

ERRATUM

In the article, "Developing Limited Proteolysis and Mass Spectrometry for the Characterization of Ribosome Topography" by Moo-Jin Suh, Soheil Pourshahian, and Patrick A. Limbach which was published in the July issue, Vol. 18, No. 7, pages 1304-1317, Figure 6 failed to print in color. The publisher regrets the error. The corrected figure is shown below, with the corresponding figure legend.

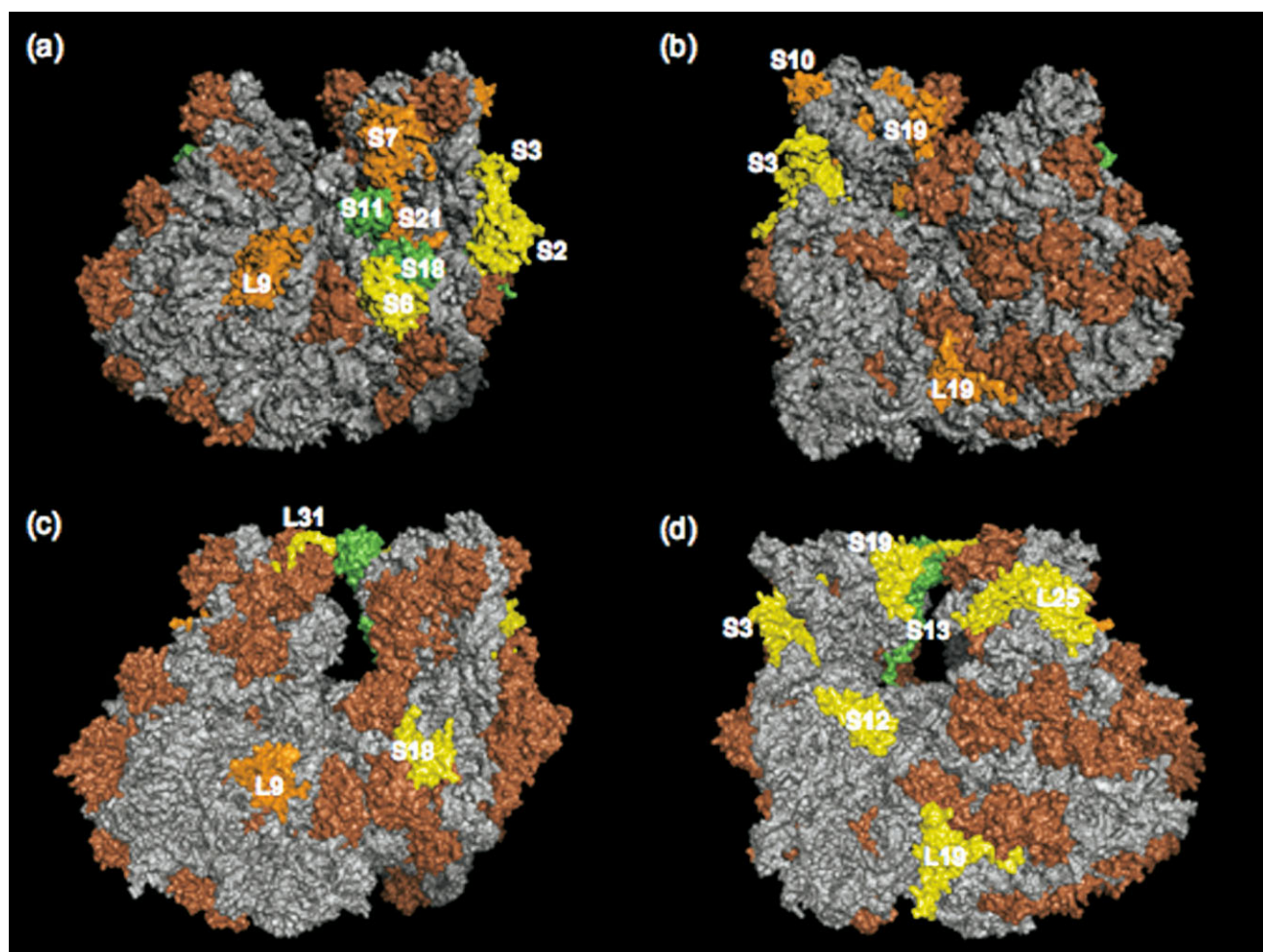


Figure 6. Surface presentation of the ribosomal proteins digested with both trypsin and Proteinase K. Color scheme: rRNA is represented in gray. Category one proteins (susceptible to digestion within 30 min) are represented in yellow. Category two proteins (susceptible to digestion within 125 min) are represented in green. Category three proteins (stable to digestion up to 1000 min) are represented in orange (see text). Proteins that were still detected intact even after 1500 min of incubation are represented in brown. (a) and (b) Views of the surfaces of the *E. coli* 70S ribosome with protease susceptible proteins identified. (a) to (b) are 180° rotations about the vertical axis. (a) View from the left-hand side with the 30S subunit on the right and the 50S subunit on the left. (b) View from the right-hand side with the 30S subunit on the left and the 50S subunit on the right. (c) and (d) Views of the surfaces of the *T. thermophilus* 70S ribosome with protease susceptible proteins identified. (c) to (d) are 180° rotations about the vertical axis with similar perspectives as for *E. coli*. Images of the 30S and 50S subunit structures were produced using MacPyMol [36] from the 3.5 Å crystal structure of *E. coli* (accession 2AVY and 2AW4) [17] and the 2.8 Å crystal structure of *T. thermophilus* (accession 2J00 and 2J01) [18].