

## Optional: Import a Partek Project from Genome Studio

An Illumina-type project file (.bsc format) can be imported in Illumina's GenomeStudio® (please note: to process 450K chips, you need GenomeStudio 2010 or newer) and exported using the Partek Methylation Plug-in for GenomeStudio. For more information on the plug-in, please see the [plug-in user guide](#). The plug-in creates six files: a Partek project file (\*.ppj), an annotation file (\*.annotation.txt), files containing intensity values (\*.fmt and \*.txt), and files containing -values (\*.fmt and \*.txt) (Figure 1).

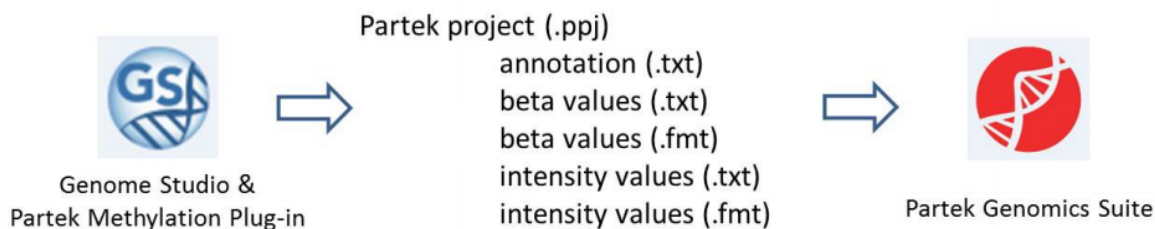


Figure 2. Output of Partek Methylation Plug-in for GenomeStudio

To load all the files automatically, open the .ppj file as follows.

- Select **Methylation** from the *Workflows* drop-down menu
- Select **Illumina BeadArray Methylation** from the *Methylation sub-workflows* section
- Select **Import Illumina Methylation Data** from the *Import* section
- Select **Load a project following Illumina GenomeStudio export** from the *Load Methylation Data* dialog

## Additional Assistance

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



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