

Supplementary Table S6. HLA-G 3'UTR genotypic polymorphisms in the offspring of women with recurrent miscarriage and uneventful pregnancy.

	<i>RM offspring (n=23)</i>		<i>Controls (n=45)</i>		<i>OR</i>	<i>95% CI</i>	<i>P</i>	<i>P_c</i>
3003 CC	1	4.3%	1	2.2%	n.c.			
	8	34.8%	10	22.2%	1.94	0.63-5.95	0.245	1.000
	14	60.9%	34	75.6%	ref.			
3010 CC	5	21.7%	11	24.4%	0.58	0.16-2.02	0.389	1.000
	15	65.2%	19	42.2%	ref.			
	3	13.0%	15	33.3%	0.25	0.06-1.04	0.057	0.969
3027 AA	0	0.0%	0	0.0%	n.c.			
	3	13.0%	3	6.7%	2.10	0.39-11.34	0.389	1.000
	20	87.0%	42	93.3%	ref.			
3035 CC	17	73.9%	35	77.8%	ref.			
	6	26.1%	10	22.2%	1.24	0.38-3.97	0.722	1.000
	0	0.0%	0	0.0%	n.c.			
3142 CC	5	21.7%	11	24.4%	0.71	0.20-2.50	0.598	1.000
	14	60.9%	22	48.9%	ref.			
	4	17.4%	12	26.7%	0.52	0.14-1.95	0.335	1.000
3187 AA	13	56.5%	20	44.4%	ref.			
	9	39.1%	21	46.7%	0.66	0.23-1.88	0.436	1.000
	1	4.3%	4	8.9%	n.c.			
3196 CC	11	47.8%	24	53.3%	ref.			
	11	47.8%	17	37.8%	1.41	0.50-4.00	0.516	1.000
	1	4.3%	4	8.9%	n.c.			
3422 CC	14	60.9%	29	64.4%	ref.			
	9	39.1%	14	31.1%	1.33	0.47-3.82	0.594	1.000
	0	0.0%	2	4.4%	n.c.			
3496 AA	1	4.3%	1	2.2%	n.c.			
	8	34.8%	10	22.2%	1.94	0.63-5.95	0.245	1.000
	14	60.9%	34	75.6%	ref.			
3509 GG	11	47.8%	23	51.1%	ref.			
	11	47.8%	17	37.8%	1.35	0.48-3.85	0.571	1.000
	1	4.3%	4	8.9%	0.52	0.05-5.25	0.581	1.000
	0	0.0%	1	2.2%	n.c.			

Data are all n (%). All univariate logistic regression analysis. Per HLA-G genotype the highest prevalence was defined as the reference group. If percentages in a group were below 5%, no calculations were performed. *P*, *p* value; *P_c*, *p* value corrected for multiple comparisons; *OR*, odds ratio; *95% CI*, 95% confidence interval; n.c., not calculated; ref, reference group.