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

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ABSTRACT

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.

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- 1 Collection and processing of human tissues.
Sample procurement: dx.doi.org/10.17504/protocols.io.bi9pkdvn
Flash freezing: dx.doi.org/10.17504/protocols.io.bi9qkdvw
- 2 sci-RNA-seq2 assay
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