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The Niefind group made effective tests for an upcoming crystallographic fragment screening campaign. Further, diffraction data to determine structures of the OCNDS-related mutant  $CK2\alpha^{Tyr50Ser}$  and of several enzyme/ligand complexes were collected successfully.

We collected several complete datasets of the membrane binding domain 3 of the ER membrane-located copper transporter RAN1 from *Arabidopsis thaliana* to a resolution of approx. 1.9 Å. We were able to solve the structure using molecular replacement. However model and data have not yet been deposited in the PDB. A manuscript describing our findings is in perparation.