

Supplementary Material

Lipase-Catalyzed Synthesis, Antioxidant Activity, Antimicrobial Properties and Molecular Docking Studies of Butyl Dihydrocaffeate

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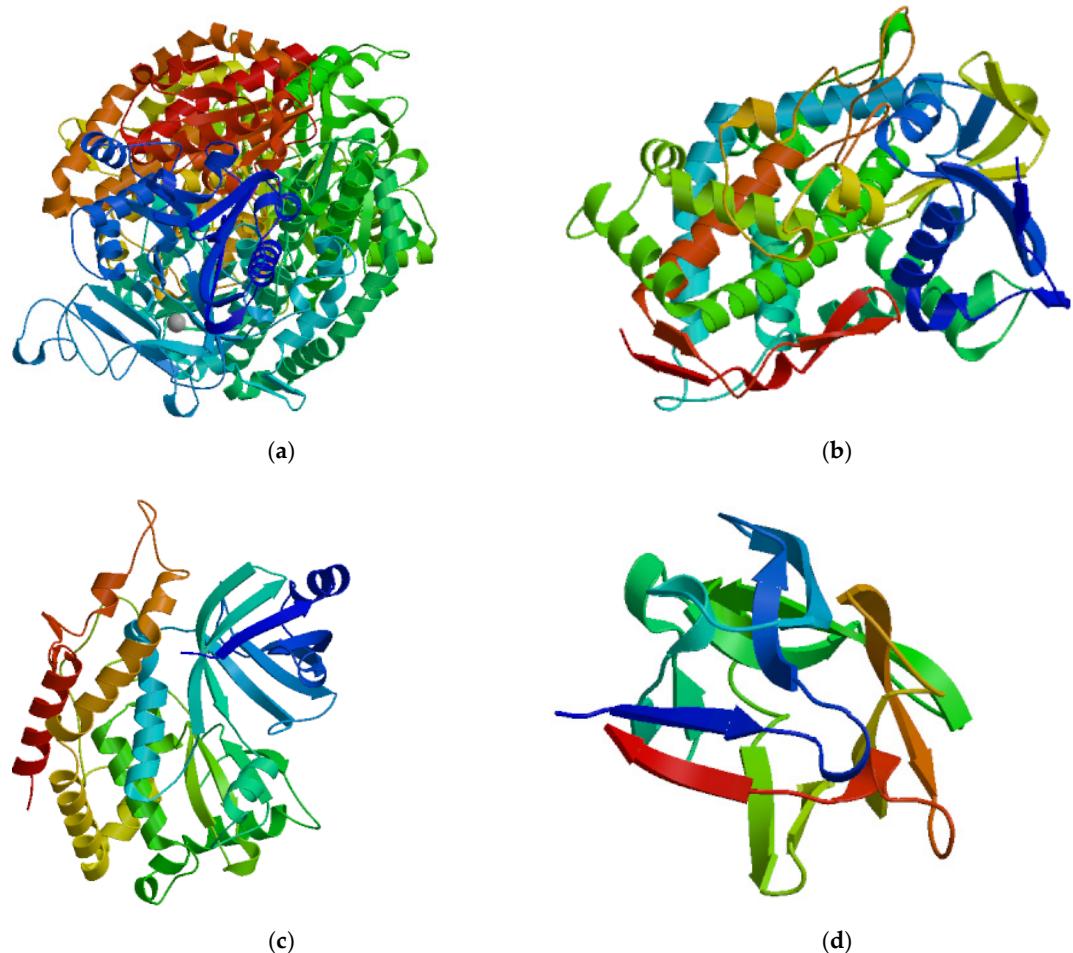


Figure S1. Structures of the modelled proteins (a) glutamine-fructose-6-phosphate transaminase, (b) 14- α sterol demethylase B, (c) invasin CotH3, (d) mucoricin.

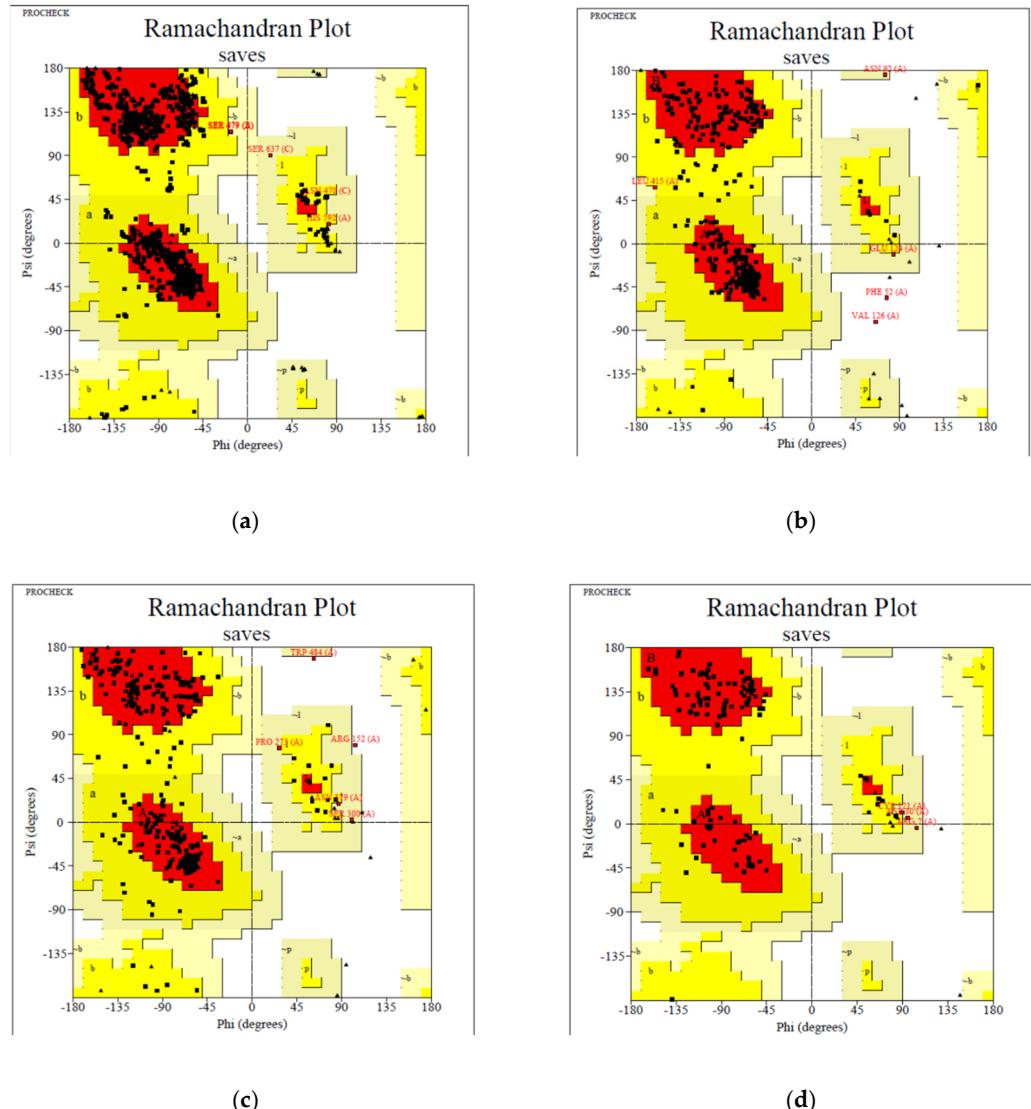


Figure S2. The Ramachandran's plots (PROCHECK) of the modelled proteins **(a)** glutamine-fructose-6-phosphate transaminase, **(b)** 14- α sterol demethylase B, **(c)** invasin CotH3, **(d)** mucorcinin.

Table S1. The estimated target-template alignment / predicted model quality indices.

	GFAT	14-α sterol demethylase B	Invasin CotH3	Mucoricin
SWISSPROT				
Template	6r4g.1.A	4uhl.1.A	5jd9.1.A	3pg0.1.A
Sequence Identity (%)	59.88	47.15	19.92	32.09
Coverage	0.98	0.86	0.91	0.97
Sequence Similarity	0.48	0.43	0.31	0.37
GMQE	0.83	0.67	0.59	0.68
QMEANDisCo Global	0.80 ± 0.05	0.70 ± 0.05	0.56 ± 0.05	0.68 ± 0.07
ERRAT				
Overall Quality Factor	95.02	92.31	86.13	87.12
PROCHECK				
Residues in Most Favoured Regions	91.70%	89.60%	84.30%	82.50%
Residues in Additional Allowed Regions	7.90%	9.20%	14.60%	15.10%
Residues in Generously Allowed Regions	0.40%	0.70%	0.60%	2.40%
Residues in Disallowed Regions	0.00%	0.50%	0.60%	0.00%

Abbreviations: GFAT - glutamine-fructose-6-phosphate transaminase, GMQE - Global Model Quality Estimate