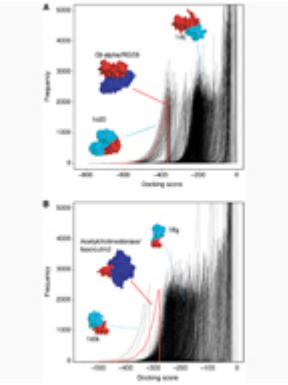
Computational programs used to predict the composition of protein complexes



This finding was possible after running more than 100,000 docking experiments using the MareNostrum supercomputer

4199.pdf
Barcelona Supercomputing Center - Centro Nacional de Supercomputación