

[Inici](#) > Crystal structure of a substrate-bound amino acid antiporter in the outward open conformation: mechanism of substrate recognition and transport

Crystal structure of a substrate-bound amino acid antiporter in the outward open conformation: mechanism of substrate recognition and transport

Authors: [Kowalczyk](#), / [Ratera, M](#) / [Paladino](#), / [Bartoccioni, Paola](#) / [Errasti](#), / [Valencia, E](#) / [Portella](#), / [Zorzano, Antonio](#) / [Fita, Ignacio](#) / [Orozco, Modesto](#) / [Carpena, Xavier](#) / [Vázquez-Ibar, JL](#) / [Palacín, M](#)

Barcelona Supercomputing Center - Centro Nacional de Supercomputación

Source URL (retrieved on 20 abr 2024 - 03:58): <https://www.bsc.es/ca/research-and-development/publications/crystal-structure-substrate-bound-amino-acid-antiporter-the>