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## Significance of Hybrid Evolutionary Computation for Ab Initio Protein Folding Prediction

DOI: 10.1007/978-3-540-73297-6\_10  
Studies in Computational Intelligence, 2007, , 241-268.

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1	DFS Based Partial Pathways in GA for Protein Structure Prediction. <i>Lecture Notes in Computer Science</i> , <b>2008</b> , 41-53	0.9	2