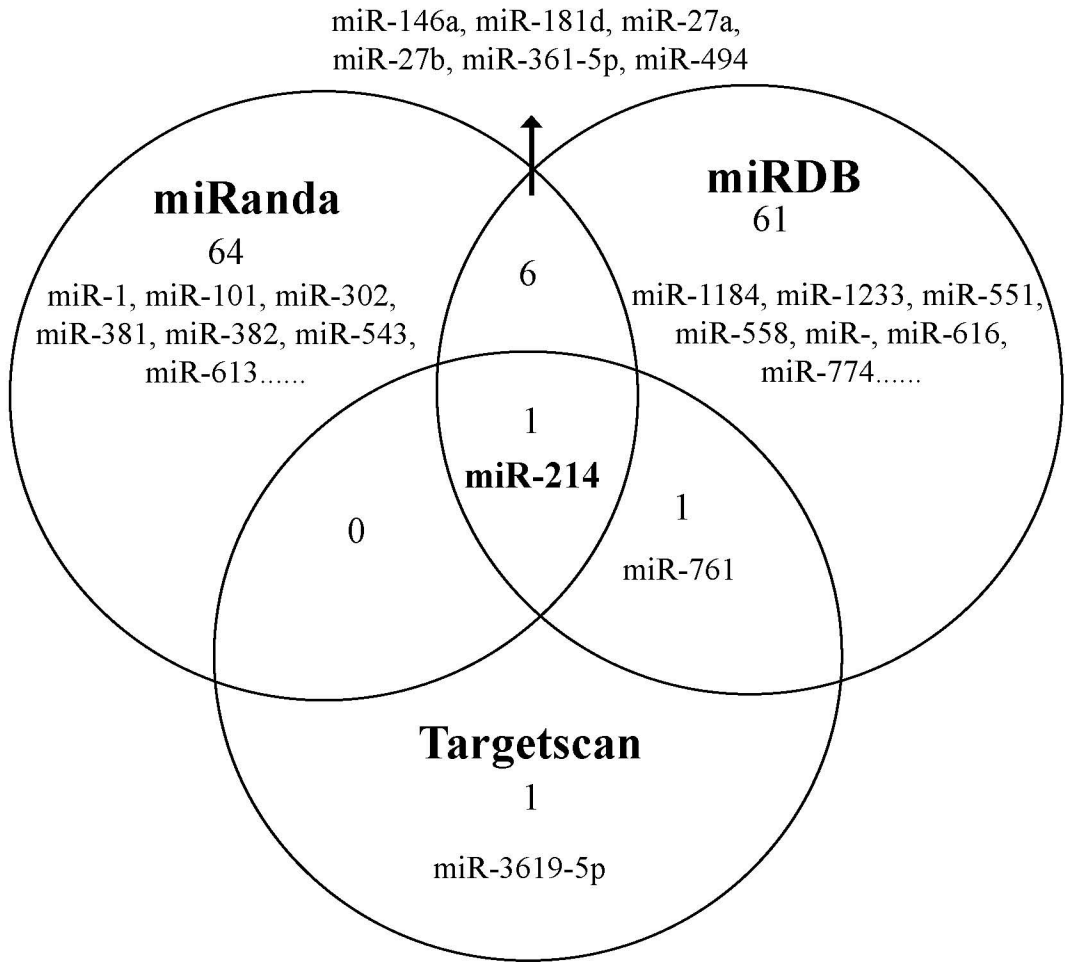


Figure S1

A Only miR-214 can be predicted by three softwares



B The conserved miR-214 binding sites in 3' UTR of RNF8 mRNA

