



reads: metagenome or
metatranscriptome



10^7 /sample

assembly



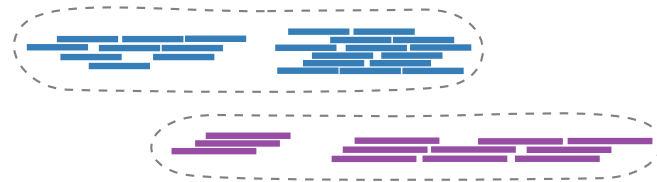
contigs



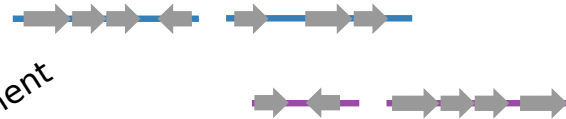
binning



genome bins



call genes



alignment



contig coverage
or
expression profiles

