

[Inicio](#) > [UEP: an open-source and fast classifier for predicting the impact of mutations in protein?protein complexes](#)[Abstract](#)[Motivation](#)[Results](#)[Availability and implementation](#)[Supplementary information](#)

[UEP: an open-source and fast classifier for predicting the impact of mutations in protein?protein complexes](#)[Abstract](#)[Motivation](#)[Results](#)[Availability and implementation](#)[Supplementary information](#)

URL: <https://academic.oup.com/bioinformatics/article/37/3/334/5881630>

Authors: [Amengual-Rigo, Pep](#) / [Fernández-Recio, Juan](#) / [Guallar, Victor](#) / [Elofsson, Arne](#)

Publication: Bioinformatics

Pagination:

Barcelona Supercomputing Center - Centro Nacional de Supercomputación

Source URL (retrieved on 14 Dic 2024 - 05:59): <https://www.bsc.es/es/research-and-development/publications/uep-open-source-and-fast-classifier-predicting-the-impact>