Highlight and select

There are many cases in which analyses of different subsets of the samples or species are required. This can be easily achieved, without the need to create large numbers of separate datasheets, by temporarily selecting subsets from a single sheet, analysing them (and thus creating new branches on the Explorer tree, with the results windows listing the selection used for any particular branch), and then restoring the full data set. There are several different ways to select subsets, described below, but it is important to keep in mind the distinction between highlighting and selection. The act of clicking on a row and/or column header *highlights* that row and/or column; it does not *select* it. Once you are happy that you have highlighted the correct set of samples (and/or variables) you can select them using the **Select>Highlighted** menu. Highlighting is just an intermediate stage, and has functions other than selection (e.g. to identify samples that need individual transformation, whilst the rest of the matrix remains unchanged - see next section). Alternatively, highlighting can be bypassed altogether and selection made by other direct choices from the **Select** menu.

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