

ID23-1 (22 April 2022)

ETR1 crystallized via LCP was screened for good diffracting crystals but none was of sufficient quality.

Several variants of CK2alpha and Ck2beta proteins were screened for high resolution data, 27 data sets collected.

From the PETases and the esterase CSS1 we have dead mutants which were either co-crystallized or soaked with their substrate with the aim to elucidate the catalytic mechanism. We collected 11 data sets.

All characterizations and data collections were native ones, so that only one energy/wavelengths was used.