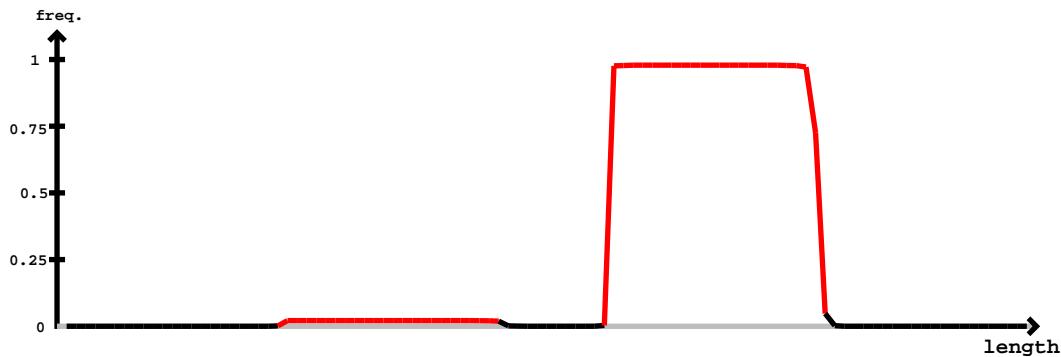miRBase precursor : dme-mir-124_MI0000373
Total read count : 76059
dme-miR-124-5p_MIMAT0020813 read count
dme-miR-124-3p_MIMAT0000351 read count
remaining reads : 0

dme-miR-124-3p_MIMAT0000351

dme-miR-124-5p_MIMAT0020813

```
5'- ucauuugguacguuuuucuccugguuauccacuguaggccuauauguauuuccaccauaaggcacgcggugaaugccaagagcgaacgcaguucuacaaau  -3'   exp
     .........(((((..(((.((((((((..((((((((..(((((.((((((.........))))))))..))))))).))))))).))).)))))...........   reads    mm         sample
     ....................ugguauccacuguaggccuau.........................................................          1       0          seq
     ....................ugAuauccacuguaggccuaua........................................................          1       1          seq
     ....................ugguauccacuguaggccuaua........................................................          10      0          seq
     ...................Agguauccacuguaggccuauau........................................................          1       1          seq
     ...................Ggguauccacuguaggccuauau........................................................          1       1          seq
     ....................ugguauccacuguaggccuauau.......................................................          11      0          seq
     ....................ugguauccacuguagUccuauaug......................................................          1       1          seq
     ....................ugguauccacCguaggccuauaug......................................................          1       1          seq
     ....................ugguauccacuguaggccuauCug......................................................          1       1          seq
     ...................Agguauccacuguaggccuauaug.......................................................          5       1          seq
     ....................ugguauccacuguaggccuauaug......................................................          69      0          seq
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     ....................ugguauccacuguaggccuauaugA.....................................................          3       1          seq
     ....................ugguauccacuguaggccuauaugu.....................................................          1       0          seq
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     .....................gguauccacuguaggccuau.........................................................          26      0          seq
     .....................gguauccacuguaggccuaua........................................................          17      0          seq
     .....................gguaCccacuguaggccuauau.......................................................          1       1          seq
     .....................gguauccacuguaggccuauaA.......................................................          6       1          seq
     .....................gguauccacuguaggccuauau.......................................................          78      0          seq
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     .....................gguauccacuguUggccuauau.......................................................          1       1          seq
     .....................gguauccacugCaggccuauaug......................................................          3       1          seq
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     ....................Aguauccacuguaggccuauaug.......................................................          3       1          seq
     .....................gguauccacuguaggccuauauA......................................................          4       1          seq
     .....................gguauccacuAuaggccuauaug......................................................          1       1          seq
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     .....................gguaGccacuguaggccuauaug......................................................          3       1          seq
     .....................gguauccacuguaggccuCuaug......................................................          1       1          seq
     ....................ggCauccacuguaggccuauaug.......................................................          4       1          seq
     ....................gAuauccacuguaggccuauaug.......................................................          4       1          seq
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dme-miR-124-3p_MIMAT0000351

dme-miR-124-5p_MIMAT0020813

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Sequence			
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gguauccacuguaggccuGuaug	1	1	seq
gguauccacuguagAccuauaug	3	1	seq
gguauccacuguaggUcuauaug	1	1	seq
gguauccacuguaggccuauaCg	3	1	seq
ggGauccacuguaggccuauaug	10	1	seq
gguauccacuguagCccuauaug	1	1	seq
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gguauccacuguaggcUuauaug	1	1	seq
gguauAcacuguaggccuauaug	4	1	seq
gguaCccacuguaggccuauaug	2	1	seq
gguauccCcuguaggccuauaug	2	1	seq
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.gGauccacuguaggccuauau.	2	1	seq
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auaaggcacgcgguAaaugcca	2	1	seq
auaaggcacgcggugaUugcca	1	1	seq
Uuaaggcacgcggugaaugcca	2	1	seq
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auaaggcacgcgCgaaugcca	1	1	seq
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uaagAcacgcggugaaug	1	1	seq
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uaaggcacgcggugCaugcc	1	1	seq
uaaggcacgcggugaaugcc	360	0	seq
uaaggcacgcAgugaaugcc	2	1	seq
Caaggcacgcggugaaugcc	2	1	seq

ucauuugguacguuuuucuccu**gguauccacuguaggccuauaug**uauuuccacca**uaaggcacgcgguggaaugccaag**agcgaacgcaguucuacaaau

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.............................................uaaggcacgcggugaaCgcc.........................  1      1    seq
.............................................uaaggcacgcggugaaugcA.........................  3      1    seq
.............................................uaaggcacgcggGgaaugcc.........................  4      1    seq
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.............................................Aaaggcacgcggugaaugcc.........................  8      1    seq
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.............................................uaaggcacgcggugaaugccG........................  75     1    seq
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.............................................uaaggcacgcggugaaugAca........................  9      1    seq
.............................................uCaggcacgcggugaaugcca........................  2      1    seq
.............................................uaaggcacgcggGgaaugcca........................  133    1    seq
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.............................................uaaggcaUgcggugaaugcca........................  11     1    seq
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Sequence	Count	Mismatch	Type
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....uaaUgcacgcggugaaugccaa.......................	25	1	seq
....uaaggcacgcggugaauUccaa.......................	44	1	seq
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....uaaggcacgcggugUaugccaa.......................	78	1	seq
....uaaggcacgcggugaaugccaa.......................	44387	0	seq
....uaaggcacgcgguCaaugccaa.......................	20	1	seq
....uUaggcacgcggugaaugccaa.......................	5	1	seq
....uaaggcUcgcggugaaugccaa.......................	2	1	seq
....uaaggcacgcgguUaaugccaa.......................	36	1	seq
....uaaggcaGgcggugaaugccaa.......................	2	1	seq
....uaaggcGcgcggugaaugccaa.......................	45	1	seq
....uaagCcacgcggugaaugccaa.......................	8	1	seq
....uaaggcacgcAgugaaugccaa.......................	94	1	seq
....Aaaggcacgcggugaaugccaa.......................	2311	1	seq
....uaaggcacgcgAugaaugccaa.......................	92	1	seq
....uaaggcaAgcggugaaugccaa.......................	17	1	seq
....uaaggcacgcggAgaaugccaa.......................	68	1	seq
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....uaaggcacgcgguAaaugccaa.......................	87	1	seq
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....uaaggcacgcggugaaugUcaa.......................	89	1	seq
....uaaggcacgcggugaaugccCa.......................	564	1	seq
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....uaaggcacgcggugaaugccaG.......................	91	1	seq
....uaaggcacgcggugaaugccaC.......................	212	1	seq
....uaaggcacgcgguaCugccaa.......................	172	1	seq
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....uaaCgcacgcggugaaugccaa.......................	9	1	seq

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....Uaggcacgcggugaaugccaa.....................	3	1	seq

ucauuugguacguuuuucuccu**gguauccacuguaggccuauaug**uauuuccacca**uaaggcacgcggugaaugccaag**agcgaacgcaguucuacaaau

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..............................................aagAcacgcggugaaugccaa......................    1    1    seq
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..............................................Uaggcacgcggugaaugccaag.....................    1    1    seq
..............................................aaggcacgcggugaaugccaaA.....................    1    1    seq
...............................................aggcacgcggugaaugcc........................    1    0    seq
..............................................aCgcacgcggugaaugcca.......................    1    1    seq
...............................................aggcacgcggugaaugcca.......................   18    0    seq
...............................................aggcacgcggGgaaugccaa......................    1    1    seq
...............................................aggcacgcggugaaugccCa......................    1    1    seq
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...............................................aggcacgcggugaaugccaa......................   66    0    seq
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