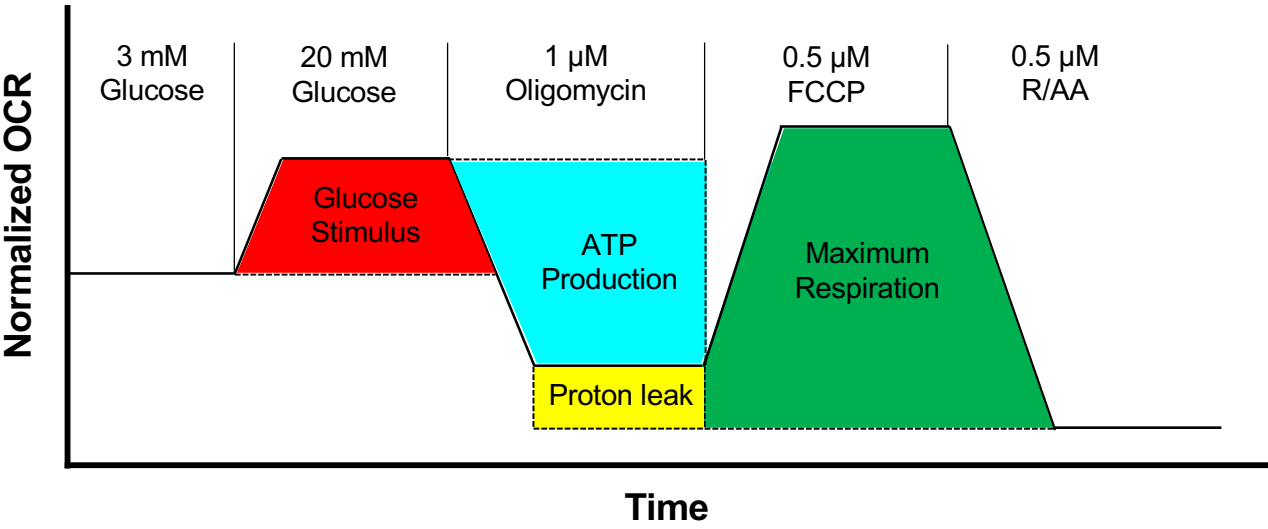
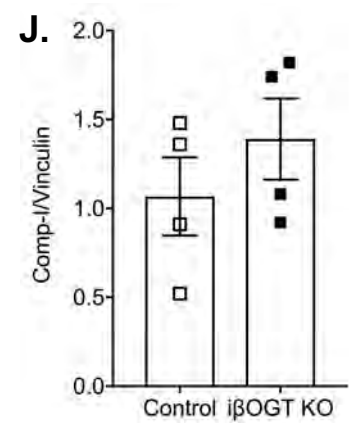
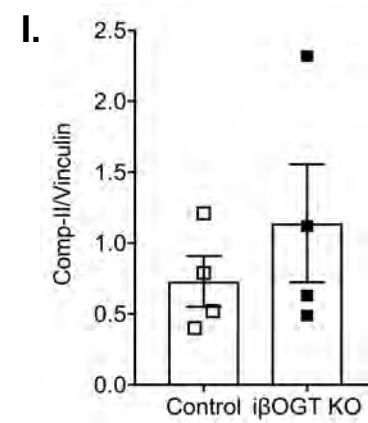
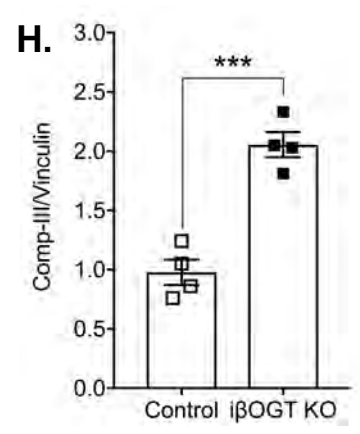
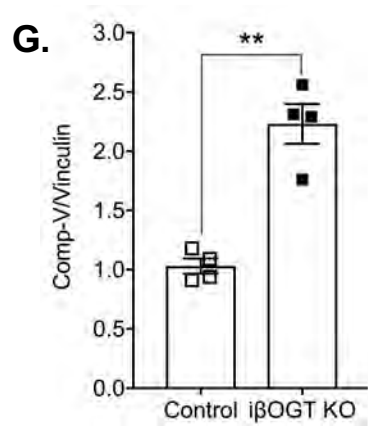
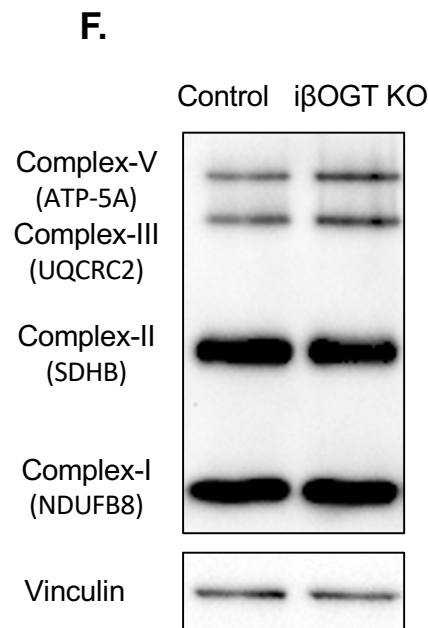
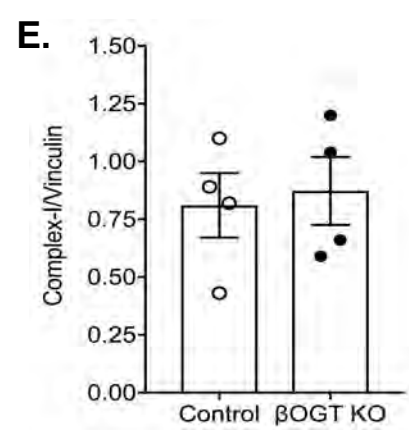
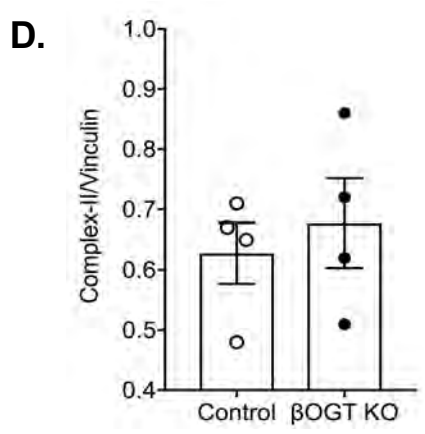
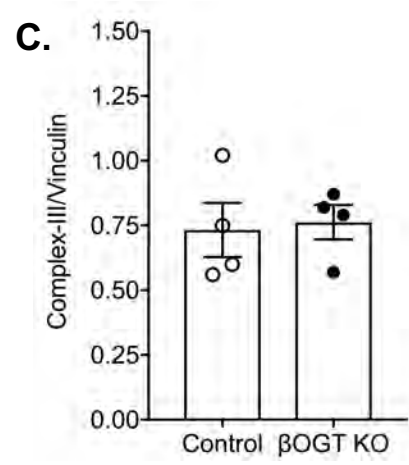
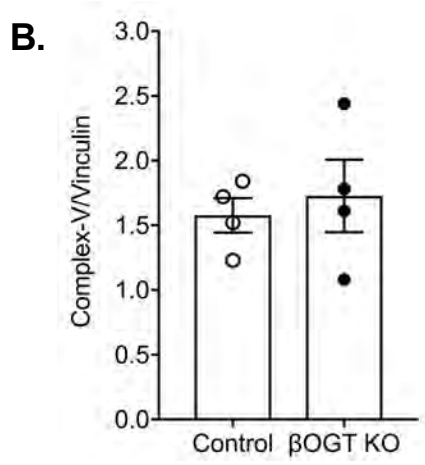
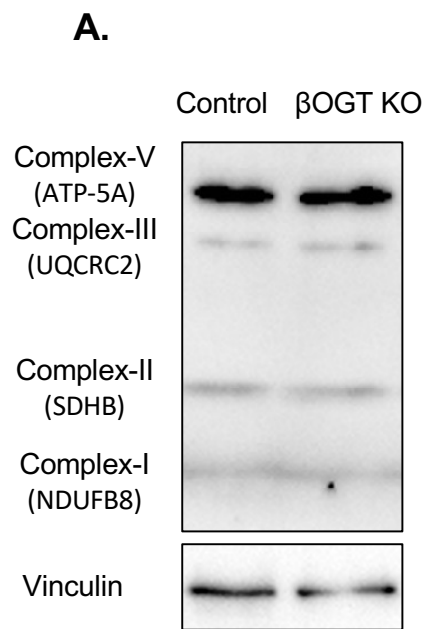


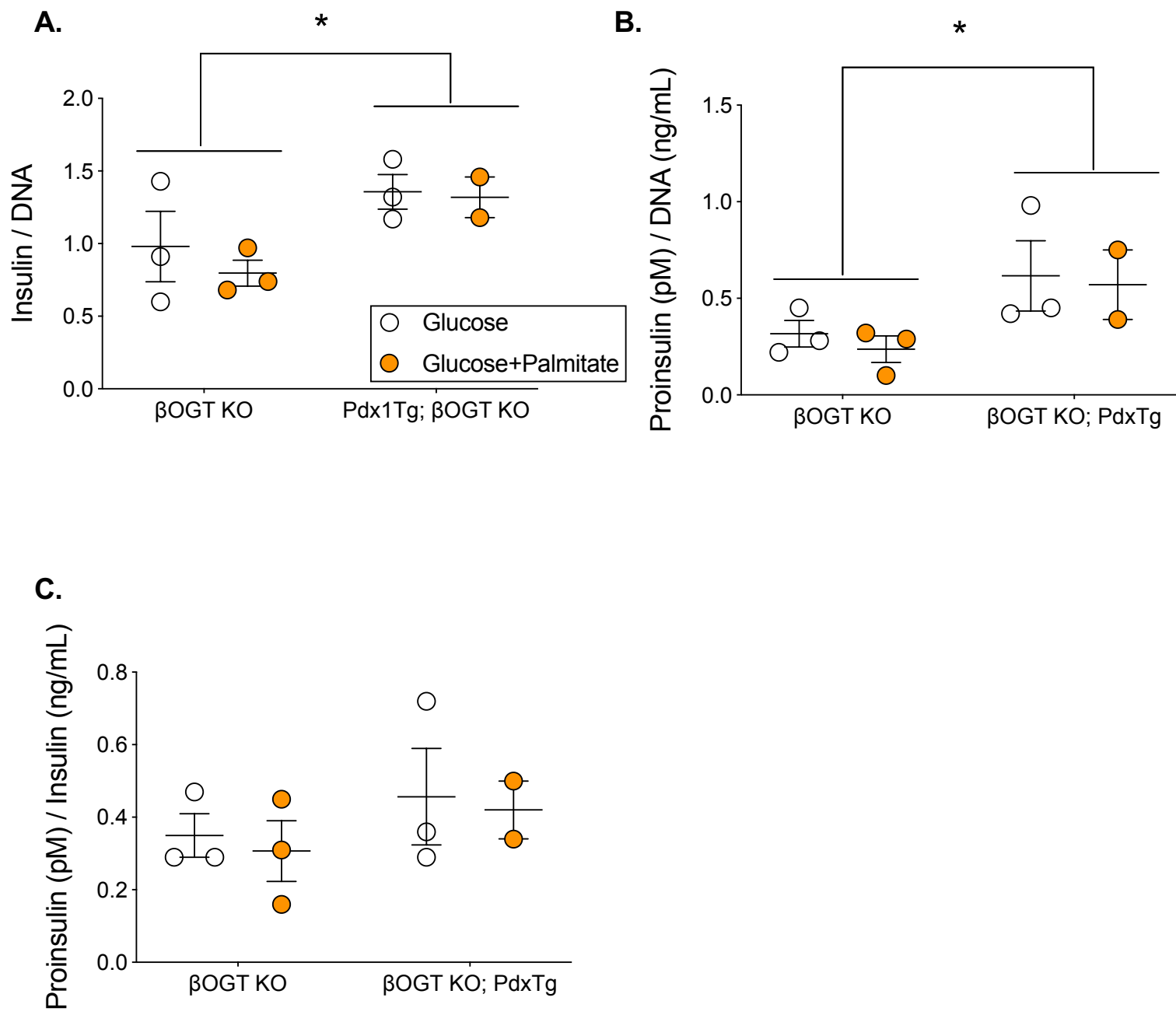
Supplementary Figure: 1



Supplementary Figure: 2



Supplementary Figure: 3



Supplementary Figure 1

(A) The schematic diagram for analyzing glucose stimulation, ATP production, proton leak and maximum respiration, spare capacitance, basal respiration.

Supplementary Figure 2

(A-J). Western blot analysis for Complex-V, IV,III and II between islet lysates from β OGTKO (A) i β OGTKO (F) and respective littermate controls. Quantification of the blots are shown in B-E for β OGTKO and G-J for i β OGTKO, n=4 and *, p-value ≤ 0.05 vs. β OGTKO and control.

Supplementary Figure 3

(A) Islet insulin content, normalized to DNA, is shown for both post-GSIS (glucose) and post-FASIS (glucose + palmitate) treated islets from β OGTKO mice (n=3 glucose, 3 glucose + palmitate) and Pdx1Tg; β OGTKO (n=3, 2). Data from the same islets was used for islet proinsulin content (B) and proinsulin/insulin ratio (C) as a measure of insulin processing capacity.