

STANDARD OPERATING PROCEDURES

CATEGORY: Laboratory Work Instructions	SOP Number: Version 1.8 Spleen_Rutgers
TITLE: Separation and Freezing Technique for Spleen Mononuclear Cells from Donor Spleens at ITN Central Cell Processing Core Facility	

AUTHOR:	Signed:
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	Date: 6.8.07

EFFECTIVE DATE: September 2007

REVISION HISTORY:			
Number	Section	Pages	Initials/Dates
001	ALL	Adapted SOP#TCD_SPLEEN V5.0 as used at Peter Heeger's lab for use at RUCDR	RH/6.8.07
002	Materials and Methods	Updated the list of reagents	SK/07.19.07
003	Appendix	Add direction for cleaning the sieve and sieve base	SK/07.19.07
004	Appendix	Add directions on how to prepare 1% vol/vol antibiotics in RPMI 1640	SK/07.19.07

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005		Add step for filtering cell suspension	SK/07.19.07
006	5.2.2	Change centrifugation conditions to 30 min at 2000 rpm	SK/07.19.07
007	5.2.4	Change centrifugation conditions to 7 min at 1500 rpm	SK/07.19.07
008	5.2.7	Change centrifugation conditions to 7 min at 1500 rpm	SK/07.19.07
009	5.5	Change final cell dilution to 20×10^6 SMNCs	SK/07.19.07
010	5.3	Add RBC lysis step to the cell counting procedure	SK/07.19.07
011	Appendix	Add ViCell setting suggested by Beckman Coulter for reference.	SK/07.19.07
012	6.3.3.3	Add ViCell set up values for control	SK / 10.15.07
013	4.2.6	Update reagents	SK / 10.15.07

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1.0	Title
	Separation and Freezing Technique for Spleen Mononuclear Cells from Donor Spleens ITN Central Cell Processing Core Facility

2.0	Purpose
	The purpose of this SOP is to describe the steps involved in the isolation and cryopreservation of spleen mononuclear cells from a donor spleen using the Ficoll method.

3.0	Definitions and Abbreviations	
	SMNC	Spleen Mononuclear Cells
	FCS	Fetal Calf Serum
	PBS	Phosphate Buffer Saline
	DMSO	Dimethyl Sulfoxide

4.0	Equipment and Reagents			
	4.1	Equipment		
		Name	Vendor	Catalog Number
	4.1.1	Sieve base	PGC Scientific	76-0161-00
	4.1.2	Sieve # 80 (8" Tall)	PGC Scientific	75-9002-39
	4.1.3	60 ml Syringe	No specific vendor required	
	4.1.4	Tweezers	No specific vendor required	
	4.1.5	Scalpel	No specific vendor required	
	4.1.6	- 80 Freezer	No specific vendor required	
	4.1.7	Conical Tube, 50 ml, Sterile, Falcon	Falcon 352074	Fisher # 14-432-22
	4.1.8	Centrifuge: Beckman Coulter Allegra 6R - Rotor Gh 3.8 or	No specific vendor required	

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			comparable		
		4.1.9	ViCell XR	Beckman Coulter	
		4.1.10	Liquid nitrogen Tank	Thermo Forma	8030
		4.1.11	Rate Controlled Cell Freezer	In place	
		4.1.12	Serological Pipette, 2 ml	Fisher	13-678-11C
		4.1.13	Pipette Aid	Drummond 4-000-100	Fisher # 13-681-19
		4.1.14	Vials, Cryogenic, Sterile, 1.8 ml	Nalgene/Nunc# 368-632	Fisher # 12-565-171N
		4.1.15	Class II Biological Safety Cabinet	Forma Scientific or equivalent	Model 1286
		4.1.16	250 ml Container (receiver for a cell culture filter)	No specific vendor required	
		4.1.17	Hematology Mixer Model M26125	Barnstead/Thermoline	Fisher # 12-814-2
		4.1.18	Set of Pipettes	No specific vendor required	
		4.1.19	Cell Strainer, 40 μ m	BD Falcon	352340
	4.2	Reagents			
			Name	Vendor	Catalog Number
		4.2.1	RPMI Human Media 1640 (1X), liquide, with Glutamax-I and HEPES buffer	Invitrogen	72400-120
		4.2.3	Dulbecco's PBS sterile, Ca ⁺⁺ , Mg ⁺⁺ Free	Cellgro 21-031-CM	VWR# 45000-436
		4.2.4	Fetal Calf Serum (FCS)	No specific vendor required	
		4.2.5	Ficoll	Axis Shield POC AS	Greiner#1114550
		4.2.6	5,000 units Penicillin/5,000 μ g Streptomycin, 100 ml	Gibco/Invitrogen	15070-063
		4.2.7	7X® Cleaning Solution 1 Gal. (M)	MP Biomedicals	097667093

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	4.2.8	ViaCheck concentration control	Polysciences, Inc.	24629
	4.2.9	ViCell™ Reagent Quad Pack	Beckman	383721
	4.2.10	VersaLyse, 100 tests	Beckman Coulter	IM3648
	4.2.11	Freezing Medium B <ul style="list-style-type: none"> • 20%DMSO • HuAB serum • Tissue culture tested • Heat inactivated Filtered	Provided by RUCDR	
	4.2.12	HuAB serum <ul style="list-style-type: none"> • Tissue culture tested • Heat inactivated • Filtered 	Provided by ITN	
	4.2.13	Trypan Blue	Fisher Biotech	BP1302-10
	4.2.14	Bleach	Clorox	

IMPORTANT:

Keep all reagents at recommended storage conditions. Allow reagents to equilibrate to room temperature before use.

Do not work with chilled reagents so as to maintain the membrane lipid fluidity of the cells.

5.0	Procedures
	IMPORTANT: All work needs to be performed under a biological safety cabinet observing biosafety regulations using sterile technique whenever working with open containers.
	Note: Typically one ~1 in ³ spleen sample will be shipped to the Spleen Core from the transplant trial sites. In the event that more than one sample is received by the Spleen Core, process all of the samples according to the steps below.
	5.1 Preparation of the spleen sample
	5.1.1 Wet the sieve with 20 ml of RPMI with 10% FCS and antibiotics. Place the sieve on top of the receiving container and then place the ~1 in ³ spleen sample into the sieve.
	5.1.2 While the ~1 in ³ spleen sample is in the sieve, slice the spleen piece into thin (~1/8 in) slices with the scalpel. Clean and discard any fatty

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		tissue from the spleen slices.
	5.1.3	Gradually press the spleen slices through the sieve with the back of the 60 ml syringe so that it all goes easily through sieve. At the same time steadily pour small incremental quantities of RPMI with 10% FCS and antibiotics using a total of 100 ml to make sure cells are washed through. You will see that fibroid parts of the spleen will stay on the sieve after this step is done.
	5.1.4	Add another 25 ml of RPMI with 10% FCS and antibiotics to the sieve and press the tissue through one more time.
	5.1.5	Put drained matter into a sterile 250 ml container/receiver. Let it sit for 5 minutes.
	5.1.6	Replace the receiving container under the sieve and pour all of the contents of the 250 ml container back into the sieve and gradually press the unsuspended white matter through the screen.
	5.1.7	Add 60 ml RPMI with 10% FCS and antibiotics to the 250 ml container in order to rinse the remaining matter from it. Then pour this suspension over the remaining white matter in the sieve.
	5.1.8	Gently press the white matter through the sieve.
	5.1.9	Aliquot the strained material from the receiving container into 50 ml tubes at a volume of 10 ml each.
	5.2	SMNC Isolation using Ficoll
	5.2.1	Underlay 10 ml Ficoll into each tube. Since 10 ml was aliquoted into each tube in Step 5.1.9 you will now have a 1:1 proportion of reagent volume to the sample volume
	5.2.2	Centrifuge at RT for 30 minutes at 2000 rpm with the <u>brake off.</u>
	5.2.3	For each prep, using a sterile serological pipette aspirate the interphase layer of cells along with the upper layer of serum and place them into a fresh 50 ml Falcon Tube. It is possible to combine 2-3 samples together at this step.
	5.2.4	For each prep, add PBS up to 45 ml mark on the tube
	5.2.5	Centrifuge this mixture for 7 minutes at 1500 rpm with the <u>brake on - low.</u>
	5.2.6	Discard the supernatant and re-suspend the cells in 10 ml of 10% FCS prepared in RPMI (see Appendix A).
	5.2.7	Centrifuge cells for 5 minutes at 1500 rpm with the <u>brake on - low.</u>

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5.3	When the centrifuge stops, take the tubes out and pour off the liquid. Re-suspend the pellet by tapping the tube.	
5.4	Add 6 ml of Human AB serum to the tube.	
5.5	Filter cell suspension through the 40uM BD Cell strainer.	
5.6	Counting Cells	
	5.6.1	Perform cell counts according to the steps in Section 6.0 of this SOP. Make sure to perform a red blood cell lysis prior to performing the cell count.
5.7	Centrifuge cells for 5 minutes at 1500 rpm with the <u>brake on - low</u> .	
5.8	While cells are in the centrifuge, label the appropriate number of 1.8 ml cryotubes per patient based upon the cell count. Each cryotube will contain approximately 20×10^6 SMNCs.	
5.9	Preparation for the Cryopreservation of Cells	
	5.9.1	When the centrifuge stops, take the tubes out adjust volume of the remaining solution to a cell concentration of 40×10^6 cells/ml of Human AB serum.
	5.9.2	Mix the SMNCs in the tube by gently tapping the side of the tube. Do not use a pipette.
	5.9.3	Add a volume of Freezing Medium B that is equal to that of the Human AB serum from step 5.9.1 to the tube containing the SMNCs. This will bring the cell concentration to 20×10^6 cells/ml.
	5.9.4	Once mixing is complete, quickly aliquot SMNC suspension into the pre-labeled 1.8 ml cryovials. Aliquot only 1.0 ml of cells suspension per cryovial. DO NOT freeze more than 20×10^6 cells per vial. NOTE: Pipette gently to minimize sheer forces!
5.10	Freeze the cryovials using the appropriate Rutgers Cell and DNA Repository protocol for effective cryopreservation. The protocol used is dependent upon the freezer used.	
	5.10.1	The <i>ThermoForma</i> freezer requires the use of Rutgers Cell and DNA Repository protocol "Loading and Starting ThermoForma Control Rate Freezer, Version T.F.2.1"
	5.10.2	The <i>Planer Kryo 10 Series III</i> freezer requires the use of one of two Rutgers Cell and DNA Repository protocols depending on which equipment will be used to store the ITN cells: "Loading and Starting Planer Kryo 10 Series III Control Rate Freezer (MR3 Computer), Version MR3.1.2" or "Loading and Starting Planer Kryo 10 Series III Control Rate Freezer (MRV Computer), Version M.R.V. 1.2"

NOTE ON CELL YIELDS:

Per 1 in³ spleen section the range of cells recovered will typically be between 5×10^7 and 1×10^9 . Many variables will affect the recovery, including the amount of time between procurement and processing and the subject's health and age.

6.0		ViCell™ XR Viability Analyzer Operation and Analysis of SMNCs	
		<p>SAFETY CAUTION</p> <p>If injuries occur due to chemical exposure or a reagent spill occurs, IMMEDIATELY notify a supervisor and REHS at 732-445-2550. If accident occurs after hours, call Rutgers Police at 6-911. Give them the following information:</p> <ul style="list-style-type: none"> *Nature of accident *Your name and location (building and room numbers) *Your telephone number <p>The ViCell™ should be left on at all times, unless it is being serviced by a Beckman Coulter service technician.</p> <p>Do not place fingers near the carousel while the machine is operating to avoid injury. If a tube becomes jammed in the carousel remove it with forceps only and never by hand.</p>	
	6.1	Wear all appropriate personal protective equipment.	
	6.2	ViCell™ XR setup	
		6.2.1	Ensure that the computer attached to the ViCell™ XR is turned on.
		6.2.2	Double click on the "ViCell XR" icon to activate the software.
		6.2.3	Listen for the pump to turn on. The system will then go to idle mode as indicated on the Instrument Status and Control window.
		6.3.4	Observe the "Runs Left" indicator: If the number of runs left on the machine is less than the number of samples-plus-three, the reagent quad pack must be changed prior to use and the liquid waste container along with the used-cup bin must be emptied (see Appendix C for this procedure.)
		6.3.5	Ensure that a control has been run within the last 24 hours by checking the ViCell™ XR Usage Log. If a control has been performed proceed to section 6.4. If it is necessary to run a control, proceed to section 6.3.
	6.3	<i>Running Control</i>	
		6.3.1	Aliquot 0.5-1.0 ml of concentration control into a sample cup and insert into the next available carousel position.
		6.3.2	Double click on the "Control" icon on the left of the screen to display the login screen.

		6.3.3	Ensure that the data in the “Current Lot” display corresponds with the control lot being used. If it does match, proceed to section 6.4. If it does not match, perform the following steps:																																				
		6.3.3.1	Single click on the “New Lot” button.																																				
		6.3.3.2	Enter the following information as below. <table border="1" data-bbox="716 550 1409 1003"> <thead> <tr> <th>Field Name</th> <th>Data to Enter</th> </tr> </thead> <tbody> <tr> <td>Name</td> <td>Concentration Control</td> </tr> <tr> <td>Cell Type</td> <td>Control</td> </tr> <tr> <td>Assay Parameter</td> <td>Total cell /ml</td> </tr> <tr> <td>Acceptance Limits</td> <td>1E6, 4E6, or 8E6</td> </tr> <tr> <td>Comments</td> <td>New Lot</td> </tr> <tr> <td>Current Lot Fields</td> <td>Information Obtained From Bottle</td> </tr> </tbody> </table> <table border="1" data-bbox="734 1047 1365 1614"> <tbody> <tr> <td>Minimum Diameter (µm)</td> <td>2</td> </tr> <tr> <td>Maximum Diameter (µm)</td> <td>50</td> </tr> <tr> <td>Minimum Circularity</td> <td>0.9</td> </tr> <tr> <td>Number of Images</td> <td>50</td> </tr> <tr> <td>Aspirate Cycles</td> <td>2</td> </tr> <tr> <td>Trypan Blue Mixing Cycles</td> <td>3</td> </tr> <tr> <td>Cell Brightness (%)</td> <td>70</td> </tr> <tr> <td>Cell Sharpness</td> <td>75</td> </tr> <tr> <td>Viable Cell Spot Brightness (%)</td> <td>55</td> </tr> <tr> <td>Viable Cell Spot Area (%)</td> <td>1.0</td> </tr> <tr> <td>De-cluster Degree</td> <td>Low</td> </tr> </tbody> </table>	Field Name	Data to Enter	Name	Concentration Control	Cell Type	Control	Assay Parameter	Total cell /ml	Acceptance Limits	1E6, 4E6, or 8E6	Comments	New Lot	Current Lot Fields	Information Obtained From Bottle	Minimum Diameter (µm)	2	Maximum Diameter (µm)	50	Minimum Circularity	0.9	Number of Images	50	Aspirate Cycles	2	Trypan Blue Mixing Cycles	3	Cell Brightness (%)	70	Cell Sharpness	75	Viable Cell Spot Brightness (%)	55	Viable Cell Spot Area (%)	1.0	De-cluster Degree	Low
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		6.3.3.3	<p>Settings for Viability Control</p> <table border="1"> <thead> <tr> <th>Field Name</th> <th>Data to Enter</th> </tr> </thead> <tbody> <tr> <td>Name</td> <td>Viability Control</td> </tr> <tr> <td>Cell Type</td> <td>Control</td> </tr> <tr> <td>Assay Parameter</td> <td>Viability</td> </tr> <tr> <td>Acceptance Limits</td> <td>0%, 50%, or 100%</td> </tr> <tr> <td>Comments</td> <td>New Lot</td> </tr> <tr> <td>Current Lot Fields</td> <td>Information Obtained From Bottle</td> </tr> </tbody> </table> <table border="1"> <tbody> <tr> <td>Minimum Diameter (µm)</td> <td>5</td> </tr> <tr> <td>Maximum Diameter (µm)</td> <td>50</td> </tr> <tr> <td>Minimum Circularity</td> <td>0.9</td> </tr> <tr> <td>Number of Images</td> <td>50</td> </tr> <tr> <td>Aspirate Cycles</td> <td>2</td> </tr> <tr> <td>Trypan Blue Mixing Cycles</td> <td>3</td> </tr> <tr> <td>Cell Brightness (%)</td> <td>85</td> </tr> <tr> <td>Cell Sharpness (%)</td> <td>100</td> </tr> <tr> <td>Viable Cell Spot Brightness (%)</td> <td>60</td> </tr> <tr> <td>Viable Cell Spot Area (%)</td> <td>3.0</td> </tr> <tr> <td>De-cluster Degree</td> <td>Low</td> </tr> </tbody> </table>	Field Name	Data to Enter	Name	Viability Control	Cell Type	Control	Assay Parameter	Viability	Acceptance Limits	0%, 50%, or 100%	Comments	New Lot	Current Lot Fields	Information Obtained From Bottle	Minimum Diameter (µm)	5	Maximum Diameter (µm)	50	Minimum Circularity	0.9	Number of Images	50	Aspirate Cycles	2	Trypan Blue Mixing Cycles	3	Cell Brightness (%)	85	Cell Sharpness (%)	100	Viable Cell Spot Brightness (%)	60	Viable Cell Spot Area (%)	3.0	De-cluster Degree	Low
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		6.3.4	Click "OK."																																				
		6.3.5	Click "Start Query."																																				
		6.3.6	<p>When the query is complete, observe the graph that is displayed on the computer screen and ensure that the data for the current date is within acceptable parameters. If the data is acceptable proceed with section.</p> <p>If it is not within acceptable parameters and troubleshooting methods recommended in the operator manual does not help, contact a Beckman Coulter service technician for service.</p>																																				
		6.3.7	Record the control run in the ViCell™ XR Usage Log.																																				

	6.4	<i>Loading and Running Samples</i>																							
		NOTE: Steps 6.4.1, 6.4.2, and 6.4.3 are performed aseptically under a class II biological safety cabinet.																							
		6.4.1	Add 0.450 ml of VersaLyse to a sample cup.																						
		6.4.2	Remove 50 μ l of SMNC suspension from the flask and add it to the sample cup.																						
		6.4.3	Incubate for 10 minutes at RT and use Vi-Cell cell counter according to manufacturer's instructions.																						
		6.4.4	Place the sample cup(s) into an empty carousel position and record the cell line number(s) in a notebook.																						
		6.4.5	Click the "Cell Type" icon in the tool bar window located on the left of the screen.																						
		6.4.6	In the tool bar to the right, select "Lymphoblastoid."																						
		6.4.6.1	<p>Ensure the following parameters are met in the Cell Type window.</p> <table border="1" data-bbox="743 1037 1373 1606"> <tr> <td>Minimum Diameter (microns)</td> <td>7</td> </tr> <tr> <td>Maximum Diameter (microns)</td> <td>20</td> </tr> <tr> <td>Number of Images</td> <td>50</td> </tr> <tr> <td>Aspirate Cycles</td> <td>2</td> </tr> <tr> <td>Trypan Blue Mixing Cycles</td> <td>3</td> </tr> <tr> <td>Cell Brightness (%)</td> <td>85</td> </tr> <tr> <td>Cell Sharpness</td> <td>100</td> </tr> <tr> <td>Viable Cell Spot Brightness (%)</td> <td>65</td> </tr> <tr> <td>Viable Cell Spot Area (%)</td> <td>3.5</td> </tr> <tr> <td>Minimum Circularity</td> <td>0.1</td> </tr> <tr> <td>De-cluster Degree</td> <td>Low</td> </tr> </table> <p>IMPORTANT: If the parameters are not acceptable, contact a supervisor before continuing.</p>	Minimum Diameter (microns)	7	Maximum Diameter (microns)	20	Number of Images	50	Aspirate Cycles	2	Trypan Blue Mixing Cycles	3	Cell Brightness (%)	85	Cell Sharpness	100	Viable Cell Spot Brightness (%)	65	Viable Cell Spot Area (%)	3.5	Minimum Circularity	0.1	De-cluster Degree	Low
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		6.4.7	Click the "Log In Sample" button to open the log in window.																						

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		6.4.7.1	On the screen, select the carousel position for the sample cup and enter the respective cell line number under "Sample ID." (This step and steps 6.4.7.2, 6.4.7.3, and 6.4.7.4 must be repeated for each sample cup being run.) Note: The following characters may not be entered in the sample ID field: [] = : /
		6.4.7.2	Set dilution factor to 10.0.
		6.4.7.3	Enter any appropriate comments into the comment field.
		6.4.7.4	If the results are to be printed or saved to a Microsoft Excel file, select the appropriate box and/or file.
		6.4.7.5	Once all positions are recorded click "Start Queue." IMPORTANT: Do not leave the ViCell™ XR while it is running. Sample cups can become jammed in the carousel and cause damage to the instrument. Use only forceps to remove jammed cups.
		6.4.8	Once queue has been processed, the results can be recorded from the Previous Run Results window or accessed by opening "file," then opening "run," and opening the desired run.
		6.4.9	After all samples have been analyzed, select "file" and "exit" to terminate the ViCell™ XR program.

APPENDIX A

1X Dulbecco's PBS (DPBS)

HuAB serum: Human AB serum (GeminiBiotech), tested for assay compatibility, heat inactivated and filtered. It can be thawed and refrozen up to 3 times or can be thawed and kept at 4°C for up to one month.

Freezing Medium B: 20% DMSO (Sigma D2650) in heat inactivated HuAB serum, filtered. It can only be thawed one time and must be used the same day.

DMSO Hybri-Max tm Sigma D2650, graded for sensitive cells.
It is not primarily the toxicity of DMSO itself, but the osmotic shock caused by DMSO rapidly that harms the cells most. Also, the viability and functionality of the cells is much better if DMSO is added at room temperature.

200 ml of RPMI with 10% FCS

20 ml FCS

180 mL RPMI Medium 1640

Preparation of the 1% vol/vol antibiotic solution in RPMI1640

Mix 1 ml aliquot of Penicillin / Streptomycin solution (purchased from Gibco/Invitrogen)

With 99 ml of RPMI 1640. This will give you final 1 % volume to volume of antibiotic solution in RPMI 1640.

For future use: Prepare 5 ml aliquots of the antibiotics solution to be used for each 500 ml RPMI bottle. Aliquots of the antibiotics should be stored frozen at -80°C until use

Appendix B

To calculate the correct centrifuge speed for a given RCF, use the following formula:

$$\text{RPM Speed Setting} = \sqrt{((\text{RCF}) \times (100,000) / (1.12) \times (r))}$$

Appendix C**Sieve and sieve base cleaning procedure**

- Put sieve and sieve base in the sink and spread with 10% bleach and leave it for 5 min
- Rinse with tap water
- Clean with detergent (**7X® CLEANING SOLUTION**) as follow:
- Prepare a 1% solution in tap water and gentle scrubbing with a soft brush.
- Drain the used solution and rinse several times with tap water (warm is better).
- Leave them submerged into a recipient with tap water for at least 1 h
- Final-rinse using distilled water.
- Place cleaned items on clean bench-paper and allow to air dry.
- Send for sterilization.

Appendix D**Installing New Quad Pack**

1. Open the reagent compartment door on the right side of the ViCell™ XR and remove the color coded caps from the empty reagent quad pack.
2. Install the color coded caps onto the new quad pack, ensuring that the color coding matches correctly.
3. Place new quad pack into the ViCell™ XR.
4. Remove the liquid waste bottle from the ViCell™ XR.
5. Install an empty 1 liter bottle onto the waste tube cap in the ViCell™ XR. Save the cap from the bottle.
6. Open the old quad pack and dispose remaining reagents into the liquid waste bottle that was just removed from the machine. Recap the full liquid waste bottle with the cap from the new empty bottle and dispose of the bottle as per the *Rutgers University Chemical Hygiene Guide*.
7. Remove and the used cup receptacle from the ViCell™ XR and empty it into the bio-hazardous waste container. Return the cup receptacle to the ViCell™ XR once empty.
8. Record quad pack change and/or prime into the ViCell™ XR Usage Log.

Appendix E**Deviations that must be recorded:**

Code	Description
	Travel
CCP-11	Sample arrived in a wrong collection tube
CCP-12	Sample has been in transit for more then 48 hrs (CPT and Ficoll)
CCP-13	Short blood draw observed in the received tube (CPT and Ficoll)
	Process at the Site
CCP-21	Received tube was not spun before shipping to the Core (CPT only)
CCP-22	No shipment notification given to the core 48 hrs prior to sample arrival
CCP-23	No shipment notification given at all
	Process at the Core
CCP-31	Lost all of the pellet while aspirating supernatant (CPT and Ficoll)
CCP-32	Layers did not separate properly while centrifuging (CPT and Ficoll)
CCP-33	ITN SOP violation noted
	Requisition form related
CCP-41	Patient information had to be corrected/added/changed per ITN confirmation
CCP-42	No collection date and/or time listed on the requisition form
CCP-43	No requisition form faxed in advance
CCP-44	Collection date/time/visit need to be confirmed
CCP-45	Visit not available on the requisition form