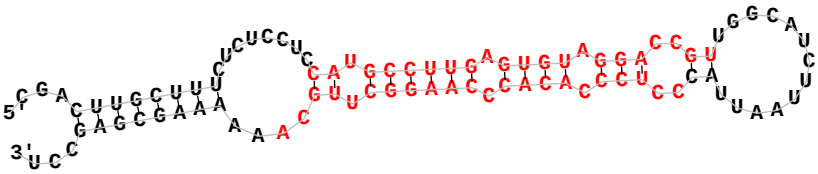
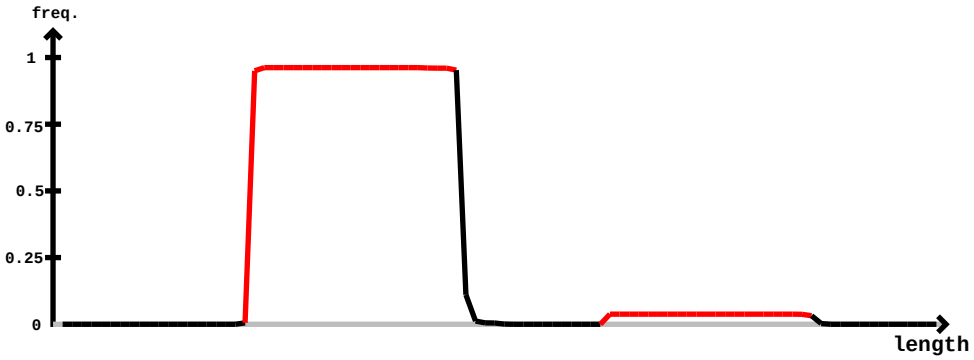
miRBase precursor : hsa-mir-532
Total read count : 1502
hsa-miR-532-5p read count : 1445
hsa-miR-532-3p read count : 57
remaining reads : 57





hsa-miR-532-3p

hsa-miR-532-5p

5'- cgacuugcuuucucuccuccaugccuugaguguaggaccguuggcaucuuaauuaccucccacacccaaggcuugcaaaaaagcgagccu -3' exp

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

reads	mm	sample
.....ccaugccuugaguguaggaccgu.....	7	0	hsa
.....ccaugccuugaguguaggaccguA.....	1	1	hsa
.....caugccuugaguguagga.....	2	0	hsa
.....caugccuugaguguaggac.....	1	0	hsa
.....caugccuugaguguaggaccg.....	11	0	hsa
.....caugccuugagCguaggaccgu.....	1	1	hsa
.....caugccuugaguguaggaUcgu.....	2	1	hsa
.....caugccuugaguguaggacUgu.....	2	1	hsa
.....cUugccuugaguguaggaccgu.....	1	1	hsa
.....caugccuugagugGaggaccgu.....	1	1	hsa
.....caugccuugaguguaUgaccgu.....	5	1	hsa
.....caugccuugGguguaggaccgu.....	2	1	hsa
.....caugccuugaguguaggaccAu.....	1	1	hsa
.....caugccuugaguAuaggaccgu.....	2	1	hsa
.....cauCccuugaguguaggaccgu.....	2	1	hsa
.....caugccuugagGguaggaccgu.....	1	1	hsa
.....caugccuugaguguaggCccgu.....	2	1	hsa
.....caugccuugaguguGggaccgu.....	4	1	hsa
.....cauUccuugaguguaggaccgu.....	5	1	hsa
.....caugccuugaCuguaggaccgu.....	1	1	hsa
.....caugccuugCguguaggaccgu.....	2	1	hsa
.....cCugccuugaguguaggaccgu.....	10	1	hsa
.....caugccuugUguguaggaccgu.....	2	1	hsa
.....caugccuCgaguguaggaccgu.....	1	1	hsa
.....caugccuugaguguagUaccgu.....	7	1	hsa
.....caugccuugaguUuaggaccgu.....	6	1	hsa
.....caugccuugaguguaggacGgu.....	3	1	hsa
.....caugccuugaguguagAaccgu.....	1	1	hsa
.....caugccuugaguguaggaGcgu.....	5	1	hsa
.....caugccuugagAguaggaccgu.....	1	1	hsa
.....caugccuugaguguaggaccgu.....	1137	0	hsa
.....caugcUuugaguguaggaccgu.....	2	1	hsa
.....caugccuuAaguguaggaccgu.....	1	1	hsa
.....caugccuugagugAaggaccgu.....	1	1	hsa

hsa-miR-532-3p

hsa-miR-532-5p

cgacuugcuuucucuccuccaugccuugaguguaggaccguuggcaucuuaauuaccccucccacacccaaggcuugcaaaaaagcgagccu

..................caugccuugaguguaggaAcgu.....	2	1	hsa
..................caugccuugagugCaggaccgu.....	2	1	hsa
..................caugccuugaguguaggaccgA.....	6	1	hsa
..................caugccCugaguguaggaccgu.....	2	1	hsa
..................caugccuugaguCuaggaccgu.....	1	1	hsa
..................caugccuuUaguguaggaccgu.....	4	1	hsa
..................caugccuugaguguCggaccgu.....	2	1	hsa
..................caugccuugaUuguaggaccgu.....	6	1	hsa
..................caugccuGgaguguaggaccgu.....	2	1	hsa
..................caugccuugaguguaggaccUu.....	5	1	hsa
..................caugccGugaguguaggaccgu.....	1	1	hsa
..................caugccuugaguguaggaccguA.....	115	1	hsa
..................caugccuugCguguaggaccguu.....	1	1	hsa
..................caugccuugaguguaggaccguC.....	1	1	hsa
..................caugccuugaguguaggaccguu.....	25	0	hsa
..................caugccuugaguguaggaccguG.....	2	1	hsa
..................caugccuugaguguaggaccguuU.....	1	1	hsa
..................caugccuugaguguaggaccguAg.....	4	1	hsa
..................caugccuugaguguaggaccguug.....	1	0	hsa
..................caugccuugaguguaggaccguuA.....	2	1	hsa
..................caugccuugaguguaggaccguuC.....	2	1	hsa
..................caugccuugaguguaggaccguugA.....	1	1	hsa
..................caugccuugaguguaggaccguuggA.....	2	1	hsa
..................caugccuugaguguaggaccguuggU.....	2	1	hsa
..................caugccuugaguguaggaccguuggc.....	1	0	hsa
..................caugccuugagugGaggaccguuggc.....	1	1	hsa
..................augccuugaguguaggaccgu.....	13	0	hsa
..................augccuuAaguguaggaccgu.....	1	1	hsa
..................augccuugaguguaggaccguA.....	3	1	hsa
..................augccuugaguguaggaccguuggca.....	1	0	hsa
....ccucccacacccaaggcuug.....	1	0	hsa
....Ucucccacacccaaggcuugc.....	1	1	hsa
....ccucccacacccaaggcuugc.....	6	0	hsa
....ccuUccacacccaaggcuugc.....	1	1	hsa
....ccucccacacccaaggcuugcC.....	1	1	hsa
....ccucccacacccaaggcuugca.....	40	0	hsa
....ccucccacacccaagUcuugca.....	1	1	hsa
....ccuUccacacccaaggcuugca.....	1	1	hsa
....ccucccacacccaaggcGugca.....	1	1	hsa
....ccucccacacccaaggcuugcaa.....	1	0	hsa
....ccucccacacccaaggcuugcaU.....	3	1	hsa