

CORRECTION

Correction: Extrapolation of Inter Domain Communications and Substrate Binding Cavity of Camel HSP70 1A: A Molecular Modeling and Dynamics Simulation Study

The *PLOS ONE* Staff

[Fig 4](#) incorrectly appears as a duplicate of Fig 5. The publisher apologizes for the error. Please find the corrected version of [Fig 4](#) here.



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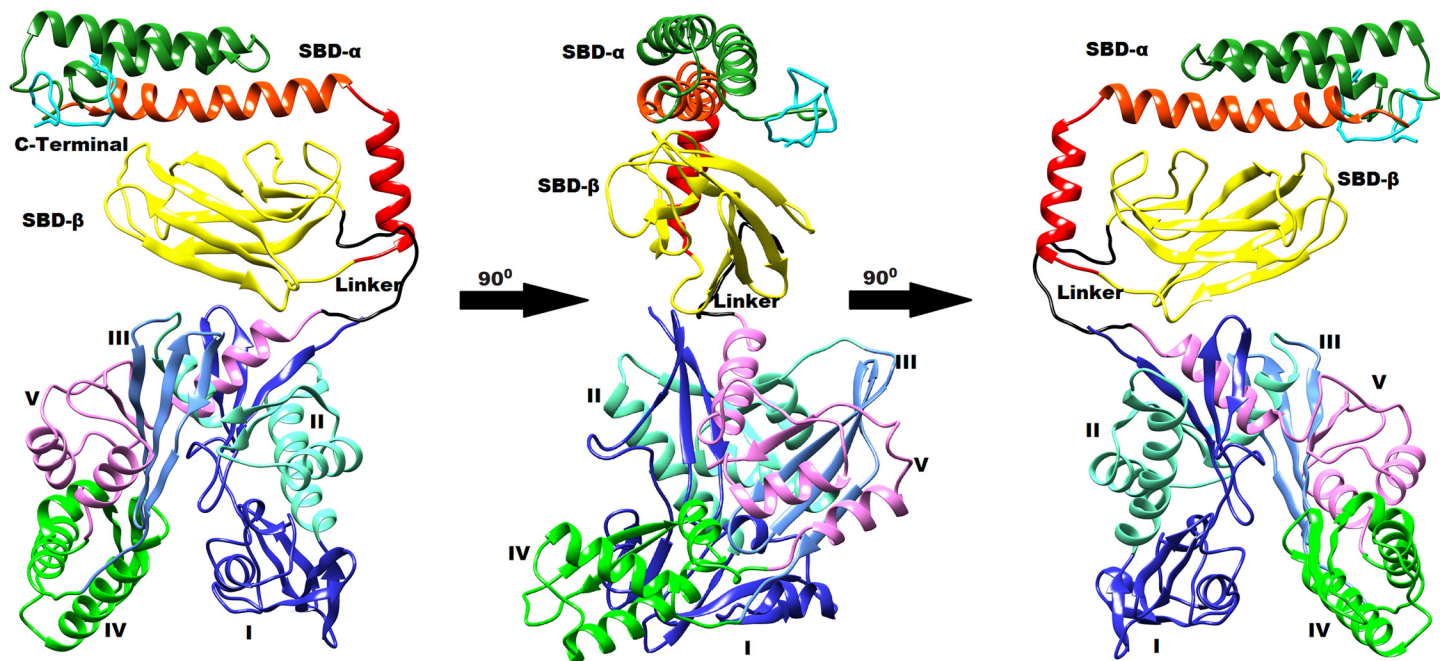


Fig 4. Typical average 3D structure represented in cartoon diagram of close state of cHSP70 rotated by 90° after relaxation through MD simulation.

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Reference

1. Gupta S, Rao AR, Varadwaj PK, De S, Mohapatra T (2015) Extrapolation of Inter Domain Communications and Substrate Binding Cavity of Camel HSP70 1A: A Molecular Modeling and Dynamics Simulation Study. PLoS ONE 10(8): e0136630. doi:[10.1371/journal.pone.0136630](https://doi.org/10.1371/journal.pone.0136630) PMID: [26313938](https://pubmed.ncbi.nlm.nih.gov/26313938/)