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Supplemental Information

mTOR Signaling and SREBP Activity

Increase FADS2 Expression

and Can Activate Sapienate Biosynthesis

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Supplementary figures and tables

Figure S1



Figure S1: Correlation of *FADS2* expression with copy number and DNA mutations status of signature genes in TCGA database, Related to **Figure 1** and **Table S4**

(A) Heat map representing the correlation between *FADS2* expression and the average copy number alteration of *PTEN* and *MYC* based on the TCGA data derived from 12,650 patients representing 33 major human cancer type. Positive correlations (p<0.05) are indicated in yellow and negative correlations (p<0.05) are indicated in blue. No significant correlations (p>0.05) are presented in white.

(B) Heat map representing the correlation between *FADS2* and the average of *TP53* and *PIK3CA* mutations. TCGA samples representing 33 major human cancer type were divided by mutation status of *TP53* and *PIK3CA* in wild type and mutated (defined as having any type of mutation) and *FADS2* expression was compared between both groups using a t-test. Correlations (p<0.05) between increased *FADS2* expression and the presence of mutations are indicated in yellow; correlations (p<0.05) between decreased *FADS2* expression and the presence of mutations are indicated in blue. No significant correlations (p>0.05) are presented in white.

ACC: Adrenocortical carcinoma, BLCA: Bladder Urothelial Carcinoma, BRCA: Breast invasive carcinoma, CESC: Cervical squamous cell carcinoma and endocervical adenocarcinoma, CHOL: Cholangiocarcinoma, COAD: Colon adenocarcinoma, DLBC: Lymphoid Neoplasm Diffuse Large B-cell Lymphoma, ESCA: Esophageal carcinoma, GBM: Glioblastoma multiforme, HNSC: Head and Neck squamous cell carcinoma, KICH: Kidney Chromophobe, KIRC: Kidney renal clear cell carcinoma, KIRP: Kidney renal papillary cell carcinoma, LAML: Acute Myeloid Leukemia, LGG: Brain Lower Grade Glioma, LIHC: Liver hepatocellular carcinoma, LUAD: Lung adenocarcinoma, LUSC: Lung squamous cell carcinoma, MESO: Mesothelioma, OV: Ovarian serous cystadenocarcinoma, PAAD: Pancreatic adenocarcinoma, PCPG: Pheochromocytoma and Paraganglioma, PRAD: Prostate adenocarcinoma, READ: Rectum adenocarcinoma, SARC: Sarcoma, SKCM: Skin Cutaneous Melanoma, STAD: Stomach adenocarcinoma, TGCT: Testicular Germ Cell Tumors, THCA: Thyroid carcinoma, THYM: Thymoma, UCEC: Uterine Corpus Endometrial Carcinoma, UCS: Uterine Carcinosarcoma, UVM: Uveal Melanoma





Figure S2: mTOR inhibition does not alter palmitate levels, Related to Figure 2 and Figure 3

(A, B) Relative palmitate level upon treatment with DMSO (A, N=12, B, N=11), rapamycin (A) (20 nM, 72 h, N=12) or Torin1 (B) (40 nM, 72 h, N=11) in WT and Tsc2^{-/-} MEFs. Bar graphs are presented as mean \pm SD from biological independent samples. Statistical testing was performed by a two-sided unpaired Student's t test.

(C, D) Relative palmitate level upon treatment with DMSO (C, N=9, D, N=12), rapamycin (C) (20 nM, 72 h, N=9) or Torin1 (D) (40 nM, 72 h, N=12) in HUH7 cancer cells. Bar graphs are presented as mean ± SD from biological independent samples. Statistical testing was performed by a two-sided unpaired Student's t test.

Figure S3



Figure S3: mTOR inhibition results in elevated polyunsaturated fatty acid levels, Related to Figure 2 and Figure 3

(A-C) Relative levels of linoleic acid (A), docosahexaenoic acid (B) and arachidonic acid (C) in the wildtype (WT) MEFs upon treatment with DMSO or Torin1 (40 nM, 72 h, A, N=3, B-C, N=6). Bar graphs are presented as mean ± SD from biological independent samples. Statistical testing was performed by a two-sided unpaired Student's t test.

(D-F) Relative levels of linoleic acid **(D)**, docosahexaenoic acid **(E)** and arachidonic acid **(F)** in the Tsc2^{-/-} MEFs upon treatment with DMSO or Torin1 (40 nM, 72 h, **D**, N=3, **E-F**, N=6). Bar graphs are presented as mean ± SD from biological independent samples. Statistical testing was performed by a two-sided unpaired Student's t test.

(G-I) Relative levels of linoleic acid (G), docosahexaenoic acid (H) and arachidonic acid (I) in the HUH7 cancer cells upon treatment with DMSO or Torin1 (40 nM, 72 h, G, N=3, H-I, N=6). Bar graphs are presented as mean \pm SD from biological independent samples. Statistical testing was performed by a two-sided unpaired Student's t test.

Figure S4



Figure S4: SREBP modulation does not alter palmitate level and regulates *Fads2* and *Scd1* transcription in hypoxia, Related to **Figure 4**

(A) Relative palmitate level was determined in U87 cells stably expressing ER-mSREBP-1 or ER-mSREBP-2. Cells were treated with 100 nM 4-OHT or solvent for 24h in medium containing 10% FBS (N=3). Bar graphs are presented as mean ± SD from biological independent samples. Statistical testing was performed by a two-sided unpaired Student's t test.

(B, C) Palmitate level upon treatment with DMSO (N=12) or fatostatin (10 μ M, 72 h, B, N=12, C, N=10) in WT and Tsc2^{-/-} MEFs (B) and in HUH7 cancer cells (C). Bar graphs are presented as mean ± SD from biological independent samples. Statistical testing was performed by a two-sided unpaired Student's t test.

(D, E) FadsS2 (D) and Scd1 (E) mRNA expression in WT and Tsc2^{-/-} MEFs cells treated with fatostatin (10 μ M, 48 h, N=3) combined with hypoxia (1% O₂). Relative mRNA levels were determined by qPCR. Bar graphs are presented as mean ± SD from biological independent samples. Statistical testing was performed by a two-sided unpaired Student's t test.

	Gene	Forward (5'-3')	Reverse (5'-3')
	FADS2	GACCACGGCAAGAACTCAAAG	GAGGGTAGGAATCCAGCCATT
	SCD1	TCTCTGCTACACTTGGGAGC	GAGCTTTGTAAGAGCGGTGG
Human	FADS2 (Site A)	TCTGCTGATCGCTGTGGAAACT	TCAGCCCTCCCGCTATGGACTTT
	FADS2 (Site B)	TGGAGGCAAAAGTCCATAGC	GATCCCTGGCTTCCCAGT
	АСТВ	GCCTCGCCTTTGCCGAT	CGCGGCGATATCATCATCC
	RPL19	ACCCCA ATGAGACCA ATG AA	CGCAAAATCCTCATTCTCCT
	Fads2	ATAGTAGCTGATGGCCCAAGC	AGCCCCTTGAGTATGGCAAG
Mouse	Scd1	CGCTCTTTACCCTTTGCTGG	AAGAACTGGAGATCTCTTGGAGC
	Rpl19	CAGGCATATGGGCATAGGGAA	TGCCTTCAGCTTGTGGATGT

Table S5: qPCR primer sequences list, Related to KEY RESOURCES TABLE