rank	motif	–log10 p–value	best match
1	2.0 1.5 1.0 0.5 0.0	34	let-7/miR-98
2	2.0 1.5 1.0 0.5 0.5	18	miR-29
3	2.0 1.5 1.0 0.5 0.0	16	miR–183
4	2.0 1.5 1.0 0.5 0.5	16	miR-200/429
5	2.0 1.5 1.0 0.5 0.5 0.5	15	miR-486
6	2.0 1.5 1.0 0.5 0.0	14	miR-520/525
7	2.0 1.5 1.0 0.5 0.0	13	miR-214-3p/761/3619