Partition Function and Base Pair Propabilities for Sequence Alignment(s)
Computing the structure with maximum expected accuracy (MEA)
Partition Function for two Hybridized Sequences as a sequence Process
Calculate Partition Functions for a Distance Based Partitioning
Computing the centroid structure
Partition Function for two Hybridized Sequences
Partition functions for locally stable secondary structures
Computing Partition Functions and Pair Propabilities
RNA Secondary Structure Prediction