



### Pi-ATAC 👄

Xingqi Chen<sup>1</sup>, Ulrike Litzenburger<sup>1</sup>, Yuning Wei<sup>1</sup>, Alicia N. Schep<sup>2</sup>, Edward L. LaGory<sup>3</sup>, Hani Choudhry<sup>4</sup>, Amato J. Giaccia<sup>3</sup>, William J. Greenleaf<sup>2</sup>, Howard Y. Chang<sup>1</sup>

<sup>1</sup>Center for Personal Dynamic Regulomes, Stanford University, Stanford, CA 94305, <sup>2</sup>Center for Personal Dynamic Regulomes, Stanford University, Stanford, CA 94305; Dept of Genetics, Stanford University, Stanford, CA 94305; Department of Applied Physics, Stanford University, Stanford, CA 94305, 3Division of Radiation and Cancer Biology, Department of Radiation Oncology, Stanford University, Stanford, CA 94305, <sup>4</sup>Department of Biochemistry, Cancer Metabolism and Epigenetic Unit, Faculty of Science, Cancer and Mutagenesis Unit, King Fahd Center for Medical Research, King Abdulaziz University, Jeddah, Saudi Arabia.

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Human Cell Atlas Method Development Community HYChangLab



Ulrike Litzenburger



#### **ABSTRACT**

Here we introduce Protein-indexed Assay of Transposase Accessible Chromatin with sequencing (Pi-ATAC) that combines singlecell chromatin and proteomic profiling. In conjunction with DNA transposition, the levels of multiple cell surface or intracellular protein epitopes are recorded by index flow cytometry and positions in arrayed microwells, and then subject to molecular barcoding for subsequent pooled analysis. PiATAC simultaneously identifies the epigenomic and proteomic heterogeneity in individual cells. Pi-ATAC reveals a casual link between transcription factor abundance and DNA motif access, and deconvolute cell types and states in the tumor microenvironment in vivo. We identify a dominant role for hypoxia, marked by HIF1A protein, in the tumor microvenvironment for shaping the regulome in a subset of epithelial tumor cells.

EXT ERNAL LINK

https://www.biorxiv.org/content/early/2018/04/27/310359

THIS PROTOCOL ACCOMPANIES THE FOLLOWING PUBLICATION

Joint single-cell DNA accessibility and protein epitope profiling reveals environmental regulation of epigenomic heterogeneity

Xingqi Chen, Ulrike Litzenburger, Yuning Wei, Alicia N. Schep, Edward L. LaGory, Hani Choudhry, Amato J. Giaccia, William J. Greenleaf, Howard Chang

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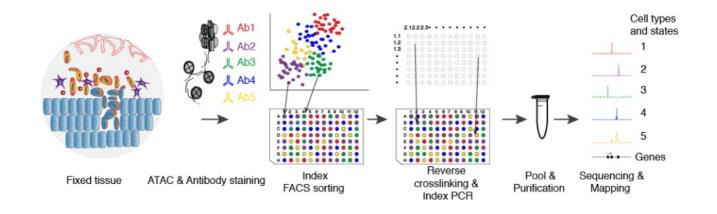
**PROTOCOL STATUS** 

#### Working

We use this protocol in our group and it is working

**GUIDELINES** 

Overview



|       | NAME ~                                             | CATALOG #   | VENDOR ~                 |  |  |  |  |
|-------|----------------------------------------------------|-------------|--------------------------|--|--|--|--|
|       | Nextera DNA library preparation kit 24 samples     | FC-121-1030 | Contributed by users     |  |  |  |  |
|       |                                                    |             |                          |  |  |  |  |
| STEPS | STEPS MATERIALS                                    |             |                          |  |  |  |  |
|       | NAME Y                                             | CATALOG #   | VENDOR ~                 |  |  |  |  |
|       | HBSS - 1% FBS (no collagenase) solution            |             | Contributed by users     |  |  |  |  |
|       | PBS                                                |             | Contributed by users     |  |  |  |  |
|       | 2%PFA                                              |             | Contributed by users     |  |  |  |  |
|       | Glycine, sodium salt                               | G-450       | Gold Biotechnology       |  |  |  |  |
|       | Nextera DNA library preparation kit 24 samples     | FC-121-1030 | Contributed by users     |  |  |  |  |
|       | 0.5M EDT A                                         | AM92606     | Contributed by users     |  |  |  |  |
|       | PBS                                                |             | Contributed by users     |  |  |  |  |
|       | Proteinase K                                       | E00491      | Thermo Fisher Scientific |  |  |  |  |
|       | NEBNext High-Fidelity 2X PCR Master Mix - 250 rxns | M0541L      | New England Biolabs      |  |  |  |  |
|       | MinElute Reaction Cleanup Kit                      | 28204       | Qiagen                   |  |  |  |  |
|       |                                                    |             |                          |  |  |  |  |

## SAFETY WARNINGS

For safety warnings and hazard information, please refer to the SDS (Safety Data Sheet).

### BEFORE STARTING

# Prepare before:

2% PFA

1.25 M Glycin

single cell suspension solution: 200 U/ml Collagenase in HBSS-FBS (1%))

Nextera Kit: Illumina #15028212 (containing Tn5, 2X TD buffer)

Permeabilization buffer: 10 mM Tris ph7.5

10 mM NaCl 3 mM MgCl2

add fresh 0.1% NP40 (NOTE: NP40 concentration/other detergents can vary from cell type to cell type)

reverse x-link Buffer: 50 mM Tris-HCl pH 8.0

0.5% Tween20 0.5% Igepal CA-630 5 ng/ml proteinase K

0.5 M EDT A

2x NEBNext High Fidelity PCR MasterMix

Nextera adapters for scATAC (Buenrostro et al, Nature volume 523, pages 486-490 (23 July 2015))

1 **▶**NOTE

Please see the "before start" section of the  $\underline{\text{Guidelines}}$  for buffers and solutions to prepare prior to start.

## Preparation

2 If using tissue: take out tissue and put on 10 cm dish filled with 10 ml PBS on ice. Weigh

## Then make a single cell suspension

3 Remove PBS from dish, but keep on ice.

## Single cell suspension

- 4 Cut tissue in small pieces using a scalpel or razor blade on ice.
- $5 \qquad \text{Add single cell dissociation solution to dish (for most tissues: 200 U/ml Collagenase in HBSS-FBS (1\%))}.$
- 6 Try to pipet with 10 ml pipet, fill into 15ml tube.
- 7 Incubate 15 ml tube in 37°C incubator rotating (depending on amount and type of tissue).

§ 37 °C Incubation

**© 00:15:00** Incubation

8 Spin down for 5 min at 1000 rpm.

**७00:05:00** Spin down



- 17 Spin down.
- Permeabilize cells using buffer containing 0.1% NP40, no incubation, immediately spin 5 min 1200 rpm.

```
©00:05:00 Spin
```

19 If staining is desired: stain with antibodies here.

### ATAC

- 20 Spin down.
- 21 ATAC: (for 50,000 cells) 25  $\mu$ l 2xTD, 22,5  $\mu$ l H<sub>2</sub>O, 2.5  $\mu$ l Tn5
  - → 30 min at 37 °C



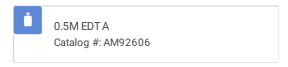




8 37 °C

**©00:30:00** 

- Nextera DNA library preparation kit 24 samples
  Catalog #: FC-121-1030
- Stop reaction by adding 40 mM EDTA: 500 mM Stock: 1:12.5



23 Spin down for 5 minutes at 1200 rpm.

```
© 00:05:00 Spin down
```

24 Resuspend in PBS – place on ice.



25~ Add Proteinase K 200  $\mu g/\mu l$  to reverse x-link buffer = 1:100 from 20 mg/ml Stock.



26 Prepare 96-well plates with 25 μl reverse x-link buffer per well.

■25 μl reverse x-link buffer (per well)

27 quick-spin the plates to move all buffer to bottom

#### FACS and reverse x-link

- 28 Set up single cell 96 or 384 well sort at a BD FACS Ariall or similar and in Diva software check index sort box.
- 29 If stained: measure controls (in bulk): e.g. unstained, FMO, uniquely stained samples. then measure and record fluorescence intensity.
- 30 Spin down plates immediately after sort.
- 31 Use heat sealer for sealing plates.
- 32~ 60 °C reverse x-link O.N. (min 4 h) in incubator or PCR cycler.

```
§ 60 °C reverse x-link O.N.

© 04:00:00 reverse x-link O.N.
```

## PCR and clean up

33 Inactivate Proteinease K for 10 min at 80°C.

```
§ 80 °C ProteinaseK inactivation

© 00:10:00 ProteinaseK inactivation
```

34 PCR: 100 wells/plate: per well: 25 μl NEBNext PCR MM, 2.5 μl Ad1, 2.5 μl Ad2 individual Primerstock concentration: 25 μM





NEBNext High-Fidelity 2X PCR Master Mix

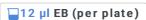
- 250 rxns

by New England Biolabs
Catalog #: M0541L

# 35 Cycling conditions:

| 72°C | 5min  |           |
|------|-------|-----------|
| 98°C | 30 s  |           |
| 98°C | 10 s  | 20 cycles |
| 63°C | 30 s  |           |
| 72°C | 1 min |           |

Cleanup using MinElute vacuum:
pool wells into a reservior,
add PB (6ml /96 well plate)
Wash with PE (10 ml/96well plate),
spin down,
elute in 12 µl EB per 96 well plate.





MinElute Reaction Cleanup Kit

by Qiagen

Catalog #: 28204

 $37 \quad \text{Library quantification using KAPA and BioAnalyzer. If required library cleanup using PAGE gel} \\$ 

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