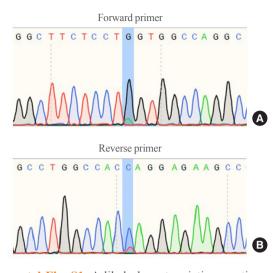
EnM



**Supplemental Fig. S1.** A likely hematopoietic somatic mosaic variant c.914G>A detected in *DNMT3A*. (A) Electropherogram obtained using forward primer. (B) Electropherogram obtained using reverse primer. This variant showed variant allele frequency of 18% in the exome sequencing data.