Corrigendum: FB5P-seq: FACS-Based 5-Prime End Single-Cell RNA-seq for Integrative Analysis of Transcriptome and Antigen Receptor Repertoire in B and T Cells

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A Corrigendum on

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In the original article, there was a mistake in **Table S3** as published. The sequence of Read2_SP, the Read2 sequencing primer, was incorrect. The corrected **Table S3** appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

TABLE S3 | Primers used in FB5P-seq.

Primer name	Function in FB5P-seq	Sequence 5'-3'
(dT)30_Smarter	Prime Reverse Transcription at 3' ends of polyadenylated mRNA and introduce PCR handle	TGCGGTATCTAAAGCGGTGAGTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTVN
TSO_BCx_UMI5_TATA	Perform template switching and introduce 8 nt. well-specific barcode & 5 nt. UMI	AGACGTGTGCTCTTCCGATCTXXXXXXXXNNNNNTATArGrGrG
Satija_PCR	Forward primer for LD-PCR amplification of cDNA libraries	AGACGTGTGCTCTTCCGATCT
SmarterR	Reverse primer for LD-PCR amplification of cDNA libraries	TGCGGTATCTAAAGCGGTGAG
i7_BC1	Forward primer for 5'-end enrichment of tagmented library and incorporation of plate-specific barcode	CAAGCAGAAGACGGCATACGAGATCCTGGTAGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
i7_BC2		CAAGCAGAAGACGGCATACGAGATTAAGCATGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
i7_BC3		CAAGCAGAAGACGGCATACGAGATAGATGTGCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
i7_BC4		CAAGCAGAAGACGGCATACGAGATGTCGAGCAGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
i7_BC5		CAAGCAGAAGACGGCATACGAGATGAATTGCTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
i7_BC6		CAAGCAGAAGACGGCATACGAGATAAGCAACTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
i7_primer	Forward primer for amplification of sequencing library	CAAGCAGAAGACGGCATACGA
Read1_SP	Read1 sequencing primer	TCGTCGGCAGCGTCAGATGTGTATAAGAGACAG
i7_SP	Index Read i7 sequencing primer	AGATCGGAAGAGCACACGTCTGAACTCCAGTCAC
Read2_SP	Read2 sequencing primer	GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT