




Publisher Correction: SAIGE-GENE+ improves the efficiency and accuracy of set-based rare variant association tests

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 Check for updates

Wei Zhou , Wenjian Bi , Zhangchen Zhao , Kushal K. Dey ,
Karthik A. Jagadeesh, Konrad J. Karczewski , Mark J. Daly,
Benjamin M. Neale  and Seunggeun Lee 

In the version of this article initially published, a footnote designating Wei Zhou, Wenjian Bi and Zhangchen Zhao as equally contributing authors was missing and has now been amended to the HTML and PDF versions of the article.

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