

Maximum-likelihood density modification

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Crystal Structure of ATP Sulfurylase from the Bacterial Symbiont of the Hydrothermal Vent Tubeworm <i>Riftia pachyptila</i> . <i>Biochemistry</i> , 2001, 40, 14509-14517.	1.2	42
2	Crystal structure of the 30 S ribosomal subunit from <i>Thermus thermophilus</i> : purification, crystallization and structure determination. <i>Journal of Molecular Biology</i> , 2001, 310, 827-843.	2.0	128
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9	Crystal structure of the bacterial cell division inhibitor MinC. <i>EMBO Journal</i> , 2001, 20, 2454-2461.	3.5	83
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74	Automated side-chain model building and sequence assignment by template matching. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2003, 59, 45-49.	2.5	94
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