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Doi:10.1093/bioinformatics ...The Improved Quality Is Due To Newly Designed Features And Careful Parameterization Of The SVM. The Features Incorporate Amino Acids Characterized By The Highest And The Lowest Propensities To Constitute Catalytic Residues, Gly ... Catalytic Residue Prediction (Petrova And Wu, ... Mar 9th, 2024.

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Algorithms On Strings, Trees And Sequences Note That The Converse Is Not True: \ ... Similar Sequences Yield Similar Structures, But Quite Different Sequences Can Produce Remarkably Similar Structures." Jan 6th, 2024.

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