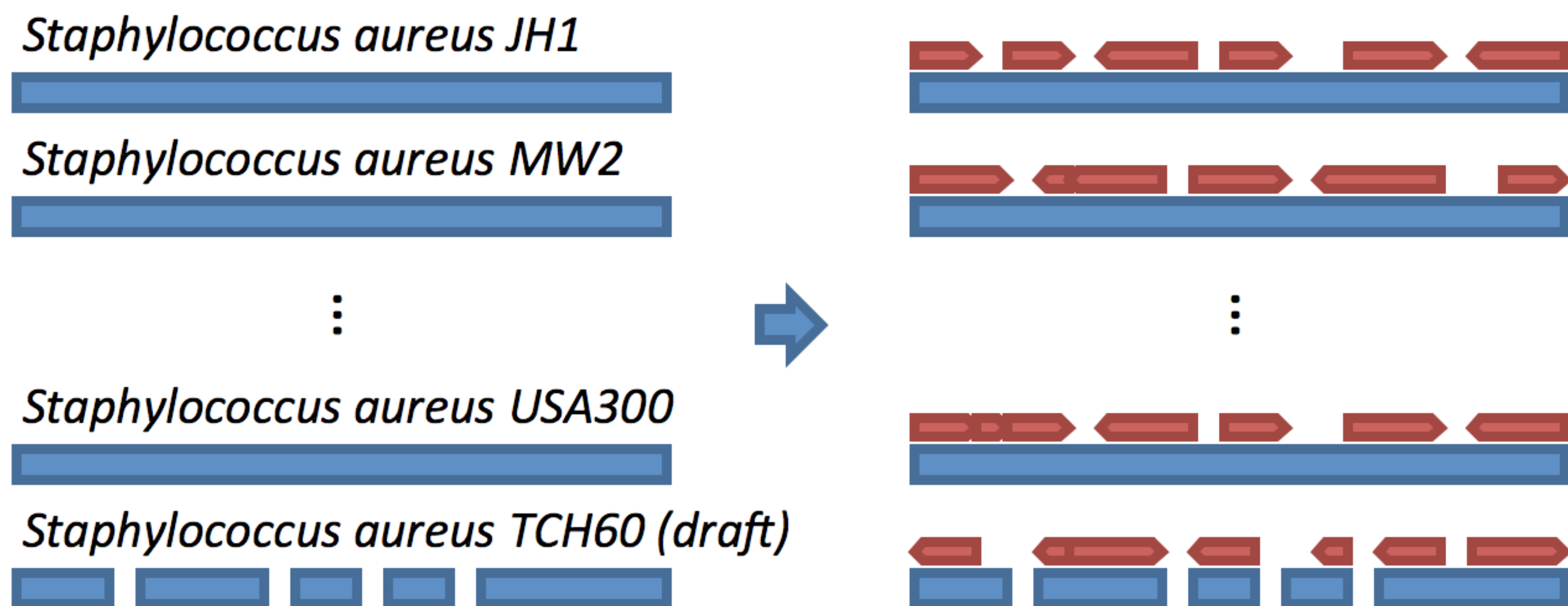
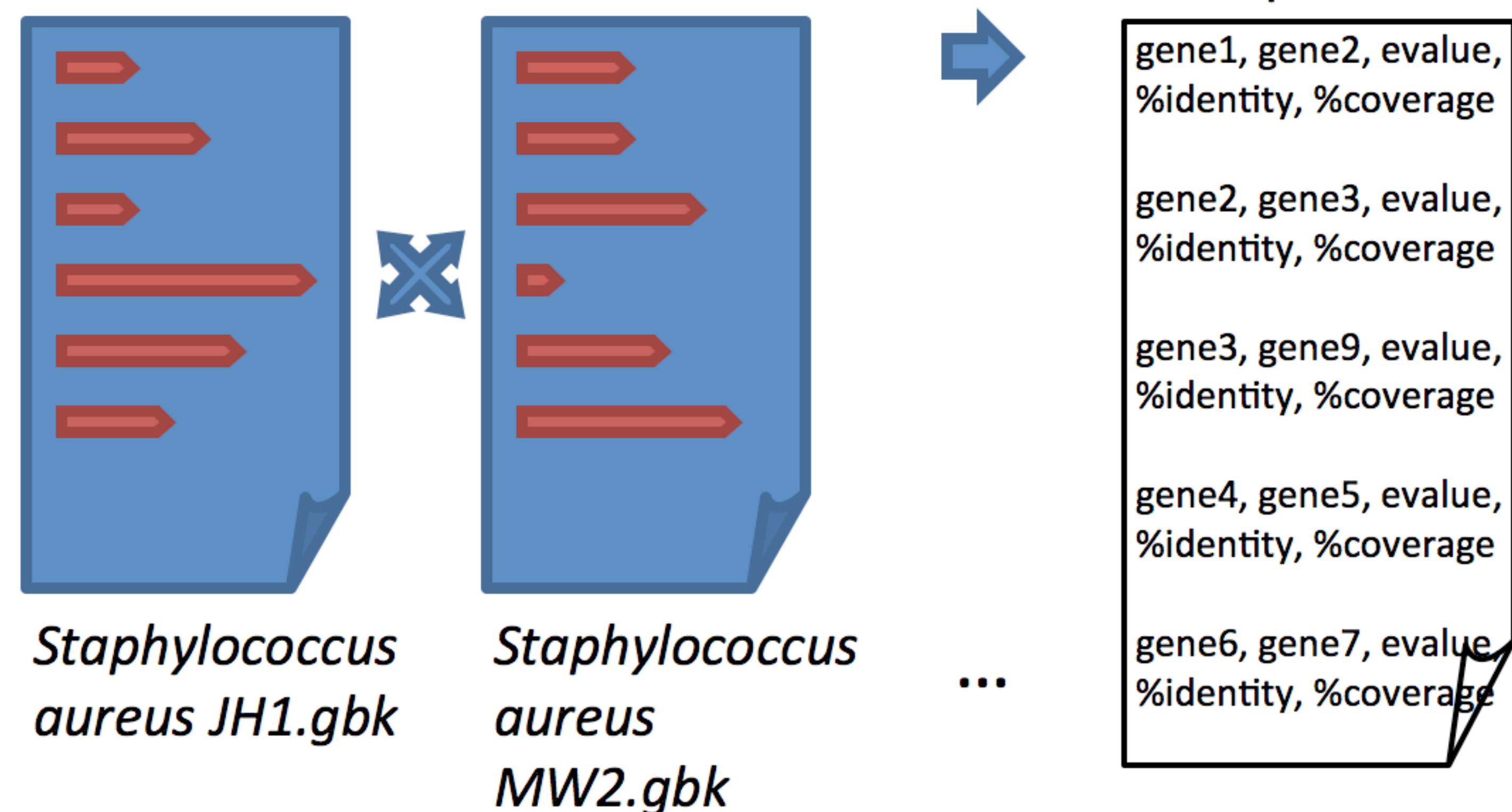


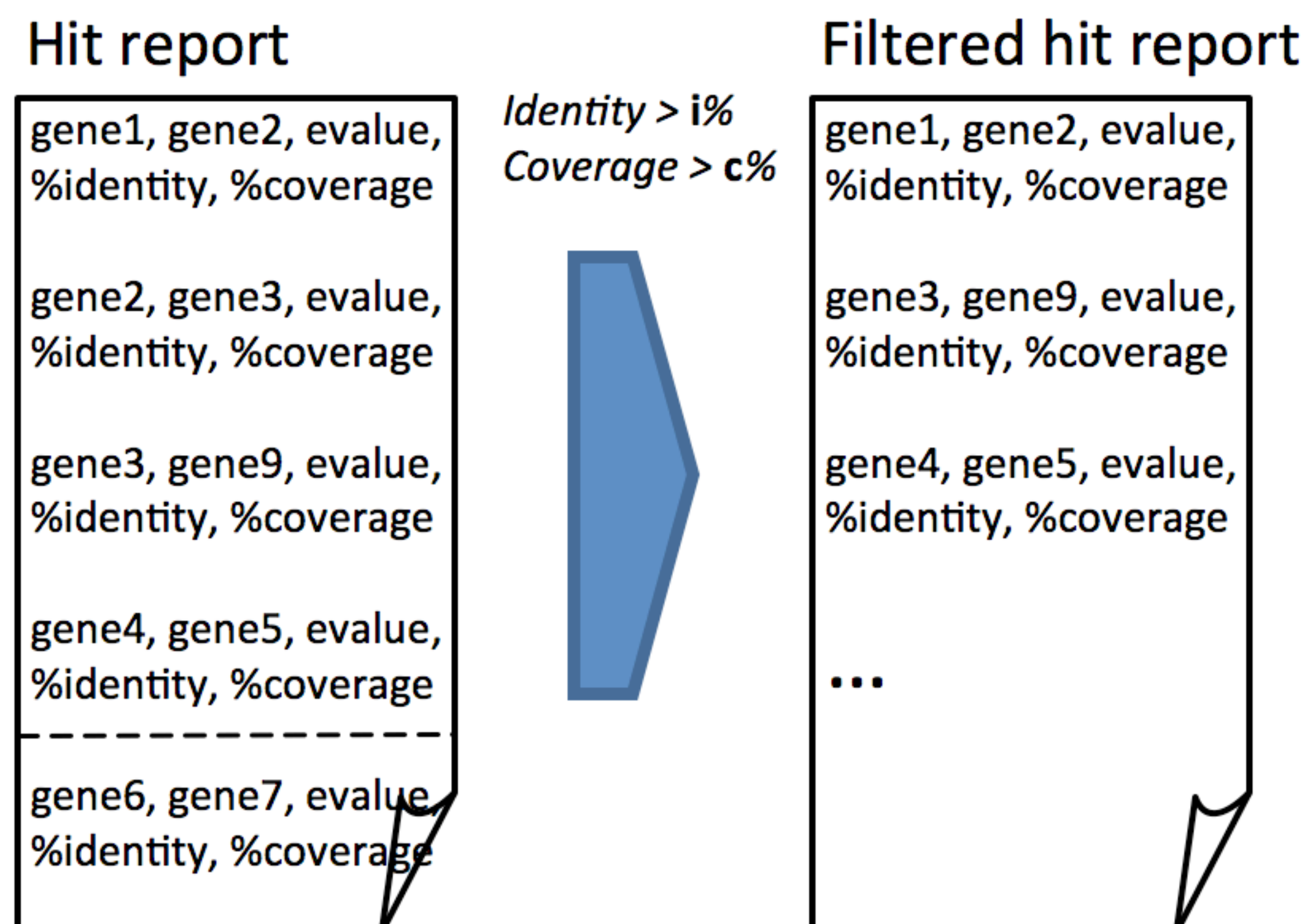
A CDS re-annotation



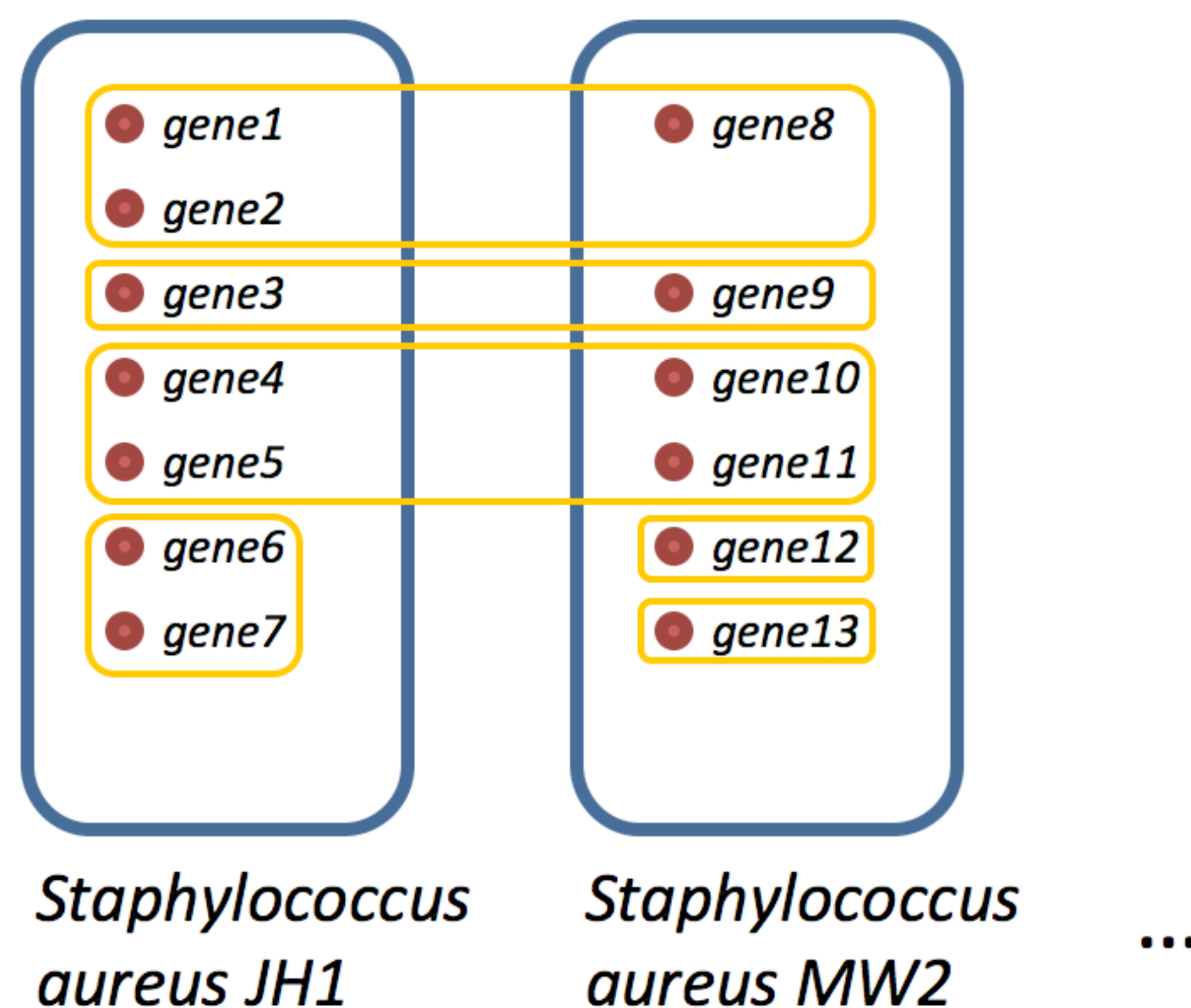
B Pairwise protein alignment



C Filtering of homologs



D Construction of gene families



E Count shared gene families

