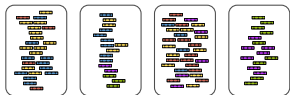


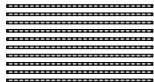
Read Libraries



Contigs

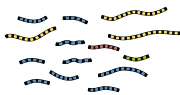


Reference Genome Sequences



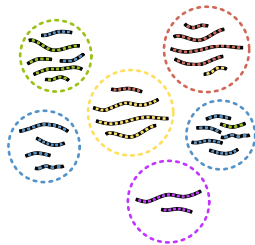
(b) enrich

Reduced Contigs



(c) select

Genome Bins



(a) (re-)cluster or (re-)classify

(d) refine

Reduced Genome Bins

