

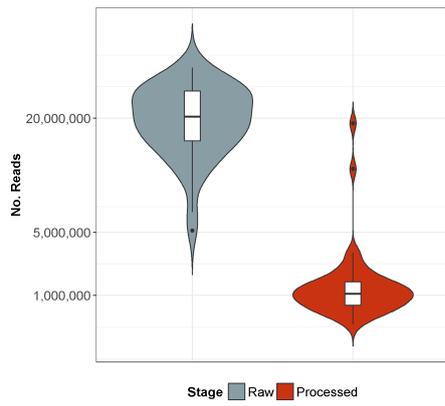
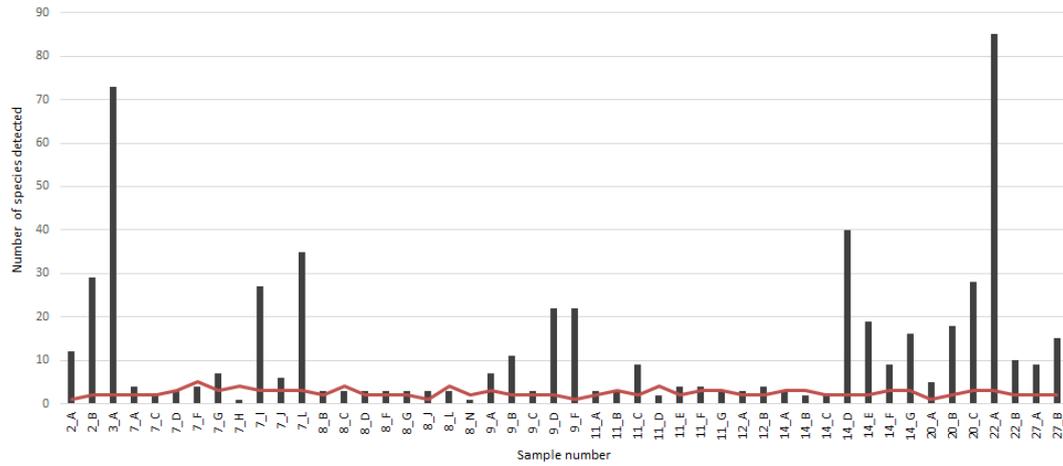
A**B**

Figure S1: Sequencing metrics and quality control. A) number of raw reads versus processed reads and B) Number of species detected by NextSeq and culture. Black bars represent number of species detected by NextSeq and red line represents the number of species detected by culture for each sample.