

**Genome Research 14:** 2279–2286 (2004)

**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  $ENC_{dif}$ . The formula should read:

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

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

The authors apologize for any confusion this may have caused.