

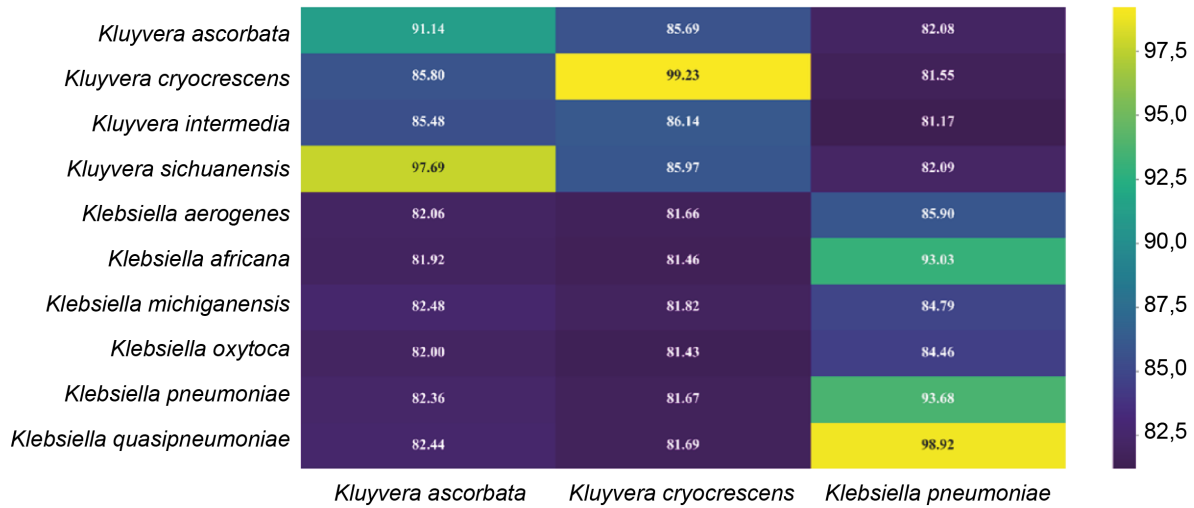


Figure 1. Average Nucleotide Identity (ANI) analysis of the assembled genomes. Heatmap showing pairwise ANI values between the three assembled genomes initially identified as *Kluyvera ascorbata*, *Kluyvera cryocrescens*, and *Klebsiella pneumoniae* and reference genomes from the genera *Kluyvera* and *Klebsiella*. The highest ANI values were observed between isolate BOG01-6 and *Kluyvera sichuanensis* (97.69%), isolate BOG02-3 and *Kluyvera cryocrescens* (99.23%), and isolate BOG01-4 and *Klebsiella quasipneumoniae* (98.92%), supporting their taxonomic reassignment. Color intensity represents ANI similarity percentages.