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By George C. Shields;Paul G. Seybold**



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Modular polyketide synthases substrate specificities of uncharacterized PKS clusters listed in Table 6 are bonafide blind predictions by our computational approach.

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Computational Approaches for Protein Function Prediction: A Survey Gaurav Pandey, Vipin Kumar and Michael Steinbach Department of Computer Science and Engineering

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Zhang, M.Q.: Computational Prediction of Eukaryotic Protein-Coding Genes. Nature Reviews Genetics 3, 698 709 (2002) View Article

Computational approaches for protein function

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Computational approaches can be developed for automatically and productivities can be used for collaboration prediction. Our approach is

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The pressing need for the prediction of protein function has led to the development of hundreds of different techniques for solving this problem by computational means.

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we summarize progress in the past two years in the areas of computational approaches to RNA tertiary
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