

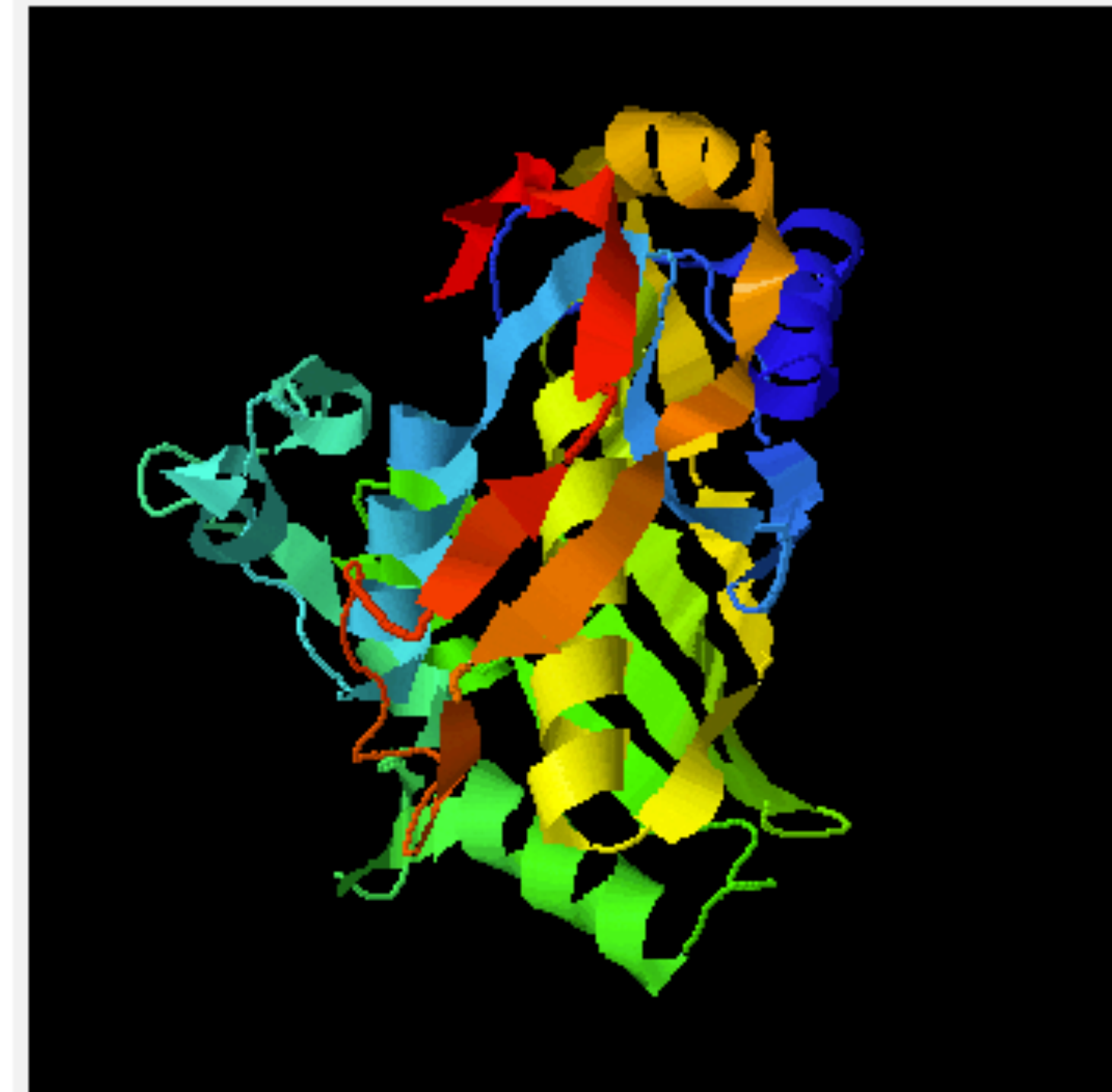
- The confidence of each model is quantitatively measured by C-score that is calculated based on the significance of threading template alignments and the convergence parameters of the structure assembly simulations. C-score is typically in the range of $[-5, 2]$, where a C-score of a higher value signifies a model with a higher confidence and vice-versa. TM-score and RMSD are estimated based on C-score and protein length...
- Is I-TASSER confident about its final model?

Top 5 final models predicted by I-TASSER

(For each target, I-TASSER simulations generate a large ensemble of structural quantitatively measured by C-score that is calculated based on the significance on C-score and protein length following the correlation observed between these rank models as seen in our benchmark tests. If the I-TASSER simulations conv

- [More about C-score](#)
- [Local structure accuracy profile of the top five models](#)

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- [Download Model 1](#)
- C-score=1.91 ([Read more about C-score](#))
- Estimated TM-score = 0.99 ± 0.04
- Estimated RMSD = $2.2 \pm 1.7 \text{ \AA}$

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- Is I-TASSER confident about its final model? Yes.

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