

A Good prognosis(n=10)/Poor prognosis(n=10)

miRNA Array

22 miRNAs upgraded

miRbase

4miRNAs:Dead miRNA Entry

18miRNAs, compared with miRNA array

qRT-PCR

16miRNAs coherent

2 miRNAs opposite

miRwalk and starBase

7miRNAs had target gene

Cell lines

3 miRNAs inhibited proliferation

miR-29b-2-5p most stable in both *in vivo* and *in vitro*

