| TIDFODIASTS | | | |
|-------------|----------------------------------------------------------------------------------------------|---------------------------------------------|-------------------------------------|
| Gene symbol | Gene title | ATF6α(1-373) versus empty fold change | XBP1(S) versus empty fold change |
| Chka | choline kinase alpha | 16.88 | 1.03 |
| Chkb | choline kinase beta | 4.63 | 1.10 |
| Lpin1 | lipin 1 | 4.58 | 3.46 |
| Fasn | fatty acid synthase | 3.92 | 1.02 |
| Lpin3 | lipin 3 | 3.73 | 8.34 |
| Cyb5r1 | cytochrome b5 reductase 1 | 2.78 | 0.98 |
| Crot | carnitine O-octanoyltransferase | 2.03 | -1.31 |
| Acsl3 | acyl-CoA synthetase long-chain family member 3 | 2.02 | 1.65 |
| Cyb5 | cytochrome b-5 | 1.79 | 3.45 |
| Agpat6 | 1-acylglycerol-3-phosphate O-acyltransferase 6 (lysophosphatidic acid acyltransferase, zeta) | 1.63 | 2.11 |
| Ppap2b | phosphatidic acid phosphatase type 2B | 1.50 | -2.40 |
| Elovl4 | elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 4 | 1.45 | -2.40 |
| Pecr | peroxisomal trans-2-enoyl-CoA reductase | -1.72 | 4.47 |
| Acatl | acetyl-CoA acetyltransferase 1 | -2.04 | -1.08 |
| Hadh2 | hydroxyacyl-Coenzyme A dehydrogenase type II | -2.07 | -1.33 |
| Elovl5 | ELOVL family member 5, elongation of long chain fatty acids (yeast) | -2.28 | -1.44 |
| Acsl4 | acyl-CoA synthetase long-chain family member 4 | -2.34 | -1.05 |
| Hadhsc | L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain | -2.62 | 1.00 |
| Fads1 | fatty acid desaturase 1 | -2.64 | 1.07 |
| Sgms1 | Sphingomyelin synthase 1 | -2.85 | 1.15 |
| Ppap2c | phosphatidic acid phosphatase type 2c | -2.95 | -2.06 |
| Acsl5 | acyl-CoA synthetase long-chain family member 5 | -3.14 | -1.14 |
| Slc44a1 | Choline transporter-like protein 1 | -3.19 | -2.22 |
| Elovl6 | ELOVL family member 6, elongation of long chain fatty acids (yeast) | -3.47 | 1.60 |
| Nsmaf | Neutral sphingomyeliase activation associated factor | -3.48 | -1.25 |

Table S1. Comparison of lipid metabolism genes regulated by enforced expression of ATF6α(1-373) and XBP1(S) in NIH-3T3 fibroblasts

Affymetrix microarray analysis revealed that a subset of genes encoding proteins that function in lipid metabolism were upregulated in ATF6 α (1-373)-transduced NIH-3T3 fibroblasts (\geq 2-fold as compared with empty vector controls; *P*<0.05).