

Adding an annotation link

While many types of data sets are automatically linked with appropriate annotation files upon import, if this does not occur, a spreadsheet can be manually linked with an annotation file.

- Right-click *Breast_Cancer.txt* in the spreadsheet tree
- Select **Properties** (Figure 1)

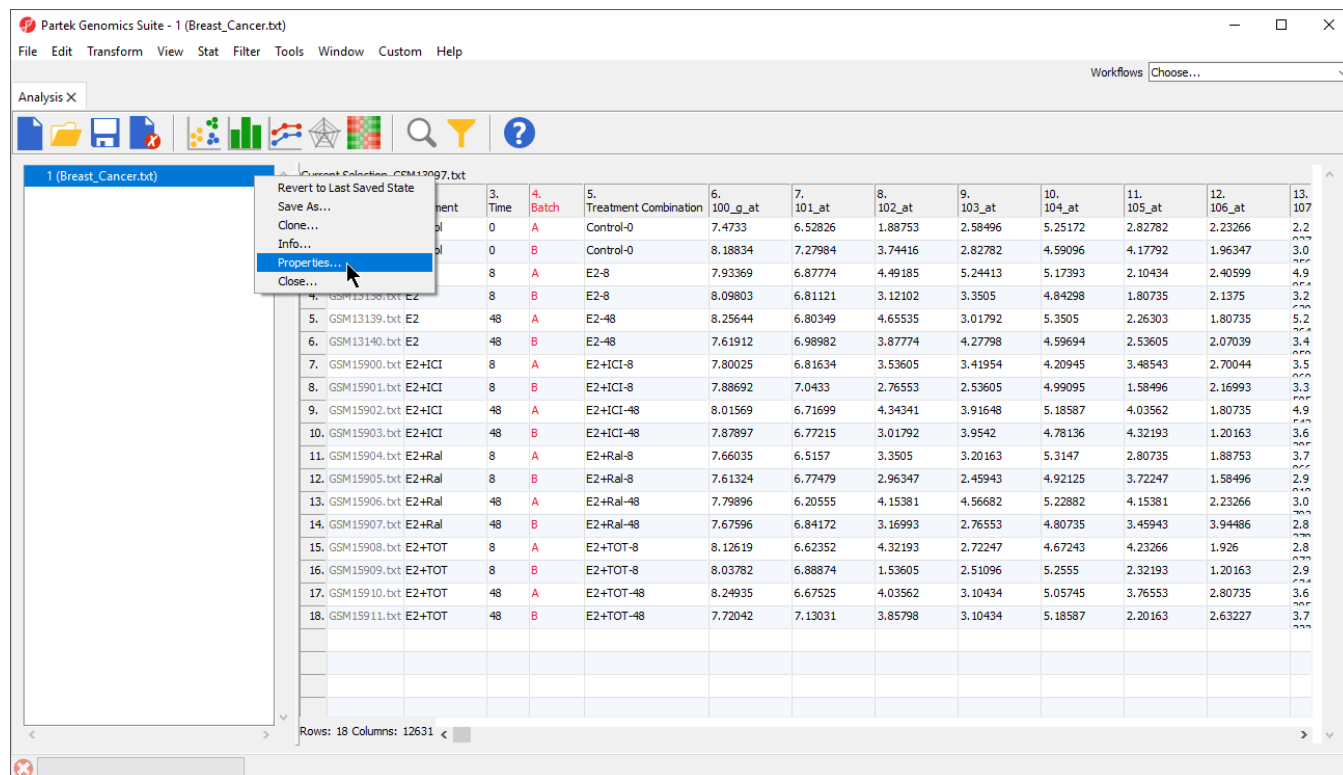


Figure 3. Selecting file properties for a spreadsheet

Configure the *Configure Genomic Properties* as shown (Figure 2) with the following steps:

- Select **Gene Expression** from the *Choose the type of genomic data* drop-down menu
- Select **Feature in column label**
- Select **Browse...**
- Select **HG_U95Av2.na36.annot.csv** from the microarray libraries folder
- Select **Set Column**
- Select **Gene Symbol** from the *Choose column containing gene symbol/microRNA name* dialog
- Select **Homo sapiens** and **hg19** from the *Species and Genome Build* drop-down menus

Figure 4. Configure the genomic properties dialog as shown

There is now an * after the spreadsheet name in the spreadsheet tree. This indicates an unsaved change has been made to the spreadsheet.

- Select () to save the changes

« [Importing the data set](#) [Exploring the data set with PCA](#) »

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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