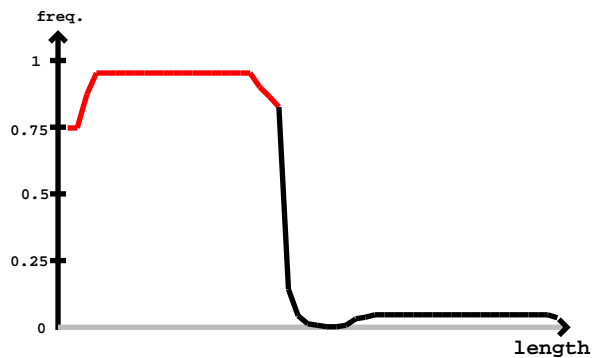
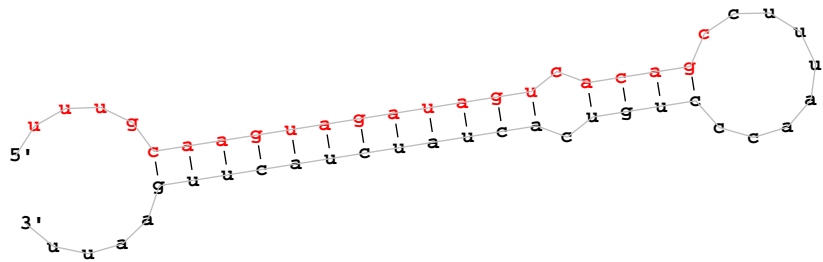


miRBase precursor : chromosome_5_19451
 Total read count : 534
 chromosome_5_19451 read x508t
 remaining reads : 26



chromosome_5_19451			
5'-	-3'	exp	
uuugcaaguagauagucacagc	uuuuaaccugucacuaucuaucugaauu	reads	mm sample
...((((((((((((((((.....))))))))))))))....			
uuugcaaguagauagucGc.....		1	1 seq
uuugcaaguagauagucAU.....		4	1 seq
Auugcaaguagauagucac.....		4	1 seq
uuugcaaguagauagucac.....		9	0 seq
uuugcaaguagauagGcaca.....		1	1 seq
uuugcaaguagauagucaca.....		1	0 seq
Auugcaaguagauagucaca.....		1	1 seq
Auugcaaguagauagucacag.....		1	1 seq
uuugcaaguagauagucacGg.....		4	1 seq
uuugcaaguagauagucacag.....		6	0 seq
uuCgcaaguagauagucacagc.....		1	1 seq
uuugcaaguagaGagucacagc.....		1	1 seq
uuugcaaguagauagucacagc.....		148	0 seq
uuugcaaguagauagCcacagc.....		1	1 seq
uuugcaagCagauagucacagc.....		1	1 seq
uuugcaaguagGuagucacagc.....		4	1 seq
uuugcaaguagauUgucacagc.....		2	1 seq
uuugcaaCuagauagucacagc.....		1	1 seq
uuugcaaguagauagucGcagc.....		2	1 seq
uuugcaaguagauagucacaUc.....		1	1 seq
Auugcaaguagauagucacagc.....		82	1 seq
uuugcaaguagauagucacagA.....		2	1 seq
uuugcaaguGgauagucacagc.....		1	1 seq
uuAgcaaguagauagucacagc.....		1	1 seq
uuuAcaaguagauagucacagc.....		1	1 seq
uuugcaaguagauagucacaAc.....		1	1 seq
uuugcaaguagauagucacagU.....		51	1 seq
Cuugcaaguagauagucacagc.....		3	1 seq
uuugcaGguagauagucacagc.....		1	1 seq
uuugcaaguagauGgucacagc.....		1	1 seq
Guugcaaguagauagucacagc.....		1	1 seq
uuugcaaguagauagucacagcU.....		7	1 seq
uuugcaaguagauagucacagcG.....		1	1 seq
uuugcaaguagauagCcacagcc.....		2	1 seq
Auugcaaguagauagucacagcc.....		18	1 seq

uuugcaaguagauagucacagccuuuaacccugucacuaucuaacuugaauu			
uuugcaaguagauagucacagcc.....	16	0	seq
uuugcaaguagauagucacagcA.....	1	1	seq
uuugcaaguagauagucacagccA.....	5	1	seq
uuugcaaguagauagucacagccC.....	1	1	seq
uuugcaaguagauagucacagccu.....	1	0	seq
Auugcaaguagauagucacagccu.....	4	1	seq
uuugcaaguagauagucacagccAu.....	3	1	seq
uuugcaaguagauagucacagccuu.....	1	0	seq
.uugcaagGagauagucac.....	1	1	seq
.uugcaaguagauagucac.....	8	0	seq
.uugcaaguagauagucaU.....	1	1	seq
.uugcaaguagauagucaca.....	12	0	seq
.uugcaaguagauagucaUa.....	1	1	seq
.uugcaaguagauagucacC.....	2	1	seq
.uugcaaguagauagucacag.....	1	0	seq
.uugcaaguagauagucacagc.....	24	0	seq
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.uugcaaguagauagucacagccu.....	1	0	seq
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.ugcaaguagauagucacagcc.....	2	0	seq
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.....cccugucacuGucuaacuugaauu	3	1	seq
.....ccugucacuaucuaUuugaauu	2	1	seq
.....ccugucacuaucuaacuugaauA	2	1	seq
.....ccugucacuaucuaacuugaauC	1	1	seq
.....ccugucacuGucuaacuugaauu	5	1	seq
.....ccugucacuaucuaacuugaauu	3	0	seq
.....cugucacuaucuaacuugaau.	1	0	seq
.....cugucacuaucuaacuugaauu	2	0	seq
.....cugucacuaucuaUuugaauu	1	1	seq
.....ugucacuaucuaacuugaaC.	1	1	seq
.....ugucacuaucuaacuugaau.	4	0	seq