

Adipose tissue dysfunction is associated with low levels of the novel Palmitic Acid Hydroxystearic Acids

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Supplemental table 1. Partial correlations controlling for adipocyte cell size

| <i>Gene vs GLUT4</i> | <i>R_p</i> | <i>p-value</i> |
|-----------------------|----------------------|----------------|
| PPAR γ | 0.50 | 0.001 |
| C/EBP α | 0.48 | 0.001 |
| Adiponectin | 0.75 | <0.001 |
| ChREBP | 0.32 | 0.034 |
| ACACA | 0.75 | <0.001 |
| FASN | 0.65 | <0.001 |
| <i>PAHSA vs GLUT4</i> | <i>R_p</i> | <i>p-value</i> |
| Total-PAHSA | 0.67 | 0.024 |
| 5-PAHSA | 0.18 | 0.61 |
| 9-PAHSA | 0.63 | 0.039 |
| 10-PAHSA | 0.54 | 0.087 |
| 12/13-PAHSA | 0.61 | 0.049 |

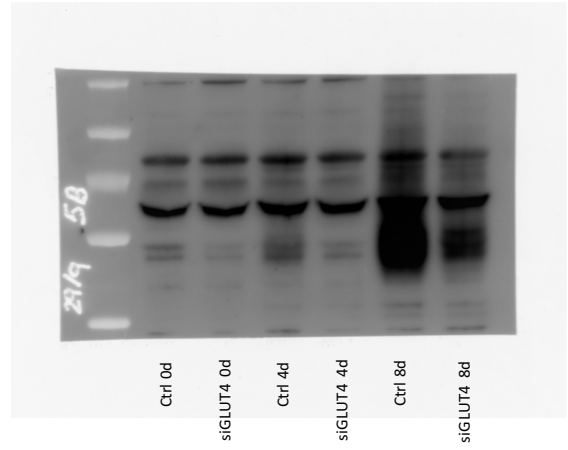
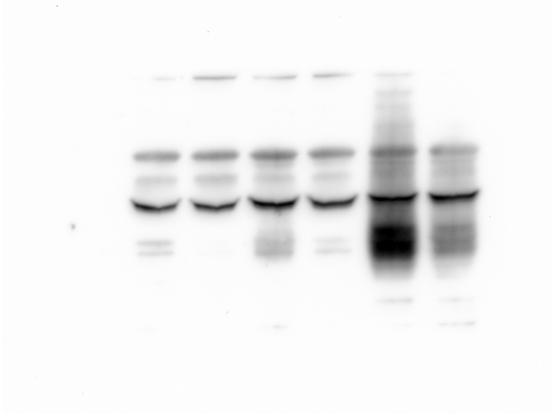
Full length captures

Full length membranes to Figure 3A

Some of the original membranes have been cut to enable blotting of multiple proteins at different molecular size.

The original tiff-files from caputre is full length.

Figure 3A GLUT4 protein



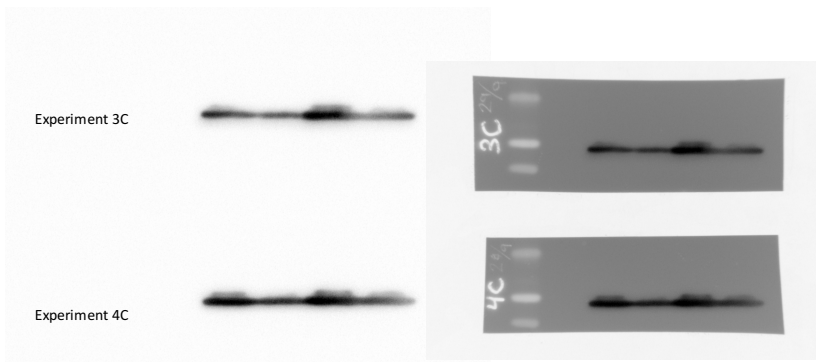
Full length membranes to Figure 3C

FABP4 protein

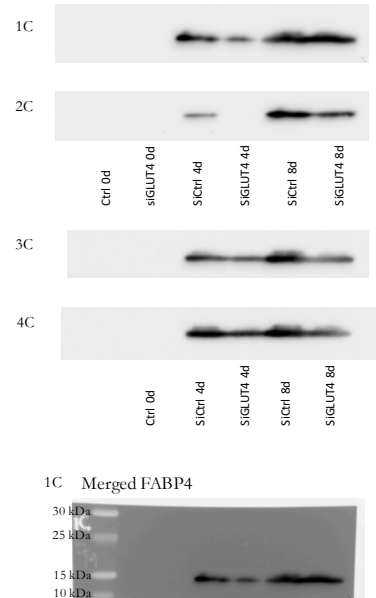
Some of the original membranes have been cut to enable blotting of multiple proteins at different molecular size.

The original tiff-files from caputre is full size.

Two independent experiments have been visualized at the same time -> both are on the original TIFF-file



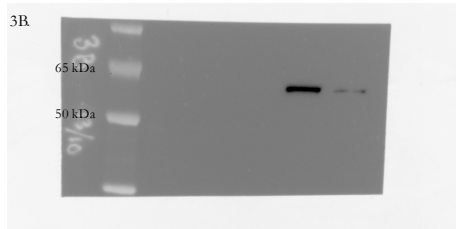
All quantified blots



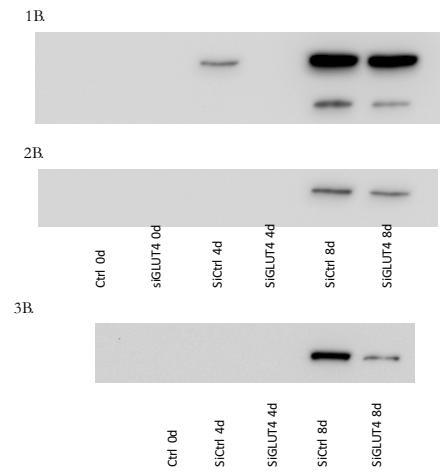
1.C, 2.C, 3C, 4C FABP4 #3544 1:1000 rabbit
NuPage 4-12%, 1mm, MOPS, BioRad transfer 30 min

Full length membranes to Figure 3C
Perilipin protein

Some of the original membranes have been cut to enable blotting of multiple proteins at different molecular size.
The original tiff-files from caputre is full length.



All quantified blots



1.B och 2.B Perilipin #9349s 1:1000 rabbit
NuPage 4-12%, 1mm, MOPS, BioRad transfer 30 min