Gene	P value	Hazard Ratio(95% CI)	
PDLIM4	< 0.001	1.218(1.083–1.37)	→
TIMP1	< 0.001	1.246(1.096-1.418)	1-0-4
FAM20C	< 0.001	1.331(1.127–1.572)	
STC1	< 0.001	1.352(1.16–1.576)	▶
FOSL1	< 0.001	1.374(1.15–1.642)	
MXRA8	< 0.001	1.46(1.193–1.786)	
OSMR	< 0.001	1.468(1.233-1.748)	
PLAUR	< 0.001	1.477(1.212-1.801)	——
C9orf64	< 0.001	1.559(1.203-2.021)	
ARSD	< 0.001	1.668(1.25-2.224)	
FES	< 0.001	1.698(1.262-2.286)	•
VDR	< 0.001	1.702(1.276-2.27)	——
PDIA4	< 0.001	1.708(1.287-2.268)	
SLC43A3	< 0.001	1.764(1.289-2.415)	•
GNS	< 0.001	1.793(1.283-2.507)	•
GZMB	< 0.001	1.917(1.395-2.634)	•
SH2D2A	< 0.001	1.922(1.331-2.775)	
SLC16A13	< 0.001	2.257(1.398-3.645)	•
KRT80	< 0.001	2.925(1.678-5.099)	•
			0 1 2 3 4 5 6 Hazard Ratio