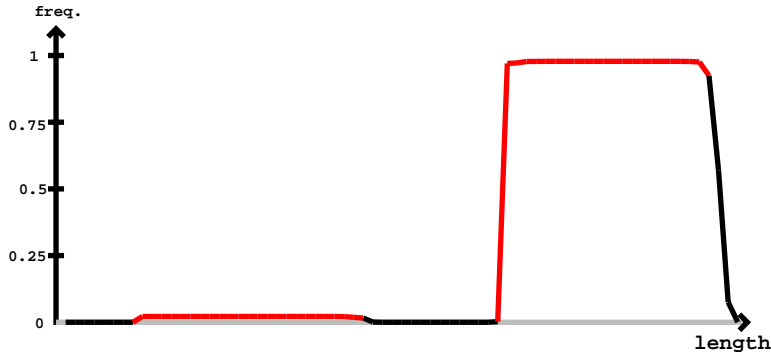
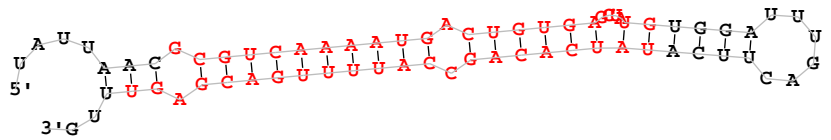


miRBase precursor : dme-mir-13b-2_MI0000135
 Total read count : 21515
 dme-miR-13b-2-5p_MIMAT0020797 read count 1630797
 dme-miR-13b-3p_MIMAT0000119 read count 20049
 remaining reads : 0



dme-miR-13b-2-5p_MIMAT0020797

dme-miR-13b-3p_MIMAT0000119

| 5' - | reads | mm | sample | -3' | exp |
|---|-------|----|--------|-----|-----|
| uauuaacgcgucaaaaugacugugagcuauguggauuugacuucuaucacagccauuuugacgaguuug | | | | | |
|(((.....(((.....(((.....(((.....))))))))))))))..... | 13 | 0 | seq | | |
|gcgucaaaaugacugugagcu..... | 50 | 0 | seq | | |
|gcgucaaaaugacugugagcuC..... | 1 | 1 | seq | | |
|gcgucaaGaugacugugagcua..... | 1 | 1 | seq | | |
|gcgucaaaaugacugugagUua..... | 1 | 1 | seq | | |
|gcgucaaaaugacugugagcuG..... | 1 | 1 | seq | | |
|gcgucaaaaugacugugagAuau..... | 1 | 1 | seq | | |
|gcgucaaaaugacugugagcuau..... | 53 | 0 | seq | | |
|gcgucaaaaugacugugagcuac..... | 1 | 1 | seq | | |
|gcgucaaaaugacuAugagcuau..... | 3 | 1 | seq | | |
|gcgucaaaaugacugugagcuacA..... | 1 | 1 | seq | | |
|gcAucaaaaugacugugagcuau..... | 1 | 1 | seq | | |
|gcgucaaaaugacugUagcuau..... | 3 | 1 | seq | | |
|gcgucaaaaugacugAgagcuau..... | 1 | 1 | seq | | |
|gcgCcaaaaugacugugagcuau..... | 1 | 1 | seq | | |
|gAgucaaaaugacugugagcuau..... | 3 | 1 | seq | | |
|gcgucaaaaugacugugagcuCug..... | 1 | 1 | seq | | |
|gcgucaaaaugacugugagcuauA..... | 2 | 1 | seq | | |
|gcgucaaaGugacugugagcuau..... | 1 | 1 | seq | | |
|gcgucaaaaugacugugagcuau..... | 297 | 0 | seq | | |
|gcgucaaaaugacugugCgcuau..... | 1 | 1 | seq | | |
|gcgucaaaauAacugugagcuau..... | 2 | 1 | seq | | |
|gcgucaaaaugacugugagcuacGg..... | 1 | 1 | seq | | |
|gcgucaaaaugacugugGgcuau..... | 1 | 1 | seq | | |
|Aogucaaaaugacugugagcuau..... | 1 | 1 | seq | | |
|gcgucaaaaugacugugagcuGug..... | 1 | 1 | seq | | |
|gUgucaaaaugacugugagcuau..... | 1 | 1 | seq | | |
|gcgGcaaaaugacugugagcuau..... | 2 | 1 | seq | | |
|gcgucaaaaugacugugagcuau..... | 1 | 1 | seq | | |
|Ucgucaaaaugacugugagcuau..... | 1 | 1 | seq | | |
|gcgUaaaaugacugugagcuau..... | 1 | 1 | seq | | |
|gcgucaaaaugacugugagcuauGA..... | 15 | 1 | seq | | |
|gcgucaaaaugacugugagcuauagu..... | 3 | 0 | seq | | |
|aaaugacugugagcuau..... | 1 | 0 | seq | | |

uauuaacgcucaaaaugacugagcuauugggauuugacucauauacacagccauuuugacgaguug

| | | | |
|----------------------------------|------|---|-----|
|Aauaucacagccauuuugacgagu.. | 1 | 1 | seq |
|auaucacagccauuuugacgag... | 8 | 0 | seq |
|auGucacagccauuuugacgagu... | 1 | 1 | seq |
|auaucacagccauuuugacgagu... | 17 | 0 | seq |
|Uuaucacagccauuuugacgagu... | 1 | 1 | seq |
|auaucacagccauuuugacgagu... | 17 | 0 | seq |
|auaucacagccauuuugacgaguA. | 4 | 1 | seq |
|uauacacagccauuuCgac..... | 1 | 1 | seq |
|uauacacagccauuuugac..... | 3 | 0 | seq |
|uauacacagcUauuuugacg..... | 1 | 1 | seq |
|uauacacagccauuuugacg..... | 7 | 0 | seq |
|uauacGcagccauuuugacga..... | 1 | 1 | seq |
|uauacacagccauuuugacga..... | 44 | 0 | seq |
|uCuacacagccauuuugacga..... | 1 | 1 | seq |
|uauacacGgccauuuugacgag... | 2 | 1 | seq |
|uauacacagccauuuugacgag... | 1009 | 0 | seq |
|uauacacagccauuuCgacgag... | 2 | 1 | seq |
|uauacacagccauuGugacgag... | 1 | 1 | seq |
|uauacacagccauuuugacgCg... | 3 | 1 | seq |
|uauacacagccauuuugacUag... | 1 | 1 | seq |
|uaGcacagccauuuugacgag... | 1 | 1 | seq |
|uauacacagccauuuugaAgag... | 2 | 1 | seq |
|uauacacagGcauuuugacgag... | 1 | 1 | seq |
|uauacacagccauuuugacAag... | 2 | 1 | seq |
|Aauacacagccauuuugacgag... | 25 | 1 | seq |
|uaCcacagccauuuugacgag... | 5 | 1 | seq |
|uauacacagccauCuugacgag... | 1 | 1 | seq |
|uauacacagccauuuugacgaA... | 5 | 1 | seq |
|uauacacagccuuuugacgag... | 1 | 1 | seq |
|uCuacacagccauuuugacgag... | 1 | 1 | seq |
|uauacacagccauuuAgacgag... | 2 | 1 | seq |
|uauacacagcUauuuugacgag... | 2 | 1 | seq |
|uauacacagccGuuuugacgag... | 1 | 1 | seq |
|uauacacagccauuuugacgGg... | 1 | 1 | seq |
|uauUacagccauuuugacgag... | 1 | 1 | seq |
|uauacacagccauuugacgag... | 1 | 1 | seq |
|uauacacagccauuuUacgag... | 2 | 1 | seq |
|uauacacagccauuuugacgUg... | 1 | 1 | seq |
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|uauacacagccauuuugCcgag... | 1 | 1 | seq |
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|uGucacagccauuuugacgag... | 1 | 1 | seq |
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|uGucacagccauuuugacgagu... | 9 | 1 | seq |
|uauacacagccauuuugacgCgu... | 14 | 1 | seq |
|uauacacagccauuAugacgagu... | 1 | 1 | seq |
|uauacacagcUauuuugacgagu... | 3 | 1 | seq |
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|uauacacagccuuuugacgagu... | 2 | 1 | seq |
|uauacacagccauuuCacgagu... | 5 | 1 | seq |
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|uauacacagccaGuuuugacgagu... | 1 | 1 | seq |
|uauacacagccUuuuugacgagu... | 2 | 1 | seq |
|Aauacacagccauuuugacgagu... | 266 | 1 | seq |
|uauacacagccauuuugacgaUu... | 2 | 1 | seq |
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|uauacacagccGuuuuugacgagu... | 5 | 1 | seq |
|uauacacGgcauuuugacgagu... | 10 | 1 | seq |
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|uauacacagcAuuuuugacgagu... | 5 | 1 | seq |
|uauacacagccauuuugaGgagu... | 2 | 1 | seq |
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|uauacacagccauGuugacgagu... | 6 | 1 | seq |
|uauacacagccauuuugacgUgu... | 3 | 1 | seq |
|uaucaAagccauuuugacgagu... | 1 | 1 | seq |

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|uaucaacagccauuuugUcgagu.. | 16 | 1 | seq |
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|uaucaacagUcauuuugacgagu.. | 4 | 1 | seq |
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|uUucaacagccauuuugacgagu.. | 1 | 1 | seq |
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|uaucaacagccauuuuugacgaCu.. | 2 | 1 | seq |
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|uCucaacagccauuuuugacgagu.. | 39 | 1 | seq |
|Caucaacagccauuuuugacgagu.. | 31 | 1 | seq |
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|uauUaacagccauuuuugacgagu.. | 5 | 1 | seq |
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|uaucaacagccauuuuugCcgagu.. | 10 | 1 | seq |
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|uauGaacagccauuuuugacgagu.. | 3 | 1 | seq |

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|uauacacagccauCuugacgaguu.. | 13 | 1 | seq |
|uauacacagccauuuugacgaguG.. | 16 | 1 | seq |
|uauacacagUcauuuugacgaguu.. | 5 | 1 | seq |
|uauacacagccauuuugacGguu.. | 22 | 1 | seq |
|uaucaAagccauuuugacgaguu.. | 1 | 1 | seq |
|Gaucacagccauuuugacgaguu.. | 8 | 1 | seq |
|uCuacacagccauuuugacgaguu.. | 18 | 1 | seq |
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|uauacacagccauGuugacgaguu.. | 10 | 1 | seq |
|uauacacagccauuuugacgaguA.. | 32 | 1 | seq |
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|uauacacagccauuuugaAgaguu.. | 7 | 1 | seq |
|Aaucacagccauuuugacgaguu.. | 220 | 1 | seq |
|uauacacagccauuuugaGgaguu.. | 1 | 1 | seq |
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|uauacacagccauuuugacgaguu.. | 9634 | 0 | seq |
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|uaGcacagccauuuugacgaguu.. | 4 | 1 | seq |
|uauacacagccaCuugacgaguu.. | 14 | 1 | seq |
|uaucGcacgccaauuuugacgaguu.. | 22 | 1 | seq |
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|uauacacagccauuuuugacgaguuCu.. | 1 | 1 | seq |
|uauacacagccauuuuugacGguuu.. | 1 | 1 | seq |
|uauacGcacgccaauuuuugacgaguuu.. | 1 | 1 | seq |
|uauacacagccauuuuugacgaguuC.. | 11 | 1 | seq |
|uauacacagccauuuuugacgaguuAg.. | 2 | 1 | seq |
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|aucacagccaAuuuugacgaguu... .. | 1 | 1 | seq |
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|ucacagccauuuuugacgag... .. | 2 | 0 | seq |
|ucGcacgccaauuuuugacgaguu... .. | 1 | 1 | seq |
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|cacagccauuuuugacgag... .. | 1 | 0 | seq |
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dme-miR-13b-2-5p_MIMAT0020797

dme-miR-13b-3p_MIMAT0000119

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|acagccauuuugacgagu.. | 1 | 0 | seq |
|acagccauuuugacgaguA. | 1 | 1 | seq |
|Aagccauuuugacgagu.. | 3 | 1 | seq |
|cagccauuuugacgagu.. | 1 | 0 | seq |