

Erratum to: Expression of myeloperoxidase and gene mutations in AML patients with normal karyotype: double *CEBPA* mutations are associated with high percentage of MPO positivity in leukemic blasts

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The original version of this article unfortunately contained some errors in Table 2 in the column headed “Amino acid changes”. The corrected table is given here.

We sincerely apologize for the errors.

The online version of the original article can be found under doi:[10.1007/s12185-011-0883-y](https://doi.org/10.1007/s12185-011-0883-y).

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Table 2 Genetic findings of the patients with *CEBPA* mutations

Patient	Category	Nucleotide changes	Amino acid changes	Comments
4	Double	218_219insC 1129_1130insATGTGGAGACGCAGC AGAAGGTGCTGGAGCTGACCAAGTGAC AATGACCCGCTGCGCAAGC	H24fsX107 K326_R327insHVETQKQKVLELTSDNDRLRK	Produces N-terminal stop codon In-frame insertion in bZIP
6	Double	200_218delinsCT 1087_1089dup	S17fsX101 K313dup	Produces N-terminal stop codon In-frame duplication in bZIP
7	Double	368_369insA 1080_1082del	F73fsX107 Q312del	Produces N-terminal stop codon In-frame deletion in bZIP
13	Double	303_316del 1062_1063insTTG	L52fsX102 K304_Q305insL	Produces N-terminal stop codon In-frame insertion in bZIP
19	Double	215_225del 1101_1102insCAGCGCAACGTGGAGACGCAGC AGAAGGTGCTGGAGCTG	P23fsX103 L317_T318insQRNVETQKQKVLEL	Produces N-terminal stop codon In-frame insertion in bZIP
22	Double	213del 1064_1129dup	P23fsX159 K326_R327msQRNVETQKQKVLELTSDNDRLRK	Produces N-terminal stop codon In-frame insertion in bZIP
27	Double	324_328dup 1062_1063insTTG	S61fsX161 K304_Q305insL	Produces N-terminal stop codon In-frame insertion in bZIP
39	Double	213del 1081_1086dup	P23fsX159 Q311_Q312dup	Produces N-terminal stop codon In-frame duplication in bZIP
47	Double	397del 1101_1102insCAGCGCAACGTGGAGACGCA GCAGAAGGTGCTGGAGCTG	Q83fsX159 L317_T131insQRNVETQKQKVLEL	Produces N-terminal stop codon In-frame insertion in bZIP
49	Double	297_304del 758del	E50fsX104 P204fsX317	Produces N-terminal stop codon Frameshift between TAD2 and bZIP; produces stop codon in bZIP
35	Single	1087_1089dup	K313dup	In-frame duplication in bZIP

Nucleotide numbering was performed according to NCBI Entrez accession no. XM_009180.3, in which the major translational start codon starts at nucleotide position 151. The locations of functional domains are derived from Mueller and Pabst.1

bZIP basic leucine zipper region, *TAD2* second transactivation domain