

ABSTRACT

In the gene sequencing analysis method and apparatus, storage medium, and computer device provided by the present disclosure, short sequences transmitted from a sequencing platform in real time are first sliced to perform corresponding slicing processing, and then the slice data is input into a memory. Thus, a pre-loaded encapsulation program in the memory can be called to perform bioinformatic analysis on the slice data, thereby obtaining a corresponding analysis result. Such a process can acquire the slice data of the sequencing of the sequencing platform in real time and perform sequencing and analysis on the slice data, rather than performing the processing analysis only after the sequencing platform has finished the sequencing and the complete sequencing results are transmitted to the corresponding platform. Therefore, the present disclosure enables the whole of sequencing and analysis to be accelerated. In addition, the data for analysis is the slice data, which can be transmitted faster and takes less time compared with the whole sequencing result.