Table S2. Summary of MDR findings for RA

No. of factor	Combination	CV* consistency	Prediction error (%)	*P* value
2	-3479T>G (*CD86*),	9/10	33.4	0.019
	-3458A>G (*CD40LG*)			
3	-3479T>G (*CD86*),	6/10	43.1	0.451
	-3458A>G (*CD40LG*),			
	+49A>G (*CTLA4*)			
4	-3479T>G (*CD86*),	6/10	44.6	0.541
	-3458A>G (*CD40LG*),			
	+49A>G (*CTLA4*)			
	-1C>T (*CD40*)			
5	-3479T>G (*CD86*),	3/10	49.4	0.946
	-3458A>G (*CD40LG*),			
	+49A>G (*CTLA4*),			
	-1C>T (*CD40*),			
	IVS3+17T>C (*CD28*)			

*Cross validation.