

Three-dimensional structure of a new member of the α -amylase family

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We describe the 3-D structure of a new α -amylase family member. According to its amino acid sequence [1], The structure was solved using the method by [2] The enzyme adopts the structure of a classical (β/α)₈-barrel fold [3], described originally in the structure of an interesting enzyme [4], etc., etc.

Acknowledgement(s)

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