Scanorama 🔗

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`1` *Works for me*   dx.doi.org/10.17504/protocols.io.9gch3sw

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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).