

Table S8. Association between gene mutations and patients' clinical course

<i>Gene</i>		NRG	RG	P.value
<i>MYD88 L265P</i>	mutant	4	4	0.678
	wild type	8	14	
<i>MYD88</i>	mutant	2	2	1
	wild type	10	16	
<i>CD79B Y196X</i>	mutant	7	10	1
	wild type	5	8	
<i>CD79B</i>	mutant	7	7	0.457
	wild type	5	11	
<i>PIM1</i>	mutant	2	6	0.419
	wild type	10	12	
<i>BTG2</i>	mutant	6	13	0.266
	wild type	6	5	
<i>TBL1XR1</i>	mutant	10	15	1
	wild type	2	3	
<i>TOX</i>	mutant	8	17	0.128
	wild type	4	1	
<i>B2M</i>	mutant	10	17	0.548
	wild type	2	1	
<i>CARD11</i>	mutant	10	15	1
	wild type	2	3	
<i>PRDM1</i>	mutant	10	17	0.548
	wild type	2	1	
<i>TNFAIP3</i>	mutant	11	16	1
	wild type	1	2	
<i>TMEM30A</i>	mutant	11	17	1
	wild type	1	1	
<i>PRKCD</i>	mutant	11	18	0.4
	wild type	1	0	

RG: recurrence group; NRG: non-recurrence group