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Supplemental Figures

Figure S1. Effects of mTOR inhibitors and SREBP1/2 knockdowns on the induction of *SREBF1, SREBF2*, and *ACACA* transcription by oncogenic PI3K and K-Ras. (**a**) Oncogene and mTORC1-dependent regulation of *SREBF1, SREBF2*, and *ACACA* gene expression. RNA was isolated from MCF10a cells (from **2b**) stably expressing empty vector, PIK3CA^{H1047R},or K-Ras^{G12V} serum starved for 16 h in the presence of vehicle, rapamycin (20 nM), or Torin1 (250 nM) for analysis by qRT-PCR, n=2. (**b**) Effect of SREBP knockdown on *SREBF1, SREBF2*, and *ACACA* gene expression. MCF10a cells from **2c** were transfected with siRNAs targeting SREBP1, SREBP2, or both. RNA was isolated from cells 72 h post-transfection following 20 h starvation, n=3. Representative data are shown as mean ± s.e.m. relative to vector-expressing cells. # P-value < 0.05 compared to vector-expressing cells; * P-value < 0.05 compared to vehicle-treated cells expressing the same oncogene.

Figure S2. mTOR inhibitors decreases the expression of SREBP and its target genes in breast cancer cells. (**a**) mTORC1-dependent *SREBF1* and *SREBF2* expression in breast cancer cells. qRT-PCR analysis of RNA isolated from cells (from **3b**) serum starved for 18 h in the presence of vehicle, rapamycin (20 nM), or Torin1 (250 nM). Data are shown as mean ± s.e.m. relative to vehicle-treated cells. * P-value < 0.05 compared to vehicle-treated cells. (**b**) Immunoblots demonstrating the specificity of SREBP1 and SREBP2 antibodies. MDA-MB-468, MDA-MB-453, and Hs578T cells were transfected with siRNAs targeting SREBP1 and SREBP2 for 72 h. Cytosolic and nuclear fractions

were collected after 18 h serum starvation to detect the cytosolic precursor (P) and the nuclear mature (M) forms of SREBP1 and 2, and the C-terminal cleaved form of SREBP2 (C). (c) Effect of long-term mTOR inhibition on FASN and SCD levels. MDA-MB-453 cells were serum starved for 96 h and rapamycin (20 nM) or Torin1 (250 nM) was added in 24 h increments.

Figure S3. Effect of SREBP1 or SREBP2 knockdown on expression of *SREBF1*, *SREBF2*, and *SCD* in breast cancer cells. Breast cancer cells were transfected with siRNAs targeting SREBP1, 2 or both. RNA was extracted for qRT-PCR analysis 72 h after transfection, following 18 h serum starvation. Representative data are shown as mean \pm s.e.m. relative to vehicle-treated cells, n=2. * P-value < 0.05 compared to cells with control siRNAs.

Figure S4. Effects of SREBP1 and SREBP2 depletion on the proliferation of breast cancer cells in full, lipid-rich serum. MDA-MB-468, MDA-MB-453, and Hs578T cells were transfected with siRNAs targeting SREBP1, SREBP2, or both. Cells cultured in full serum were counted every 24 h, starting 24 h post-transfection (t = 0 h). Data are shown as mean \pm s.e.m., n=3. * P-value < 0.05 compared to cells with control siRNAs at that time point.

Figure S5. Validation of antibodies for use in tissue array dot blots. MDA-MB-468 cells were serum starved for 20 h in the presence of vehicle or Torin1 (250 nM). Dilutions of protein extract from these cells were spotted onto nitrocellulose membranes and immunoblotted for the indicated proteins.









С



MDA-MB-453	
Torin1 (H)	0 24 48 72 96
P-S6K	-
S6K	
FASN	
SCD	8
Actin	

Figure S2





Figure S4

