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Sequence Results: We will examine the quality of the DNA sequence from each sample and use two DNA sequences (forward and reverse) taken from the same sample to verify accurate sequencing results. Identify similar sequences: Using BLAST, we will search for similar (related) DNA sequences from a database of known sequences. 3th, 2024

DNA Structure: A-, B- And Z-DNA Helix Families

TA	TC	TG	TT
-3.82	-11.85	-13.48	-9.45
36.0	36.9	34.5	35.6
Average	-7.92 ± 2.57	35.7 ± 8.0	-12.60 ± 3.2

DNA Structure: A-, B- And Z-DNA Helix Families 4

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Use Of Uracil-DNA Glycosylase Enzyme To Reduce DNA-related ... Predicts Sensitivity To Specific Treatments [2]. Correct And Precise Mutation Detection Is An Essential Step In Precision Medicine [3]. Although The Benefits Of NGS Are Unquestionable, The Complexities Of These Method-ologies And The Req 1th, 2024.

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