Rickettsia japonica 1.000 -Shared genome fraction 0.999 0.998 0.997 0.996 0.995 0.994 0.993 99.980 99.985 99.990 99.995 100.000 99.970 99.975 Average nucleotide identity (%) non-parametric KDE 160 Midpoint(s) 140 120 Density 100 80 60

