## Supplementary Figure 1









**Supplementary Figure 1.** TLS composition and immune features of human PDA. **A**, CD8 T cell frequency and number was determined by flow cytometry. Green circles indicate TLS<sup>+</sup> tumors. Grey box (left panel) demarcates T cell frequency range in blood that does not directly correlate with T cell frequency in tumors. **B**, Example of a preinvasive intraepithelial lesion in Nadj pancreas obtained from a resected patient sample. **C**, Frequency of TLS in Nadj pancreas, PNET and PDA specimens. **D**, CD20 staining of Nadj pancreas, PNET, spleen and PDA demarcate TLS. Inset, rare CD20<sup>+</sup> B cells in Nadj pancreas and spleen. Arrows, TLS. Scale bar represents 100  $\mu$ m. **E**, Cellular composition of lymphoid follicles in spleen and tumoral TLS. CA, central arteriole. Scale bar represents 25  $\mu$ m. **F**, Quantification of IHC staining from Supplementary Figure 1E and was determined by digital analysis of scanned slides. Spl, spleen. **G**, Example of a TLS associated with cytokeratin (CK)<sup>+</sup> tumor cells and surrounded by  $\alpha$ SMA<sup>+</sup> fibroblasts. Scale bar represents 25  $\mu$ m.

#### **Supplementary Figure 2**



**Supplementary Figure 2.** Gating strategy for analyzing CD244 (2B4) and Tim3 expression on CD8 T cells in blood of normal individuals (NI), in blood of PDA patients (PDA blood) and in tumors. The far left panel is first gated on CD45<sup>+</sup> cells. The right three panels are gated on the CD8<sup>+</sup> T cell gate as shown.

## **Supplementary Figure 3**

# Β



Supplementary Figure 3. Representative human PDA section stained with multiplex immunohistochemical panel. A, Individual antibody stains (left) and positive cell subsets outlined (right).B, Same image stamp from (A) with combined antibodies.





**Supplementary Figure 4.** Example of multiplex IHC analyses of human PDA using HALO software. Top row (**A**, **B**) shows representative staining of tumor cells (magenta), immune cells and PD-L1 (green). Bottom row (**C**, **D**) stromal and tumor regions outlined and cell numbers quantified. Tumor cell nuclei are shown in blue. Light pink outlines PD-L1<sup>+</sup> cells.



CD163 PD-1 PD-L1 DAPI

**Supplementary Figure 5**. PD-L1<sup>+</sup> stromal cells in PDA include CD163<sup>+</sup> macrophages (Figure 6A) as well as an additional CD163<sup>-</sup> cell subset shown here.

**Supplementary Table 1**. Antibodies used for tissue multiplex in Figure 6.

Position	Antibody	Clone/Host	Company	Concentration	<b>OPAL Fluor</b>	
1	CD8	144B/Mouse	DAKO	0.2 µg/ml	540	
2	CD66b	G10F5/Mouse	BD Biosciences	1.25 µg/ml	520	
3	PD-1	D4W2J/Rabbit	Cell Signaling	0.06 µg/ml	570	
4	PD-L1	E1L3N/Rabbit	Cell Signaling	0.5 µg/ml	620	
5	CD163	EP324/Rabbit	Bio SB	0.25 µg/ml	650	
6	AE1/AE3	AE1/AE3/Mouse	DAKO	0.34 µg/ml	690	

щ1	Diagnosia	Differentiation	Change	Neediwent	Survival	Cause of	TIC
#-	Diagnosis	Status/Grade	Stage	Neoadjuvant	(Days)	death	ILS
PC-1	PDA	Moderate	T3N1	N	Alive	Alive	Y
PC-2	PDA	Poor	T1N1	N	864	PDA	Y
PC-3	PDA	Well	T3N0	N	ND <sup>2</sup>	-	Y
PC-4	PDA	Poor	T3N1	Ν	478	Unknown	Ν
PC-5	PDA	Well	T3N0	Ν	ND	-	Y
PC-6	PDA	Moderate	T3N1	N	343	Metastatic	Ν
						angiosarcoma	
PC-7	PDA	Well	T3N0	Ν	ND	-	Ν
PC-8	PDA	Well	T2N1	Ν	ND	-	Ν
PC-9	PDA	Poor	T3N0	Ν	702	Unknown	Ν
PC-10	PDA	Moderate	T3N1	Ν	696	PDA	Y
PC-11	PDA	Moderate	T3N1	Ν	1033	Unknown	Y
PC-12	PDA	Moderate	T3N1	Ν	626	Unknown	Y
PC-13	PDA	Moderate	T3N1	Ν	755	Unknown	Y
PC-14	PDA	Moderate	T3N0	Ν	ND	-	Y
PC-15	PDA	Moderate	T3N0	Y (Chemorads <sup>3</sup> )	983	Unknown	Ν
PC-16	PDA	Moderate	T3N0	Y (Chemo <sup>4</sup> )	ND	-	Y
PC-17	PDA	No PDA <sup>5</sup>	NA <sup>6</sup>	Y (Chemorads)	624	PDA	Ν
PC-18	PDA	Moderate	T0N1	Y (Folfirinox)	961	PDA	Ν
PC-19	PDA	Poor	T3N0	Y (Folfirinox)	ND	-	Ν
PC-20	PDA	Moderate	T2N0	Y (Chemorads <sup>7</sup> )	ND	-	Ν
PC-21	PNET	High	T2N1	Ν	ND	-	Ν
PC-22	PNET	Low	T2N0	Y (Chemo <sup>8</sup> )	Alive	Alive	Ν
PC-23	PNET	Low	T2N1	Ν	Alive	Alive	Ν
PC-24	PNET	Intermediate	T2N0	Ν	ND	-	Ν
PC-25	PNET	Low	T3N0	Ν	ND	-	Ν
PC-26	PNET	High <sup>9</sup>	T3N1	Ν	ND	-	Y
PC-27	PNET	Low	T3N1	Y (Chemorads <sup>10</sup> )	ND	-	Ν

#### Supplementary Table 2. Patient characteristics.

<sup>1</sup>Sample #PC1-PC9 correspond to the same tumors #PC1-PC9 as described in Supplementary Table 3 <sup>2</sup>ND, not determined

<sup>3</sup>Gemcitabine/Taxotere/Oxaliplatin/Capecitabine/XRT

<sup>4</sup>Gemcitabine/Taxotere/Capecitabine

<sup>5</sup>Pancreas with atrophy, fibrosis, dilated ducts, and residual islet cell islands

<sup>6</sup>NA, not applicable <sup>7</sup>Folfirinox/Capecitabine/XRT <sup>8</sup>Capecitabine /Temozolomide <sup>9</sup>Mixed neuroendocrine and acinar features

<sup>10</sup>Capecitabine/XRT

	_							-	Мах
			#		#	%			Freq
Sample	Tissue	# Seq	Unique	# Prod	Unique	Prod	Entropy	Clonality	(%)
PC-1	Blood	22683	18915	17596	14718	77.6	13.56	0.02	0.69
PC-2	Blood	22616	16801	18317	13573	81.0	12.92	0.06	1.19
PC-3	Blood	34536	24286	28189	20166	81.6	12.64	0.12	7.91
PC-4	Blood	40341	19349	32245	15554	79.9	11.00	0.21	7.13
PC-5	Blood	55151	45949	45332	37807	82.2	14.84	0.02	0.42
PC-6	Blood	41850	35797	34637	29492	82.8	14.69	0.01	0.20
PC-7	Blood	257543	211787	205772	167410	79.9	16.23	0.06	3.13
PC-1	Tumor	9369	4572	7069	3402	75.5	10.49	0.11	2.31
PC-2	Tumor	8782	4069	7126	3264	81.1	10.31	0.12	2.74
PC-3	Tumor	1149	922	873	708	76.0	9.13	0.04	4.26
PC-4	Tumor	11053	8100	8705	6443	78.8	12.06	0.05	1.44
PC-5	Tumor	13490	7696	11145	6252	82.6	11.66	0.08	1.16
PC-6	Tumor	23728	11148	20217	9237	85.2	11.73	0.11	3.22
PC-7	Tumor	8095	5418	6226	4152	76.9	11.28	0.06	1.54
PC-8	Tumor	86957	14772	69736	11968	80.2	9.44	0.30	8.53
PC-9	Tumor	22738	7735	17322	5756	76.2	10.57	0.15	1.83

**Supplementary Table 3**. TCR<sup>β</sup> CDR3 sequence data in blood and tumors from PDA patients.

TCRβ CDR3 sequence data is located at Adaptive Biotechnologies immunoSEQ ANALYZER: https://clients.adaptivebiotech.com/pub/stromnes-2017-cancerimmunologyresearch