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Hydrogen bonds are related to the thermal stability of 16S rRNA					JBiSE Subscription	
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Author(s) Hiroshi Nakashima, Ai Fukuoka, Yuka Saitou ABSTRACT The number of base pairs in the 16S rRNA secondary structures of 51 bacterial sequences was					About JBiSE News	
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highly	bunted, and the number of hydrogen bonds was estimated. The number of hydrogen bonds was ghly correlated with the optimal growth temperature (OGT) rather than with the $G+C$ content. Sired and unpaired nucleotides in mesophiles were compared to those in thermophiles. OGT exhibited relationship with paired nucleotides but not with unpaired nucleotides. The total number of paired as lell as unpaired nucleotides in mesophiles was very similar to that in thermophiles. However, the emponents in base pairs in mesophiles significantly differed from those in thermophiles. As compared ith mesophiles, the number of $G-C$ base pairs in thermophiles was high whereas that of $A-C$ base				Recommend to Peers	
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pairs was low. In this study, we showed that hydrogen bonds are important for stabilizing 16S rRNAs at high temperatures.				Downloads:	452,882	
KEYWORDS Optimal Growth Temperature; 16S Ribosomal RNA; G + C Content; Hydrogen Bonds; Base Pairs; Nucleotide				Visits:	1,291,950	
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