

CORRECTION

Correction: TCGA based integrated genomic analyses of ceRNA network and novel subtypes revealing potential biomarkers for the prognosis and target therapy of tongue squamous cell carcinoma

The PLOS ONE Staff

There are errors in the Funding statement. The publisher apologizes for the errors. The correct Funding statement is as follows: This study was supported by the Fundamental Research Funds for the Central Universities of Central South University (1053320182836). The funder had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

## Reference

 Li Z, Jiang C, Yuan Y (2019) TCGA based integrated genomic analyses of ceRNA network and novel subtypes revealing potential biomarkers for the prognosis and target therapy of tongue squamous cell carcinoma. PLoS ONE 14(5): e0216834. https://doi.org/10.1371/journal.pone.0216834 PMID: 31141819



GOPEN ACCESS

Citation: The *PLOS ONE* Staff (2019) Correction: TCGA based integrated genomic analyses of ceRNA network and novel subtypes revealing potential biomarkers for the prognosis and target therapy of tongue squamous cell carcinoma. PLoS ONE 14(6): e0218987. https://doi.org/10.1371/journal.pone.0218987

Published: June 20, 2019

Copyright: © 2019 The PLOS ONE Staff. This is an open access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.