"To *B* or not to *B*" in Nucleic Acids Chemistry

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In this presentation, I will provide an overview of the basic concepts, methods, and applications of predicting the stabilities of nucleic acid structures. We explain the theory of the most successful prediction method based on a nearest-neighbor (NN) model. To improve the versality of prediction, corrections for various solution conditions considered hydration have been investigated. I also describe advances in the prediction of non-canonical structures. Finally, studies of intracellular analysis and prediction are discussed for the application of NN parameters

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