

# Carboxypeptidase E mediates palmitate-induced $\beta$ -cell ER stress and apoptosis

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Obesity is a principal risk factor for type 2 diabetes, and elevated fatty acids reduce  $\beta$ -cell function and survival. An unbiased proteomic screen was used to identify targets of palmitate in  $\beta$ -cell death. The most significantly altered protein in both human islets and MIN6  $\beta$ -cells treated with palmitate was carboxypeptidase E (CPE). Palmitate reduced CPE protein levels within 2 h, preceding endoplasmic reticulum (ER) stress and cell death, by a mechanism involving CPE translocation to Golgi and lysosomal degradation. Palmitate metabolism and  $\text{Ca}^{2+}$  flux were also required for CPE proteolysis and  $\beta$ -cell death. Chronic palmitate exposure increased the ratio of proinsulin to insulin. CPE null islets had increased apoptosis *in vivo* and *in vitro*. Reducing CPE by  $\approx 30\%$  using shRNA also increased ER stress and apoptosis. Conversely, overexpression of CPE partially rescued  $\beta$ -cells from palmitate-induced ER stress and apoptosis. Thus, carboxypeptidase E degradation contributes to palmitate-induced  $\beta$ -cell ER stress and apoptosis. CPE is a major link between hyperlipidemia and  $\beta$ -cell death pathways in diabetes.

2D difference gel electrophoresis proteomics | free fatty acids | hyperproinsulinemia | mechanisms of  $\beta$ -cell lipotoxicity | type 2 diabetes

There is a strong association between type 2 diabetes and obesity. High levels of circulating lipids, including free fatty acids, are a prominent clinical feature of type 2 diabetes and represent an important risk factor for this disease (1, 2). But exactly how elevated lipids might lead to diabetes remains unresolved. Fatty acids increase basal insulin secretion (3) and the relative levels of circulating proinsulin (4). Chronic exposure to the free fatty acid palmitate has been shown to impair glucose-stimulated insulin release (i.e., lipotoxicity) (5–10).  $\beta$ -Cell apoptosis can be initiated by high levels of palmitate (6, 7, 11–14), which may account in part for alterations in insulin secretory function (13). A number of studies have established palmitate targets in the  $\beta$ -cell, including lipid metabolism (15, 16), mitochondrial function (17–23), and prosurvival transcription factors such as Pdx1 (24, 25). Recently, a role for endoplasmic reticulum (ER) stress in lipotoxicity has been demonstrated in multiple cell types, including  $\beta$ -cells (11, 26, 27). The effects of palmitate on  $\beta$ -cell survival are likely mediated by a number of mechanisms.

In the present study, we conducted unbiased proteomic screens using human islets and MIN6  $\beta$ -cells to elucidate targets of palmitate. Carboxypeptidase E (CPE) was the most significantly changed protein in both screens. Mice lacking CPE develop hyperproinsulinemia and hyperglycemia (28), but the involvement of this protein in  $\beta$ -cell apoptosis has not been reported. Palmitate caused the rapid intracellular redistribution and degradation of CPE via mechanisms that required palmitate metabolism,  $\text{K}_{\text{ATP}}$  channel closure,  $\text{Ca}^{2+}$  influx, and protease activity. We further showed that CPE levels control  $\beta$ -cell ER stress and apoptosis. Thus, CPE is a critical link between saturated free fatty acids, a major type 2 diabetes risk factor, and  $\beta$ -cell dysfunction.

## Results

**Analysis of the  $\beta$ -Cell Proteome During Palmitate-Induced Apoptosis.** Palmitate, but not oleate, induced dose-dependent apoptosis, illustrated by DNA laddering [supporting information (SI) Fig. S1A]. A high-throughput, real-time assay of  $\beta$ -cell death demonstrated that palmitate-induced propidium iodide incorporation was significant at  $\approx 12$  h. Neither the rate nor the amplitude of palmitate-induced death differed between 5 mM glucose and 25 mM glucose (Fig. S1B). Palmitate did not require high glucose to induce ER stress in MIN6 cells (Fig. S1C) or human islets (Fig. S1D; see also Fig. S4B). We have shown previously that palmitate could induce apoptosis in mouse islets cultured in 5 mM glucose (29). To better understand the mechanism of  $\beta$ -cell lipotoxicity, we undertook proteomic analyses of MIN6 cells and human pancreatic islets exposed to palmitate. Strikingly, the gel features most reduced by palmitate in both MIN6 cells and human islets were identified as CPE (Fig. 1 and Figs. S2 and S3). Significant changes were also found in protein spots identified as key metabolic enzymes (NADH dehydrogenase, phosphoglycerate mutase, ATP citrate lyase), and components of the proteasome (UCHL1, proteasome subunit  $\alpha 3$ ) (Tables S1 and S2). Because of its known role in insulin processing and diabetes, we focused primarily on CPE in this study.

**Palmitate Causes Rapid CPE Degradation.** The reduced protein spots identified as CPE in the 2-D gels were the approximate size of full-length CPE, rather than fragments. To exclude the possibility that the observed decrease in relative abundance of the gel features was due solely to posttranslational modifications that alter the isoelectric point, we performed Western blot analysis. These experiments demonstrated that palmitate significantly and dose dependently decreased total CPE protein in both MIN6 cells and human islets (Fig. 2A and Fig. S4A and B), with the strongest inhibition seen at a 6:1 palmitate-to-albumin ratio, a dose similar to what would be expected in obese or diabetic patients (30, 31). Similar results were seen when GAPDH or tubulin were used as loading controls (not shown). A loss of CPE was also detected with antibodies to the C-terminal and N-terminal regions (Fig. S4C), suggesting loss of the entire protein. A related enzyme, carboxypeptidase D, was not reduced by palmitate in low glucose (Fig. S4D and F). Prohormone convertase 1/3 and prohormone convertase 2 were

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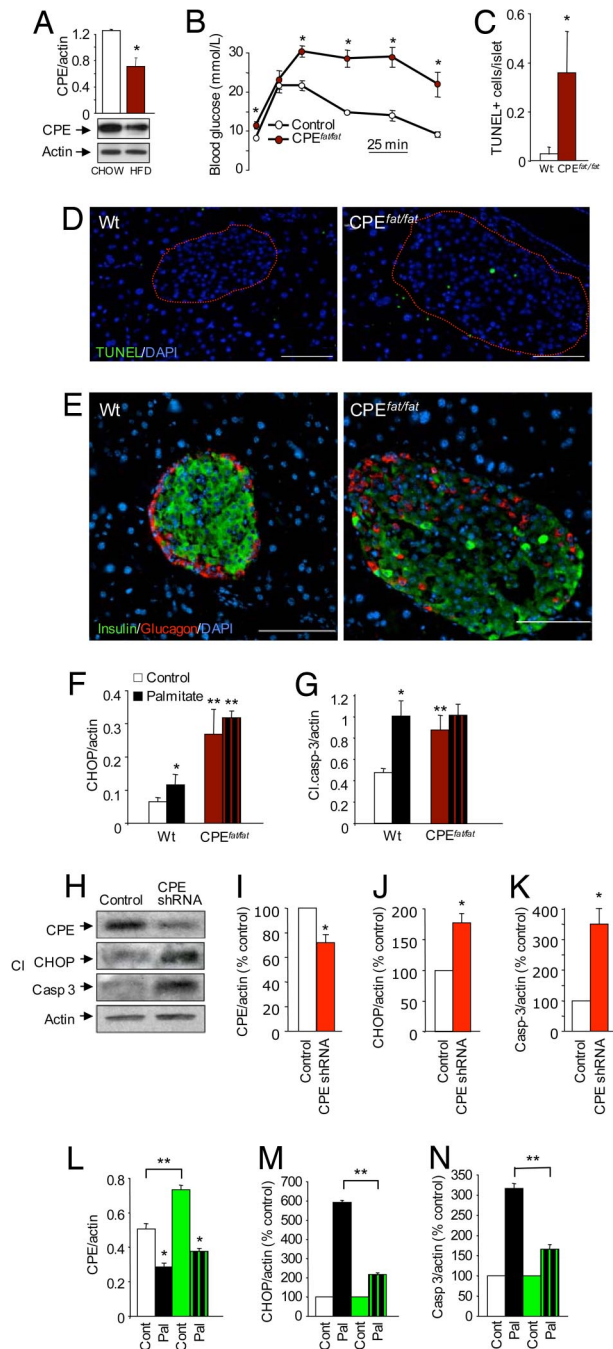
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**Fig. 4.** *In vivo* and *in vitro* role of CPE in  $\beta$ -cell death. The ratio of CPE-processed mature insulin to total insulin was dramatically decreased in islets from high-fat-fed mice and in MIN6 cells treated with palmitate. This decrease in mature insulin was comparable to that seen in mice lacking CPE (Fig. S6). (A) Reduced CPE protein expression in isolated islets from C57BL/6J mice fed high-fat diet (40% calories from fat) for 6 months ( $n = 3$ ). (B) Homozygous null  $CPE^{fat/fat}$  mice were significantly heavier ( $28 \pm 2$  g vs.  $41 \pm 3$  g) and exhibited significantly impaired i.p. glucose tolerance compared with littermate controls ( $n = 4$ ). (C and D) Increased TUNEL staining in islets of pancreatic sections of  $CPE^{fat/fat}$  mice compared with wild-type controls ( $n = 3$ ). (Scale bar,  $100 \mu\text{m}$ .) (E) Representative insulin (green), glucagon (red), and DAPI (blue) staining in the islets of  $CPE^{fat/fat}$  mice and wild-type controls. Islets from mutant mice had weaker and more heterogeneous insulin staining, as well as disrupted architecture. (F and G) CHOP and cleaved caspase-3 protein were quantified in isolated islets from control and  $CPE^{fat/fat}$  mice treated as indicated for 24 h in 20 mM glucose ( $n = 3$ ). (H–K) Significant small hairpin RNA-mediated reduction in CPE protein increases CHOP expression and caspase-3 cleavage, compared with MIN6 cells transfected with a scrambled shRNA control ( $n = 3$ ). (L–N) MIN6 cells were transfected with a CPE plasmid (green bars) with palmitate for

translocation to lysosomes. However, both E64D and another protease inhibitor (ALLM; data not shown) caused ER stress on their own, likely because of a backlog of undegraded proteins, making interpretation of their effects on palmitate-induced  $\beta$ -cell death difficult. An inhibitor of UCHL1, a component of the ubiquitin system and the 2-D gel feature most increased by palmitate in the proteomic analysis, prevented the loss of CPE in response to palmitate in high glucose, but not low glucose (Fig. S7B). In our previous study, ALLM and specific deletion of calpain-10 partially reduced apoptosis in primary mouse islets in long-term cultures containing palmitate and 5 mM glucose (29). However, the link between the calpain-10 pathway and the ER stress-associated cascade that CPE participates in remains unclear.

Our results demonstrated that palmitate, but not bromopalmitate, increased the proinsulin-to-insulin ratio at physiological glucose concentrations. This indicates another important functional consequence of CPE degradation. An elevated proinsulin-to-insulin ratio is a well established clinical finding in diabetes and it has also been suggested to play a possible role in prediabetic states (35, 36). Our results provide a plausible mechanism for this defect. While our work demonstrates effects of FFAs on CPE, a previous study found reduced PC2 and PC1/3 posttranslational processing and increased proinsulin-to-insulin ratio in MIN6 cells treated with FFA for 7 days (4). Interestingly, CPE is required for the processing of prohormone convertases, and hyperproinsulinemic mice lacking CPE have reduced prohormone convertase (PC) 1/3 and 2 activity (55). CPE also reduces the ability of prohormone convertase products to inhibit PC1/3 and 2 (56). Thus, CPE may regulate PC1/3 and PC2 at multiple levels. Because we did not observe significant effects of palmitate on PC1/3 or PC2 at 24 h, it is likely that the loss of CPE precedes the loss of the prohormone convertases. Thus, palmitate may target multiple insulin-processing enzymes, perhaps CPE first, ultimately leading to a relative increase in the secretion of proinsulin, which has only 10% of the biological activity of insulin (57).

In conclusion, it was established in this study that *in vitro* treatment of both MIN6 cells and human islets with palmitate and *in vivo* exposure to a high-fat diet led to a reduction in CPE protein. Previously, CPE has been considered to be a “housekeeping” enzyme; here we demonstrate that CPE can be degraded in response to an extracellular signal in  $\beta$ -cells. Both loss-of-function and gain-of-function approaches suggested that CPE positively controls  $\beta$ -cell survival, via effects on ER stress. We propose a model for  $\beta$ -cell lipotoxicity, enabled by our unbiased proteomic screens (Fig. S8). Together, these results provide evidence that CPE is a key link between hyperlipidemia/FFAs, insulin processing, and  $\beta$ -cell apoptosis pathways in type 2 diabetes.

## Experimental Procedures

**Reagents and Animals.** A detailed list of reagents can be found in the *SI Text*.  $CPE^{fat/fat}$  mice on the C57BL/6J background were from Jackson Laboratories. Intraperitoneal glucose tolerance tests (IPGTT; 2 g glucose/kg body weight) were performed on littermate males after a 12-h fast. In some studies, male C57BL/6J mice were fed a high-fat diet (40% of calories from fat, TD88137; Harlan Teklad) for 6 months and compared to littermates fed normal chow. Baseline phenotypes of these mice are detailed in the Supplement. All studies were approved by the University of British Columbia Animal Care Committee.

**Cell Culture.** Our human islet, mouse islet, and MIN6 cell culture methods have been described (25, 29, 58) and are detailed in the *SI Text*. Hormone release was assessed by static incubation or perfusion (25). Insulin and proinsulin were assayed with Linco RIA or ELISA, respectively. Immunofluorescent staining is described in *SI Text*.

24 h at 25 mM glucose ( $n = 3$ ). The asterisk denotes significance between palmitate and control; the double asterisk denotes significance between vectors within the same treatment.

**Cell Death and Apoptosis Assays.** The incorporation of propidium iodide was monitored in the incubated chamber of a KineticScan Reader (Cellomics). Propidium iodide fluoresces brightly once it passes through the compromised plasma membrane of dying cells. Additional details on this assay and the TUNEL assay are outlined in the *SI Text*.

**Immunoblot and Proteomics.** We used standard methods for immunoblots, and our approach to 2-D-DIGE proteomics of human islets has been described (see *SI Text* for expanded details). Briefly, lysates from control and palmitate-treated cells were labeled with Cy dyes and analyzed in the same 2-D gel. Individual "spots" were quantified and considered significantly different if they were >2 standard deviations outside of the normalized distribution of gel feature intensities. Significantly different spots were sequenced by using mass spectroscopy.

**shRNA-Mediated CPE Knockdown and CPE Overexpression.** CPE levels were reduced in MIN6 cells by using a vector expressing shRNA sequences designed against CPE (see *SI Text*) transfected with Lipofectamine 2000 (Invitrogen). GFP-positive cells were sorted by fluorescence-activated cell sorting (BD FACS Vantage

SE/DIVA). CPE plasmids, one under the control of the CMV promoter, and another under the control of the rat insulin promoter, were kindly supplied by Dr. L. D. Fricker. CPE DNA (1  $\mu$ g) or empty vector, was transfected into MIN6 cells by using Lipofectamine 2000 (Invitrogen).

**Statistics.** Statistical analysis was performed with SigmaStat (Systat). ANOVA (Student–Newman–Keuls) or t test was used where appropriate.  $P < 0.05$  was considered significant.

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