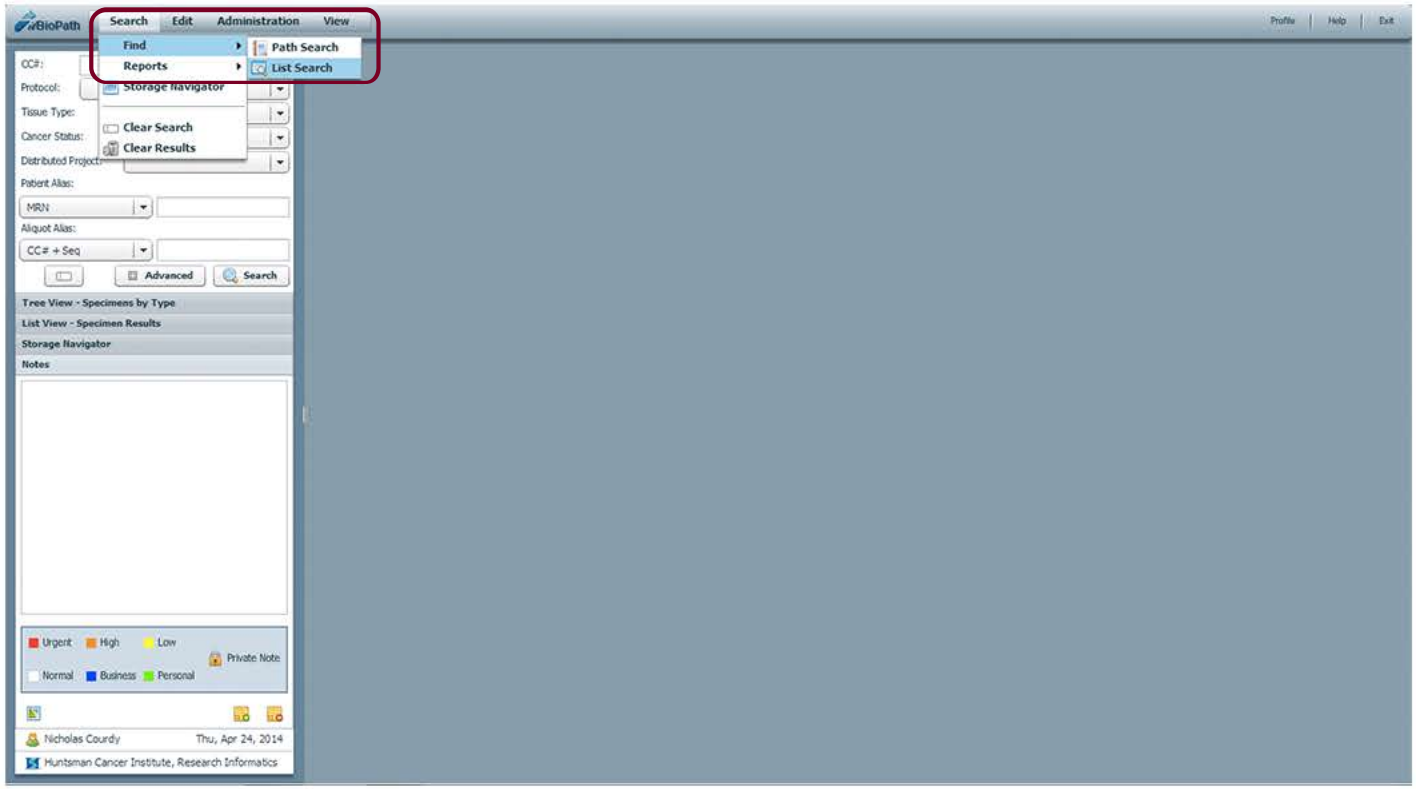
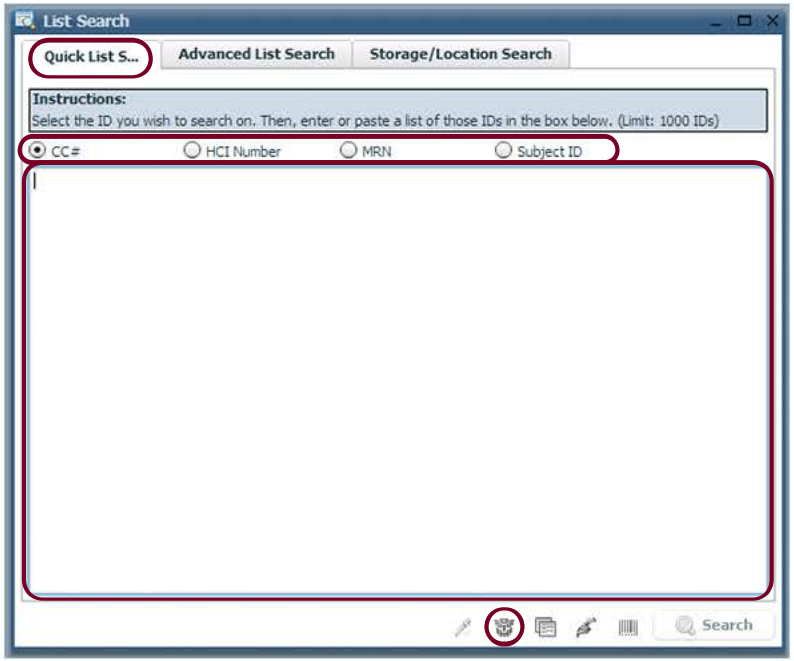
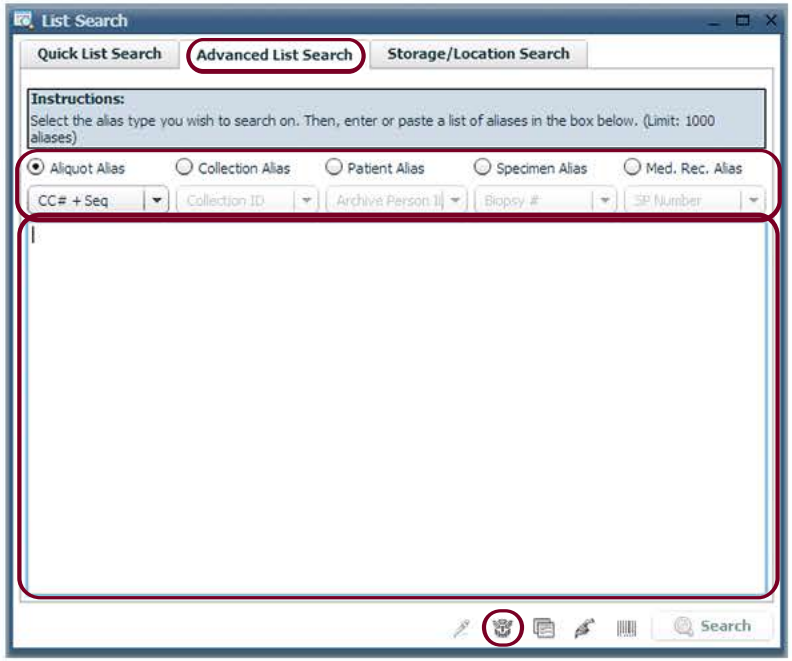
itBioPath HOW-TO

Performing a Batch Transformation

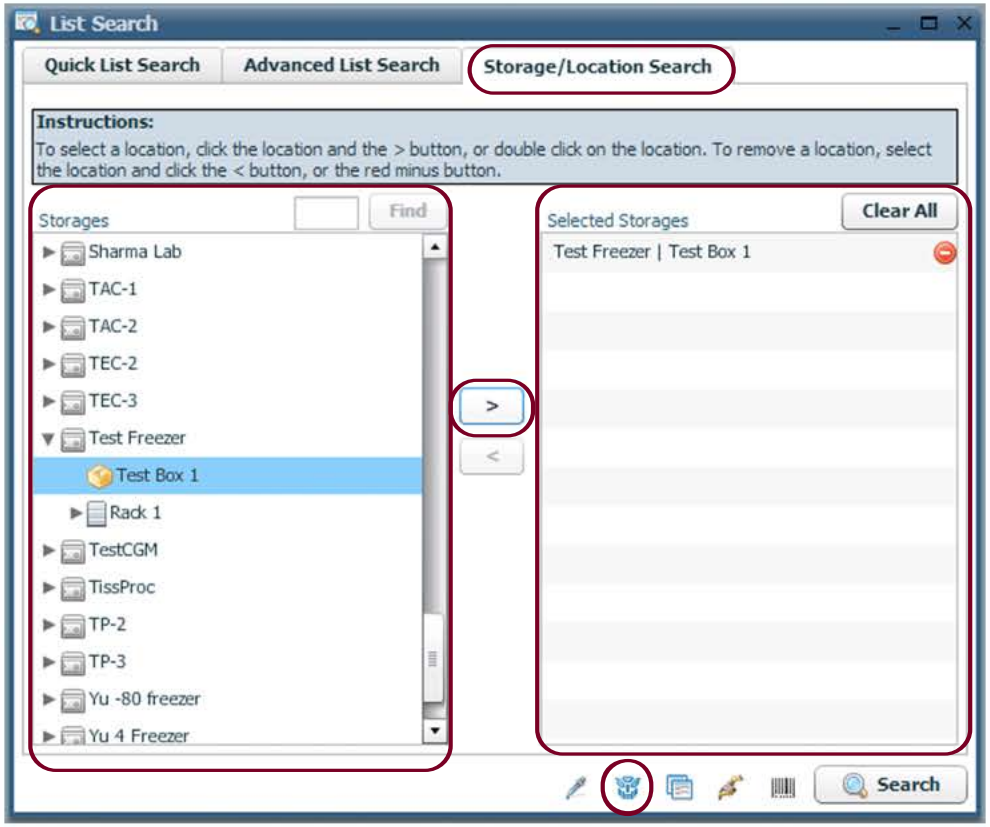


The first method for searching data in itBioPath for a batch process is the List Search. Navigate to Search > Find > List Search. A new window will open.

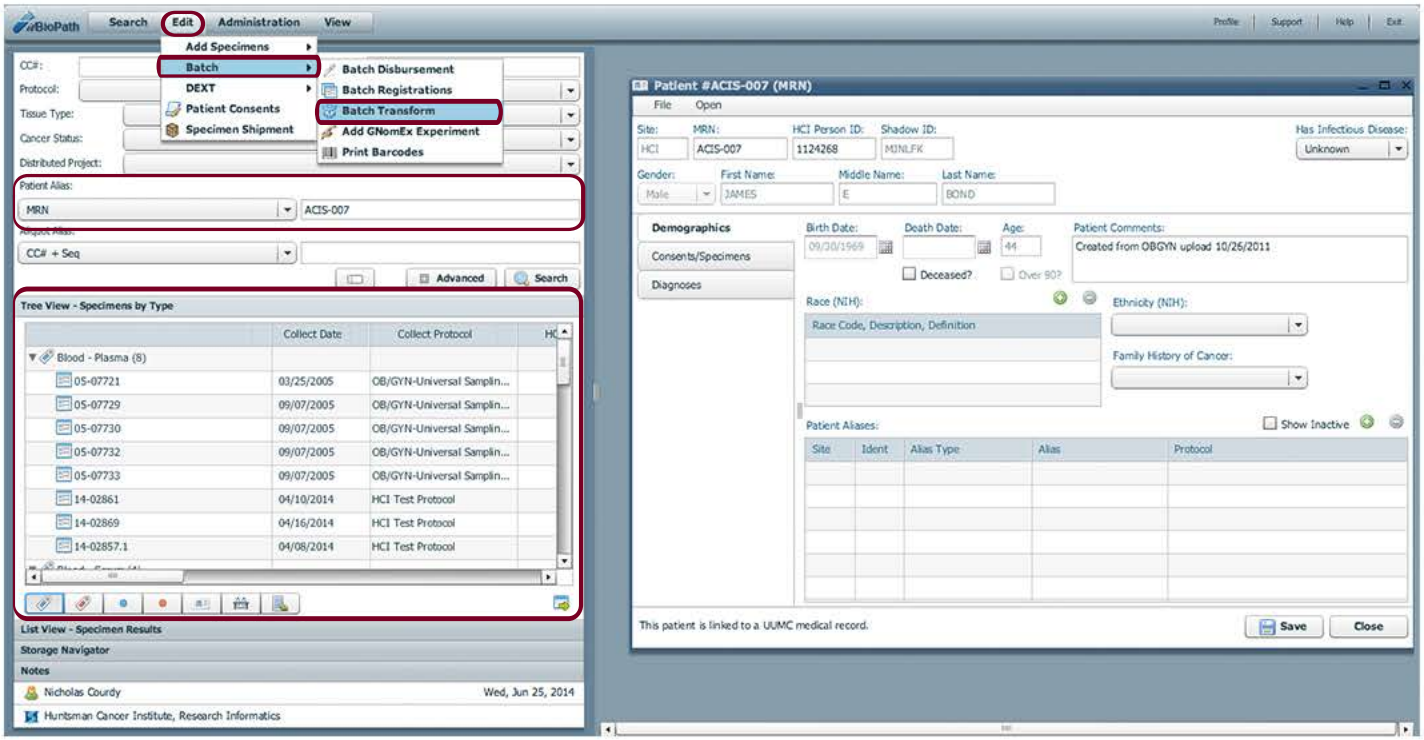




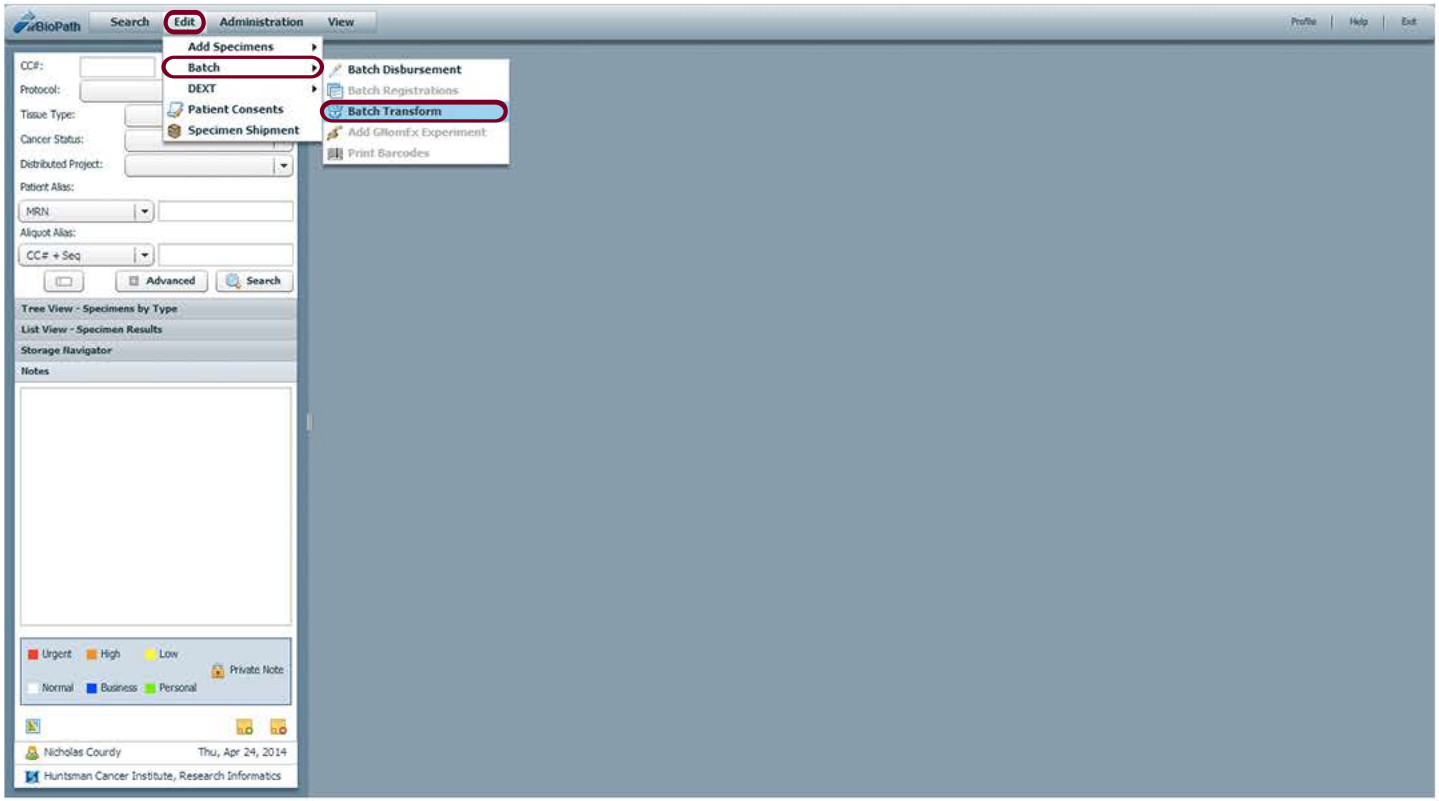
The list search window has 3 tabs. The ‘Quick List Search’ and ‘Advanced List Search’ make it possible to choose from multiple selections and enter specific identifiers in the text field. When the information is correct click on the flower icon to start a batch transformation with the information entered.



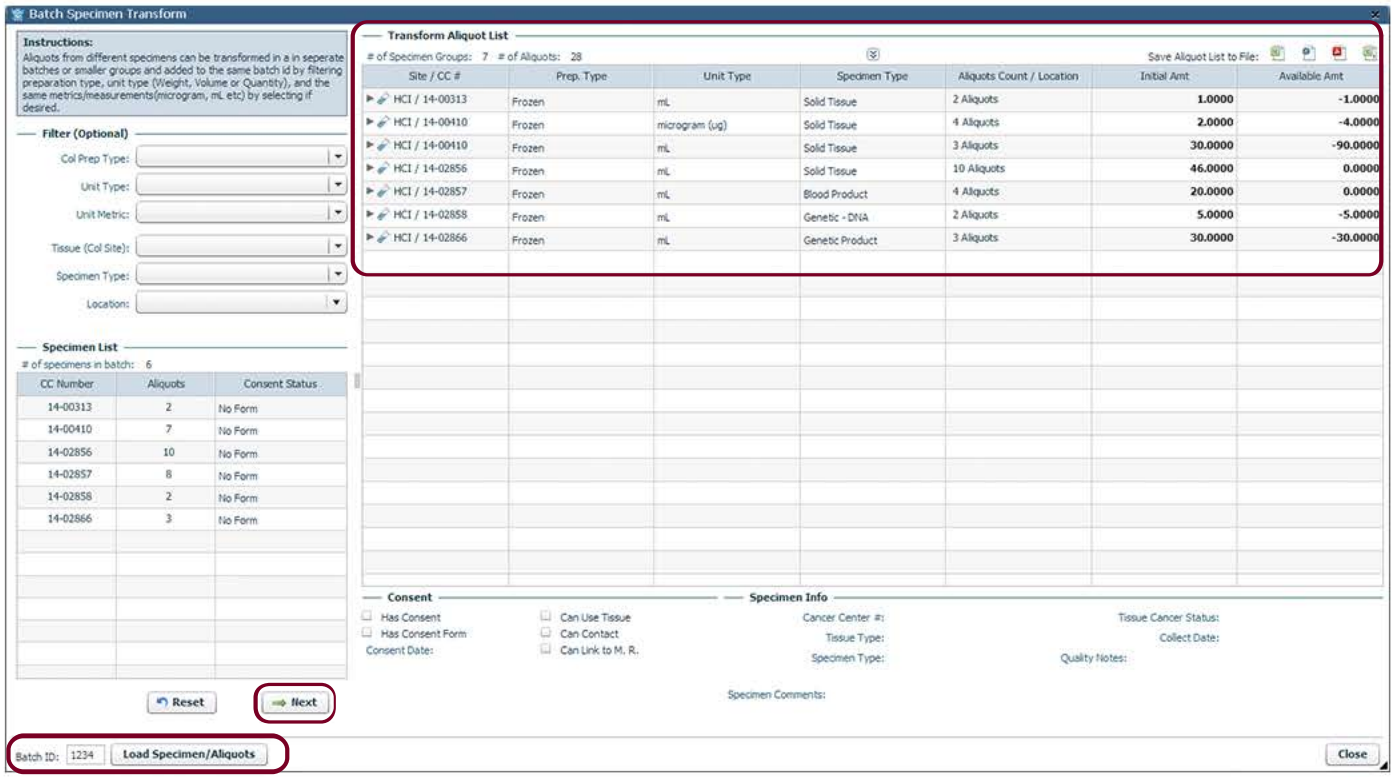
The third tab of the List Search window is the 'Storage/Location Search'. This tab makes it possible to search by a box or boxes in various freezers. Find the specific box ready for transformation and click the arrow icon to move it into the 'Selected Storage' column. When all the boxes have been moved, click on the icon to start a batch transformation of the specimens in the boxes selected.



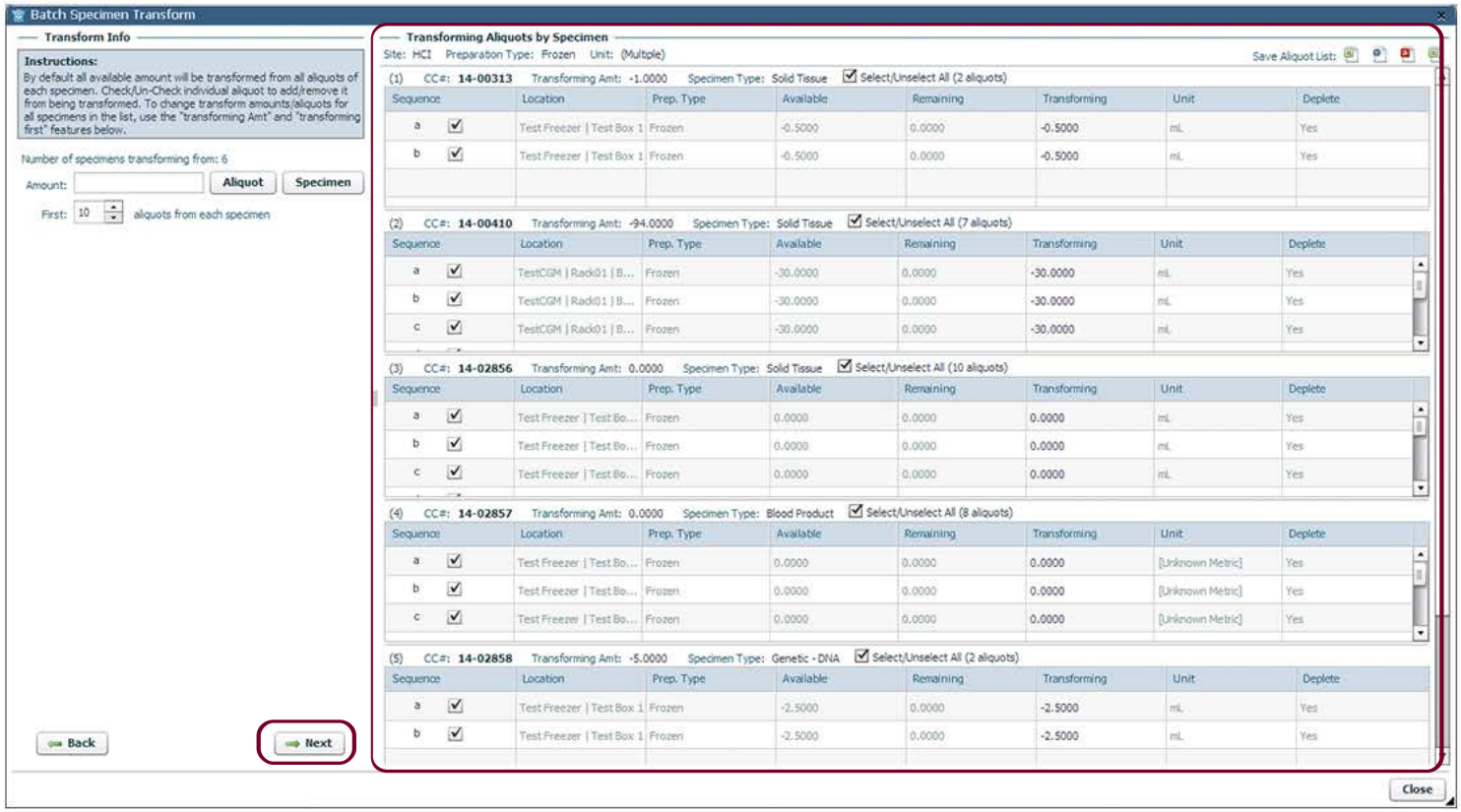
The second method for searching data in itBioPath for a batch process is the Basic/Advanced search on the Dashboard. After a search is conducted and information is retreived navigate to Edit > Batch > Batch Transform.
In this example a search by Patient MRN was conducted. itBioPath found the specimens associated to the patient, and now a transformation of these specimens can be initiated.



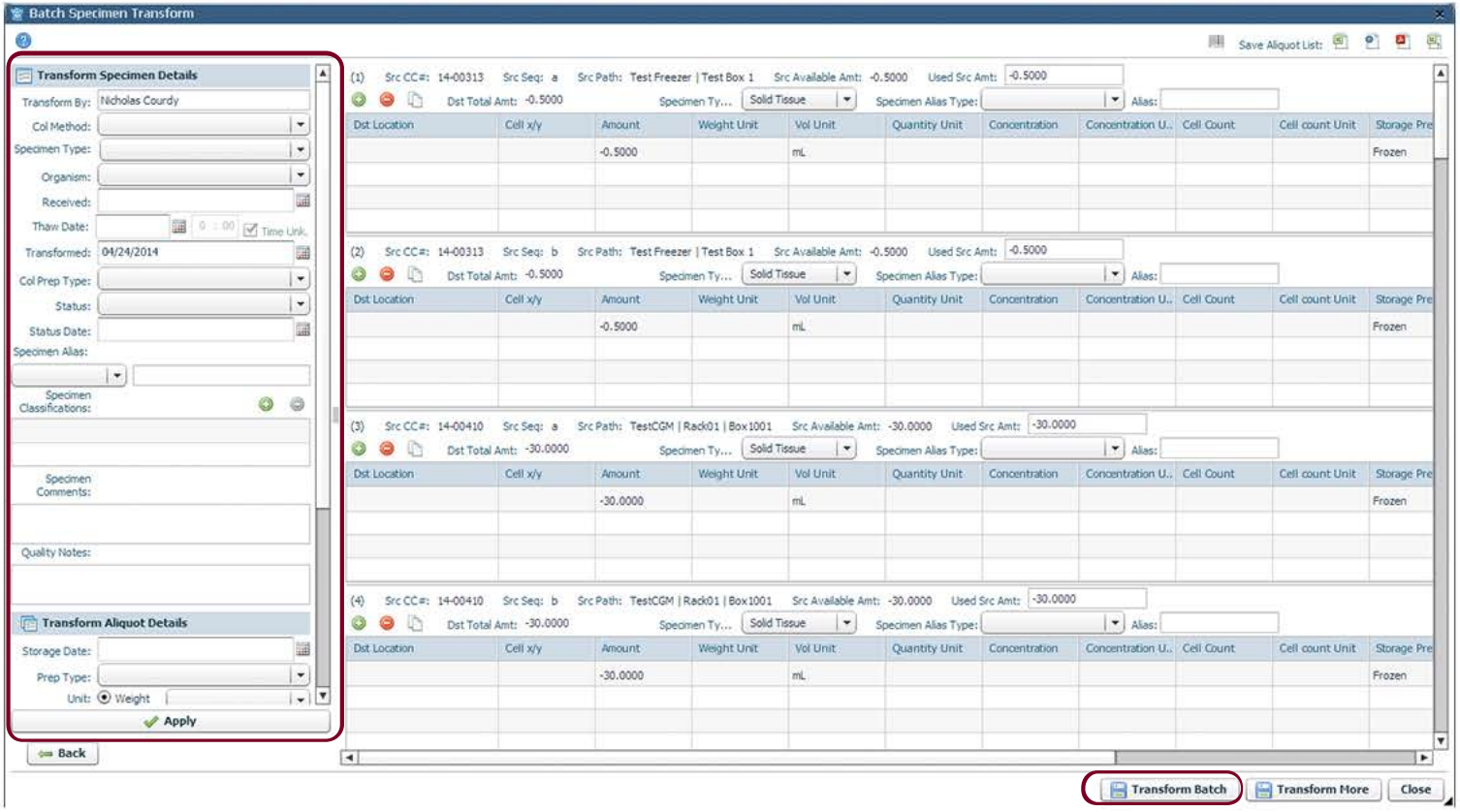
The third method for searching data in itBioPath for a batch process is a Batch ID. This method can only be used if a search has not already been conducted in itBioPath.
If itBioPath is blank, navigate to Edit > Batch > Batch Transform.
A new window will open.



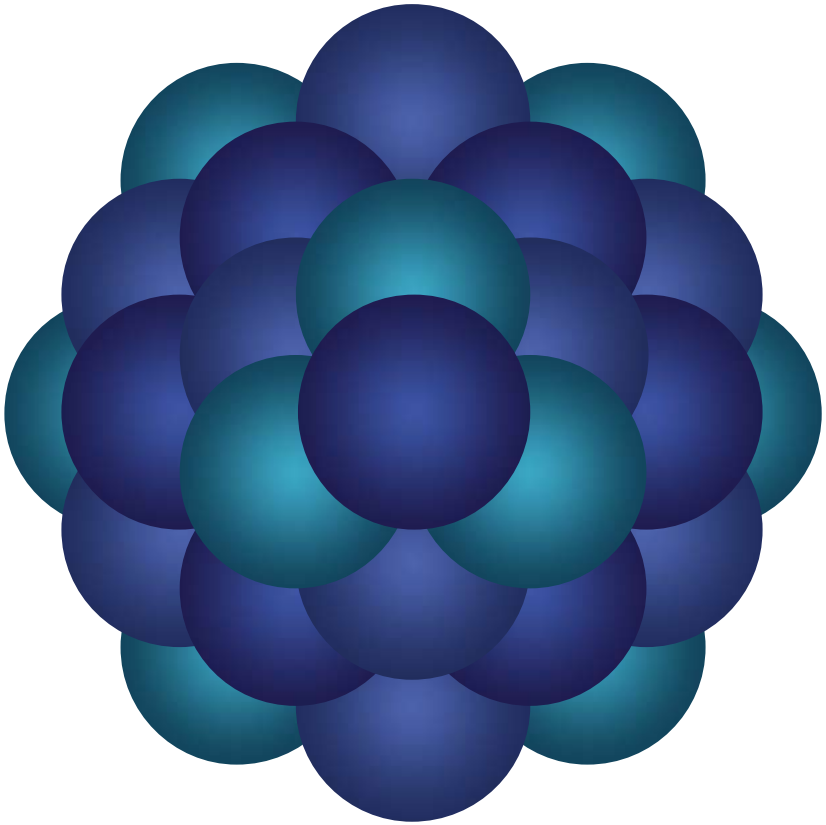
In the bottom left of the ‘Batch Specimen Transform’ window you can load specimens by a Batch ID. A Batch ID is a specimen alias that can be assigned to multiple specimens. Available aliquots for transformation will appear. *This window will look similar if transforming using the List Search or Dashboard search methods. When the Aliquots have been loaded click ‘Next’.



The second step of the Batch Transformation determines what sequences will be transformed. Once the aliquot information is correct, click 'Next'.



The third and final step of the transformation specifies details about the transformation. This includes specifying aliquot amounts that are being used for the transformation. When the information is correct, click ‘Transform Batch’.



Questions?

Email ‘Informatics - itBioPath Support’ email group