rank	motif	–log10 p–value	best match
1	2.0 1.5- 1.0 0.5 0.0 <b>ASUGUGAA</b>	123	miR–27a/b
2	2.0 1.5 1.0 0.5 0.0	103	miR-15/16/195/322/497
3	2.0 1.5- 1.0 0.5 0.0	98	let–7/miR–98
4	2.0 1.5 1.0 0.5 0.0	86	miR-24
5	2.0 1.5- 1.0- 0.5 0.0	52	miR-4707
6	2.0 1.5- 1.0 0.5 0.0	29	miR-17/20/93/106
7	2.0 1.5 1.0 0.5 0.0 GGCCAGUA	26	miR-193-3p
8	2.0 1.5 1.0 0.5 0.0 SAGUGGU	5	hsa–miR–7–1