



Figure S1. Maximum parsimony phylogenetic tree of partial 16S rDNA sequences of isolates in the ILSI NA *Cronobacter* collection and relevant type strains. Strain designations include Cornell University Food Safety Lab designations that start with “FSL” as well as 4-digit Health Canada designations. Relevant type strain sequences were obtained from the Ribosomal Database project. Each type strain sequence is denoted by genus and species; strain; GenBank accession number. The *Escherichia coli* ATCC11775 type strain sequence was included as an outgroup. Numerical node labels represent the percentage of bootstrap replicates (n=500) that supported the respective node. Only bootstrap values greater than 50 are shown. Scale represents estimated substitutions per site.