Supplementary Figure 2



Supplementary Figure 2. APOE is highly expressed by tumor-associated macrophages. (A) Dot plot of lineage markers used to define identified cell populations in orthotopic KPC tumors (n=2). Color of the dot represents average expression. Size of the dot represents expression frequency. (B) Co-immunofluorescence staining on normal mouse pancreas (N Panc) and orthotopic KPC tumor for APOE (green), F4/80 (red) and DAPI (blue). White arrowhead denotes APOE and F4/80 colocalization. White box represents higher magnification. Scale bars, 100 µm. (C) Violin plot of normalized ApoE expression in mouse normal pancreas (grey) and tumor (navy) samples. (D) Bright field images of bone marrow cells cultured with either HPNE (top) or 7940b (bottom) conditioned media for 5 days. Scale bar, 400 µm. Inset represents higher magnification of boxed area. (E) Experimental design schematic. (F) qRT-PCR analysis of *Apoe* and (G) *Tnfa* mRNA levels relative to *Cyclophilin A* housekeeping in untreated macrophages (control), and macrophages treated with either HPNE or 7940b conditioned media for 24 hours. Statistical significance was determined using one-way ANOVA with Tukey's test for multiple correction. (H) iKras* experimental design. (I) Immunohistochemical staining for APOE in mouse normal pancreas, pancreatic tissue after 3 weeks of oncogenic *Kras* expression (3W ON), and extinguishment of oncogenic *Kras* expression for 3 days (3d OFF) and 1 week (1W OFF). Scale bars, 50 µm. (J) Quantitation of positive APOE staining as percent area in a 40x field. 5 images per mouse were averaged. Control (n=2), 3W ON (n=3), 3d OFF (n=3), and 1W OFF (n=3). W=week. d=day. Statistical significance was determined using one-way ANOVA with Tukey's test for multiple correction.