



Supplemental_Fig_S23: Differential expression analysis at single cell resolution identifies expression changes in Type 2 diabetic and non-diabetic exocrine cell types.

Additional scatter plots depicting the \log_2 fold change of gene expression observed in Type 2 diabetic versus non-diabetic bulk intact islet transcriptomes on the y-axis and the respective \log_2 fold change values of gene expression observed in those of Type 2 diabetic versus non-diabetic single cells. Points highlighted in black represent genes that were detected to be significantly differentially expressed (FDR <0.05) in acinar, stellate, or ductal exocrine cells at single cell resolution. In all plots, points highlighted in grey were not significantly differentially expressed (FDR > 0.05). No significant changes in gene expression were found between the Type 2 diabetic and non-diabetic PP/gamma cell transcriptomes. γ = PP/gamma, FC = fold change, T2D = Type 2 diabetic, ND = non-diabetic.