

# Three-dimensional structure of a new member of the $\alpha$ -amylase family

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We describe the 3-D structure of a new  $\alpha$ -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  $(\beta/\alpha)_8$ -barrel fold [3], described originally in the structure of an interesting enzyme [4], etc., etc.

## Acknowledgement(s)

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