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Heterogeneous data analysis for annotation of microRNAs and novel genome assembly

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      ATGGCGCGGGATTGTGAGGGATTAGGAACCAATTCTGGATCGTGACTTCG
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      TTCTCCCTGCAGCCCCCTGCCGGTGTTCGTGATAGGTGAAAATAAACCATGGAATTC
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      TCAATTCGTGTTTC
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      TTTTCTCT
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      CTGACTTCAGCAGAGTATCTC
      TCGGATTC

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