Gene	P value	Hazard Ratio(95% CI)	
SLC31A2	< 0.001	0.506(0.356-0.719)	
OCLN	< 0.001	0.593(0.457-0.769)	10-1
MAN2A1	< 0.001	0.652(0.528-0.804)	
MANSC1	< 0.001	0.674(0.577-0.787)	
COBLL1	< 0.001	0.693(0.592-0.811)	
EGR3	0.029	0.859(0.749-0.984)	
EGR1	0.001	0.883(0.819-0.951)	
CCND1	0.034	0.952(0.909-0.996)	+
SCN9A	0.014	1.19(1.04–1.38)	I →1
DNAJB4	0.041	1.22(1.01–1.48)	
HSPB8	0.036	1.23(1.01–1.49)	
SCN3A	< 0.001	1.23(1.11–1.36)	
ZBED1	0.004	1.39(1.11–1.73)	
NTF3	0.005	2(1.24–3.23)	
SOX11	< 0.001	2.65(2.08-3.4)	
RGS7	0.002	3.71(1.6-8.63)	
		•	0 1 2 3 4 5 6 7 8 9 Hazard Ratio