

Beaumont

Soft Lab End User Training

Specimen Tracking: Basics of Specimen Receiving & Archiving

Objectives

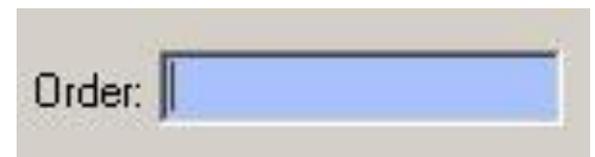
- After viewing the following tutorial, the user will have visited the following topics:
 - Specimen receiving functions
 - How to place specimens into archiving
 - Removing a stored specimen from archiving

Specimen Receiving Functions

- Specimen that have been collected, must be received by the lab staff in SOFT LAB.
- Specimens can be received individually
 - Order Entry screen (seen in past tutorials)
- Specimens can also be received in batches
 - Specimen Receiving Icon
- Specimen Tracking is the SOFT Lab application that tracks the location and time of specimen receipt in the lab

Receiving Specimens in Batches

- To receive multiple specimens at one time:
 - Click the Specimen Receiving icon from the Soft Lab home screen
- *Have the Batch Receiving function box checked
- Place your cursor in the Order box (becomes blue), and begin scanning your specimen barcodes

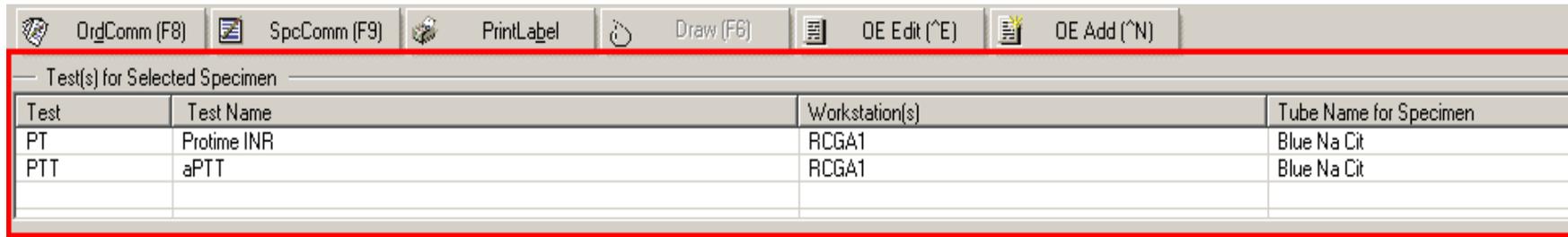


Receiving Specimens in Batches

- As you scan in specimens, **always** verify that patient information scanned matches information on the specimen
- While you scan the center of the screen populates with Order Information
- The bottom of the screen displays test information

The screenshot shows a software window titled "Receiving" with a menu bar (File, Edit, View, Specimens, Assign Collection, Label Printing, Collection, Receiving, Status, Utilities, Tools, Window, Help) and a toolbar. Below the toolbar, there are tabs for "1 List Receiving", "2 Non-List Receiving", "3 Scanner Receiving", and "5 Upload From Scanner Receiving". An "Order:" field and a "Batch Receiving" checkbox are visible. The main area contains a table with the following columns: Order Information (Order, Prior, Spc, Wk, MRN, Barcode, Name, DT, OC, SC), Collection (Y, N, C, By), C DATE, C TIME, Receiving (Y, N, R, X, R, By), R DATE, R TIME, and Order in. A red box highlights the "Order Information" section. At the bottom, there is a section titled "Test(s) for Selected Specimen" with columns for Test, Test Name, Workstation(s), and Tube Name for Specimen. A red box highlights this section.

Receiving Specimens in Batches



Test	Test Name	Workstation(s)	Tube Name for Specimen
PT	Protime INR	RCGA1	Blue Na Cit
PTT	aPTT	RCGA1	Blue Na Cit

- The bottom portion of the screen displays which tests are ordered on the tube that was scanned in the **Order** box above
- There can be multiple tests ordered on the same tube
 - Example: PT and PTT as shown above were on the same tube

Receiving Specimens in Batches

Order information										Collection				Receiving								
Order	Prior	Spc	Wk	MRN	Barcode	Name	DT	OC	SC	Y	N	C By	C DATE	C TIME	Y	N	R	X	R By	R DATE	R TIME	
262600007	R	BOU	RCGA1	6000158	B6	TEST, ALLERGY				✓		AMARM	01/26/2012	00:00	✓				AMARM	01/26/2012	14:02	

- The top portion of the screen populates with collection and receipt information
- *If there is **no collector information** you must type **COLLI** as the collector ID (Inpatient Collector)
 - Enter the collection time printed at bottom of tube
- *However, If there is no collection time printed on tube:
 - Enter time as **00:00**

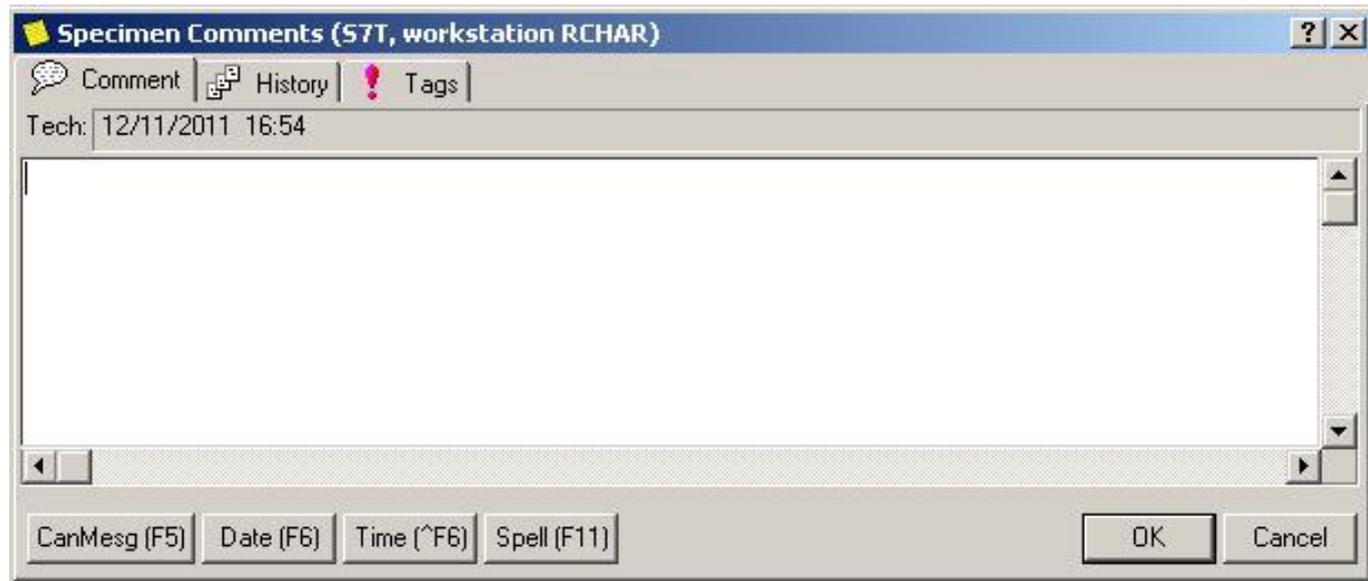
Adding Comments to Received Specimens

- There are cases where you must write a comment on the specimen(s) received
 - Example: Receiving specimens in STAT lab that will then be sent to the Core lab via pneumatic tube system
- **OrdComm**-Order Comment
 - Comment pertains to all specimens within that Order
- **SpcComm**-Specimen Comment
 - Comment pertains to only that specimen (tube, vial etc)

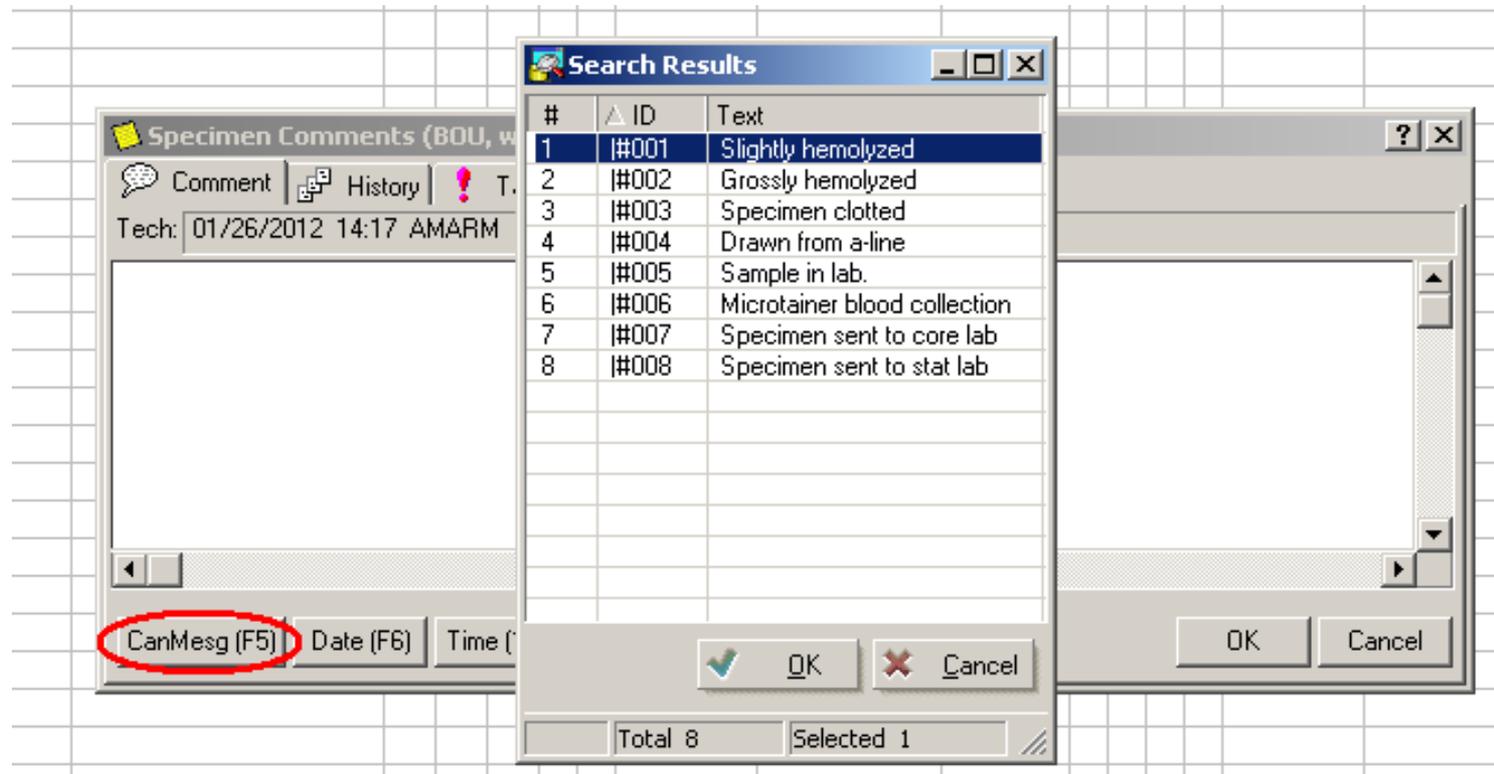


Adding Comments to Received Specimens

- Click either **OrdComm** or **SpcComm**
- Type your own message in the Text box



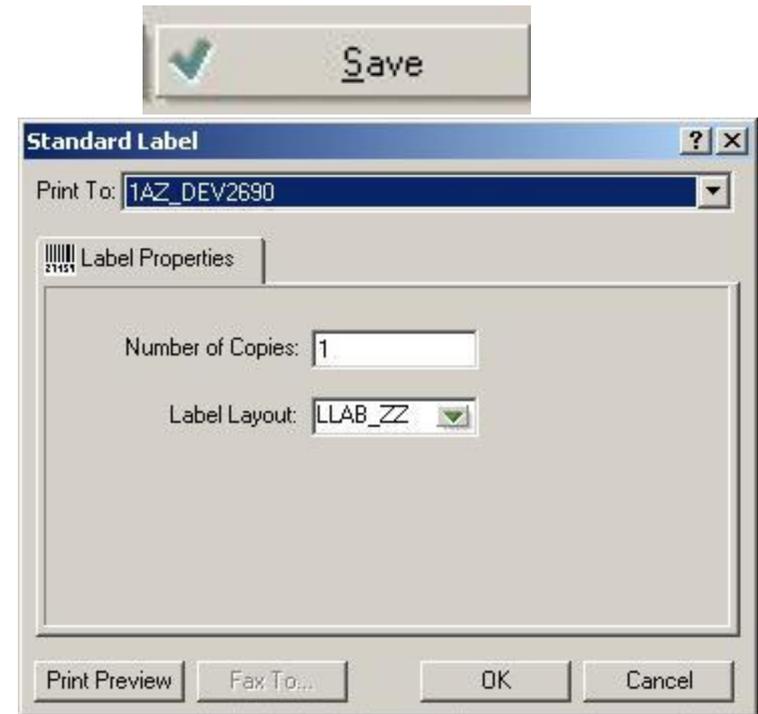
Adding Comments to Received Specimens



- You can also choose a comment from one of the **Canned Messages** available
- Recall that you must proceed comment with (?) to keep message internalized

Finalizing Specimen Receipt

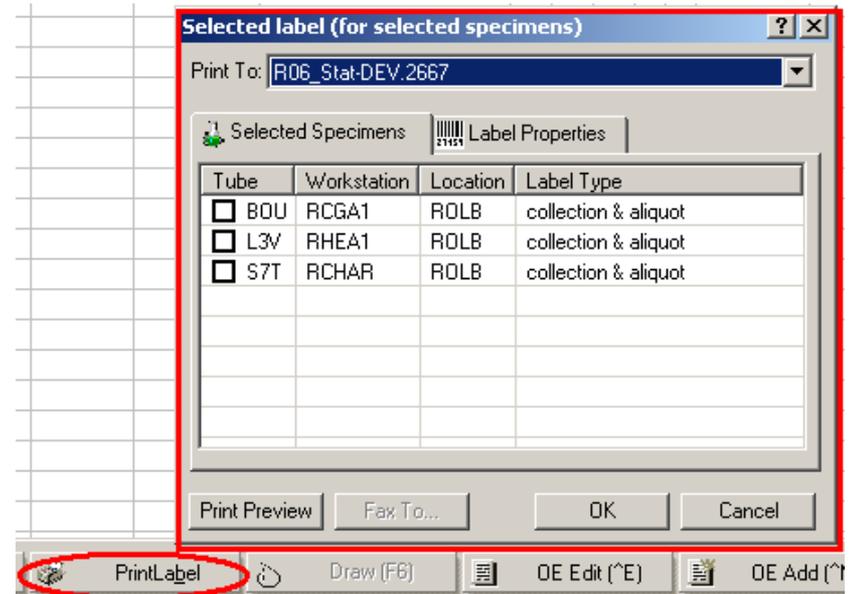
- To finish the process, Click the **Save** button at the bottom right of screen
- A Standard Label printer box will pop-up
- Click **Cancel**



Printing Additional Labels

- If you wish to print additional labels for the received specimens, you may do so **before** pressing the **SAVE** button
- Click the **PrintLabel** button at the middle of the screen
- Check the boxes for the specimen types which you wish to print extra labels

Example BOU=Blue top tube



Placing a Specimen into Archiving

- Archiving is done through the Specimen Tracking function, a.k.a. “little blue feet”
- Click on the Tracking Icon from the home screen



Placing a Specimen into Archiving

- Specimen Tracking Function Menu will appear
- The function you will want is **Storage**
- Select the correct **Storage** rack for the specimen type you wish to archive, click **OK**

Specimen Tracking Function Menu

Terminal Name: A0934

#	Status	Location	Place	Type	Actions
1	D - Ordered	D - Ordering Location	RSTAT		
2	C - Collected	C - Collection Center	RSTAT		
3	R - Received	R - Receiving	RSTAT		
4	X - Cancelled/Error	X - Cancelled/Error	RSTAT		
A	S - Storage	Z - Stat Lab Walk-In	SLCHEM		R(###-###,+1)
B	S - Storage	Z - Stat Lab Walk-In	SLHEMO		R(###-###,+1)
C	S - Storage	Z - Stat Lab Walk-In	SLCOAG		R(###-###,+1)
D	S - Storage	Z - Stat Lab Walk-In	SLALIQ		R(###-###,+1)
E	F - Removed from Storage	D - Testing Department	RSTAT		

OK Cancel

Storage Rack Options

- Available storage rack options at the RO Core Lab archiving station:
 - RO Chemistry Remco
 - RO Coagulation Remco
 - RO Hematology Remco
 - Chemistry Walk In –(FLD) Fluids
 - Chemistry Walk In –(UCUP) Urine Cups

Placing a Specimen into Archiving

- Once you have selected the correct rack, scan your specimen
- Type your specimen tag number into the Tag space
- Then Click **Add Specimen** to begin archiving your rack

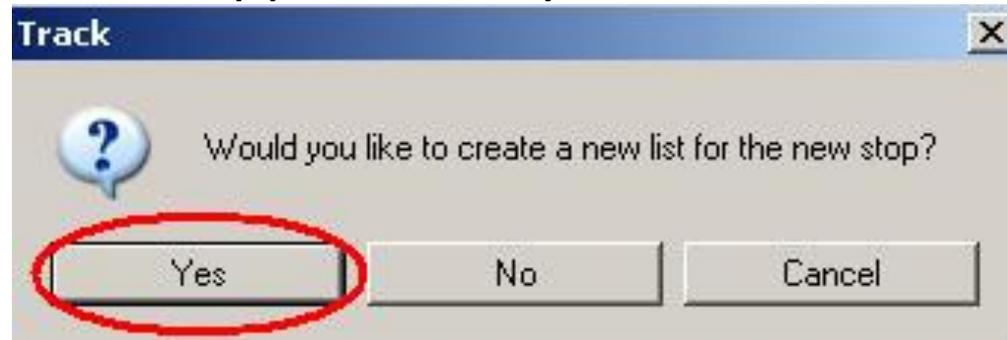
The screenshot shows a software interface with two main input panels. The left panel has fields for 'Place: RSTAT', 'Comment:', and 'Tag:'. The right panel has fields for 'Place: SLCHEM', 'Comment:', and 'Tag: 078-001'. The 'Tag: 078-001' field is circled in red. To the right of these panels is a 'Printer Name:' field and three buttons: 'History', 'Lists', and 'Add Specimen'.

Rack Tag Information

- Rack Tag number assignments follow this pattern: ###-###
- There are two parts to the Tag number:
 - First being the rack number
 - Second part of the tag is the position in the rack
- There are 108 total spots in each rack

Placing a Specimen into Archiving

- Continue to scan specimens until you have entered all the specimens you wish to archive
- *Scan carefully and verify that the specimen tag matches the physical spot in the rack where you are placing the specimen
- When finished, Click **FILE** then select **SAVE**
- A **Track** box will appear; always select **YES**

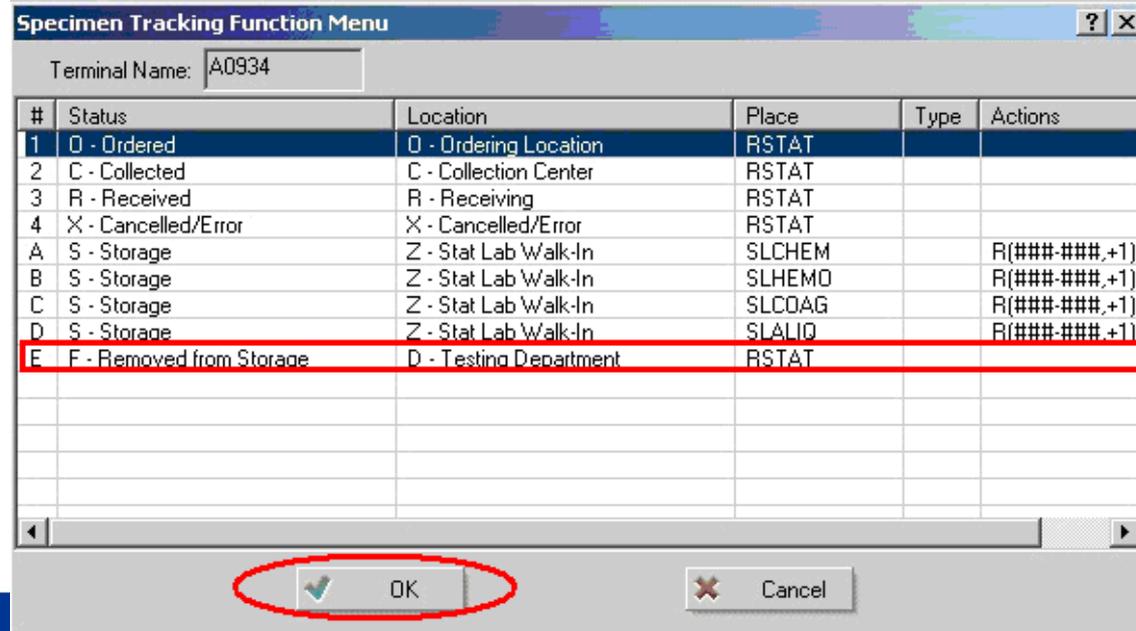


Placing a Specimen into Archiving

- Each time you scan specimens into Storage you must print a Tracking list
- When printing be sure to change the printer from **Modem** to your local printer by using the drop down menu
- Select **OK** to print your list
- Once list has printed, write the specimen Rack Number on first page

Removing a Specimen from Archiving

- If you wish to “pull” a specimen from storage:
- Go to Specimen Tracking icon
- At Tracking Function menu, select
 - **F-Removed from Storage** option



Removing a Specimen from Archiving

- Take your specimen and scan the barcode in the box
- The “New Stop” for that specimen will become “Removed from Storage”
- In the **Comment** field, enter a reason for pulling the specimen
 - Example: Added-on for Lipase

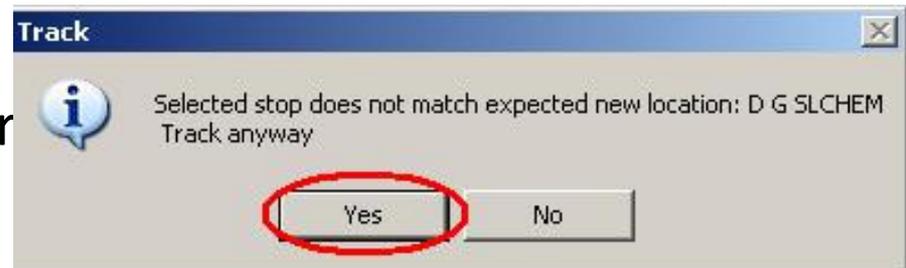
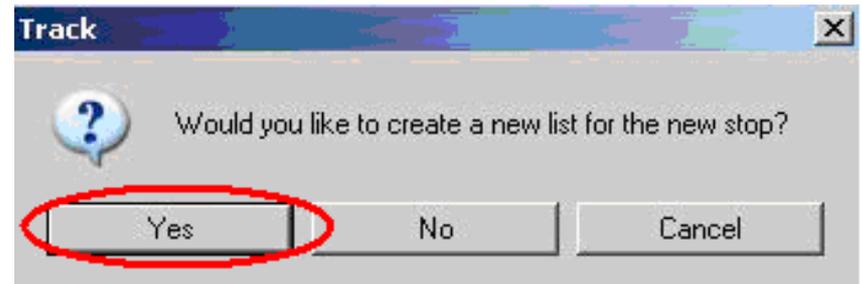


The image shows a screenshot of a software interface titled "New Stop". It contains several input fields:

- Status: A dropdown menu with the letter "F" selected, and a text box containing "Removed from Storage".
- Location: A dropdown menu with the letter "D" selected, and a text box containing "Testing Department".
- Place: A text box containing "RSTAT".
- Comment: A text box containing "Comment Here", which is circled in red.
- Tag: An empty text box.

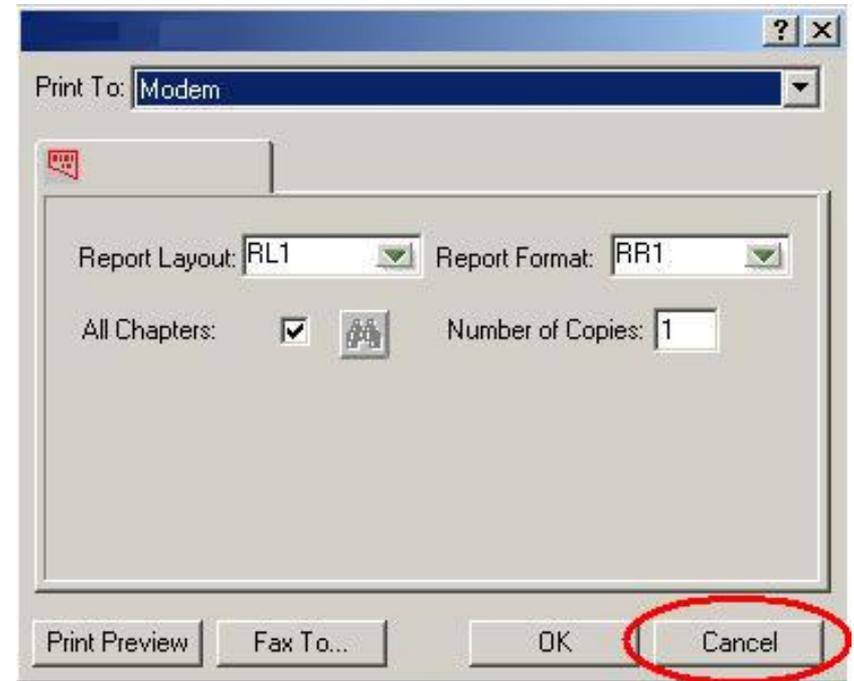
Removing a Specimen from Archiving

- After you have written a comment, click on **File>SAVE**
- The tracking pop-up window will appear
- Click **YES**
- A second pop-up window (Image at Right) will appear
- Once again, click **YES**



Removing a Specimen from Archiving

- This time it is not necessary to print a list with your tracking action
- At the print pop-up window, click **CANCEL**



Credits

- Material from this tutorial is credited to the following individuals:
 - Daniel Kirchhoff
 - Ronald Raese
- Questions on content and/or input or critique on required modifications should be addressed to:
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