

**Supplementary Table 1**

<b>cloning primers</b>	
<b>name</b>	<b>sequence</b>
5'-cREL TS1-Spel	<u>ggactagtc</u> cagaaggcagttgaagtgag
3'-cRELTS1-Sacl	cgagctcgaagagatgatggtggcttgg
5'-cREL TS2-Spel	<u>ggactagtg</u> aactctacttgtgcactgg
3'-cREL TS2-Sacl	cgagctccaggacattcaggtagcatg
5'-RELA-Spel	<u>ggactagtg</u> aagccctccaaaagcacttac
3'-RELA-Sacl	cgagctcgaaagagcaagagtccaagtg
5'-NFKBIA-Spel	<u>ggactagtg</u> atgactgtgtgttggagg
3'-NFKBIA-Sacl	cgagctcgtacaccatttacaggaggg
5'-IKBKB-Spel	<u>ggactagtc</u> gcttcctcagcagctgtgac
3'-IKBKB-Sacl	cgagctccttagatgctgtgagagacag
5'-IKBKG-Spel	<u>ggactagtg</u> cgagatacatgtcatggagtg
3'-IKBKG-Sacl	cgagctcgtccatcaatagtgactgg
5'-TAB1-Spel	<u>ggactagtc</u> cagtagagtgtgtgagtgcag
3'-TAB1-Sacl	cgagctcggtagctgtgtctcaaagaacc
5'-TAB2-Spel	<u>ggactagtg</u> aaggtgtgacaagatgggtg
3'-TAB2-Sacl	cgagctccaaacactgtgttgtgtcacc
5'-TRAF2-Spel	<u>ggactagtc</u> cttcatcaaggccattgtgg
3'-TRAF2-Sacl	cgagctccctaggaatgctcccttctctc
5'-BCL10-Spel	<u>ggactagtg</u> acccttactaggaagaacg
3'-BCL10-Sacl	cgagctcggaaactccaaaatagcactcac
5'-PIK3CB-Spel	<u>ggactagtc</u> caagtcagtagacaaaccaacc
3'-PIK3CB-NaeI	gccggcggcattgtgagtaaagagcacag
5'-MALT1-Spel	<u>ggactagtc</u> acttccaacttactctaggtc
3'-MALT1-Sacl	cgagctccaactatgtatcaactgtgaatac
5'-PLCG1-Spel	<u>ggactagtg</u> cttagacagaaaccaagcc
3'-PLCG1-Sacl	cgagctcaggatgatttatttggcagtcagatcttaagagg
5'-CD3E-Spel	<u>ggactagtc</u> cctgaatcagagacgcac
3'-CD3E-Sacl	cgagctcggagatgcaaatgaccatccag
<b>mutagenesis primers</b>	
<b>name</b>	<b>sequence</b>
5'-CD3E-BS2mut	ggactagtcctgaatcagagacgcacatctgaccctctggagaaTCGCGAGtcccgcctggcccagg
3'-CD3E-BS2mut	cctgggcccagcgggaCTCGGAGttctccagagggtcagatgctctctgattcaggactagtcc
5'-PIK3CB-mut	gtttggagaatggtgaaTCGCGAGaggaagaaatggatac
3'-PIK3CB-mut	gtatccatttcttctcctCTCGGAGttcaccattctccaaac
5'-PLCG1-BS1mut	cagcacacacaaaacctTCGCGAGcacatttgggctcag
3'-PLCG1-BS1mut	ctgagcccaaatgtgCTCGGAGaggtttgtgtgtgctg
5'-PLCG1-BS2mut	ccctcttaagatctgAGCGCTGaaataaatcatcctcagctcgcg

3'-PLCG1-BS2mut	cgagctcgaggatgatttatttCAGCGCTcagatcttaag aggg
5'-TAB2-mut	gaaagacttggttgccTCGCGATtaactgtgtacagtg
3'-TAB2-mut	cactgtacacagttaaTCGCGAaggcaaccaagtctttc
5'-NFKBIA-mut	cttatatccacactgcaTCGCGACTggccaaaacgtc
3'-NFKBIA-mut	gacgttttgggcccagTCGCGAtgcagtgatataag

