

Supplementary Table S1. Hardy-Weinberg analyses for HLA-G 3'UTR genotypes in women with RM and controls.

SNP	<i>Recurrent miscarriage (n=23)</i>			<i>Controls (n=46)</i>		
	common <i>P</i>	homozygotes <i>P</i>	heterozygote <i>P</i>	common <i>P</i>	homozygotes <i>P</i>	heterozygotes <i>P</i>
14-bp	0.1444	0.3020	0.3020	0.1220	0.3119	0.2870
+3003	#	0.9370	&	0.5888	0.7548	0.6054
+3010	0.7171	0.7637	0.7526	0.1379	0.2975	0.2953
+3027	#	0.9653	&	0.8602	0.9474	0.8626
+3035	0.6481	0.7965	0.6619	0.7823	0.9045	0.7874
+3142	0.7171	0.7637	0.7526	0.2322	0.3997	0.3987
+3187	0.5487	0.6508	0.5836	0.3077	0.3077	0.2440
+3196	0.4238	0.5347	0.4688	0.2315	0.3908	0.2679
+3422	0.9250	0.9520	0.9292	0.5149	0.7170	0.5321
+3496	#	0.8996	&	0.5888	0.7548	0.6054
+3509	0.5855	0.6588	0.6280	0.2938	0.4428	0.3332

All Hardy-Weinberg analyses. *P*, p value. # Too many parameters for chi-square test. & Too few cases to calculate p-value.