
A unique serine-specific elongation factor Tu found in nematode mitochondria

Takashi Ohtsuki, Aya Sato, Yoh-ichi Watanabe & Kimitsuna Watanabe
Nat. Struct. Biol. 9, 669–673 (2002).

Figure 3a in this paper contained a mistake. In the tRNA sequence depicted in this panel, the 47th and 48th positions are C and U, respectively. We apologize for any inconvenience this may have caused.

ADDENDA

The crystal structure of the proprotein processing proteinase furin explains its stringent specificity

Stefan Henrich, Angus Cameron, Gleb P Bourenkov, Reiner Kiefersauer, Robert Huber, Iris Lindberg, Wolfram Bode & Manuel E Than
Nat. Struct. Biol. 10, 520–526 (2003).

During the proofing stage of this paper, a publication presenting the crystal structure of Kex2, the homolog of furin in yeast, in complex with an inhibitor (Holyoak *et al. Biochemistry* 42, 6709–6718; 2003) appeared online. We regret not having mentioned this work as a 'Note added in proof'. The full reference for Holyoak *et al.* is included here.

Holyoak, T. *et al.* 2.4 Å resolution crystal structure of the prototypical hormone-processing protease Kex2 in complex with an Ala-Lys-Arg boronic acid inhibitor. *Biochemistry* 42, 6709–6718 (2003).