

## Supplementary Material

Supplementary Table S1 Primer sequences for qRT-PCR analyses.

qRT-PCR primer sequences	
mouse Actb forward primer	5'- GATCTGGCACCACACCTTCT -3'
mouse Actb reverse primer	5'- GGGGTGTTGAAGGTCTCAAA -3'
mouse Nfyav1 forward primer	5'- AAGTCCAGACCCTCCAGGTAGT -3'
mouse Nfyav1 reverse primer	5'- GATGGGTTGGCCTGTTGAT -3'
mouse Nfyav2 forward primer	5'- GCCATGGAGCAGTATACGACA -3'
mouse Nfyav2 reverse primer	5'- CCTGGACCTGCTGCTGAA -3'
mouse Pck1 forward primer	5'- CCTTTGGAAGCGGATATGGT -3'
mouse Pck1 reverse primer	5'- TTGCCTTCGGGGTTAGTTATG -3'
mouse G6pc forward primer	5'- ACTGTGGGCATCAATCTCCTCT -3'
mouse G6pc reverse primer	5'- GGGCGTTGTCCAAACAGAA -3'
human ACTB forward primer	5'- ACCAACTGGGACGACATGGAGAAA -3'
human ACTB reverse primer	5'- TAGCACAGCCTGGATAGCAACGTA -3'
human PCK1 forward primer	5'- GACATTGCCTGGATGAAGTTTG -3'
human PCK1 reverse primer	5'- TTCTTCTGGATGGTCTTGATGG -3'
human SLC1A5 forward primer	5'- TCCGCTTCTTCAACTCCTTCA -3'
human SLC1A5 reverse primer	5'- AAACCCACATCCTCCATCTCC -3'
human GLS forward primer	5'- TTCTCAGGGCAGTTTGCTTTC -3'
human GLS reverse primer	5'- TTGCCCATCTTATCCAGAGGA -3'
human GLUD1 forward primer	5'- AAGGCAAAGCCCTATGAAGGA -3'
human GLUD1 reverse primer	5'- CATTGGCACCTTCAGCAATG -3'
human GOT2 forward primer	5'- ACCGGGATGATAATGGAAAGC -3'
human GOT2 reverse primer	5'- CTTGCAAAATTCAGCCAGTCC -3'
human OGDH forward primer	5'- GCAGATGTGCAACGATGACC -3'
human OGDH reverse primer	5'- TGGAAGAAGTTGCCAGGAGTG -3'
human PCK1 forward primer for CUT&RUN assay	5'- GGTGCATCCTTCCCATGAAC -3'
human PCK1 reverse primer for CUT&RUN assay	5'- CTGGTTGGCAAAACACCACA -3'

## Tsujimoto and Ito et al. Supplementary Figure S1



Numl low 114 high 63 37 21

12 5 Number at risk low 219 86 high 152 48

1

9 3 3 0 6 1 1 1

19 16 2

**Supplementary Figure S1.** The expression of PCK1 and G6PC predicts better patient survival only for gluconeogenic organ cancer. Kaplan-Meier plots of overall survival of patients with gluconeogenic organ cancer (A; for PCK1, B; for G6PC) and non-gluconeogenic organ cancer (C; for PCK1, D; for G6PC). Data were obtained from the Kaplan-Meier plotter online tool. The hazard ratio (HR) and respective log-rank p-values identifying the high-expression group (red) and the low-expression group (black) are shown.



## Tsujimoto and Ito et al. Supplementary Figure S2

**Supplementary Figure S2.** NFYAv2 enhances the anti-tumor effects of gluconeogenesis. **(A)** Western blot analysis of the expression levels of PCK1 and NFYA in NFYAv1OE SK-Hep1 cells under normal or glucose deprivation conditions after 10 hours culture. **(B)** Representative fluorescence images of living cells (green) detected with Calcein-AM and dead cells (red) detected with Propidium iodide (PI) in wild-type, NFYAv1OE, and NFYAv2OE SK-Hep1 cells under normal or glucose deprivation conditions after 10 hours culture. **(C)** The bar graph shows the percentage of dead cells among the more than 1,500 cells counted each condition. Data are shown as mean and SEM. Unpaired two-tailed t-test was performed. (n.s.) not significant; (\*\*\*) P<0.001.



## Tsujimoto and Ito et al. Supplementary Figure S3

**Supplementary Figure S3.** Glutaminolysis is normal in NFYAv2OE SK-Hep1 cells. (A) A diagram illustrating the reaction of glutaminolysis and gluconeogenesis. (B) qRT-PCR analysis of the expression levels of SLC1A5, GLS, GLUD1, and GOT2 in wild-type and NFYAv2OE SK-Hep1 cells under normal or glucose deprivation conditions after 5 hours of culture. (C-E) The intracellular glutamine (C), glutamate (D), and a-KG (E) levels in wild-type and NFYAv2OE SK-Hep1 cells under normal or glucose deprivation conditions after 5 hours of culture. (F) qRT-PCR analysis of the expression levels of OGDH in wild-type and NFYAv2OE SK-Hep1 cells under normal or glucose deprivation conditions after 5 hours of culture. (F) qRT-PCR analysis of the expression levels of OGDH in wild-type and NFYAv2OE SK-Hep1 cells under normal or glucose deprivation conditions after 5 hours of culture. (F) qRT-PCR analysis of the expression levels of OGDH in wild-type and NFYAv2OE SK-Hep1 cells under normal or glucose deprivation conditions after 5 hours of culture. (F) qRT-PCR analysis of the expression levels of OGDH in wild-type and NFYAv2OE SK-Hep1 cells under normal or glucose deprivation conditions after 5 hours of culture. (G) Western blot analysis of Nfya protein levels in the liver tissue of  $Nfyav1^{+/+}$ ,  $Nfyav1^{+/-}$ , and  $Nfyav1^{-/-}$  mice. (H) Pyruvate tolerance test (PTT) in  $Nfyav1^{+/+}$  (n=5) and  $Nfyav1^{-/-}$  (n=4) mice. A bar graph shows the area under curve. All data are shown as mean and SEM. Unpaired two-tailed t-test was performed. (n.s.) not significant; (\*) P<0.05; (\*\*) P<0.01; (\*\*\*) P<0.001.





**Supplementary Figure 4.** NFYAv2 enhances gluconeogenesis by transcriptional activation of PCK1. (A) qRT-PCR analysis to validate the knockdown of *PCK1* gene expression in NFYAv2OE SK-Hep1 cells under normal or glucose deprivation conditions after 5 hours of culture. Data are shown as mean and SEM. Unpaired two-tailed t-test was performed. (n.s.) not significant; (\*\*\*) P<0.001. (B) Representative fluorescence images of living cells (green) detected with Calcein-AM and dead cells (red) detected with Propidium iodide (PI) in wild-type and NFYAv2OE SK-Hep1 cells knocked down PCK1 gene expression under normal or glucose deprivation conditions after 10 hours culture. (C) The bar graph shows the percentage of dead cells among the more than 600 cells counted

each condition. Data are shown as mean and SEM. Unpaired two-tailed t-test was performed. (n.s.) not significant; (\*) P<0.05. (**D**, **E**) Using RNA-seq data from TCGA, 297 human hepatocellular carcinomas were classified into two groups according to logFC of VIM/CDH1: one group predominantly expressing NFYAv1 ( $1 < \log$ FC: n=74) and another group predominantly expressing NFYAv2 (logFC < -1: n=83). The expression levels (log2 counts) of NFYA (D) and PCK1 (E) in each group were calculated. Unpaired two-tailed t-test was performed. (n.s.) not significant; (\*\*\*) P<0.001.