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In this webinar tutorial, we will discuss the following topics: Best practices for bioinformatic analyses using the versatile QIAGEN CLC Genomics Workbench; QC, adapter trimming and strategies for host read depletion and viral content enrichment; Simple pipelines for assembly and strain identification

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Training Webinars & Tutorials To facilitate access to the latest bioinformatics tools and resources, the OSTR has developed license agreements with a variety of bioinformatics companies that provide software packages to analyze genomic and proteomic data.

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Bioinformatics Training and Education Program. Home » Training » Class Archives. December 2020. ... Tutorial & Discussion. BTEP bulk RNA-Seq Weeks: Qiagen Ingenuity Pathway Analysis (IPA) Single-cell RNA-Seq Analysis on NIDAP – Tutorial & Discussion.

Class Archives - Bioinformatics Training and Education Program

The NIH Library Bioinformatics Support Program is presenting three unique sessions about QIAGEN software on May 8–10 in the NIH Library Training Room, Building 10. For more information about the classes and to register, use the links below. Ingenuity Variant Analysis (IVA) and Human Gene Mutation Database (HGMD) May 8, 9:30 a.m.–12:30 p.m.

NIH Library Bioinformatics Classes: QIAGEN | NIH Library

QIAGEN Bioinformatics was established in 2014, based on the acquisitions of Ingenuity, CLC bio, and BIOBASE. Today, the three legacy companies are joined by OmicSoft and N-of-One, working closely together to help our users innovate, integrate and translate basic discoveries into clinical care.

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