

Association between *SLC19A1* gene polymorphism and high dose methotrexate toxicity in childhood acute lymphoblastic leukaemia and non Hodgkin malignant lymphoma: introducing a haplotype based approach

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SUPPLEMENTARY TABLE 1. Characteristics of the selected *SLC19A1* SNPs

SNP	Nucleotide change	AA / Aa / aa	Allele frequency (MAF)**	Location on chromosome 21	
rs1131596 *	T > C	TT / CT / CC	0.479	46957916	5' UTR
rs1051266	G > A	GG / GA / AA	0.491	46157794	exon 2
rs2838958	T > C	TT / TC / CC	0.409	46948567	intron 4
rs2838956	T > C	TT / TC / CC	0.473	46945024	intron 5
rs17004785	C > G	CC / CG / GG	0.118	46932618	3' UTR
rs12483377	C > T	CC / CT / TT	0.055	46931109	3' UTR
rs2838951	C > G	CC / CG / GG	0.333	46929720	3' UTR

* Also known as rs3177999; ** Data from the SNP database; AA = homozygous normal genotype, Aa = carrier of one polymorphic allele, aa = homozygous polymorphic genotype