

	pvalue	Hazard ratio
SLC16A1-AS1	<0.001	3.781(2.310–6.189)
PCED1B-AS1	<0.001	1.184(1.096–1.278)
MMP25-AS1	<0.001	1.315(1.151–1.504)
LINC00894	<0.001	2.991(2.069–4.324)
DUXAP8	<0.001	4.211(2.554–6.943)
LINC01426	<0.001	1.084(1.047–1.123)
AL022322.1	<0.001	1.272(1.129–1.434)
ASMTL-AS1	<0.001	1.098(1.054–1.144)
AC005785.1	<0.001	1.589(1.299–1.945)
AC079322.1	<0.001	5.491(2.823–10.679)
PVT1	<0.001	1.287(1.184–1.399)
AL031714.1	<0.001	2.116(1.472–3.042)
USP30-AS1	<0.001	1.169(1.066–1.282)
MIR155HG	<0.001	1.105(1.061–1.152)
AC107021.2	0.001	1.128(1.048–1.213)
LINC01355	<0.001	1.874(1.561–2.250)
TRG-AS1	<0.001	1.681(1.277–2.211)
FAM13A-AS1	<0.001	1.397(1.158–1.685)
AL021707.6	0.001	1.115(1.045–1.190)
LINC00893	<0.001	1.686(1.281–2.220)
PELATON	<0.001	1.212(1.082–1.358)
AC007098.1	<0.001	2.126(1.479–3.056)
AL137186.2	<0.001	2.169(1.594–2.952)
LINC02609	<0.001	1.465(1.226–1.751)
HLA-DQB1-AS1	0.003	1.045(1.015–1.075)
ARHGAP27P1-AS1	<0.001	1.359(1.159–1.584)
MYG1-AS1	<0.001	1.752(1.511–2.031)
PRKAR1B-AS1	<0.001	1.186(1.104–1.274)
MALAT1	<0.001	1.011(1.005–1.016)
LINC00342	<0.001	1.263(1.167–1.367)

