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