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Shared requirements for key residues in the antibiotic resistance enzymes ErmC and ErmE suggest a common mode of RNA recognition

DOI: 10.1074/jbc.RA120.014280 Journal of Biological Chemistry, 2020, 295, 17476-17485.

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5	Decoding the Mechanism of Specific RNA Targeting by Ribosomal Methytransferases.		
4	Three critical regions of the erythromycin resistance methyltransferase, ErmE, are required for function supporting a model for the interaction of Erm family enzymes with substrate rRNA. <i>Rna</i> , 2021 ,	5.8	О
3	Crystal structure and functional analysis of mycobacterial erythromycin resistance methyltransferase Erm38 reveals its RNA binding site <i>Journal of Biological Chemistry</i> , 2022 , 101571	5.4	О
2	Decoding the Mechanism of Specific RNA Targeting by Ribosomal Methyltransferases <i>ACS Chemical Biology</i> , 2022 ,	4.9	О
1	Macrolide, lincosamide, glycopeptide, and other antibacterial antibiotics. 2023 , 157-213		O