

Features



Predefined workflows for Genome variant detection (genome, exome and targeted sequencing), Transcriptome profiling, ChIPSeq analysis, *De novo* genome and Transcriptome assembly.



Flexibility to create customized pipelines and perform batch runs.



Interactive and graphically enhanced visualization of complex data.



Two versions:
SanGeniX Titanium-server based and SanGeniX Platinum-cloud based.



Multiple Linux platforms:
Ubuntu, CentOS and RHEL.

Applications



Plant
Genomics



Animal
Genomics



Clinical Diagnostics and
Personalized Medicine



Microbial
Genomics

Our Collaborators:

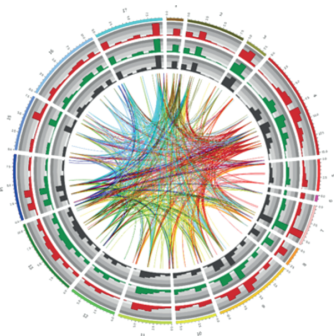
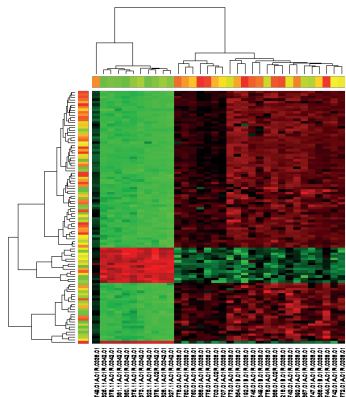
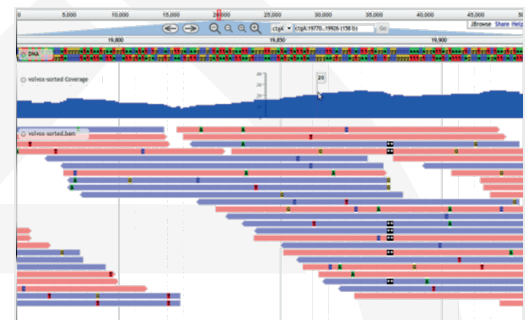


Funded by:

Comparison Matrix

Features	CLC Genomics	DNA Nexus	SeqMan	Strand NGS	Seven Bridges	i-OMICS	Globus Genomics	Partek Genomics	Next Gene	SanGeniX
Supports all major workflows	✓		✓	✓		✓		✓	✓	✓
Interactive and enhanced visualization for all workflows			✓	✓		✓		✓	✓	✓
Custom workflows	✓				✓	✓	✓	✓	✓	✓
Supports multiple sequencers	✓	✓	✓	✓		✓		✓	✓	✓
Data, workflow and results sharing					✓	✓	✓	✓		✓
Cost Effective							✓			✓
Mobile App										✓

Visualizations:



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