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10x ATAC Genomics Sample Processing

Allen Institute for Brain Science¹¹Allen Institute

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dx.doi.org/10.17504/protocols.io.14egn7ndmv5d/v1**Allen Institute for Brain Science**

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Protocol is used to profile all the open chromatin regions at a single nuclei level through the rapid generation of NGS-ready libraries from a pool of transposed nuclei.

[MB0203_10x_ATAC_Genomics_Sample_Processing.docx](#)

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protocols.io<https://dx.doi.org/10.17504/protocols.io.14egn7ndmv5d/v1> protocol ,

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