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**Codon usage bias from tRNA's point of view: Redundancy, specialization, and efficient decoding for translation optimization**

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In the first paragraph of the first column on page 2281 and in Figure 1, there is a typo in the definition of  $\text{ENC}_{\text{dif}}$ . The formula should read:

$$\text{ENC}_{\text{dif}} = -(\text{ENC}'_{\text{RP}} - \text{ENC}'_{\text{All}})/\text{ENC}'_{\text{All}}$$

Thus, positive values of  $\text{ENC}_{\text{dif}}$  indicate codon usage bias in ribosomal proteins as mentioned throughout the text.

The authors apologize for any confusion this may have caused.