Streptococcus salivarius 1.00 Shared genome fraction 0.95 0.90 0.85 0.80 0.75 99 95 96 97 98 100 Average nucleotide identity (%) 1.2 non-parametric KDE Midpoint(s) 1.0 8.0 Density 0.6 0.4 0.2 0.0 96 95 97 98 99 100 Average nucleotide identity (%)