

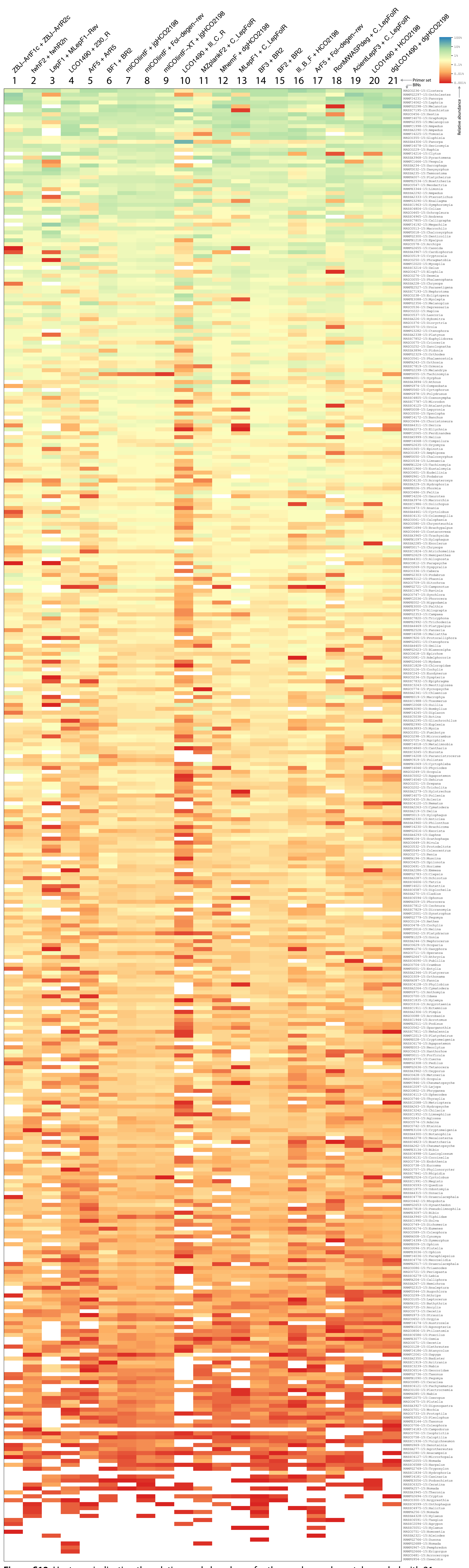


Figure S12: Heatmap indicating the relative read abundance for the mock sample metabarcoded with 21 different primer sets. Reads are subsampled to 100,000 reads per sample (reads with abundance below 0.001% where discarded). BINs are sorted by average relative abundance. See Table S1 for the raw data.