SUPPLEMENTARY TABLE 1. Compilation of c-MYC missense mutations in Burkitt's and other lymphomas.

Control Cont		on of c-MYC missense mutations in Burkitt's and other lymphomas.		
LLS binard properties PSST SQUELTEY, CORREST SQUELTEY, SQU	Source	Mutation(s)	Publication (PMID)	Notes
CA66 on line (BL) QUARL CORPUS LAST PROVIDED AND REST REST SHEET MAY COUNTY AND	, ,			
Life	LL3 bursal lymphoma cells	P63T	6322173	Westaway et al., 1984. Chicken Myc
BL1	CA46 cell line (BL)	V5I, P57S, P60H, L179V, Q306H	3921823	Showe et al., 1985.
BL2	Ly65 cell line (BL)	Q33H, Q36H, E39D, K51R, 62P, S64R, C70W, frameshift 136–159	3517879	Murphy et al., 1986.
Bit SSSP PR33,94T SSSP P	BL1	A44V, S46N, T58N	8220424	Bhatia et al., 1993. These are patient BL samples. Sequenced second exon only
BLT	BL2	P59S	8220424	Bhatia et al., 1993. These are patient BL samples. Sequenced second exon only
BL7	BL4	S62P, P63L,S64T	8220424	Bhatia et al., 1993. These are patient BL samples. Sequenced second exon only
BLT	BL5	Q34H, P63S	8220424	Bhatia et al., 1993. These are patient BL samples. Sequenced second exon only
BLT	BL7	A44G. D48N. L56P. T58A. P63L	8220424	Bhatia et al., 1993. These are patient BL samples. Sequenced second exon only
BL10				
BL11 BL12 NBSS B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL12 NBSS B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL14 C375. FL905. ARV 1788 B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL17 C375. E305. ARV 1788 B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL19 BL19 PSTS B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL20 BL19 BL20 PSTS B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL20 PSTS B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL20 PSTS B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL22 SSGA B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL23 SSGA B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL23 SSGA B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL23 SSGA B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL24 SSGA B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL25 SSGA B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL26 SSGA B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL27 SSGAPPA5A B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL28 SSGAPPA5A B20424 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL29 SSGAP B20444 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL29 SSGAPPA5A B20444 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL29 SSGAPPA5A B20444 Bhata et al., 1903. These are patient BL samples. Sequenced second conor only BL29 SSGAPPA5A B20444 B10444 B10544 B105444 B105444 B105444 B105444 B105				
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B.1.3				
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BL17 G37E_B300_A4V_T851 8220424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL18 B148 et al., 1963. These are patient BL amples. Sequenced second exon only BL10 P57S_P60H 8220424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL20 P57S_P60H 8220424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL21 N86T 8220424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL22 S18C S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL24 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL24 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL24 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL24 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL25 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL amples. Sequenced second exon only BL26 S200424 Bhatta et al., 1963. These are patient BL26				
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BL13 D118V 16094360 Hemann et al., 2005. BL patient samples. Full Myc sequencing				
	BL13	D118V	16094360	Hemann et al., 2005. BL patient samples. Full Myc sequencing

SUPPLEMENTARY TABLE 1 (cont'd). Compilation of c-MYC missense mutations in Burkitt's and other lymphomas.

	I (cont'd). Compilation of c-MYC missense mutations in Burkitt's and other lymphomas.	D 11: 1: (D111D)	N. f
Source	Mutation(s)	Publication (PMID)	Notes
BL14	D26G, F93L, F138T, V145L, S182G	16094360	Hemann et al., 2005. BL patient samples. Full Myc sequencing
BL17	F31L, P191S	16094360	Hemann et al., 2005. BL patient samples. Full Myc sequencing
BL18	P60S	16094360	Hemann et al., 2005. BL patient samples. Only first 90 aa of Myc sequenced
BL19	P57T	16094360	Hemann et al., 2005. BL patient samples. Only first 90 aa of Myc sequenced
BL20	P57S	16094360	Hemann et al., 2005. BL patient samples. Only first 90 aa of Myc sequenced
BL21	T58A	16094360	Hemann et al., 2005. BL patient samples. Only first 90 aa of Myc sequenced
BL23	P60S	16094360	Hemann et al., 2005. BL patient samples. Only first 90 aa of Myc sequenced
BL25	P57S	16094360	Hemann et al., 2005. BL patient samples. Only first 90 aa of Myc sequenced
BL26	T58A	16094360	Hemann et al., 2005. BL patient samples. Only first 90 aa of Myc sequenced
BL2	Q35H, S175N	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
BL30	Q35H, T58I, G90D, S203N	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
BL41	C25W, L149V, D251N	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
BL58	P2S, V5A, S6T, R10K, E39D, P42S, A44V,	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
BL70	T58I	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
CA46	V5I, P60H, L178V, Q306H	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
Raji cell line	S6T, F7I, R10K, R66H, G88D, S114N, D120G, L149V, C171S, D180N, I189M, S203R, S230A, P245S, S249T	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2386	S249N	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2389	F115L , S249N	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2392	P290H	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2402	A154G	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2406	E39D, S64R, L69V, S172C	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2430	P60A, E104D, F138C	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2431	C70W	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2432	Y22H, S62P	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2433	E39D, T58N	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2443	F138S, C171Y, L240F	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2448	S250R	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2523	F138S, F124L	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2524	A44T, A150D, S175N	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2531	L149V	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
SLN2539	P60A	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
Thomas cell line	T8S, Q36L, T95P, A183V	22885699	Schmitz et al. 2011. Derived from RNAseq and validated by Sanger
BL-1	S175N, S373R	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-4	T248I	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-5	D97G, S175R	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-8	H359R, N360S	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-13	V5G, E29D	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-14	V5A, S64R, V111A, S114N, N127K, E243R	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-15	S6T, T8A, E39D, T95S, V111M, I125M, Q131H, F138C, K148T, V194L,S203N	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-19	F138S	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-20	P59A	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-21	V5L, F93I, F138V, L149V, Q179H, S182R, S204T	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-25	N4K	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-26	S249R	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-28	Y74H, L149V, del 203–214, P245A	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
BL-29	S6N, F138S	23091298	Giulino-Roth et al. 2012. Targeted genome resequencing of BL patient samples
Namalwa cell line	S249R	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
JiJoye cell line	Y12S, Q98H	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
EB3 cell line	F138S	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
1102	S18F	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
667	S18R	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
513	S6R, C188W	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
744	S6R	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
1059	L14V	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
1060	T58N, S64N, D221N	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
479	T58N	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
672	P72A, S175T	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
475	V75A	23143597	Love et al. 2012. Exome resequencing and validated by Sanger
674	A76V	23143597	Love et al. 2012. Exome resequencing and validated by Sanger

SUPPLEMENTARY TABLE 1 (cont'd). Compilation of c-MYC missense mutations in Burkitt's and other lymphomas.

COLI ELIMENTARY TABLE I (CONTA). Compilation of C-in Commissions in Barriers and other symphomas.						
Source	Mutation(s)	Publication (PMID)	Notes			
1090	F138S, A154T	23143597	Love et al. 2012. Exome resequencing and validated by Sanger			
515	F138S	23143597	Love et al. 2012. Exome resequencing and validated by Sanger			
516	F138S	23143597	Love et al. 2012. Exome resequencing and validated by Sanger			
668	F138S, S249N, V330P, F374R	23143597	Love et al. 2012. Exome resequencing and validated by Sanger			
675	F138S	23143597	Love et al. 2012. Exome resequencing and validated by Sanger			
676	L176F, S203G, S249N, Q256S	23143597	Love et al. 2012. Exome resequencing and validated by Sanger			
698	S203T	23143597	Love et al. 2012. Exome resequencing and validated by Sanger			
510	S249N	23143597	Love et al. 2012. Exome resequencing and validated by Sanger			