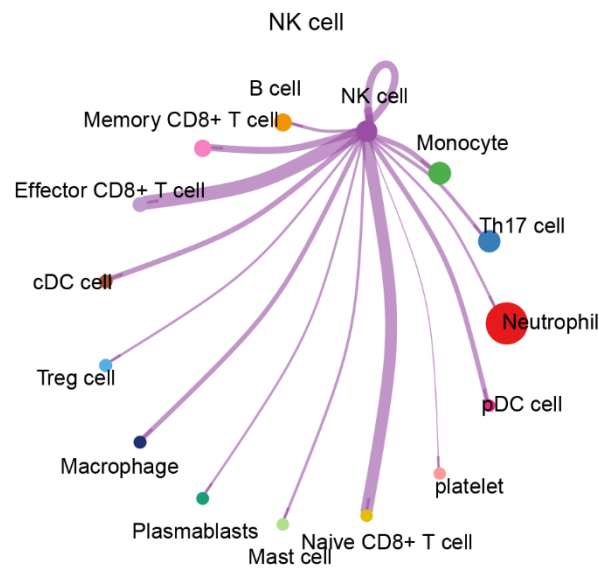
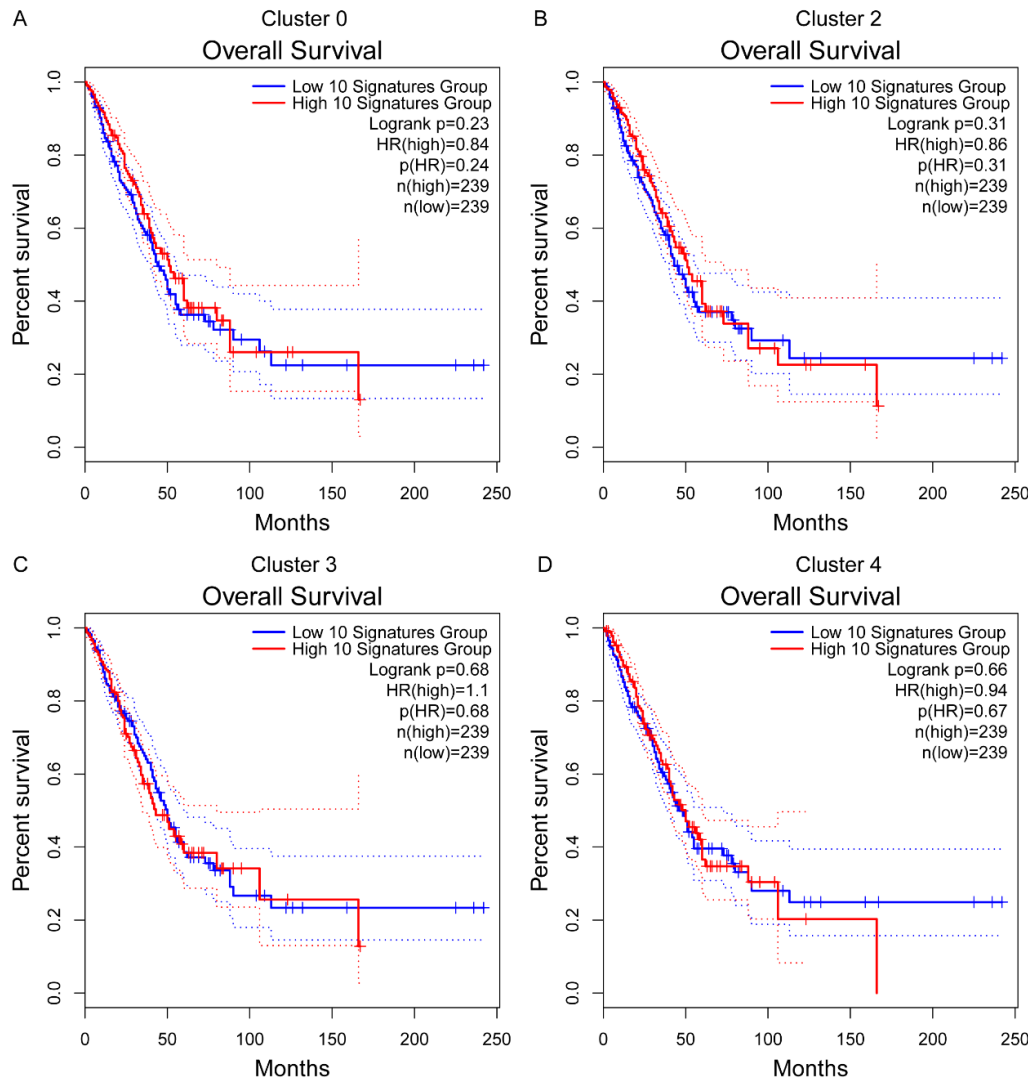


Supplementary figure 1. The weight of NK cell communication with other immune subsets.



Supplementary figure 2. Kaplan-Meier survival curves of high vs. low expression of 10 prognostic signatures in cluster 0, 2, 3, 4.



Supplementary figure 3. Longitudinal copy number variation (CNV) landscape in matched baseline and progressive tumor samples.

	Chr1	Chr4	Chr6	Chr7	Chr8	Chr9	Chr10	Chr12	Chr14	Chr15	Chr17	Chr19	Chr20	Chr21	Chr22
Gain		PDGFRA	ROS1	EGFR	CCNE2			CCND2			CDK12		AURKA	ERG	
								ARID2			ERBB2		GNAS	TMPRSS2	
Loss	BLC9				MYC	PSIP1	TET1	ERBB3	HIF1A	BUB1B	BRCA1	KEAP1			EP300
					PPP2R2A	CDKN2A	PTEN				TP53	DNM2			
					FGFR1	CDKN2B	FAS				PER1	CALR			
						JAK2									
						CD274									
					PDCD1LG2										
	Baseline	Disease progression													

Supplementary figure 4. Gene Set Enrichment Analysis (GSEA) of immune-related pathways.

