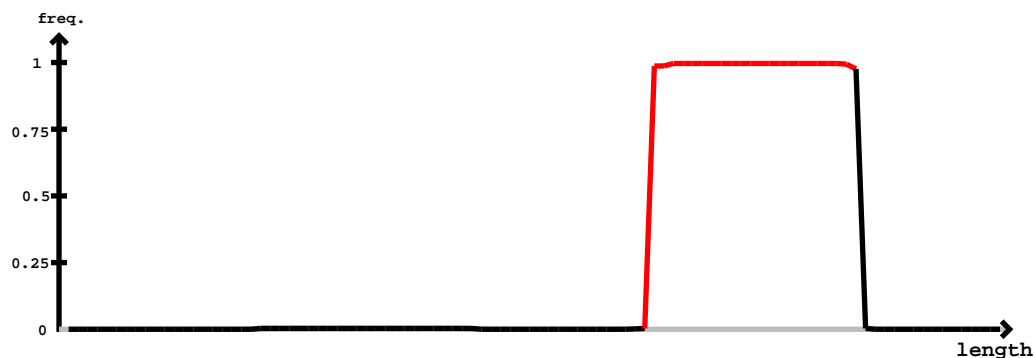| miRBase precursor | : cel-mir-36 |
Total read count	: 14535
cel-miR-36 read count	: 14483
remaining reads	: 52

cel-miR-36

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

	reads	mm	sample
..................cgccaauuuucgcuucagCgc..	1	1	seq
..................cgccaauuuucgcuucagugcu...	2	0	seq
..................cgccaauuuucgcuucaguUcu...	1	1	seq
..................cgAcaauuuucgcuucagugcua..	1	1	seq
..................cgccaauuuucgcuucaUugcua..	3	1	seq
..................cgccaauuuucgcuucagugcua..	37	0	seq
..................cgccaauuuucgcuucagGgcua..	1	1	seq
..................cgccaauuuucgcuucagCgcua..	1	1	seq
..................cgccaauuuucgcuucagugcuaU...	2	1	seq
...................gccaauuuucgcuucaAugcua...	1	1	seq
...................gccaCuuucgcuucagugcua..	1	1	seq
....cuaucaccgggCgaaaauucgca.......................	1	1	seq
....Aaucaccgggugaaaauucgcaug......................	19	1	seq
....uCucaccgggugaaaauucgcaug......................	1	1	seq
....Cucaccgggugaaaauucgcaug.......................	10	1	seq
....aucaAcgggugaaaauucgcaug.......................	1	1	seq
....Uucaccgggugaaaauucgcaug.......................	1	1	seq
....aucaccgggugaaaauucgcaug.......................	12	0	seq
....ucaccgggugaaaauucgc..........................	1	0	seq
....ucacAgggugaaaauucgca.........................	2	1	seq
....ucaccggGgaaaauucgca..........................	2	1	seq
....ucaccgggugaaaauucgUa.........................	1	1	seq
....ucaccgggugaUaauucgca.........................	1	1	seq
....ucaccgggugaaaauuAgca.........................	1	1	seq
....ucaccgggugUaaaauucgca.........................	1	1	seq
....ucaccgggCgaaaauucgca.........................	1	1	seq
....ucaccgggugaaaauucgca.........................	25	0	seq
....ucacUgggugaaaauucgcau.......................	1	1	seq
....ucaAcgggugaaaauucgcau.......................	1	1	seq
....ucaccgggugaaaauucgcau.......................	229	0	seq
....ucaccgggugaaaauucUcau.......................	1	1	seq
....uGaccgggugaaaauucgcau.......................	1	1	seq
....ucaccgggugaaaauucgcCu.......................	1	1	seq
....ucaccgggGgaaaauucgcau.......................	3	1	seq
....ucacAgggugaaaauucgcau.......................	6	1	seq

cel-miR-36

caccgcugucggggaaccgcgccaauuuucgcuucagugcuagaccauccaaagugucuaucaccgggugaaaauucgcaugggucccgacgcgga

Sequence	Count	Edit	Type
....ucaccgggugaaaauuGgcau................	2	1	seq
....ucaccgUgugaaaauucgcau................	1	1	seq
....ucaccgggugUaaauucgcau................	1	1	seq
....uAaccgggugaaaauucgcau................	1	1	seq
....ucaccgggugaaaauucAcau................	1	1	seq
....ucaccgggugaaaauucgcaG................	1	1	seq
....ucaccgggugaaaauuAgcau................	6	1	seq
....ucaccgCgugaaaauucgcau................	1	1	seq
....ucaccgggugaaGauucgcau................	1	1	seq
....ucaccggguUaaaauucgcau................	2	1	seq
....ucaccgggugaGaauucgcaug................	12	1	seq
....ucaccgggugaaaauucgUaug................	18	1	seq
....uGaccgggugaaaauucgcaug................	14	1	seq
....ucaccgggGgaaaauucgcaug................	251	1	seq
....ucaccgggCgaaaauucgcaug................	157	1	seq
....ucaccggUugaaaauucgcaug................	10	1	seq
....ucGccgggugaaaauucgcaug................	6	1	seq
....ucacAgggugaaaauucgcaug................	298	1	seq
....ucaccggguAaaaauucgcaug................	15	1	seq
....ucaccgggugaaaauGcgcaug................	8	1	seq
....ucaccgggugaaaGuucgcaug................	2	1	seq
....ucaccgCgugaaaauucgcaug................	66	1	seq
....Ccaccgggugaaaauucgcaug................	9	1	seq
....ucaAcgggugaaaauucgcaug................	59	1	seq
....ucaccggAugaaaauucgcaug................	19	1	seq
....ucaccgggugaaaGucgcaug................	10	1	seq
....ucaccgggugaaaauucgcauU................	316	1	seq
....ucaccgggugaaUauucgcaug................	5	1	seq
....ucaccgggugaaGauucgcaug................	16	1	seq
....ucaccgAgugaaaauucgcaug................	24	1	seq
....Gcaccgggugaaaauucgcaug................	16	1	seq
....ucaccgggugaaaauuUgcaug................	30	1	seq
....ucaccgggugaaCauucgcaug................	30	1	seq
....ucaccgggugaaaauucgcaCg................	9	1	seq
....ucaccggguUaaaauucgcaug................	13	1	seq
....ucaccgggugaaaauucgcaug................	11571	0	seq
....ucaccgggugaaaauucgcaGg................	36	1	seq
....ucaccgggugaaaauucgcauA................	34	1	seq
....ucaccgggugCaaauucgcaug................	47	1	seq
....Acaccgggugaaaauucgcaug................	7	1	seq
....ucaccgggugaaaauuCcaug................	13	1	seq
....ucaccgggugaaaCuucgcaug................	10	1	seq
....ucaccUggugaaaauucgcaug................	20	1	seq
....ucaccgggAgaaaauucgcaug................	143	1	seq
....ucacUgggugaaaauucgcaug................	29	1	seq
....ucaccgggugaaaauucgcUug................	20	1	seq
....ucaccggguguaaauucgcaug................	25	1	seq
....ucaccgUgugaaaauucgcaug................	9	1	seq
....ucCccgggugaaaauucgcaug................	10	1	seq
....ucaccgggugaaaauCcgcaug................	12	1	seq
....ucaccgggugaaaauucgcauC................	22	1	seq
....uUaccgggugaaaauucgcaug................	29	1	seq
....ucaccgggugaaaauucgGaug................	3	1	seq
....ucaUcgggugaaaauucgcaug................	16	1	seq
....ucacGgggugaaaauucgcaug................	8	1	seq
....ucaccgggugaaaauucgCgug................	8	1	seq
....ucaccgggugaaaCuucgcaug................	11	1	seq
....ucaccggguCaaaauucgcaug................	12	1	seq
....ucaccgggugaaaauAcgcaug................	7	1	seq
....ucaccgggugaaaauucgcaAg................	6	1	seq
....ucaccgggugaaaauuCUcaug................	66	1	seq
....ucaccgggugaaaUuucgcaug................	17	1	seq
....ucaccggCugaaaauucgcaug................	20	1	seq
....ucaccAggugaaaauucgcaug................	21	1	seq
....ucaccggguggaaauucgcaug................	32	1	seq
....ucaccgggugaaaauucgAaug................	19	1	seq
....ucaGcgggugaaaauucgcaug................	21	1	seq
....ucaccCggugaaaauucgcaug................	10	1	seq
caccgcugucggggaaccgcgccaauuuucgcuucagugcuagaccauccaaagugucuaucaccgggugaaaaAucgcaug.gucccgacgcgga	10	1	seq
....ucaccgggugaaaauuGgcaug................	14	1	seq

```
caccgcugucggggaaccgcgccaauuuucgcuucagugcuagaccauccaaagugucuaucaccgggugaaaauucgcaugggguccccgacgcgga
```

Sequence	Count	Flag	Label
....ucaccgggugaaaauucAcaug..............	27	1	seq
....ucaccgggugaCaauucgcaug..............	19	1	seq
....ucaccgggugaaaauucgcCug..............	22	1	seq
....ucaccgggugaUaauucgcaug..............	13	1	seq
....uAaccgggugaaaauucgcaug..............	26	1	seq
....ucaccgggugaaaauuAgcaug..............	95	1	seq
....ucaccgggugaaaauucgcaGgg..............	1	1	seq
....ucaccgggugaaaauucgcaugC..............	2	1	seq
....ucaccgggugaaaauucgcaugU..............	36	1	seq
....ucaccgggugaaaauucgcaugA..............	10	1	seq
....ucaccgggugaaaauucgcauggC.............	1	1	seq
....caccgggugaaaCuucgcaug..............	1	1	seq
....cacUgggugaaaauucgcaug..............	1	1	seq
....caccgggugaaaauucgcaug..............	3	0	seq
....caccgggGgaaaauucgcaug..............	2	1	seq
....accgggugaaaauucgca.................	1	0	seq
....accgggugaaaauucgcau...............	3	0	seq
....accgggugaaaauGcgcaug..............	2	1	seq
....accgggCgaaaauucgcaug..............	1	1	seq
....accgggugaaaauucgcauC..............	1	1	seq
....accgggugaaaAucgcaug..............	1	1	seq
....accgggugaaaauucUcaug..............	2	1	seq
....accgggugaaaauucCcaug..............	1	1	seq
....accgggugaaaauucgcaCg..............	1	1	seq
....accggguCaaaauucgcaug..............	2	1	seq
....accggAugaaaauucgcaug..............	1	1	seq
....accgggugaaaauuAgcaug..............	2	1	seq
....accCggugaaaauucgcaug..............	2	1	seq
....acUgggugaaaauucgcaug..............	1	1	seq
....accgggugaaaauucgcauU..............	2	1	seq
....accgggugaaaauucgcaug..............	104	0	seq
....accgAgugaaaauucgcaug..............	1	1	seq
....accgggAgaaaauucgcaug..............	1	1	seq
....accgggGgaaaauucgcaug..............	4	1	seq
....cgggugaaaauucgcaug..............	1	0	seq