

## Supplementary Material

### Network and Computational Drug Repurposing Analysis for c-Myc Inhibition in Burkitt Lymphoma

Yongmin Lee, Seungyeon Nam\*

\* Correspondence: Seungyeon Nam: nams@gachon.ac.kr

#### Supplementary Tables

**Table S1.** In network analysis, expression levels (log2 fold change) for each cell line regarding genes that showed down-regulation patterns (JQ1-treated vs. untreated and c-Myc knock-down vs. shControl) during c-Myc inhibition relative to control in at least two cell lines.

Down-regulation genes			
Gene	Daudi	ST486	Raji
CDK4	-0.129	-0.033	-0.331
FGFR1	-0.021	-0.128	-0.252
GRB2	-0.029	-0.013	-0.022
MAP2K1	-0.009	-0.045	-0.035
MAPK1	-0.010	-0.028	0.053
MAPK3	NA	-0.073	-0.695
HRAS	NA	-0.051	-0.126
ROCK2	NA	-0.023	-0.157
AKT1	NA	-0.008	-0.224
AKT2	NA	-0.005	-0.104
KRAS	-0.003	NA	-0.188
MYLK3	-0.116	-0.023	NA
CBLC	-1.381	-0.114	NA
ARHGEF6	-0.044	NA	-0.650
CRLF2	NA	-0.001	-0.140
RASGRP2	-0.137	NA	-0.389

<i>CBL</i>	NA	-0.122	-0.143
<i>STMN1</i>	-0.092	NA	-0.449
<i>ELK1</i>	NA	-0.044	-0.148

**Table S2.** In network analysis, expression levels (log2 fold change) for each cell line regarding genes that showed up-regulation patterns (JQ1-treated vs. untreated and c-Myc knock-down vs. shControl) during c-Myc inhibition relative to control in at least two cell lines.

Up-regulation genes			
Gene	Daudi	ST486	Raji
<i>PIK3CB</i>	0.090	0.008	0.053
<i>JAK1</i>	0.107	-0.042	0.488
<i>AKT3</i>	0.452	0.018	-0.079
<i>TYK2</i>	0.233	0.024	0.069
<i>SOS1</i>	0.062	0.010	0.156
<i>MYL12B</i>	0.036	NA	0.310
<i>FGFR2</i>	NA	0.037	0.117
<i>STAT2</i>	0.138	0.097	NA
<i>TPR</i>	NA	0.008	0.087
<i>CACNA1H</i>	0.366	NA	0.155
<i>RASGRP3</i>	NA	0.068	0.169
<i>CCND1</i>	0.738	0.060	NA
<i>CACNG5</i>	NA	0.137	0.033
<i>STAT1</i>	NA	0.048	0.482
<i>SOCS7</i>	NA	0.008	0.055
<i>PIK3CA</i>	NA	0.018	0.442
<i>PIAS3</i>	NA	0.015	0.093
<i>FGF11</i>	0.256	0.034	NA
<i>PIP5K1C</i>	NA	0.025	0.129
<i>RRAS2</i>	0.010	NA	0.136

**Table S3.** Based on the network analysis results through PATHOME-Drug, drug information related to genes in MAPK signaling pathway that contains many genes with consistent expression patterns relative to control (JQ1-treated vs. untreated and c-Myc knock-down vs. shControl).

Gene	Drug	Gene	Drug
<i>ARAF</i>	Adenosine Triphosphate	<i>FOS</i>	Nadroparin
<i>BRAF</i>	Dabrafenib		Pseudoephedrine
	Regorafenib	<i>GRB2</i>	Pegademase Bovine
	Sorafenib	<i>IGF1R</i>	Insulin Glargine
	Vemurafenib		Insulin Lispro
<i>CCND1</i>	Arsenic Trioxide		Insulin Regular
<i>EGFR</i>	Afatinib		Mecaserman
	Cetuximab		Porcine Insulin
	Erlotinib	<i>MAP2K1</i>	Bosutinib
	Gefitinib		Trametinib
	Lapatinib	<i>MAPK1</i>	Arsenic Trioxide
	Lidocaine		Isoprenaline
	Panitumumab	<i>MAPK3</i>	Sulindac
	Trastuzumab		Vorinostat
	Vandetanib	<i>PDGFRB</i>	Becaplermin
<i>FGF4</i>	Pentosan Polysulfate		Dasatinib
<i>FGFR1</i>	Palifermin		Imatinib
	Ponatinib		Pazopanib
	Regorafenib		Regorafenib
	Sorafenib		Sorafenib
<i>FGFR2</i>	Palifermin		Sunitinib
	Ponatinib	<i>RAF1</i>	Dabrafenib
	Regorafenib		Regorafenib
	Thalidomide		Sorafenib