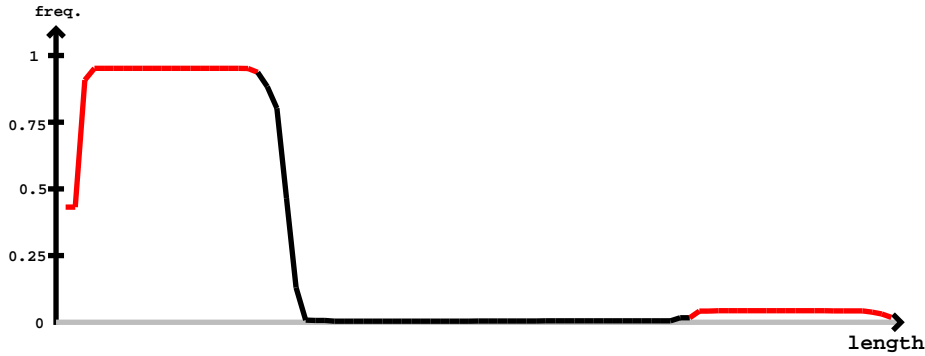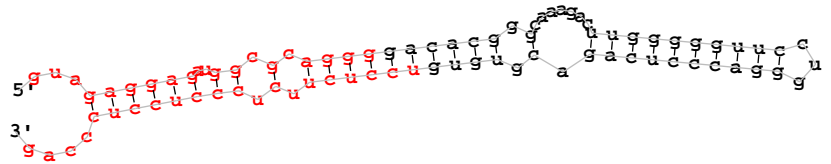miRBase precursor : hsa-mir-877
Total read count : 1478
hsa-miR-877-5p read count : 1403
hsa-miR-877-3p read count : 63
remaining reads : 12

hsa-miR-877-5p

hsa-miR-877-3p

```
5'- guagaggagauggcgcagggg gacacgggcaaagacuuggggguuccugggacccucagacgugug uccucuucucccuccucccag   -3'    exp
    ...((((((..((.(.((((((((((((.(.......((((((((((((...))))))))))).).))))))))))).).))))))))....   reads   mm      sample
    guagaggagauggcgcagggg.........................................................................   1       0       Z3_
    guagaggagauggcgcaggggA........................................................................   1       1       Z3_
    guagaggagauggcgcaggggg........................................................................   1       0       Z3_
    guagaggagauggcgcagggga........................................................................   10      0       Z3_
    guagaggagauggcgcUgggga........................................................................   1       1       Z3_
    guagaggagauggcAcagggga........................................................................   1       1       Z3_
    guagaggagauggcgcaggggU........................................................................   1       1       Z3_
    guagaggagGuggcgcagggggac......................................................................   1       1       Z3_
    guagaggagauggcgcaggggggA......................................................................   8       1       Z3_
    guagaggagauggcgcaggggac.......................................................................   7       0       Z3_
    guagaggagauggcgcagggAac.......................................................................   1       1       Z3_
    guagaggagauggcgcaggggacG......................................................................   1       1       Z3_
    guagaggagauggcgcaggggaca......................................................................   10      0       Z3_
    .uagaggagauggcgcagggg.........................................................................   1       0       Z3_
    .uagaggagauggcgcaggggA........................................................................   1       1       Z3_
    .uagaggagauggcgcagggga........................................................................   24      0       Z3_
    .uagaggagauggcgcaggggaU.......................................................................   3       1       Z3_
    .uagaggagauggcgcaggggaA.......................................................................   19      1       Z3_
    .uagaggagauggcgcaggggac.......................................................................   5       0       Z3_
    .uagaggagauggcgcaggggaAa......................................................................   2       1       Z3_
    ..agaggagauggcgcaggggac.......................................................................   1       0       Z3_
    ..agaggagauggcgcaggggacU......................................................................   1       1       Z3_
    .......................acgggcaaagacuugggggguuccugggacccucagacgugug.....................        1       0       Z3_
    ........................cgggcaaagacuugggggguuccugggacccucagacgugug.....................        5       0       Z3_

    guagaggagauggcgcagg...........................................................................   7       0       Z1_
    guagaggagauggcgcaggg..........................................................................   7       0       Z1_
    guagaggagauggcgcaggA..........................................................................   9       1       Z1_
    guagaggagauggcgcagggU.........................................................................   3       1       Z1_
    guagaggagauggcgcagggA.........................................................................   10      1       Z1_
    guagaggagauggcgcaggggg........................................................................   8       0       Z1_
    guagaggagauggcgcagggga........................................................................   59      0       Z1_
    guagaggagauggcgcaggggU........................................................................   1       1       Z1_
    guagaggagauggcgcaggggG........................................................................   1       1       Z1_
```

guagaggagauggcgcagggggacacgggcaaagacuuggggguuccugggacccucagacguguguccucuucucccuccucccag

Sequence	Count		Sample
guagaggagauggcgcaggggUa..	5	1	Z1_
gGagaggagauggcgcagggggga...	1	1	Z1_
guagaggagGuggcgcagggggga...	1	1	Z1_
guagGggagauggcgcagggggga...	2	1	Z1_
guagaggagauggcgcaggggAa..	3	1	Z1_
guGgaggagauggcgcagggggga...	1	1	Z1_
guagaggagauggcgcagggggac...	21	0	Z1_
guagaggagauggcgcagggggaG...	4	1	Z1_
guagaggagauggcgcagggggaA...	27	1	Z1_
guagGggagauggcgcagggggac...	1	1	Z1_
guagaggagauggcgcagggggaU...	15	1	Z1_
guagaggagauggcgcagggggaca..	28	0	Z1_
guagaggagauggcAcagggggaca..	2	1	Z1_
guagaggagauggcgcagggggaAa..	1	1	Z1_
guagaggagauggcgcagggggacU..	13	1	Z1_
.uagaggagauggcgcaggU...	2	1	Z1_
.uagaggagauggcgcaggA...	5	1	Z1_
.uagaggagauggcgcaggg...	3	0	Z1_
.uagaggagauggcgcagggA..	5	1	Z1_
.uagaggagauggcgcagggU..	4	1	Z1_
.uagaggagauggcgcaggggg...	9	0	Z1_
.uagaggagaCggcgcaggggg...	1	1	Z1_
.uagaggGgauggcgcaggggg...	1	1	Z1_
.uGgaggagauggcgcagggggga...	1	1	Z1_
.uagaggaUauggcgcagggggga...	1	1	Z1_
.uagaggagaugUcgcagggggga...	1	1	Z1_
.uagaggaAauggcgcagggggga...	2	1	Z1_
.uagaggagauggcgcaggggU...	2	1	Z1_
.uagaggagauggcgcagggggga...	87	0	Z1_
.uagGggagauggcgcagggggga...	1	1	Z1_
.uagaggagauggcgcagggggAa...	10	1	Z1_
.uagaggagauggcgcagggggac...	21	0	Z1_
.uagaggagauggcgcagggggaA...	47	1	Z1_
.uagaggagauggcgcagggggaU...	16	1	Z1_
.uagaggagauggcgcagggggacG..	4	1	Z1_
.uagaggagauggcgcagggggaca..	7	0	Z1_
.uagaggagauggcgcagggggaAa..	10	1	Z1_
.uagaggagauggcgcagggggacU..	16	1	Z1_
.uagaggagauggcgcagggggacaU...	1	1	Z1_
..agaggagauggcgcagggggga...	5	0	Z1_
..agaggagauggcgcagggggaU...	3	1	Z1_
..agaggagauggcgcagggggac...	6	0	Z1_
..agaggagGuggcgcagggggac...	1	1	Z1_
..agaggagauggcgcagggggaA...	2	1	Z1_
..agaggagauggcgcagggggaUa..	1	1	Z1_
..agaggagauggcgcagggggaca..	5	0	Z1_
..agaggagauggcgcagggggacacgC...	2	1	Z1_
..agaggagauggcgcagggggacacgU...	2	1	Z1_
....gacccucagacguguguccucuucucccuc.......	1	0	Z1_
....uguccucuucucccuccucc...	2	0	Z1_
....uguccucuucucccuccucU...	1	1	Z1_
....uguccucuucucccuccucccc..	3	0	Z1_
....uguccucuucucccuccuccA..	1	1	Z1_
....uccucuucucccuccuccc..	2	0	Z1_
....uccucuucucccuccucccU.	2	1	Z1_
....uccucuucucccuccucccC.	3	1	Z1_
....uccucuucucccuccuccca.	5	0	Z1_
....Cccucuucucccuccucccag	1	1	Z1_
....uccucuucucccuccucccag	5	0	Z1_
....uccucuucucccuccucccaA	2	1	Z1_
guagaggagauggcgcagg...	6	0	GDM
guagaggagauggcgcagg...	6	0	GDM
guagaggagauggcgcagA...	3	1	GDM
guagaggagauggcgcagC...	2	1	GDM
guagGggagauggcgcaggg..	3	1	GDM
guagGggagauggcgcaggg...hsa-miR-877-3p........	3	1	GDM
guagaggagauggcgcaggA...	26	1	GDM
guagaggagauggcgcaggg..	8	0	GDM

guagaggagauggcgcagggacacgggcaaagacuuggggguuccugggacccucagacgugug<span>uccucuucucccuccucccag

Sequence	Count	Mism	Group
guagaggagauggcgcaggg..	8	0	GDM
guagaggagauggcgcaggA..	26	1	GDM
guagaggagauggcgcaggA..	26	1	GDM
guagaggagauggcgcaggg..	8	0	GDM
guagaggagauggcgcagAg..	5	1	GDM
guagaggagauggcgcagggA...	24	1	GDM
guagaggagauggcgcaggggg..	12	0	GDM
guagaggagauggcgcaggggg..	12	0	GDM
guagaggagauggcgcagggA...	24	1	GDM
guagaggagauggcgcagggA...	24	1	GDM
guagGggagauggcgcaggggg..	1	1	GDM
guagaggagauggcgcaggggg..	12	0	GDM
guagaggagGuggcgcaggggg..	1	1	GDM
guagaggagauggcgcagggU...	2	1	GDM
gGagaggagauggcgcaggggga...	1	1	GDM
guagaggagauggcgcaggggUa...	1	1	GDM
guagaggagauggcgcaggggga...	70	0	GDM
guagaggagauggcgcaggggU..	3	1	GDM
guagaggagauggcgcaggggG..	2	1	GDM
guagaggagauggcgcagggAa..	16	1	GDM
guagaggagauggcgcagggAa..	16	1	GDM
guagaggagauggcgcagggAa..	16	1	GDM
guGgaggagauggcgcaggggga...	2	1	GDM
guagaggagauggcgcaggggU..	3	1	GDM
guagaggagauggcgcaggggga...	70	0	GDM
guUgaggagauggcgcaggggga...	1	1	GDM
guagaggagauggcgcaggggga...	70	0	GDM
guagaggagauggcgcaggggga A...	31	1	GDM
guagaggagauggcgcaggggga U...	15	1	GDM
guagaggagauggcgcaggggga U...	15	1	GDM
guagaggagauggcgcUgggac..	1	1	GDM
guagaggagauggcgcaggggga U...	15	1	GDM
guagaggagauggcgcaggggac...	28	0	GDM
guagaggagauggcgcaggggga A...	31	1	GDM
guagaggagauggcgcaggggac...	28	0	GDM
guagaggagauggcgcaggggac...	28	0	GDM
guagaggagauggcgcaggggga A...	31	1	GDM
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guagaggagauggcgcaggggacU..	32	1	GDM
guagaggagauggcgcaggggaGa..	1	1	GDM
guagaggagauggcgcaggggacU..	32	1	GDM
guagaggagauggcgcaggggaca..	6	0	GDM
guagaggagauggcgcaggggacU..	32	1	GDM
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.uagaggagauggcgcagggA..	1	1	GDM
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.uagGggagauggcgcaggggga..	1	1	GDM
.uagaggagauggcgcagggAa...	6	1	GDM
.uagaggagauggcgcaggggga..	121	0	GDM
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.uagaggagauggcgcaggggga..	121	0	GDM
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.uagaggagauggcgcagggAa...	6	1	GDM
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.uagaggagauggcgcaggggU..	1	1	GDM
.uagaggagauggcgcaggggG..	4	1	GDM
.uagaggagGuggcgcaggggga..	2	1	GDM
.uagaggagauggcgcaggggga A..hsa-miR-877-3p.......	85	1	GDM
.uagaggagauggcgcAcaggggac...uccucuucucccuccucccag	1	1	GDM
.uagaggagauggcgcaggggac..	44	0	GDM

hsa-miR-877-5p

hsa-miR-877-3p

guagaggagauggcgcagggg gacacgggcaaagacuuggggguuccugggacccucagacguguguccucuucucccuccucccag

Sequence	n1	n2	Group
.uagaggagauggcgcagggggaA..	85	1	GDM
.uUgaggagauggcgcagggggac...	1	1	GDM
.uagaggagauggcgcagggggaG...	1	1	GDM
.uagaggagauggcgcagggggaU...	34	1	GDM
.uagaggagauggcgcagggggaU...	34	1	GDM
.uagaggagauggcgcagggggac...	44	0	GDM
.uagaggagauggcgcagggggaU...	34	1	GDM
.uagaggagauggcgcagggggaA...	85	1	GDM
.uagaggagauggcgcagggggac...	44	0	GDM
.uagaggagauggcgcagggggacU..	14	1	GDM
.uagaggagauggcgcagggggaca...	7	0	GDM
.uagaggagauggcgcagggggaca...	7	0	GDM
.uagaggagauggcAcagggggaca...	1	1	GDM
.uagaggagauggcgcagggggacU..	14	1	GDM
.uagaggagauggcgcagggggacU..	14	1	GDM
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.uagaggagauggcgcagggggaUa...	1	1	GDM
.uagaggagauggcgcagggggaAa...	3	1	GDM
.uagaggagauggcgcagggggaAa...	3	1	GDM
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..agaggagauggcgcaggggga...	7	0	GDM
..agaggagauggcgcaggggga...	7	0	GDM
..agaggagauggcgcaggggga...	7	0	GDM
..agaggagauggcgcaggggU...	3	1	GDM
..agaggagauggcgcaggggU...	4	1	GDM
..agaggagauggcgcaggggU...	4	1	GDM
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..agaggagauggcgcagggggac..	7	0	GDM
..agaggagauggcgcagggggac..	7	0	GDM
..agaggagauggcgcagggggacU...	3	1	GDM
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..agaggagauggcgcagggggaca..	8	0	GDM
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....uguccucuucucccuccucc...	1	0	GDM
....uguccucuucucccuccuccU..	2	1	GDM
....uguccucuucucccuccuccU..	2	1	GDM
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....uguccucuucucccuccuccca.	1	0	GDM
....uguccucuucucccuccuccAag	1	1	GDM
....uccucuucucccuccuccU..	1	1	GDM
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....uccucuucucUcuccuccca.	1	1	GDM
....uccucuucucccuccuccccaA	4	1	GDM
....uccucuucucccuccucccag	12	0	GDM
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....uccucuucucccuccucccag	12	0	GDM
....uccucuucucccuccucccag	12	0	GDM
....cucuucucccuccucccag	2	0	GDM
guagaggagauggcgcaA..	1	1	Z2_
guagaggagauggcgcagA...	1	1	Z2_
guagaggagauggcgcaggg..	2	0	Z2_
guagaggagauggcgcaggA..	4	1	Z2_
guagaggagauggcgcaggAg...	1	1	Z2_
guagaggagauggcgcagggg...	4	0	Z2_
guagaggagauggcgcaggggU..	3	1	Z2_
guagaggagauggcgcaggggC..	1	1	Z2_
guagaggagauggcgcaggggga...	13	0	Z2_
guagaggagauggcgcaggggAa...	3	1	Z2_
guagaggagauggcgcagggggac..	9	0	Z2_
Uuagaggagauggcgcagggggac..	1	1	Z2_
guagaggagauggcgcagggggaA..	9	1	Z2_
guagaggagauggcgcagggggaU..	2	1	Z2_
guagaggagauggcgcagggggacU...hsa-miR-877-3p.......	3	1	Z2_
.uagaggagauggcgcagggg...	3	0	Z2_
.uagaggagauggcgcaggggU..	1	1	Z2_

hsa-miR-877-3p

guagaggagauggcgcaggggacacgggcaaagacuuggggguuccugggacccucagacgugugccucuucucccuccucccag

```
.uagaggagauggcgcagggAa...........................................................    1     1     Z2_
.uagaggagauggcgcaggggG...........................................................    1     1     Z2_
.uagaggagauggcgcagggga...........................................................   14     0     Z2_
.uagaggagauggcgcaggggaA..........................................................    7     1     Z2_
.uagaggagauggcgcaggggaU..........................................................    4     1     Z2_
.uagaggagauggcgcaggggac..........................................................    2     0     Z2_
..agaggagauggcgcaggggga..........................................................    1     0     Z2_
..agaggagauggcgcaggggaca.........................................................    1     0     Z2_
...........................................................uccucuucucccuccuccU..    1     1     Z2_
```