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## FlowFISH with PrimeFlow

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**Protocol status:** Working

**We use this protocol and it's working**

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## Abstract

### Detect RNA expression in single cell with PrimeFlow

## Materials

ThermoFisher PrimeFlow RNA Assay Kit (88-18005-210)

positive control probe: RPL13A (Thermo VA4-13187-PF)

probe against gene of interest: MESDC1 (Thermo VA1-3010837-PF)

## Troubleshooting



## Before start:

- 1 Make sure the heat block temp is set to 40°C using the digital monitor. The heat block should be ON at least a day in advance to stabilize temperature
- 2 Turn on Hyb oven (use to pre-warm target probe diluent)
- 3 Make FACS staining buffer: 0.5% BSA in PBS, filter sterilized
  - 10% BSA stock: 3g BSA (light sensitive powder in 4°C MISC) + 30mL PBS
  - Store at 4°C
- 4 Set vortex to ~1500RPM

## A. Fixation and permeabilization

- 5
  - leave ~100µL when removing buffer, vortex to resuspend pellet
  - invert to mix when adding 1mL and tap the bottom if some cells stuck
  - pipette up and down to mix for fixation, permeabilization, and hybridization
  - may be in bulk, use volumes that cells don't exceed 10M cells/mL
  - pre-warm PrimeFlow RNA **Wash Buffer** to room temperature
- 6 Aliquot 5-10M cells in PBS per sample
- 7 Add 1mL PBS per sample, pipette to mix, and spin down at 500g at 4°C for 5min, discard supernatant, resuspend in ~100µL by vortexing gently
- 8 Prepare **Fixation Buffer 1** by mixing equal parts of PrimeFlow RNA **Fixation Buffer 1A** and PrimeFlow RNA **Fixation Buffer 1B**
  - need 1mL per sample: 500µL Buffer 1A + 500µL Buffer 1B
  - mix gently by inverting, don't vortex or vigorously shake, prepare fresh
  - keep at 4°C
- 9 Add 1mL **Fixation Buffer 1**, **pipette** to mix, incubate in dark (put cardboard box on it) rotating on rotator for 30min at 2–8°C in cold room
  - fixation time is critical, do NOT overshoot since the cells will lose integrity
- 10 Prepare 1X RNA **Permeabilization Buffer** with RNase Inhibitors by diluting PrimeFlow RNA **Permeabilization Buffer** (10X) and **RNase Inhibitors** (100X) with RNase-free water
  - need 2mL per sample: 200µL Perm Buffer + 1.78mL H<sub>2</sub>O + 20µL RNase Inhibitor



- mix gently by inverting, don't vortex or vigorously shake, prepare fresh
  - keep at 4°C
- 11 Spin down cells at 800g at 4°C for 5min, discard supernatant
- speed is critical here: slower spin will lose a lot of cells
- 12 Add 1mL RNA **Permeabilization Buffer** with RNase Inhibitors, **pipette** to mix, and spin down at 800g at 4°C for 5min, discard supernatant
- 13 Repeat wash with 1mL RNA **Permeabilization Buffer**
- set centrifuge to room temp
- 14 Prepare 1X RNA **Fixation Buffer 2** by diluting PrimeFlow RNA **Fixation Buffer 2** (8X) with PrimeFlow RNA **Wash Buffer**
- need 1mL per sample: 125µL Fix Buffer 2 + 875µL RNA Wash Buffer
  - mix gently by inverting, don't vortex or vigorously shake, prepare fresh
  - keep at room temp
- 15 Add 1mL RNA **Fixation Buffer 2**, pipette to mix, and incubate for 60min in the dark at room temperature while rotating
- 16 Spin down cells at 800g at room temp for 5min, discard supernatant, resuspend (~100µL) by vortexing gently
- 17 Add 1mL PrimeFlow RNA **Wash Buffer**, invert to mix, and spin down at 800g at room temp for 5min, discard supernatant, resuspend (~100µL) by vortexing
- if in bulk, the cells should be transferred to the 1.5mL tubes from kit
- 18 Repeat wash with 1mL RNA **Wash Buffer**
- \* can store samples overnight at 4°C, then the last wash with RNase Inhibitors

## B. Target Probe hybridization

- 19 Thaw Probe Sets (20X), including positive control (RPL13A, Type 4) on ice; pre-warm PrimeFlow **RNA Target Probe Diluent** to 40°C
- 20 take 2µL cells from "unstained" sample into 18µL PBS → measure on countess with trypan blue and record cell amount and % live
- **critical that the residual volume after all washes be as close to 100µL as possible, use the markings on the 1.5mL tubes to assist**
- 21 Dilute **Probe Sets** (20X) 1:20 in PrimeFlow RNA **Target Probe Diluent**, mix thoroughly by **pipetting** up and down

- need 100µL per sample: 5µL Target Probe + 5µL RPL13A Probe + 90µL Diluent
- for unstained sample: 100µL Diluent
- keep at 40°C

- 22 Add 100µL diluted **Target Probe(s)** into the cell suspension (~100µL), pipette to mix, briefly vortex, then incubate for 2h at 40°C
- do not pipette solutions onto the walls of the tubes, and samples should be mixed well before incubating
  - vortex samples to mix every 30min
  - temp is critical for hybridization

- 23 Add 1mL PrimeFlow RNA **Wash Buffer**, invert to mix, and spin down at 800g for 5min, discard supernatant, resuspend (~100µL) by vortexing gently

- 24 Prepare PrimeFlow RNA **Wash Buffer** with RNase Inhibitors by diluting **RNase Inhibitors** (100X) with RNA **Wash Buffer**

- need 1mL per sample: 10µL RNase Inhibitor + 990µL Wash Buffer
- mix gently by inverting, prepare fresh
- keep at room temp

- 25 Add 1mL PrimeFlow RNA **Wash Buffer** with RNase Inhibitors, **invert** to mix, and spin down at 800g for 5min, discard supernatant, resuspend (~100µL) by vortexing gently
- Store samples overnight in the dark at 4°C

## C. Signal amplification

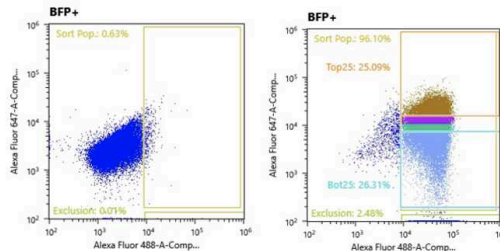
- 26
- Pre-warm samples and PrimeFlow RNA **Wash Buffer** to room temperature
  - Pre-warm PrimeFlow RNA **PreAmp Mix**, PrimeFlow RNA **Amp Mix**, and PrimeFlow RNA **Label Probe Diluent** to 40°C
- 27 Add 100µL PrimeFlow RNA **PreAmp Mix** into the cell suspension (~100µL), **pipette** to mix, briefly vortex, then incubate for 1.5h at 40°C
- do not pipette solutions onto the walls of the tubes, and samples should be mixed well before incubating
- 28 Add 1mL PrimeFlow RNA **Wash Buffer**, **invert** to mix, and spin down at 800g for 5min, discard supernatant, resuspend (~100µL) by vortexing gently
- 29 Repeat wash two times with 1mL RNA **Wash Buffer**, for a total of three washes
- 30 Add 100µL PrimeFlow RNA **Amp Mix** into the cell suspension (~100µL), **pipette** to mix, briefly vortex, then incubate for 1.5h at 40°C

- do not pipette solutions onto the walls of the tubes, and samples should be mixed well before incubating
- 31 Add 1mL PrimeFlow RNA **Wash Buffer**, **invert** to mix, and spin down at 800*g* for 5min, discard supernatant, resuspend (~100µL) by vortexing gently
- 32 Repeat wash with 1mL RNA **Wash Buffer**
- 33 Dilute PrimeFlow RNA **Label Probes** (100X) 1:100 in PrimeFlow RNA **Label Probe Diluent**
  - need 100µL per sample: 1µL Label Probes + 99µL Diluent
  - keep at 40°C
- 34 Add 100µL diluted **Label Probes** into the cell suspension (~100µL), **pipette** to mix, briefly vortex, then incubate for 1h at 40°C
  - do not pipette solutions onto the walls of the tubes, and samples should be mixed well before incubating
- 35 Add 1mL PrimeFlow RNA **Wash Buffer** at room temp, **invert** to mix, and spin down at 800*g* for 5min, discard supernatant, resuspend (~100µL) by vortexing gently
- 36 Repeat washes with warm (35°C) 1mL RNA **Wash Buffer** 5 times
- 37 Add 1mL **FACS staining buffer** (or PrimeFlow RNA **storage buffer**), **invert** to mix, and spin down at 800*g* for 5min, discard supernatant, resuspend (~100µL) by vortexing gently
  - Samples can be stored in the dark at 4°C for up to three days before analysis
  - QC: take stained vs unstained samples and check fluorescence with microscope

## D. Flow cytometric analysis

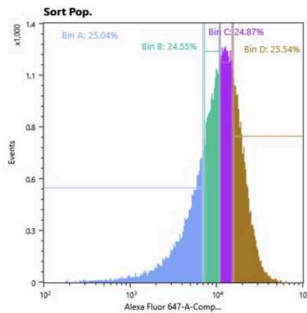
- 38 Add 100µL FACS buffer to cells, transfer all to filter, quick spin
- 39 Add 100µL FACS buffer to cap to wash remaining cells from filter, quick spin
- 40 FACS:
  - 400-500µL of an optimally concentrated sample (20M cells/mL) take ~30min
  - when low volume left, add ~200µL staining buffer can recover more cells
  - setup:
    - o sort population should be <10% of unstained sample

- o exclusion should be <5% of stained sample
  - § if more than 5%, increase voltage or decrease compensation
- o compensate for each gene: AF647 over AF488
  - § RPL13A (AF488) expression = total RNA, compensate gene of interest expression (AF647) relative to total RNA
  - § average AF488 for Top & Bottom 25% AF647 should be within 10% of average AF488 for all sort population



Sort Pop.	64,557	96.10%	64.56%	
Alexa Fluor 488-A-Comp...				44,080
Alexa Fluor 647-A-Comp...				12,052
Top25	16,858	25.09%	16.86%	
Alexa Fluor 488-A-Comp...				45,768
Alexa Fluor 647-A-Comp...				21,142
Bot25	17,678	26.31%	17.68%	
Alexa Fluor 488-A-Comp...				43,948
Alexa Fluor 647-A-Comp...				4,742

- sort:



## E. gDNA Extraction after FlowFISH

- 41 - Homemade **ChIP Lysis Buffer**:
  - o 1% SDS, 10mM EDTA, 50mM Tris-HCl, pH7.5
  - o store at 4°C
  - o for 50mL
  - Make Lysis Buffer fresh or warm at room temp 30-60min to solubilize precipitate
  - Always include a no cell control until PCR to check contamination
  - Set thermocycler:
    - o 65°C Hold → 65°C 10min → 37°C Hold → 37°C 30min → 65°C Hold → 65°C 2h → 95°C 20min → 4°C Hold
  - Protocol for **1M** cells



- 42 Spin cells down for 10min at 800*g* at 4°C, remove supernatant
- 43 Resuspend cells in 70μL **ChIP Lysis Buffer**, transfer to 96-well plate
- 44 Incubate at 65°C for 10min
- 45 When samples cool to 37°C, add 2μL **RNase Cocktail** (Invitrogen AM2286) mix well by pipetting
- 46 Incubate at 37°C for 30min
- 47 Add 10μL **Proteinase K** (Thermo 25530049), mix well by pipetting
- 48 Incubate at 65°C for 2h, then 95°C for 20min
- 49 Store at 4°C (can store overnight)
- 50 Beads clean with 0.7X AMPure beads: 70μL AMPure beads for ~100μL samples
  - a. warm AMPure beads to room temp
  - b. add beads, pipette to mix, and bind DNA for 2min
  - c. wash 3 times with 150μL 70% EtOH
  - d. let beads dry for 5-10min after 3<sup>rd</sup> wash
  - e. elute with ~40-50μL H<sub>2</sub>O, elute for 2min, transfer to a new plate/tube





## Protocol references

ThermoFisher PrimeFlow RNA Assay (**88-18005-210**)

Manual: [https://www.thermofisher.com/document-connect/document-connect.html?url=https://assets.thermofisher.com/TFS-Assets%2FMSG%2Fmanuals%2FMAN0019788\\_PrimeFlowRNAAssay\\_UG.pdf](https://www.thermofisher.com/document-connect/document-connect.html?url=https://assets.thermofisher.com/TFS-Assets%2FMSG%2Fmanuals%2FMAN0019788_PrimeFlowRNAAssay_UG.pdf)

Adapted from: Fulco CP, Nasser J, Jones TR, Munson G, Bergman DT, Subramanian V, Grossman SR, Anyoha R, Doughty BR, Patwardhan TA, Nguyen TH, Kane M, Perez EM, Durand NC, Lareau CA, Stamenova EK, Aiden EL, Lander ES, Engreitz JM. Activity-by-contact model of enhancer-promoter regulation from thousands of CRISPR perturbations. Nat Genet. 2019 Dec;51(12):1664-1669. doi: 10.1038/s41588-019-0538-0