Supplementary Materials for

Single-cell atlas of peripheral blood mononuclear cells from pregnant women

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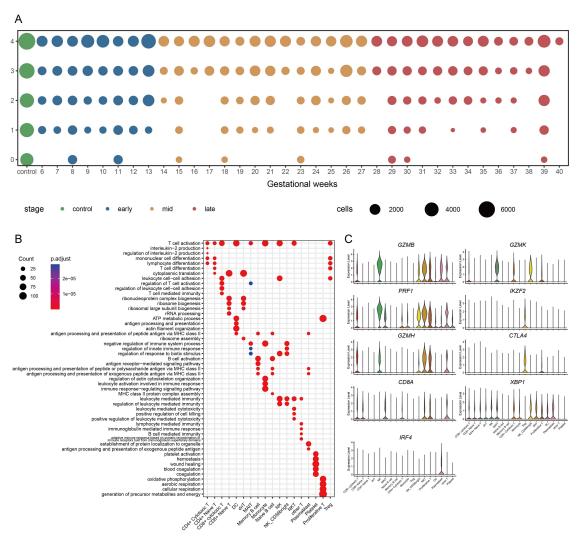


FIGURE S1 Single cell data display for each library and GO term for all cell types (related to Figure 1). (A) Schematic diagram of sample collection. The abscissa axis indicates gestational weeks (GW). Control means subjects who are not pregnant (n=5). Each point represents a subject. The size of the point indicates the number of PBMC collected by a subject, and the color indicates the pregnancy period of the subject. early= the first trimester (GW6-13) (n=34), mid= the second trimester (GW14-27) (n=52), late= the third trimester (GW28-40) (n=45). (B) GO term enrichment for differentially expressed genes in all cell types. The p adjusted values lower, point redder. Point size represents the number of DEGs involved in specific GO terms. (C) Violin plots of expression values for other cell type-specific marker genes.

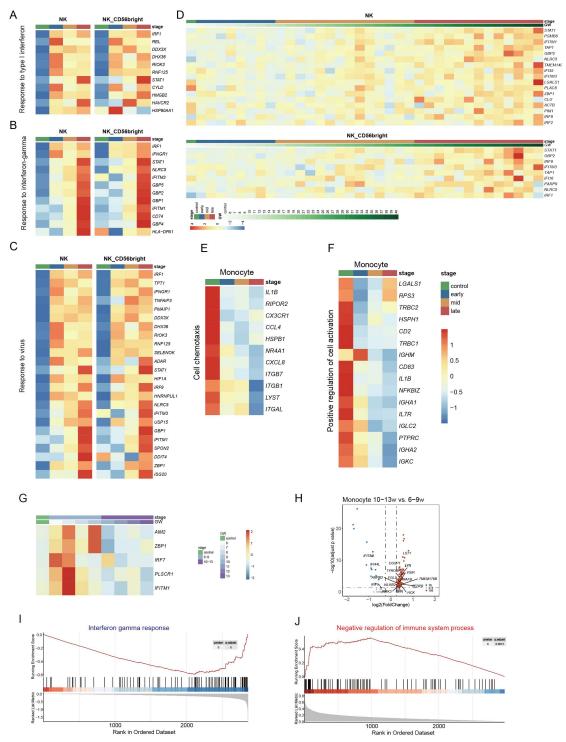


FIGURE S2 Transcriptomic profiling of NK cells and monocytes during pregnancy (related to Figure 3). (A) Heatmap of DEGs enriched in "response to type I interferon", (B) "response to interferon-gamma", (C) "response to virus" in NK/NK_CD56bright cells across four conditions. Colors represent scaled genes expression level, and higher expression is red and lower expression is blue. (D) Heatmap of ISGs across four stages in NK/NK_CD56bright cells. Colors represent scaled genes expression level, and higher expression is red and lower expression is blue. (E-F) Heatmap of DEGs enriched in "cell chemotaxis", "positive regulation of cell activation" across four stages in monocytes. (G) Heatmap of the ISGs, including *AIM2*, *ZBP1*, *IRF7*, *PLSCR1* and *IFITM1* in monocytes derived from GW6-13 and pre-pregnant control samples. (H) Volcano

plots of DEGs of monocyte in GW6-9 and GW10-13. (I-J) GSEA results of "interferon gamma response" and "negative regulation of immune system process".

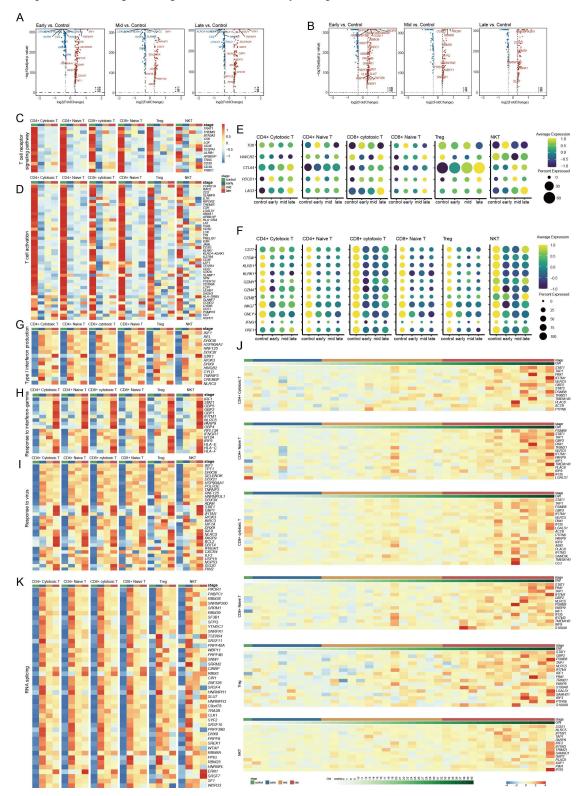


FIGURE S3 Transcriptomic profiling of T cells throughout gestation (related to Figure 4). (A) Volcano plots of differentially expressed genes in T cells. Interferon-related genes and T cell activation-related genes are labeled. (B) Volcano plots of differentially expressed genes in T cells. RNA splicing related genes are labeled. (C-D) Heatmap of DEGs enriched in "T cell receptor

signaling pathway", "T cell activation" in T cells across four conditions. (E-F) Dot plot showing expression of genes associated with exhausted and cytotoxic in T cells across four conditions. (G-I) Heatmap of DEGs enriched in "type I interferon production", "response to interferon-gamma", "response to virus" in T cells across four conditions. (J) Heatmap of ISGs among every gestational week. Colors represent scaled genes expression level, and higher expression is red and lower expression is blue. (K) Heatmap of DEGs enriched in "RNA splicing" in T cells across four conditions.

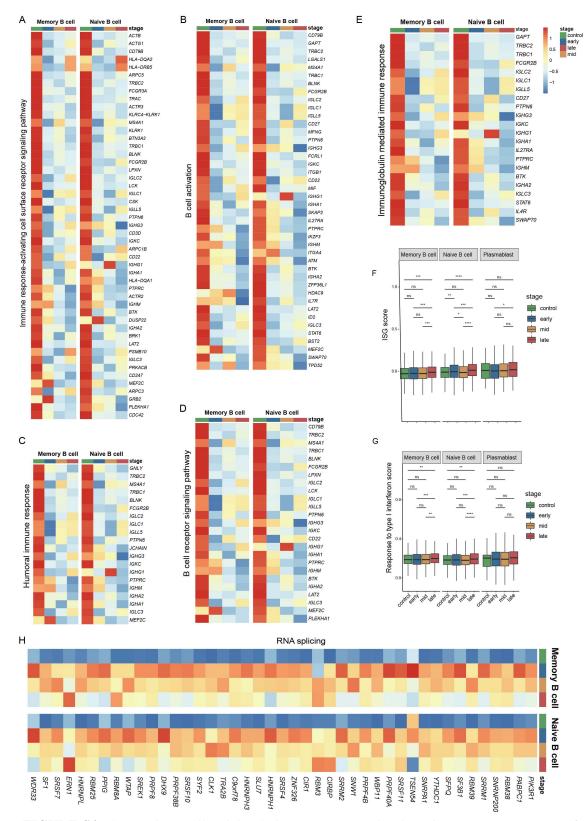


FIGURE S4 Changes in B cell activity during pregnancy (related to Figure 5). (A) Heatmap of DEGs enriched in "immune response-activating cell surface receptor signaling pathway", (B) "B cell activation", (C) "humoral immune response", (D) "B cell receptor signaling pathway", (E) "immunoglobulin mediated immune response" in Naïve/Memory B cells across four conditions. (F) Box plots of the collected ISGs scores across different clusters and conditions. Statistical significance of difference in pairs among four pregnant stages were labeled at the top of box plot.

Wilcoxon rank-sum test was applied. (G) Box plots of the cell scores of GO biological process terms (response to type I interferon) in B cells across four conditions. Wilcoxon rank-sum test was applied. All differences with p < 0.05 are indicated. *p < 0.05, **p < 0.01, ****p < 0.001, ns= not significant. (H) Heatmap of DEGs enriched in "RNA splicing" in Naïve/Memory B cells across four conditions.

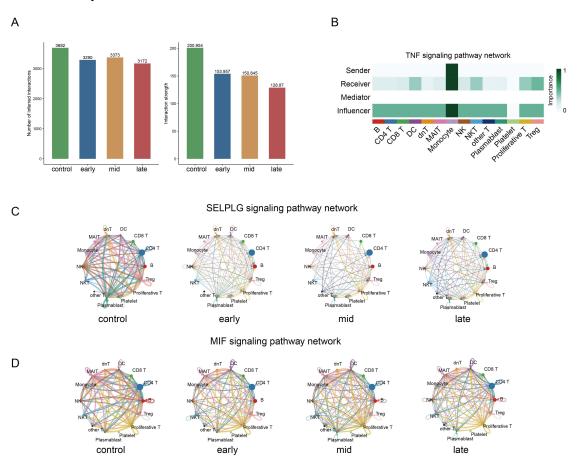


FIGURE S5 PBMC cell communication in pregnancy and non-pregnancy (related to Figure 6). (A) The bar chart shows that the total number of interactions of the inferred cell-cell communication networks from different conditions. (B) Heatmap shows centrality scores of TNF signaling pathway network. (C-D) SELPLG signaling pathway network and MIF signaling pathway network in different conditions.

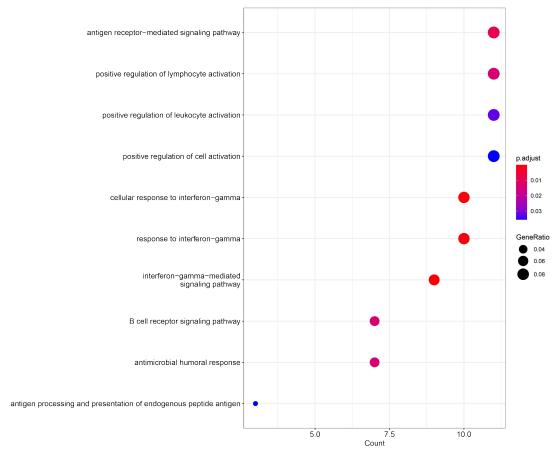


FIGURE S6 Enriched GO terms for genes involved in the five cell-type-specific models toward gestational age prediction (related to Figure 7)

Supplementary Table 1. Basic information of the enrolled subjects

	Control	Early	Mid	Late
	non-pregnancy	the first trimester (GW6-13)	the second trimester (GW14-27)	the third trimester (GW28-40)
	N=5	N=34	N=52	N=45
Demographics				
Age, years	25±1.9	29.7±2.7	28.7±2.6	28.7±3.2
BMI, kg/m ²	22.85±3.05	20.65±2.01	22.51±1.95	25.62±2.47
Previous births, No. (%)				
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)
Abortion, drug abortion, and ectopic pregnancy, No. (%)				
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)
Pregnancy complications, No. (%)				
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)
Immune system diseases, No. (%)				
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)
Appears during pregnancy, No. (%)				
Subclinical hypothyroidism				
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)
ICP		` ′	` ′	· /
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)
GDM	. ,	` '	` ′	` ′
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)
HDP				
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)
Gestational thrombocytopenia	. ,	` '	` ′	` ′
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)
FGR	- (/	- (/	- (/	
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)
Macrosomia	7(220)	- (()	. ()	- ()
Yes	0(0)	0(0)	0(0)	0(0)
No	5(100)	34(100)	52(100)	45(100)

ICP: intrahepatic cholestasis of pregnancy, GDM: gestational diabetes mellitus, HDP: hypertensive disorders of pregnancy, FGR: fetal growth restriction

Supplementary Table 2. Gestational weeks and cell ratio of all subjects

Supp		ntary Gestational		2. Ges	cD8+	CD8+				of all	•							Deoliforotico	
Sample	Stage	weeks	Cvtotoxic 7	Γ Naive T	cvtotoxic T	Naive T DC	dnT	MIMI	Memory B cell	Monocyte	B cell		NK_CD56bright				Platelet	Proliferative T	reg
control1_ control1_		control control	2.39% 3.01%	17.12% 13.84%	35.53% 38.28%	14.85% 0.68% 13.84% 0.36%	7.35%	2.46% 2.37%	2.54% 3.96%	0.80%	2.50% 3.22%	8.94% 9.35%	1.29% 1.07%	0.95% 0.87%	0.04%	0.34% 0.23%	0.23% 0.18%	0.53% 0.46%	0.87% 1.30%
control2_ control2		control	2.28% 2.56%	19.94% 19.17%	19.37% 15.44%	28.47% 0.57% 28.42% 0.59%	14.19% 17.99%		2.00%	0.29%	1.06%	4.45% 4.52%	1.14%	1.26%	0.00%	0.61%	0.24%	0.12%	1.26%
control3		control	2.43% 2.51%	16.13% 17.67%	22.91% 24.00%	16.88% 0.65% 17.24% 0.72%		4.53% 5.65%	2.66% 2.61%	3.23% 1.50%	5.28% 4.78%	6.50% 5.84%	0.51% 0.58%	0.56%	1.54% 0.00%	0.33% 0.19%	0.05%	0.33%	1.82% 1.21%
control4_ control4	1 control	control	3.03%	13.64%	33.59%	17.55% 0.12%	9.61%	2.79%	1.95%	0.28%	1.72%	10.93%	1.32%	1.32%	0.00%	0.28%	0.04%	0.24%	1.60%
control5_	l control	control	3.59% 2.06%	14.53% 14.06%	28.44% 24.35%		9.65% 17.91%	2.75%	2.58%	0.06% 0.95%	2.84%	10.49% 7.32%	1.18% 0.49%	1.07% 0.82%	0.06%	0.17% 0.13%	0.06% 0.26%	0.34% 0.42%	2.02%
control5_ NP150	2 control 6	control early	2.06% 3.15%	14.95% 16.61%	21.92% 28.23%		12.76%	2.27%	3.03% 1.49%	0.83% 4.28%	4.59% 0.96%	6.65% 5.68%	0.55% 1.14%	1.22%	0.00%	0.00% 0.35%	0.05%	0.23% 0.61%	2.06% 1.22%
NP152 NP78	6 6	early early	1.34% 2.21%	16.68% 18.38%	11.12% 21.21%	26.27% 0.96% 15.83% 0.96%	15.82% 9.69%	4.31% 3.50%	2.49% 3.41%	4.99% 1.92%	4.31% 5.61%	6.33% 11.37%	2.11% 0.72%	0.29% 1.01%	0.00%	0.77% 0.91%	0.00%	0.48% 1.10%	1.73% 2.16%
NP87 NP115	6 7	early early	2.00% 1.17%	15.91% 12.37%	21.69% 25.91%	16.69% 1.11% 16.97% 1.04%		5.23% 4.99%	2.00%	2.22%	3.23% 4.53%	11.35% 7.12%	0.89% 1.10%		0.22%	0.11% 1.36%	0.44%	0.89% 0.26%	1.11%
NP120 NP124	7	early early	2.51% 1.70%	13.28% 18.93%	20.87% 25.33%	20.15% 0.53% 21.28% 0.26%	11.76%	6.27%	3.24% 1.44%	1.32%	4.36% 1.31%	11.56% 8.88%	1.25% 0.52%	0.59%	0.07%	0.33% 0.52%	0.00% 0.26%	0.33%	1.59% 1.57%
NP76 NP103	7 8	early early	2.30% 2.46%	18.92% 15.02%	19.81% 30.48%	17.68% 0.51% 18.71% 0.44%			1.45%	6.26% 3.55%		10.52% 4.14%	1.11% 0.64%	0.60%	0.00%	0.47%	0.04%	1.36%	2.17%
NP111 NP117	8	early	2.75%	19.08%	27.62%	11.58% 0.67% 20.13% 0.40%	8.69%	5.35%	4.83%	5.20%	2.75%	6.98%	1.56%		0.00%	0.89%	0.00%	0.30%	0.74%
NP130 NP75	8	early early	2.01%	17.16% 19.96%	34.74% 24.83%	8.37% 1.04% 18.77% 1.26%	7.89%	3.25%	2.84%	3.74%	3.46% 0.44%	10.24%	0.35%	1.11%	1.04%	0.69%	0.07%	0.42%	1.59%
NP54	8	early early	2.13%	30.98%	23.11%	15.57% 1.15%	11.64%	0.98%	1.33% 2.30%	1.48%	1.31%	5.40% 4.59%	1.48%	0.66%	0.00%	0.66%	0.00%	0.16%	1.80%
NP65 NP74	9	early early	1.68% 2.84%	9.02% 14.48%	29.64% 32.25%		7.17% 10.73%		3.36% 2.57%	3.64% 0.89%	6.16% 1.06%	16.13% 6.47%	0.84% 1.12%		5.27% 0.00%	0.22% 0.24%	0.00%	2.07% 0.21%	0.84% 2.04%
NP79 NP104	9 10	early early	2.27% 2.55%	30.24% 22.90%	17.03% 24.36%	12.90% 0.83% 20.28% 1.02%	6.91% 12.55%	4.60%	4.44% 1.82%	0.93% 0.44%	2.19%	11.15% 3.21%	0.00% 0.44%	0.31% 0.88%	0.10%	0.31% 0.15%	0.00%	2.06% 0.88%	0.00% 1.75%
NP108 NP6	10 10	early early	3.11% 1.80%	23.94% 21.82%	27.98% 19.73%	19.07% 0.85%	13.09%	6.29% 7.12%	2.21% 2.47%	1.67% 1.71%	1.71%	12.80% 6.26%	0.87% 1.42%	1.04%	0.00%	0.39%	0.03%	0.42%	1.44%
NP85 NP107	10 11	early early	4.56% 3.05%	26.03% 19.06%	23.68% 30.43%	11.25% 0.95%	12.79% 9.97%	3.97% 4.32%	3.82% 1.02%	1.76% 3.49%	1.47% 0.32%	3.09% 10.93%	0.29% 1.27%	0.44% 1.08%	0.00%	0.44% 0.64%	0.00%	0.74% 1.08%	1.76% 1.08%
NP123 NP56	11 11	early early	2.18% 2.08%	17.49% 15.74%	25.45% 26.94%	22.97% 0.15% 22.77% 0.72%	12.61%		1.65% 2.21%	1.20%	1.50%	5.33% 6.70%	1.13% 0.78%	0.75%	0.00%	0.38%	0.08%	0.15% 0.85%	1.65%
NP57 NP94	11 11	early early	4.79% 2.43%	18.34% 19.61%	20.60%	17.18% 0.82% 13.18% 0.54%	13.69% 8.75%	1.98%	3.83%	1.23% 0.81%	4.18%	9.17% 4.70%	0.75% 0.27%	0.62%	0.00%	1.03%	0.34%	0.21%	1.23%
NP110 NP121	12 12	early early	1.61% 3.35%	16.83% 22.04%	26.58% 25.82%	20.76% 0.91% 17.16% 0.36%			1.33%	0.28%	1.19%	4.70% 2.69%	0.84%	0.42%	0.00%	1.68%	0.00%	0.35%	3.02% 1.45%
NP49 NP92	12 12	early early	3.01% 4.83%	15.88% 23.57%	23.63% 22.69%	16.42% 1.46% 16.92% 0.65%			1.09%	2.92% 0.84%	3.10% 3.67%	7.66% 4.60%	1.00% 1.12%	0.82%	0.00%	0.09%	0.00%	0.27%	1.09%
NP119 NP52	13 13	early early	3.10%	18.43% 18.25%	29.59%	18.55% 0.31% 24.71% 0.57%	13.32%	2.12%	2.87%	0.28%	3.77%	3.22%	0.79% 0.86%	1.34%	0.00%	0.20%	0.04%	0.43%	1.65%
NP77 NP80	13 13	early early	3.40% 2.09%	21.69% 19.04%	30.48% 40.14%	16.94% 0.44% 9.50% 0.44%		6.65%	4.00%	0.42% 1.85%	2.12% 2.78%	3.45% 5.72%	0.53% 0.56%	0.75%	0.00%	0.18%	0.13%	0.40%	1.37%
NP126	14	mid	5.49%	20.91%	31.78%	7.83% 0.58%	11.33%	3.04%	2.69%	0.35%	0.70%	5.84%	3.86%	1.75%	0.00%	1.05%	0.23%	0.70%	1.87%
NP21 NP70	14 14	mid mid	2.24% 2.32%	18.16% 16.86%	20.85% 30.74%	21.30% 0.22% 14.96% 0.98%	10.88%	3.56%	1.35% 3.61%	4.93% 3.40%	4.48% 3.04%	8.30% 2.94%	1.12% 0.83%	0.45% 1.03%	0.00% 3.20%	0.90% 0.15%	0.00% 0.15%	0.67% 0.41%	1.12% 0.93%
NP109 NP116	15 15	mid mid	1.13% 2.37%	17.80% 16.70%	21.60% 23.52%	20.20% 0.67% 21.74% 0.89%	15.61%	1.98%	3.93% 3.06%	1.27% 0.49%	6.53% 2.37%	5.33% 8.40%	0.27% 0.30%	0.40%	0.00%	0.60% 0.49%	0.13% 0.20%	0.47% 0.30%	2.20% 1.19%
NP153 NP154	15 15	mid mid	3.05% 1.63%	11.07% 15.53%	30.73% 29.86%	11.45% 1.15% 19.45% 0.36%	5.34% 9.69%	3.44% 1.08%	1.72% 1.20%	16.03% 3.55%	1.72% 0.96%	8.59% 9.51%	0.76% 0.60%	1.34% 0.90%	0.38% 1.69%	0.19% 0.42%	0.19% 0.06%	2.10% 2.41%	0.76% 1.08%
NP36 NP24	15 16	mid mid	1.15% 1.43%	12.01% 22.81%	24.52% 10.90%	21.58% 0.38% 25.68% 0.14%			1.28% 0.43%	7.28% 5.16%	1.02% 1.29%	6.51% 7.75%	1.40% 1.29%	0.38% 1.43%	0.26% 0.00%	1.02% 0.14%	0.00% 0.14%	0.77% 0.29%	1.53% 1.87%
NP55 NP122	16 17	mid mid	3.87% 2.64%	13.80% 24.09%	32.21% 20.01%	14.73% 0.70% 18.48% 0.41%		5.45% 4.13%	5.87% 2.72%	0.33%	4.10% 1.53%	6.57% 4.46%	0.56% 1.16%	1.49%	0.00%	0.42%	0.05%	0.61% 1.65%	0.98%
NP82 NP12	17 18	mid mid	2.48% 3.85%	12.88% 19.46%	22.38% 19.27%	25.14% 0.55% 25.82% 0.96%	14.81%		2.13%	1.03%	4.89%	2.96%	0.62%	0.62%	0.00%	0.48%	0.07%	0.28%	1.17%
NP142 NP145	18 18	mid mid	4.19%	10.03% 17.44%	33.50% 15.68%		8.50%	3.05%	1.78%	6.22%	1.90%	9.90% 8.57%	1.52%	0.76%	4.19%	1.02%	0.00%	0.76%	1.14%
NP146 NP64	18 18	mid mid	2.73%	17.99% 17.73%	24.86% 25.64%	22.90% 0.65% 19.80% 1.27%			0.22%	4.47%	0.76%	3.60%	1.42% 0.52%		0.00%	0.87%	0.33%	1.09%	0.65%
NP100	19 19	mid	0.76%	15.37%	26.94%	20.70% 2.59%		0.61%	0.00%	5.18%	0.76%	6.85% 8.88%	1.22% 0.52%	0.76%	0.00%	1.52%	0.00%	0.30%	1.83%
NP102 NP131 NP156	19 19 19	mid mid	1.97%	18.93% 10.59% 14.16%	25.33% 40.11% 35.84%	21.28% 0.26% 13.12% 0.28% 16.69% 0.65%	9.93%	4.87%	1.97%	1.83% 5.72% 2.54%	1.12%	5.53%	0.19% 0.90%	0.39% 1.12% 0.74%	0.00%	0.52% 0.84% 0.65%	0.26% 0.09% 0.16%	0.19% 0.90%	1.41%
NP125 NP128	20 20	mid mid	2.07% 3.98%	18.59%	27.42% 28.35%	11.75% 0.38% 10.84% 0.80%	8.83%	5.15% 7.19%	2.30% 3.17%	6.68%	0.82% 0.84% 1.18%	5.56% 11.14% 6.77%	0.61% 1.23%	0.77%	0.00%	0.54% 0.47%	0.31%	0.69% 0.43%	1.92% 1.09%
NP45	20	mid mid	1.52%	16.90% 26.03%	27.98%	16.27% 0.87%	5.86%	5.42%	1.08%	6.58% 3.04%	1.30%	6.94%	0.43%	0.87%	0.00%	0.43%	0.00%	0.87%	0.87%
NP53 NP147	20 21	mid mid	3.59% 1.52%	11.95% 13.52%	35.61% 27.15%	24.51% 0.51%	18.76%		3.49% 0.90%	1.80% 2.14%	4.05% 0.85%	5.54% 1.07%	0.21% 0.39%		0.00%	0.26% 0.62%	0.10%	0.41% 0.51%	1.28% 1.86%
NP148 NP149	21 21	mid mid	2.53% 2.86%	18.62% 13.79%	19.54% 32.91%	28.58% 0.69% 20.53% 0.18%	9.21%	0.97%	2.30% 2.69%	0.77% 0.53%	1.76% 2.47%	5.59% 7.27%	0.69% 1.81%	0.46% 0.48%	0.00%	0.84% 0.75%	0.08%	0.61% 1.85%	1.07%
NP28 NP101	21 22	mid mid	2.17% 2.26%	17.83% 18.38%	32.17% 25.64%	17.83% 0.50% 19.91% 0.55%	13.12%	4.62%	2.33% 2.22%	0.50% 4.06%	2.33% 1.71%	5.17% 3.60%	0.17% 0.65%	0.28%	0.00%	0.67% 0.46%	0.00% 0.05%	1.00% 0.83%	1.00% 1.66%
NP14 NP132	22 23	mid mid	2.90% 1.10%	10.41% 11.51%	38.52% 34.99%	16.85% 0.86% 10.22% 1.47%			2.68% 2.12%	1.50% 5.80%	1.82% 2.58%	5.58% 4.51%	0.97% 2.30%	0.97% 1.38%	0.00%	0.00% 1.29%	0.43% 0.18%	0.86% 1.20%	1.18% 0.64%
NP26 NP47	23 23	mid mid	2.32% 1.79%	16.49% 15.65%	13.49% 13.50%	23.57% 0.27% 35.36% 0.72%		1.36%	1.23%	3.00% 1.19%	1.77%	13.76% 5.26%	0.95% 0.48%	1.23%	0.00%	0.27% 0.12%	0.00% 0.12%	0.41%	1.50% 1.31%
NP50 NP99	23 23	mid mid	1.62% 2.06%	17.90% 21.90%	22.07% 11.81%	17.77% 1.62% 20.99% 0.57%	11.71% 16.06%		1.08%	6.06% 9.17%	1.62% 1.72%	10.50% 4.93%	0.67% 1.61%	1.21%	0.00%	1.21% 1.61%	0.27% 0.57%	0.81% 0.46%	2.42% 3.21%
NP106 NP58	24 24	mid mid	1.87% 1.66%	15.16% 13.82%	18.59% 28.30%	24.79% 0.24% 19.57% 0.39%	19.92% 10.45%		1.93%	0.72%	3.79%	7.04%	0.90% 1.00%	0.36%	0.00%	0.66% 1.22%	0.06%	0.96%	1.44%
NP72 NP96	24 24	mid mid	2.15%	10.24%	23.11%	22.15% 0.35%			1.87%	1.80% 4.95%	3.11%	6.37% 4.52%	1.80% 0.22%	0.42%	0.00%	0.48%	0.21%	0.55%	1.11%
NP31 NP61	25	mid mid	2.99%	24.31%	29.42% 19.20%	19.19% 0.21% 19.53% 0.49%	10.23%	2.13%	0.64%	0.85%	1.49%	3.20%	0.64% 0.98%	0.64%		0.64%	0.00%	1.49%	1.92%
NP88	25	mid	2.56%	29.14%	21.68%	15.15% 1.17%	9.79%	0.47%	3.03%	3.73%	1.86%	3.96%	0.23%	0.00%	0.00%	1.17%	0.93%	3.26%	1.86%
NP97 NP105	25 26	mid mid	2.02%	29.29% 18.00%	16.16% 23.15%	16.92% 0.00% 23.64% 0.86%	10.48%	4.81%	0.00% 2.87%	5.30%	0.25%	2.27% 5.28%	0.51% 0.53%	1.52% 0.56%	0.00%	0.30%	0.00%	1.01% 0.53% 0.52%	4.55% 1.19%
NP30 NP86	26 26	mid mid	1.15%	10.80%	30.43% 60.99%	20.36% 0.10% 6.46% 0.08%	5.20%	0.87%	3.46% 2.21%	0.63% 1.89%	3.36% 5.67%	5.46% 2.52%	0.31%		0.24%	0.52%	0.00%	0.39%	0.00%
NP89 NP128	26 27	mid mid	3.40% 2.13%	19.96% 15.30%	30.77% 19.75%	12.39% 0.56% 23.30% 0.68%	12.03%	6.27%	2.95% 5.81%	1.44% 1.82%	2.07% 5.95%	3.13%	0.74% 0.64%	1.02% 0.45%	0.23%	0.39% 0.32%	0.07%	0.46% 0.09%	0.77% 2.00%
NP139 NP39	27 27	mid mid	3.86% 1.87%	16.77% 13.14%	15.13% 29.93%	24.04% 0.74% 18.09% 0.24%	15.49%	3.08%	1.93% 2.11%	1.04% 0.81%	1.78% 1.22%	9.94% 8.92%	0.45% 0.73%	1.46%		0.45% 0.57%	0.00% 0.24%	1.19% 0.81%	0.59% 1.22%
NP4 NP118	27 28	mid late	1.85% 2.35%	16.96% 14.04%	17.06% 43.58%	29.82% 0.58% 11.69% 0.65%	5.50%	3.45%	2.05% 2.75%	1.56% 3.00%	4.19% 1.50%	4.48% 6.10%	0.58% 1.30%	0.49% 1.35%	0.05%	0.19% 0.70%	0.00%	0.39% 0.50%	1.07%
NP140 NP129	28 29	late late	2.14% 1.81%	13.34% 13.48%	29.26% 27.86%	23.52% 0.77% 16.02% 0.41%	15.37%	8.05%	1.80%	1.71% 3.04%	0.86% 1.64%	0.86% 7.48%	0.77% 0.66%		0.00%	0.34% 0.41%	0.00% 0.16%	1.45% 0.33%	1.11%
NP136 NP137	29 29	late late	1.14% 1.63%	10.91% 15.07%	27.52% 23.61%	16.37% 1.80% 19.75% 0.30%	21.31%	5.12%	3.30% 2.60%	11.75% 0.82%	1.56% 1.86%	5.70% 3.27%	0.18% 0.67%	0.36% 0.82%	0.30%	0.60% 0.45%	0.42% 0.52%	0.90% 1.04%	0.42% 1.19%
NP5 NP90	29 29	late late	1.68% 3.12%	16.97% 24.97%	29.72% 15.61%	18.17% 0.72% 19.81% 0.97%			3.25% 1.94%	3.85% 2.15%	4.09% 0.32%	3.01% 8.18%	0.84% 1.40%	0.24% 0.75%		0.72% 0.75%	0.36% 0.43%	1.44% 1.72%	1.44%
NP17 NP20	30 30	late late	1.81% 2.21%	11.33% 15.35%	25.98% 42.90%	17.82% 0.00% 11.67% 0.42%			1.36% 0.53%	7.10% 3.58%	3.78% 1.37%	2.57% 4.52%	1.21% 0.84%	0.45% 0.74%		0.30% 0.32%	0.00% 0.21%	0.91% 0.32%	1.36%
NP34 NP73	30 30	late late	2.22% 2.66%	18.95% 20.67%	20.16% 18.01%	25.60% 0.20% 18.92% 1.37%	16.73%	1.61%	1.41%	2.82% 4.33%	1.21%	4.03% 10.94%	0.20%	1.01% 0.30%	0.00%	0.20%	0.20% 0.15%	1.21% 0.76%	2.22% 2.51%
NP81 NP113	30 31	late late	2.05% 1.91%	17.05% 18.41%	23.75% 23.95%	19.70% 0.15% 16.54% 0.49%			1.35% 3.10%	1.65%	1.60% 9.49%	8.20% 2.32%	0.70% 0.65%	0.50%	0.00%	0.30% 0.81%	0.30%	0.95% 0.16%	2.00% 1.06%
NP135 NP25	31 31	late late	1.90%	13.58%	24.09% 31.58%	14.45% 2.48% 15.63% 0.54%	12.99%	5.55%	4.96%	6.42% 5.14%	1.31%	6.42%	0.00% 0.75%	0.58%	0.58%	0.73%	0.15%	3.36% 1.18%	0.44%
NP38	31	late	3.77%	15.53%	33.57%	18.31% 0.18%	8.98%	2.42%	1.71%	4.58%	2.96%	4.76%	0.54%	0.27%	0.00%	0.72%	0.00%	0.45%	1.26%
NP59 NP60	32 32	late late	2.37%	24.54% 16.54%	19.27% 19.77%	23.55% 0.08% 20.34% 0.38%	13.88%	1.90%	2.14%	1.99% 2.85%		1.99%	0.69% 0.57%	0.00%		0.54%	0.08%	0.69%	1.15%
NP71 NP19	32 33	late late	3.59% 3.09%	20.27% 16.67%	18.21% 19.14%	20.84% 0.67% 16.67% 1.23%	11.11%	5.86%	6.28% 3.70%	2.40% 6.48%	1.29% 2.78%	3.88% 8.33%	0.53% 0.62%		0.31%	0.57% 0.31%	0.05% 0.62%	1.01% 1.54%	0.96% 1.23%
NP27 NP63	33 33	late late	1.33% 3.22%	12.09% 18.56%	29.41% 23.71%	25.41% 0.41% 22.00% 0.52%	13.52%	2.34%	1.43% 2.39%	3.59% 1.35%	0.92% 1.46%	3.59% 5.36%	0.82% 0.57%		1.20%	0.41% 0.62%	0.51% 0.26%	0.61% 0.99%	0.82% 1.40%
NP83 NP37	33 34	late late	3.58% 2.13%	15.69% 13.55%	17.59% 31.09%	26.93% 0.58% 13.37% 0.00%	8.86%	1.24%	2.92% 1.51%	2.34% 3.99%	1.53% 4.07%	3.50% 7.97%	0.66% 0.62%	0.71%		0.51% 0.18%	0.07%	0.88% 1.15%	1.82% 0.97%
NP43 NP8	34 34	late late	1.70% 2.37%	16.27% 14.88%	27.72% 21.74%	12.58% 0.38% 18.54% 0.19%	11.80%	3.59%	2.74%	5.87% 5.32%	3.60% 4.62%	6.43% 9.17%	0.66% 1.28%		0.19%	0.95% 0.19%	0.28%	1.51% 0.58%	1.80% 2.05%
NP112 NP127	35 35	late late	2.63% 1.44%	17.94% 13.83%	20.26% 33.35%	16.27% 0.80% 19.14% 0.57%			1.99% 3.78%	3.59% 1.10%		10.29% 4.93%	1.83% 0.67%	0.56% 0.24%		0.32% 0.77%	0.56% 0.43%	0.96% 0.72%	1.12% 2.15%
NP40 NP44	35 35	late late	3.90% 2.73%	17.14% 14.77%	25.97% 30.91%	14.81% 0.26% 17.95% 0.23%	10.39% 12.50%	2.86% 2.05%	3.38% 2.27%	4.42% 3.41%	2.60% 1.36%	10.13% 6.59%	0.26% 0.45%	0.26% 0.45%	0.00%	0.00% 0.91%	0.00%	2.34% 2.27%	1.30% 1.14%
NP32 NP35	36 36	late late	0.44% 1.45%	10.19% 12.40%	33.92% 29.34%	10.63% 0.53% 15.29% 0.41%	14.59%	0.53%	1.23%	5.71% 4.55%	6.41%	10.98% 15.50%	0.79% 0.21%	0.53%		0.88% 0.62%	0.26% 0.83%	0.26%	1.58% 1.24%
NP48 NP66	36 37	late late	1.76% 1.45%	13.56% 13.41%	22.54% 47.74%	17.96% 0.18%	14.08%	4.40%	1.41% 1.81%	4.23% 1.72%		11.97%	0.70% 0.45%	0.88% 2.54%	0.35%	0.18% 0.54%	0.18%	0.70% 0.82%	2.11% 0.72%
NP69 NP7	37 37	late late	0.28% 3.46%	17.20% 14.90%	17.34% 15.33%	30.65% 0.55% 22.68% 0.22%	15.95%	5.55%	0.83% 2.81%	3.47% 4.54%	0.42%	3.88% 12.31%	1.11% 0.65%	0.42% 0.43%	0.00%	0.42% 0.22%	0.14% 0.00%	0.55% 1.30%	1.25% 2.59%
NP9 NP155	37 38	late late	1.77% 2.80%	16.59% 18.53%	26.11% 38.02%	25.66% 0.00% 11.89% 0.26%	17.48%	2.88%	0.88%	2.88% 2.36%	1.11%	2.21%	0.22% 0.79%	0.44% 1.40%	0.00%	0.00%	0.66% 0.61%	0.22%	0.88% 1.75%
NP41 NP10	38 39	late late	1.56% 2.93%	21.25% 16.41%	36.06% 31.45%	14.62% 0.78% 19.73% 1.56%	9.36%		0.39%	4.48% 1.37%	0.58%	5.07% 8.40%	0.19% 0.59%	0.58% 1.17%	0.00%	0.00%	0.19%	1.56% 0.78%	2.92% 0.98%
NP114 NP159	39 39	late late	3.59% 3.10%	21.16% 20.89%	29.06% 21.92%	17.35% 0.53% 16.38% 0.22%	10.33%	5.76%	2.53% 1.99%	1.55% 1.62%	1.73% 0.74%		0.62% 1.11%	0.84% 0.52%	0.00%	0.44% 0.15%	0.22% 0.15%	0.80% 0.96%	2.17% 2.14%
NP67 NP91	39 39	late late	2.77%	13.61% 19.17%	24.33% 33.28%	21.96% 0.35% 15.45% 0.59%	16.60%	1.84%	3.47% 1.90%	1.54%	4.26%	3.73% 2.75%	0.97% 0.94%	0.83%	0.53%	0.57%	0.40%	0.57%	1.67%
NP158	40	late	2.07%	17.97%	28.57%	22.00% 0.12%			1.15%	1.61%	2.19%		0.58%	0.46%		0.69%	0.12%	1.73%	2.42%

$Supplementary\ Table\ 3A.\ The\ number\ of\ single\ cells\ and\ pseudocells\ in\ the\ training\ dataset\ and\ the\ independent\ testing\ dataset,$ with the number of features selected by Lasso

C II.	T 4 1 1 11 (N)	Training	g dataset	Independent t	esting dataset		
Cell type	Total single cells(N)	Single cells (N)	Pseudocells (N)	Single cells (N)	Pseudocells (N)	Selected Features by Lasso (N)	
CD8+ cytotoxic T	47043	32930	3234	14113	131	1615	
CD8+ Naive T	31258	21880	2131	9378	131	1335	
CD4+ Naive T	30180	21126	2055	9054	131	1284	
dnT	21240	14868	1427	6372	131	1044	
NK	10875	7612	710	3263	131	600	
MAIT	6775	4742	434	2033	131	447	
Monocyte	4490	3143	283	1347	131	309	
Memory B cell	4460	3122	290	1338	129	325	
Naive B cell	4426	3097	295	1328	129	275	
CD4+ Cytotoxic T	4348	3043	271	1305	131	286	
Treg	2587	1810	171	777	129	185	
NK_CD56bright	1431	995	125	427	120	139	
NKT	1372	957	130	411	123	132	
Proliferative T	1268	884	131	380	126	147	

Supplementary Table 3B. RMSE values calculated by 14 cell-type-specific models with the optimal hyperparameters determined by the 5-fold cross validation in the training dataset

Call tune		RMSE				
Cell type	n_estimators	min_samples_split	min_samples_leaf	max_depth	mean	std
CD8+ cytotoxic T	100	2	2	190	14.730	0.407
CD8+ Naive T	100	2	1	65	14.871	0.327
CD4+ Naive T	100	2	1	65	15.293	0.372
dnT	70	2	2	195	17.097	0.781
NK	100	2	1	65	15.975	0.328
MAIT	100	2	2	190	20.134	1.826
Monocyte	100	2	1	15	21.252	1.495
Memory B cell	80	5	1	165	23.483	1.612
Naive B cell	55	5	2	30	25.051	1.414
CD4+ Cytotoxic T	100	2	1	65	20.875	0.728
Treg	85	5	2	175	25.349	0.996
NK_CD56bright	85	5	2	175	25.871	3.671
NKT	30	2	2	160	26.990	3.161
Proliferative T	60	2	4	20	33.395	2.372

^{*}hyperparameter space:

n_estimators = [10, 20, 30, ..., 100], max_depth = [10, 15, 20, ..., 205],

min_samples_split = [2, 5, 10], min_samples_leaf = [1, 2, 4],

All other hyperparameters were configured with default values in the scikit-learn libaray.