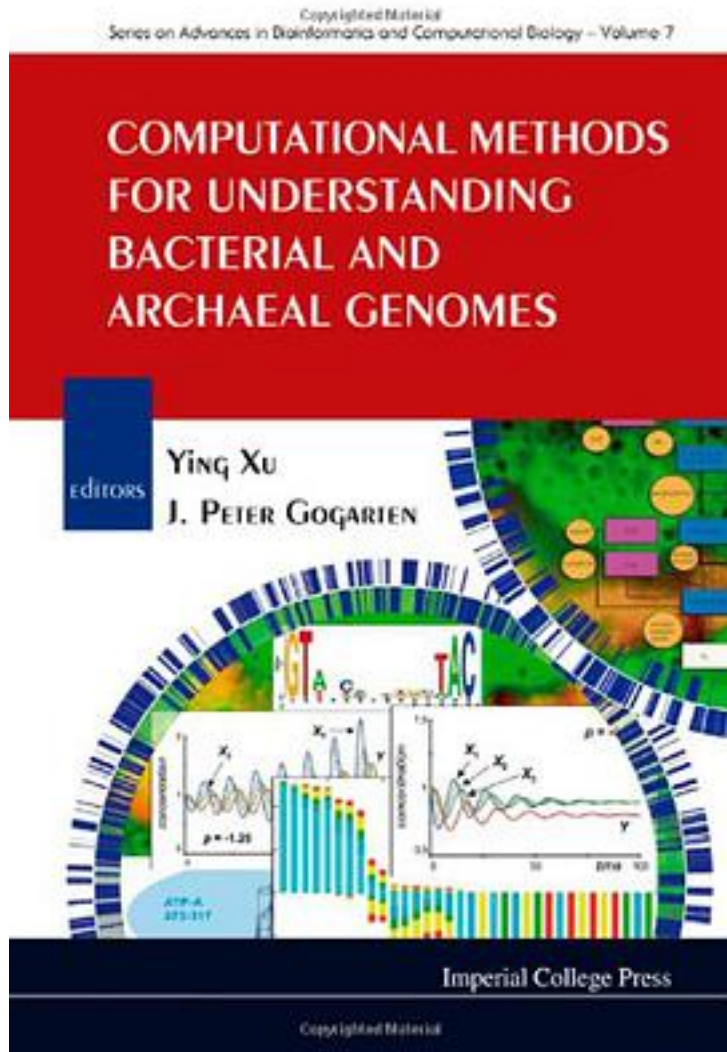


Computational Methods for Understanding Bacterial and Archaeal Genomes



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