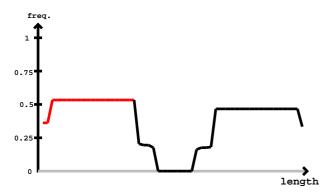
miRBase precursor : chromosome_11_51004

Total read count : 210 chromosome_11_51004 read &bant remaining reads : 98





chromosome_11_51004

-	aacguccuuuuuaaauguagauaauauaaauuaucuacauuuaaaacggacguu	-3'	exp	
	(((((((((((((((((((((((((((((((((((((((reads	mm.	sample
	aacguccCuuuuaaaugua	2	1	seq
	aacguccuuuuuaaaugua	42	0	seq
	Uacguccuuuuuaaaugua	1	1	seq
	aacguccuCuuuaaaugua	1	1	seq
	aacguccuuuuuaaauguG	1	1	seq
	aacguccuuuCuaaaugua	1	1	seq
	aacguccuuuGuaaaugua	1	1	seq
	Gacguccuuuuuaaaugua	4	1	seq
	aacguccuuuuuaaauguag	1	0	seq
	aacguccuuuuuaaauguagau	2	0	seq
	Gacguccuuuuuaaauguagaua	2	1	seq
	aacguccuuuAuaaauguagaua	1	1	seq
	aacguccuuuuuaaauguagauG	1	1	seq
	aacguccuuuuuGaauguagaua	1	1	seq
	aacguccuuuuGaaauguagaua	1	1	seq
	aacguccuuuuuaaauguagaua	13	0	seq
	aacgucAuuuuuaaauguagaua	1	1	seq
	.acguccuuuuuaaaugua	10	0	seq
	.Gcguccuuuuuaaaugua	3	1	seq
	.acguccuuuuuaaaugCa	2	1	seq
	.acgucGuuuuuaaaugua	1	1	seq
	.acguccuCuuuaaauguag	1	1	seq
	.acguccuuuuuaaauguagau	1	0	seq
	.acguccuuuuuaaauguagGu	1	1	seq
	.acAuccuuuuuaaauguagaua	1	1	seq
	.acguccuuuuCaaauguagaua	2	1	seq
	.acguGcuuuuuaaauguagaua	1	1	seq
	.acguccuuuuuaaauguagaua	13	0	seq
	uaucuacauuuaaaaAggacgu.	15	1	seq
	uaucuacauuuaaaaAggacguu	19	1	seq
	aucuacauuuaaaaAggacgu.	1	1	seq
	aucuacauuuaaaaAggacguu	2	1	seq
	cuacauuuaaaaAggacguu	1	1	seq
	uacauuuaaaaAggacgu.	12	1	seq
	uacauuuaaaaGggacguu	2	1	seq

chromosome_11_51004

aacguccuuuuuaaauguagauaauauaaauuaucuacauuuaaaacggacguu

.....uacauuuaaaaAggacguu 46 1 seq