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Author Correction: The Molecular Tumor Board Portal supports clinical decisions and automated reporting for precision oncology

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In this article, the Code availability section has been updated to clarify conditions for access and to revise the repository link. The new statement now reads “Aggregated results shown in this article were created using Python (>3.5) NumPy, Pandas and Matplotlib open-source libraries. The code for the MTBP components reproducing the variant annotation framework discussed in the manuscript has been deposited to Zenodo with access restricted for research use only (see <https://doi.org/10.5281/zenodo.6376428> for terms of use details). Access to other components available in MTBP production systems can be provided upon agreement with the copyright holders.” The change has been made in the HTML and PDF versions of the article.

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