

Volume 75 (2019)

Supporting information for article:

Structure of the dihydrolipoamide succinyltransferase catalytic domain from *Escherichia coli* in a novel crystal form: a tale of a common protein crystallization contaminant

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Figure S1 (a) Content of the asymmetric unit in the *I*4 crystal form which consists of six monomers. (b) A sample of the electron density map $(2F_o - F_c)$ contoured at 1 r.m.s.d.. The area shown is region 4 of the active site (Knapp *et al.*, 2000). It has been suggested that His375 is the catalytic residue. (c) A sample of the electron density map shown along the three-fold NCS axis of one trimer.