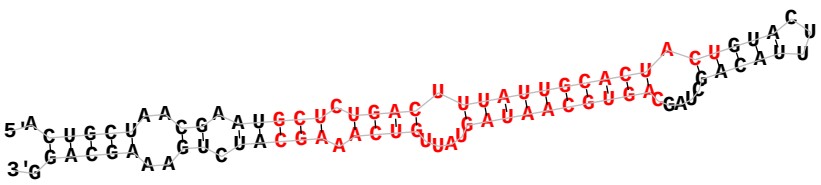
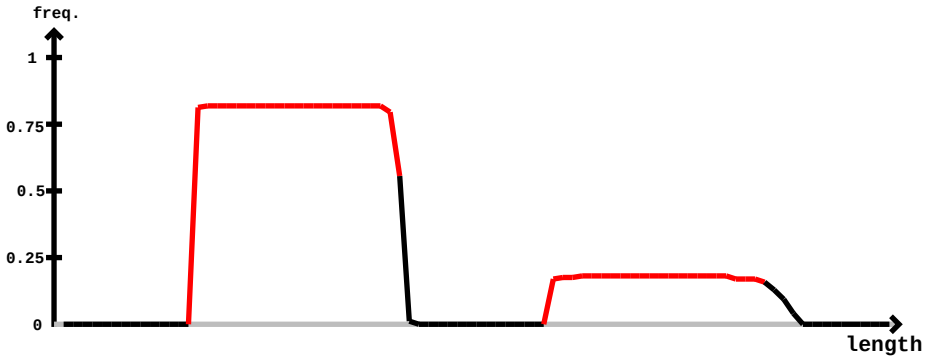
miRBase precursor : hsa-mir-301a
Total read count : 171
hsa-miR-301a-5p read count : 140
hsa-miR-301a-3p read count : 31
remaining reads : 31





hsa-miR-301a-3p

hsa-miR-301a-5p

5'- acugcuaacgaaugcucugacuuuauugcacuacuguacuuuacagcuagcagugcaauaguauugucaaagcaucugaaagcagg -3' exp

.((((((..((.((((((.((((.((((((((((.((((((...)))))).....)))))))))).....))))).)))))).))..)))))).

reads	mm	sample
.....gcucugacuuuauugcaAua.....	1	1	hsa
.....gcucugacuuuauugcacua.....	3	0	hsa
.....gcucGgacuuuauugcacuac.....	1	1	hsa
.....gcucugacuuuauugcacuac.....	32	0	hsa
.....gcucugacuuuauugcacuaU.....	3	1	hsa
.....gcucugacuAuauugcacuac.....	1	1	hsa
.....gcucugacuGuauugcacuac.....	2	1	hsa
.....gcucugacuuuauuUcacuac.....	1	1	hsa
.....gcucAgacuuuauugcacuacu.....	1	1	hsa
.....gcucugCcuuuauugcacuacu.....	1	1	hsa
.....gcucCgacuuuauugcacuacu.....	1	1	hsa
.....gcucugacuuuauuAcacuacu.....	1	1	hsa
.....gcucugacuuuauugcacuaAu.....	1	1	hsa
.....gcucugacuuuauugcacGacu.....	1	1	hsa
.....gcucugacuuuauugcacuacA.....	1	1	hsa
.....gcuUugacuuuauugcacuacu.....	1	1	hsa
.....gcucuUacuuuauugcacuacu.....	1	1	hsa
.....gcucugacuuuauuUcacuacu.....	1	1	hsa
.....gcucugacuuuauugcacuacu.....	83	0	hsa
.....gcucugacuuuauugcacuacug.....	2	0	hsa
.....cucugacuuuauugcacuac.....	1	0	hsa
.....cCgugcaauaguauuguca.....	1	1	hsa
.....cagugcaauaguauuguca.....	1	0	hsa
.....cagugcaauaguauugucaaaA.....	1	1	hsa
.....cagugcaauaguauugucaaag.....	1	0	hsa
.....cagugcaauaguauuUucaaagc.....	1	1	hsa
.....cagugcaauaguauugucaaagc.....	4	0	hsa
.....cagugcaauaguauugucaaagca.....	6	0	hsa
.....cagugcaauaguauugucaaagcau.....	9	0	hsa
.....cagugcaauaguauugucaaagcauU.....	5	1	hsa
.....agugcaauaguauugucaaagcauU.....	1	1	hsa
.....ugcaauaguauugucaaagcauU.....	1	1	hsa