rank	motif	–log10 p–value	best match
1	2.0 1.5 1.0 0.5 0.0	336	miR-194
2	2.0 1.5 1.0 0.5 0.0	320	let-7/miR-98
3	2.0 1.5 1.0 0.5 0.0	274	miR-200/429
4	2.0 1.5 1.0 0.5 0.0	150	miR-29
5	2.0 1.5 1.0 0.5 0.0 ACUUGAA	108	miR-26-5p/1297/4465
6	2.0 1.5 1.0 0.5 0.0	98	miR-17/20/93/106/519/526
7	2.0 1.5 1.0 0.5 0.0	43	miR-210-3p
8	2.0 1.5 1.0 0.5 0.0	39	miR-4291