

Rosetta™

Rosetta™ is a molecular modeling software package for understanding protein structures, protein design, protein docking, protein-DNA and protein-protein interactions. The Rosetta software contains multiple functional modules, including RosettaAbinitio, RosettaDesign, RosettaDock, RosettaAntibody, RosettaFragments, RosettaNMR, RosettaDNA, RosettaRNA, RosettaLigand, RosettaSymmetry, and more. For a list of current features, see [current release notes](#).

PyRosetta can be found [here](#).

For more information: <http://www.rosettacommons.org/>

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Available at GitHub at: <https://github.com/RosettaCommons/rosetta/blob/main/LICENSE.md>

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