rank	motif	–log10 p–value	best match
1	2.0 1.5 1.0 0.5 0.0	582	let-7/miR-98
2	2.0 1.5 1.0 0.5 0.0	499	miR-29
3	2.0 1.5 1.0 0.5 0.0	459	miR-200/429
4	2.0 1.5 1.0 0.5 0.0	285	miR–194
5	2.0 1.5 1.0 0.5 0.0	258	miR-17/20/93/106/519/526
6	2.0 1.5 1.0 0.5 0.0	134	miR-191
7	2.0 1.5 1.0 0.5 0.0 ACUUGAA	133	miR-26/1297/4465
8	2.0 1.5 1.0 0.5 0.0	88	miR-22