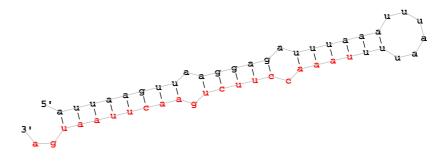
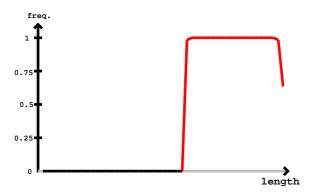
miRBase precursor : chromosome\_10\_50002

 $\begin{array}{lll} \mbox{Total read count} & : 63 \\ \mbox{chromosome\_10\_50002 read} \mbox{ } \pmb{66} \mbox{unt} \\ \mbox{remaining reads} & : 0 \end{array}$ 





## chromosome\_10\_50002

5'- auuaaguuaaggagauuuaaauuuaauuu <mark>uaaaccuucugaacuuaauga</mark>	-3'	exp	
(((((((((((((((((((((((((((((((((((((((	reads	mm	sample
uaaaccuucugaacuuaau	1	0	seq
Aaaaccuucugaacuuaaug.	1	1	seq
uaaaccuucugaacuuaauA.	2	1	seq
uaaaccuucugaacuuaaug.	19	0	seq
uaaaccuucugaacuuaaCga	1	1	seq
uaaaccuucugaacuuaauga	33	0	seq
uaGaccuucugaacuuaauga	1	1	seq
uaaaccuCcugaacuuaauga	1	1	seq
Aaaaccuucugaacuuaauga	1	1	seq
uaaaccuucugaacuuaaugG	1	1	seq
uaaaccuucugaGcuuaauga	1	1	seq
aaaccuucugaacuuaauga	1	0	sea