



Supplemental Figure 2. Quantification of *cld* genes per 16S rRNA gene with depth in the laboratory-scale bioreactor. Dark gray bars represent relative gene quantities for the ~60% perchlorate-removal condition; light gray bars represent relative gene quantities for the ~20% perchlorate-removal condition. Error bars represent the propagated standard deviations of triplicate *cld* and 16S rRNA qPCR assays for the pooled DNA samples.