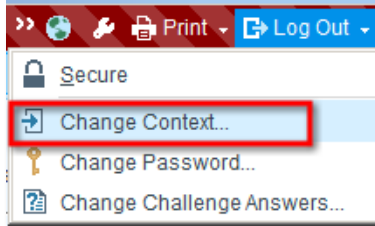


# Flow: LTA and Tech Chimerism Guide:


## Receiving the Chimerism Specimen:

1. **Flow LTA:** From the upper right-hand corner of the AP Beaker screen, click on the arrow next to the Log Out button. Choose Change Context...



2. **Flow LTA:** Change the context of the POC- login from “Lab Surgical Path” to “Lab SIC” (You are now in CP Beaker):



3. **Flow LTA:** On your Beaker Lab Dashboard – OHSU, click on the **Specimen Receiving** link or click on the  Receiving tab near the top left of your screen.

4. **Flow LTA:** In the **Receiving** activity, make sure the **Scan** button is selected, and scan in your specimen to receive it. You can also click the Specimen Lookup button and type the ID manually.



5. **Flow LTA:** After you receive the specimen, click on the **Summary** tab for this specimen. Scroll down the **Summary** pane until the section where the order questions are showing is visible. Under **Specify Sorted Populations/Fractions**, there should be a comment as to what populations or fractions the physician is interested in. Write down the requested populations on the specimen label for the Flow Techs to process. Give the sample and the label to the Flow Techs.

Receiving - OHSU All Specimens to Receiving - 1 of 1 selected

Specimen: 17IM-340F0001 List: Chimerism Sorted Cells, DNA, Blood Patient: Beaker, Hank MRN: 03438074 List: 17-IN811 2

**Summary** Details


Order: 162736255 12/06/2017 1301 by Kathryn Stinson  
 Enc. Department: LAB SURG  
 Dx code:   
 Diagnoses:   
 Date/Time: 12/6/2017 01:09 PM  
 Collector:   
 Department:   
 Draw type:   
 Type: Blood  
 Source: Blood  
 Lab comments:

**CHIMERISM SORTED CELLS, DNA, BLOOD**

Order Question: Specify Sorted Populations/Fractions Answer: CD3 and CD33

Supplemental Indications:  
 Donor Name/ID Number:  
 Gender of Donor:  
 Date of Transplant:  
 Shipping Instructions:  
 Immuno: whole bld rocker

## Processing the Chimerism Specimens:

- Flow Tech:** When you are ready to result the Cells Sorted and the Purity, go to the **Outstanding List** under the **Outstanding List – Core Lab** view. To change your view, click  Views button and pick the view you want. Double click on the specimen from the **Outstanding List** to open up **Result Entry**. Click on the **Edit** button.

**Result Entry**

Views - Open Remove All Comm Log Labels & Docs Tracking Result Correction Verify

Specimens  
 Templates  
 CHIMERISM SORTED  
 17IM-298F0001  
 CHIMERISM SORTED  
 17IM-340F0001  
 CHIMERISM SORTED

**Summary**

**17IM-340F0001**

**Beaker, Hank (MRN 03438074)**  
 M, 23 yrs, 1/1/1994

**Blood, Blood**  
 Collected 12/6/2017 1309  
 Container: 3 GREEN NaHep-4mL


Specimen Follow Ups

Edit Save Next Clear Changes Repeat Method Mnemonic Keys Scan

(# Rst) Component	Value	A	D	L	R	Units	Ref. R
1 ANTIBODY SORTED - 1	cd3						
2 CELLS SORTED - 1	50000						
3 PURITY - 1	91.0					%	
4 ANTIBODY SORTED - 2	cd33						
5 CELLS SORTED - 2	150000						
6 PURITY - 2	99.0					%	
7 ANTIBODY SORTED - 3							
8 CELLS SORTED - 3							

- Flow Tech:** Enter the Antibody Sorted, the Cells Sorted and the Purity information. Click the **Save** button. As soon as you save the results, you will get the following message:

Result Entry - Reflex Actions

 **The following reflex actions were taken based on the results entered**

1. Test CHIMERISM SORTED CELLS, DNA, BLOOD was ordered. A specimen was created and received.

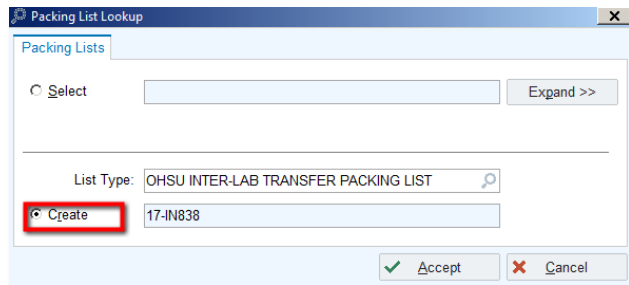
OK

This message means that the reflex KDL test was ordered. Click the **Verify** button and final verify the

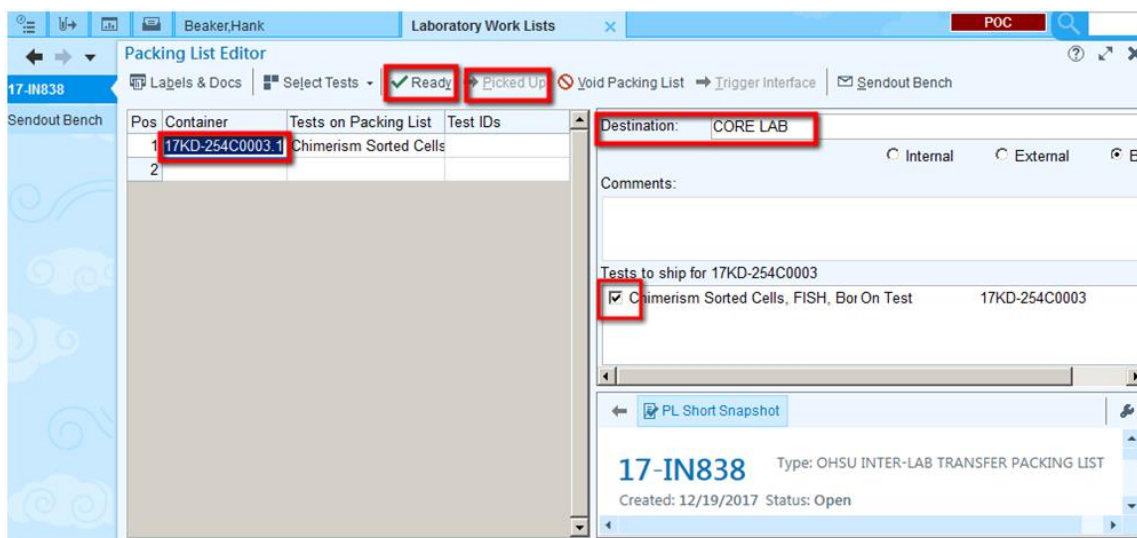
chimerism sort. A label for the reflex should print. Give the printed label and the specimens to the Flow LTA.

## Creating the Packing List to Core Lab:

- Flow LTA:** A packing list needs to be created to send the specimen to Core Lab who will then forward the sample to KDL. Log onto Beaker as the context of “**Lab SIC**”. Click on the **Packing List Editor** from the **Beaker Lab Dashboard-OHSU**. Choose the **Create** option.



- Flow LTA:** In the Destination field, type in “**Core Lab**”. Scan in the printed KDL label to add the specimen to the packing list. Make sure that in the **Tests to ship for <SpecimenID>** field that you check off the test. This should default to “checked off” if you scanned in the .1 aliquot label, but if you scan any other aliquots, this may not be checked off.



- Flow LTA:** Click the **Ready** button, and the **Picked Up** button to close out the packing list. Send the specimens, the remaining KD labels, and the “Send to KDL” paper to Core Lab via tube system.

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