

[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 27 Mar 2023 - 00:31):** <https://www.bsc.es/es/research-and-development/publications/uep-open-source-and-fast-classifier-predicting-the-impact>