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

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## Optimising sample multiplexing oligos by flow cytometry

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#### **ABSTRACT**

Optmisation of sample multiplexing oligos by scRNA-Seq is costly and time consuming. A cheaper and faster method is to use a flow cytometry read-out with fluorescent detection oligonucleotides.

This method can also be used to mix samples with different fluorescently labelled oligos and investigate signal swapping.

#### **IMAGE ATTRIBUTION**

Made with Biorender.com



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GUIDELINES

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**Keywords:** scRNA-Seq, facs, sample multiplexing, hashtag

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#### **MATERIALS**

- Phosphate Buffered Saline, without magnesium and calcium
- Sample multiplexing oligos (MULTI-Seq, CellPlex or Hashtag Antibody)
- 10ug/mL DAPI
- 30% BSA stock solution
- Fluorescent detection oligonucleotides

A	В	С
Name	Sequence	Modification
A647_FB2_detect	/5Alex647N/CCTTAGCCGCTA ATAGGTGAGC	5' Alexa 647 modification
A594_FB2_detect	/5Alex594N/TTGCTAGGACCG GCCTTAAAGC	5' Alexa 594 modification
A594_FB1_detect	/5Alex594N/TTGCTAGGACCG GCCTTAAAGC	5' Alexa 594 modification
A647_Total-SeqC_detect	/5Alex647N/CTGTCTCTTATA CACATCTCCG	5' Alexa 647 modification
AF647_oligo_dT_detect	/5Alex647N/TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	5' Alexa 647 modification

Fluorescent detection oligonucleotides. Order with HPLC modification

MULTI-Seq barcoding oligos. I substituted the poly-A tail for 10x Genomics feature barcode 2 sequence.

A	В	С
Name	Sequence	Barcode
multiSeq_FB2_BC 2	CCTTGGCACCCGAGAATTCCA CCACAATGGCTCACCTATTAG CGGCTAAGG	CCACAA TG
multiSeq_FB2_BC	CCTTGGCACCCGAGAATTCCA TGAGACCTGCTCACCTATTAG CGGCTAAGG	TGAGAC CT
multiSeq_FB2_BC	CCTTGGCACCCGAGAATTCCA GCACACGCGCTCACCTATTAG CGGCTAAGG	GCACAC GC
multiSeq_FB2_BC 5	CCTTGGCACCCGAGAATTCCA AGAGAGAGGCTCACCTATTAG CGGCTAAGG	AGAGA GAG
multiSeq_FB2_BC	CCTTGGCACCCGAGAATTCCA TCACAGCAGCTCACCTATTAG CGGCTAAGG	TCACAG CA
multiSeq_FB2_BC 7	CCTTGGCACCCGAGAATTCCA GAAAAGGGGCTCACCTATTAG	GAAAA GGG

А	В	С
	CGGCTAAGG	
multiSeq_FB2_BC 8	CCTTGGCACCCGAGAATTCCA CGAGATTCGCTCACCTATTAG CGGCTAAGG	CGAGAT TC
multiSeq_FB2_BC 9	CCTTGGCACCCGAGAATTCCA GTAGCACTGCTCACCTATTAG CGGCTAAGG	GTAGCA CT
multiSeq_FB2_BC 10	CCTTGGCACCCGAGAATTCCA CGACCAGCGCTCACCTATTAG CGGCTAAGG	CGACC AGC
multiSeq_FB2_BC 11	CCTTGGCACCCGAGAATTCCA TTAGCCAGGCTCACCTATTAG CGGCTAAGG	TTAGCC AG
multiSeq_FB2_BC 12	CCTTGGCACCCGAGAATTCCA GGACCCCAGCTCACCTATTAG CGGCTAAGG	GGACC CCA
multiSeq_FB2_BC 13	CCTTGGCACCCGAGAATTCCA CCAACCGGGCTCACCTATTAG CGGCTAAGG	CCAACC GG

#### SAFETY WARNINGS

Please follow all Manufacturer safety warnings and recommendations.

# Prepare multiplexing reagent

- 1 For CellPlex and Hashtag antibody the reagent comes ready to use.
  - 1.1 Prepare a dilution series for titration if desired

## **MULTI-Seq oligo preparation**

2 Mix anchor and barcode strands in 1:1 molar ratio in PBS (without FBS or BSA at 2 μM concentration (10X stock).

## Sample preparation

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4 I use suspension cell lines to titrate cell multiplexing oligos so the sample preparation is easy. 4.1 Prepare a single-cell suspension of the sample to be tested. 4.2 Wash cells once in plain PBS without additives. I centrifuge suspension cell lines at 400xg. Resuspend in PBS. 4.3 Count cells and transfer 100k to 1M cells (preferably 500k) into a 1.5mL tube for labelling Labelling samples with multiplexing oligos 5 This is largely based on original protocols but with the addition of a fluorescent secondary oligo. 5.1 Resuspend cells in 180uL of plain PBS. 5.2 Add 20 µL 10X Anchor:Barcode solution and pipette gently to mix 10 – 15 times. 5.3 Incubate on ice for 5 minutes.

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5.4 Add 20 µL Co-Anchor solution and pipette gently to mix. 5.5 Incubate 5 minutes longer on ice. 5.6 Add 1.5mL of 1% BSA in PBS (ice cold) to quench. 5.7 Add 1.5mL of 1% BSA in PBS (ice cold) to quench. 5.8 Add 1mL of 1% BSA in PBS (ice cold) to quench. 5.9 Centrifuge cells at 4°C 400xg 5min. 5.10 Resuspend cells in the remaining 100uL of supernatant then transfer to a new 1.5mL tube. 5.11 Add 1.9mL 1% BSA in PBS and spin 5 minutes 400g at 4°C.

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Analyse with relevant software.

6.4

