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Scanorama [↗](#)

Brian Hie¹, Bryan Bryson¹, Bonnie Berger¹

¹Massachusetts Institute of Technology

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Works for me

[dx.doi.org/10.17504/protocols.io.9gch3sw](https://doi.org/10.17504/protocols.io.9gch3sw)



Brian Hie ⚡

ABSTRACT

Scanorama enables efficient integration of heterogeneous scRNA-seq data sets.

EXTERNAL LINK

<http://scanorama.csail.mit.edu>

THIS PROTOCOL ACCOMPANIES THE FOLLOWING PUBLICATION

Brian Hie, Bryan Bryson & Bonnie Berger. Efficient integration of heterogeneous single-cell transcriptomes using Scanorama. Nature Biotechnology 37:685–691 (2019).



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