

May 17, 2024

Version 2

10x Multiome Sample Processing V.2

DOI

<https://dx.doi.org/10.17504/protocols.io.bp2l61mqrvqe/v2>

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Protocol status: Working

We use this protocol and it's working

Created: May 17, 2024

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Protocol Integer ID: 100005

Keywords: open chromatin region, 10x multiome sample processing protocol, single nuclei level, transposed nuclei, nuclei, national institutes of health

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_ChromiumN

ex...

5MB



MB0204_10x_Multiome

...

1.7MB

