10Xv2 RNASeq Sample Processing

ZengU19 BICCN Grant[1]

[1]Allen Institute

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ABSTRACT

Allows for rapid generation of 3' transcriptomic-NGS-ready- single-cell-libraries from pools of cells.

DOCX

10Xv2 RNASeq Sample
Processing.docx

PROTOCOL STATUS

Working

We use this protocol in our group and it is working