

## **Supplemental Material**

Table S1-S3

Figure S1-8

**Supplementary Table 1**

Species	# of cells	% of reads mapped to genome	Mean reads/cell	Median genes/cell	Total genes detected
Mouse	3013	78.1	80508	952	15380
Rat	3935	82.5	54456	1113	14944
Pig	4671	90.9	61638	1065	14589
Human	2868	91.7	112080	878	17658

**Supplementary Table 2**

Cell Type	Gene	P value	Cell Type	Gene	P value
Resident macrophages	C1qa	0.00E+00	Unknown	Cd7	2.06E-179
Resident macrophages	C1qb	0.00E+00	Unknown	Xcl1	4.04E-179
Resident macrophages	<b>C1qc</b>	0.00E+00	Unknown	Cxcr6	3.33E-120
Resident macrophages	Cd81	0.00E+00	Unknown	Trbc2	8.37E-104
Resident macrophages	Apoe	0.00E+00	Unknown	Gimap4	4.19E-103
Resident macrophages	H2-Eb1	0.00E+00	Unknown	Gimap6	2.55E-79
Resident macrophages	Ctss	0.00E+00	Unknown	Ltb	1.36E-76
Resident macrophages	Ms4a7	7.14E-240	Unknown	Trbc1	3.35E-70
Resident macrophages	Cd74	3.25E-229	Unknown	Ccl4	1.03E-07
Resident macrophages	Mmp12	1.48E-135	Unknown	Ccl3	9.89E-04
cDC2 DCs	Cd209a	0.00E+00	Innate Lymphoid Cells	Gata3	2.94E-279
cDC2 DCs	<b>Clec10a</b>	1.12E-274	Innate Lymphoid Cells	<b>Cxcr6</b>	6.75E-200
cDC2 DCs	Tnip3	2.53E-210	Innate Lymphoid Cells	Lztf1l	9.02E-150
cDC2 DCs	Cfp	8.55E-188	Innate Lymphoid Cells	Ly6a	2.01E-121
cDC2 DCs	H2-Oa	4.82E-186	Innate Lymphoid Cells	Skap1	1.66E-113
cDC2 DCs	H2-DMb2	1.42E-146	Innate Lymphoid Cells	Ltb	6.25E-99
cDC2 DCs	S100a6	6.01E-125	Innate Lymphoid Cells	Hs3st1	3.74E-89
cDC2 DCs	Napsa	1.67E-123	Innate Lymphoid Cells	Phlda1	3.44E-87
cDC2 DCs	Tnfsf9	2.13E-112	Innate Lymphoid Cells	Lmo4	2.29E-35
cDC2 DCs	Socs3	8.42E-69	Innate Lymphoid Cells	Junb	8.94E-29
NK cells	Gzma	0.00E+00	cDC1 DCs	Xcr1	0.00E+00
NK cells	<b>Nkg7</b>	0.00E+00	cDC1 DCs	Ifi205	4.49E-219
NK cells	Gzmb	0.00E+00	cDC1 DCs	Cd24a	6.31E-214
NK cells	Klre1	0.00E+00	cDC1 DCs	Cxx1b	5.95E-160
NK cells	Klra8	0.00E+00	cDC1 DCs	Cxx1a	7.09E-150
NK cells	Klra4	2.76E-302	cDC1 DCs	Tnni2	3.39E-122
NK cells	Klra9	1.27E-297	cDC1 DCs	<b>Naaa</b>	4.87E-96
NK cells	Ccl5	6.02E-284	cDC1 DCs	Ppt1	9.94E-58
NK cells	AW112010	2.24E-199	cDC1 DCs	Cst3	1.01E-51
NK cells	Lgals1	1.74E-190	cDC1 DCs	Irf8	9.37E-51
Ly6chi macrophages	Chil3	4.23E-160	B cells	Ly6d	0.00E+00
Ly6chi macrophages	Plac8	2.50E-138	B cells	Cd79a	0.00E+00
Ly6chi macrophages	Ms4a4c	5.04E-127	B cells	Iglic3	0.00E+00
Ly6chi macrophages	Lyz1	2.13E-118	B cells	Ms4a1	0.00E+00
Ly6chi macrophages	Ly6c2	3.23E-111	B cells	Ebf1	0.00E+00
Ly6chi macrophages	Ms4a6c	2.64E-89	B cells	Iglic2	3.04E-153
Ly6chi macrophages	Ifitm3	1.09E-77	B cells	Iglic1	7.77E-150
Ly6chi macrophages	Lyz2	4.71E-77	B cells	Cd79b	5.07E-90
Ly6chi macrophages	<b>Irf7</b>	1.35E-67	B cells	Ighm	2.80E-39
Ly6chi macrophages	Gbp2	1.89E-60	B cells	Igkc	3.51E-39
Ly6clo macrophages	Ear2	4.26E-255	Neutrophils	Retnlg	0.00E+00
Ly6clo macrophages	Fabp4	9.84E-246	Neutrophils	Slpi	6.02E-283
Ly6clo macrophages	Eno3	8.06E-228	Neutrophils	S100a9	3.84E-209
Ly6clo macrophages	Treml4	1.63E-216	Neutrophils	<b>S100a8</b>	1.07E-206
Ly6clo macrophages	Pglyrp1	8.26E-167	Neutrophils	Hp	1.90E-144
Ly6clo macrophages	<b>Cebpb</b>	7.06E-130	Neutrophils	G0s2	7.00E-124
Ly6clo macrophages	Msrb1	6.57E-127	Neutrophils	Cxcl2	4.22E-116
Ly6clo macrophages	Nr4a1	6.65E-126	Neutrophils	Msrb1	1.36E-57
Ly6clo macrophages	Gngt2	1.72E-117	Neutrophils	Il1b	3.03E-47
Ly6clo macrophages	Smpdl3a	1.50E-115	Neutrophils	S100a11	2.10E-38

Supplementary Table 3

Infiltrating macrophage	Resident macrophage		
Cell number	Reads	Cell number	Reads
R1_C12	1,492,862	R2_C03	2,134,946
R1_C15	2,227,789	R2_C05	2,303,807
R1_C16	6,460,052	R2_C06	738,061
R1_C23	1,050,069	R2_C10	159,872
R1_C24	2,596,836	R2_C12	253,938
R1_C25	3,425,398	R2_C13	3,578,235
R1_C32	3,406,075	R2_C14	2,304,633
R1_C34	3,262,353	R2_C21	2,807,416
R1_C38	1,247,509	R2_C23	3,143,572
R1_C40	258,583	R2_C28	1,657,015
R1_C52	3,180,335	R2_C30	1,896,495
R1_C53	1,822,262	R2_C32	476,005
R1_C54	2,280,668	R2_C34	4,799,911
R1_C55	2,294,715	R2_C36	4,347,918
R1_C56	3,071,868	R2_C39	1,542,320
R1_C58	4,297,437	R2_C42	2,068,422
R1_C60	3,016,114	R2_C46	1,865,383
R1_C63	3,895,750	R2_C47	1,067,191
R1_C65	1,038,025	R2_C53	91,473
R1_C69	653,524	R2_C55	4,186,470
R1_C70	4,737,539	R2_C56	2,602,547
R1_C71	2,564,324	R2_C60	183,934
R1_C77	216,424	R2_C61	4,719,640
R1_C78	2,581,007	R2_C64	1,628,547
R1_C81	9,560	R2_C73	1,259,852
R1_C83	3,635,582	R2_C74	3,947,573
R1_C84	2,171,672	R2_C78	4,059,899
R1_C85	644,928	R2_C79	1,172,156
R1_C87	2,828,777	R2_C80	1,811,933
R1_C90	3,485,523	R2_C81	206,613
R1_C91	1,703,711	R2_C82	6,985,563
R1_C93	941,270	R2_C84	2,860,895
R1_C95	2,109,498	R2_C90	1,543,593
		R2_C91	2,093,469
		R2_C92	2,839,715
		R2_C93	2,515,540

**Figure S1.** tSNE projection and violin plot showing expression of *Cd79a* in mouse renal innate immune cells.

**Figure S2** Gating strategy used to isolate infiltrating and resident macrophages based on the canonical Cd11b and F/480 approach. Infiltrating and resident macrophages were collected and subjected to C1 Fluidigm single cell RNA sequencing.

**Figure S3** tSNE projections and violin plots showing a top DEG that was used to identify **(A)** neutrophils (*S100a8*), **(B)** NK cells (*Nkg7*), **(C)** innate lymphoid cells (ILCs, *Cxcr6*), **(D)** infiltrating macrophages (*Irf7*, *Cebpb*), **(E)** and dendritic cells (*Naaa*, *Clec10a*) in single cell data from a rat kidney.

**Figure S4** tSNE projections and violin plots showing a top DEG that was used to identify **(A)** neutrophils (*S100A8*), **(B)** NK cells (*NKG7*), **(C)** innate lymphoid cells (ILCs, *CXCR6*), **(D)** infiltrating macrophages (*IRF7*, *CEBPB*), and **(E)** dendritic cells (*NAAA*) in single cell data from a pig kidney. Unfortunately, the *Clec10a* orthologue has not been identified in the pig.

**Figure S5** tSNE projections and violin plots showing a top DEG that was used to identify **(A)** neutrophils (*S100A8*), **(B)** NK cells (*NKG7*), **(C)** innate lymphoid cells (ILCs, *CXCR6*), **(D)** infiltrating macrophages (*IRF7*, *CEBPB*), **(E)** and dendritic cells (*NAAA*, *CLEC10A*) in single cell data from a human kidney.

**Figure S6** Heat map of top 100 DEGs in lineage negative CD45<sup>+</sup> innate immune cells from human kidney tissue identified using Fluidigm C1 single cell RNA sequencing.

**Figure S7** Gating strategy used to identify resident macrophages in multiple species using CD74 and CD81 antibodies.

**Figure S8** Quantification of the percent chimerism of B/T cells, infiltrating monocytes/macrophages, and neutrophils in the blood of mice undergoing parabiosis. \*\*\*  
P<0.0001.

Figure S1.

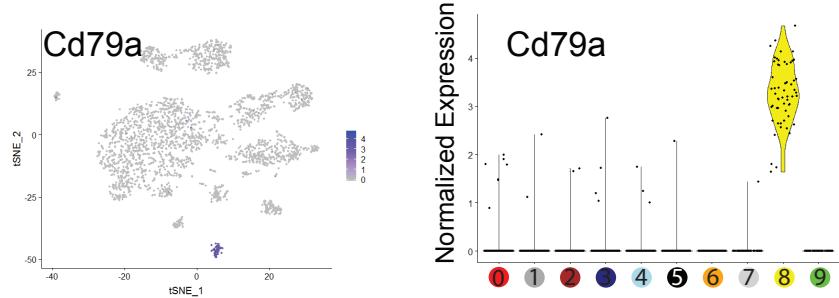
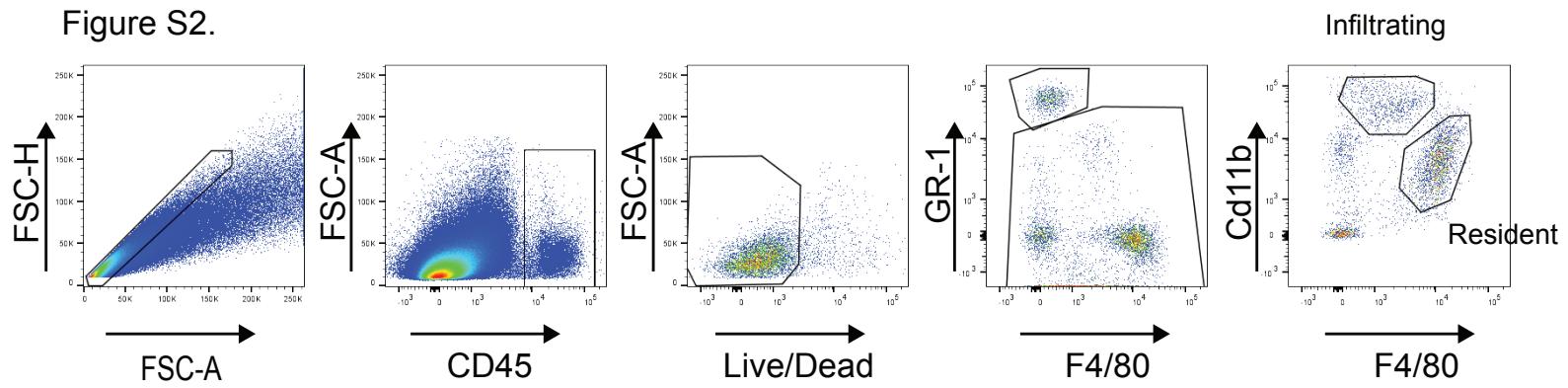


Figure S2.



**Figure S3**

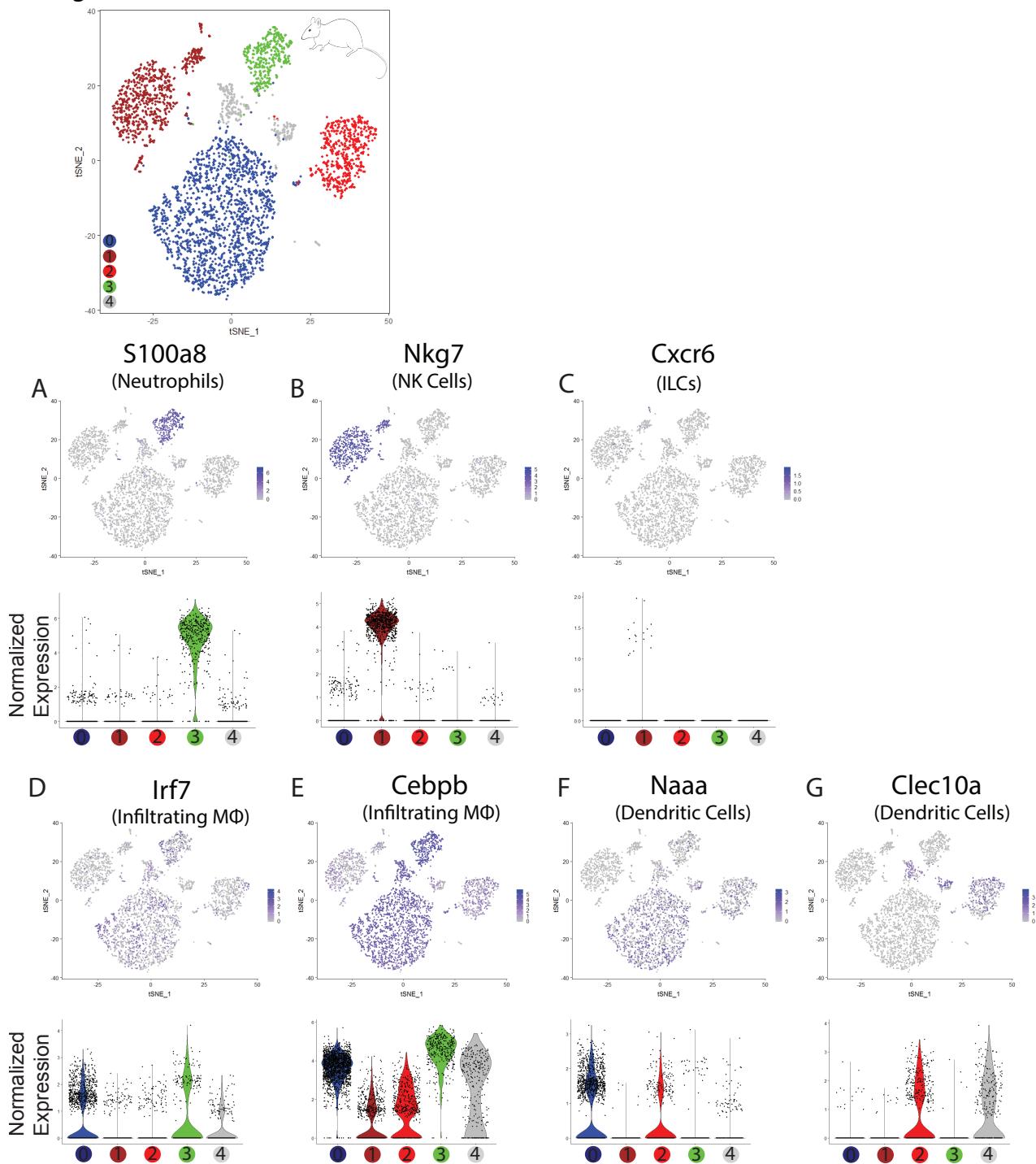


Figure S4

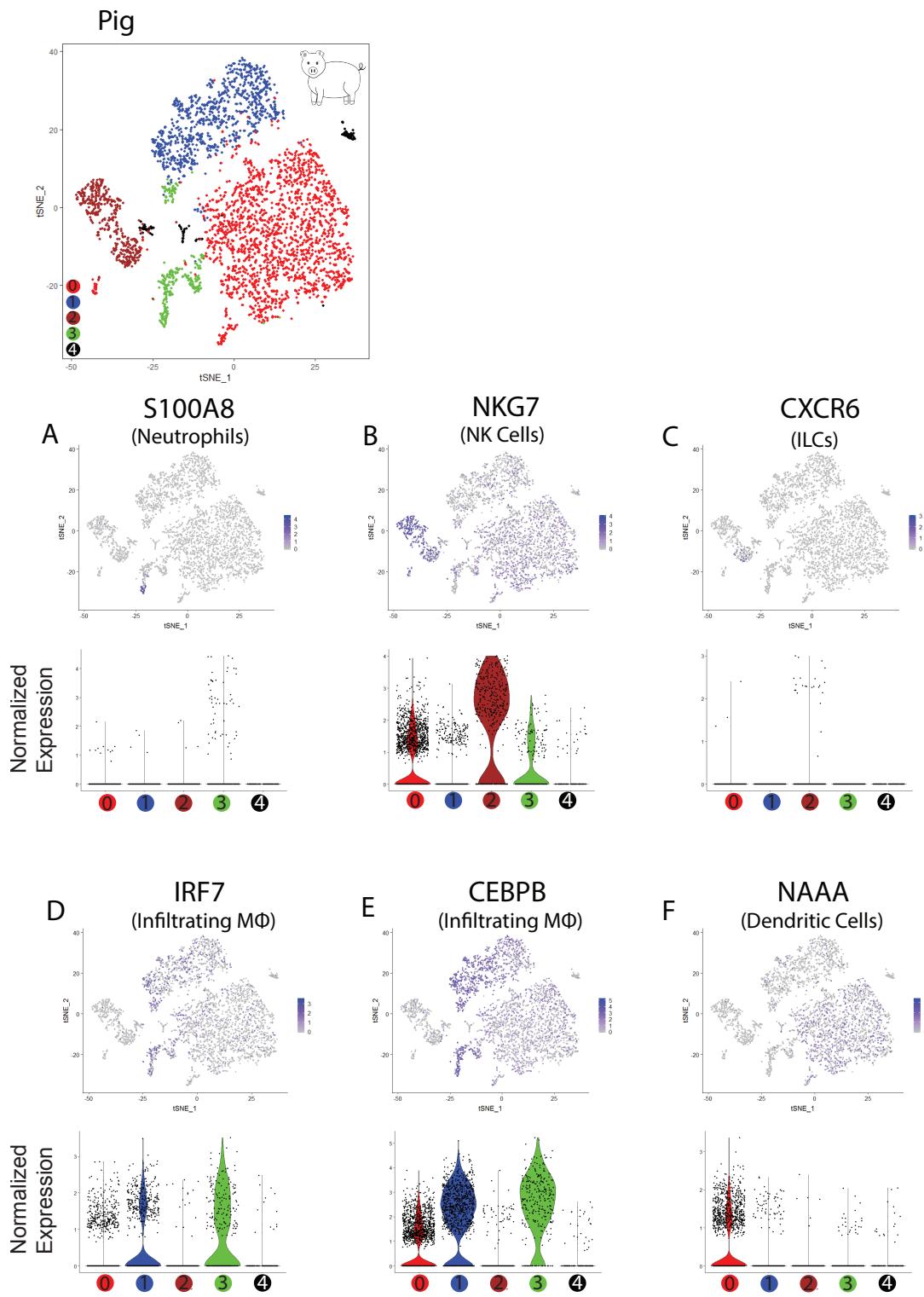


Figure S5

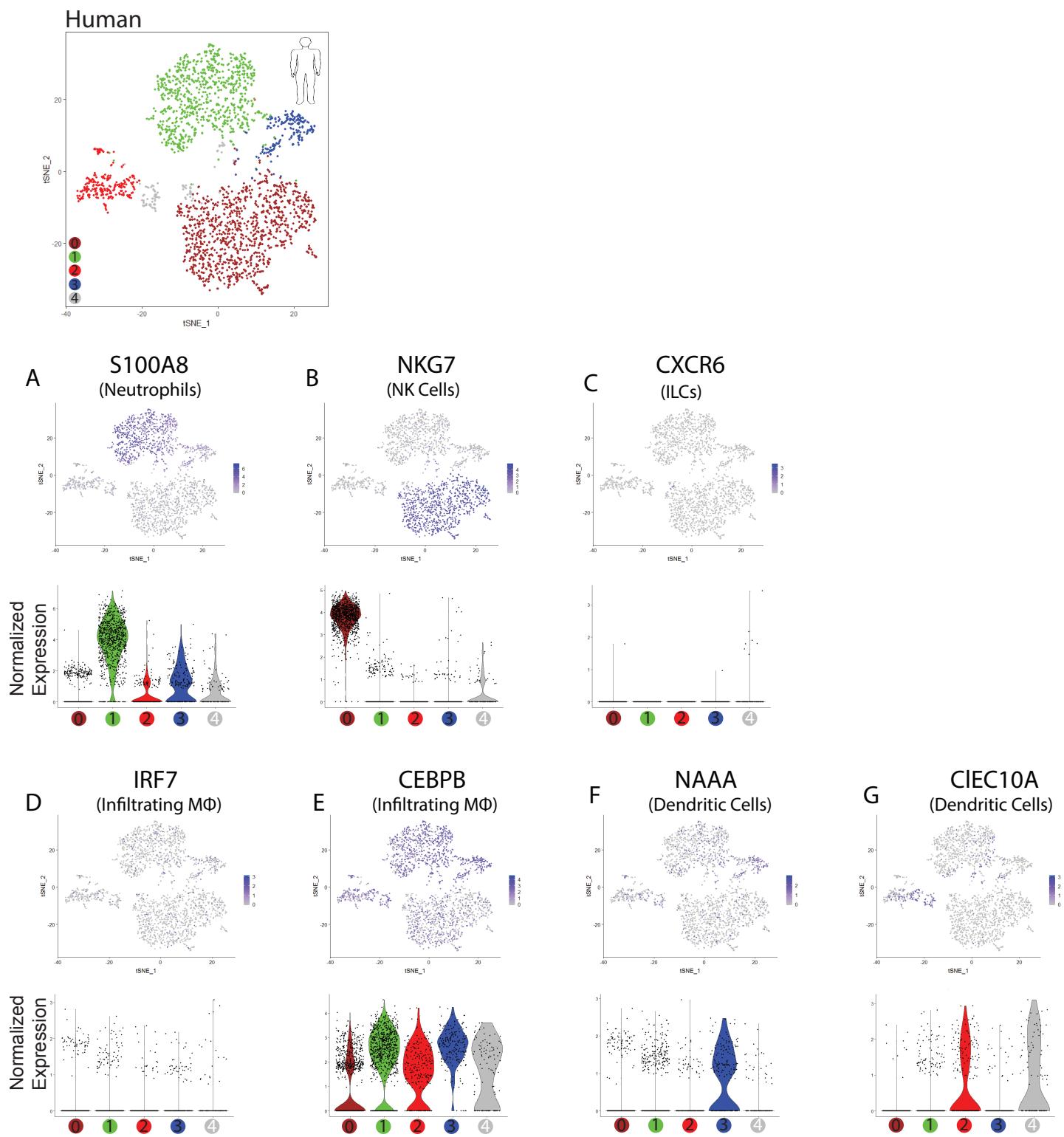


Figure S6

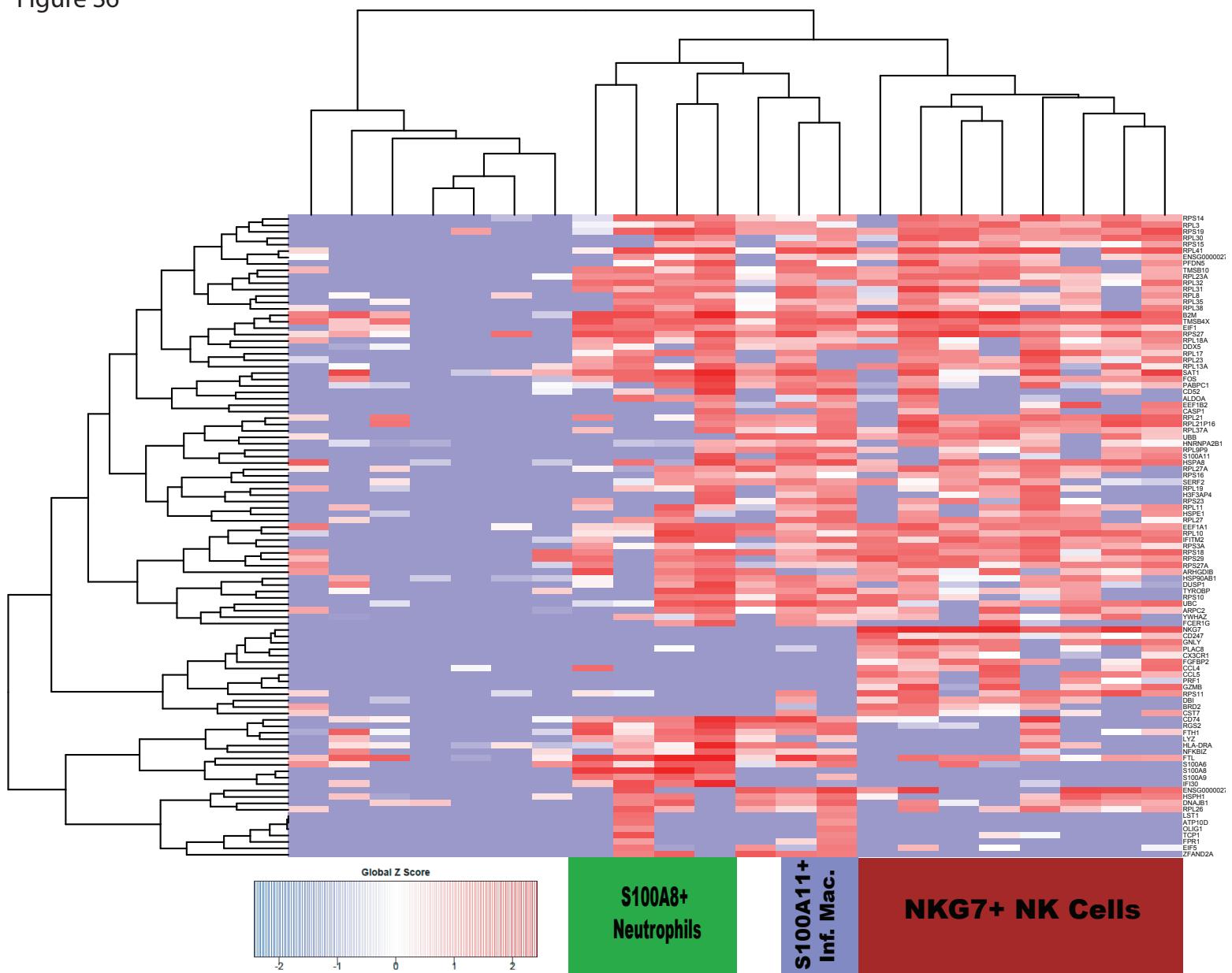


Figure S7

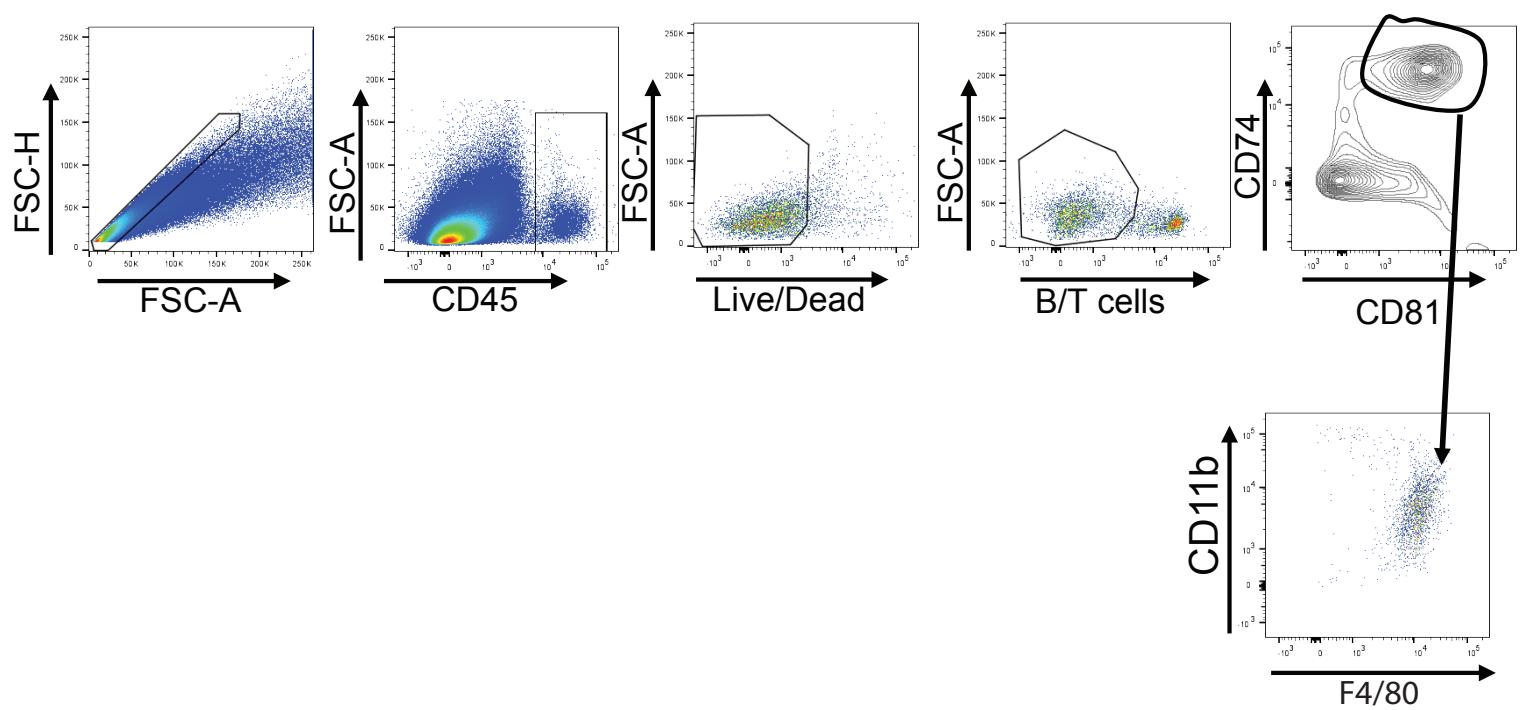


Figure S8

