



**FIGURE S2. ANI percentage identity of sequenced *Duganella*, *Pseudoduganella* and *Janthinobacterium* strains.** The ANIm was constructed using genomic data published or extracted from <http://www.jgi.doe.gov/> using the IMG software at <https://img.jgi.doe.gov/cgi-bin/er/main.cg>. Genome information was obtained from the following genome files deposited at GenBank and the respective publications. For strain information consult material and methods, “Bacterial strains and growth”.