

SUPPLEMENTARY MATERIAL

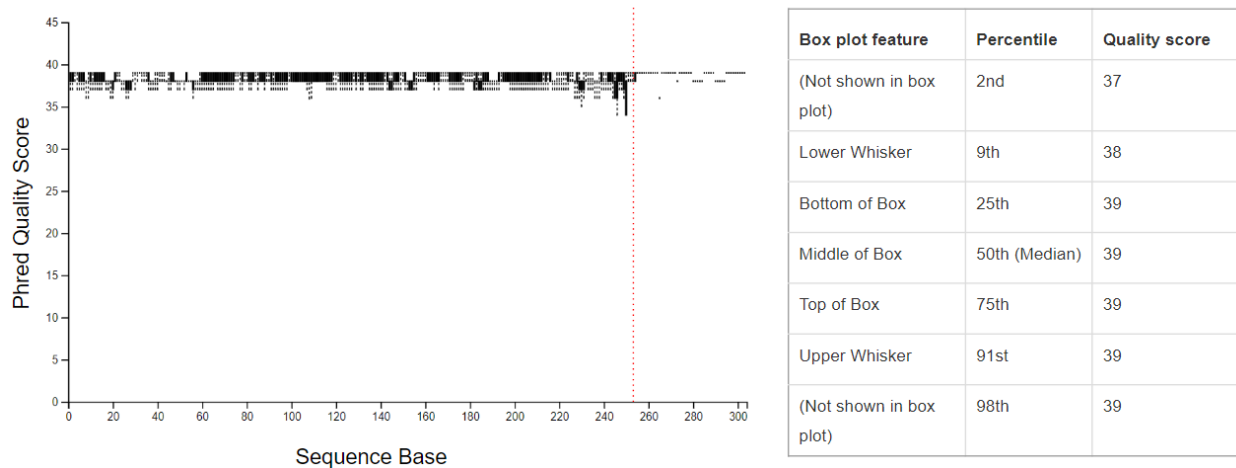


FIG. S1 Quality score and statistics of demultiplexed sequences. Left: Plot of the phred quality score of the forward sequences in the McClorry et al. dataset. The red dotted line indicates the truncation length we used of 253 nucleotides. Right: percentile values for the quality score at 253 nucleotides, with a median Phred score of 39.