



Pairwise gANI

saccharibacter_am169	100	99.3	99.3	99.3	99.3	99.2	98.8	83	76	71.9	71.9	saccharibacter_am169
parasaccharibacter_apium	99.3	100	99.4	99.4	99.4	99.3	98.9	83	76	71.9	71.9	parasaccharibacter_apium
parasaccharibacter_apium_a2	99.3	99.4	100	100	100	99.4	98.9	83	76	71.9	71.8	parasaccharibacter_apium_a2
parasaccharibacter_apium_b8	99.3	99.4	100	100	100	99.4	98.9	83	76	71.9	71.8	parasaccharibacter_apium_b8
parasaccharibacter_apium_c6	99.3	99.4	100	100	100	99.4	98.9	83	76	71.9	71.8	parasaccharibacter_apium_c6
saccharibacter_3.A.1	99.2	99.3	99.4	99.4	99.4	100	98.9	83	76	71.9	71.8	saccharibacter_3.A.1
saccharibacter_m18	98.8	98.9	98.9	98.9	98.9	98.9	100	83.1	76.1	71.9	71.8	saccharibacter_m18
bombella_intestini	83	83	83	83	83	83	83.1	100	76.4	71.9	71.5	bombella_intestini
parasaccharibacter_apium_as	76	76	76	76	76	76	76.1	76.4	100	71.9	70.5	parasaccharibacter_apium_as
saccharibacter_floricola	71.9	71.9	71.9	71.9	71.9	71.9	71.9	71.9	71.9	100	70.4	saccharibacter_floricola
gluconobacter_oxydans	71.9	71.9	71.8	71.8	71.8	71.8	71.8	71.5	70.5	70.4	100	gluconobacter_oxydans