

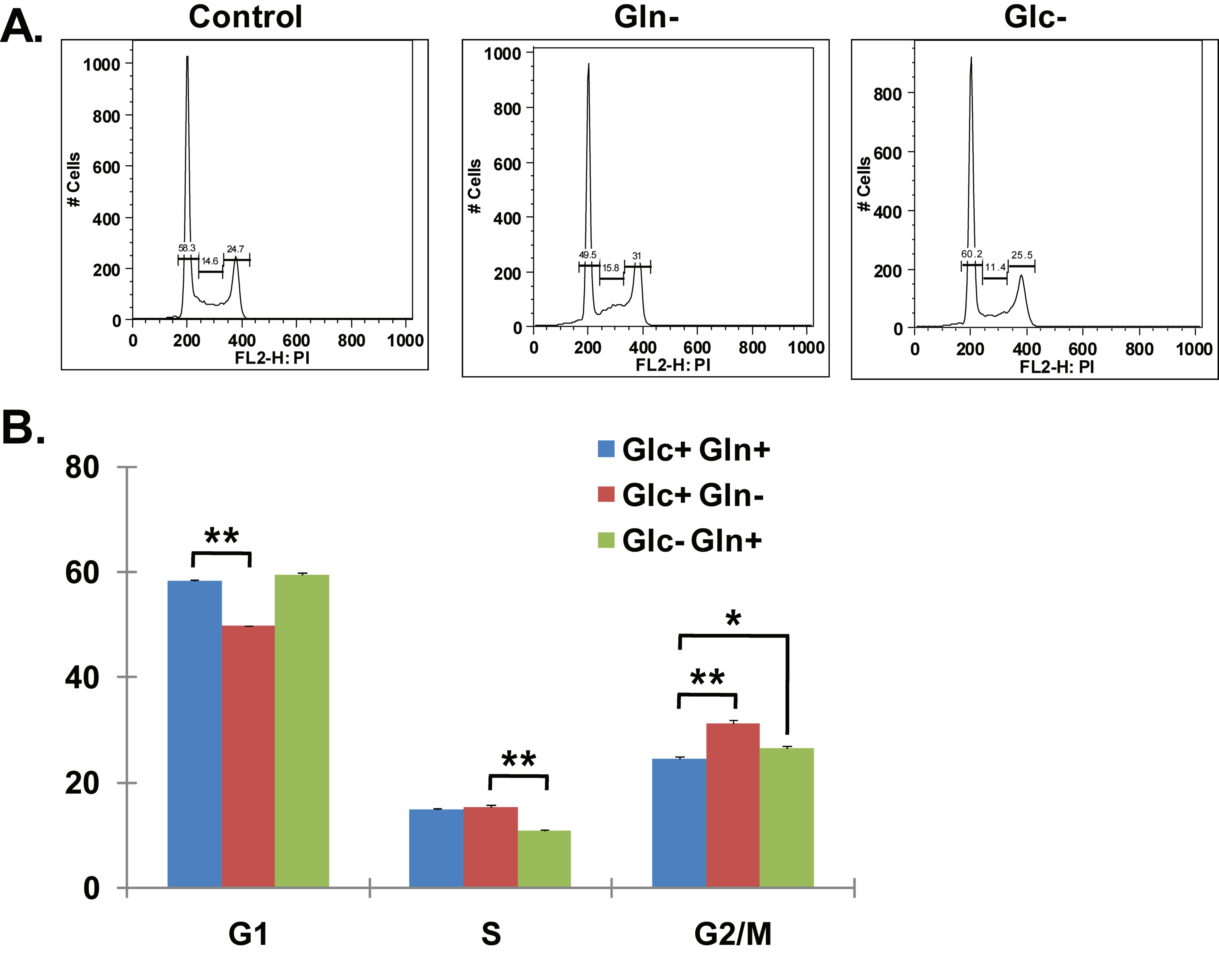
Supplemental Material to:

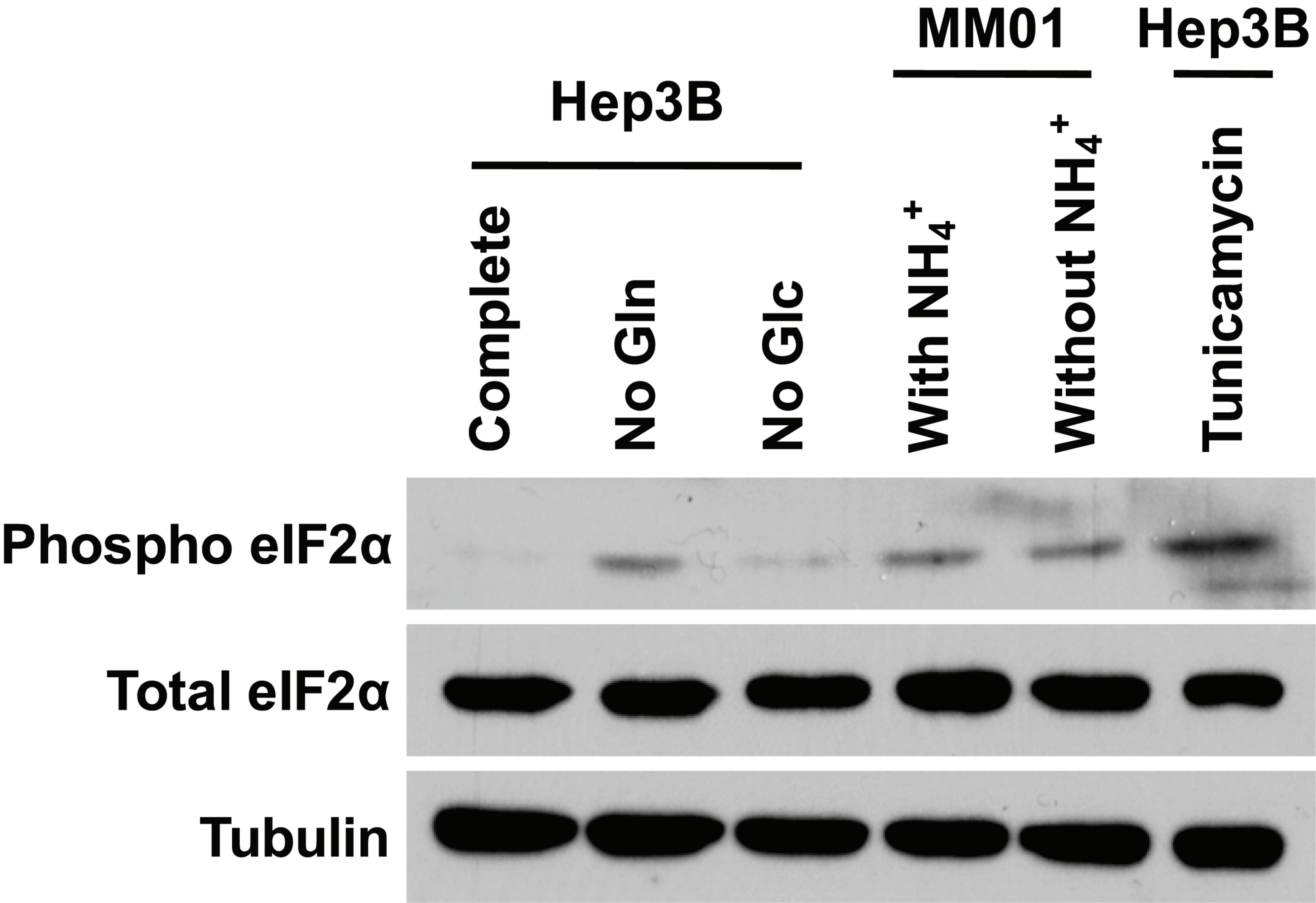
Shuo Qie, Dongming Liang, Chengqian Yin, Weiting Gu, Meng Meng, Chenguang Wang and Nianli Sang

Glutamine depletion and glucose depletion trigger growth inhibition via distinctive gene expression reprogramming

2012; 11(19) http://dx.doi.org/10.4161/cc.21944

http://www.landesbioscience.com/journals/cc/article/21944





Supplemental Table 1 (In a separated Excel file.)

The list of detected genes in Hep3B cells treated with either Gln- or Glc-depletion or MM01 cells compared to Hep3B cells in complete medium. (List based on Gln-depleted/Control with Log_2 Ratio >3 & <-3).

Supplemental Table 2 (In a separated Excel file.)

The lists of genes for Figure 2E as following:

	Sheet No.	Group
Gln-depletion vs Control cells	Sheet 1	Cell Cycle Control of Chromosomal Replication
	Sheet 2	Cell Cycle Regulation by BTG Family Proteins
	Sheet 3	Cell Cycle: G1/S Checkpoint Regulation
	Sheet 4	Cyclins and Cell Cycle Regulation
	Sheet5	Mitotic Roles of Polo-Like Kinase
	Sheet 6	Cell Cycle: G2/M DNA Damage Checkpoint Regulation
Glc-depletion vs Control cells	Sheet 7	Cell Cycle Control of Chromosomal Replication
	Sheet 8	Cell Cycle Regulation by BTG Family Proteins
	Sheet 9	Cell Cycle: G1/S Checkpoint Regulation
	Sheet 10	Cyclins and Cell Cycle Regulation
	Sheet 11	Mitotic Roles of Polo-Like Kinase
	Sheet 12	Cell Cycle: G2/M DNA Damage Checkpoint Regulation

Supplemental Table 3 Regulation Directions of TCF3 Target Genes Upon Gln-Depletion

Target Genes	Log₂Ratio	Published Findings	Prediction
WBP5	-3.343	Downregulates (1)	Activated
TYMS	-2.308	Downregulates (1)	Activated
TOP2A	-2.881	Downregulates (1)	Activated
TCF3	1.616	Upregulates (11)	Activated
SLC12A2	-1.115	Downregulates (1)	Activated
PGK1	-2.324	Downregulates (1)	Activated
NKX2-5	1.114	Upregulates (1)	Activated
MKI67	-1.600	Downregulates (1)	Activated
MAD2L1	-3.492	Downregulates (1)	Activated
KLHDC2	-2.862	Downregulates (1)	Activated
FGFR2	-2.384	Downregulates (3)	Activated
EPRS	-5.584	Downregulates (1)	Activated
DAB2IP	2.447	Upregulates (2)	Activated
CDH1	-3.429	Downregulates (4)	Activated
CCND3	1.020	Upregulates (1)	Activated
CCNA2	-2.753	Downregulates (1)	Activated
CA2	-1.991	Downregulates (1)	Activated
BCL2L11	1.057	Upregulates (3)	Activated
ATF3	-1.833	Downregulates (1)	Activated
ANLN	-2.477	Downregulates (1)	Activated
AATF	-4.070	Regulates (1)	
ACACB	1.779	Regulates (8)	
AP3D1	-2.799	Regulates (1)	
ARSA	1.258	Regulates (1)	
BID	-1.330	Regulates (1)	
CASP3	-3.254	Regulates (1)	
CASP6	-5.129	Regulates (1)	
CCNE1	-6.109	Regulates (1)	
CHRNA5	-1.648	Regulates (1)	
CPT1B	1.717	Regulates (1)	
CYP11A1	2.416	Regulates (1)	
DOCK1	-2.450	Regulates (1)	
E2F4	-4.240	Regulates (1)	
ETS2	1.267	Regulates (1)	
GADD45A	-3.341	Regulates (1)	
GATA2	-1.815	Regulates (1)	
GATA3	1.218	Regulates (2)	
GPAM	1.204	Regulates (1)	
ID2	-3.763	Regulates (1)	

IFNAR1	1.050	Regulates (1)	
JUND	1.293	Regulates (1)	
LRBA	-3.714	Regulates (1)	
MLLT4	1.394	Regulates (1)	
MSMO1	-1.107	Regulates (1)	
MYC	1.471	Regulates (1)	
NFIL3	1.909	Regulates (2)	
PAFAH1B3	-2.704	Regulates (1)	
PLCG2	2.595	Regulates (1)	
PTPRU	2.551	Regulates (1)	
RORA	1.148	Regulates (1)	
TF	-5.361	Regulates (1)	
TMEM180	2.032	Regulates (1)	
XBP1	-6.097	Regulates (1)	

^{*} TCF3 function predicted based on directions of target gene expression.

Supplemental Table 4 Regulation Directions of XBP1 Target Genes Upon Glc-Depletion

Genes in dataset	Log₂ Ratio	Findings	Prediction*
SRP68	1.159	Upregulates (1)	Activated
SERP1	1.667	Upregulates (2)	Activated
SEC24D	1.230	Upregulates (1)	Activated
SEC22B	1.050	Upregulates (1)	Activated
PIGA	1.179	Upregulates (1)	Activated
PDIA6	1.317	Upregulates (2)	Activated
MCFD2	1.227	Upregulates (1)	Activated
HYOU1	1.543	Upregulates (1)	Activated
HSPA13	2.207	Upregulates (1)	Activated
HMOX1	2.244	Upregulates (1)	Activated
HM13	1.191	Upregulates (1)	Activated
HERPUD1	2.349	Upregulates (1)	Activated
GOLPH3L	1.055	Upregulates (1)	Activated
GOLGA4	1.064	Upregulates (1)	Activated
EDEM1	1.292	Upregulates (7)	Activated
DNAJC1	1.357	Upregulates (1)	Activated
DNAJB9	3.093	Upregulates (6)	Activated
DERL1	1.290	Upregulates (1)	Activated
DDIT3	2.642	Upregulates (1)	Activated
COG6	1.033	Upregulates (1)	Activated
BFAR	1.241	Upregulates (1)	Activated
BET1	1.171	Upregulates (1)	Activated
ATF6	1.423	Upregulates (1)	Activated
ARFGAP3	1.811	Upregulates (1)	Activated
ALG2	1.010	Upregulates (1)	Activated
TTR	-1.702	Upregulates (1)	Inhibited
SERPINA1	-2.051	Upregulates (1)	Inhibited
RCN3	-1.208	Upregulates (1)	Inhibited
FKBP1B	-1.063	Upregulates (1)	Inhibited
FKBP11	-1.271	Upregulates (1)	Inhibited
CAT	-1.089	Upregulates (16)	Inhibited
APOA1	-1.039	Upregulates (1)	Inhibited
DNAJC3	1.068	Regulates (6)	
MYC	1.752	Regulates (2)	
TOP1	1.580	Regulates (1)	

^{*} XBP1 function predicted based on directions of target gene expression.

Supplemental Table 5 Regulation Directions of PPAR α Target Genes in MM01

Target Genes	Log(Ratio)	Findings	Prediction*
SORD	-1.221	Downregulates (1)	Activated
SOCS2	-1.220	Downregulates (2)	Activated
SELENBP1	-1.424	Downregulates (1)	Activated
PAH	-1.766	Downregulates (1)	Activated
NPC1L1	-1.295	Downregulates (2)	Activated
MBL2	-1.427	Downregulates (2)	Activated
KHK	-1.794	Downregulates (1)	Activated
INSIG1	1.118	Upregulates (2)	Activated
HPD	-1.749	Downregulates (1)	Activated
HNF4A	-1.284	Downregulates (2)	Activated
G0S2	1.614	Upregulates (1)	Activated
FTCD	-2.096	Downregulates (1)	Activated
FGG	-1.214	Downregulates (2)	Activated
FGB	-3.597	Downregulates (6)	Activated
FGA	-1.488	Downregulates (3)	Activated
EDN1	-3.070	Downregulates (2)	Activated
CEBPD	-1.241	Downregulates (1)	Activated
CDKN2A	1.221	Upregulates (8)	Activated
C6	-1.971	Downregulates (1)	Activated
C1S	-2.261	Downregulates (1)	Activated
C1R	-1.641	Downregulates (1)	Activated
ARG1	-4.019	Downregulates (1)	Activated
APOC3	-2.498	Downregulates (13)	Activated
APOC2	-1.155	Downregulates (2)	Activated
ACTA1	-3.146	Regulates (2)	
APOM	-1.507	Regulates (5)	
CFI	-1.277	Regulates (1)	
ECI2	-1.072	Regulates (1)	
FABP5	-1.361	Regulates (3)	
GLUD1	-2.143	Regulates (1)	
GPT	-1.644	Regulates (1)	
HPX	-3.111	Regulates (1)	
IFITM3	-1.042	Regulates (1)	
IGFBP1	-1.422	Regulates (7)	
IGFBP3	1.583	Regulates (1)	
SULT2A1	-2.190	Regulates (3)	
UGT2B4	-2.320	Regulates (1)	

^{*} PPAR α function predicted based on directions of target gene expression.