rank	motif	–log10 p–value	best match
1	2.0 1.5 1.0 0.5 0.0	252	let-7/miR-98
2	2.0 1.5- 1.0- 0.5 0.0	152	miR–1278
3	2.0 1.5 1.0 0.5 0.0	122	miR-194
4	2.0 1.5 1.0 0.5 0.0 UCGGAAGC	85	miR-516
5	2.0 1.5 1.0 0.5 0.0 UAGCA	76	miR–27
6	2.0 1.5- 1.0 0.5 0.0	71	miR-29
7	2.0 1.5- 1.0 0.5 0.0	61	miR-210
8	2.0 1.5 1.0 0.5 0.0	47	miR-383