OPEN ACCESS Freely available online

PLOS ONE

Correction

Correction: Archaeal Tuc1/Ncs6 Homolog Required for Wobble Uridine tRNA Thiolation Is Associated with Ubiquitin-Proteasome, Translation, and RNA Processing System Homologs

The *PLOS ONE* Staff

File S1 is corrupted and will not open. Please view the correct File S1 here.

Supporting Information

File S1. Supporting Information. File S1 includes: **Table S1**. Strains, plasmids used in this study; **Table S2**. Oligonucleotide primers used in this study; **Figure S1**. Dendrogram analysis of *Haloferax volcanii* NcsA and homologs of the α hydrolase (ANH) superfamily from archaea, eukaryotes and bacteria; **Figure S2**. 3D-structural model of *Haloferax volcanii* NcsA; **Figure S3**. Organization of *ncsA* and its targeted deletion on the genome of *Haloferax volcanii*; **Figure S4**. Growth of *Haloferax volcanii ΔncsA* mutant compared to parent strain H26 at optimum growth temperature; **Figure S5**. Lys204 residue of NcsA is found isopeptide linked to SAMP2; **Figure S6**. Detection of E1-like UbaA and PAN-A/1 ATPase in NcsA-StrepII pull-down fractions.
(PDF)

Reference

1. Chavarria NE, Hwang S, Cao S, Fu X, Holman M, et al. (2014) Archaeal Tuc1/Ncs6 Homolog Required for Wobble Uridine tRNA Thiolation Is Associated with Ubiquitin-Proteasome, Translation, and RNA Processing System Homologs. PLoS ONE 9(6): e99104. doi:10.1371/journal.pone.0099104

Citation: The *PLOS ONE* Staff (2014) Correction: Archaeal Tuc1/Ncs6 Homolog Required for Wobble Uridine tRNA Thiolation Is Associated with Ubiquitin-Proteasome, Translation, and RNA Processing System Homologs. PLoS ONE 9(12): e115719. doi:10.1371/journal.pone.0115719

Published December 16, 2014