

Resource Summary Report

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Scanorama

DOI:10.17504/protocols.io.9gch3sw

Type: Protocol

Proper Citation

Brian Hie, Bryan Bryson, Bonnie Berger 2019. Scanorama. protocols.io
[dx.doi.org/10.17504/protocols.io.9gch3sw](https://doi.org/10.17504/protocols.io.9gch3sw)

Protocol Information

URL: <https://dx.doi.org/10.17504/protocols.io.9gch3sw>

Authors: Brian Hie, Bryan Bryson, Bonnie Berger

Group: Human Cell Atlas Method Development Community

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

Affiliations: Massachusetts Institute of Technology, Massachusetts Institute of Technology, Massachusetts Institute of Technology

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

Version: 1

Publication Date: 2019

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Record Creation Time: 20210329T220533+0000

Record Last Update: 20210329T221009+0000

Data and Source Information

Source: