

Supplemental Table S1. The frequency distribution of *IL10* gene polymorphisms for DLBCL and FL included in the current meta-analysis

Surname	Year	Country	Ethnicity	Source	Genotype method	Subtype	Case				Control			
							CC	CT	TT	All	CC	CT	TT	All
-819C>T (rs1800871) polymorphism														
Lech-Maranda	2004	France	Caucasian	HB	PCR-RFLP	DLBCL	107	81	11	199	53	46	13	112
Lan	2006	USA	Mixed	PB	TaqMan	DLBCL	81	67	7	155	329	211	34	574
Lan	2006	USA	Mixed	PB	TaqMan	FL	53	51	5	109	329	211	34	574
Wang	2006	USA	Mixed	PB	TaqMan	DLBCL	187	142	22	351	514	339	81	934
Wang	2006	USA	Mixed	PB	TaqMan	FL	157	94	17	268	514	339	81	934
Lech-Maranda	2007	France	Caucasian	HB	PCR-RFLP	FL	56	34	10	100	53	46	13	112
Purdue	2007	Australia	Caucasian	PB	TaqMan	DLBCL	113	54	5	172	295	170	23	488
Purdue	2007	Australia	Caucasian	PB	TaqMan	FL	123	71	9	203	295	170	23	488
Hosgood	2013	Asia	Asian	HB	TaqMan	DLBCL	562	469	89	1120	2033	1700	364	4097
Hosgood	2013	Asia	Asian	HB	TaqMan	FL	116	106	30	252	2033	1700	364	4097
-572C>A (rs1800872) polymorphism														
Lech-Maranda	2004	France	Caucasian	HB	PCR-RFLP	DLBCL	107	81	11	199	53	46	13	112
Lan	2006	USA	Mixed	PB	TaqMan	DLBCL	82	62	8	152	331	189	43	563
Lan	2006	USA	Mixed	PB	TaqMan	FL	55	44	10	109	331	189	43	563
Wang	2006	USA	Mixed	PB	TaqMan	DLBCL	185	141	22	348	515	342	81	938
Wang	2006	USA	Mixed	PB	TaqMan	FL	155	95	17	267	515	342	81	938
Lech-Maranda	2007	France	Caucasian	HB	PCR-RFLP	FL	56	34	10	100	53	46	13	112
Purdue	2007	Australia	Caucasian	PB	TaqMan	DLBCL	113	54	5	172	297	169	23	489
Purdue	2007	Australia	Caucasian	PB	TaqMan	FL	123	72	9	204	297	169	23	489
Hosgood	2013	Asia	Asian	HB	TaqMan	DLBCL	566	460	89	1115	2041	1685	362	4088
Hosgood	2013	Asia	Asian	HB	TaqMan	FL	118	104	32	254	2041	1685	362	4088

PB, Population based; HB, Hospital based; PCR-RFLP, Polymorphism chain reaction-restriction fragment length polymorphism; DLBCL, Diffuse large B-cell lymphomas; FL, Follicular lymphoma.