#### **Supplemental Figure Legends**

**Supplemental Figure 1:** (A) Maximum tumor diameter measured at the time of IRE or sham placebo surgery. PBS/Sham n = 13, IRE n = 14, β-glucan/Sham n = 13, and β-glucan + IRE n = 17, respectively. (B) Quantified mRNA expression of TNF- $\alpha$ , IL-1 $\beta$ , iNOS, IL-12p35, TGF- $\beta$ , IL-10, HK-1, and CCL2 in CD11b+ cells measured by RT-PCR. PBS n=5, IRE n=5, β-glucan n=5, β-glucan + IRE n=5, respectively. Data are presented as mean ± SEM. Significance: ns= not significant; \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001.

**Supplemental Figure 2:** (A) Myeloid cell gating strategy. (B) Frequency of CD45+ leucocytes, CD11b+ meloid cells, CD11b+F4/80+ macrophages, CD11b+Ly6G-Ly6C+ monocytes, and CD11b+Ly6G+Ly6C-neutrophils within KPC tumors treated by PBS,  $\beta$ -glucan, IRE, or  $\beta$ -glucan + IRE. n = 5 per group. Data are presented as mean  $\pm$  SEM. Significance: ns= not significant; \*\*p <0.01.

**Supplemental Figure 3:** (A) Representative flow countour plots of Ly6C+TNF- $\alpha$ + cells late in disease progression from KPC tumors within PBS, β-glucan, IRE, and β-glucan + IRE treated mice. (B) Summarized CD11b+Ly6C+TNF- $\alpha$ + percent and MFI. PBS n = 4, IRE n = 4, β-glucan n = 4, β-glucan + IRE n = 5. (C) Representative flow countour plots of CD11b+F4/80+TNF- $\alpha$ + cells late in disease progression in KPC tumors from PBS, β-glucan, IRE, and β-glucan + IRE treated mice. (D) Summarized CD11b+F4/80+TNF- $\alpha$ + percent and MFI. PBS n = 4, IRE n = 4, β-glucan n = 5, β-glucan + IRE n = 5 Data are presented as mean ± SEM. Significance: ns= not significant

**Supplemental Figure 4:** (A) Quantified frequency of CD69, TIM-3, and LAG-3 expression on CD8+ T cells within PBS, β-glucan, IRE, or β-glucan + IRE treated KPC tumors. (B) Quantified frequency of PD-1, Granzyme B, CD69, TIM-3, and LAG-4 expression on CD4+ T cells within each group. PBS n = 5, β-glucan n = 5, IRE n = 4, and β-glucan + IRE n = 5, respectively. Data are presented as mean  $\pm$  SEM. Significance: ns= not significant; \*p < 0.05, \*\*p < 0.01.

**Supplemental Figure 5:** (A) Representative flow contour plots of CD11b+TNF- $\alpha$ + cells from the pancreas of WT mice fed PBS or increasing doses of oral β-glucan (0.5mg, 1mg, 2mg) daily for one week. (B) Quantified percent CD11b+TNF- $\alpha$ + cells are shown, PBS n =2, 0.5mg β-glucan n = 5, 1mg β-glucan n = 5, 2mg β-glucan n = 5, respectively. Data are presented as mean ± SEM. Significance: ns= not significant; \*\*p <0.01, \*\*\*\*p <0.001, \*\*\*\*p <0.0001.

**Supplemental Figure 6:** (A) WT mice were treated with PBS, one IP β-glucan injection (1mg/mouse), or oral β-glucan (1mg/mouse/day) for 1 week. The mice were then euthanized, pancreases harvested, processed to single cell suspension, and analyzed by flow cytometry. Representative flow contour plots of CD11b+F4/80+TNF- $\alpha$ + expressing cells. (B) Quantified percent and MFI of F4/80+TNF- $\alpha$ + cells are shown. (C) Representative flow contour plots comparing CD11b+Ly6C+TNF- $\alpha$ + expressing cells. (D) Quantified percent and MFI of Ly6C+TNF- $\alpha$ + cells are shown. (E) Representative flow contour plots comparing CD11b+Ly6G+TNF- $\alpha$ + expressing cells. (F) Quantified percent and MFI of Ly6G+TNF- $\alpha$ + cells are shown. (G) Representative flow contour plots comparing CD11b+CD11c+TNF- $\alpha$  expression. (H) Quantified percent and MFI of CD11c+TNF- $\alpha$ + cells are shown. PBS n = 4, IP β-glucan n = 5, and oral β-glucan n = 5, respectively. Data are presented as mean ± SEM. Significance: ns= not significant; \*p < 0.05, \*\*p < 0.01, \*\*\*p<0.001, \*\*\*\*p<0.0001.

**Supplemental Figure 7:** (A) Heatmap scaled expression of cell surface markers within each cell cluster as determined by CyTOF workflow analysis.













