Supplemental Figure 1. Benchmarking for elapsed runtime and memory for HAYSTAC, Sigma, Kraken2/Bracken, Krakenuniq and MALT when analysing samples of 10 K, 100 K and 1 M reads against a database of 500 species. Memory remains constant as sample size increases, and runtime in most methods (other than Kraken2/Bracken and Krakenuniq) scales with the database size rather than input sample size.)