

[Inici](#) > A systematic analysis of scoring functions in rigid-body protein docking: The delicate balance between the predictive rate improvement and the risk of overtraining

---

## **A systematic analysis of scoring functions in rigid-body protein docking: The delicate balance between the predictive rate improvement and the risk of overtraining**

**URL:** <http://onlinelibrary.wiley.com/doi/10.1002/prot.25289/abstract>

**UPCommons Handle URL** <http://upcommons.upc.edu/handle/2117/106838>

**Authors:** [Barradas-Bautista, Didier](#) / [Moal, Iain](#) / [Fernández-Recio, Juan](#)

**Publication:** Proteins: Structure, Function, and Bioinformatics

**Volume / Number / Pagination:** 85 / 7 / 1287 - 1297

Barcelona Supercomputing Center - Centro Nacional de Supercomputación

---

**Source URL (retrieved on 25 abr 2024 - 08:11):** <https://www.bsc.es/ca/research-and-development/publications/systematic-analysis-scoring-functions-rigid-body-protein-0>