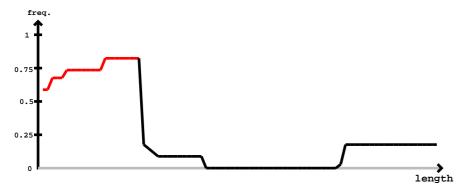
miRBase precursor : chromosome_9_39646

Total read count : 34 chromosome_9_39646 read x2\(\textit{fint}\) remaining reads : 9





chromosome_9_39646

$\verb 5'-ccuuuguagagaaacagucaaaucaaaauggggccauuggugccuauagccccguauggauuugacuuuuucucuacaaauggggccauuggugccuauagccccguauggauuugacuuuuucucuacaaauggggccauuggugccuauagccccguauggauuugacuuuuucucuacaaauggggccauuggugccuauagccccguauggauuugacuuuuucucuacaaauggggccauuggugccuauagccccguauggauuugacuuuuucucuacaaauggggccauuggugccuauagccccguauggauuugacuuuuucucuacaaauggggccauuuggugccuauagccccguauggauuugacuuuuucucucuacaaauggggccauuuggugccuauagccccguauggauuugacuuuuucucucuacaaauggggccauuuggugccuauagccccguauggauuugacuuuuucucucuacaaa$	agg -3	•	exp		
((((((((((((((((((((((((((((((((((((((())) r	eads		mm	sample
ccuuuguagagaaacaguca	4	ŀ		0	seq
Ucuuuguagagaaacaguca	1			1	seq
cUuuuguagagaaacaguca	6	i		1	seq
ccuuuguagagaaacGguca	1			1	seq
ccuuuAuagagaaacaguca	1			1	seq
ccuuugCagagaaacaguca	6	i		1	seq
ccuuuguagagaaacagucaa	1			0	seq
.cuuugCagagaaacaguca	1			1	seq
.cCuuguagagaaacaguca	2	!		1	seq
uguagagaaacaguGaaa	1			1	seq
uguagagaaacaguGaaau	1			1	seq
aacagucaaaucaaaaCgggg	3	1		1	seq
uugacuGuuucucuacaa	agg 1			1	seq
ugacuGuuucucuacaa	agg 5	;		1	seq