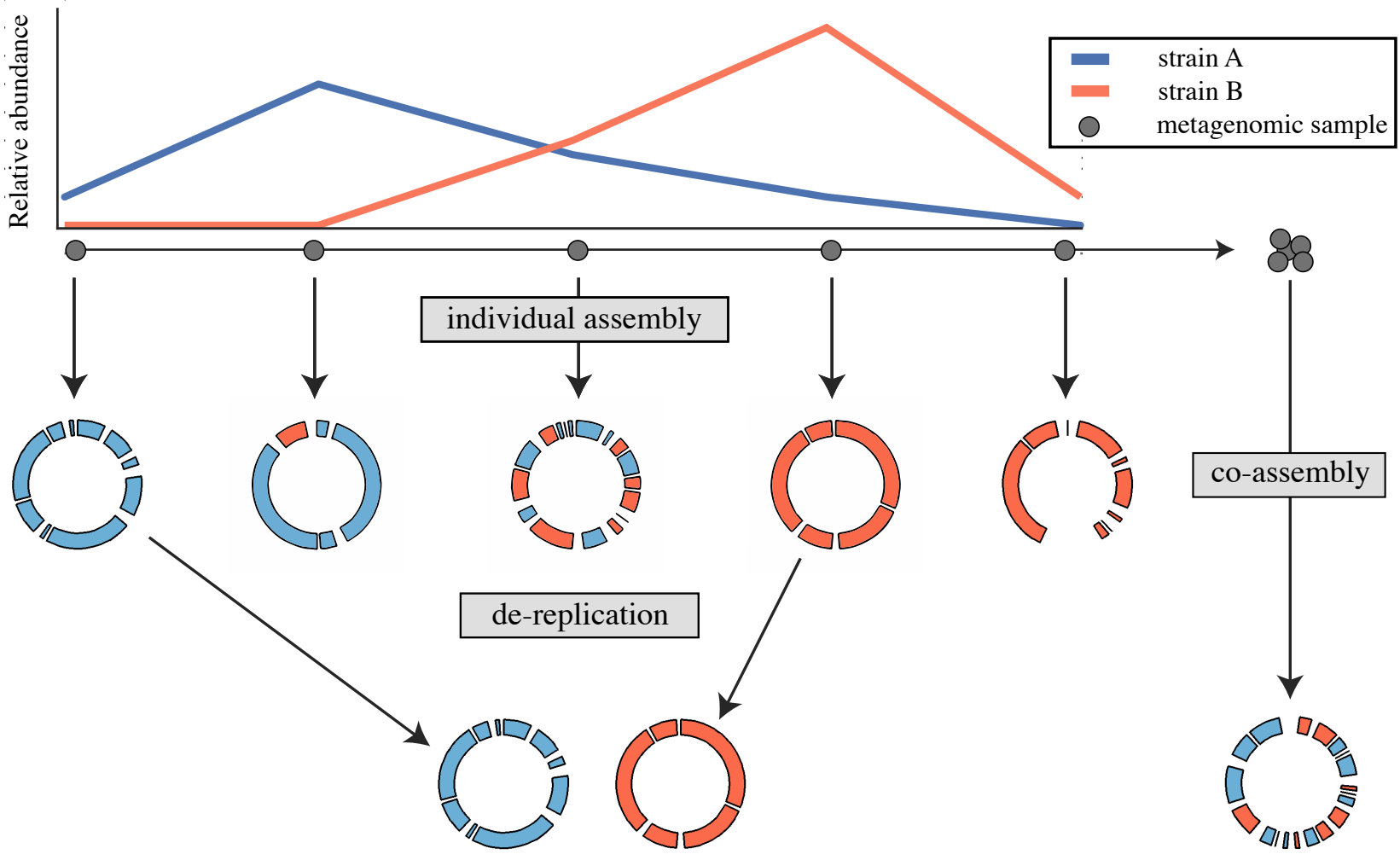
Relative abundance
strain A
strain B
metagenomic sample
individual assembly
co-assembly
de-replication