

ANALYSIS



CONCLUSION

The structure of the protein is highly similar to that of the protein from *Escherichia coli*, which is a well-studied enzyme. The structure is a dimeric protein, with each subunit consisting of a beta-barrel domain and an alpha-helical domain. The beta-barrel domain is formed by eight beta-strands, and the alpha-helical domain is formed by two alpha-helices and a loop.

REFERENCES

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