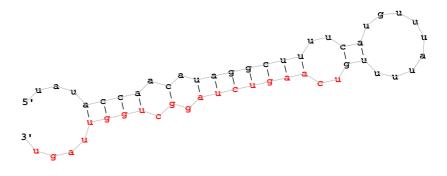
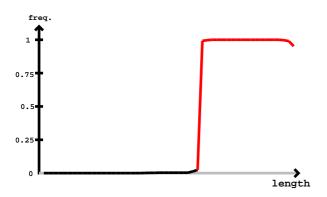
: chromosome\_4\_11169
Total read count : 868
chromosome\_4\_11169 read x86fit
remaining reads : 3





## chromosome\_4\_11169

- uauaccaacauaggcuuuucauguuuauuuugucaagucuaggcugguuagu	-3' exp	<b>,</b>	
((((.(((((((()))))))))))))	reads	mm	sample
aCguuuauuuugucaagucuaggcugguua	1	1	seq
uguuuauuuugucaagucuaggcugguuag.	1	0	seq
uuuauuuugucaagucuaggcugguuagu	1	0	seq
ugucaagucuaggcugguu	1	0	seq
ugucaagucuaggcugguuagu	5	0	seq
ugucaagucuaggcugguuagA	1	1	seq
ugucaagucuaggcuggCuagu	1	1	seq
gucaagucuaggcugguC	2	1	seq
gucaagucuaggcugguuag.	1	0	seq
gucaagucuaggcugguuaUu	2	1	seq
gucaagucuaggcugguuagu	5	0	seq
ucaagucuaggcugguua	4	0	seq
ucaagucuaggcGgguua	1	1	seq
ucaagucuaggcugguuaU.	4	1	seq
ucaagucuaggcugguuaA.	3	1	seq
ucaagucuaggcugguuag.	24	0	seq
ucaagucuaggcugguCagu	1	1	seq
ucGagucuaggcugguuagu	2	1	seq
ucaagucuaUgcugguuagu	1	1	seq
ucaagucuaggcCgguuagu	6	1	seq
ucaagucuaggcugguuagG	1	1	seq
ucaagucuaggcuAguuagu	1	1	seq
ucaGgucuaggcugguuagu	2	1	seq
ucaaAucuaggcugguuagu	1	1	seq
ucaagucuaggcugguuaUu	1	1	seq
ucaaguUuaggcugguuagu	1	1	seq
ucaagucuaggcugguuagu	669	0	seq
ucaaguAuaggcugguuagu	2	1	seq
Gcaagucuaggcugguuagu	1	1	seq
ucaagucuaggcugguuagC	65	1	seq
ucaagucuaggUugguuagu	1	1	seq
ucaagucuaggcugguuGgu	1	1	seq
ucaagCcuaggcugguuagu	4	1	seq
ucaagucuaggcugguuCgu	1	1	seq
ucaagucuaggcugguuUgu	6	1	seq

## chromosome\_4\_11169

uauaccaacauaggcuuuucauguuauuuug <mark>ucaagucuaggcugguuagu</mark>			
ucaagucCaggcugguuagu	4	1	seq
ucaagucuaggcugAuuagu	1	1	seq
ucaagucuaggcugguuagA	16	1	seq
Ccaagucuaggcugguuagu	6	1	seq
ucCagucuaggcugguuagu	2	1	seq
ucaagucuaggcGgguuagu	1	1	seq
ucaagucuGggcugguuagu	2	1	seq
ucaagucuaggcuggCuagu	5	1	seq
caagucuaggcugguuagu	3	0	seq
caagucuaggcugguuagG	1	1	seq
aGgucuaggcugguuagu	1	1	seq

seq

0

.....aagucuaggcugguuagu 2