



**Supplemental Figure S1.** Histograms of pairwise ANI values between 2,218 *B. cereus* group genomes with N50 > 20 Kbp. For each of 18 published species and 3 proposed putative species, ANI values calculated between two genomes for which one or more genome(s) shared  $\geq 95$  ANI with the species type strain or reference genome are colored in pink. ANI values calculated between two genomes for which both genomes shared < 95 ANI with the species type strain/reference genome are colored in gray. See Supplemental Table S1 for details regarding species type strain genomes.