



**Additional File 3.** Maximum likelihood tree of SGAir strains, 55 complete genomes of *Bacillus cereus* retrieved from NCBI, with outgroups, *Escherichia coli* 157 H7 str. Sakai, *Bacillus subtilis* subsp. *subtilis* str., and *Bacillus subtilis* subsp. *spizizenii* TU-B-10.