



Cancer Genomics: Chapter 10. Genomic Resource Projects

Matthew Parker, Erin Hedlund, Jinghui Zhang

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The vast amount of genomic data being produced by the research community is becoming readily accessible to biomedical researchers and clinicians to apply to their cancer(s) of interest. The major cancer genome projects, among others, The Cancer Genome Atlas (TCGA), the International Cancer Genome Consortium (ICGC) and the Pediatric Cancer Genome Project (PCGP) are contributing to this genomic data goldmine by sequencing hundreds to thousands of cancer genomes and supplementing these data with analyses such as gene expression and methylation. In addition to the raw data that are being made available through large data warehouses, “Data Portals” are becoming the norm for accessing and analyzing these data by third parties. We describe key features of some of these portals and other tools for the analysis of next-generation sequencing and other genomic data.

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