

CITE-seq assay scheme – 10x Genomics single cell 3' v2 chemistry

cDNA production from 10x single cell 3' v2 chemistry ([From 10x Genomics](#))



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Gel bead oligo primer annealed to antibody-conjugated oligo – **partial small RNA read 2 handle**

10x gel bead oligo
 5' -CTACACGACGCTCTCCGATCT-NNNNNNNNNNNNNNNN-NNNNNNNNNN-TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTVN-3'
 3' AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAABxxxxxxxxxxxxx**ACCTTAAGAGCCACGGTTCC**

antibody-conjugated CITE-seq oligo



double stranded cDNA after RT (and cDNA PCR with additive) – 112 nt:

ADT cDNA additive primer: 5' -**CCTTGGCACCCGAGAATT*C*C**-3'

5' -CTACACGACGCTCTCCGATCT-NNNNNNNNNNNNNNNN-NNNNNNNNNN-TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTVNxxxxxxxxxxxxx**TGGAATTCTCGGGTGCCAAGG**-3'
 3' -GATGTGCTGCGAGAAGGCTAGA-NNNNNNNNNNNNNNNN-NNNNNNNNNN-AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAABxxxxxxxxxxxxx**ACCTTAAGAGCCACGGTTCC**-5'

Library PCR – addition of **P5** and **P7** ends. SI-PCR (**P5**) and RPI-x (**P7**) – shown with 6 nt i7 index

5' -CTACACGACGCTCTCCGATCT-NNNNNNNNNNNNNNNN-NNNNNNNNNN-TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTVNxxxxxxxxxxxxx**TGGAATTCTCGGGTGCCAAGG**-3'
 3' -GATGTGCTGCGAGAAGGCTAGA-NNNNNNNNNNNNNNNN-NNNNNNNNNN-AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAABxxxxxxxxxxxxx**ACCTTAAGAGCCACGGTTCC**-5'

AATGATACGGCGACCACCGA-GATCTACACTCTTCCCTACACGACGCTC-3'

3' -**ACCTTAAGAGCCACGGTTCCTTGAGGTCAGT**Gxxxxxx**TAGAGCATACGGCAGAAGACGAAC**

CITE-seq ADT final library – 190 nt (192 nt with 8 nt i7 index)

READ 1 -->
 AATGATACGGCGACCACCGA-GATCTACACTCTTCCCTACACGACGCTCTCCGATCT-NNNNNNNNNNNNNNNN-NNNNNNNNNN-TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTVNxxxxxxxxxxxxx**TGGAATTCTCGGGTGCCAAGGACTCCAGTCAC**xxxxxx**ATCTCGTATGCCGCTCTTCGCTGT**
 TTAGTATGCCGCTGGTGGCT-CTAGATGTGAGAAGGATGTGCTGCGAGAAGGCTAGA-NNNNNNNNNNNNNNNN-NNNNNNNNNN-AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAABxxxxxxxxxxxxx**ACCTTAAGAGCCACGGTTCCTTGAGGTCAGT**Gxxxxxx**TAGAGCATACGGCAGAAGACGAAC**
 <-- READ 2
 Ab barcode (e.g. 12 nt)

CITE-seq assay scheme: 10x Genomics single cell 3' v2 chemistry. Marlon Stoekius & Peter Smibert. Contact: nyqctech@gmail.com