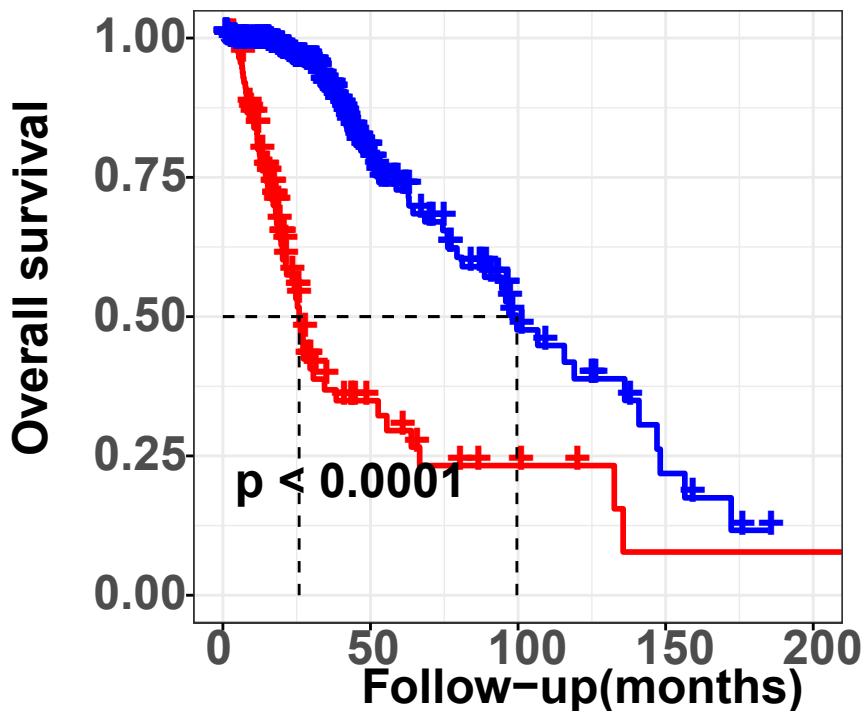


TCGALGG cohort(n=511)

High methylation IEC Low score



Number at risk

