

## ERRATA

# Erratum: “Characterization of a New RNase HII and Its Essential Amino Acid Residues in the Archaeon *Sulfolobus tokodaii* Reveals a Regulatory C-Terminus” [*Biochemistry (Moscow)*, 75, 930 (2010)]

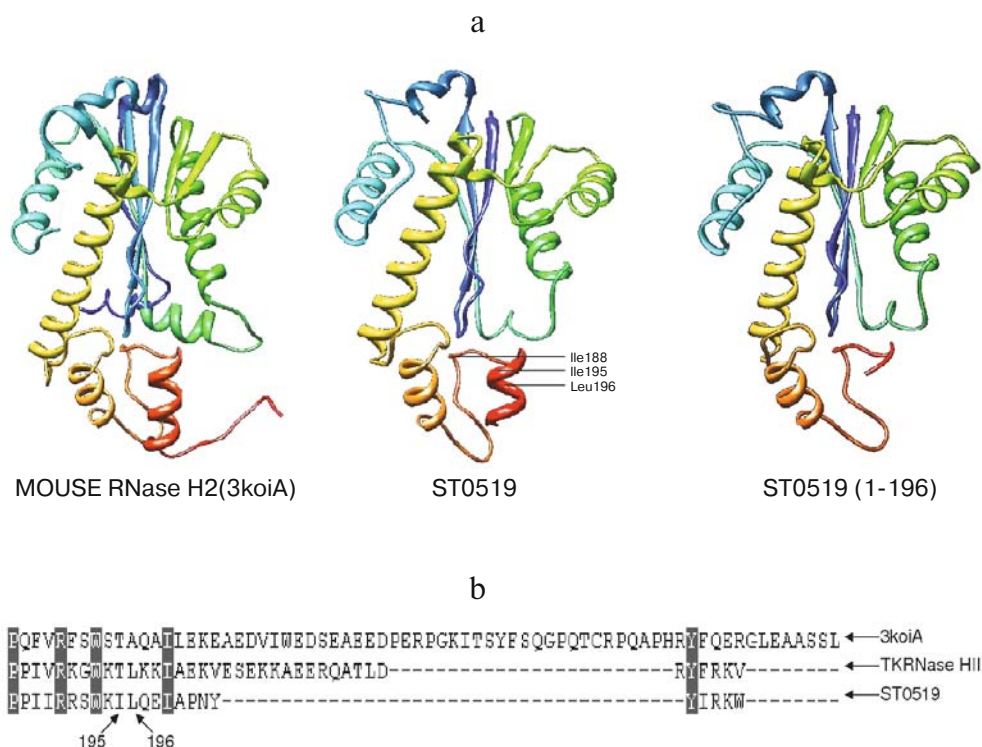
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Received March 30, 2010  
Revision received May 14, 2010

DOI: 10.1134/S0006297910100159

1. The figure below is color insert to the article of Ke Zhan and Zheng-Guo He "Characterization of a new RNase HII and its essential amino acid residues in the archaeon *Sulfolobus tokodaii* reveals a regulatory C-terminus". This color insert was mistakenly attached to a different article. We apologize for this mistake.



**Fig. 5.** (Ke Zhan and Zheng-Guo He) Schematic diagram of the *S. tokodaii* ST0519 structure. a) The structure obtained using the homology modeling method. The structures for mouse RNase H2 and ST0519 or St0519(1-196) are indicated. b) The C-terminal sequence alignments of ST0519, TkrNase HII, and mouse RNase H2. TkrNase HII represents ST0753 from *Thermococcus kodakaraensis*. The conserved L196 is indicated.