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Author Correction: Cancer LncRNA Census reveals evidence for deep functional conservation of long noncoding RNAs in tumorigenesis

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Correction to: *Communications Biology* <https://doi.org/10.1038/s42003-019-0741-7>, published online 05 February 2020.

In the published version of this paper, the members of the Pan-Cancer Analysis of Whole Genomes (PCAWG) Consortium were listed in the Supplementary Information; however, these members should have been included in the main paper. The original Article has been corrected to include the members and affiliations of the PCAWG Consortium in the main paper; the corrections have been made to the HTML version of the Article but not the PDF version. In the PCAWG Drivers and Functional Interpretation Group, the affiliation for Erik Larsson has also been changed from ‘Computational Biology Center, Memorial Sloan Kettering Cancer Center, New York, NY 10065, USA’ to ‘Institute of Biomedicine, Sahlgrenska Academy at University of Gothenburg, Gothenburg, Sweden’. Additional corrections to affiliations have been made to the PDF and HTML versions of the original Article for consistency of information between the PCAWG list and the main paper.

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