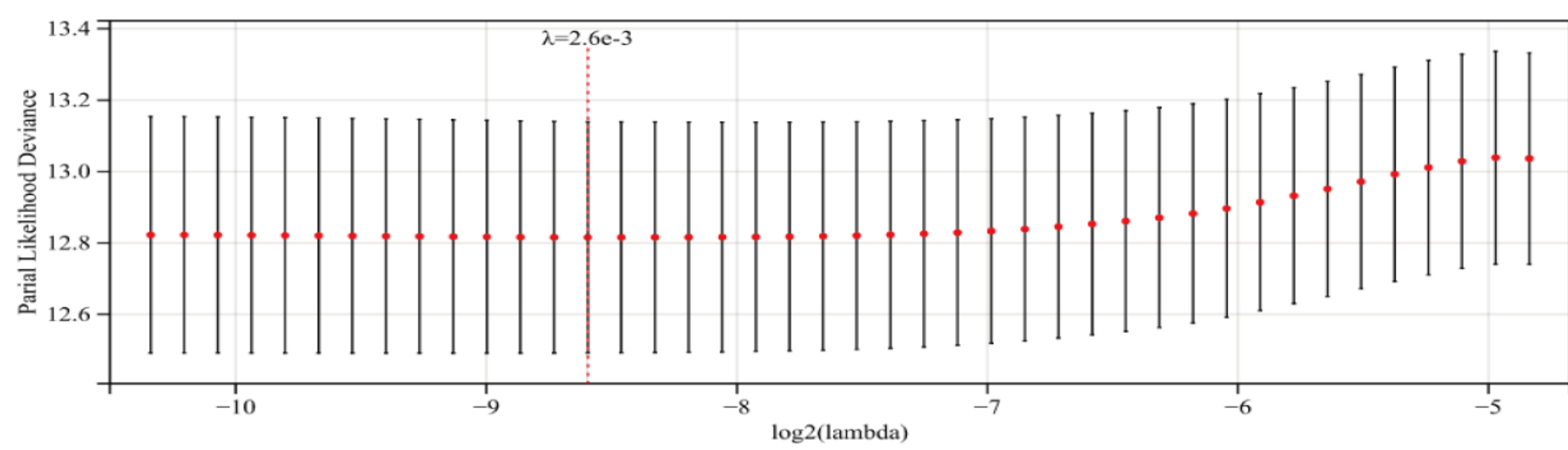
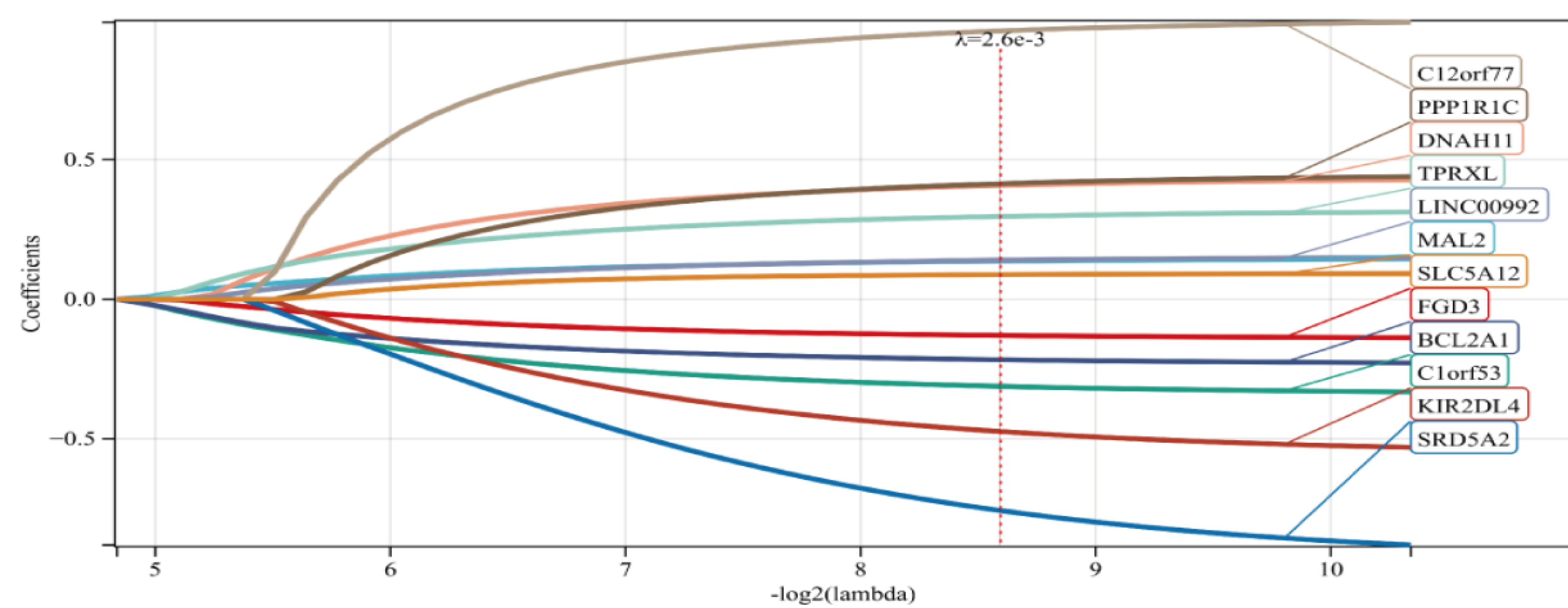
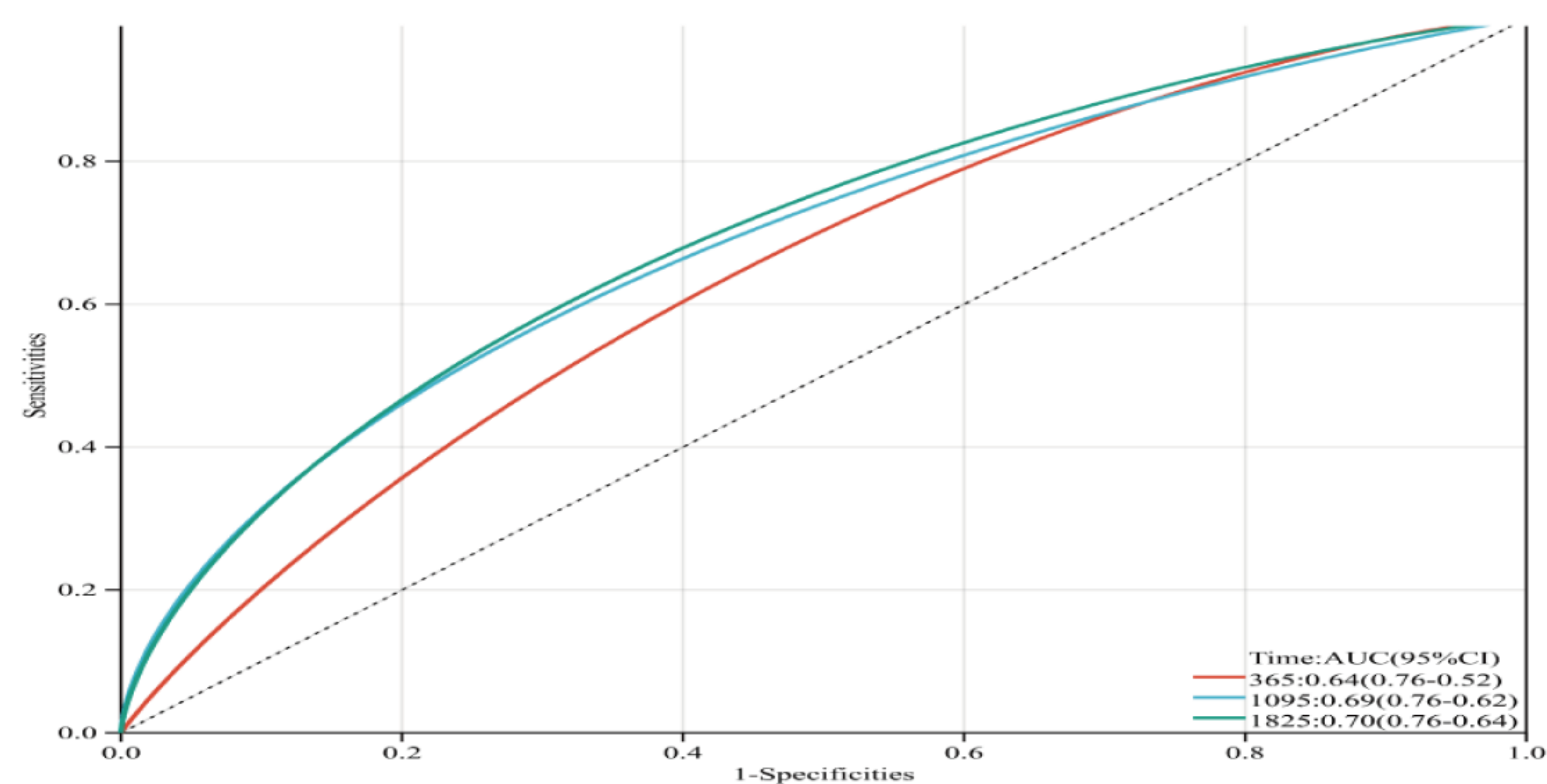
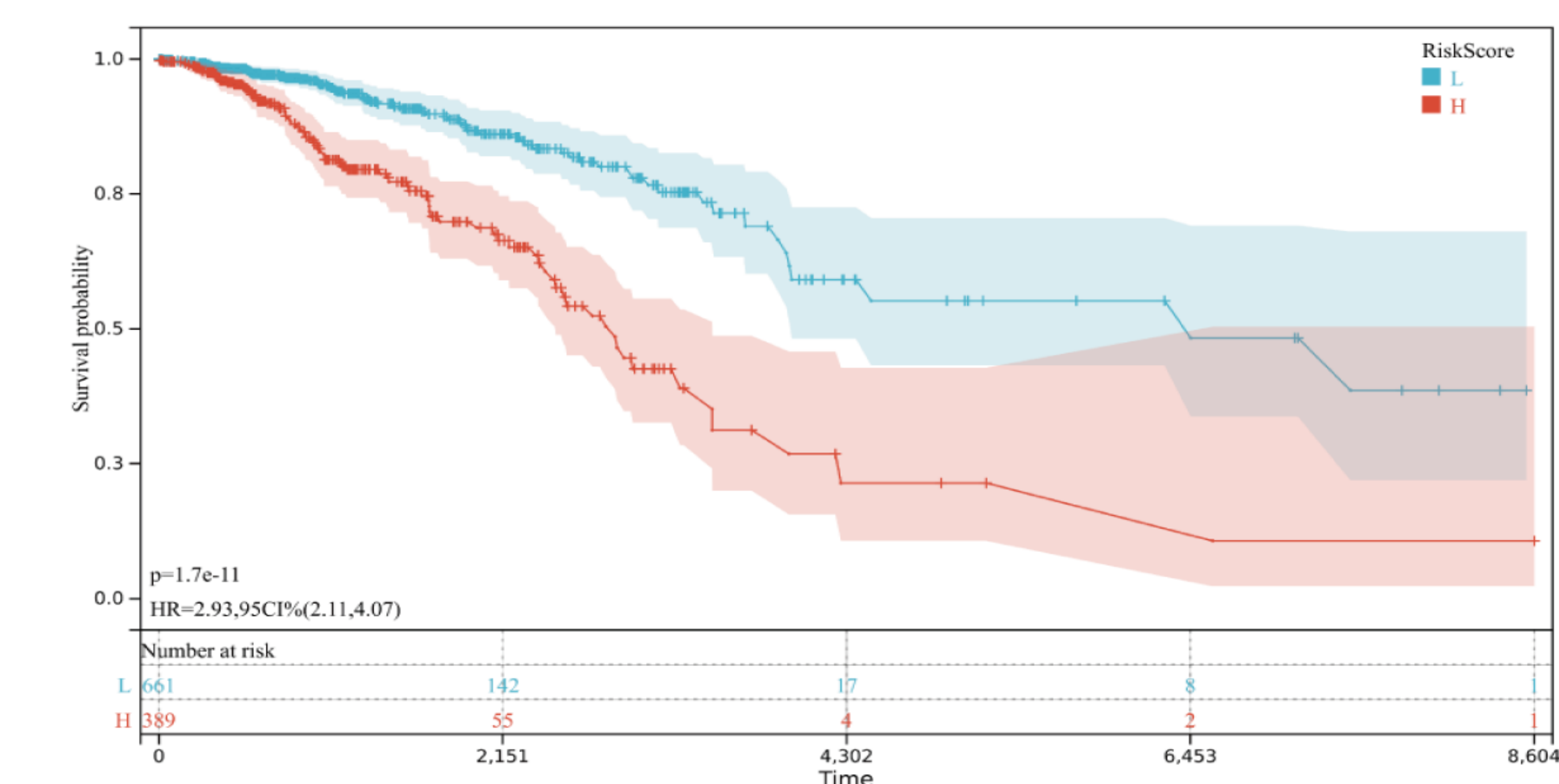


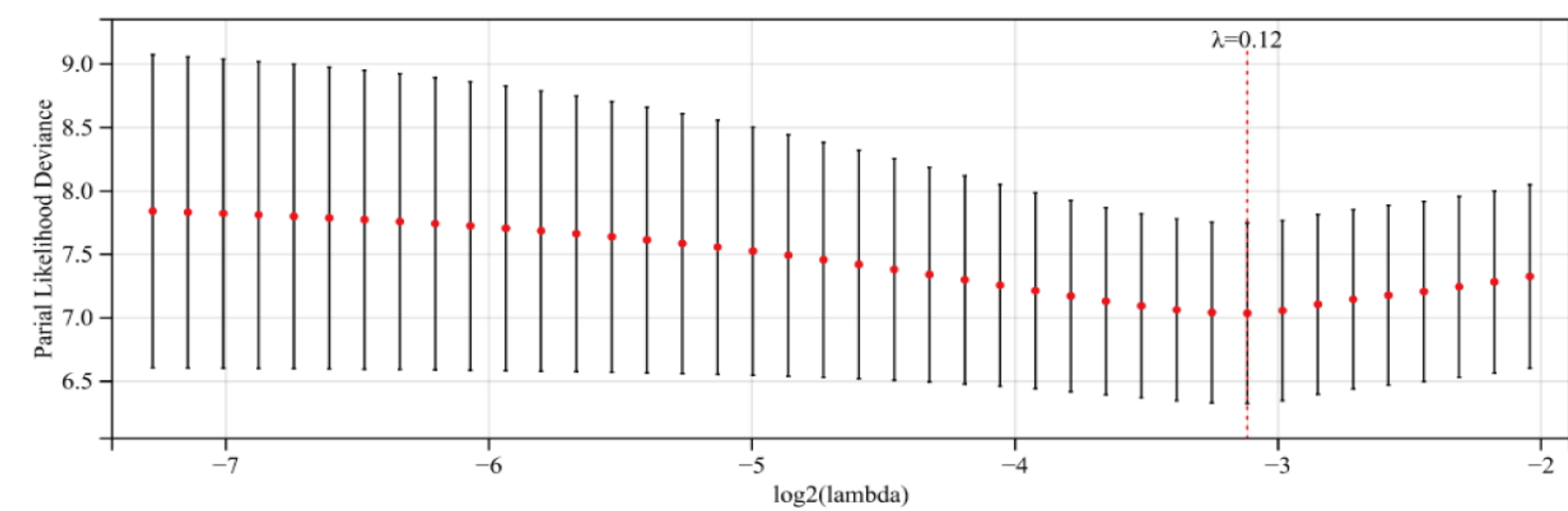
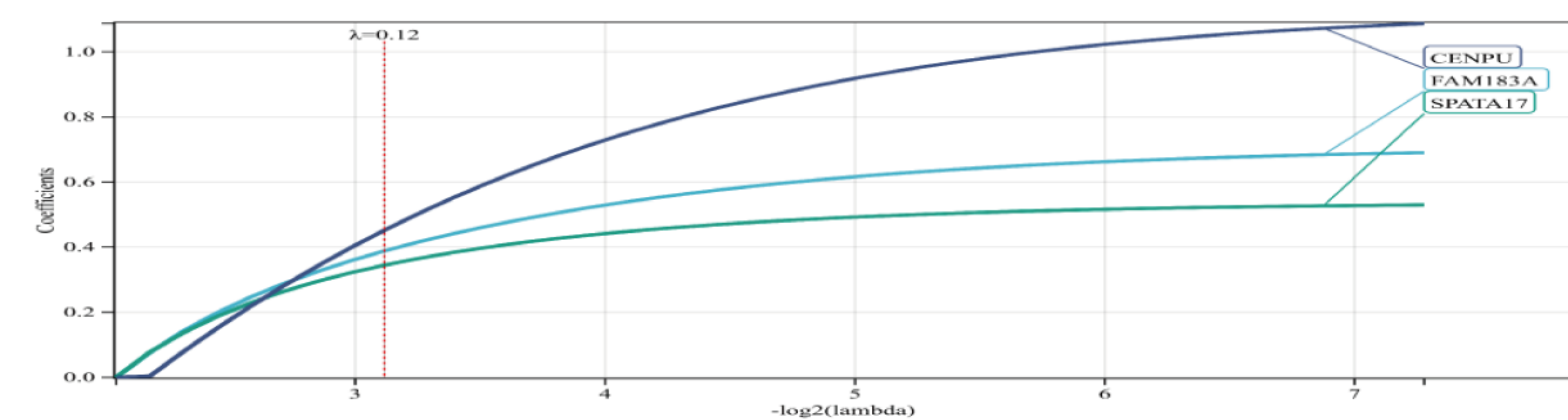
BRCA



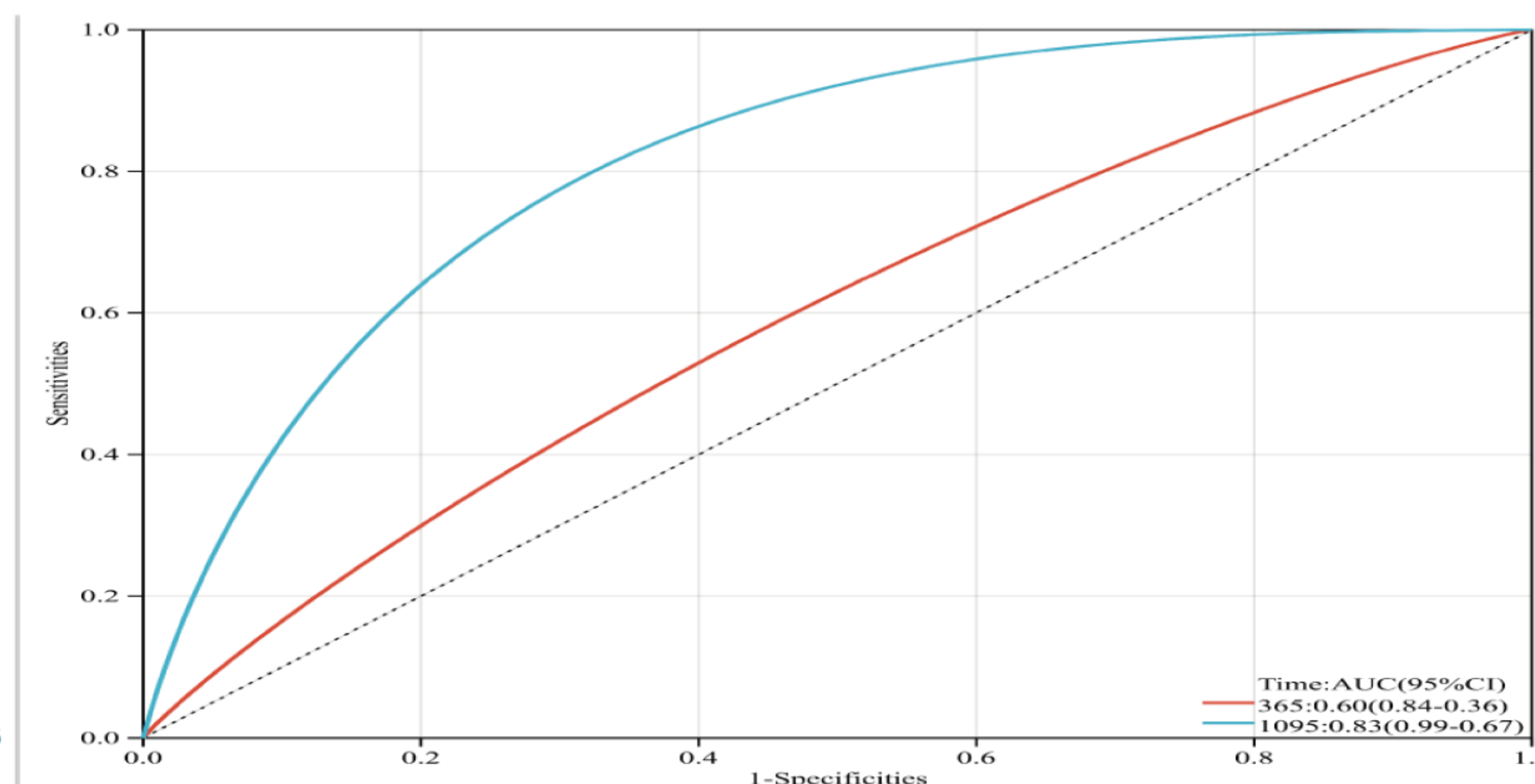
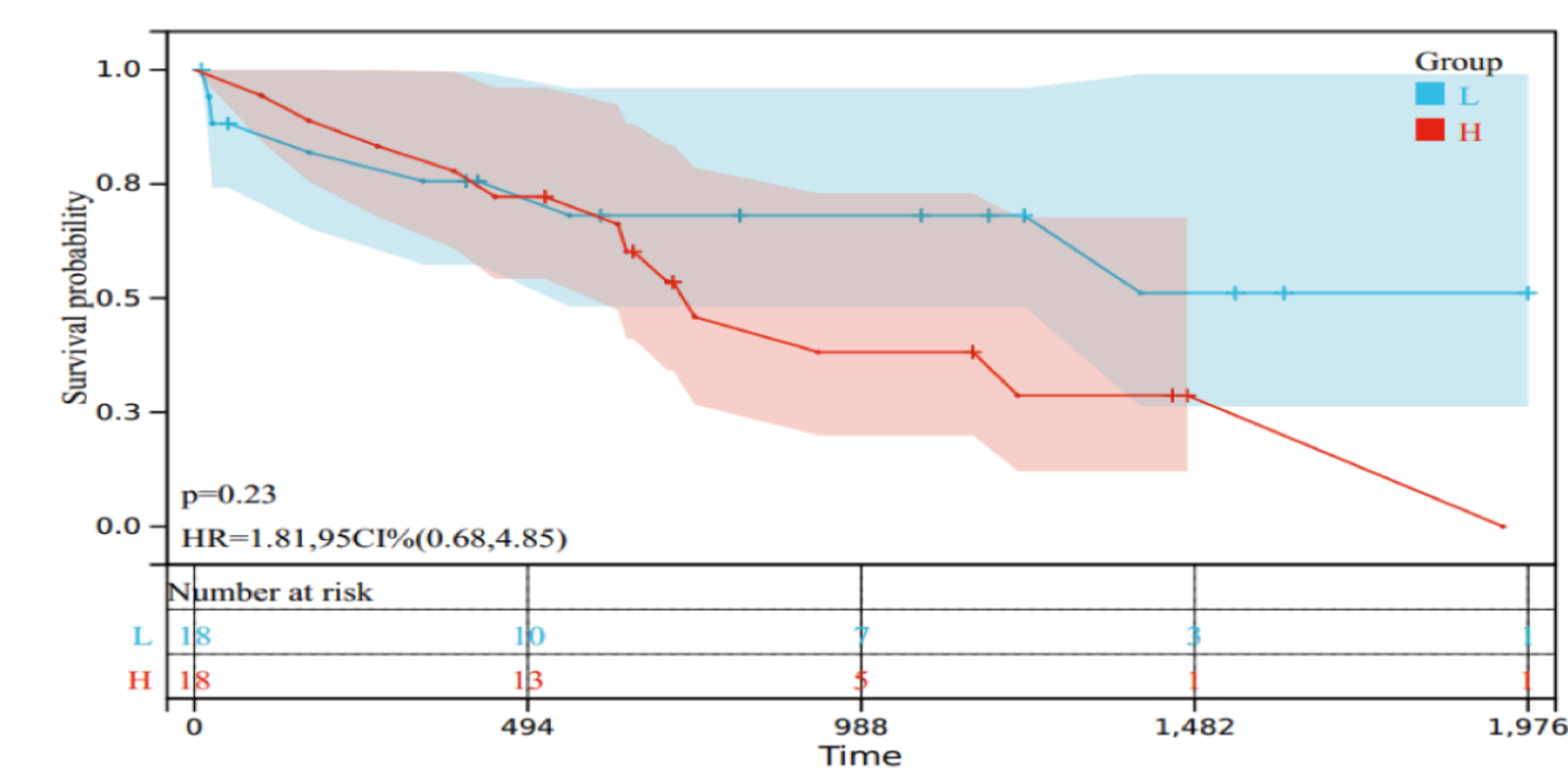
Features	p-value	Hazard Ratio(95%CI)
C1orf53	4.0e-3	0.71(0.56-0.90)
TPRXL	0.02	1.37(1.06-1.78)
PPP1R1C	0.02	1.57(1.08-2.28)
BCL2A1	0.03	0.79(0.64-0.97)
FGD3	0.03	0.87(0.76-0.99)
DNAH11	0.04	1.54(1.02-2.35)



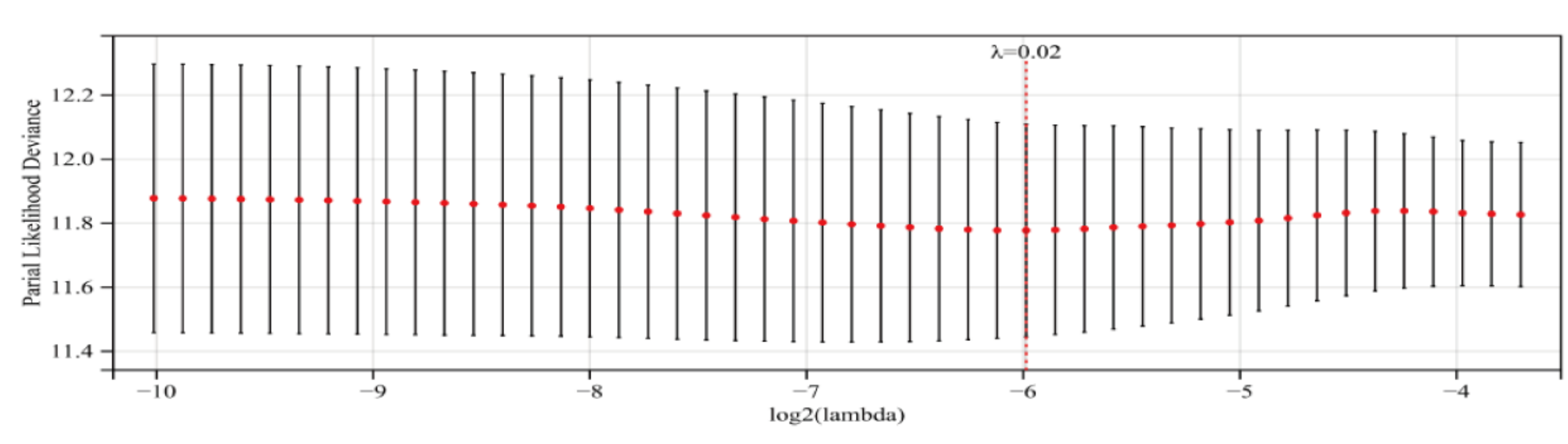
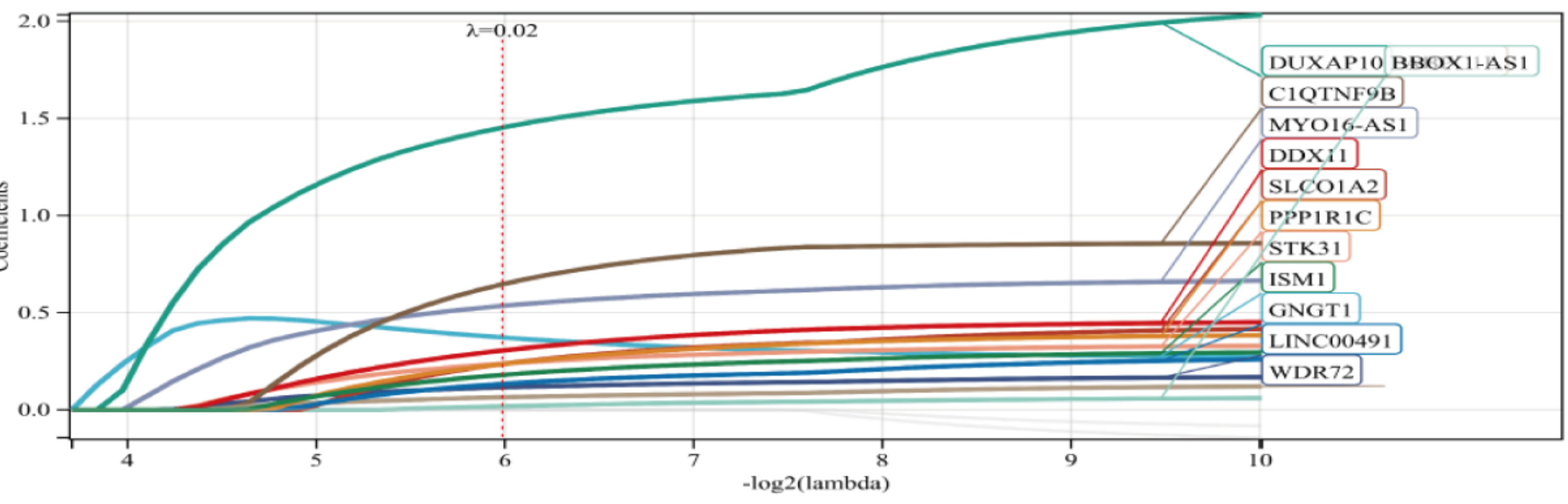
CHOL



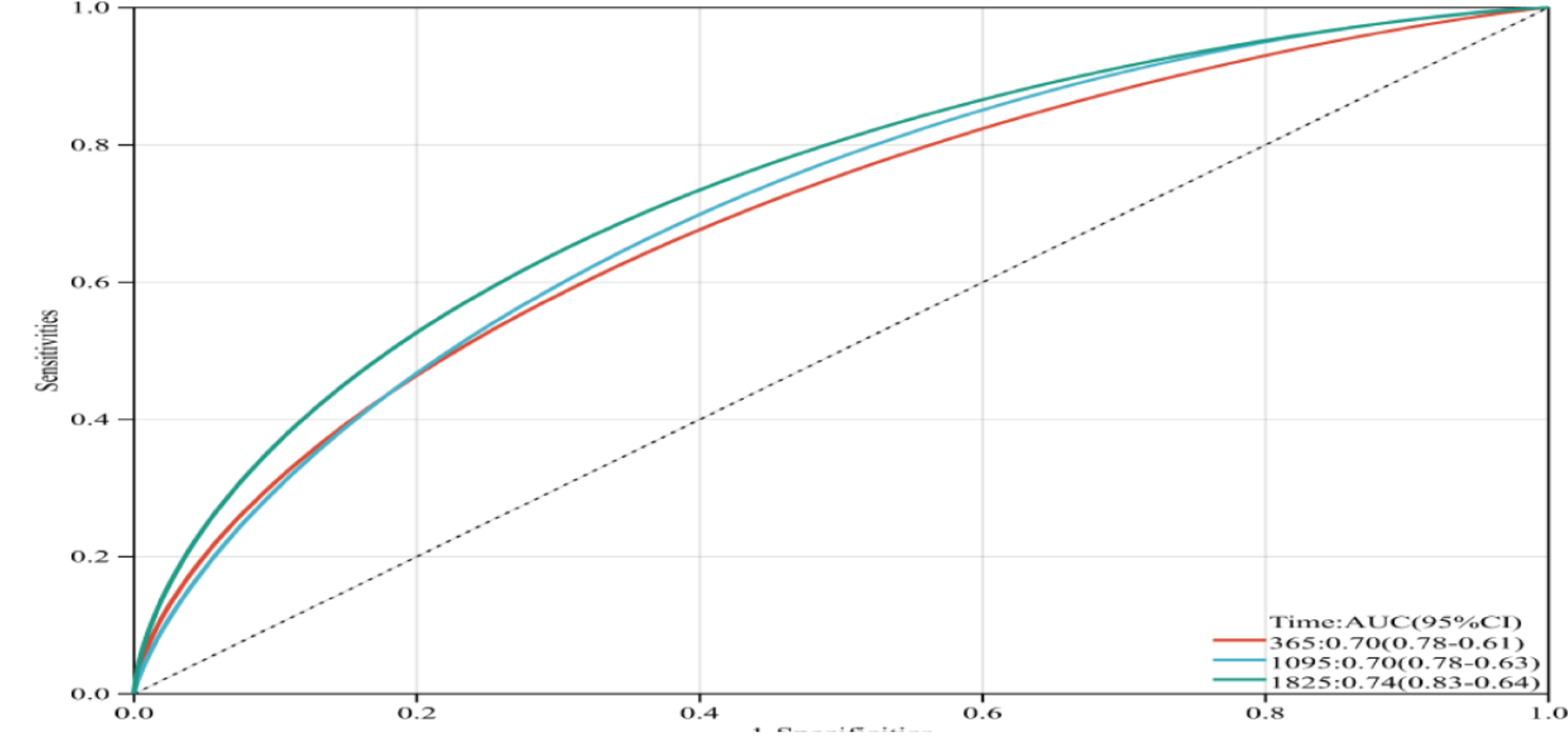
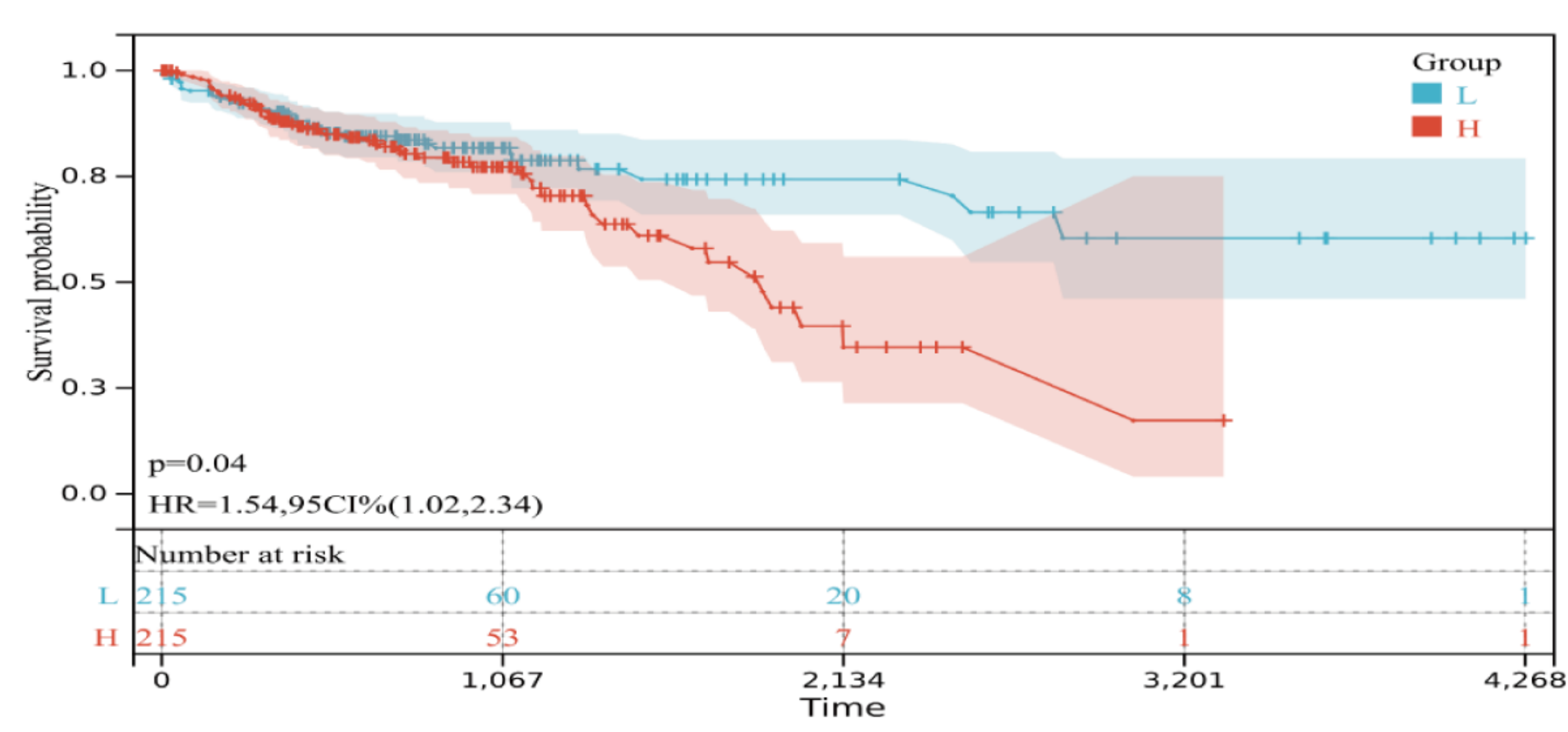
Features	p-value	Hazard Ratio(95%CI)
CENPU	0.04	3.11(1.04-9.27)



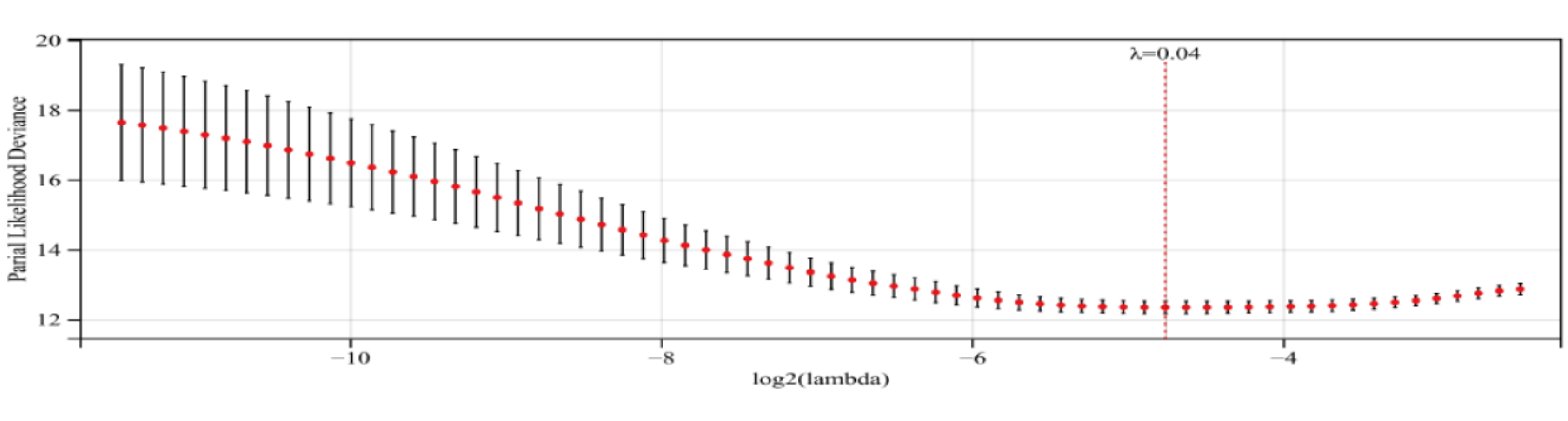
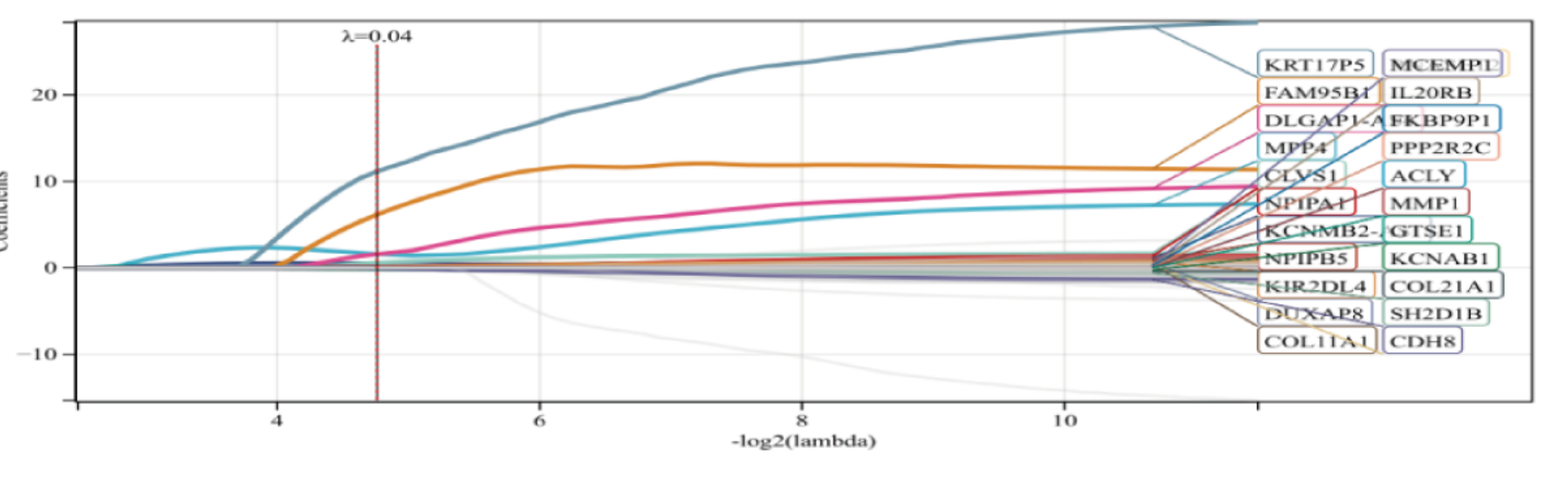
COAD



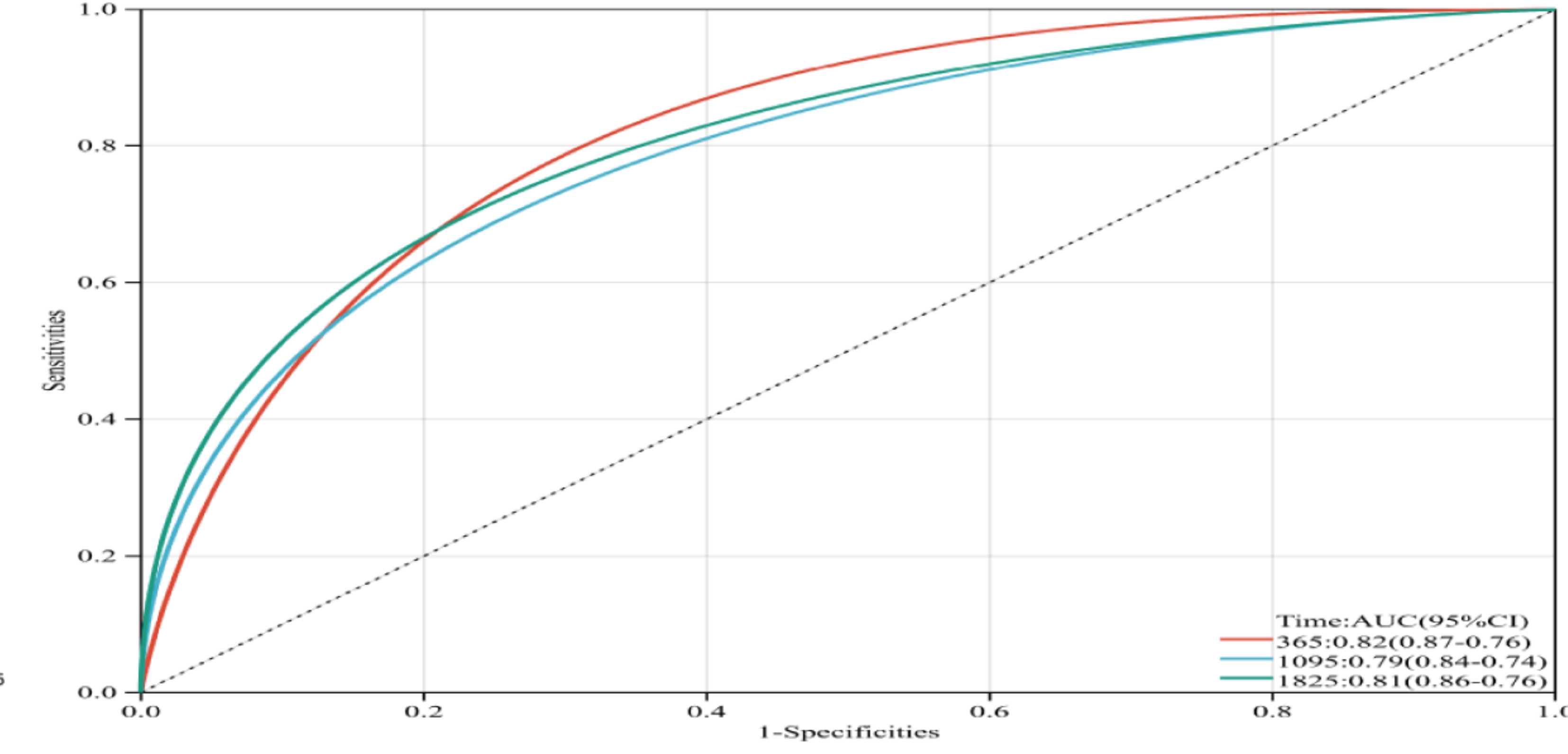
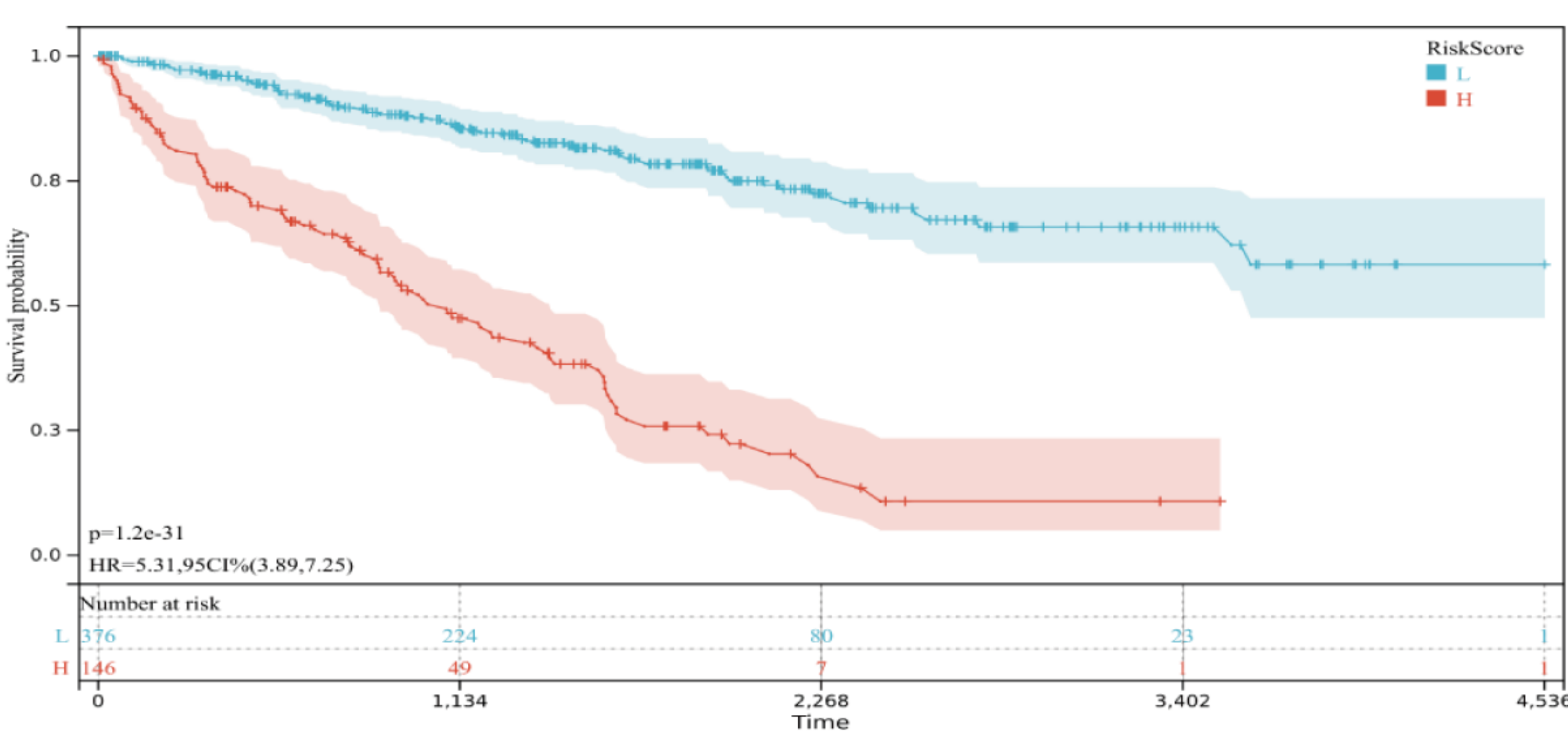
Features	p-value	Hazard Ratio(95%CI)
MYO16-AS1	0.01	1.92(1.14-3.24)
DUXAP10	0.03	5.63(1.17-27.18)
DDX11	0.04	1.59(1.02-2.49)
ISM1	0.05	1.33(1.00-1.77)



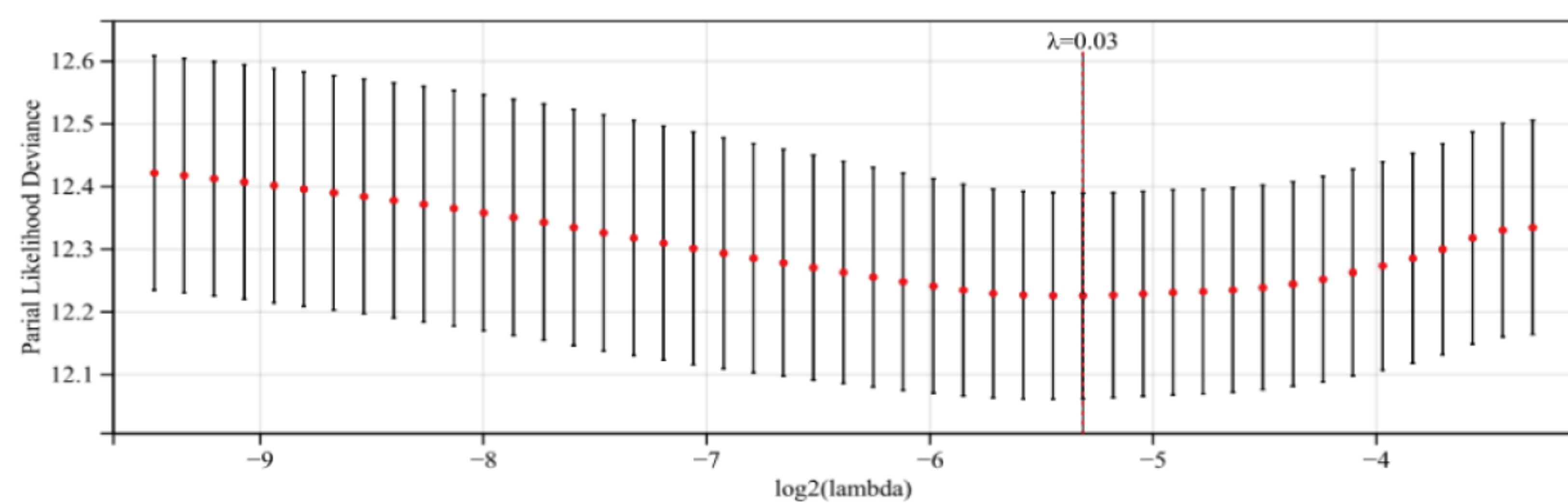
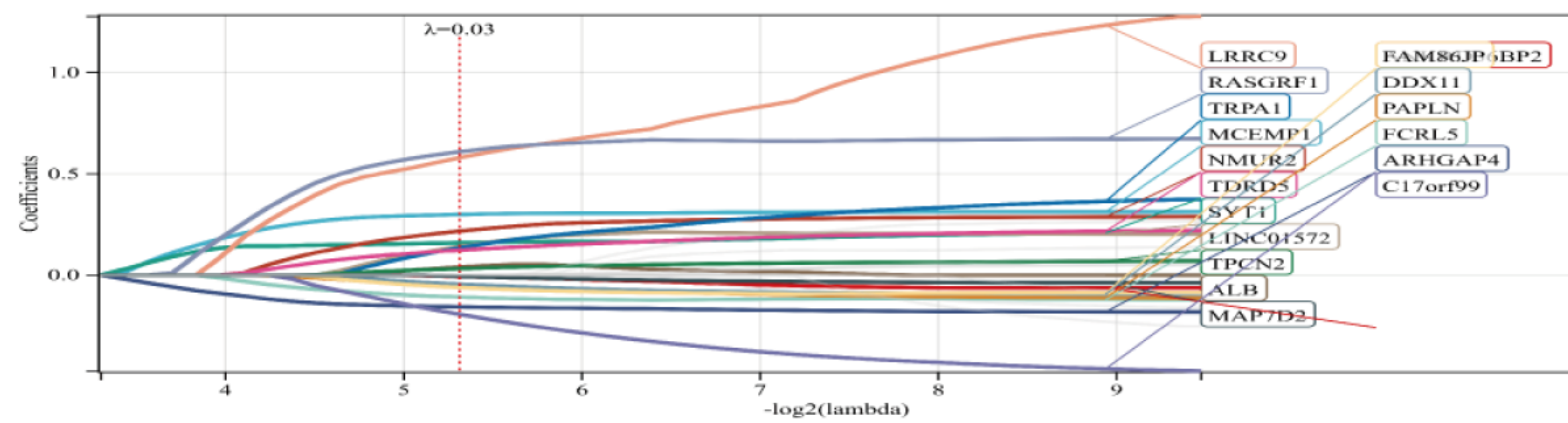
KIRC



Features	p-value	Hazard Ratio(95%CI)
COL11A1	1.2e-6	1.50(1.27-1.77)
FAM95B1	3.8e-5	4260242.16(2980.88-6088689972.53)
CLVS1	6.4e-5	4.75(2.21-10.22)
RNASET2	3.4e-3	1.22(1.07-1.39)
KRT17P5	0.01	267778935.00(74.25-965784601966810.00)
PPP2R2C	0.02	1.36(1.05-1.74)
DLGAP1-AS4	0.02	89.83(2.14-3769.12)
KIR2DL4	0.02	1.75(1.10-2.79)
COL21A1	0.02	0.77(0.62-0.97)
SH2D1B	0.03	0.52(0.29-0.92)
DUXAP8	0.04	1.49(1.02-2.17)
CDH8	0.05	0.41(0.17-0.98)



HNSC



Features	p-value	Hazard Ratio(95%CI)
RASGRF1	0.02	2.12(1.13-3.98)
TDRD5	0.03	1.22(1.02-1.45)
MCEMP1	0.03	1.38(1.03-1.86)
ARHGAP4	0.04	0.84(0.71-0.99)

