

OpenCB Home

Welcome to OpenCB!

The Computational Biology (OpenCB) open-source software initiative implements different high-performance and scalable open-source projects for the analysis of high-throughput genomic data.

OpenCB projects are developed and maintained by researchers of different universities and big projects such as University of Cambridge and Genomics England. OpenCB is open-source and freely available at <https://github.com/opencb>

Overview

OpenCB provides a high-performance and scalable platform for the analysis of high-throughput genomic data.

Contact

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Projects

CellBase

OpenCGA

IVA

Genome Maps

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Main Contributors

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