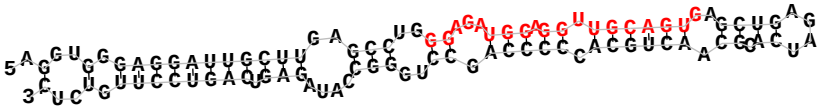
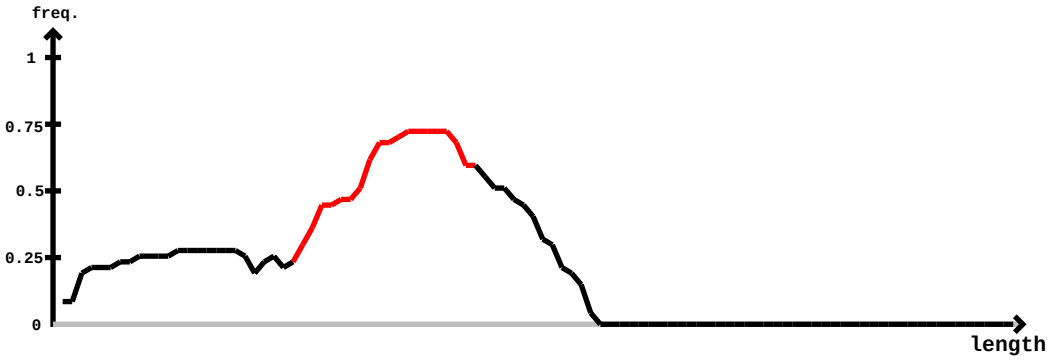
miRBase precursor	: hsa-mir-1273f
Total read count	: 47
hsa-miR-1273f read count	: 10
remaining reads	: 37





hsa-miR-1273f

```
5'- aggugggaggauugcuugagccuggGGAGAUGGAGGUUGCAGUGagcugagaucacgcaacugcacccccagccugggccauagagucaguccuugucuc  -3'   exp
    ......((((((((((..(((..((...(((.((.((((((..((((....)).)).)))))).)))))).))..)))....))).)))))).....
```

	reads	mm	sample
aggugggaggauugcuug	1	0	hsa
aggugggaggauugcuuga	1	0	hsa
aggugggaggauuAcuugagc	1	1	hsa
aggCgggaggauugcuugagcc	1	1	hsa
.gguggggaggauugcuuga	2	0	hsa
.gguggggaggauugcuugag	1	0	hsa
.gUuggggaggauugcuugagcc	1	1	hsa
.gguggggaggauCgcuugagcc	1	1	hsa
..gugggaggauCgcuugagcc	1	1	hsa
.....ggaggauugcuugagGcu	1	1	hsa
.......aggauCgcuugagccugggaga	1	1	hsa
..........uugcuugagccugggagaCg	1	1	hsa
..................ccugggagGuggagguugca	2	1	hsa
..................ccugggagGuggagguugcag	1	1	hsa
..................cugggagGuggagguugcag	1	1	hsa
..................cugggagGuggagguugcaguga	1	1	hsa
..................ugggagauggagAuugcag	1	1	hsa
..................ugggagGuggagguugcagugagcu	1	1	hsa
....................gggagGuggagguugcag	1	1	hsa
....................gggagGuggagguugcagugagcuga	1	1	hsa
....................ggagauggaggCugcaguga	1	1	hsa
....................ggagaCggagguugcagugagc	1	1	hsa
....................ggagGuggagguugcagugagcuga	1	1	hsa
......................gagGuggagguugcagug	1	1	hsa
......................gagCuggagguugcagug	1	1	hsa
......................gagGuggagguugcagugagc	1	1	hsa
......................agGuggagguugcagugagcug	1	1	hsa
......................agGuggagguugcagugagcuga	1	1	hsa
......................agGuggagguugcagugagcugaga	2	1	hsa
........................Guggagguugcagugagcuga	1	1	hsa
........................uggagguugcagugagcug	1	0	hsa
........................gCagguugcagugagcugag	1	1	hsa
........................Agagguugcagugagcugagaucac	1	1	hsa
........................gagguugcagugagcCgaga	1	1	hsa
........................gagguugcagugagcugagauc	1	0	hsa

hsa-miR-1273f

aggugggaggauugcuugagccuggagauggagguugcagugagcugagaucacgcaacugcacccccagccugggccauagagucaguccuugucuc

..........gagguugcagugagcugagaucG..........	4	1	hsa
..........agguugcagugagcCgaga..........	1	1	hsa
..........agguugcagugagcugagaucG..........	1	1	hsa
..........agguugcagugagcugagaucGc..........	1	1	hsa
..........guugcagugagcCgagau..........	1	1	hsa
..........uugcagugagcCgagauc..........	1	1	hsa