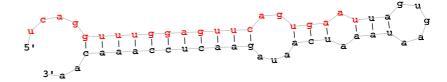
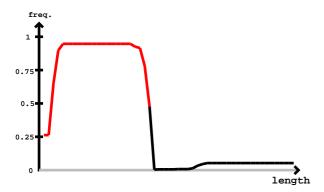
: chromosome_4_12853
Total read count : 510
chromosome_4_12853 read x482t
remaining reads : 28





chromosome_4_12853

٠-	ucagguuuggaguucagugaauuagugaauaaaucaauagaacuccaaacaa	-3' e	exp	
	((((((((((((((((()))).)))))))))	reads	mm	sample
	Gcagguuuggaguucagu	2	1	seq
	ucCgguuuggaguucagu	1	1	seq
	ucagguuuggaguucagu	8	0	seq
	ucagguuuggaguucagug	1	0	seq
	ucagguuuggaguucaguga	7	0	seq
	Acagguuuggaguucaguga	3	1	seq
	Acagguuuggaguucagugaa	1	1	seq
	ucagguuuggaguucagugaa	15	0	seq
	ucaggCuuggaguucagugaa	2	1	seq
	ucagguuuggUguucagugaau	1	1	seq
	Ccagguuuggaguucagugaau	2	1	seq
	ucagguuAggaguucagugaau	1	1	seq
	ucagguuuggaguucagugaau	73	0	seq
	ucaggAuuggaguucagugaau	2	1	seq
	ucagguuuggaguucagCgaau	1	1	seq
	ucagguuuggaguucGgugaau	1	1	seq
	uAagguuuggaguucagugaau	1	1	seq
	ucagguuuggaguucUgugaau	1	1	seq
	ucaggGuuggaguucagugaau	1	1	seq
	Acagguuuggaguucagugaau	6	1	seq
	ucCgguuuggaguucagugaau	2	1	seq
	ucagguuuggaguucagugaauG	1	1	seq
	ucagguuuggaguucagugaauCagugaa	1	1	seq
	.cagguuuggaguucagug	2	0	seq
	.Uagguuuggaguucagug	2	1	seq
	.cagguuuggagCucaguga	1	1	seq
	.cagguuuggagGucaguga	1	1	seq
	.cagguuuCgaguucaguga	1	1	seq
	.Vagguuuggaguucaguga	9	1	seq
	.cagguuuggaguucaAuga	2	1	seq
	.cagguuuggaguucagCga	1	1	seq
	.cagguuuggaguCcaguga	1	1	seq
	.cagguuuggaguucaguga	30	0	seq
	.Aagguuuggaguucaguga	1	1	seq
	.cagguuuggaguucGguga	1	1	seq

ucagguuuggaguucagugaauuagugaauaaaucaauagaacuccaaacaa			
.cagguuuggaguucagugUa	1	1	seq
.Aagguuuggaguucagugaa	2	1	seq
.cagguuuggaguucagugaG	2	1	seq
.cagguuuggaguucagugaa	66	0	seq
.cagguuuggaguucGgugaa	1	1	seq
.cagguuuggagCucagugaa	1	1	seq
.Uagguuuggaguucagugaa	7	1	seq
.cagguuuggagGucagugaau	1	1	seq
.Aagguuuggaguucagugaau	2	1	seq
.Gagguuuggaguucagugaau	2	1	seq
.cagguuuggaguucagugaaC	3	1	seq
.cagguuuggUguucagugaau	2	1	seq
.cagguuuggaguucagugGau	2	1	seq
.cagguuuggaguucGgugaau	3	1	seq
.cagguuuggaguucagugCau	1	1	seq
.cagguuuggaguucagugaau	43	0	seq
.Uagguuuggaguucagugaau	6	1	seq
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agguuugUaguucaguga	1	1	seq
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Ggguuuggaguucagugaa	5	1	seq
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agguuuggaguucagugCa	1	1	seq
Ugguuuggaguucagugaa	5	1	seq
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	1	1	seq
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Ugguuuggaguucagugaau			seq
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agguuuggaguucagugaaG	1	1	seq
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Ggguuuggaguucagugaau	9	1	seq
Aguuuggaguucagugaa	4	1	seq
Cguuuggaguucagugaa	3	1	seq
Uguuuggaguucagugaa	1	1	seq
Aguuuggaguucagugaau	2	1	seq
Uguuuggaguucagugaau	11	1	seq
gguuuggaguucagugaau	4	0	seq
agugaauaaaucaauagaacuccaaacaa	2	0	seq
aUuaaaucaauagaacuccaaacaa	1	1	seq
uaaaucaauagaacuccaaacaa	1	0	seq
aaaucaauagaacuccaaacaa	2	0	seq
aUaucaauagaacuccaaacaa	1	1	seq
aaucaauagaacuccaaacaa	10	0	seq
aucaauagaacuccaaacaa	5	0	seq
Gucaauagaacuccaaacaa	1	1	seq

.....ucaauagaacuccaaacaa 4 0 seq