# Supplementary materials



Figure S1 SROC curve for 20 sets of single miRNAs



Figure S2 SROC curve for 7 sets of combined miRNAs



Figure S3 SROC curve for 13 sets of single lncRNAs

 Figure S4 SROC curve for 5 sets of combined lncRNAs

森林图（去掉13、17、19、25）调整

Figure S5 Forest plots after excluding the 13th、17th、19tn、25th sets of miRNAs

figure 8

Figure S6 Diagrams of miRNAs: (a)Goodness-of-fit, (b)Bivariate normality, (c)Influence analysis, (d)Outlier ditection. Goodness-of-fit and Bivariate normality showed that random effects model is suitable. Influence analysis showed studies weight more than others, and Outlier diection is responsible for heterogeneity.

森林图（去掉15、16、17）调整

Figure S7 Forest plots after excluding the 15th、16th、17th sets of lncRNAs

figure 11

Figure S8 Diagrams of lncRNAs: (a)Goodness-of-fit, (b)Bivariate normality, (c)Influence analysis, (d)Outlier ditection.