rank	motif	−log10 p−value	best match
1		72	miR−4667−3p
2		42	miR−26/1297/4465
3		32	miR−21
4		26	miR−629
5		16	miR−3183/4723/6769
6		14	miR−4640−3p
7 *		3	hsa−miR−4753−3p