rank	motif	-log10 p-value	best match
1	2.0 1.5 1.0 0.5 0.0	4	hsa-miR-593
2	2.0 1.5 1.0 0.5 0.0	4	hsa-miR-1281
3	2.0 1.5 1.0 0.5 0.0 <b>AAGA</b>	3	hsa–miR–1237
4	2.0 1.5 1.0 0.5 0.0	3	hsa-miR-4530
5 *	2.0 1.5 1.0 0.5 0.0 SACASA	1	hsa-miR-3686