

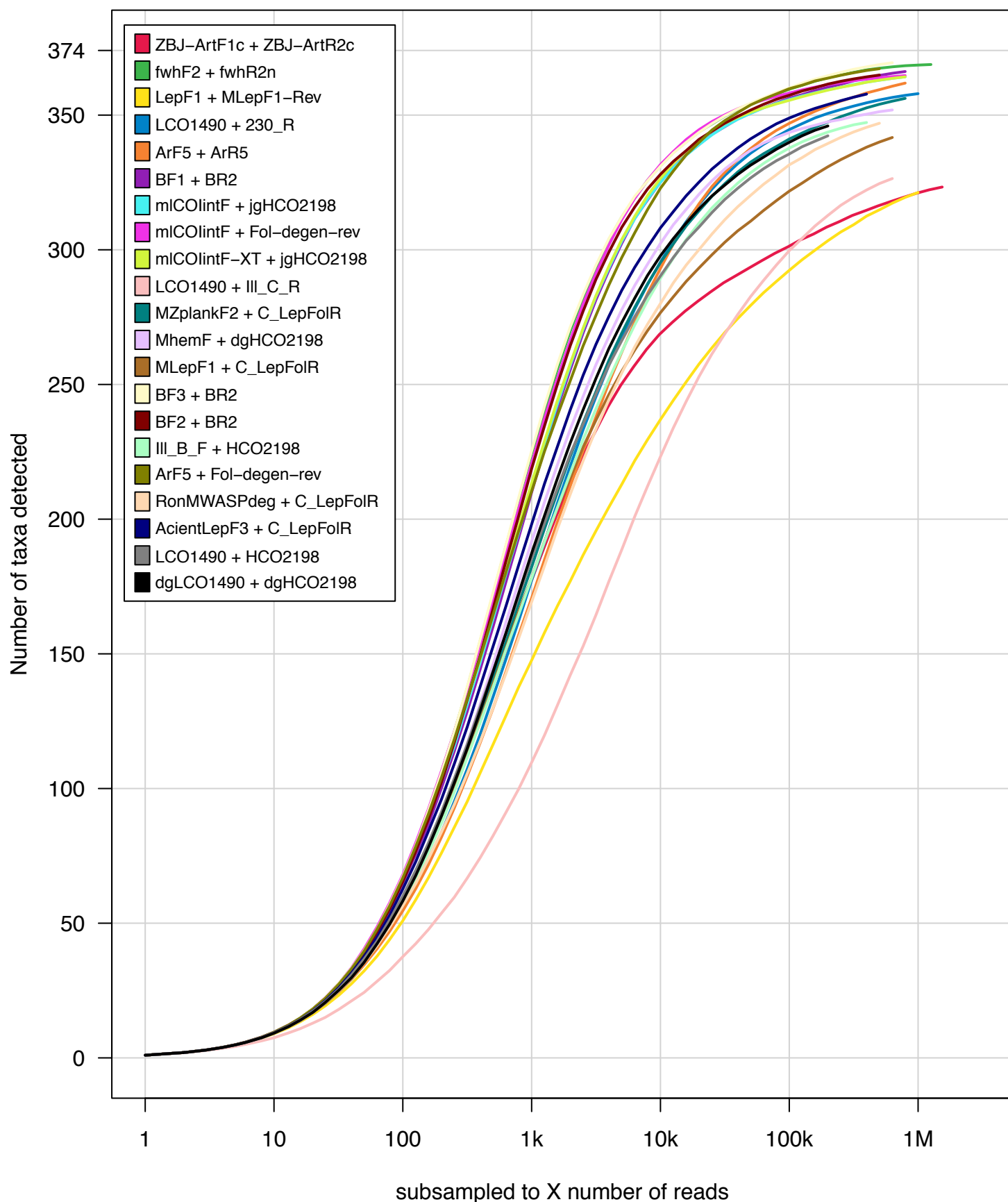


Figure S11: Rarefaction curve showing the amount of taxa recovered at different sequencing depth for metabarcoding the mock sample with all 21 primer sets. For each data point 1000 subsamples were taken.