95997 bp
U.ohnoi AP018696.1 tufA complete genome 103313 bp
U.lacinulata MT160635.1 strain U16 tufA
U.lacinulata MT160636.1 strain U80 tufA
U.lacinulata MT160637.1 strain U82 tufA
U.lacinulata MT160639.1 strain U84 tufA
U.lacinulata MT160641.1 strain U86 tufA
U.lacinulata MT160642.1 strain U85 tufA
U.lacinulata MT160647.1 strain U90 tufA
U.lacinulata MT160652.1 strain U97 tufA
U.lacinulata MT160653.1 strain U96 tufA
U.lacinulata MT160654.1 strain U11 tufA as U.laetevirens
U.lacinulata MT160655.1 strain U99 tufA
U.lacinulata MT160659.1 strain U12 tufA
U.lacinulata MT160661.1 strain U14 tufA
U.lacinulata MT160665.1 strain U20 tufA
U.lacinulata MT160669.1 strain U103 tufA
U.lacinulata MT160680.1 strain U35 tufA
U.lacinulata MT160687.1 strain U2 tufA
U.lacinulata MT160690.1 strain U5 tufA
U.lacinulata MT160692.1 strain U7 tufA
U.lacinulata MT160693.1 strain U8 tufA
U.lacinulata MT160697.1 strain U41 tufA
U.lacinulata MT160699.1 strain U43 tufA
U.lacinulata MT160704.1 strain U49 tufA
U.lacinulata MT160706.1 strain U108 tufA
U.lacinulata MT160709.1 strain U104 tufA
U.lacinulata MT160714.1 strain U55 tufA
U.lacinulata MT160718.1 strain U59 tufA
U.lacinulata MT160724.1 strain U110 tufA
U.lacinulata MT160726.1 strain U62 tufA
U.lacinulata MT160729.1 strain U63 tufA
U.lacinulata MT160731.1 strain U65 tufA
U.lacinulata MT160735.1 strain U71 tufA
U.lacinulata MT160739.1 strain U75 tufA
U.lacinulata MT160740.1 strain U74 tufA
U.lacinulata MT160657.1 strain U98 tufA
U.lacinulata MT160658.1 strain U13 tufA
U.lacinulata MT160689.1 strain U4 tufA
U.lacinulata MT160720.1 strain U58 tufA
U.lacinulata MW543061.1 tufA lectotype L0054997 complete genome 10724
U.sp. A AF-2021 MT160638.1 tufA strain U81
U.sp. A AF-2021 MT160644.1 tufA strain U87
U.sp. A AF-2021 MT160645.1 tufA strain U89
U.sp. A AF-2021 MT160646.1 tufA strain U91
U.sp. A AF-2021 MT160648.1 tufA strain U93
U.sp. A AF-2021 MT160656.1 tufA strain U10
U.sp. A AF-2021 MT160660.1 strain U15 tufA
U.sp. A AF-2021 MT160663.1 strain U19 tufA
U.sp. A AF-2021 MT160666.1 strain U22 tufA
U.sp. A AF-2021 MT160667.1 strain U21 tufA
U.sp. A AF-2021 MT160668.1 strain U24 tufA
U.sp. A AF-2021 MT160673.1 strain U27 tufA
U.sp. A AF-2021 MT160675.1 strain U29 tufA
U.sp. A AF-2021 MT160677.1 strain U30 tufA
U.sp. A AF-2021 MT160681.1 strain U34 tufA
U.sp. A AF-2021 MT160683.1 strain U36 tufA complete cds
U.sp. A AF-2021 MT160688.1 strain U3 tufA
U.sp. A AF-2021 MT160694.1 strain U9 tufA
U.sp. A AF-2021 MT160696.1 strain U42 tufA
U.sp. A AF-2021 MT160710.1 strain U51 tufA
U.sp. A AF-2021 MT160711.1 strain U50 tufA
U.sp. A AF-2021 MT160717.1 strain U56 tufA
U.sp. A AF-2021 MT160723.1 strain U111 tufA
U.sp. A AF-2021 MT160730.1 strain U66 tufA
U.sp. A AF-2021 MT160734.1 strain U69 tufA
U.sp. A AF-2021 MT160738.1 strain U72 tufA
U.sp. A AF-2021 MT160741.1 strain U77 tufA
U.sp. A AF-2021 MT160742.1 strain U76 tufA
U.sp. A AF-2021 MT160743.1 strain U79 tufA
U.gigantea MT160649.1 strain U92 tufA
U.gigantea MT160672.1 strain U101 tufA
U.gigantea MT160676.1 strain U31 tufA
U.gigantea MT160682.1 strain U37 tufA
U.gigantea MT160698.1 strain U44 tufA
U.gigantea MT160712.1 strain U53 tufA
U.gigantea MT160713.1 strain U52 tufA
U.gigantea MT160715.1 strain U54 tufA
U.gigantea MT160716.1 strain U57 tufA
U.gigantea MT160727.1 strain U61 tufA
U.flexuosa KX579943.1 tufA complete genome 89414 bp
U.linza KX058323.1 isolate QD08 tufA complete genome 86726 bp
U.prolifera KX342867.1 tufA complete genome 93066
U.sp. KP720616.1 UNA00071828 tufA complete genome
U.sp. MW699788.1 strain Q253 tufA comple genome 88878 bp