

User Manual

Partek Genomics Suite is a comprehensive suite of advanced statistics and interactive data visualization specifically designed to reliably extract biological signals from noise. Designed for high-dimensional genomic studies containing thousands of samples, Partek Genomics Suite is fast, memory efficient and will analyze large data sets on a personal computer. It supports a complete workflow including convenient data access tools, identification and annotation of important biomarkers, and construction and validation of predictive diagnostic classification systems.

- [Lists](#)
- [Annotation](#)
- [Hierarchical Clustering Analysis](#)
- [Gene Ontology ANOVA](#)
- [Visualizations](#)
- [Visualizing NGS Data](#)
- [Chromosome View](#)
- [Methylation Workflows](#)
- [Trio/Duo Analysis](#)
- [Association Analysis](#)
- [LOH detection with an allele ratio spreadsheet](#)
- [Import data from Agilent feature extraction software](#)
- [Illumina GenomeStudio Plugin](#)

Additional information can be found in the manual for [Partek Genomics Suite version 6.6](#).

Additional Assistance

If you need additional assistance, please visit [our support page](#) to submit a help ticket or find phone numbers for regional support.



Your Rating:  Results:  47 rates