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# 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

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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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.

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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	TGCGGTATCTAAAGCGGTGAGTTTTTTTTTTTTTTTTTT
TSO_BCx_UMI5_TATA	Perform template switching and introduce 8 nt. well-specific barcode & 5 nt. UMI	AGACGTGTGCTCTTCCGATCTXXXXXXXNNNNNTATArGrGrG
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	CAAGCAGAAGACGGCATACGAGATCCTGGTAGGTGACTGGAG TTCAGACGTGTGCTCTTCCGATCT
i7_BC2		CAAGCAGAAGACGGCATACGAGATTAAGCATGGTGACTGGAG TTCAGACGTGTGCTCTTCCGATCT
i7_BC3		CAAGCAGAAGACGGCATACGAGATAGATGTGCGTGACTGGAG TTCAGACGTGTGCTCTTCCGATCT
i7_BC4		CAAGCAGAAGACGGCATACGAGATGTCGAGCAGTGACTGGAG TTCAGACGTGTGCTCTTCCGATCT
i7_BC5		CAAGCAGAAGACGGCATACGAGATGAATTGCTGTGACTGGAG TTCAGACGTGTGCTCTTCCGATCT
i7_BC6		CAAGCAGAAGACGGCATACGAGATAAGCAACTGTGACTGGAG TTCAGACGTGTGCTCTTCCGATCT
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