

DEC 29, 2023

# O DoTA-seq V3.1 Forked from DoTA-seq V3.1

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DoTA-seq



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## OPEN ACCESS



#### DOI:

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**Protocol status:** Working We use this protocol and it's working

Created: Aug 24, 2023

Last Modified: Dec 29,

2023

### **ABSTRACT**

This protocol describes the process of DoTA-seq generating a single cell sequencing library from a cell suspension. This workflow can be performed in two days, with the PCR step happening overnight. Before beginning this workflow make sure to have:

- 1. The necessary microfluidics devices prepared and ready to go
- 2. The multiplex DoTA-seq target primers validated to work together without generating large molecular weight primer dimers.

Please read the publication for further details.

### **GUIDELINES**

**Strongly recommend** all pre-PCR steps (setting up reagents, washing gels) to be done in a PCR Clean hood. This has two purposes:

- 1. Reduce PCR contamination of templates which can strongly effect single-cell PCR reactions.
- 2. Reduce dust contamination of reagents which can clog devices and cause failures.

### **MATERIALS**

ddPCR Supermix for probes (no dUTP) BioRad
Sciences Catalog #1863024

MetaPolyzyme Sigma
Aldrich Catalog #MAC4L-5MG

Lysozyme from chicken egg white Sigma
Aldrich Catalog #L6876

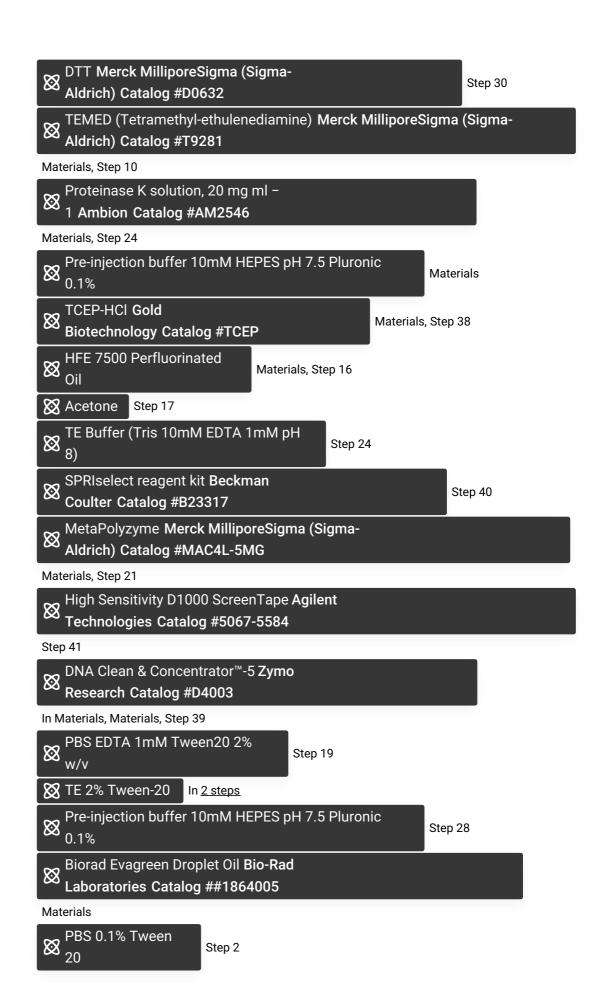
# **PROTOCOL integer ID:** 86909

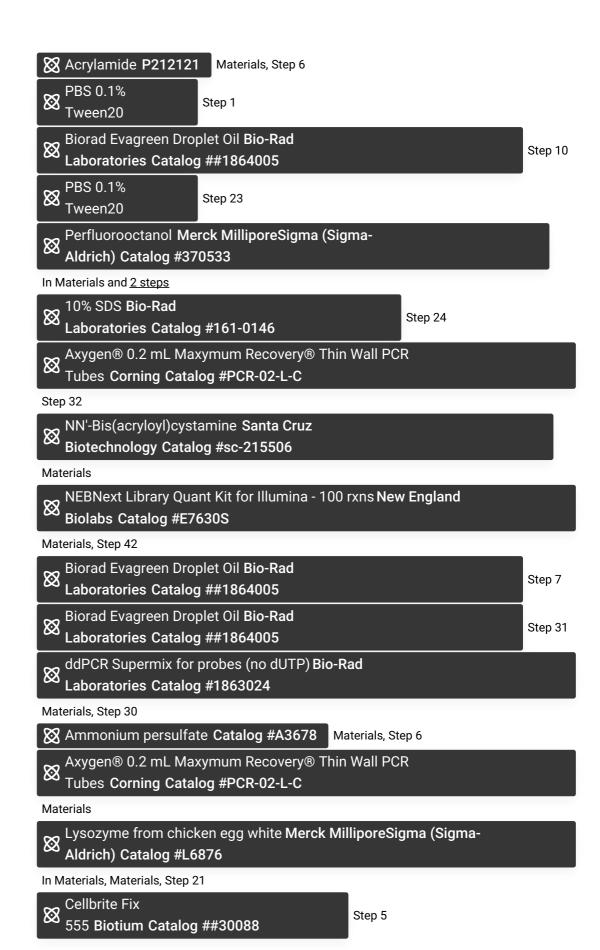
HFE 7500 Perfluorinated Oil Perfluorooctanol Sigma Aldrich Catalog #370533 TCEP-HCI Gold **Biotechnology Catalog #TCEP** NN'-Bis(acryloyl)cystamine Santa Cruz Biotechnology Catalog #sc-215506 Ammonium persulfate Catalog #A3678 🔀 Acrylamide P212121 TEMED (Tetramethyl-ethulenediamine) Sigmaaldrich Catalog #T9281 Biorad Evagreen Droplet Oil BioRad Sciences Catalog ##1864005 DNA Clean & Concentrator™-5 Zymo Research Catalog #D4003 DNA Clean & Concentrator™-5 Zymo Research Catalog #D4003 Axygen® 0.2 mL Maxymum Recovery® Thin Wall PCR Tubes Corning Catalog #PCR-02-L-C NEBNext Library Quant Kit for Illumina - 100 rxns New England Biolabs Catalog #E7630S SYBR Green Thermo Fisher Scientific Proteinase K solution, 20 mg ml -1 Ambion Catalog #AM2546 Lysozyme from chicken egg white Sigma Aldrich Catalog #L6876 Pre-injection buffer 10mM HEPES pH 7.5 Pluronic 0.1%

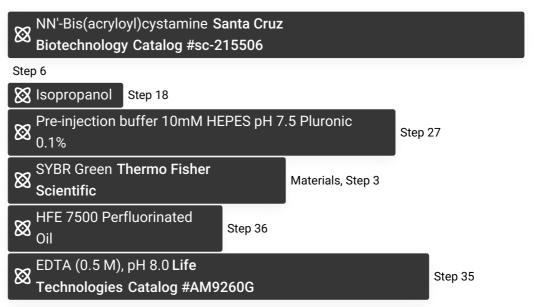
### Safety information

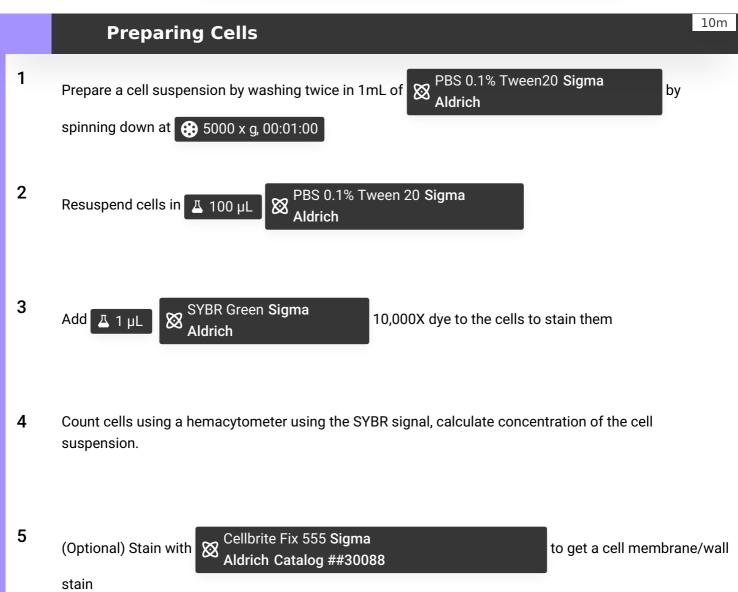
Unpolymerized Acrylamide is toxic, handle with care and dispose according to regulations

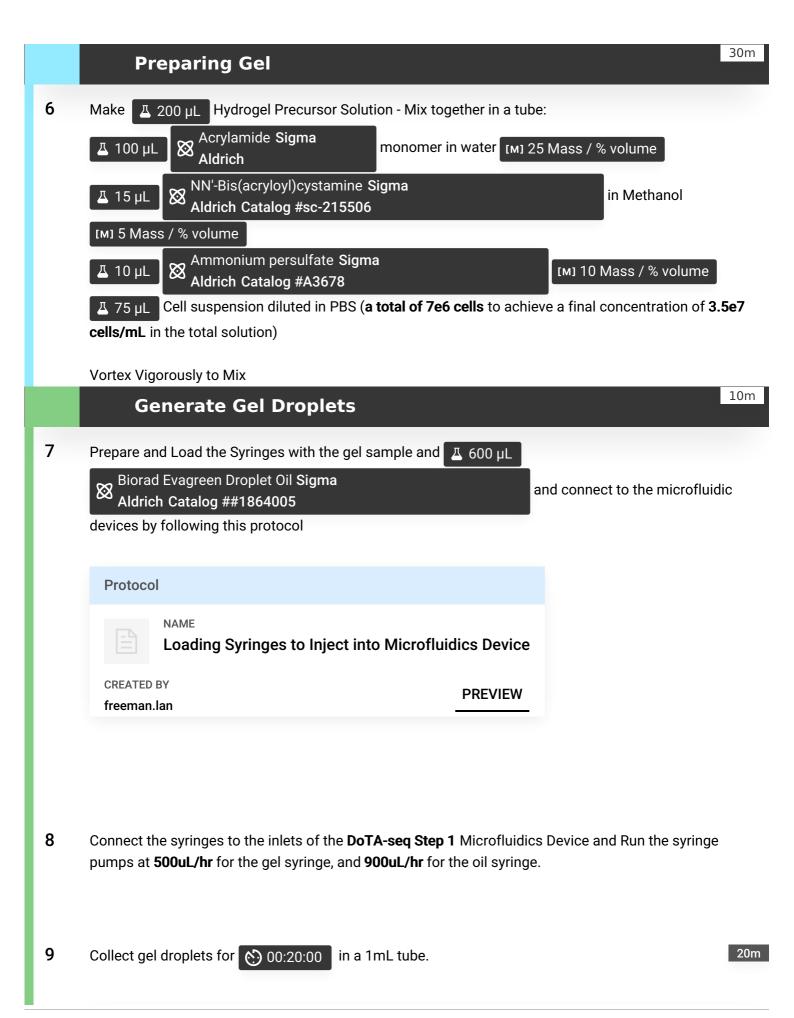
### PROTOCOL MATERIALS











#### Note

Sometimes the initial droplet formation produces polydisperse droplets. In this case, wait 2 min for the bad emulsion to leave the outlet tubing into a waste tube, then begin collecting in the collection tube.

10 Make Δ 200 μL Gel Polymerization Oil - mix together in a tube:



Aldrich Catalog #T9281

Add the Gel polymerization oil to the collected droplets, invert slowly 3 times to mix, and Incubate the 10m tube containing droplets at 37 °C for 00:10:00 to complete polymerization of the gel matrix.

### Note

You can now look at the emulsion under the microscope using

© Countess slides Thermo Fisher Scientific Catalog #C10228 to determine the encapsulation ratio of your cells. SYBRGreen and CF555 signal should be concordant and correspond to cells.

### Breaking out gels from emulsion

- Pulse spin the emulsion in a centrifuge to close pack the emulsion and drain the oil to the bottom of the tube.
- Use a pipette to remove the oil from the bottom of the tube, leaving just the emulsion

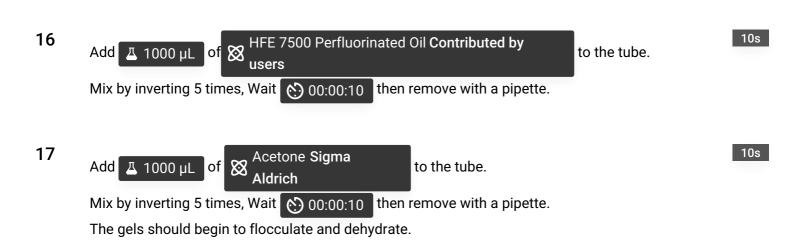
Add 200 µL Perfluorooctanol Sigma to break the emulsion

1111

Vortex, then Wait 00:01:00 for the emulsion to break.

Pulse spin again and remove the oil in the bottom of the tube with a pipette.

15



Add Δ 1000 μL of Sigma Isopropanol Sigma Aldrich

Mix by inverting 5 times. Wait Δ 00:00:10, then remove with a pipette

Mix by inverting 5 times, Wait 00:00:10 then remove with a pipette.

The gels should dehydrate and become hard.

Note: Do not wait too long as it could cause the gels to irreversibly aggregate into clumps.



Note

You can now look at the gels under the microscope using

© Countess slides Thermo Fisher Scientific Catalog #C10228 to determine the encapsulation ratio of your cells.

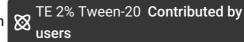
For **SYBR Staining**, you should first wash a small aliquot in PBS 2% Tween to remove background SYBR+ oil droplets before visualization on the microscope.

You should see some loss in CF555 signal as the acetone and alcohol wash removes some bacterial membranes.

### Lysing Bacteria



These gels can be stored at 4 °C in



for several days

without impacting DoTA-seq results.

#### Note

You can now look at the gels under the microscope using

Countess slides Thermo Fisher Scientific Catalog #C10228 to determine the encapsulation ratio and lysis efficiency of your cells.

It is advised to restain with SYBR and CF555 to get best signal. Lysed cells should exhibit SYBR signal but no CF555 Signal.

### **Barcoding the Cells**

7m

27 Wash the gels three times in  $\boxed{A}$  1000  $\mu$ L

Pre-injection buffer 10mM HEPES pH 7.5 Pluronic 0.1% **Sigma**Aldrich

28 Resuspend gels in Δ 100 μL

Pre-injection buffer 10mM HEPES pH 7.5 Pluronic 0.1% Sigma

Load the gels into a syringe following the protocol described in this excellent visual protocol.

### **CITATION**

Demaree B, Weisgerber D, Lan F, Abate AR (2018). An Ultrahigh-throughput Microfluidic Platform for Single-cell Genome Sequencing.. Journal of visualized experiments: JoVE.

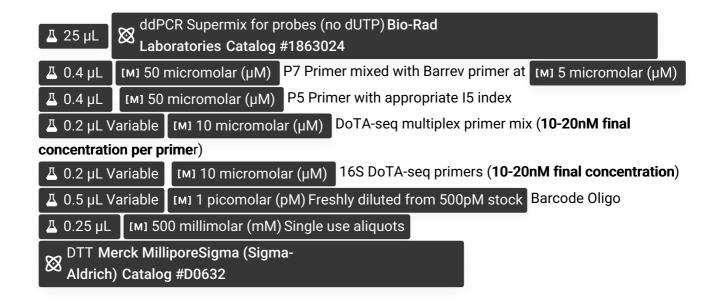
LINK

https://doi.org/10.3791/57598

Alternatively, for a simpler version, you can also use a P200 pipette to directly pipette the gel into a syringe backfilled with HFE7500.

30 Generate a PCR Master Mix (This mix gives about ~10,000 cells per library - Scale up as required)

20m



### Note

The ratio of 16S to DoTA-seq target primers mix can be varied depending on the relative amplification efficiencies. The best way to determine is to start from equal concentrations, then adjust based on the sequencing results (do most cells contain more 16S reads than target reads?)

### Note

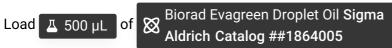
### IMPORTANT - BARCODE CONCENTRATIONS MAY NEED TO BE MEASURED

Typically, 0.5uL of 1pM barcode will give approximately 1 barcode for every 10 droplets. However, it is best to measure the barcode encapsulation rate by making PCR droplets containing the barcodes at the expected dilution and P7 and Barrev primers targeting the barcode for amplification. Visualize the resulting PCR emulsion using SYBRgreen staining under the microscope to obtain the real encapsulation ratio. Typically, the real barcode concentration can be ~5 fold off from the expected concentration based on manufacturer's labelling.

### Note

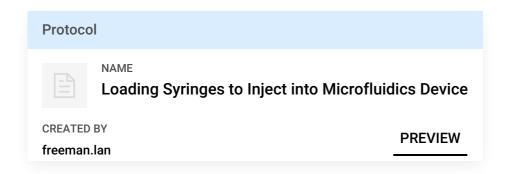
Barcode oligos should always be freshly diluted from 500pM to 1pM before use, as we have found gradual loss of barcodes over time in a 1pM solution.

Load the PCR mastermix into the syringe following this protocol



into a syringe

following this protocol



Connect the syringes to the **DoTA-seq Step 2** microfluidics device.

7m

Run the syringe pumps at **200uL/hr** for the gel and PCR mastermix, and **800uL/hr** for the oil syringe. Collect droplets in an



- Use a pipette to remove the oil in the PCR tube, leaving just the emulsion layer (it's okay to have a little bit of oil remaining).
- Thermocycle the PCR emulsion as follows:

4h



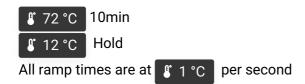
40 cycles of:

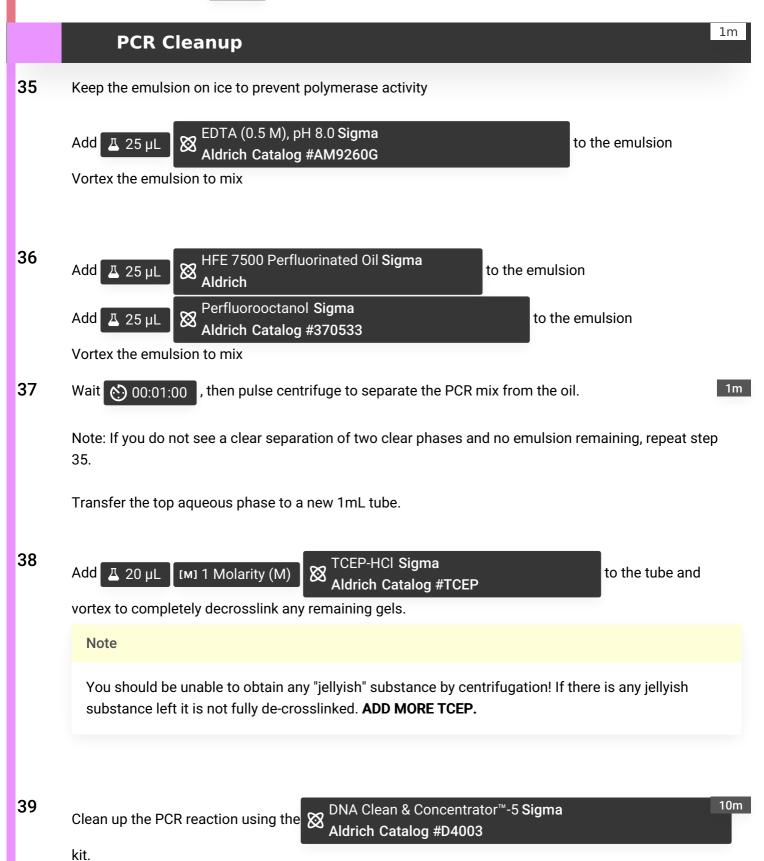
- \$\cdot\$ 95 °C
   30s

   \$\cdot\$ 72 °C
   10s

   \$\cdot\$ 60 °C
   5 min
- \$ 72 °C 30s

Final incubation of:





Elute in 🗸 50 µL Elution Buffer.

40 Remove primer dimers and free barcodes using the

10m

SPRIselect reagent kit Sigma
Aldrich Catalog #B23317

with 0.7X volume of beads.

41 Check the resulting library for primer dimers using

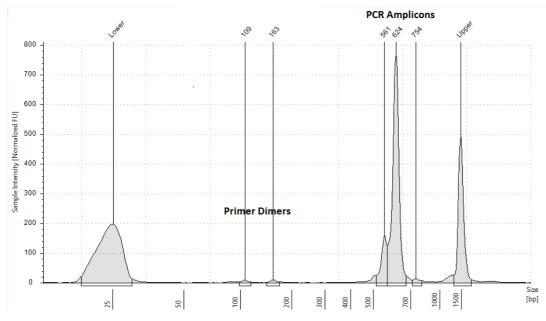
10m



with a High Sensitivity D1000 ScreenTape Sigma Aldrich Catalog #5067-5584

Other high sensitivity capillary electrophoresis methods will also work.

There should be minimal primer dimers on the trace. Below is an example of an acceptable trace.



Example of an acceptable Tapestation trace.

42 Quantify the library using a qPCR library quantification kit such as

1h

NEBNext Library Quant Kit for Illumina - 100 rxns Sigma
Aldrich Catalog #E7630S

### Note

Note that you must use a PCR based library quantification kit as not all amplicons contain all the adaptors for sequencing and therefore will throw off sequence non-specific forms of quantification!

43 Sequence the library on an Illumina sequencer using Custom Sequencing Primers listed here.

DoTA-seq-Oligo-Sequences.xlsx