A straightforward SPSS: the k-mer set itself {ATAA, TAAC, AACA, ACAA, CAAT, AATT, ATTG, ACAC, CACC} 28 nucleotides

ACAC→CACC→...

de Bruijn graph (node-centric)

ACAA→CAAT→AATT→ATTG ...→ATAA→TAAC→AACA