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.