## **ERRATUM**

## Post-transcriptional nucleotide modification and alternative folding of RNA

## Mark Helm

Nucleic Acids Res., (2006) 34, 721-733

The author wishes it to be known that there was a print error in Figure 4 of the above mentioned article. A corrected figure is shown below with its legend.



**Figure 4.** Rearrangement on the secondary structure level induced by double methylation of human cytosolic tRNA<sup>Asn</sup>. The calculated structure of the unmodified transcript on the left side features an aberrant D-stem without G10, but including G26. Double methylation on *N6* of G26 impedes its Watson–Crick pairing with Cytidines and thus renders the base pair G26-C11, which is highlighted by a box, impossible. The fully modified tRNA<sup>Asn</sup> may thus adopt the classical cloverleaf structure as shown on the right. U? denoted an unknown modified uridine, likely a derivative of ribothymidine, at position 54.

© 2007 The Author(s)

This is an Open Access article distributed under the terms of the Creative Commons Attribution Non-Commercial License (http://creativecommons.org/licenses/ by-nc/2.0/uk/) which permits unrestricted non-commercial use, distribution, and reproduction in any medium, provided the original work is properly cited.