rank	motif	-log10 p-value	best match
1	2.0 1.5- 1.0- 0.5- 0.0-	18	miR-4768/6833-3p
2	2.0 1.5 1.0 0.5 0.0 AASGOSS	15	miR-4448
3	2.0 1.5 1.0 0.5 0.0	9	miR-4495
4	2.0 1.5 1.0 0.5 0.0 GARGARGA	9	miR-877
5 *	2.0 1.5 1.0 0.5 0.0 CUGCU	3	miR–195