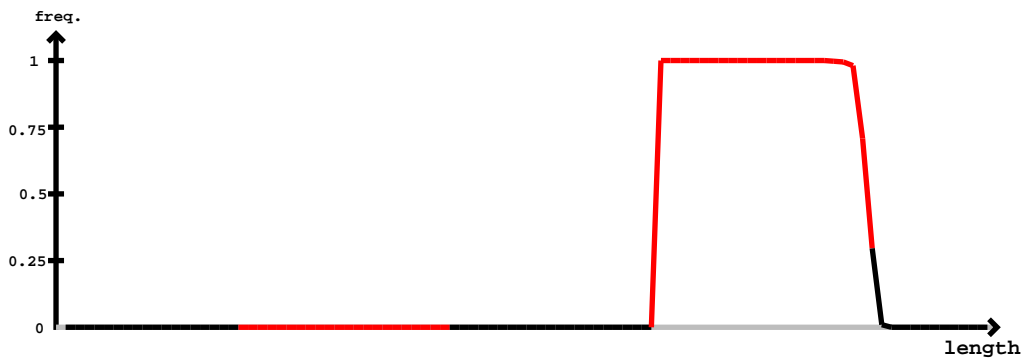
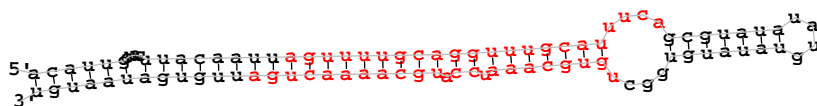


miRBase precursor : hsa-mir-19b-2
 Total read count : 1461
 hsa-miR-19b-3p read count : 1461
 hsa-miR-19b-2-5p read count 0
 remaining reads : 0



hsa-miR-19b-3p

hsa-miR-19b-2-5p

5' -	reads	mm	sample
acauugcuacuacaauuaguuuugcaggguugcauuucagcguaauaauuguaauugggcugugcaaauc <u>caugcaaaacuga</u> uuuguaaauugu			
.....ugugcaaauc <u>caugUaaa</u>	3	1	GDM
.....ugugcaaauc <u>caugUaaa</u>	3	1	GDM
.....ugugcaaaGc <u>caugcaaaa</u>	1	1	GDM
.....ugugcaaauc <u>caugcaaaa</u>	3	0	GDM
.....ugugcaaauc <u>caugUaaaac</u>	8	1	GDM
.....ugugcaaauc <u>caugcaaaaac</u>	5	0	GDM
.....ugugcaaauc <u>caugUaaaac</u>	8	1	GDM
.....ugugcaaauc <u>caugcaaaaac</u>	5	0	GDM
.....ugugcaaauc <u>cauUcaaaaacu</u>	2	1	GDM
.....ugugcaaauc <u>caugcaaaGcu</u>	2	1	GDM
.....ugugcaaauc <u>caugUaaaacu</u>	91	1	GDM
.....ugugcaaauc <u>caugcaaaaacu</u>	118	0	GDM
.....ugugcaaauc <u>caugcaaaaacu</u>	118	0	GDM
.....ugugcaaG <u>ucaugcaaaaacu</u>	1	1	GDM
.....ugugcaaauc <u>caugcaaaaacC</u>	9	1	GDM
.....ugugcaaauc <u>caugcGaaaacu</u>	2	1	GDM
.....ugugcaaauc <u>caugcaaaaacC</u>	9	1	GDM
.....ugugcaaauc <u>caugcaaaaUcu</u>	1	1	GDM
.....ugugcaaauc <u>caugUaaaacu</u>	91	1	GDM
.....ugugcaaauc <u>cauUcaaaaacu</u>	2	1	GDM
.....ugugcaaauc <u>caugcaaaaacu</u>	118	0	GDM
.....ugugcaaauc <u>caugcaaaaacC</u>	9	1	GDM
.....ugugcaaauc <u>caugcaaaaacA</u>	2	1	GDM
.....ugugcaaauc <u>caugUaaaacu</u>	91	1	GDM
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.....ugugcaaauc <u>caugUaaaacug</u>	43	1	GDM
.....ugugcaaauc <u>caugcaaaaUug</u>	1	1	GDM
.....ugugcaaauc <u>caugcaGaacug</u>	2	1	GDM
.....ugugcaaauc <u>caugUaaaacug</u>	43	1	GDM
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.....ugugcaaauc <u>caCgcaaaaacug</u>	1	1	GDM
.....ugugcaaauc <u>caugcaaaaacug</u>	240	0	GDM
.....ugugcaaauc <u>caugcaaaaacuA</u>	6	1	GDM
.....ugugcG <u>aauc</u> caugcaaaaacug.....	2	1	GDM

acauugcuacuuaacaauuaguuuugcagguuugcauuucagcguaauaauuguaauauguggcugugcaaauc caugcaaaacuga uugugauaaugu

.....ugugcaaauc caugcaaaacGg.....	13	1	GDM
.....ugCgcaaauc caugcaaaacug.....	2	1	GDM
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.....ugugcaaauc caugcaaaacugU.....	24	1	GDM
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.....ugugcaaauc caugUaaacuga.....	37	1	GDM
.....ugugcaaauc caugUaaacuga.....	37	1	GDM
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.....ugugcaaauc caugUaaacuga.....	37	1	GDM
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hsa-miR-19b-2-5p

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