




Author Correction: Proteogenomic analysis reveals RNA as a source for tumor-agnostic neoantigen identification

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 Check for updates

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With respect to the original “Data availability” statement, since the publication of this work, to ensure patient data privacy protection per request of the authors’ institute *Technical University of Munich*, the RNA-seq dataset from sorted CD8+ TILs originally deposited to the European Nucleotide Archive (ENA) with study accession number [PRJEB61429](https://www.ebi.ac.uk/ena/browser/view/PRJEB61429) has been transferred to the European Genome-phenome Archive (EGA) under restricted access with the study identifier [EGAS00001006706](https://www.ebi.ac.uk/ega/browser/view/EGAS00001006706) (along with the other sequencing datasets generated in the study) and the dataset identifier [EGAD50000000193](https://www.ebi.ac.uk/ega/browser/view/EGAD50000000193). The “Data availability” section of the Article file (PDF and HTML versions) and the Data section of the Reporting Summary have been updated to reflect these changes.

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