Gene	P value	Hazard Ratio(95% CI)	
GPC2	< 0.001	0.675(0.552-0.826)	
PLXNB1	< 0.001	0.716(0.624-0.821)	
CASKIN1	< 0.001	0.739(0.625-0.874)	
DOC2A	< 0.001	0.756(0.65-0.88)	
C14orf37	< 0.001	0.757(0.657-0.873)	
CCNJL	< 0.001	0.795(0.703-0.9)	
PALM	< 0.001	0.809(0.724-0.903)	
CYP2E1	< 0.001	0.815(0.728-0.913)	
KIF26A	< 0.001	0.832(0.755-0.916)	+
STOX2	< 0.001	0.836(0.763-0.916)	
HTR7	< 0.001	1.18(1.09–1.28)	
SORT1	< 0.001	1.2(1.09–1.33)	•••
BVES	< 0.001	1.21(1.1–1.34)	
METTL7B	< 0.001	1.22(1.11–1.35)	
ITGAX	< 0.001	1.31(1.13–1.52)	• • • • •
GNGT2	< 0.001	1.37(1.16–1.62)	
JPH4	< 0.001	1.39(1.16–1.66)	
TFEB	< 0.001	1.45(1.2–1.75)	
S100A4	< 0.001	1.47(1.21–1.77)	
MVP	< 0.001	1.64(1.25–2.15)	
			0 0.2 0.4 0.6 0.8 1 1.2 1.4 1.6 1.8 2 2.2 Hazard Ratio