The Role of TRAPy/SSR3 in Preproinsulin Translocation Into the Endoplasmic Reticulum

## Xu et al., Supplement

**Supplemental Figure S1:** Preproinsulin synthesized in TRAPγ/SSR3-KO 293T cells is not oxidized, whereas proinsulin, which enters the ER lumen, is oxidized. Control 293T cells or TRAPγ/SSR3-KO 293T cells were transfected to express recombinant WT human preproinsulin. At 48 h post-transfection, the cells were pulse-labeled, lysed, immunoprecipitated with anti-insulin, and the immunoprecipitates analyzed by SDS-PAGE under reducing conditions (DTT "+") or nonreducing conditions (DTT "-"), followed by phosphorimaging.

Supplemental Figure S2: TRAP $\gamma$ /SSR3 re-expression rescues proinsulin protein level in TRAP $\gamma$ /SSR3-KO 293T cells. A. TRAP $\gamma$ /SSR3-KO 293T cells were transfected to express WT human preproinsulin plus plasmid encoding either empty vector ("EV") or each individual subunit of the TRAP/SSR complex as indicated at the top of the Figure. At 48 h post-transfection, the cells were lysed and analyzed by reducing SDS-PAGE and immunoblotting with the antibodies indicated. **B.** Quantitation (mean  $\pm$  S.D.) of proinsulin and TRAP/SSR subunit protein levels from three independent experiments identical to that shown in panel A; \*p < 0.05.

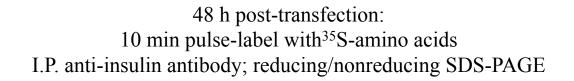
Supplemental Figure S3. Effect of TRAP $\gamma$ /SSR3 deficiency on the mRNA levels encoding *INS* and other TRAP/SSR subunits. INS1E cells were transfected with scrambled oligo ("Ctrl") or TRAP $\gamma$ /SSR3-targeted siRNA for knockdown ("SSR3i"). A. At 72 h post-transfection, total insulin (*Ins1* + *Ins2*) mRNA levels were analyzed by rt-qPCR and quantitation (mean ± S.D.) is shown from five independent experiments; \*p < 0.05. B. At 72 h post-transfection, cells like those in panel A were preincubated in RPMI containing 2.5 mM glucose for 1.5 h followed by a 4 h incubation in 2.5, 11.1, or 25 mM glucose, respectively, as in Figure 4. Insulin (*Ins1* + *Ins2*) and TRAP/SSR complex subunit mRNA levels were analyzed by rt-qPCR and quantitation (mean ± S.D.) is shown from three independent experiments; no significant acute glucose-dependent changes were seen for any mRNA.

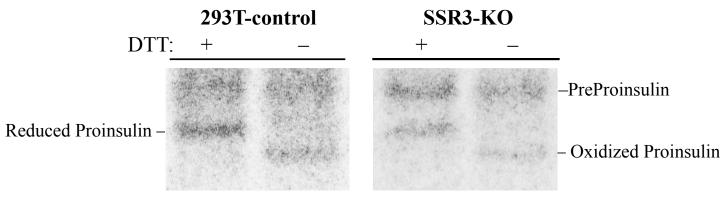
Supplemental Figure S4. TRAP $\gamma$ /SSR3 protein expression in the INS832/13  $\beta$ -cell line. Immunofluorescence with anti-TRAP $\gamma$ /SSR3 (red) stains all INS832/13 cells. Nuclei were counter-stained with DAPI (blue); scale bar is indicated.

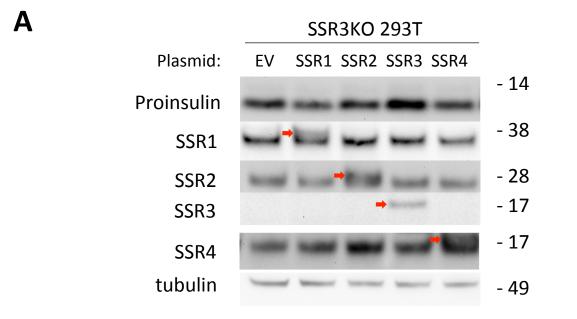
Supplemental Figure S5. Expression of TRAP $\gamma$ /SSR3 restores proinsulin levels in SSR3-KD  $\beta$ -cells, but not in SSR1-KD  $\beta$ -cells. INS832/13 cells were transfected with scrambled oligo ("Ctrl") or TRAP $\alpha$ /SSR1-targeted siRNA ("SSR1-KD") or TRAP $\gamma$ /SSR3-targeted siRNA ("SSR3-KD"). At 24 h post-transfection, cells were re-plated and transfected with empty vector ("–") or plasmid encoding flagtagged TRAP $\gamma$ /SSR3 ("+"). After a further 48 h, cells were lysed and analyzed by reducing SDS-PAGE and immunblotting with the indicated antibodies.

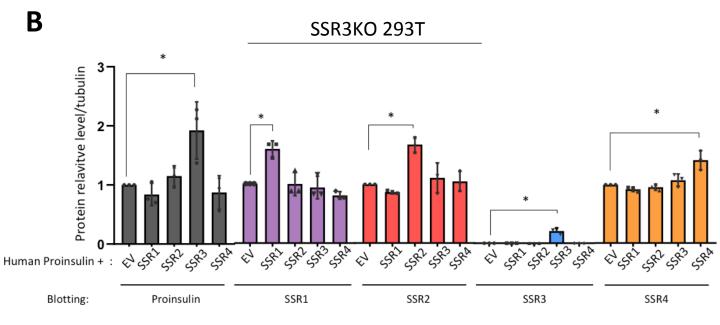
**Supplemental Figure S6. Overexpression of** TRAPα/SSR1 **does not increase proinsulin levels in otherwise normal (control) INS-832/13 cells.** INS-832/13 cells were infected with a replication-deficient adenoviral vector driving the expression of human TRAPα/SSR1 at 50 PFU/cell, as indicated. After 48 hours post-infection, cell lysates were subjected to SDS-PAGE and immunoblotted with anti-proinsulin, anti-TRAPα/SSR1, and anti-tubulin (a loading control).

Supplemental Figure S7. Effects of high glucose stimulation on proinsulin and TRAP $\gamma$ /SSR3 in human islets. Quantitation (mean ± S.D.) of proinsulin and TRAP $\gamma$ /SSR3 protein levels from three technical replicates of each condition in human islets including those shown in Figure 6G. CHX = cycloheximide (100 µg/mL) during the 2 h high-glucose exposure; \* p < 0.05.



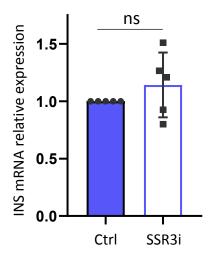


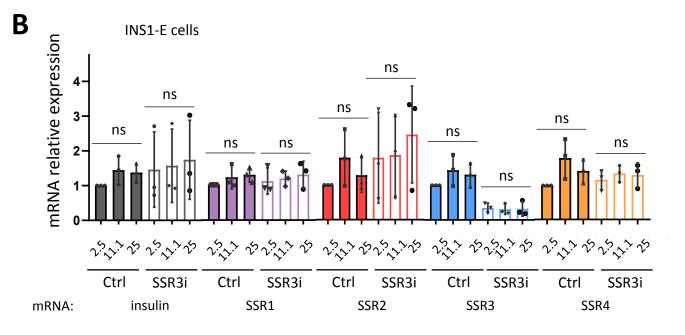




INS1-E cells

Α



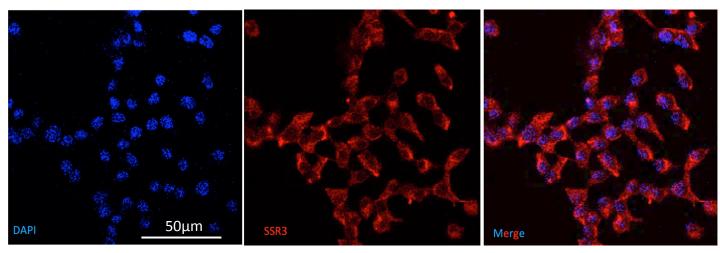


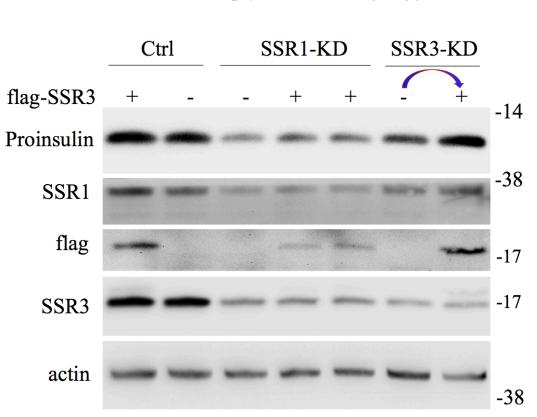
# INS-832/13 cells

### DAPI

Endogenous SSR3

Merge





#### INS832/13 cells (SSR1-KD, SSR3-KD, or Control) Transfected with empty vector or flag-tagged SSR3

