

<u>Inicio</u> > UEP: an open-source and fast classifier for predicting the impact of mutations in protein?protein complexesAbstractMotivationResultsAvailability and implementationSupplementary information

UEP: an open-source and fast classifier for predicting the impact of mutations in protein?protein complexesAbstractMotivationResultsAvailability and implementationSupplementary information

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

Authors: <u>Amengual-Rigo</u>, Pep / <u>Fernández-Recio</u>, Juan / <u>Guallar</u>, Victor / <u>Elofsson</u>, Arne

Publication: Bioinformatics

Pagination:

Barcelona Supercomputing Center - Centro Nacional de Supercomputación

Source URL (retrieved on 27 *Abr 2024 - 01:04*): <u>https://www.bsc.es/es/research-and-</u> development/publications/uep-open-source-and-fast-classifier-predicting-the-impact