

Author Correction: Analyses of non-coding somatic drivers in 2,658 cancer whole genomes


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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. Additional minor 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. An additional affiliation has been added for Husen M. Umer (Department of Oncology-Pathology, Science for Life Laboratory, Karolinska Institute, Stockholm, Sweden). The affiliation for Erik Larsson has been changed from Computational Biology Center, Memorial Sloan Kettering Cancer Center, New York, NY, USA to Institute of Biomedicine, Sahlgrenska Academy at University of Gothenburg, Gothenburg, Sweden.

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