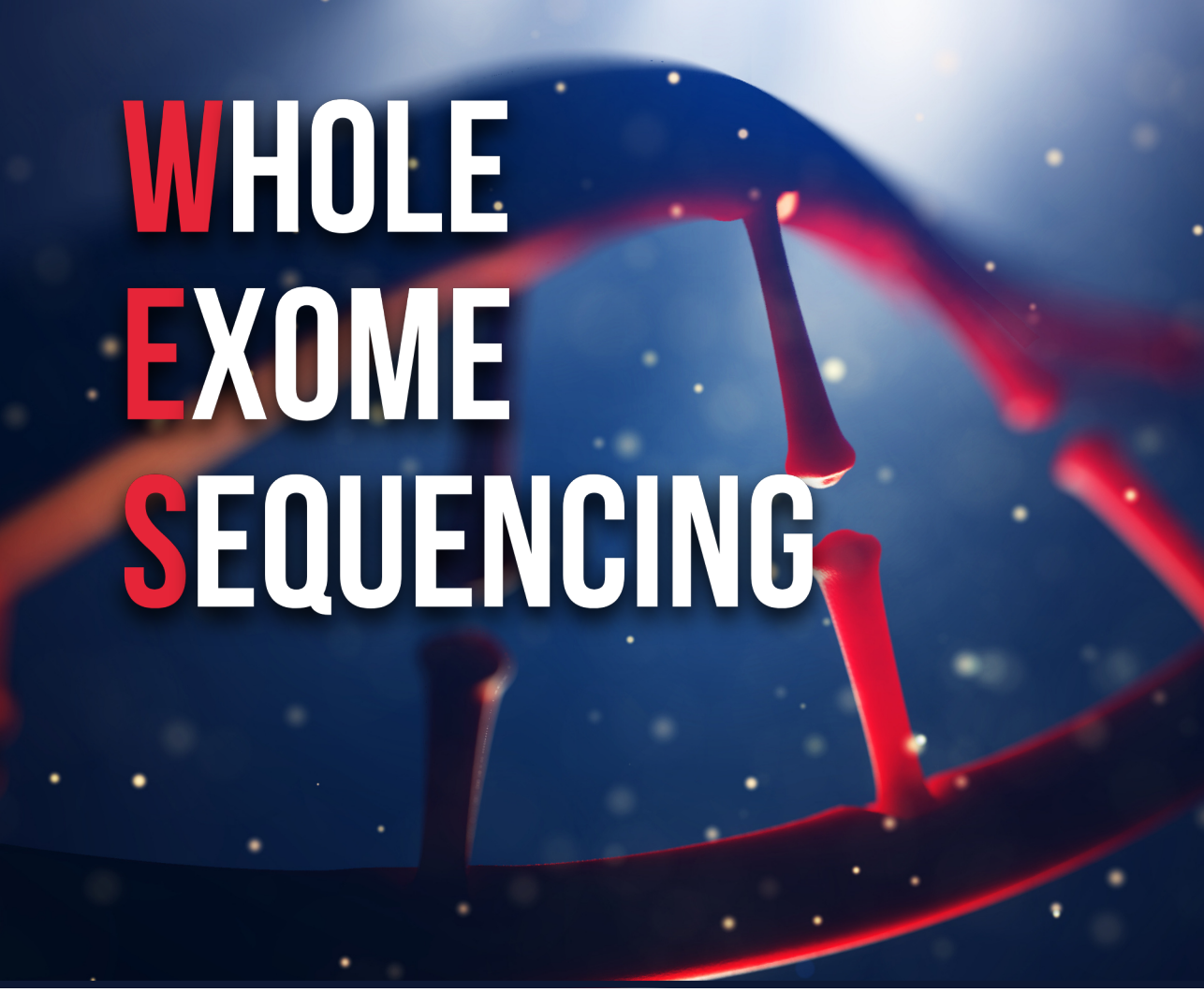

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Whole Exome Sequencing (WES) is one of the most comprehensive methods for diagnosing genetic diseases. Through this method, all coding DNA regions of approximately 20,500 genes in the human genome are sequenced and meaningful formations or changes that may be associated with diseases can be detected. This method can be used in different targets such as undiagnosed patients, cases that require investigation of a large number of genes for diagnosis, and in huge studies such as premarital carrier screening. In Whole Exome Sequencing, in addition to the genes that exist in the Clinical Exome, so many other genes that might be associated with different diseases and their association is not confirmed yet, are screened. The storage of these data may also provide a rapid diagnosis opportunity for the patient in the evaluation of the diseases that may occur in the future.

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