rank	motif	−log10 p−value	best match
1		123	miR−27a/b
2		103	miR−15/16/195/322/497
3		98	let−7/miR−98
4		86	miR−24
5		52	miR−4707
6		29	miR−17/20/93/106
7		26	miR−193−3p
8 *		5	hsa−miR−7−1