

User friendly and full support for Nanopore data.

A university Lab Director at an recommend CLC Genomics Workbench for this reason:

“ User-friendly GUI with advanced bioinformatics analysis. Incorporation of third generation sequencing technologies like Nanopore improve application for bioinformatic novices.

Source: Lab Director, Educational Institution



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TechValidate
by SurveyMonkey

Based on a response of 10 to the question "On a scale of 0-10, how likely would you be to recommend QIAGEN Bioinformatics?"