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BIG Data in Biology: A Review

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Keywords BIG Data; Bioinformatics; MapReduce; MPI; GPU. **ABSTRACT:** With significant advances in high-throughput sequencing technologies and consequently the exponential expansion of biological data, bioinformatics encounters difficulties in storage and analysis of vast amounts of biological data. The gap between sequencing throughput and computer capabilities in dealing with such big data is growing (Almeida, 2012). Considering the annual growth of data generation, the digital universe - data we generate annually - will reach 44 zettabytes, or 44 trillion gigabytes by the year 2020, which is ten times the size of the digital universe in 2013 (Kashyap,2014). Given the urgency of establishing a new computational framework, high-performance computing has become extremely important for large-scale data analysis. In this paper, we review various technologies such as MapReduce, message passing interface (MPI) and graphics processing unit (GPU) in context of bioinformatics and biological domain and also the work conducted by our lab in the area of network based therapeutics (Jaisri and Rawal et al 2016). © 2016 iGlobal Research and Publishing Foundation. All rights reserved.

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