

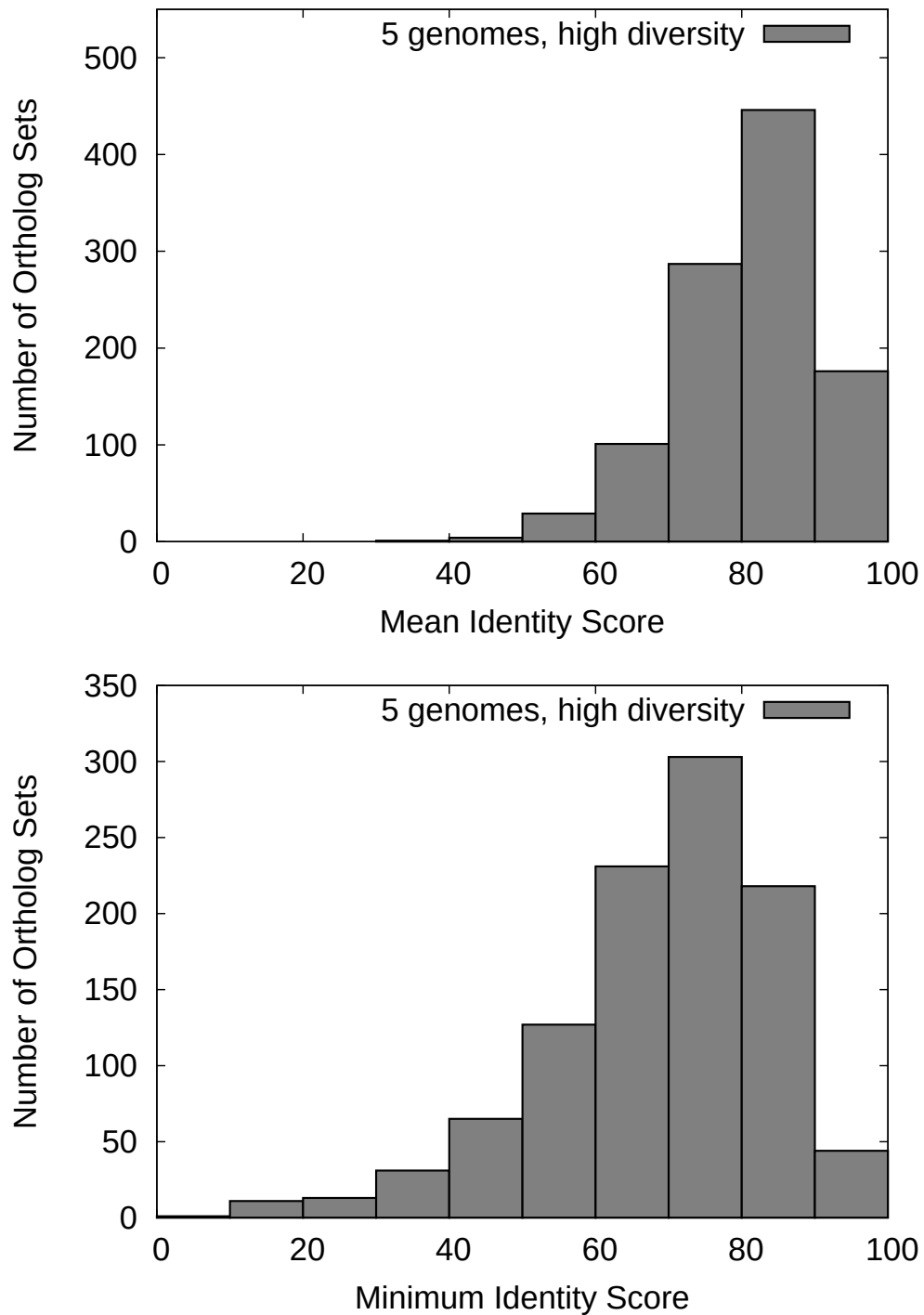


Figure S3: Histogram of mean (top) and minimum (bottom) identity score between genes in ortholog sets derived from the high diversity, 5 genome set.