

**Supplementary Table 2 Haplotype analysis of 2 SNPs**

CHR	SNP	BP	MAF			P-value					Gene	rel.loci		
			A1	A2	ADR <sup>a</sup>	Non-ADR <sup>b</sup>	allelic	dominant	recessive	minimum	OR	95% CI		
2	rs3820706	152957411	A	G	0.34	0.43	8.26E-05	1.07E-01	8.13E-09	8.13E-09	1.47	(1.21-1.79)	CACNB4	-1818
2	rs16830728	152981335	G	T	0.33	0.41	1.11E-04	6.16E-02	7.24E-08	7.24E-08	1.47	(1.21-1.78)	STAM2	0

Haplotype/SNP allele(s)	Haplotype frequencies	ADR, Non-ADR frequencies	P value	OR*	Effect
GT	0.58	0.65, 0.37	5.91E-05	1.48	Risk
AG	0.38	0.32, 0.40	2.00E-04	0.69	Protective
AT	0.03	0.02, 0.03	0.571	0.84	Protective
GG	0.01	0.01, 0.01	0.6	0.77	Protective

SNP, single nucleotide polymorphism; ADR, adverse drug reaction; OR, odds ratio.

\*ORs are calculated using the nonrisk haplotype as reference

<sup>a</sup>Individuals who developed grade 2 alopecia.

<sup>b</sup>Individuals who did not developed any ADRs after chemotherapy.