

Resource Summary Report

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sci-RNA-seq2 Pipeline for the Human Biomolecular Atlas Program (HuBMAP)

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Protocol Information

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Authors: Shin Lin, Cole Trapnell, Jay Shendure

Summary: This document provides an overview of the protocols used by the Caltech-UW TMC to perform sci-RNA-seq2. This involves sequencing of the transcriptomes of dissociated single cells in a massively parallel manner.

Affiliations: University of Washington, University of Washington, University of Washington

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