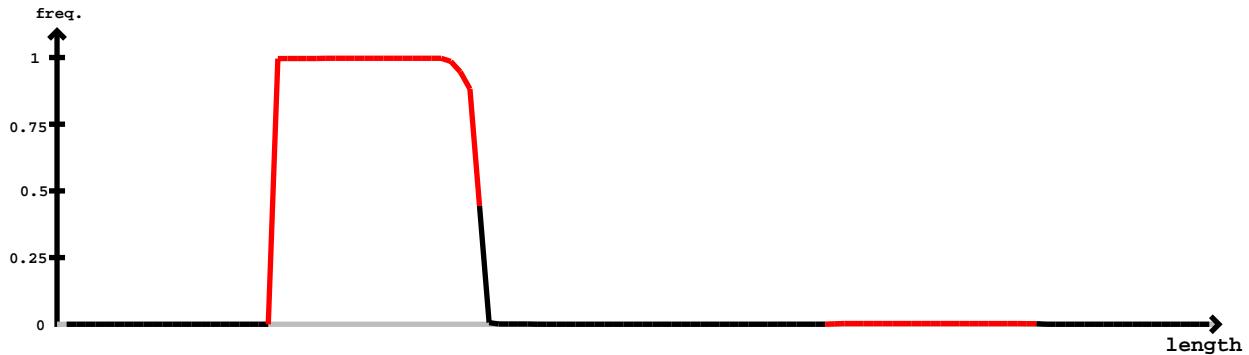miRBase precursor   : hsa-mir-98
Total read count   : 6720
hsa-miR-98-5p read count : 6700
hsa-miR-98-3p read count : 18
remaining reads   : 2

hsa-miR-98-5p

hsa-miR-98-3p

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     .((((..(((((((((((((((.(((((((((((((((((.((((((((((...........))))))..........))))).)))))))))))))))))))))))))))).)))).)))).    reads       mm        sam
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hsa-miR-98-5p

hsa-miR-98-3p

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Sequence	Count	Flag	Group
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...................ugagguaguaaguuguauuguA..	35	1	GDM
...................ugagguUguaaguuguauuguu..	2	1	GDM
...................ugUgguaguaaguuguauuguu..	1	1	GDM
...................ugagguaguaaagCguauuguu..	4	1	GDM
...................ugagguaguaGguuguauuguu..	84	1	GDM
...................ugGgguaguaaguuguauuguu..	5	1	GDM
...................ugagguaguGaguuguauuguu..	8	1	GDM
...................ugaAguaguaaguuguauuguu..	3	1	GDM
...................ugagguaguaaguugCauuguu..	3	1	GDM
...................ugagguaguaaguuguauuguG..	3	1	GDM
...................Ggagguaguaaguuguauuguu..	2	1	GDM
...................ugagguaguaagAuguauuguu..	3	1	GDM
...................ugagguaguaaguuguauuAuu..	1	1	GDM
...................ugaggAaguaaguuguauuguu..	1	1	GDM
...................ugagguaguGaguuguauuguu..	8	1	GDM
...................ugagguaguaaguuguUuuguu..	1	1	GDM
...................ugagguaguaaguuguauCguu..	3	1	GDM
...................ugagguaguaaguuguGuuguu..	13	1	GDM
...................ugagguaguaaguuguGuuguu..	13	1	GDM
...................ugagguaguaaguCguauuguu..	5	1	GDM
...................ugagguaguaaguuguauuguA..	35	1	GDM
...................ugGgguaguaaguuguauuguu..	5	1	GDM
...................Ggagguaguaaguuguauuguu..	2	1	GDM
...................ugagguaguaaguuguauuguu..	1174	0	GDM
...................ugagguaguaaguuguauuguuU...	11	1	GDM
...................ugagguaguaaguuguauuguuU...	11	1	GDM
...................ugagguaguaaguuguauugCug...	1	1	GDM
...................ugagguaguaaguuguauuguug...	2	0	GDM
...................ugagguaguaaguuguauuguuA...	4	1	GDM
...................ugagguaguaaguuguauuguuU...	11	1	GDM
...................ugagguaguaaguuguauAguug...	1	1	GDM
...................ugagguaguaaguuguauuguuA...	4	1	GDM
...................ugagguaguaaguuguauuguuUu..	1	1	GDM
.....................uaguaaguuguauuguuguggg...	2	0	GDM
....cuauacaacuuacuacuuuccU..................	5	1	GDM
....uauacaacuuacuacuuuc....................	3	0	GDM
....uauacaacuuacuacuuuccU..................	1	1	GDM
...................ugagguaguaUguuguau..	1	1	Z3_
...................ugagguaguaaAuuguau..	4	1	Z3_
...................ugagguaguaaguuguauC...	1	1	Z3_
...................ugagguaguaGguuguauu...	1	1	Z3_
...................ugagguaguaaguuguauu...	12	0	Z3_
...................ugagguaguaaguuguauug..	23	0	Z3_
...................ugagguaguaaguuguauuA..	1	1	Z3_
...................ugagguaguaGguuguauug..	6	1	Z3_
...................ugagguaguaaguuguauAg..	1	1	Z3_
...................ugagguagCaaguuguauug..	1	1	Z3_
...................ugagguaguaaguuguauugA...	2	1	Z3_
...................Cgagguaguaaguuguauuguu..	1	1	Z3_
...................ugagguaguaaguugCauugu...	2	1	Z3_
...................ugagguaguaaguuguauugC...	18	1	Z3_
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...................ugagguaguaaguuguauAgu...	17	1	Z3_
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...................ugaAguaguaaguuguauugu...	1	1	Z3_
...................ugagguaguaGguuguauugu...	10	1	Z3_
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...................ugagguaguaaguuguauGguu..	1	1	Z3_
...................Agagguaguaaguuguauuguu..	2	1	Z3_
...................ugagguaguaaguuguauuguu..	211	0	Z3_
aggauucugcucaugccaggg...ugagguaguaGguuguauuguu...guggguagggauauuaggccccaauuagaagauaa...cuauacaacuuacuacuuuccc...uggugugguggcauauuca	12	1	Z3_
...................ugagguaguaaguuguauuguA..	6	1	Z3_

hsa-miR-98-5p

hsa-miR-98-3p

aggauucugcucaugccagggugaggguaguaaguuguauuguugugggguagggauauuaggccccaauuagaagauaacuauacaacuuacuacuuucccuggugguguggcauauuca

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....................ugaggguaguaaguuguauuguG...........................................................    1     1    Z3_
....................ugaggguaguaaguuguGuuguu...........................................................    2     1    Z3_
....................ugaggguaguaaguuguauugCu...........................................................    5     1    Z3_
....................ugaggguaguaaguCguauuguu...........................................................    1     1    Z3_
....................ugagguGguaaguuguauuguu...........................................................    2     1    Z3_
....................ugaggguaguaaguuguauAguu...........................................................   23     1    Z3_
....................ugaggguaguaaguuguauuguC...........................................................   19     1    Z3_
....................Cgaggguaguaaguuguauuguu...........................................................    1     1    Z3_
....................ugaggguagCaaguuguauuguu...........................................................    2     1    Z3_
....................ugaggguaguaaguuguauuguuU...........................................................    2     1    Z3_
....................ugaggguaguaaguuguauuguuA...........................................................    1     1    Z3_
.......................................................................cuauacaacuuacuacuuuccU..................    3     1    Z3_
........................................................................uauacaacuuacuacuuuccU..................    2     1    Z3_
```

hsa-miR-98-5p

hsa-miR-98-3p

aggauucugcucaugccagggugaggguaguaaguuguauuguugugggguagggauauuaggccccaauuagaagauaacuauacaacuuacuacuuucccuggugguguggcauauuca