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Clustering and graph mining techniques for classification of complex structural variations in cancer genomes

URL: https://www.nature.com/articles/s41598-022-07211-6

Authors: Gomez-Sanchez, Gonzalo / Delgado-Serrano, Luisa / Carrera, David / Torrents, David / Berral, Josep Lluis

Publication: Scientific Reports

Volume:

Barcelona Supercomputing Center - Centro Nacional de Supercomputación

Source URL (retrieved on *18 des 2024 - 07:50*): https://www.bsc.es/ca/research-and-development/publications/clustering-and-graph-mining-techniques-classification-complex