



May 16, 2024 Version 2

10x Multiome Sample Processing V.2

DOI

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Protocol status: Working

We use this protocol and it's working

Created: May 26, 2022

Last Modified: May 16, 2024

Protocol Integer ID: 100005

Abstract

Protocol is used to profile all the open chromatin regions at a single nuclei level through the rapid generation of NGS-ready libraries, and for rapid generation of 3' transcriptomic-NGS-ready-single-cell-libraries from the same pool of transposed nuclei.

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Attachments



CG000338 ChromiumNe

X...

5MB



MB0204 10x Multiome .

..

1.7MB

