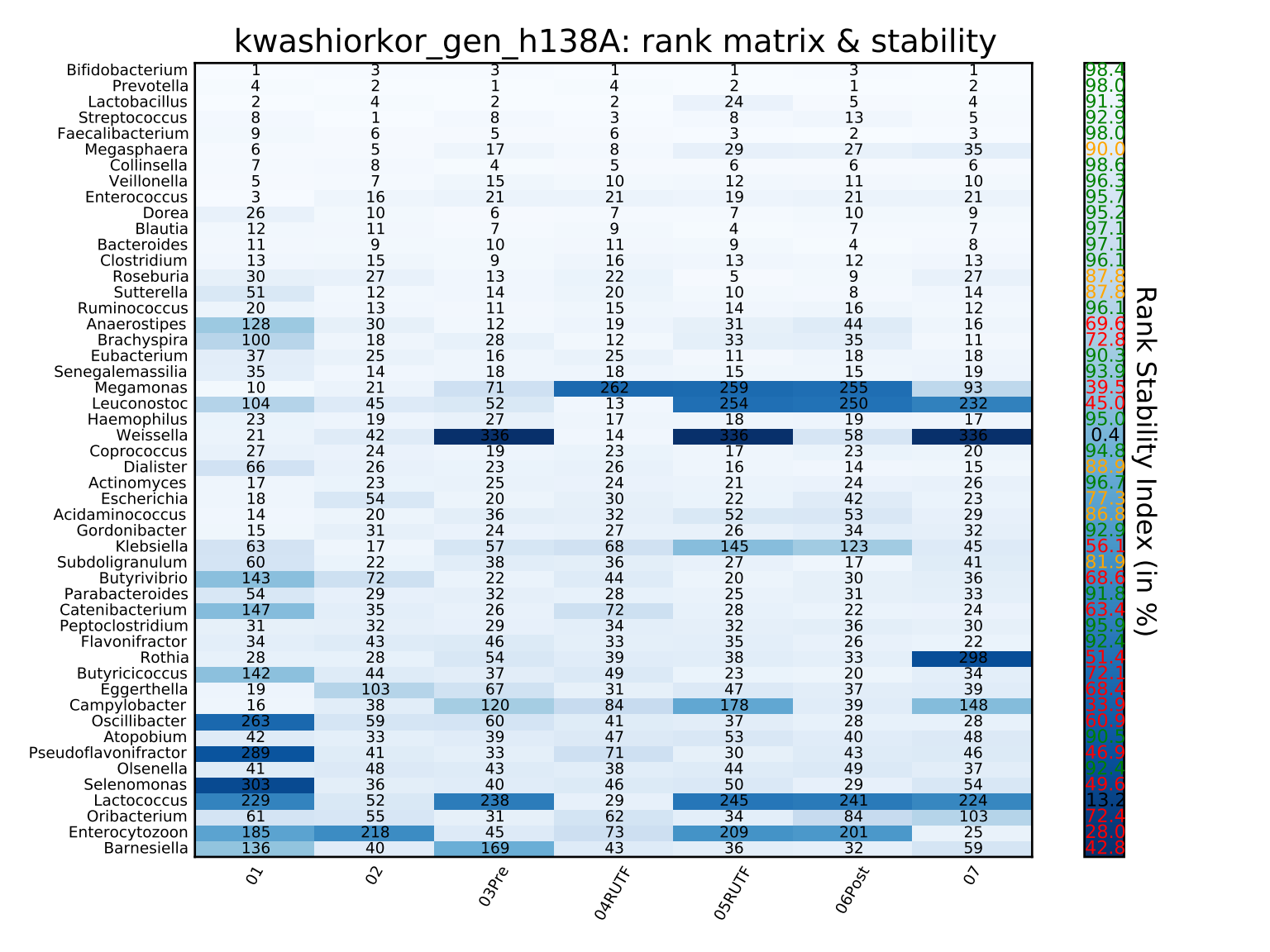
kwashiorkor_gen_h138A: rank matrix & stability
Bifidobacterium 1 3 3 1 1 3 1 98.4
Prevotella 4 2 1 4 2 1 2 98.0
Lactobacillus 2 4 2 2 24 5 4 91.3
Streptococcus 8 1 8 3 8 13 5 92.9
Faecalibacterium 9 6 5 6 3 2 3 98.0
Megasphaera 6 5 17 8 29 27 35 90.0
Collinsella 7 8 4 5 6 6 6 98.6
Veillonella 5 7 15 10 12 11 10 96.3
Enterococcus 3 16 21 21 19 21 21 95.7
Dorea 26 10 6 7 7 10 9 95.2
Blautia 12 11 7 9 4 7 7 97.1
Bacteroides 11 9 10 11 9 4 8 97.1
Clostridium 13 15 9 16 13 12 13 96.1
Roseburia 30 27 13 22 5 9 27 87.8
Sutterella 51 12 14 20 10 8 14 87.8
Ruminococcus 20 13 11 15 14 16 12 96.1
Anaerostipes 128 30 12 19 31 44 16 69.6
Brachyspira 100 18 28 12 33 35 11 72.8
Eubacterium 37 25 16 25 11 18 18 90.3
Senegalemassilia 35 14 18 18 15 15 19 93.9
Megamonas 10 21 71 262 259 255 93 39.5
Leuconostoc 104 45 52 13 254 250 232 45.0
Haemophilus 23 19 27 17 18 19 17 95.0
Weissella 21 42 336 14 336 58 336 0.4
Coprococcus 27 24 19 23 17 23 20 94.8
Dialister 66 26 23 26 16 14 15 88.9
Actinomyces 17 23 25 24 21 24 26 96.7
Escherichia 18 54 20 30 22 42 23 77.3
Acidaminococcus 14 20 36 32 52 53 29 86.8
Gordonibacter 15 31 24 27 26 34 32 92.9
Klebsiella 63 17 57 68 145 123 45 56.1
Subdoligranulum 60 22 38 36 27 17 41 81.9
Butyrivibrio 143 72 22 44 20 30 36 68.6
Parabacteroides 54 29 32 28 25 31 33 91.8
Catenibacterium 147 35 26 72 28 22 24 63.4
Peptoclostridium 31 32 29 34 32 36 30 95.9
Flavonifractor 34 43 46 33 35 26 22 92.4
Rothia 28 28 54 39 38 33 298 51.4
Butyricicoccus 142 44 37 49 23 20 34 72.1
Eggerthella 19 103 67 31 47 37 39 68.4
Campylobacter 16 38 120 84 178 39 148 33.9
Oscillibacter 263 59 60 41 37 28 28 60.9
Atopobium 42 33 39 47 53 40 48 90.5
Pseudoflavonifractor 289 41 33 71 30 43 46 46.9
Olsenella 41 48 43 38 44 49 37 92.4
Selenomonas 303 36 40 46 50 29 54 49.6
Lactococcus 229 52 238 29 245 241 224 13.2
Oribacterium 61 55 31 62 34 84 103 72.4
Enterocytozoon 185 218 45 73 209 201 25 28.0
Barnesiella 136 40 169 43 36 32 59 42.8
01
02
03Pre
04RUTF
05RUTF
06Post
07
Rank Stability Index (in %)