

The HADDOCK2.4 web server for integrative modeling of biomolecular complexes

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Figure S1 – Histogram of HADDOCK2.4 web server run time execution, based on statistics from January 2020 to June 2023. Many factors affect the run time, such as the size of the system and number of components being modelled, to the number of restraints defined, the overall load on our server and the distributed HTC resources used for the computations.

