Electronic Supplementary Material (ESI) for Toxicology Research.

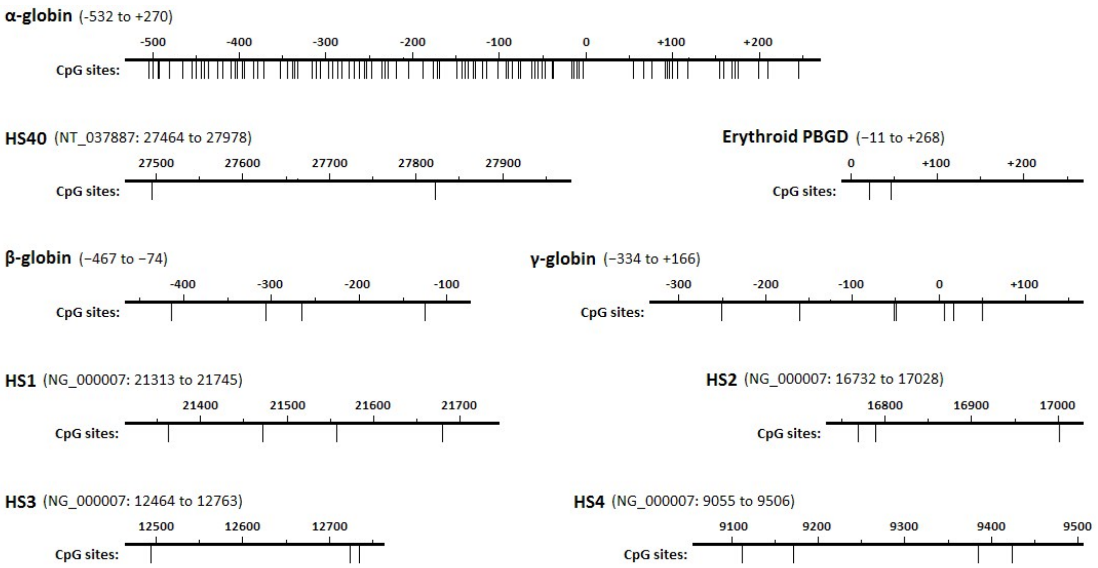

Fig. S1 The erythroid genes and their far-upstream regulatory elements in which the DNA methylation levels were analyzed in this study. The black line indicated the length of analyzed sequence of each genes or regulatory elements. The below black bars indicated CpG sites in analyzed sequences.