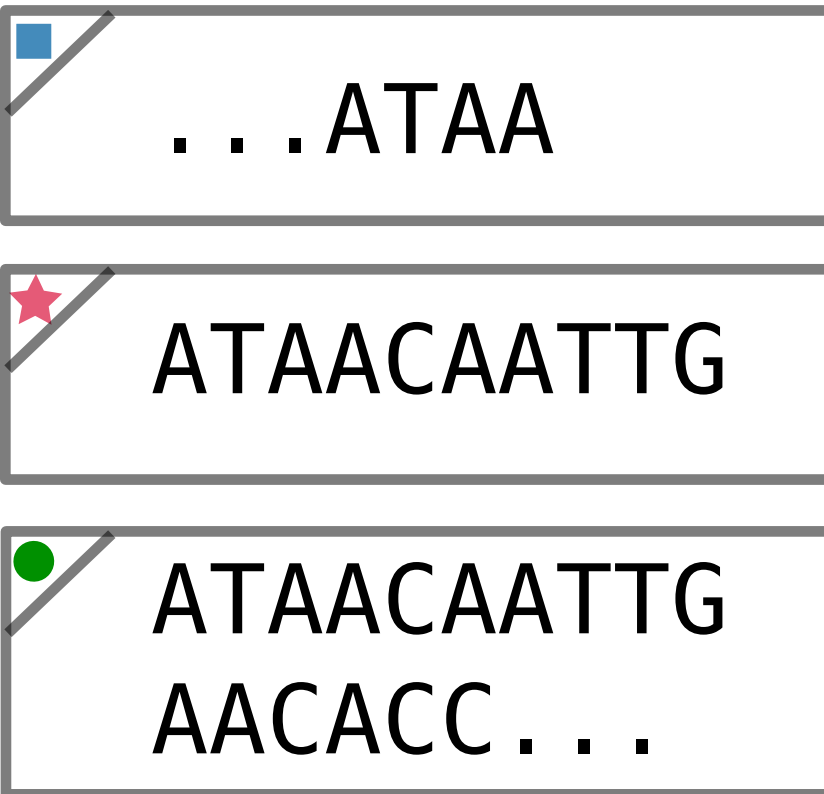
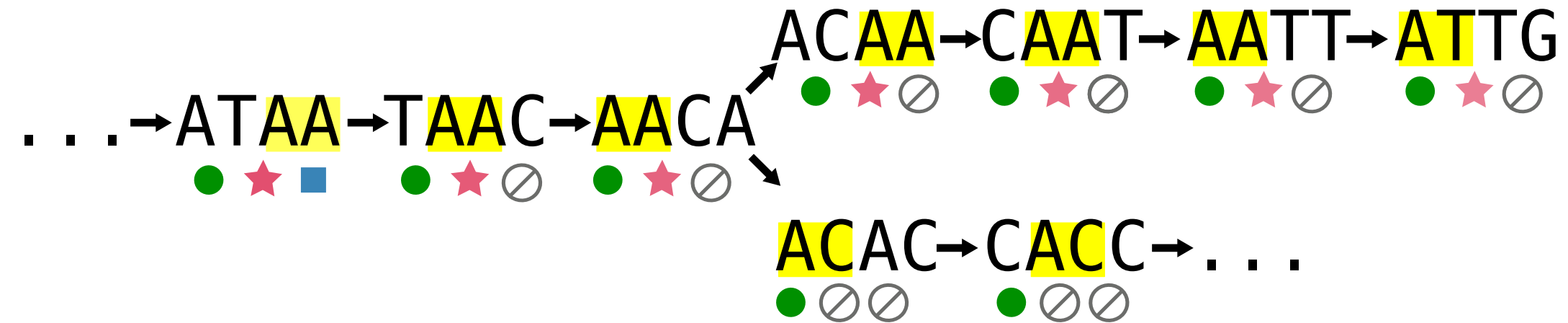


datasets: ● ★ ■



reads

de Bruijn graph (node-centric)



■ minimizer (m=2)

⊘ absence from a dataset

A straightforward SPSS: the k-mer set

{ATAA, TAAC, AACCA, ACAA, CAAT, AATT, ATTG, ACAC, CACC} 28 nucleotides