White papers

White papers found in this section provide detailed information about Partek Genomics Suite functionality.

- Allele Intensity Import
- Allele-Specific Copy Number
- Calculating Genotype Likelihoods
- ChIP-Seq Peak Detection
- Detect Regions of Significance
- Genomic Segmentation
- Loss of Heterozygosity Analysis
- Motif Discovery Methods
- Partek Genomics Suite Security
- Reads in RNA-Seq
- RNA-Seq Methods
- Unpaired Copy Number Estimation

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: 35 rates

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