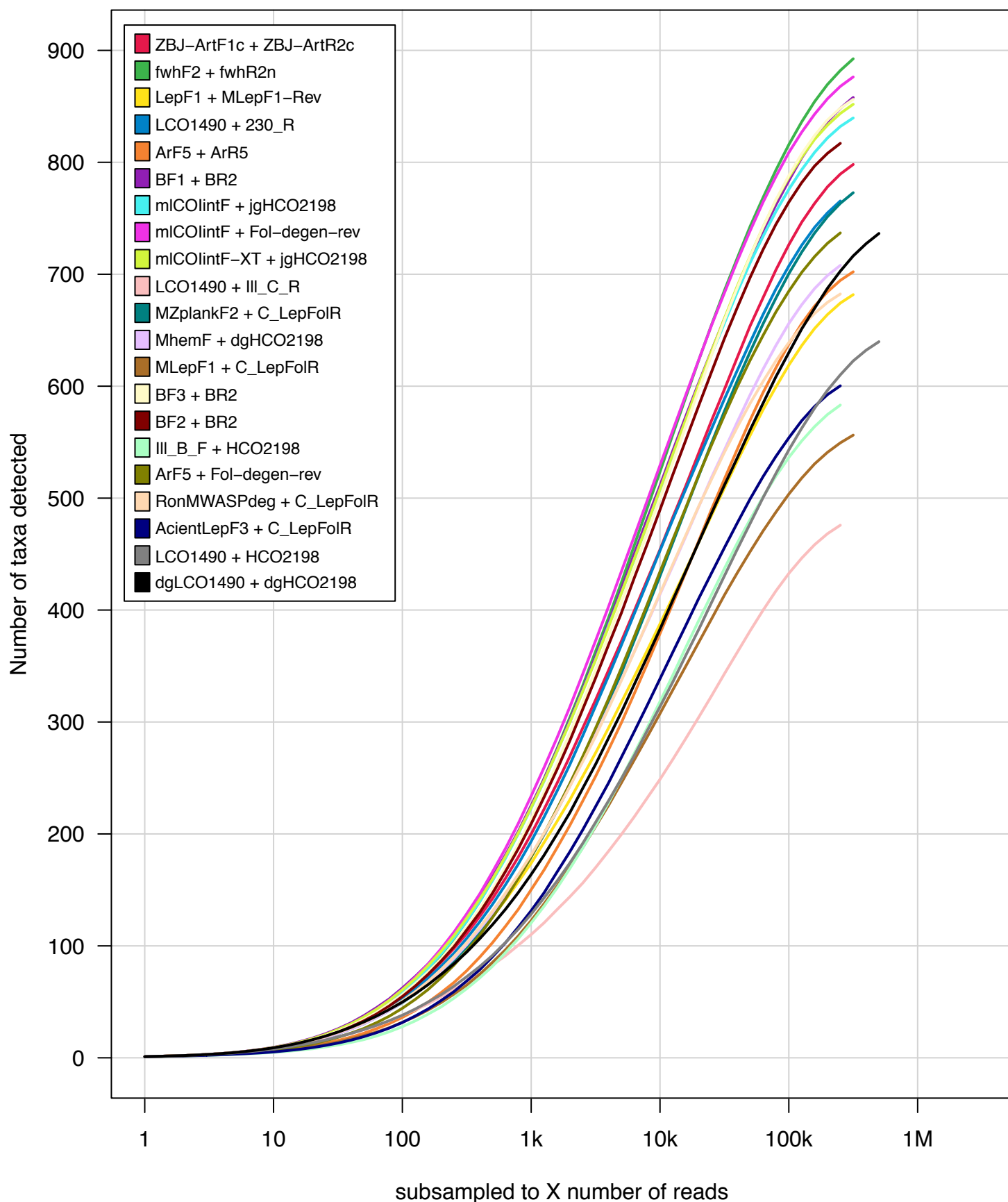




**Figure S19:** Rarefaction curve showing the amount of taxa recovered from the malaise trap sample at different sequencing depth for all 21 primer sets. For each data point 1000 subsamples were taken.