The NIH CIT Consortium Chemistry Manufacturing Controls Monitoring Committee:

J. Ansite, A.N. Balamurugan, B. Barbaro, J. Battle, D. Brandhorst, J. Cano, X. Chen, S. Deng, D. Feddersen, A. Friberg, T. Gilmore, J.S. Goldstein, E. Holbrook, A. Khan, T. Kin, J. Lei, E. Linetsky, C. Liu, X. Luo, K. McElvaney, Z. Min, J. Moreno, D. O'Gorman, K.K. Papas, G. Putz, C. Ricordi, G. Szot, T. Templeton, L. Wang, J.J. Wilhelm, J. Willits, T.

Wilson, X. Zhang

The NIH CIT Consortium

Emory University: J. Avila, B. Begley, J. Cano, S. Carpentier, E. Holbrook, J. Hutchinson, C.P. Larsen, J. Moreno, M. Sears, N.A. Turgeon, D. Webster

Massachusetts General Hospital: S. Deng, J. Lei, J.F. Markmann

NIAID: N.D. Bridges, C.W. Czarniecki, J.S. Goldstein, G. Putz, T. Templeton, T. Wilson

NIDDK: T.L. Eggerman

Northwestern University: P. Al-saden, J. Battle, X. Chen, A. Hecyk, H. Kissler, X. Luo, M. Molitch, N. Monson, E. Stuart, A. Wallia, L. Wang, S. Wang, X. Zhang

University of Alberta, Edmonton: D. Bigam, P. Campbell, P. Dinyari, T. Kin, N.

Kneteman, J. Lyon, A. Malcolm, D. O'Gorman, C. Onderka, R. Owen, R. Pawlick, B.

Richer, S. Rosichuk, D. Sarman, A. Schroeder, P.A. Senior, A.M.J. Shapiro, L. Toth, V. Toth, W. Zhai

University of California-San Francisco: K. Johnson, J. McElroy, A.M. Posselt, M.

Ramos, T. Rojas, P.G. Stock, G. Szot

K. Peterson, C. Ricordi, J. Szust, X. Wang

Schwieger, J. Willits, J. Yankey

University of Illinois, Chicago: B. Barbaro, J. Martellotto, J. Oberholzer, M. Qi, Y. Wang University of Iowa (Data Coordinating Center): L. Bayman, K. Chaloner, W. Clarke, J.S. Dillon, C. Diltz, G.C. Doelle, D. Ecklund, D. Feddersen, E. Foster, L. G. Hunsicker, C. Jasperson, D-E Lafontant, K. McElvaney, T. Neill-Hudson, D. Nollen, J. Qidwai, H. Riss, T.

University of Miami: R. Alejandro, A.C. Corrales, R. Faradji, T. Froud, A.A. Garcia, E. Herrada, H. Ichii, L. Inverardi, N. Kenyon, A. Khan, E. Linetsky, J. Montelongo, E. Peixoto,

University of Minnesota: M.H. Abdulla, J. Ansite, A.N. Balamurugan, M.D. Bellin, M. Brandenburg, T. Gilmore, J. V. Harmon, B.J. Hering, R. Kandaswamy, G. Loganathan, K. Mueller, K.K. Papas, J. Pedersen, J.J. Wilhelm, J. Witson

University of Pennsylvania: C. Dalton-Bakes, H. Fu, M. Kamoun, J. Kearns, Y. Li, C. Liu, E. Luning-Prak, Y. Luo, E. Markmann, Z. Min, A. Naji, M. Palanjian, M. Rickels, R. Shlansky-Goldberg, K. Vivek, A.S. Ziaie

University of Wisconsin: L. Fernandez, D.B. Kaufman, L. Zitur **Uppsala University:** D. Brandhorst, A. Friberg, O. Korsgren

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- At Northwestern University, U01Al089316.
- At the University of Alberta, Edmonton: U01Al065191.
- At the University of California, San Francisco, U01DK085531.
- At the University of Illinois, Chicago, 5U01DK070431-10.
- At the University of Iowa, U01DK070431.
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- At the University of Miami: 1UL1TR000460.
- At the University of Minnesota: 5M01-RR000400 and UL1TR000114.
- At the University of Pennsylvania: UL1TR000003.

Address correspondence to: Camillo Ricordi MD, Chairman, CIT Steering Committee, ricordi@miami.edu

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Purified Human Pancreatic Islets Master Production Batch Record, Part 1 University of Illinois, Chicago & University of Miami (Product Codes PHPI-A-01, PHPI-E-01, PHPI-L-01) – Standard Operating Procedure of the NIH Clinical Islet Transplantation Consortium

CellR4 2017; 5 (2): e2284

DAIT, NIAID, NIH

SOP Attachment



Document No. SOP 3101, B02-1 Revision No.

Effective Date 04 September 2009 Supersedes Date 21 July 2009

Page 1 of 46

Document Title:

PURIFIED HUMAN PANCREATIC ISLETS MASTER PRODUCTION BATCH RECORD, PART 1 UNIVERSITY OF ILLINOIS, CHICAGO & UNIVERSITY OF MIAMI (PRODUCT CODES PHPI-A-01, PHPI-E-01, PHPI-L-01)

1.0 MASTER PRODUCTION BATCH RECORD APPROVAL

Signature on file	_ Date:	- 52
Camillo Ricordi, M.D. University of Miami, Miami, Florida	39 Pa	20.
onversity of whatte, mane, rionda		
Signature on file	Date:	
Jose Oberholzer, M.D.		
University of Illinois, Chicago		
Signature on file	Date:	
James P. Markmann, M.D., Ph.D.		
Massachusetts General Hospital, Boston, Massachusetts		
Signature on file	Date:	
Christine W. Czamiecki, Ph.D.		
DAIT, NIAID, NIH, Bethesda, Maryland		

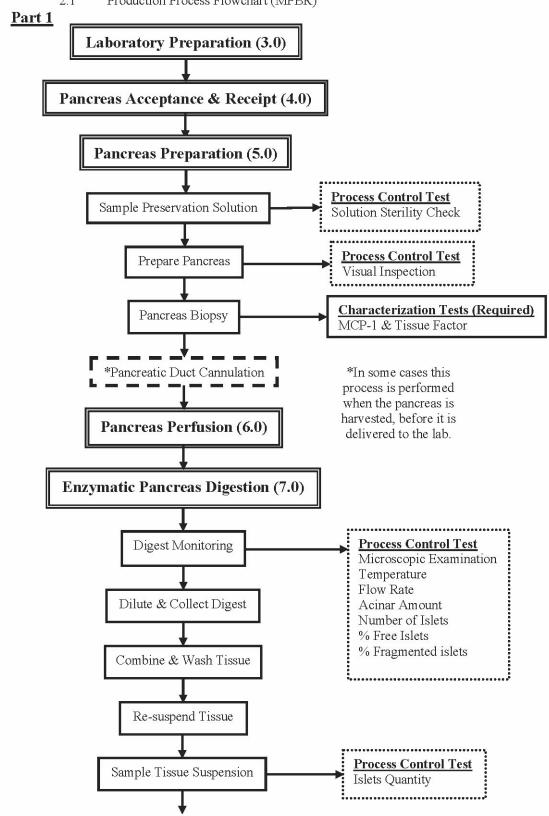
Changes to this Master Production Batch Record must be proposed to the Chief, Regulatory Affairs, DAIT, NIAID, NIH, and approved by all the original signatories, or their successors, before implementation.

Islets Lot Number:	

Document No.	Revision No.	Effective Date	Supersedes Date	Page 2 of 46
SOP 3101, B02-1	04	04 September 2009	21 July 2009	
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)				

2.0 FLOWCHART AND SAMPLING TABLE

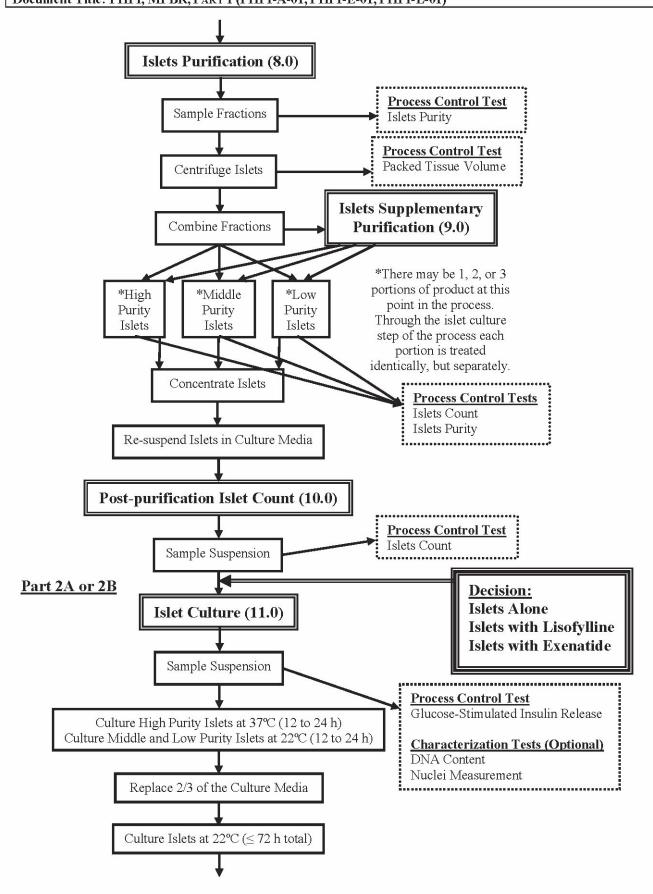
2.1 Production Process Flowchart (MPBR)



Islets Lot Number:

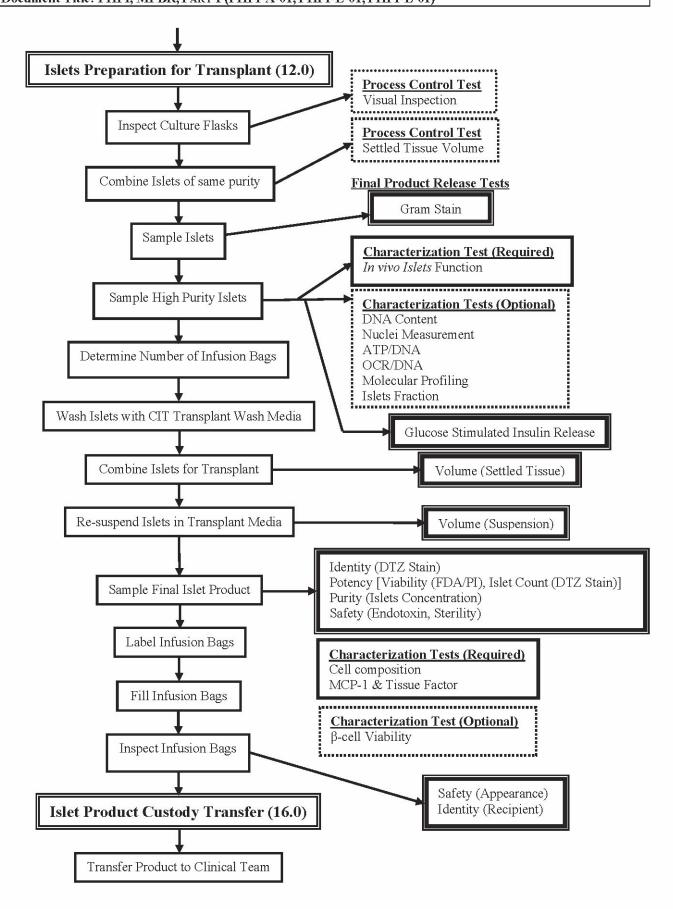
Document No. SOP 3101, B02-1 Revision No. Effective Date 04 September 2009 Supersedes Date 21 July 2009

Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)



Document No. Revision No. Effective Date SOP 3101, B02-1 04 04 September 2009 21 July 2009

Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)



Document No.	Revision No.	Effective Date	Supersedes Date	Page 5 of 46
SOP 3101, B02-1	04	04 September 2009	21 July 2009	
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-I-01)				

2.2 Samples and Tests

MPBR	SAMPLE TYPES & QUANTITIES	
SECTION	PROCESS CONTROL TESTS	TESTS
5.1	Preservation Solution, 3 mL	Sterility
7.1.3	Pancreas Digest, ≤ 1-2 mL periodically	Acinar Amount, # of Islets, % Free Islets, % Fragmented
7.5.1	Diluted Pancreas Digest, 100 μL	Islets Count
8.3.7	Purification Fractions, 0.5 mL/each of 12 fractions & 0.5 mL of W1 fraction, each COBE Run	Islets Purity (%)
8.4.3	Supplementary Purification Islets, 100 μL	Islets Count
9.1.3.6 or 9.2.21	Purification Fractions, 0.5 mL/each of 12 fractions & 0.5 mL of W1 fraction	Islets Purity (%)
10.2	Purified Islets, 2 X 100 μL, High, Middle, Low Purity Levels	Islets Count
12.10	Cultured Islets, All Measured, High, Middle, Low Purity Levels	Settled Tissue Volume
12.13	Cultured Islets, 2 X 100 µL, High, Middle, Low Purity Levels	Post-culture Islets Count
	INTERIM CERTIFICATE OF ANALYSIS	
11.1	Suspension, 400 IEQ, High Purity Islets	Glucose Stimulated Insulin Release
	INTERIM & FINAL	
	CERTIFICATES OF ANALYSIS	
12.11.6	Supernatant above cultured islets, volume according to institution's procedure, High, Middle, Low Purity Levels	Gram Stain
12.18.1	Combined Islets, All Measured, High, Middle, Low Purity Levels	Settled Tissue Volume
12.18.2	Suspension, 2 X 100 μL/Each Final Product T-75 Flask	Islets Identity, Quantity, Concentration
12.18.2	Suspension, 100 IEQ/Each Final Product T-75 Flask	Viability
12.18.2	Supernatant, 1 mL/Each Final Product T-75 Flask	Endotoxin
	FINAL CERTIFICATE OF ANALYSIS	
12.14	Suspension, 400 IEQ, High Purity Islets	Glucose Stimulated Insulin Release
12.18.2	Suspension, 3 mL/Each Final Product T-75 Flask	Sterility, 21 CFR 610.12
	REQUIRED PRODUCT CHARACTERIZATION TESTS	
	FOR INFORMATION ONLY	
5.7	Superficial biopsy of approximately 3 mm X 3 mm X 3 mm	MCP-1 and Tissue Factor
12.14	Suspension, 4,000 IEQ, High Purity Islets	In vivo (Nude Mouse) Islets Function
12.18.2	Suspension, 1,000 IEQ/Each Final Product T-75 Flask	Cell Composition
12.18.2	Suspension, 500 to 1,000 IEQ/Each Final Product T-75 Flask	MCP-1 and Tissue Factor
	OPTIONAL PRODUCT CHARACTERIZATION TESTS	
	FOR INFORMATION ONLY	
11.1	Suspension, 3 X 100 IEQ, High Purity Islets	DNA Content
11.1	Suspension, 3 X 100 IEQ, High Purity Islets	Nuclei Measurement
12.14	Suspension, 3 X 100 IEQ, High Purity Islets	DNA Content
12.14	Suspension, 3 X 100 IEQ, High Purity Islets	Nuclei Measurement
12.14	Suspension, 500 IEQ, High Purity Islets	ATP/DNA Ratio
12.14	Suspension, 5,000 IEQ, High Purity Islets	OCR/DNA
12.14	Suspension, 5,000 IEQ, High Purity Islets	Molecular Profiling
12.14	Suspension, 500 IEQ, High Purity Islets	Islets Fraction
12.18.2	Suspension, 2,000 IEQ/Each Final Product T-75 Flask	β-cell Viability

Document No.	Revision No.	Effective Date	Supersedes Date	Page 6 of 46
SOP 3101, B02-1	04	04 September 2009	21 July 2009	
Document Title: PHPL MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)				

Note: Materials used in this process may transmit infectious agents. Therefore, each person participating in this process must be trained in, and follow, the institution's procedures for handling potentially infectious agents. All waste materials from this process that may have contacted the pancreas or the islets must be discarded as Biohazardous Waste.

Note: It is extremely important to protect the pancreas and the islets from contamination by adventitious microorganisms and pyrogenic agents. Reagents and equipment that may contact the pancreas or islets must be sterile, pyrogen-free, and single-use whenever possible. The institution's procedures for aseptic technique must be followed throughout the execution of this Production Batch Record. All "open" procedure steps must be performed in a clean and disinfected Certified Class II area or Biological Safety Cabinet (BSC).

Note If, at any time during the execution of this Production Batch Record, you observe:

- 1) potential discrepancies in the identification of the pancreas or islets,
- 2) unusual appearance of any materials,
- 3) unusual, or improper performance of any equipment, or
- *inadvertent deviations from the process as defined in this Production Batch Record or the institution's established procedures;*

you must notify the Laboratory Director, or designee, immediately.

The Laboratory Director, or designee, must investigate the observation, and write, sign and date a report giving the details of the observation and its resolution according to the institution's procedures. The occurrence of the event is documented in this Production Batch Record by writing "See Report #X" at the location in the Batch Record where the observation occurred. When allowed by the institution's procedures the report, or a copy, must be filed with this Batch Record. When not allowed, it must be traceable through the unique identification number ("Report #X") written in the Batch Record. The process for reporting a deviation to the CMCMC as defined in DAIT SOP 3200 must also be followed.

3.0 LABORATORY PREPARATION

3.1	lentification of Institution, Personnel, Raw Materials and Purchased Reagents, Sterilized Items
	quipment and Disposable Items

3.1.1	Institution Manufacturing Purified Human Pancreatic Islets Product
	Name of Institution:

3.1.2 Personnel

Attach to this Batch Record a list of the names of all personnel directly involved in the execution of this Batch Record and their signatures and initials, or have them sign and initial the table below.

Document No.	Revision No.	Effective Date	Supersedes Date	Page 7 of 46
SOP 3101, B02-1	04	04 September 2009	21 July 2009	
Document Title: PHPL MPBR PART 1 (PHPL-A-01, PHPL-E-01, PHPL-L-01)				

PRINTED NAME	SIGNATURE	Initials

3.1.3 Raw Materials and Purchased Reagents

Below is a list of the raw materials and purchased reagents used in this procedure, including their catalog numbers and suppliers, where specific Catalog Numbers and Suppliers are required. Record in the table the Catalog Number and Supplier, where not already specified, and the lot number and expiration date of each material used.

	RAW MATERIAL AND PURCHASED REAGENTS	Catalog Number	SUPPLIER	Lot Number	EXPIRATION DATE
1.	CMRL 1066, Supplemented, CIT Modifications				
2.	CMRL 1066 Transplant Media, contains Hepes and without Sodium Bicarbonate				
3.	Hanks' Balanced Salt Solution (HBSS), 1X				
4.	Heparin Sodium Injection USP, Preservative Free		Units/mL		
5.	HEPES Buffer, 1 M				
6.	Gradient Stock Solution				
7.	Phase I Solution				
8.	Cold Storage/Purification Stock Solution				
9.	Albumin Human USP, 25% Solution				
10.	Hydrochloric Acid NF, 1 N				
11.	Insulin-like Growth Factor-1 (IGF-1), 1.0 mg/vial	CM001	Cell Sciences		

Document No. SOP 3101, B02-1	Revision No.	Effective Date 04 September 2009	Supersedes Date 21 July 2009	Page 8 of 46	
301 3101, 102-1	04	04 September 2009	21 July 2009		
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)					

RAW MATERIALS AND PURCHASED REAGENTS (Continued)

RAW MATERIAL AND PURCHASED REAGENTS	CATALOG NUMBER	SUPPLIER	LOT NUMBER	EXPIRATION DATE
12. Insulin Human Injection USP, Recombinant				
13a. Collagenase NB 1 GMP Grade	17452	SERVA/Nordmark		
13b. Neutral Protease NB GMP Grade	30303	SERVA/Nordmark		
14a. Collagenase NB 1 Premium Grade	17455	SERVA/Nordmark		
14b. Neutral Protease NB	30301	SERVA/Nordmark	alle and a state of the state o	
15a. CIzyme Collagenase HA	001-1000	VitaCyte LLC		
15b. CIzyme Thermolysin	002-1000	VitaCyte LLC		
16. OptiPrep	1114542	Nycomed		
17. Trimming Solution				
18. Human Pancreas, Deceased Donor	See Section 4.2 and SOP 3108			
19. PentaStarch, 10% Solution				
20. Povidone Iodine USP, 10%				
21. Pulmozyme (dornase alpha), 2.5 mL/vial, 1 mg/mL	NDC No. 50242-100-40	Genentech		
22. RPMI 1640 with L-Glutamine				
23. Sterile Water for Injection USP				
24. Viaspan (UW Solution)	1000-46-06	Duramed Pharmaceuticals		
25. Biocoll Separating Solution, Density 1.100	L6155	Biochrome AG/ Cedarlane		
26. Calcium Chloride USP (Dihydrate) (CaCl ₂ 2 H ₂ O)				
27. Cefazolin Sodium USP				
28. Lisofylline, 60 mg/mL	Formula # 0109-00	DiaKine Therapeutics		
29. Exenatide, 250 μg/mL	NDC # 66780-210-07	Amylin Pharmaceuticals		
30. Ricordi Infusion Bag	IB-01	Biorep Technologies, Inc.		

Verified	by:		Date:	
,	IB-01	Biorep Technologies, Inc.		
L	66780-210-07	Pharmaceuticals		

Islets Lot Number: _

Document No.	Revision No.	Effective Date	Supersedes Date	Page 9 of 46	
SOP 3101, B02-1	04	04 September 2009	21 July 2009		
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)					

3.1.4 Sterilized Items

Attach a list of all items used in this process that have been sterilized, the sterilizer load
numbers and dates, and verify that the sterilizations were performed within the time
period validated by the institution.

	Verified by:	Date:
3.1.5	Equipment	
	Attach a list of all equipment used in the mannumbers, serial numbers, etc.	ufacturing process, including identification
	Verified by:	Date:
3.1.6	Disposable Items	
	Attach a list of all disposable items used in the number, and the expiration date.	is process, the supplier of each, the lot
	Verified by:	Date:
Biologic	al Safety Cabinet and Laboratory Preparation	
to the in	the laboratory, including the Biological Safety stitution's procedure(s) and record the prepara or copy, with this Batch Record.	
Verified	l by:	Date:

3.3 Dilution Media Preparation

3.2

3.3.1 Equilibrate RPMI 1640 for digest dilution to room temperature prior to use for approximately 1 to 2 hours.

3.3.2 Prepare four 1L containers ahead of time and store at 2°C to 8°C before use:

REQUIRED	USED
1 st Container	
400 mL of RPMI 1640	mL
200 mL of Albumin Human USP, 25% Solution	mL
200 Units of insulin (final concentration: 0.2 Units/mL)	Units
10,000 Units of heparin (final concentration: 10 Units/mL)	Units
2 nd Container	
400 mL of RPMI 1640	mL
200 mL of Albumin Human USP, 25% Solution	mL
200 Units of insulin (final concentration: 0.2 Units/mL)	Units
10,000 Units of heparin (final concentration: 10 Units/mL)	Units

lsl	ets	Lot	N	um	ber:						

Effective Date Supersedes Date Document No. Revision No. Page 10 of 46
 SOP 3101, B02-1
 04
 04 September 2009
 21 Ju

 Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)
 21 July 2009

mL
mL
Units
Units
mL
mL
Units
Units

		Performed by:	Date:
		Verified by:	Date:
	3.3.3	Fill as many additional containers as needer Solution each to provide a final concentration	
		Number of additional containers:	_
		Volume of additional containers:	_mL
		Volume collected in each additional contain	ner: mL
		Volume of Albumin Human USP, 25% Sol	ution in each additional container ml
		Performed by:	Date:
		Verified by:	Date:
4.0	PANCREAS A	ACCEPTANCE AND RECEIPT	
	4.1 Time	of pancreas receipt in the lab:	_ (Record all times using the 24-hour clock)
	Recei	ved by:	Date:

Islets Lot Number:

Document No.	Revision No.	Effective Date	Supersedes Date	Page 11 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009			
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)						

4.2 Pancreas Donor Qualification Record (NA = Not Available)

The	REQUIREMENTS A qualified donor must have "Yes" responses to all of the Inclusion Criteria (A), and "No" responses to all of the Exclusion Criteria (B & C). tainer Label must specify Human Pancreas, and a UNOS or DDD number must be present. Organ Procurement Organization (OPO) must be identified.	Yes	No	NA
The	and "No" responses to all of the Exclusion Criteria (B & C). tainer Label must specify Human Pancreas, and a UNOS or DDD number must be present.	Yes	No	NTA
The	tainer Label must specify Human Pancreas, and a UNOS or DDD number must be present.			NA
The		1		
	organ recurrence organization (or o) must be identified.			\vdash
	nclusion Criteria (The donor or pancreas must meet these criteria.)			
	Pancreas Preservation in (i) UW, (ii) PF/UW, (iii) HTK, or (iv) PF/HTK Solution(s)			
	Maximum 12 hour cold ischemia time			\vdash
	Donor age 15-65 years			
	Cause and circumstances of death acceptable to the transplant team			\vdash
_	Exclusion Criteria (Is there evidence of the following conditions?)		6	
STATE OF THE PARTY	History or biochemical evidence of Diabetes mellitus Type 1 or 2 (Transplant teams may			
5000	consider donor HbA1C > 6.1% in the absence of transfusions in the week prior to death as an			
	indication for exclusion, with discretion for donors who have received transfusions.)			
2.	Pancreas from non-heart-beating cardiac death donors.			
3.	Malignancies, other than resected basal squamous cell carcinoma or intracranial tumor as the cause of death			
4.	Suspected or confirmed sepsis			
5.	Evidence of clinical or active viral Hepatitis [A, B (HBcAg), C]. HBsAb+ is acceptable, if there is a history of vaccination.			
	Acquired Immunodeficiency Syndrome (AIDS)			
-	HIV seropositivity (HIV-I or HIV-II), or HIV status unknown*			
	HTLV-I or HTLV-II*			
	Syphilis (RPR or VDRL positive)*			
_	Active viral encephalitis or encephalitis of unknown origin			
-	TSE or Creutzfeldt-Jacob Disease			
	Suspected Rabies Diagnosis			
	Treated or Active Tuberculosis			
14.	Individuals who have received pit-hGH (pituitary growth hormone)			
_	Any medical condition that, in the opinion of the transplant team, precludes a reasonable			
	possibility of a favorable outcome of the islet transplant procedure			
	Clinical history and/or laboratory testing suggestive of West Nile Virus, Vaccinia, or SARS			
100	Exclusion Criteria – Behavioral Profiles (Is there evidence of the following conditions?)			
	High-risk sexual behavior within 5 years prior to time of death: men who have had sex with			
	men, individuals who have engaged in prostitution, and individuals whose sexual partners have engaged in high-risk sexual behavior			
	Non-medical intravenous, intramuscular, or subcutaneous drug use within the past five years			—
	Persons with hemophilia or related clotting disorders who have received human-derived clotting factor concentrates			
20.	Findings on history or physical examination consistent with an increased risk of HIV exposure			
	Current inmates of correctional systems and individuals who have been incarcerated for more than 72 consecutive hours during the previous 12 months			

^{*}Test results for Exclusion Criteria B. 7, 8, and 9 are required by FDA regulation.

Document No.	Revision No.	Effective Date	Supersedes Date	Page 12 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009			
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)						

	Is donor qualified as pancreas	source?	Yes	No	(Circle One)
	Recorded by:			Date:	
	Review by:			Date:	
4.3	Examine the container in which and labeled with the UNOS of donor records present?				
	Yes	No	(Circle	One)	
	Is the product packaged prope	rly?			
	Yes	No	(Circle	One)	
	Comments:				
	Examined by:			Date:	
4.4	Record the following informa	tion from dono	r records provide	d by the	e OPO:

PANCREAS DONOR INFORMATION (NA = Not Available)

		Acc	EPTAB	LE?
	Observed	Yes	No	NA
UNOS or DDD Number				
Name and Location of OPO				
OPO Unique Identifier (if applicable)				
Donor Consent for Islets Transplant Present				
Donor's Date of Birth				
Donor's Gender				
Donor's ABO				
Donor's Weight				
Donor's Height				
Donor's Body Mass Index				
Extent of Hemodilution (See Flowchart & Worksheet at the end of this document)				
Donor's CMV Status				

Recorded by:	Date:	

ls	lets	Lot.	Num	ber:			

Document No.	Revision No.	Effective Date	Supersedes Date	Page 13 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009			
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)						

5.0 PANCREAS PREPARATION

5.1	In-proc	In-process Samples for Sterility Testing of Preservation Solution						
	Preserv	Preservation Method:						
	a 3 mL label th and fun	Using sterile technique, open the pancreas container in a Class 100 area. Aseptically take at least a 3 mL sample of the preservation solution in which the pancreas was transported. Prepare and label the sample according to the institution's procedure and submit for sterility (21 CFR 610.12) and fungal testing to the appropriate laboratory. Attach a copy of the requisition form to the Production Batch Record.						
	Sample	e Collected by:	Date:					
	Record	the test results, when available, in Sec	tion 17.1.					
*******	******	********************	********	************				
after the po be made ar	ancreas is nd filed wi	pancreas cleaning and cannulation are procured and before it is delivered to t ith this Production Batch Record. **************	the lab. In these	e cases, records of these activities will				
5.2		he pancreas to a cold tray containing Tanove excess tissue.	rimming Soluti	ion plus 1 g/L Cefazolin Sodium USP				
	Process	s Start time:						
	Perfor	med by:	Date:					
5.3	Examir	ne the cleaned pancreas and record obse	ervations in the	table below.				
_	Check	only one line in each category.						
		Clean		None				
	Fat -	Average	Edema	Interstitial Edema				
	rai [Patchy Infiltration	Edema	Slight Overall Swelling				
		Heavily Infiltrated	1	Overly Distended				
	Elvalo	Well Flushed		Very Soft				
	Flush -	Poorly Flushed	1	Soft				
			Texture	Firm (normal)				
				Many Firm Areas (Fibrotic)				
				Rigid Throughout				
		Blood on Capillaries		Intact				
	Blood	Blood in Intra-Parenchymal	Pancreas Condition	Capsular Damage				
		No Blood Present		Parenchymal Damage				

Islets Lot Number: _

Document No.	Revision No.	Effective Date	Supersedes Date	Page 14 of 46
SOP 3101, B02-1	UDI MDDD DADT	04 September 2009 1 (PHPI-A-01, PHPI-E-01,	21 July 2009	
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	Gross pathology ob	served? Yes	No (Circ	cle One)
	Comments:			
	Examined by:		Date:	
	Prepare the CIT Dig preparation with thi	gestion Solution according to s Batch Record.	DAIT SOP 3106, B01, and	file the record of
	Performed by:		Date:	
		<u> </u>		
5.5	Optional Pancreas S	Surface Decontamination		
	Cefazolin Sodium U with 400 mL of plai	pancreas in 250 mL of Hanks JSP, or in 250 mL of 10% Por in HBSS 1X, transfer it to a not re the original pan and instrun- uments.	vidone Iodine USP solution ew container of 400 mL of p	Rinse the pancreas plain HBSS 1X, and
	Pancreas surface de	contamination method:		
	Documented by: _		Date:	
5.6	Pancreas Cannulation	on and Biopsy		
	tail. Cut the pancre 16 to 22 gauge cann	e perfused in a controlled man as to separate the head and tan hula, one at the head and one a ct from the head of the pancre cess.	il, and cannulate the main p at the tail. You may use a s	ancreatic duct with mall cannula as a
	Performed by:		Date:	
	the main duct of the	l biopsy of approximately 3 m donor pancreas for required sample and ship it according n 17.3.	product characterization Mo	CP-1 and tissue factor
	Performed by:		Date:	
5.8	Pancreas Weight			
		rs before each step, weigh the I tissue after perfusion. Reco		

Islets Lot Number:

6.0

Islets Lot Number: _

Document No.	Revision No.	Effective Date	Supersedes Date	Page 15 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009			
Document Title: PHPL MPBR, PART 1 (PHPL-A-01, PHPL-E-01, PHPL-I-01)						

A. Cannulated Pancreas Weight (before Perfusion)	g
B. Weight of Cannulae, Sutures, and Trimmed Tissue	g
C. Trimmed Pancreas Weight (C = A – B)	g
D. Undigested Tissue Weight (Section 7.3)	g
E. Digested Tissue Weight $(E = C - D)$	g

	Recor	ded by: Date: _	
	Verific	ed by: Date: _	
Comn	nents on p	ancreas receipt and preparation:	
Verifi	ed by:	Date: _	
5.9	CIT E1	nzyme Solution Preparation (Cross out lines not used.)	1
	5.9.1	Prepare the CIT Enzyme Solution – SERVA Enzyme B11, and file the record of preparation with this Batc	
	OR		
	5.9.2	Prepare the CIT Enzyme Solution – Vitacyte Enzym B13, and file the record of preparation with this Bate	
	5.9.3	CIT Enzyme Solution (SERVA or VitaCyte Enzyme	s)
		Collagenase Activity actually used:	(Specify Units)
		Neutral Protease Activity actually used:	Units
		Thermolysin Activity actually used:	Fluorescence Units
		CIT Enzyme Solution Volume actually used:	mL
		Verified by:	Date:
PANO	CREAS P	ERFUSION	
6.1	Assem	ble perfusion equipment according to the institution's p	procedure.
	Perfor	med by: Date: _	

Document No.	Revision No.	Effective Date	Supersedes Date	Page 16 of 46
SOP 3101, B02-1	04	04 September 2009	21 July 2009	
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)				

- 6.2 Perfuse the pancreas with the CIT Enzyme Solution.
 - If indicated by the institution's procedures, prime the perfusion circuit by pumping HBSS, 1X, through it. Confirm the absence of leaks or loose connections, and drain the perfusion circuit.
 - Add CIT Enzyme Solution (Section 5.5) at 4°C to 8°C to the chamber and refill the perfusion circuit with it. Remove all air bubbles.
 - Connect the stopcock and perfusion tubing to the cannula and perfuse the pancreas for 4 to 10 minutes at 60 to 80 mm Hg, followed by 4 to 6 minutes (8 minutes maximum in case of poor distension) at 160 to 180 mm Hg at 4°C to 14°C. Note the Desired Pressure in the table below depending on when the pressure is increased.
 - Record the Perfusion Start Time (enzyme solution enters the pancreas) in the table below.
 - Monitor temperature and pressure during pancreas perfusion and record in the table below.
 - Stop perfusion after 10 minutes (12 minutes in the case of poor distension). If perfusion time exceeds 12 minutes, attach to this record a justification for the additional time.

Pancreas Perfusion Pressures and Temperatures

			Start Time:	,	Ø.
Desired Temp. (°C)	Desired Pressure (mm Hg)	Time (min)	Head Observed Pressure (mm Hg)	<u>Tail</u> Observed Pressure (mm Hg)	Observed Temp. (°C)
4-14	60 – 80	2			
4 – 14	60 – 80	4			
4-14		6			
4 – 14		8			
4-14		10			
4 – 14					
4-14					
4 – 14	160 – 180	Finish Perfusion			
Pe	erfusion comp	letion	Finish time:	Finish time:	
Total P	erfusion Time	e (Minutes)			
	Solution rem rfusion (Section			g or mL (Circle One)	
I	Distention Qu (Circle One	•	Excellent Good Partial	Excellent Good Partial	
	nts on pancreatial distention				
Perfusion	n Method:	Αι	ıtomated	Manual (Ci	rcle One)
Data rec	orded by:			Date:	

Continue to clean the pancreas during perfusion. Save all removed non-pancreatic tissue in the container from Section 5.9.

Document No.	Revision No.	Effective Date	Supersedes Date	Page 17 of 46
SOP 3101, B02-1	04	04 September 2009	21 July 2009	ruge 17 or 10
Document Title: PHPL MPBR, PART 1 (PHPL-A-01, PHPL-E-01, PHPL-L-01)				

	Post-perfusion trim fin	ish time:			
	Performed by:		Date: _		
6.3	Trimmed Pancreas We	ight			
		pleted, weigh all removed 5.9. Record this weight i ight.			
	Performed by:		Date: _		
6.4	600 mL Ricordi Diges	digestion equipment according Chamber (Biorep Tecordel No. 600-mDUR-03, v	hnologies, In	c., Model No	
	Performed by:		Date: _		
6.5	Pancreas Preparation f	or Digestion			
	pieces in a Ricordi dig chamber and add CIT 533 µm woven stainles	even to eleven similar size estion chamber. Place 6 to Enzyme Solution up to the ss steel screen on top of th erly to prevent leaking.	o 8 marbles (Se point where	See Section 7 the screen is	7.0) into the digestion to be placed. Place a
	Performed by:		Date: _		
6.6	Pancreas Processing T	Pancreas Processing Times			
	Pancreas Preparation T	out the pancreas processir ime (Process Start Time,			
		Time (Cross Clamp Time record these in the table be	, from donor	o Perfusion S records, to P	Start Time, Section 6.2), erfusion Start Time,
	from Section 6.2) and		, from donor	o Perfusion S records, to P	Start Time, Section 6.2), erfusion Start Time, Time
	A. Cross Clamp	record these in the table be	, from donor	o Perfusion S records, to P	erfusion Start Time,
	A. Cross Clamp (Donor Records) B. Process Start	record these in the table be	, from donor	o Perfusion S records, to P	erfusion Start Time,
	A. Cross Clamp (Donor Records) B. Process Start (Section 5.2) C. Perfusion Start	record these in the table be	, from donor	o Perfusion S records, to P	erfusion Start Time,
	A. Cross Clamp (Donor Records) B. Process Start (Section 5.2)	Date Date D. Pancreas Preparati	on Time	records, to P	erfusion Start Time,
	A. Cross Clamp (Donor Records) B. Process Start (Section 5.2) C. Perfusion Start	Date Date D. Pancreas Preparati (D = C - B) E. Cold Ischemia Time	on Time	records, to P	Time
	A. Cross Clamp (Donor Records) B. Process Start (Section 5.2) C. Perfusion Start (Section 6.2) *Cold Ischemia Time:	Date Date D. Pancreas Preparati (D = C - B)	on Time e* If the Cold Iso	Hours	Timeminutesminutes
	A. Cross Clamp (Donor Records) B. Process Start (Section 5.2) C. Perfusion Start (Section 6.2) *Cold Ischemia Time:	D. Pancreas Preparati (D = C - B) E. Cold Ischemia Time (E = C - A) must be 12 hours or less. site principal investigator	on Time e* If the Cold Iso	Hours	minutes e is more than 12 hours,
	A. Cross Clamp (Donor Records) B. Process Start (Section 5.2) C. Perfusion Start (Section 6.2) *Cold Ischemia Time: immediately notify the	Date Date D. Pancreas Preparati (D = C - B) E. Cold Ischemia Time (E = C - A) must be 12 hours or less. site principal investigator	on Time e* If the Cold Iso	Hours	minutes minutes e is more than 12 hours,

Islets Lot Number: _

Document No.	Revision No.	Effective Date	Supersedes Date	Dogo 19 of 46
SOP 3101, B02-1	04	04 September 2009	21 July 2009	Page 18 of 46
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)				

If the site principal investiga	or is notified, complete the following:
Name of Person notified: _	
Notified by:	
Date & Time Notified:	

7.0 ENZYMATIC PANCREAS DIGESTION

SERVA Enzymes Pancreas Digestion Parameters

CANNULATED PANCREAS WEIGHT (g) (SECTION 5.9)	CHAMBER SIZE (mL)	CIT ENZYME SOLUTION VOLUME (mL)	Marble Number	DIGESTION FLOW RATE	DILUTION FLOW RATE (mL/min)
< 100	600	350		First 5 minutes: 210 – 250	
100 - 125	600	400		mL/min	
126 – 150	600	450	6 – 8		210 - 250
> 150	600, or divide the pancreas into two portions and perform two digestions.	500		After first 5 min, 90 – 130 mL/min	

VitaCyte enzymes use one vial of each enzyme in 350 mL of CIT Enzyme Solution.

7.1	Pancreas	Digaetion
1.1	I alluluas.	DIECSUOI

7.1.1	Add any remaining residual CIT Enzyme Solution to the recirculation flask for
	introduction into the digestion circuit.

Volume of Pulmozyme (1 mg/mL) added: _____ mL

Add 0 to 5 mL of Pulmozyme (2.5 mL/ampule,1 mg/mL) to the Ricordi Digestion Chamber

Performed by:	Date:

7.1.2 Start pumping the solution at a rate of 230 ± 20 mL/min to fill the system. Record this as the Digestion Start Time in the table in Section 7.1.3. Add as much CIT Digestion Solution to the recirculation flask as needed to fill the system and to completely eliminate air from the circuit.

Immediately begin recording the temperature inside the chamber, and the flow rate in the table in Section 7.1.3.

Rock the chamber gently for the first 5 minutes and then decrease the flow rate to 110 ± 20 mL/min. Start shaking the chamber after 5 minutes. It takes approximately 3 - 5 minutes for the chamber to reach a target temperature of 32 to 38°C.

[7 • [•]]	D 4
varitied by:	Date:
Verified by:	Date.

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Document No.	Revision No.	Effective Date	Supersedes Date	Page 19 of 46			
SOP 3101, B02-1	04	04 September 2009	21 July 2009				
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-F-01, PHPI-I-01)							

7.1.3 When tissue is observed in the circulating digest, take a 1-2 mL sample of the digest from the sampling port with a syringe. Place the digest sample in a 35 mm dish and add dithizone (DTZ) stain solution. Observe the digest under a microscope. Repeat this sampling (taking the same sample volume each time) and examination every 1-2 minutes during the digestion. Record the digestion chamber temperature, the flow rate and your observations on the stained sample in the table below. Maintain temperature between 32°C and 38°C, based on digest quality, considering the following factors that help in determining when to stop digestion and start dilution:

Factors	Ranges for Switching from Digestion to Dilution*
Amount of acinar tissue	3 to 6
Number of islets	> 45 islets
% free islets	> 50%
% fragmented (over-digested) islets	< 10%

^{*}See definitions in Note, below.

Note:

Criteria for evaluating the digest and determining the end of digestion

- Estimate the amount of tissue by centering the tissue in the dish, viewing the mass with a microscope at 40X power, and estimating the amount of the visual field covered (6 = tissue covers entire visual field, 3 = tissue covers about 1/2 of the visual field, 0 = no tissue).
- Estimate the number of islets (a rough visual count, 10-20, 30-50, 80-90 islets, etc.).
- Estimate the % free islets (free islets versus the total number of islets, 25%, 50%, 90%, etc.). Free islets have less than 25% of the border attached to acinar tissue.
- Estimate the % fragmented islets (number of fragmented islets versus the total number of islets, 10%, 15%, 50%, etc.). Fragmented islets are those with a ragged border due to damage by overexposure to the enzyme (Over-digested).

Document No.	Revision No.	Effective Date	Supersedes Date	Page 20 of 46			
SOP 3101, B02-1	04	04 September 2009	21 July 2009				
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)							

	Digestion I							
Digestion	n Start Time	e:						
Time (min)	Desired Temp. (°C)	Observed Temp. (° C)	Desired Flow Rate (mL/min)	Observed Flow Rate (mL/min)	Acinar Amount (0 – 6)	# of Islets (Range)	% Free Islets	% Frag- mented Islets
0	X		210 – 250				X	
1			210 – 250					
2			210 - 250					
3			210 – 250					
4			210 - 250					
5	32 – 38		90 – 130					
6	32 – 38		90 – 130					
7	32 – 38		90 – 130					
8	32 – 38		90 – 130					
	≤30		210 – 250					
	≤ 30		210 – 250					
			210 – 250					
	<u>≤</u> 30		210 – 250					
	≤30		210 – 250					
			210 – 250					

Record Desired Temperatures and Desired	Flow Rates in vacant cells	based on Digestion Stop Time.					
Dilution Start Time = Digestion Stop Time	Digestion Time: minutes						
Dilution Stop Time:	_ minutes						
Comments:							
Recorded by: Date:							

Is.	lets .	Lot .	Num	ber: _			

7.2

7.3

Document No.	Revision No.	Effective Date	Supersedes Date	Page 21 of 46				
SOP 3101, B02-1	04	04 September 2009	21 July 2009	1 age 21 01 40				
Document Title: PHPL MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)								

7.1.4 When the decision to stop digestion is made, start dilution and collection of islets. Record the Dilution Start Time (= Digestion Stop Time) at the end of the table in Section 7.1.3 and calculate the Total Digestion Time. Decided by: Verified by: ____ Date: _____ Dilution and Collection of Islets Adjust the flow rate to 230 ± 20 mL/min, and continue shaking the digestion chamber. Add fresh RPMI 1640 at room temperature to the intake container as needed. Adjust the temperature of the chamber to ≤ 30 °C during dilution and collection. Collect the digest into the 1L containers prepared in 3.3.2. Gently swirl each container periodically as it fills. When it reaches a volume of 1L, immediately decant the solution into 250 mL conical tubes for centrifugation at 170 X g and 2°C to 8C° for 3 to 4 minutes,. Periodically take 1 to 2 mL samples of the diluted digest from the sample port with a syringe. Stain with dithizone (DTZ) solution and observe the stained sample under a microscope. Record your observations in the table in Section 7.1.3. When no islets are observed in the stained samples and little tissue remains in the chamber, discontinue the addition of media to the system, collect the media remaining in the system, and stop the circulation pump. Record the Dilution Stop Time at the end of the table in Section 7.1.3, and calculate the Total Dilution Time. Date: ____ Verified by: Remove the undigested pancreas material from the digestion chamber, weigh it, record the weight below, and in the table in Section 5.9. Calculate the weight of digested tissue in the table in Section 5.9. Examine the undigested pancreas material remaining in the digestion chamber, and estimate the percentages of pancreatic tissue and connective tissue (should equal 100%). Record these estimates below. Weight of undigested tissue remaining in chamber: _____ g Estimate of undigested pancreatic tissue: ______% Estimate of connective tissue: _______%

7.4 Tissue Recovery and Washing

Performed by: _____

7.4.1 Prior to the end of digestion prepare CIT Purification Solution and CIT Wash Solution according to DAIT SOP 3106, B02, and B12, respectively. Attach the record of preparation to this Production Batch Record and keep both solutions at 2°C to 8°C until used.

Date:

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Document No.	Revision No.	Effective Date	Supersedes Date	Dog 22 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009	Page 22 of 46		
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)						

- 7.4.2 As tissue is collected during dilution, transfer it to 250 mL conical tubes for the first four liters and centrifuge at 170 X g and 2°C to 8°C for 3 to 4 minutes, to pellet the tissue.
- 7.4.3 Decant all of the supernatant and transfer pellets with a wide mouth 10 mL pipet to a 1 L container containing 900 mL of CIT Wash Solution (keep cold).

NOTE: Be sure the flask is kept level during recombination to avoid tissue aggregation and hypoxic conditions.

- 7.4.4 If residual tissue remains, wash it with 3 to 5 mL of CIT Wash Solution.
- 7.4.5 After dilution is completed and all the tissue has been recombined into the CIT Wash Solution, mix the flask thoroughly by gentle swirling and transfer the contents into as many 250 mL sterile conical tubes as required. Centrifuge each tube at 170 X g and 2°C to 8°C for 3 to 4 minutes.
- 7.4.6 Wash the recombined tissue with CIT Wash Solution until the extracellular debris and DNA strings have been minimized. As the washing progresses, reduce the number of conical tubes to two, then one by combining tissue.

NOTE: If, during collection, DNA stings are observed after centrifugation with loose pellet formation, transfer the suspension portion of those tubes containing the majority of cells into one separate 250 mL conical tube, and keep it lying flat on the bench for 5 minutes after adding up to 200 mL of CIT Wash Solution and 200 μ L (1 μ g/mL) of Pulmozyme. After re-centrifugation, when the DNA strings have disappeared, recombine with other pellets.

Verifie	d by: Date:	
	Total Suspension Volume or Weight: mL or g (Circle One)	
7.4.8	Bring the total re-suspended islets to 200 to 250 g or mL with CIT Purification Solu Ensure that there are no clumps (dissolve if necessary). Record the volume or weigh	
	Total Packed Tissue Volume: mL	
7.4.7	After the washing is complete, visually estimate the total packed tissue volume in the final 250 mL container. Aspirate the supernatant down to the pellet.	Э

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Document No. Revision No.		Effective Date	Supersedes Date	Dogo 22 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009	Page 23 of 46		
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)						

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7.5	Pre-nu	rification	ISIETS	Collini
1.0	IIC pu	11110441011	TOTOGO	Count

- 7.5.1 Re-suspend tissue evenly. Take one 100 µL sample for one pre-purification islets count.
- 7.5.2 Perform pre-purification count according to the institution's procedure and record the data in the table below or attach spreadsheet to Production Batch Record.

Sample volume:	actor:
----------------	--------

Pre-purification Islets Count & Calculations

Pre-purification Islets Count & Calculations							
Islets Diameter (µm)	Count	Factor	IEQ				
50 – 100		0.167					
101 – 150		0.648					
151 – 200		1.685					
201 – 250		3.500					
251 – 300		6.315					
301 – 350		10.352					
> 350		15.833					
% Trapped Islets		Sample Total IEQ					
% Fragmented Islets		Suspension Total IEQ					
Technician's Initials							

Additional records are necessary if magnification calibration factors are used for individual microscopes.

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Verified by:	Date:

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totalogistics in the color and an analysis are between the color and the	

Document No.	Revision No.	Effective Date	Supersedes Date	Page 24 of 46	
SOP 3101, B02-1	04	04 September 2009	21 July 2009		
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-I-01)					

7.5.3 The maximum tissue volume for purification is 25 mL per COBE run. If the tissue volume is $<\!25$ mL, centrifuge the islets suspension and re-suspend the tissue in 100 mL of CIT Purification Solution. If the tissue volume is $>\!25$ mL, using the Packed Tissue Volume from Section 7.4.8, calculate the number of COBE runs required to process $\le\!25$ mL of packed tissue per run. Divide the tissue evenly into separate sterile 250 mL conical tubes and fill each to the 100 mL mark with additional CIT Purification Solution. During purification of the first tube, the additional conical tubes should be kept in the cold room or refrigerator for subsequent COBE runs (keep tube lying flat and mix occasionally to avoid tissue aggregation) until ready to be loaded into the COBE.

Number of conical tubes and COBE runs:	
Volume of tissue distributed into each tube:	mL
Calculated by:	Date:
Verified by:	Date:

7.5.4 When ready to load the first COBE run, add 20 mL of Albumin Human USP, 25% Solution to the tissue and mix well. Continue to Section 8.2.11.

For subsequent COBE runs, centrifuge the conical tube at 170 X g and 2°C to 8°C for 3-4 minutes. Remove the supernatant, add 20 mL of Albumin Human USP, 25% Solution to the tissue and mix well to re-suspend. Bring the tissue suspension to 120 mL in a 250 mL tube or beaker with CIT Purification Solution. Continue to Section 8.2.11.

8.0 ISLETS PURIFICATION

8.1 COBE 2991 Preparation

Set up the COBE according to the Operational Manual and the institution's procedures. The COBE must be refrigerated or placed in a cold room.

- Prepare High (1.10 g/mL) and Low (1.06 g/mL) CIT Purification Density Gradients according to SOP 3106, B10, and file the records of their preparation with this Production Batch Record.
- Label 13 X 250 mL conical tubes with the COBE run number, and "W1" and fraction numbers 1 through 12 (See tables in Section 8.3). Label a 14th 250 mL conical tube with the COBE run number and "Bag."
- Fill tubes 1 through 12 with 225 mL of CMRL 1066, Supplemented, and store at 2°C to 8°C.

Verified by:	Date:
· · · · · · · · · · · · · · · · · · ·	

- 8.2 COBE 2991 Procedure Gradient and Tissue Loading
 - 8.2.1 Assemble the COBE bag onto COBE cell processor according to institution's procedure. Place clamps near the main line on all colored tubing except one line to be used for loading the COBE bag.
 - 8.2.2 Place gradient-maker on magnetic stir plate and aseptically connect one end of size 16 tubing to gradient-maker and the other end to green tubing of the COBE bag.

Islets Lot Number:	

Document No.	Revision No.	Effective Date	Supersedes Date	Page 25 of 46
SOP 3101, B02-1	04	04 September 2009	21 July 2009	
Document Title: Pl	HPI, MPBR, PART	1 (PHPI-A-01, PHPI-E-01,	PHPI-L-01)	

- 8.2.3 Place a sterile stir bar into the left chamber (next to outlet) and turn on the stir plate.
- 8.2.4 Run tubing through pump and set pump to 60 mL/min.
- 8.2.5 Sanitize the exterior of all solution bottles before placing in the hood.
- 8.2.6 Pour 120 mL of the High Density Gradient into the left chamber of the gradient maker.
- 8.2.7 Start to pump High Density Gradient (1.10 g/mL) into COBE bag. Once this gradient reaches the bag, start the COBE at 1800 2000 rpm.
- 8.2.8 Once the entire 120 mL of High Density Gradient (1.10 g/mL) is loaded, remove excess air from the COBE bag by pressing Superout while unclamping the red tubing. Press the Hold button once the Bottom Gradient has reached the T (junction of red/green tube). Re-clamp the red tubing line and press the Stop/Reset button.
- 8.2.9 Wait for the final centrifugation of the digest tissue and then begin loading the continuous density gradient into the COBE bag (Section 7.5.4).
 - Pour 125 mL High Density Gradient (1.10 g/mL) in the left chamber (nearest the outlet) of the gradient maker. Open and close the port between the two chambers just enough to fill the opening.
 - Pour 125 mL Low Density Gradient (1.06 g/mL) in the right chamber of gradient maker (away from outlet)
 - Start the COBE and ensure that the centrifuge speed is between 1800 and 2000 rpm.

Centrifuge Speed: rpm	
Recorded by:	Date:
Open the port between the chambers, set pu	mp to 20 mL/min and load gradient up to
the T of the COBE bag tubing. Stop the put	mp when the gradient has reached the T-

NOTE: Observe the gradient maker to ensure that gradients are mixing during the continuous gradient loading.

connection.

- 8.2.10 Load the continuous gradient by unclamping the green tubing and starting the pump. Load the entire 250 mL of continuous gradient at 20 mL/minute.
- 8.2.11 When all of the gradient has been loaded, stop the pump just as the last portion of the gradient enters the tubing attached to the gradient maker.

NOTE: COBE must remain spinning during the rest of the purification process. If abnormal signs appear from rotating seal (e.g. leak, unusual noise, burnt smell, etc.), replace COBE bag and make new density gradients.

- 8.2.12 Aseptically remove the tubing from gradient maker port and move it to the beaker with tissue. Reverse the pump to purge the air.
- 8.2.13 Load the tissue with the pump at a setting of 20 mL/min. Gently swirl the beaker to keep the tissue well-suspended during the loading.

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	ment No. 3101, B02-1	Revision No. 04	Effective Date 04 September 2009	Supersedes Date 21 July 2009	Page 26 of 46
Досш	ment Title: P	HPI, MPBR, PART	1 (PHPI-A-01, PHPI-E-01,	PHPI-L-01)	

8.2.14 To ensure tissue does not back-up on the gradient (a heavy tissue line observed on the gradient), periodically turn the pump off allowing tissue to enter the gradient and then turn the pump back on again. Repeat as necessary every 1 to 2 minutes.

NOTE: As an alternate, turn the pump off for 30 seconds, followed by loading tissue for 45 seconds.

- 8.2.15 As soon as the tissue is loaded, add 30 mL of additional CIT Purification Solution to the 250 mL beaker to rinse. Load this rinse onto the COBE.
- 8.2.16 After the last portion of the rinse has entered the COBE bag, stop the pump.
- 8.2.17 Vent the system by carefully unclamping the red tubing. Re-clamp the tubing when liquid (capping solution) is approximately one inch above the ceramic seal.

NOTE: Air left in the ceramic rotating seal can cause seal failure which may lead to leaking, seal occlusion and possible system shutdown due to overpressure during Superout.

8.2.18 Clamp the green line and allow the COBE to spin for 3 minutes. Record data on Purification Data Log for each COBE run, below.

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vermed by.	Date.

- 8.3 COBE 2991 Procedure Tissue Collection
 - 8.3.1 During the 3 minute spin disconnect tubing from the pump. Prepare for collection of tissue fractions.
 - 8.3.2 Verify that the Superout Rate is set at 100 mL/min.
 - 8.3.3 After 3 minute spin slowly remove the blue clamp on the green line and quickly press the Superout button.
 - 8.3.4 Collect the first 150 mL of effluent into the conical tube labeled "W" and 12 X 25 mL fractions into the numbered conical tubes each pre-filled with 225 mL CMRL 1066, Supplemented, as described on the Purification Data Log for each respective COBE run.
 - 8.3.5 Once the fractions are collected, stop the COBE and aseptically collect the contents of the COBE bag into a 250 mL conical tube labeled "bag." Discard the COBE bag and tubing.
 - 8.3.6 Dilute the COBE bag contents up to 200 mL with CMRL 1066, Supplemented. Take a 200 μ L sample and place it into 35 mm dish. Stain the sample with dithizone according to the institution's procedure and examine it for the presence of islets. If a significant number of free islets are present keep the diluted COBE bag contents at 2°C to 8°C for further processing as instructed in Section 8.4.1. If there are not a significant number of free islets, discard the COBE bag contents.
 - 8.3.7 To evaluate each COBE fraction quickly, gently but thoroughly mix each fraction from Section 8.3.4, then quickly transfer a 0.5 mL sample to one well of a 12-well microtiter plate and 0.5 mL of the W fraction to a 35 mm dish.
 - 8.3.8 Stain each sample with dithizone according to the institution's procedure and observe for islets. Record Islets Purity (%) and disposition of each fraction on the Purification Data Log for each COBE run.

Document No.	Revision No.	Effective Date	Supersedes Date	Page 27 of 46
SOP 3101, B02-1	04	04 September 2009	21 July 2009	
Document Title: Pl	HPI, MPBR, PART	1 (PHPI-A-01, PHPI-E-01,	PHPI-L-01)	

- 8.3.9 Centrifuge the 250 mL tubes for 3 minutes at 140 X g and 2°C to 8°C. Record Packed Tissue Volumes of each COBE fraction on the Purification Data Log for each respective COBE run. Discard supernatant.
- 8.3.10 Combine the islets fractions by transferring the pellets with 10 mL pipets into four labeled 250 mL conical tubes containing 100 mL of CMRL 1066, Supplemented, to obtain the following purity levels after recombination:
 - High Purity (≥ 70%) (H),
 - Middle Purity (40% to 69%) (M),
 - Low Purity (30% to 39%) (L), and
 - Supplementary Purification Islets (<30%) (S).

Discard fractions (D) that contain little or no tissue. Keep the conical tubes flat on the bench at room temperature until the tissue of all COBE runs has been combined into the respective conical tubes.

NOTE: There will be one 250 mL conical tube for each Purity Level (High, Middle, Low Purity Islets), and one 250 mL conical tube for the Supplementary Purification Islets.

8.3.11 Repeat steps 8.2.1 to 8.3.10 for each COBE purification run. Combine fractions of similar purity into the 250 mL conical tubes prepared in Section 8.3.10.

NOTE: Scoring Guidelines for purified layers in Purification Data Logs:

- Packed Tissue Volume: estimate of the tissue volume in the individual conical tubes after they have centrifuged for 3 minutes at 140 X g and 2°C to 8°C.
- % Purity: estimate relative amount (%) of islets to total tissue.
- H M L S D: This is the disposition for each conical tube as defined in the column header.

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Document No.	Revision No.	Effective Date	Supersedes Date	Page 28 of 46
SOP 3101, B02-1	04	04 September 2009	21 July 2009	
Document Title: P	HPI, MPBR, PART	1 (PHPI-A-01, PHPI-E-01,	PHPI-L-01)	

Repeat this purification process for each of the tubes.

Purification Data Log, COBE Run #1:

Layer	Medium			
Capping Layer	CIT Purification Solution	30 mL		
Tissue Layer	20 mL of Albumin Human CIT Purification Solution 120 g			
Density	Low Density Gradient (1.06 g	mL) 125 g		
Gradients	High Density Gradient (1.10 g	/mL) 125 g		
Bottom	High Density Gradient (1.10 g	/mL) 120 g		
Centrifuge	Start Time Ce	entrifuge Stop Time		

#	CMRL 1066, Supplemented Pre-fill Vol. (mL)	Fraction Volume Collected (mL)	Packed Tissue Volume (mL)	Comments	Islet Purity (%)	Disposition: H: High, M: Middle, L: Low, S: Supplementary, D: Discard (Circle One)
W	0	150 mL				H M L S D
1	225	25				H M L S D
2	225	25				H M L S D
3	225	25				H M L S D
4	225	25				H M L S D
5	225	25				H M L S D
6	225	25				H M L S D
7	225	25				H M L S D
8	225	25				H M L S D
9	225	25				H M L S D
10	225	25				H M L S D
11	225	25				H M L S D
12	225	25				H M L S D
Bag	0	95				S D

Comments on purification:					
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and the same					
Verified by:	Date:				

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Document No.	Revision No.	Effective Date	Supersedes Date	Page 29 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009			
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)						

Layer	Medium			
Capping Layer	CIT Purification Solution 30			
Tissue Layer	mL packed tissue in this COBE Run, plus 20 mL of Albumin Human USP, 25% Solution, and q.s. to 120 g with CIT Purification Solution			
Density	Low Density Gradient (1.06 g/mL)	125 g		
Gradients	High Density Gradient (1.10 g/mL)			
Bottom	High Density Gradient (1.10 g/mL) 120 g			
Centrifuge	Start Time Centrifuge Stop Time			

#	CMRL 1066, Supplemented Pre-fill Vol. (mL)	Fraction Volume Collected (mL)	Packed Tissue Volume (mL)	Comments	Islet Purity (%)	Disposition: H: High, M: Middle, L: Low, S: Supplementary, D: Discard (Circle One)
W	0	150				H M L S D
1	225	25				H M L S D
2	225	25				H M L S D
3	225	25				H M L S D
4	225	25				HMLSD
5	225	25				H M L S D
6	225	25				HMLSD
7	225	25				H M L S D
8	225	25				H M L S D
9	225	25				H M L S D
10	225	25				H M L S D
11	225	25				H M L S D
12	225	25				H M L S D
Bag	0	95				S D

Comments on purification:_			
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Recorded	by:	Date:	
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Document No.	Revision No.	Effective Date	Supersedes Date	Page 30 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009			
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)						

Layer	Medium				
Capping Layer	CIT Purification Solution				
Tissue Layer	mL packed tissue in this COBE Run, plus 20 mL of Albumin Human USP, 25% Solution, and q.s. to 120 g with CIT Purification Solution				
Density	Low Density Gradient (1.06 g/mL)				
Gradients	High Density Gradient (1.10 g/mL)				
Bottom	High Density Gradient (1.10 g/mL)				
Centrifuge	Start Time	Centrifuge Stop Time			

#	CMRL 1066, Supplemented Pre-fill Vol. (mL)	Fraction Volume Collected (mL)	Packed Tissue Volume (mL)	Comments	Islet Purity (%)	Disposition: H: High, M: Middle, L: Low, S: Supplementary, D: Discard (Circle One)
W	0	150				H M L S D
1	225	25				HMLSD
2	225	25				H M L S D
3	225	25				H M L S D
4	225	25				H M L S D
5	225	25				H M L S D
6	225	25				H M L S D
7	225	25				H M L S D
8	225	25				H M L S D
9	225	25				H M L S D
10	225	25				H M L S D
11	225	25				H M L S D
12	225	25				H M L S D
Bag	0	95				S D

Comments on purification:					
Recorded by:	Date:				
Verified by:	Date:				

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Document No. Revision No. SOP 3101, B02-1 04		Effective Date 04 September 2009	Supersedes Date 21 July 2009	Page 31 of 46						
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)										

Layer	Medium	Amount			
Capping Layer	Ding Layer CIT Purification Solution				
Tissue Layer	mL packed tissue in this COBE Run, plus 20 mL of Albumin Huma USP, 25% Solution, and q.s. to 120 g with CIT Purification Solution				
Density	Low Density Gradient (1.06 g/mL)	125 g			
Gradients	High Density Gradient (1.10 g/mL)	125 g			
Bottom	High Density Gradient (1.10 g/mL)	120 g			
Centrifuge	Start Time Centrifuge Stop Time				

#	CMRL 1066, Supplemented Pre-fill Vol. (mL)	Fraction Volume Collected (mL)	Packed Tissue Volume (mL)	Comments	Islet Purity (%)	Disposition: H: High, M: Middle, L: Low, S: Supplementary, D: Discard (Circle One)		
W	0	150				H M L S D		
1	225	25				H M L S D		
2	225	25				H M L S D		
3	225	25				H M L S D		
4	225	25				H M L S D		
5	225	25				H M L S D		
6	225	25				H M L S D		
7	225	25				H M L S D		
8	225	25				H M L S D		
9	225	25				H M L S D		
10	225	25				H M L S D		
11	225	25				H M L S D		
12	225	25				H M L S D		
Bag	0	95				S D		

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Document No.	Revision No.	Effective Date	Supersedes Date	Page 32 of 46						
SOP 3101, B02-1	04	04 September 2009	21 July 2009							
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)										

Layer	Medium	Amount					
Capping Layer	CIT Purification Solution						
Tissue Layer	mL packed tissue in this COBE Run, plus 20 mL of Albumin Human USP, 25% Solution, and q.s. to 120 g with CIT Purification Solution						
Density	Low Density Gradient (1.06 g/s	mL) 125 g					
Gradients	High Density Gradient (1.10 g/	mL) 125 g					
Bottom	High Density Gradient (1.10 g/	mL) 120 g					
Centrifug	Start Time Ce	ntrifuge Stop Time					

#	CMRL 1066, Supplemented Pre-fill Vol. (mL)	Fraction Volume Collected (mL)	Packed Tissue Volume (mL)	Comments	Islet Purity (%)	Disposition: H: High, M: Middle, L: Low, S: Supplementary, D: Discard (Circle One)
W	0	150				H M L S D
1	225	25				H M L S D
2	225	25				H M L S D
3	225	25				H M L S D
4	225	25				HMLSD
5	225	25				H M L S D
6	225	25				H M L S D
7	225	25				H M L S D
8	225	25				H M L S D
9	225	25				H M L S D
10	225	25				H M L S D
11	225	25				HMLSD
12	225	25				H M L S D
Bag	0	95	_			S D

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Document No. Revision No. SOP 3101, B02-1 04		Effective Date 04 September 2009	Supersedes Date 21 July 2009	Page 33 of 46						
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)										

Note: If this purification process did not yield a sufficient number of, and/or sufficiently pure, islets for transplant, and there is a substantial number of impure islets in the remaining tissue, follow one of the procedures in Section 9.0, for Supplementary Purification.

- 8.4 Supplementary Purification Fractions and COBE Bag Contents Processing
 - 8.4.1 If, upon examination of the COBE bag contents, a significant number of islets is present (See Section 8.3.6), centrifuge the 250 mL conical tube containing the diluted COBE bag contents at 140 X gravity and 2°C to 8°C for three minutes, and transfer the packed tissue to the Supplementary Purification Islets 250 mL conical tube.
 - 8.4.2 Bring the volume of the Supplementary Purification Islets 250 mL conical tube to 100 mL with CMRL 1066, Supplemented.
 - 8.4.3 Take a 100 μ L sample for counting. Dilute the Supplementary Purification Islets to approximately 250 mL with CMRL 1066, Supplemented. Lay the tube on its side at 2°C to 8°C while counts are performed.

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8.4.4 Count islets according to the institution's procedure in the Supplementary Purification Islets sample and record counts in the table below and attach spreadsheet. Indicate if the tissue will be re-purified. Supplementary Purification may be indicated if there are a significant number of islets (greater than 50,000 IEQ). If Supplementary Purification is to be performed, proceed to Section 9.0.

Supplementary Purification Islets Counts & Calculations

Supplementary rurinea	1			
Sample Volume				μL
Total Volume	mL			
Dilution Factor				
Diameter, Factor	Counts		IPN (Avg.)	IEQ
50 – 100, 0.167				
101 – 150, 0.648				
151 – 200, 1.685				
201 – 250, 3.500				
251 – 300, 6.315				
301 – 350, 10.352				
> 350, 15.833				
Total				
% Trapped				
Technicians' Initials				

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cument No. P 3101, B02-1	Revi	sion No. 04	Effective Date 04 September 2009	Supersedes Date 21 July 2009	Page 34 of 46	
	PHPI, N		1 (PHPI-A-01, PHPI-E-01		- 1	
1	1		47			
comments on Su	ppieme	ntary Purifica	ation:			
	Recorded by:		_	Date:		
	Verifie	ed by:		Date:		
Decided by:				Date:		
8.5	Tissue	Preparation:	for Re-purification			
	8.5.1	5.1 If the decision in Section 8.4, is to perform a Supplementary Purification of the islets, centrifuge the 250 mL conical tube containing all the supplementary Purification Islets a 140 X gravity and 2°C to 8°C for three minutes. Remove and discard the supernatant.				
	8.5.2	Purification for 30 to 5	Supplementary Purification I n Solution and gently re-sus O minutes while preparation the Supplementary Purificat	pend them. Seal the tube for Supplementary Purific	and place it at 2°C to 8°C	
	Verifie	ed by:		Date:		
.0 ISLETS	SUPP	LEMENTAR	RY PURIFICATION			
purified	by the		y purified by the procedure of plementary Purification Procedure 5 on 9.2.			
Describe	e the sup	oplementary	purification procedure to be	used.		
Approx	ad by			Date:		

Islets Lot Number:		

Site Principal Investigator, or Designee

Document No.	Revision No.	Effective Date	Supersedes Date	Page 35 of 46	
SOP 3101, B02-1	U4	04 September 2009	21 July 2009	5	
Document Title: PHPL MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-I-01)					

9.1 OptiPrep Supplementary Purification Procedure

9.1.1 COBE 2991 Preparation

Set up the COBE according to the Operational Manual and the institution's procedures. The COBE must be refrigerated or placed in a cold room.

- Prepare High (1.10 g/mL) and Low (1.06 g/mL) CIT Purification Density Gradients according to SOP 3106, B10, and file the records of their preparation with this Production Batch Record.
- Label 13 X 250 mL conical tubes with the COBE run number and "W1" and fraction numbers 1 through 12 (See tables in Section 8.3). Label a 14th 250 mL conical tube with the COBE run number and "Bag."
- Fill tubes 1 through 12 with 225 mL of CMRL 1066, Supplemented, and store at 2°C to 8°C.

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- 9.1.2 COBE 2991 Procedure Gradient and Tissue Loading
 - 9.1.2.1 Assemble the COBE bag onto COBE cell processor according to institution's procedure. Place clamps near the main line on all colored tubing except one line to be used for loading the COBE bag.
 - 9.1.2.2 Place gradient-maker on magnetic stir plate and aseptically connect one end of size 16 tubing to gradient-maker and the other end to green tubing of the COBE bag.
 - 9.1.2.3 Place a sterile stir bar into the left chamber (next to outlet) and turn on the stir plate.
 - 9.1.2.4 Run tubing through pump and set pump to 60 mL/min.
 - 9.1.2.5 Sanitize the exterior of all solution bottles before placing in the hood.
 - 9.1.2.6 Pour 120 mL of the High Density Gradient into the left chamber of the gradient maker.
 - 9.1.2.7 Pump the bottom layer into the COBE Bag then stop the pump.
 - 9.1.2.8 Remove excess air from the COBE bag by pressing Superout while unclamping the red tubing. Press the Hold button once the Bottom Gradient has reached the T (junction of red/green tube). Re-clamp the red tubing line and press the Stop/Reset button.
 - 9.1.2.9 Begin loading the continuous density gradient into COBE bag.
 - Pour 125 mL High Density Gradient (1.10 g/mL) in the left chamber (nearest the outlet) of the gradient maker. Open and close the port between the two chambers just enough to fill the opening.
 - Pour 125 mL Low Density Gradient (1.06 g/mL) in the right chamber of gradient maker (away from outlet)
 - Open the port between the chambers, set pump to 20 mL/min and load gradient up to the T of the COBE bag tubing. Stop the pump when the gradient has reached the T-connection.

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Document No.	Revision No.	Effective Date	Supersedes Date	Dogo 36 of 46	
SOP 3101, B02-1	04	04 September 2009	21 July 2009	Page 36 of 46	
Document Title: PHPI MPRR PART (PHPI-A-01 PHPI-E-01 PHPI-I-01)					

Document Ti	itle: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-	E-01, PHPI-L-01)
NOTE:	Observe the gradient maker to ensure th gradient loading.	at gradients are mixing during the continuous
	9.1.2.10 Start the COBE and ensur	re the centrifuge speed is 1800 to 2000 rpm.
	Centrifuge Speed:	_ rpm
	Recorded by:	Date:
		lient by unclamping the green tubing and starting the 50 mL of continuous gradient at 20 mL/minute.
		has been loaded, stop the pump just as the last portion tubing attached to the gradient maker.
NOTE:		rest of the purification process. If abnormal signs sual noise, burnt smell, etc.), replace COBE bag and
	9.1.2.13 Aseptically remove the two with tissue. Reverse the page 1.2.13 Reverse the page 2.2.13 Rever	ubing from gradient maker port and move to the beaker pump to purge the air.
		Purification Islets (Section 8.5) with the pump at a ently swirl the beaker to keep the tissue well suspended
	on the gradient), periodic	back-up on the gradient (a heavy tissue line observed eally turn the pump off allowing tissue to enter the pump back on again. Repeat as necessary every
		oaded, add 30 mL of additional CIT Purification eaker to rinse. Load this rinse onto the COBE.
	9.1.2.17 After the last portion of the	he rinse has entered the COBE bag, stop the pump.
		ully unclamping the red tubing. Re-clamp the tubing ution) is approximately one inch above the ceramic
NOTE:	Air left in the ceramic rotating seal can cocclusion and possible system shutdown	cause seal failure which may lead to leaking, seal due to overpressure during Superout.
	1 0	allow the COBE to spin for 3 minutes. Record data on purification COBE run, below.

Islets Lot Number:	
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Date: _____

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Document No.	Revision No.	Effective Date	Supersedes Date	Page 37 of 46	
SOP 3101, B02-1	04	04 September 2009	21 July 2009		
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-I-01)					

- 9.1.3 COBE 2991 Procedure Tissue Collection
 - 9.1.3.1 During the 3 minute spin disconnect tubing from the pump. Prepare for collection of tissue fractions.
 - 9.1.3.2 Verify that the Superout Rate is set at 100 mL/min.
 - 9.1.3.3 After 3 minute spin, slowly remove the blue clamp on the green line and quickly press the Superout button.
 - 9.1.3.4 Collect the first 150 mL of effluent into the conical tube labeled "W1" (waste) and 12 X 25 mL fractions into the numbered conical tubes each pre-filled with 225 mL CMRL 1066, Supplemented, as described on the Purification Data Log for each respective COBE run.
 - 9.1.3.5 Once the fractions are collected, stop the COBE and discard the COBE bag and tubing.
 - 9.1.3.6 To evaluate each COBE fraction quickly, gently but thoroughly mix each fraction from step 9.1.3.4, then quickly transfer a 0.5 mL sample to one well of a 12-well microtiter plate and 0.5 mL of the W fraction to 35 mm dish.
 - 9.1.3.7 Stain each sample with dithizone according to the institution's procedure and observe for islets. Record observations on the Re-purification Data Log.
 - 9.1.3.8 Centrifuge the 250 mL tubes for 3 minutes at 140 x g and 2℃ to 8℃. Record Packed Tissue Volumes of each COBE fraction on the Re-purification Data Log. Discard the supernatant.

NOTE: Scoring Guidelines for purified layers in Purification Data Logs:

- Packed Tissue Volume: estimate of the tissue volume in the individual conical tubes after they have centrifuged for 3 minutes at 140 x g and 2°C to 8°C.
- % Purity: estimate relative amount (%) of islets to total tissue.
- H M L D: This is the disposition for each conical tube as defined in the column header.

Document No.	Revision No.	Effective Date	Supersedes Date	Page 38 of 46	
SOP 3101, B02-1	04	04 September 2009	21 July 2009		
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)					

OptiPrep Supplementary Purification Data Log

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Layer	Medium		
Capping Layer	CIT Cold Storage Solution		
Tissue Layer	mL packed tissue in this COBE Run, plus 20 mL of Albumin Human USP, 25% Solution, and q.s. to 120 g with CIT Cold Storage Solution		
Density	Low Density Gradient (1.06 g/mL)		
Gradients	High Density Gradient (1.10 g/mL) 125 g		
Bottom	High Density Gradient (1.10 g/mL)		
Centrifuge Start Time		Centrifuge Stop Time	

#	CMRL 1066, Supplemented Pre-fill Vol. (mL)	Fraction Volume Collected (mL)	Packed Tissue Volume (mL)	Comments	Islet Purity (%)	Disposition: H: High, M: Middle, L: Low, D: Discard (Circle One)
W	0	150				HMLD
1	225	25				HMLD
2	225	25				HMLD
3	225	25				HMLD
4	225	25				HMLD
5	225	25				HMLD
6	225	25				HMLD
7	225	25				HMLD
8	225	25				HMLD
9	225	25				HMLD
10	225	25				HMLD
11	225	25				HMLD
12	225	25				HMLD
Bag	0	95				D

Comments on supplementary purification:	
Recorded by:	Date:
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Verified by:	Date:

Document No.	Revision No.	Effective Date	Supersedes Date	Page 39 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009			
Document Title: PHPL MPBR PART 1 (PHPL-A-01, PHPL-E-01, PHPL-L-01)						

9.1.4 Combine fractions with purity of 30% or greater with the complimentary fractions from Section 8.3.10, and record the disposition of each fraction in the OptiPrep Supplementary Purification Data Log in Section 9.1.3.8. Discard fractions < 30% pure.

NOTE:		At this point there will be one 250 mL conical tube for each Purity Level (High, Middle, Low Purity Islets).						
	Perfor	med by:	y:					
	Verifi	ed by:		Date: _				
9.2	Biocol	Biocoll Supplementary Purification Procedure						
	9.2.1		the tissue by adding 150 om Section 8.3.10.	mL of UW Solution	n to the Supplementary Purification			
Note:		e can be			edure, up to 45mL of packed tissue ry important not to overload the			
Note:		olume of d tissue.	UW Solution for each r	un remains constan	t, regardless of the volume of the			
		Volum	e of UW Solution used fo	or each COBE run: _	mL			
		Total F	acked Tissue Volume: _	mL				
		Numbe	r of COBE runs:					
		Packed	Tissue Volume prepared	l for each COBE run	:mL			
		Perfor	med by:		Date:			
	9.2.2		e the tissue in UW solution tator (or mix the tissue in		ice or in the cold room, using the g every 5 minutes).			
		Perfor	med by:		Date:			
	9.2.3		tion of Biocoll Heavy (49 19% UW Solution mixed)		Solution mixed) and Light (30%			
		9.2.3.1	-		erile bottle. Label this Bottle with and time of preparation, and initials			
		9.2.3.2			er sterile bottle. Label this Bottle date and time of preparation, and			
		9.2.3.3	Pipette 63.7 mL of 1.10 "Heavy Gradient" and		nt Solution into the bottle labeled to mix properly.			

Islets Lot Number: _

Document No.	Revision No.	Effective Date	Supersedes Date	Page 40 of 46			
SOP 3101, B02-1	04	04 September 2009	21 July 2009				
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-I-01)							

9.2.3.4 Pipette 42.0 mL of 1.10 g/mL Ficoll Gradient Solution into the bottle labeled "**Light Gradient**" and quickly swirl bottle to mix.

	Performed by: Date:			
9.2.4	Set the COBE at 1500 rpm and Superout at 0. Press Start to start the COBE.			
9.2.5	Add $110~\mathrm{mL}$ of $1.10~\mathrm{g/mL}$ Biocoll Gradient Solution to the first (front) beaker and start the peristaltic pump on the maximum setting.			
9.2.6	After all the Biocoll Gradient Solution is loaded onto the COBE, press Superout, turn off the pump, unclamp the pump head, and turn Superout to 100 .			
9.2.7	When the Biocoll Gradient Solution reaches the beaker, quickly re-clamp the pump head. Stop the COBE and turn Superout back to 0. Change the COBE speed to $3{,}000\mathrm{rpm}$. All air should now be out of the system.			
9.2.8	Add 130 mL of Heavy Gradient to the front beaker. Unclamp the line between the beakers briefly and re-clamp to get all air out.			
9.2.9	Add 140 mL of Light Gradient to the second (rear) beaker.			
9.2.10	Turn the pump speed down to $20~\text{mL/min}$ on the peristaltic pump and turn magnetic stirrer on the lowest setting. Start the COBE. Start pump. Unclamp the line between the beakers.			
9.2.11	When nearly all the Biocoll is loaded onto the COBE, tilt the magnetic stirrer forward to ensure all Biocoll is loaded. Before the last bit of Ficoll is loaded, stop the stirrer and begin to slowly add the suspended islets to the front beaker.			
9.2.12	When all tissue has been added, rinse the conical which contained the suspended islets with 50 mL of HBSS, 1X, and add this volume to the front beaker.			
9.2.13	When everything has been loaded onto the COBE, clamp the tubing above the bag, press Super-Out (set at 0), turn off the pump and unclamp the pump head.			
9.2.14	SLOWLY, unclamp the clamp above the COBE bag and start the timer.			
	Performed by: Date:			
9.2.15	Centrifuge for 5 minutes.			
9.2.16	Prepare collection rod and line for fraction collection.			
9.2.17	Prepare 12×250 mL conical tubes. Label them #1 through #12. Leave Tube #1 empty, and pre-fill Tubes #2 through #12 with 220 mL each of CMRL 1066, Supplemented.			
	Performed by: Date:			
9.2.18	After 5 minutes, slowly adjust the Superout up to 100 and begin collecting tissue into the conical tubes.			
9.2.19	Collect 150 mL of effluent in Tube #1. Collect 30 mL of effluent in Tubes #2 through #12, to a total volume of 250 mL in each tube.			

Islets Lot Number: _

Document No.	Revision No.	Effective Date	Supersedes Date	Page 41 of 46			
SOP 3101, B02-1	04	04 September 2009	21 July 2009	1 age 41 01 40			
Document Title: Pl	Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)						

9.2.20 When all effluent has been collected, press Stop on the COBE.

Performed by:	Date:

- 9.2.21 To evaluate each COBE fraction quickly, gently but thoroughly mix each fraction from Section 9.2.19, then quickly transfer a 0.5 mL sample to one well of a 12-well microtiter plate.
- 9.2.22 Stain each sample with dithizone according to the institution's procedure and observe for islets. Record observations on the Biocoll Supplementary Purification Data Log for each COBE run, below.
- 9.2.23 Centrifuge the 250 mL tubes for 3 minutes at 140 X g and 2°C to 8°C. Record the Packed Tissue Volumes of each COBE fraction on the Biocoll Supplementary Purification Data Log for each respective COBE run. Discard supernatant.

NOTE: Scoring Guidelines for purified layers in Purification Data Logs:

- Packed Tissue Volume: estimate of the tissue volume in the individual conical tubes after they have centrifuged for 3 minutes at 140 x g and 2°C to 8°C.
- % Purity: estimate relative amount (%) of islets to total tissue.
- H M L D: This is the disposition for each conical tube as defined in the column header.

Islets Lot Number: _		
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Document No.	Revision No.	Effective Date	Supersedes Date	Page 42 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009	1 age 42 01 40		
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-I,-01)						

Biocoll Supplementary Purification Data Log, COBE Run #1:

#	CMRL 1066, Supplemented Pre-fill Vol. (mL)	Fraction Volume Collected (mL)	Packed Tissue Volume (mL)	Comments	Islet Purity (%)	Disposition: H: High, M: Middle, L: Low, D: Discard (Circle One)
1	0	150				HMLD
2	220	30				H M L D
3	220	30				HMLD
4	220	30				HMLD
5	220	30				HMLD
6	220	30				HMLD
7	220	30				H M L D
8	220	30				H M L D
9	220	30				H M L D
10	220	30				HMLD
11	220	30				HMLD
12	220	30				HMLD
	Centrifuge Star	t Time		Centrifuge Stop	Time	

Comments on purification:	
•	
Recorded by:	Date:
Verified by:	Date:

9.2.24 Repeat all steps for each COBE run.

Comments on purification:_

Document No.	Revision No.	Effective Date	Supersedes Date	Page 43 of 46		
SOP 3101, B02-1	04	04 September 2009	21 July 2009			
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)						

Biocoll Supplementary Purification Data Log. COBE Run #2:

#	CMRL 1066, Supplemented Pre-fill Vol. (mL)	Fraction Volume Collected (mL)	Packed Tissue Volume (mL)	Comments	Islet Purity (%)	Disposition: H: High, M: Middle, L: Low, D: Discard (Circle One)
1	0	150				HMLD
2	220	30				HMLD
3	220	30				HMLD
4	220	30				HMLD
5	220	30				HMLD
6	220	30				HMLD
7	220	30				HMLD
8	220	30				HMLD
9	220	30				HMLD
10	220	30				HMLD
11	220	30				HMLD
12	220	30				HMLD
	Centrifuge Star	t Time		Centrifuge Stop	Time	

Record	led by:	Date:	
Verifie	d by:	Date:	
9.2.25		tubes by adding 100 mL of CMRL 1066, Purity," "Middle Purity," and "Low Purit	
9.2.26	labeled 250 mL conical tubes a Middle Purity (69% to 40%), a pure. Keep the conical tubes f	y transferring the pellets with 10 mL pipel according to their purity level: High Purit and Low Purity (39% to 30%). Discard fr lat on the bench at room temperature unti d into the respective conical tubes.	y (≥ 70%), ractions < 30%
	Performed by:	Date:	
	Verified by:	Date:	

Islets Lot Number:

Document No.	Revision No.	Effective Date	Supersedes Date	Dogo 44 of 46	
SOP 3101, B02-1	04	04 September 2009	21 July 2009	Page 44 of 46	
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)					

10.0 POST-PURIFICATION ISLETS COUNT

10.1	After all islets are combined into the three Purity Culture Media prepared according to DAIT SOP to settle for 3 to 5 minutes. After the tissue has s final tissue in 100 to 200 mL of CIT Culture Medisolation date and Purity Level identification.	3106, B04. Allow the tissue in the conical tubes ettled, remove the supernatant and re-suspend the
	Verified by:	Date:
10.2	Gently mix each Purity Level and take two 100 μ Count. Enter the count data in the table below or Number (IPN) and Total IEQ. The contents of the Culture, Section 11.	attach spreadsheet, and calculate the Total Islet
	Sampled by:	Date:

Post-purification Islets Counts

		High Purity				Middle Purity		Low Purity				
Sample Volume				μL				μL				μL
Total Volume				mL				mL				mL
Dilution Factor												
Diameter, Factor	Cou	unts	Avg.	IEQ	Со	unts	Avg.	IEQ	Со	unts	Avg.	IEQ
50 – 100, 0.167												
101 – 150, 0.648												
151 – 200, 1.685												
201 – 250, 3.500												
251 – 300, 6.315												
301 – 350, 10.352												
> 350, 15.833												
Total												
% Trapped						6						
% Fragmented												
% Purity												
Islet Quality Grade*												
Technicians' Initials												

Islets Lot Number:		

Document No.	Revision No.	Effective Date	Supersedes Date	Dogo 45 of 46	
SOP 3101, B02-1	04	04 September 2009	21 July 2009	Page 45 of 46	
Document Title: PHPI, MPBR, PART 1 (PHPI-A-01, PHPI-E-01, PHPI-L-01)					

Post-purification Islets Calculations

	High Purity	Middle Purity	Low Purity	Total
Post-purification IPN				
Post Purification IEQ				
Pre-purification IEQ (Section 7.5.2)				
IEQ Recovery (%) (from Pre-purification IEQ)				
Total IEQ/g of trimmed pancreas (Section 5.8)				
Comments				

^{*}See Note, below, for Islets Quality Grade guidelines

Calculated by:	Date:
Verified by:	Date:

Note: Islets Quality Grade

Grade the quality of the islets based on these parameters and criteria:

Parameter	0 Points	1 Point	2 Points
Shape (3D)	flat/planar	in between	spherical
Border (2D)	irregular	in between	well-rounded
Integrity	fragmented	in between	solid/compact
Single Cells	many	a few	almost none
Diameter	all < 100 μm	a few > 200 μm	$> 10\% > 200 \mu m$

Add up the points for each sample to obtain the following grades:

- \circ 9 to 10 points = A
- \circ 7 to 8 points = B
- \circ 4 to 6 points = C
- \circ 2 to 3 points = D
- \circ 0 to 1 point = F

Document No.	Revision No.	Effective Date	Supersedes Date	Page 46 of 46	
SOP 3101, B02-1	04	04 September 2009	21 July 2009		
Document Title: PHPL MPBR, PART 1 (PHPL-A-01, PHPL-E-01, PHPL-L-01)					

Will the CIT Culture Media, the CIT Transplant Wash Media, and the CIT Transplant Media for this batch contain no drug, Lisofylline, or Exenatide?

Circle one of the following:

Islets Alone (Clinical Protocols 03 – 07)

Islets with Lisofylline (Clinical Protocol 02 only)

Islets with Exenatide (Clinical Protocol 02 only)

Approved by:	Date:
*********	********

If "Islets Alone" is circled above, continue recording the manufacturing process in Part 2A (SOP 3101, B02-2A).

If "Islets with Lisofylline" or "Islets with Exenatide" is circled above, continue recording the manufacturing process in Part 2B (SOP 3101, B02-2B).

NOTE: Part 1 of the Production Batch Record must be combined with either Part 2A or Part 2B for review and approval.