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Version 3

10x Multiome Sample Processing V.3

DOI

<https://dx.doi.org/10.17504/protocols.io.bp2l61mqrvqe/v3>

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Protocol status: Working

We use this protocol and it's working

Created: March 07, 2025

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Protocol Integer ID: 124016

Keywords: open chromatin region, chromatin, 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

