

## PROTEIN MODELLING

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2. Explosion of Biological Sequence and Structure Data. Proteins have traditionally being studied individually. A protein of interest had its coding

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The purpose of this server is to make protein modelling accessible to all life science researchers worldwide. 4 Tertiary Protein Structure and Folds

Building a Better Fragment Library for De Novo Protein ...

Fragment-based approaches are the current standard for de novo protein structure prediction. These approaches rely on accurate and reliable fragment libraries to