



Figure S1: Development of standard curves for *C. coli* (Panel A) and *C. jejuni* (Panel B) generated based on amplification of DNA from increasing number of cells (A–F: 10^7 , 10^6 , 10^5 , 10^4 , 10^3 , and 10^2 cells mL^{-1}) using species-specific TaqMan probe and SYBR Green-based quantitative real-time PCR assays.