

**Table S1:GEO datasets information**

GEO ID	Platform	Organism	Tissue	Expression Profiling	Samples size	Recipient age (mean ± SD)	Recipient gender (M/F)	Donor source							PGF	DGF	IGF	Average follow-up time
								LD	DBD	DCD	AKI	non-AKI	Nephrectomies					
GSE21374	GPL570	Homo sapien	NM kidney biopsy	Microarray	282	NM	65/40	NM	NM	NM	NM	NM	NM	NM	NM	NM	NM	774
GSE43974	GPL10558	Homo sapien	NM kidney biopsy	Microarray	554	NM		NM	27	82	38	NM	NM	NM	NM	NM	NM	NM
GSE37838	GPL570	Homo sapien	NM kidney biopsy	Microarray	78	53±13	43/27	NM	53	NM	NM	NM	NM	8	NM	NM	NM	NM
GSE10419	GPL887	Homo sapien	NM kidney biopsy	Microarray	30	48±11		NM	15	NM	14	NM	NM	NM	NM	NM	NM	NM
GSE90861	GPL18573	Homo sapien	NM kidney biopsy	RNA-Seq	46	54±10		16/7	NM	16	7	NM	NM	NM	NM	NM	11	12
GSE53769	GPL16686	Homo sapien	NM kidney biopsy	Microarray	36	48±11	12/22	NM	NM	NM	16	NM	NM	20	NM	NM	NM	NM
GSE61739	GPL13158	Homo sapien	NM kidney biopsy	Microarray	96	53±13		NM	NM	NM	NM	48	48	NM	NM	NM	NM	NM
GSE30718	GPL570	Homo sapien	NM kidney biopsy	Microarray	47	NM		NM	NM	NM	NM	28	NM	8	NM	NM	NM	NM

DBD, donors after brain death; DCD, donors after cardiac death; LD, living donors; AKI, acute kidney injury; PGF, primary graft function; DGF, delayed graft function; IGF, immediate graft function; NM, not mentioned

**Table S2: GEO datasets information of four subclasses**

	GSE10419	GSE21374	GSE30718	GSE37838	GSE43974	GSE53769	GSE61739	GSE90861
PCD-A	7	114	6	15	48	12	32	11
PCD-B	8	23	0	17	92	0	13	14
PCD-C	7	49	3	21	23	2	16	4
PCD-D	7	96	2	17	40	4	35	17

**Table S3: The RiskScore formula**

PCD pattern	Formula	
Lysosome-dependent cell death	-0.959*CTNS+1.475*CTSO+1.22*DNASE2+0.855*GGA3+1.033*HEXB+1.596*STXBP1 -1.407*CPTP+1.45*FEZ2+-0.788*KLHL3+1.931*NOD1+-1.444*ORMDL3	
Autophagy	3.41*ITGA6+0.862*ITM2C+-1.895*MADD+2.592*POLB+0.869*TGFB1+1.363*TNFRSF12A	
Apoptosis	0.844*BAX+-1.128*BRAF+3.244*ITGA6+1.182*ATF4+1.817*PIK3R3+0.703*MUC1+-0.426*NQO1+3.089*EEF1A1+1.003*RHOB	
Anoikis	-0.241*PCK2+-1.426*MTOR+0.269*GZMB+1.606*CTNNA1	
Entosis	Methoisis	1.984*RAC1+-1.108*GIT1+-1.597*MTOR
Paraptosis	-1.41*TAAR5+2.074*NT5C+-0.794*SSTR3+-0.814*ATP23+0.287*CFD+0.538*XBP1+0.881*TNFRSF19	
Immunogenic cell death	1.318*BAX+0.639*CD8A+0.757*EIF2AK3+3.642*HMGB1+-0.922*IFNG+0.947*IL1B+-0.791*IL6+1.095*NT5E	
NETosis	1.393*HMGB1+-1.132*AGER+0.556*TGFB1	
Oxeiptosis	-1.305*KEAP1+-0.415*AIFM1	
Entoticcelldeath	1.797*CTNNA1+0.377*CYBB	
Parthanatos	-0.847*MIF+-0.551*OGG1	
Cuproptosis	-0.655*GLS+0.976*DBT	
Necroptosis	-1.385*FTH1+1.234*BAX+1.983*CAPN2+0.78*IL1B+1.796*STAT6+-1.886*VDAC2+-3.169*CHMP7	
Ferroptosis	1.85*FANCD2+1.533*G6PD+-0.421*NQO1+-1.363*PCBP2+-0.954*PEBP1+-2.775*PGD+-0.964*TFRC	
Pyroptosis	0.827*BAX+1.258*CHMP4C+-4.006*CHMP7+2.663*HMGB1+-1.059*IL1A+1.983*NOD1+2.086*PLCG1+-0.836*PRKACA+1.205*TNF	