Solution structure of the lipoyl domain of the chimeric
fromNeisseria meningitidis

FEBS Journal

268, 4908-4917

DOI: 10.1046/j.0014-2956.2001.02422.x

Citation Report

#	Article	IF	Citations
1	Proteome analysis ofNeisseria meningitidis serogroup A. Proteomics, 2004, 4, 2893-2926.	1.3	57
2	Function, Attachment and Synthesis of Lipoic Acid in Escherichia coli. Advances in Microbial Physiology, 2005, 50, 103-146.	1.0	118
3	The enlargement of the hormone immune deprivation concept to the blocking of TGFÎ±-autocrine loop: EGFR signaling inhibition. Cancer Immunology, Immunotherapy, 2006, 55, 628-638.	2.0	5
4	Protein Biotinylation Visualized by a Complex Structure of Biotin Protein Ligase with a Substrate. Journal of Biological Chemistry, 2008, 283, 14739-14750.	1.6	46
5	The role of loop and Î²-turn residues as structural and functional determinants for the lipoyl domain from the Escherichia coli 2-oxoglutarate dehydrogenase complex. Biochemical Journal, 2008, 409, 357-366.	1.7	15
6	Biotin and Lipoic Acid: Synthesis, Attachment, and Regulation. EcoSal Plus, 2008, 3, .	2.1	20
7	Nuclear Magnetic Resonance Approaches in the Study of 2-Oxo Acid Dehydrogenase Multienzyme Complexesâ€"A Literature Review. Molecules, 2013, 18, 11873-11903.	1.7	9
8	Biotin and Lipoic Acid: Synthesis, Attachment, and Regulation. EcoSal Plus, 2014, 6, .	2.1	54
9	Structure of the native pyruvate dehydrogenase complex reveals the mechanism of substrate insertion. Nature Communications, 2021, 12, 5277.	5.8	39