

CORRIGENDUM

A novel fluorescence-based multiplex PCR assay for rapid simultaneous detection of *CEBPA* mutations and *NPM* mutations in patients with acute myeloid leukemias

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Since the publication of the above paper, the authors have noticed an error in the sequence of primers used for TAD1, TAD2 and bZIP amplification in Table 1.

The correct Table 1 is shown below:

The authors apologize for any inconvenience caused.

Table 1

Gene	Region	Sequence, 5'–3'	Position (nt)	Concentration (nM)	Amplicon (bp)
<i>CEBPA</i>	TAD1	Forward: VIC-5'-TCGGCCGACTTCTACGAGGC-3'	598–617	50	326
		Reverse: 5'-GCGCCCCGGGTAGTCAAAGT-3'	923–905	50	
	TAD2	Forward: NED-5'-TACCTGGACGGCAGGCTGGA-3'	1003–1022	75	251
		Reverse: 5'-TGCAGGTGCATGGTGGTCT-3'	1253–1235	75	
	bZIP	Forward: FAM-5'-AGAAGTCGGTGGACAAGAACAGCAA-3'	1415–1439	50	246
		Reverse: 5'-AGTTGCCCATGGCCTTGAC-3'	1660–1642	50	

Abbreviation: CEBPA, CCAAT/enhancer-binding protein-alpha.

Note: Sequence numbering is according to GenBank accession no. U34070 for *CEBPA*.