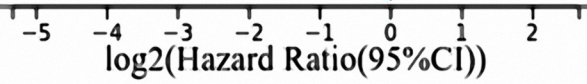CancerCode	pvalue	Hazard Ratio(95%CI)
TCGA-ACC(N=77)	6.8e-4	2.36(1.42,3.92)
TCGA-SARC(N=254)	0.02	1.35(1.06,1.72)
TCGA-KIRP(N=276)	0.02	1.65(1.08,2.54)
TCGA-LUAD(N=490)	0.08	1.15(0.99,1.34)
TCGA-KICH(N=64)	0.08	2.26(0.92,5.54)
TCGA-PAAD(N=172)	0.12	1.30(0.93,1.81)
TCGA-COAD(N=278)	0.15	1.28(0.91,1.79)
TCGA-UCS(N=55)	0.19	1.34(0.86,2.09)
TARGET-NB(N=151)	0.25	1.16(0.90,1.48)
TCGA-HNSC(N=509)	0.27	1.11(0.92,1.32)
TCGA-DLBC(N=44)	0.27	1.93(0.59,6.29)
TCGA-LAML(N=209)	0.28	1.13(0.90,1.41)
TCGA-LIHC(N=341)	0.31	1.12(0.90,1.40)
TCGA-BLCA(N=398)	0.32	1.10(0.91,1.32)
TCGA-MESO(N=84)	0.32	1.19(0.84,1.68)
TCGA-PRAD(N=492)	0.35	1.71(0.56,5.20)
TARGET-ALL(N=86)	0.35	1.23(0.80,1.90)
TARGET-LAML(N=142)	0.37	1.16(0.84,1.59)
TCGA-SKCM-P(N=97)	0.38	1.23(0.78,1.93)
TCGA-OV(N=407)	0.52	1.05(0.91,1.20)
TCGA-ESCA(N=175)	0.53	1.11(0.81,1.52)
TCGA-COADREAD(N=368)	0.54	1.10(0.81,1.51)
TCGA-THCA(N=501)	0.54	1.31(0.55,3.13)
TCGA-UVM(N=74)	0.74	1.12(0.58,2.15)
TARGET-WT(N=80)	0.78	1.08(0.63,1.87)
TCGA-STES(N=547)	0.87	1.01(0.85,1.22)
TCGA-PCPG(N=170)	0.96	1.03(0.35,3.02)
TCGA-STAD(N=372)	1.00	1.00(0.80,1.25)
TCGA-LGG(N=474)	6.9e-7	0.56(0.44,0.70)
TCGA-GBMLGG(N=619)	8.9e-4	0.75(0.63,0.89)
TCGA-LUSC(N=468)	0.04	0.86(0.75,0.99)
TCGA-READ(N=90)	0.15	0.55(0.24,1.25)
TCGA-UCEC(N=166)	0.17	0.75(0.50,1.13)
TCGA-KIPAN(N=855)	0.17	0.90(0.77,1.05)
TCGA-KIRC(N=515)	0.17	0.87(0.71,1.06)
TCGA-TGCT(N=128)	0.24	0.25(0.02,2.61)
TCGA-BRCA(N=1044)	0.26	0.89(0.74,1.09)
TCGA-GBM(N=144)	0.34	0.85(0.62,1.18)
TCGA-THYM(N=117)	0.41	0.61(0.19,1.95)
TCGA-SKCM(N=444)	0.42	0.93(0.78,1.11)
TCGA-SKCM-M(N=347)	0.44	0.93(0.76,1.12)
TCGA-CESC(N=273)	0.47	0.89(0.65,1.21)
TCGA-CHOL(N=33)	0.72	0.84(0.34,2.11)
TARGET-ALL-R(N=99)	0.76	0.96(0.72,1.27)