Corrigenda

Genome Research 28: 285–294 (2018)**Corrigendum: 3' UTR lengthening as a novel mechanism in regulating cellular senescence**

Meng Chen, Guoliang Lyu, Miao Han, Hongbo Nie, Ting Shen, Wei Chen, Yichi Niu, Yifan Song, Xueping Li, Huan Li, Xinyu Chen, Ziyue Wang, Zheng Xia, Wei Li, Xiao-Li Tian, Chen Ding, Jun Gu, Yufang Zheng, Xinhua Liu, Jinfeng Hu, Gang Wei, Wei Tao, and Ting Ni

The authors would like to correct Figure 3, panel J, in which the rightmost upper image of SA- β -gal stained 293T cells following short hairpin RNA (shRNA)-mediated knockdown of *RRAS2* with sh769 (*RRAS2*-KD-sh769) was inadvertently, and due to a labeling error, taken from the same original source image presented in the middle upper panel, which shows increased SA- β -gal activity following *RRAS2* knockdown by a different shRNA (sh646). This correction does not affect any of the conclusions of the article. The corrected image representative of *RRAS2*-KD-sh769 is provided below, and Figure 3 has been updated in the article online.

Figure 3. Panel J, rightmost upper image.

The authors thank Ning Yuan Lee for bringing this error to their attention and apologize for any confusion this may have caused.

Additionally, the authors have provided a revised Supplemental Figure S7 file in which the redundant successive Supplemental figure files have been removed. This can be found in the Revised Supplemental Material online.

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Genome Research 30: 1060–1072 (2020)**Corrigendum: Functional annotation of human long noncoding RNAs via molecular phenotyping**

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The authors would like to correct the misspelling of an author's name and the inadvertent omission of two affiliations for that author, which are as follows: Christopher J.F. Cameron, Department of Biochemistry, Rosalind and Morris Goodman Cancer Research Center, McGill University, Montréal, Québec H3G 1Y6, Canada and Department of Molecular Biophysics and Biochemistry, Yale University, New Haven, Connecticut 06510, USA.

These updates are reflected in the revised manuscript online.

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Corrigendum: Functional annotation of human long noncoding RNAs via molecular phenotyping

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Jordan A. Ramilowski, Chi Wai Yip, Saumya Agrawal, et al.

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