rank	motif	–log10 p–value	best match
1	2.0 1.5 1.0 0.5 0.0	832	let-7/miR-98
2	2.0 1.5 1.0 0.5 0.0	566	miR-29
3	2.0 1.5 1.0 0.5 0.0	512	miR-200/429
4	2.0 1.5 1.0 0.5 0.0	232	miR-24
5	2.0 1.5 1.0 0.5 0.0	195	miR-17/20/93/106/519/526
6	2.0 1.5 1.0 0.5 0.0	170	miR–194
7	2.0- 1.5- 1.0- 0.5- 0.0- ACUUGAA	169	miR-26/1297/4465
8	2.0 1.5 1.0 0.5 0.0	145	miR–147b