rank	motif	–log10 p–value	best match
1	2.0 1.5 1.0 0.5 0.0	235	let-7/miR-98
2	2.0 1.5 1.0 0.5 0.0	141	miR-29
3	2.0 1.5 1.0 0.5 0.0	95	miR-519-3p
4	2.0 1.5 1.0 0.5 0.0	78	miR-200/429
5	2.0 1.5 1.0 0.5 0.0	51	AG repeat
6	2.0 1.5 1.0 0.5 0.0	40	miR-25-3p/32-5p/92-3p/ 363-3p/367-3p
7	2.0 1.5 1.0 0.5 0.0 <b>ACUUUA</b>	20	miR-142-5p/5590-3p
8	2.0 1.5 1.0 0.5 0.0	15	miR-21-5p/590-5p