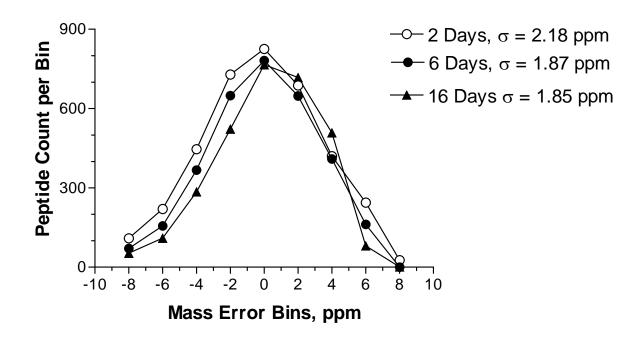
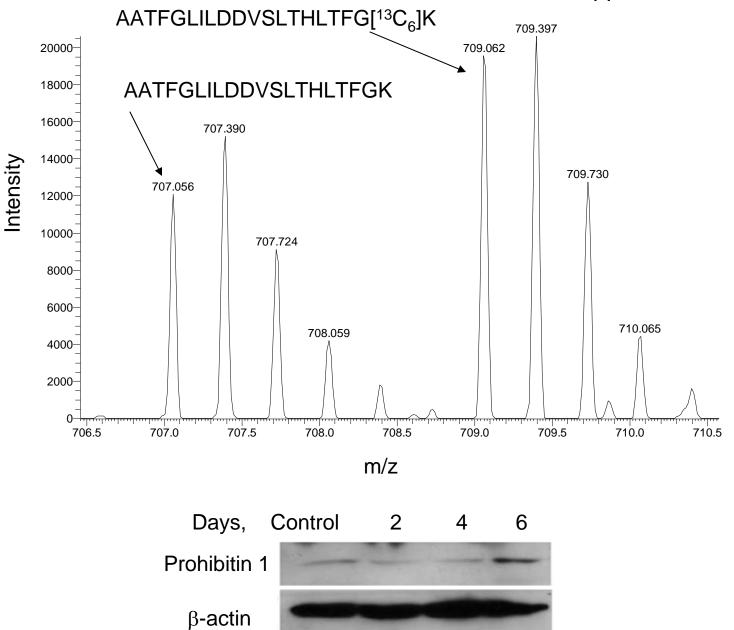
**Supplemental Fig. 1- Distribution of Mass Errors and Estimation of Statistical Mass Accuracy.** At each time point, the calibrated mass errors (ppm) for all the peptides identified from the mitochondrial fractions by Mascot were obtained from the MSQUANT peptide report. Mass errors were binned to a width of 2 ppm and the number of peptides per bin counted. Data shows the Gaussian nature of the distribution of mass deviations and the average dispersion was 1.97± 0.19 ppm.

Supplemental Fig. 2- Hyperglycemia Increased Prohibitin 1. SCs were incubated with 5.5 mM (K0) or 30 mM (K6) glucose for 6 days. (A) Spectra show MS1 scan indicating a 1.6 fold increase for the prohibitin peptide AATFGLILDDVSLTHLTFGK. (B) The expression of prohibitin 1 from a heavy mitochondrial fraction obtained from SCs subjected to hyperglycemia for the indicated time.  $\beta$ -Actin was used as a loading control.

## Supplemental Fig. 1



## Supplemental Fig. S2



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