

VICell Product Information and Testing - Amended

Product Information

Product Name	WA27			
Lot Number	WB0138			
Parent Material	WB0130			
Depositor	WiCell			
Banked by	WiCell			
Thaw Recommendation	Thaw 1 vial into 1 well of a 6 well plate. WiCell recommends thawing using ROCK Inhibitor for best results.			
Culture Platform	Feeder Independent			
	Medium: E8 plus PVA			
	Matrix: Recombinant Human Vitronectin			
Protocol	WiCell Feeder Independent E8 Medium Protocol			
Passage Number	p13			
	These cells were cultured for 12 passages prior to freeze. WiCell adds +1 to the passage number at freeze so that the number on the vial best represents the overall passage number of the cells at thaw.			
Date Vialed	20-April-2012			
Vial Label	WB0138 WA27 P13 20APR12 JJ			
Biosafety and Use Information	Appropriate biosafety precautions should be followed when working with these cells. The end user is responsible for ensuring that the cells are handled and stored in an appropriate manner. WiCell is not responsible for damages or injuries that may result from the use of these cells. Cells distributed by WiCell are intended for research purposes only and are not intended for use in humans.			

Lot Specific Testing Performed by WiCell The following tests were performed on this specific lot.

Test Description	Test Provider	Test Method	Test Specification	Result
Post-Thaw Viable Cell Recovery	WiCell	SOP-CH-305	 ≥ 15 Undifferentiated Colonies, ≤ 30% Differentiation 	Pass
Identity by STR	UW Molecular Diagnostics Laboratory	PowerPlex 16 HS System by Promega	Consistent with known profile	Pass
Sterility - Direct transfer method	Apptec	30744	Negative	Pass
Mycoplasma	Bionique	M250	No contamination detected	Pass
Karyotype by G-banding	WiCell	SOP-CH-003	Normal karyotype	Pass



Cell Product Information and Testing - Amended

General Cell Line Testing Performed by WiCell The following tests were performed on the cell line. The tests do not apply to any particular lot.

Test Description	Test Provider	Test Method
Differentiation Potential by Teratoma	WiCell	SOP-CH-213 SOP-CH-214
HLA	UW Histocompatibility Laboratory	High resolution sequencing method with Celera reagents on the ABI 3100 instrument
ABO	New York Blood Center	For ABO: Olsson ML, Chester MA. A rapid and simple ABO genotype screening method using a novel B/O2 versus A/O2 discriminating nucleotide substitution at the ABO locus. Vox Sang 1995; 69(3):242-7. For RHD: Singleton BK, Green CA, Avent ND, Martin PG, Smart E, Daka A, Narter-Olaga EG, Hawthorne LM, Daniels G. The presence of an RHD pseudogene containing a 37 base pair duplication and a nonsense mutation in Africans with the Rh D-negative blood group phenotype. Blood 2000; 95(1): 12-8.
Growth Curve (Doubling Time)	WiCell	Varies by culture platform
Flow Cytometry for ESC Marker Expression	WiCell	SOP-CH-024
Comprehensive Human Virus Panel	Charles River	ID 91/0

Amendment(s):

Reason for Amendment	
CoA updated to include copyright information.	
CoA amended to include additional product information and removal of foonotes.	
Amended STR test method and HLA test provider and test method.	
Original CoA.	

Date of Lot Release	Quality Assurance Approval		
16-August-2012	1/3/2014 X AMC Quality Assurance Signed by:		



University of Wisconsin Hospital and Clinics

Short Tandem Repeat Analysis*

Sample Report: 10459-STR

Label on Tube: 10459-STR

Sample Date: 05/11/12 Lab Received 05/11/12

Requestor: WiCell Research Institute Test Date: 05/16/12

File Name: 120517

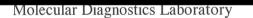
Report Date: 05/21/12

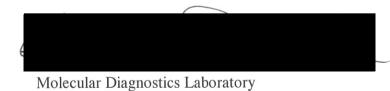
Sample Name: (label on tube) 10459-STR

Description: WI Cell Research Institute provided genomic DNA 262 ug/mL 260/280=1.97

Locus	Repeat #	STR Genotype
D168539	5,8-15	12,13
D7S820	6-14	10,10
D13S317	7-15	11,11
D5S818	7-15	11,12
CSF1PO	6-15	11,13
TPOX	6-13	8,10
Amelogenin	NA	X,X
TH01	5-11	6,8
vWA	11, 13-21	16,18

Comments: Based on the 10459-STR DNA submitted by WI Cell dated and received on 05/11/12, this sample (Label on Tube Cap only: 10459-STR) exactly matches the STR profile of the human stem cell line WA27 comprising 14 allelic polymorphisms across the 8 STR loci analyzed. No STR polymorphisms other than those corresponding to the human WA27 stem cell line were detected and the concentration of DNA required to achieve an acceptable STR genotype (signal/ noise) was equivalent to that required for the standard procedure (~1 ng/amplification reaction) from human genomic DNA. This result suggests that the 10459-STR DNA sample submitted corresponds to the WA27 stem cell line and was not contaminated with any other human stem cells or a significant amount of mouse feeder layer cells. Sensitivity limits for detection of STR polymorphisms unique to either this or other human stem cell lines is ~5%.





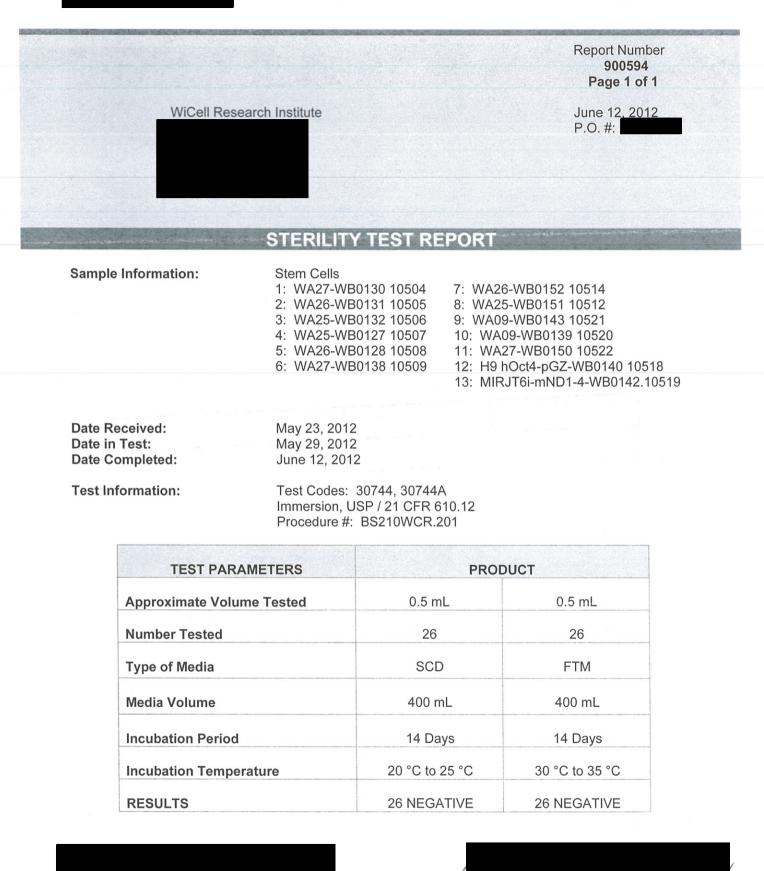
* Testing to assess engraftment following bone marrow transplantation was accomplished by analysis of human genetic polymorphisms at STR loci. This methodology has not yet been approved by the FDA and is for investigational use only.

QA Reviewer

(

This report is confidential. No part may be used for advertising or public announcement without written permission. Results apply only to the sample(s) tested.





Testing conducted in accordance with current Good Manufacturing Practices.

Technical Reviewer

Date

Date



BIONIQUE[®] TESTING LABORATORIES, INC.

MYCOPLASMA TESTING SERVICES

APPENDIX

Document ID #:	DCF9002F
Title:	QUALITY ASSURANCE REPORT - GMP
Effective Date:	11/2/11
Edition #:	03

QUALITY ASSURANCE REPORT - G M P

Test Performed	PROCEDURAL REFERENCE		Test Performed		PROCEDI	JRAL REF	ERENCE	
M-250 M-300 M-350	SOP's 3008, 3011, 3013 SOP's 3008, 3014 SOP's 3008, 3014, 3015		☐ M-700 ☐ M-800			008, 3009 008, 301		
Bionique Sample ID	#(s)_69816			s	*:	x	1 KB	
		N						

This testing procedure was performed in compliance with the FDA's Current Good Manufacturing Practice (cGMP) standards (to the extent that the regulations pertain to the procedures performed) as specified in the Code of Federal Regulations, Title 21 Parts 210 and 211 [21 CFR 210 & 211]. All related records derived from the test procedures have been reviewed by the Quality Assurance Department. The individual's signature below verifies that the methods and procedures referenced above have been followed and that the Final Report accurately reflects the raw data generated during the course of the procedures. All records, including raw data and final reports are archived on site for a minimum of seven years.

The specified test's procedures determine the intervals at which samples are inspected. The medium used for testing must pass quality control mycoplasmal growth promotion testing and sterility testing. Traceability of all of the components used is assured and supporting documentation can be supplied upon request.

Quality Assurance Review Date:	6/6	12	 1 a - 1 2
Reviewed By QA	Assistant:		
$x_{\mu} = x = 0$	18 72		

NOTE:

- 1. Prior to receipt at Bionique[®] Testing Laboratories, Inc., the stability of the test article is the responsibility of the company submitting the sample. Bionique Testing Laboratories Inc. will assume responsibility for sample stability following receipt and prior to being placed on test.
- 2. This test is for the detection of microbiological growth and does not require statistical validation.

BIONIQUE[®] TESTING LABORATORIES, INC.

ADD	FN	DIX
UI I	L/IN.	

Document ID #:	DCF9002F
Title:	QUALITY ASSURANCE REPORT - GMP
Effective Date:	11/2/11
Edition #:	03

REFERENCES

Regulatory:

- 1. Department of Health and Human Services, Food and Drug Administration (USA) [FDA]. Code of Federal Regulations [CFR], Title 21 CFR Part 210, Current Good Manufacturing Practice in Manufacturing, Processing, Packing, or Holding of Drugs; General. FDA. Office of the Federal Register, National Archives and Records Department.
- 2. Department of Health and Human Services, Food and Drug Administration (USA) [FDA]. Code of Federal Regulations [CFR], Title 21 CFR Part 211, Current Good Manufacturing Practice for Finished Pharmaceuticals. FDA. Office of the Federal Register, National Archives and Records Department.
- 3. Department of Health and Human Services, Food and Drug Administration (USA) [FDA]. Points to Consider in the Characterization of Cell Lines Used to Produce Biologicals, Director, Center for Biologics Evaluation and Research, FDA. May, 1993. Docket No. 84N-0154.
- 4. Department of Health and Human Services, Food and Drug Administration (USA) [FDA]. Code of Federal Regulations [CFR], Title 21 CFR Part 610.30, General Biological Products Standards; Subpart D, Test for Mycoplasma. FDA. Office of the Federal Register, National Archives and Records Department.

General:

- 1. Barile MF, Kern J. Isolation of Mycoplasma arginini from commercial bovine sera and its implication in contaminated cell cultures. Proceedings of the Society for Experimental Biology and Medicine, Volume 138, Number 2, November 1971.
- Chen, T.R. In situ detection of mycoplasma contamination in cell cultures by fluorescent Hoechst 33258 stain. Experimental Cell Research, 104: 255-262, 1977.
- Carolyn K. Lincoln and Daniel J. Lundin. Mycoplasma Detection and Control. U. S. Fed. for Culture Collections Newsletter, Vol. 20, Number 4, 1990.
- Fetal Bovine Serum; Proposed Guideline. National Committee For Clinical Laboratory Standards (NCCLS), Vol. 10, Number 6, 1990. (NCCLS publication M25-P).
- 5. McGarrity GJ, Sarama J, Vanaman V. Cell Culture Techniques. ASM News, Vol. 51, No. 4, 1985.
- 6. Tully JG, Razin S. Methods in Mycoplasmology, Volumes I and II. Academic Press, N.Y., 1983.
- 7. Barile MF, Razin S, Tully JG, Whitcomb RF. The Mycoplasmas, Volumes 1-4. Academic Press, N.Y., 1979.
- 8. <u>http://www.bionique.com/</u> Safe Cells Insights



BIONTOUE TESTING LABORATORIES TNC

MYCOPLASMA TESTING SERVICES

APPENDIX IV

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Document#:	DCF3013D
Edition#:	10
	07/15/2003
Effective Date:	0771572003
Title:	M-250 FINAL REPORT SHEET

M-250 FINAL REPORT

Direct Specimen Culture Procedure 3008, 3011, 3013

TO: WiCell QA WiCell Research Institute

BTL SAMPLE ID#:	69816	. ×	P.Ó.#:	DATE REC'D:	05/08/2012
TECH (CONTROL AD				2	

TEST/CONTROL ARTICLE:

WA27-WB0138 #10459

NA LOT#:

DIRECT CULTURE SET-UP (DAY 0)	DATE	: 05/09/2012	
INDICATOR CELL LINE (VERO)	SEE DNA FLUOROCH	ROME RECORD SHEET	
			DATE
THIOGLYCOLLATE BROTH	DAY 7 +	Θ	05/16/2012
	DAY 28 +	-	06/06/2012
BROTH-FORTIFIED COMMERCIAL			
0.5 mL SAMPLE	DAY 7 +	Θ	05/16/2012
6.0 mL BROTH	DAY 28 +	Θ	06/06/2012
BROTH-MODIFIED HAYFLICK			
0.5 mL SAMPLE	DAY 7 +	Θ	05/16/2012
6.0 mL BROTH	DAY 28 +	Θ	06/06/2012
BROTH-HEART INFUSION			
0.5 mL SAMPLE	DAY 7 +	Θ	05/16/2012
6.0 mL BROTH	DAY 28 +	Θ	06/06/2012
(See Reverse)			

Page 1 of 2

APPENDIX IV

	DGD20125	<u></u>				
Document#:	DCF3013E	J				
Edition#:	10					
Effective Date:	07/15/20	03				
Title:	M-250 FI	NAL REPORT	SHEET			
SAMPLE ID#: 698	16		AERO	DBIC	MICROAEROPHILIC	DATE
AGAR PLATES-FORTIF: COMMERCIAL	IED	DAY 7 DAY 14 DAY 21	+ + +	000	+ (05/16/2012 05/23/2012 05/30/2012
AGAR PLATES-MODIFIN HAYFLICK	ED	DAY 7 DAY 14 DAY 21	+ + +	8	$\begin{array}{c} + \\ + \\ + \\ + \end{array}$	05/16/2012 05/23/2012 05/30/2012
AGAR PLATES-HEART INFUSION		DAY 7 DAY 14 DAY 21	+ + +		+ () + () + ()	05/16/2012 05/23/2012 05/30/2012
BROTH SUBCULTURES	(DAY 7)		DATE	: 05	5/16/2012	
AGAR PLATES-FORTIF: COMMERCIAL	- The American Contraction	DAY 7 DAY 14 DAY 21	+ + +		+ (-) + (-) + (-)	05/23/2012 05/30/2012 06/06/2012
AGAR PLATES-MODIFIN HAYFLICK	ED	DAY 7 DAY 14 DAY 21	+ + +		$\begin{array}{c} + & - \\ + & - \\ + & - \\ + & - \end{array}$	05/23/2012 05/30/2012 06/06/2012
AGAR PLATES-HEART INFUSION		DAY 7 DAY 14 DAY 21	+ + +		+ () + () + ()	05/23/2012 05/30/2012 06/06/2012

No detectable mycoplasmal contamination RESULTS:

6/6/12 Date

ADDITIONAL COMMENTS:

M-250 Procedural Summary: The objective of this test is to ascertain whether or not detectable mycoplasmas are present in an in vitro cell culture sample, be it a primary culture, hybridoma, master seed stock or cell line. This procedure combines an indirect DNA staining approach to detect non-cultivable mycoplasmas with a direct culture methodology utilizing three different mycoplasmal media formulations. The indirect approach involves the inoculation of the sample into a mycoplasma-free VERO (ATCC) indicator cell line and performing a DNA fluorochrome assay after 72-120 hours of incubation. The direct culture aspect of the test utilizes three different mycoplasmal media forluditing both broth and agar formulations. The sample is inoculated into each of the 3 broth formulations and also onto duplicate plates (0.1 mL/plate) for each of the 3 agar formulations. Subculture from broth to fresh agar plates is carried out after 7 days incubation. Agar plates are incubated aerobically and microaerophillically in order to detect any colony forming units morphologically indicative of mycoplasma contamination. Issuance of the final microaerophillically in order to detect any colony forming units morphologically indicative of mycoplasma contamination. Issuance of the final microaerophillically in order to detect any colony forming units morphologically indicative of mycoplasma contamination. Issuance of the final microaerophillically in order to detect any colony forming units morphologically end concurrently with the test sample(s) as detailed in the referenced SOPs and that all test conditions have been found to meet the required acceptance criteria for a valid test, including the appropriate results for the positive and negative controls.



MYCOPLASMA TESTING SERVICES

5/10/1

BIONIQUE® TESTING LABORATORIES, INC.

	Document ID #: Title: Effective Date: Edition #:	DCF3008. DNA FLU 3/24/10 07	A orochrome A	SSAY RESULTS		en e			
				DROCHROME A rocedures 3008, 30					
	Sample ID # 69	816	<u>M-250</u>	Date Rec'd:	<u>05/08</u>	8/2012	P.O. #		
	Indicator Cells Inoc	culated:	Date/Initials:	5/10/12	/	nuk			
	Fixation:		Date/Initials:	5/14/12	- 1	J.			
	Staining:		Date/Initials:	5/14/12	_ /	13			
	TEST/CONTROL		·	······································			_		
gir D 5/10/12	WA27-WB0	138 1 3 0 #10459	_						
	LOT# <u>NA</u>								
	<u>WiCell QA</u> WiCell Resea	arah Institu	140	×					
	wiceli Kesea	aren mstitt							
		New York And And And							
	DNA FLUORO	CHROMI	E ASSAY RES	SULTS:					
	NEGAT	IVE:		vith staining limi		the nuclea	ar region,	which indic	ates no
	POSITI	VE:		t amount of extr l contamination		ar staining	g which st	rongly sugg	ests

INCONCLUSIVE:

A significant amount of extranuclear staining consistent with low - level mycoplasmal contamination or nuclear degeneration.

A significant amount of extranuclear staining consistent with bacterial, fungal or other microbial contaminant or viral CPE. Morphology not consistent for mycoplasmal contamination.

COMMENTS: 3 IL_Results Read by: П Date:

_____Date of Review: <u>5/11//12____</u>Reviewed=by: Set



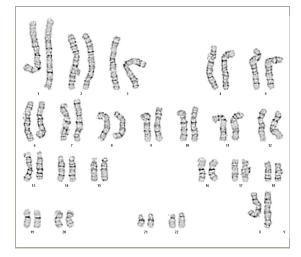
This report is updated to include the results of counting an additional twenty cells in this analysis, as requested by

Report Date: May 25, 2012 Updated Report Date: June 26, 2012 Cell Line: WA27-WB0138 10459

Passage #: 15

Date of Sample: 5/7/2012 Date Completed: 5/25/2012

Results: 46,XX



Specimen: hESC on Vitronectin Cell Line Gender: Female **Reason for Testing:** lot release testing Investigator: , Core

S02-22 Cell: Slide: 2-R1(19)KARYOTYPE Slide Type: Karyotyping

of Cells Counted: 20; an additional 20 cells were counted for this update # of Cells Karyotyped: 4 # of Cells Analyzed: 8 Band Level: 450-550

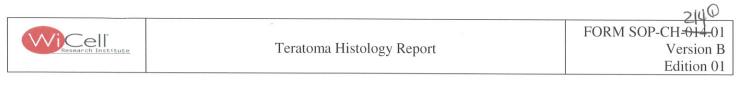
Interpretation:

No abnormalities were detected at the stated band level of resolution. Updated Interpretation: Chromosomes from an additional twenty cells were counted; there were no abnormalities found.

Completed by Completed by CG(ASC Reviewed and interpreted by	CP), on 5/21/2012 , PhD, FACMG, on 5/25/2012
A signed copy of this report is available upon request.	,,
Date:	Sent To:
Sent By:	QC Review By:

Limitations: This assay allows for microscopic visualization of numerical and structural chromosome abnormalities. The size of structural abnormality that can be detected is >3-10Mb, dependent upon the G-band resolution obtained from this specimen. For the purposes of this report, band level is defined as the number of G-bands per haploid genome. It is documented here as "band level", i.e., the range of bands determined from the four karyograms in this assay. Detection of heterogeneity of clonal cell populations in this specimen (i.e., mosaicism) is limited by the number of metaphase cells examined, documented here as "# of cells counted".

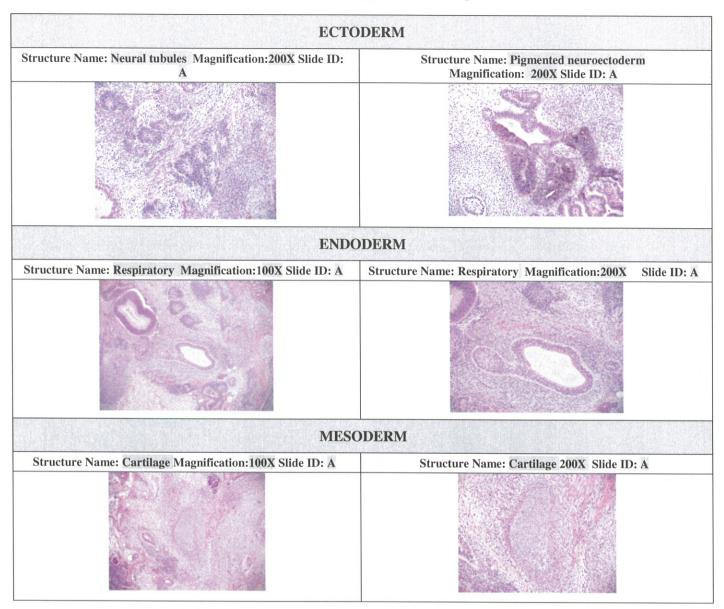
This assay was conducted solely for listed investigator/institution. The results may not be relied upon by any other party without the prior written consent of the Director of the WiCell Cytogenetics Laboratory. The results of this assay are for research use only. If the results of this assay are to be used for any other purpose, contact the Director of the WiCell Cytogenetics Laboratory.



Cell Line: WA27

Cell Lot Number: NA

Sample Number: 10447-A



Comments: Structures identified include Ectoderm (2), Mesoderm (1) and Endoderm (1)

Sample(s) were assessed for the presence of differentiation into cell types characteristic of the three embryonic germ layers, which, if present in the sample(s) examined, are represented in the photographs above. The individual's signature below verifies that this report accurately reflects the pathology observed.

Pathologist (By/Date): 6/27/2012

WHealth

University of Wisconsin Hospital and Clinics

Name: MRN: DOB: HLA#:	WICELL, 10406_HLA OS000184 WICELL	Hospital: Physician: , Category:	
	Bone Marrow (Case Histocompatibility Summary 301417-DT	

HLA Typing Results

Patient	Relation	Hap <u>A*</u>	<u>B*</u>	<u>C*</u>	DRB1*	<u>DRB3*</u>	DRB4* DRB5*	<u>DQB1*</u>	DPB1*	Tested Date Collect Date
WICELL, 10406_HLA		02:01	41:02	05:01	03:01					03/12/12
OS000184 / WICELL	Patient	66:01	44:02:01G	17:01:01G	04:01					03/01/12

HLA typings performed by sequencing, SSO, SSP or a combination. For low-resolution testing, results are reported by Serologic Equivalents. A "+" in the HLA allele designation indicates that the typing was performed by low/mid-resolution molecular method and that additional alleles are possible. Only the most frequent allele is listed.

HLA DNA-Based Typing

Name HLA / MR# Received WICELL, 10406_HLA OS000184 / WICELL	<u>Method</u> <u>Test Date</u> SEQ	<u>A*</u> 02:01 66:01	<u>B*</u>	<u>C*</u>	DRB1*	DRB3*	DRB4*	<u>DRB5*</u>	DQB1*
03/01/2012	03/20/2012		e database.	IMGT 3.7.0 2	012-01-12				
				101 0.7.0 2	012-01-12				
	SEQ		41:02 44:02:0	1G					
03/01/2012	03/20/2012								
		The report	ted allele gr	IMGT 3.7.0 2 oup B*44:02:0 te of exons 2		e following all B*44:19N	eles, which sł	nare identical s	sequences in the
				05:01					
	SEQ			17:01:0	1G				
03/01/2012	03/20/2012		a databasa:	IMGT 3.7.0 2	012 01 12				
		The report antigen re The follow	ted allele gr cognition si ving allele co	te of exons 2 combination(s)		:01 C*17:02 alleles are liste	C*17:03		sequences in the v committee as
					03:01				
00/01/0010	SEQ				04:01				
03/01/2012	03/20/2012			IMGT 3.7.0 2 are allele DRB	012-01-12 31*03:68N, first	identified in A	ugust 31, 201	1.	

Comments

WHealth

University of Wisconsin Hospital and Clinics

Name:	WICELL, 10406_HLA	
MRN:	OS000184	
DOB:		
HLA#:	WICELL	

Hospital: Physician: , Category:

Bone Marrow Case Histocompatibility Summary 301417-DT

This test was developed and its performance characteristics determined by this laboratory. It has not been cleared or approved by the U.S. Food and Drug Administration. The FDA has determined that such clearance or approval is not necessary. This test is used for clinical purposes. It should not be regarded as investigational or for research. This laboratory is certified under the Clinical Laboratory Improvement Amendments of 1988 (CLIA-88) as qualified to perform high complexity clinical laboratory testing.

Electronically signed by Kathleen J. Meuer, MT(ASCP),	03/25/2012 12:18
Director or Delegate, HLA Laboratory	Date/Time

Histocompatibility Laboratory, Room D4/231, 600 Highland Ave., Madison, WI 53792-2472 Teresa Darcy, MD, Medical Director :: Thomas M. Ellis, PhD, D(ABHI) Laboratory Director Lab: 608.263.8815 (option 3); Fax: 608.263.9610 ASHI: 01-4-WI-03-2, CLIA: 52DO661997

Printed Date: 03/25/2012 UWHC 301417-DT

Bone Marrow Case Summary Report w/Test Results



Laboratory of Immunohematology and Genomics 45-01 Vernon Blvd., Long Island City, N.Y. 11101 718-752-4771 • Fax 718-752-4747

March 20, 2012

WiCell Research Institute Attn: Quality Assurance

SAMPLE: DNA WA27 #10406 (MA#168-12)

Date Received: 03/08/12 Sample Date: 03/01/12

HISTORY: DNA from cell line.

TESTING REQUESTED: Genotype for *ABO* and common *RH*

TESTING PERFORMED: *ABO:* Polymerase chain reaction-restriction fragment length polymorphism (PCR-RFLP) testing for nucleotide (nt) positions 261 (O¹), 467 (A²), 703 (B), and 1096 (B and O²). *RH:* Multiplex PCR-RFLP for *RHD* and *RHCE*C/c.* PCR-RFLP for RHCE Exon 5 (676C>G for E/e).

DNA MOLECULAR RESULTS: *ABO:* PCR-RFLP testing indicates the presence of a nt261 deleted G, characteristic of O^1 alleles, and the presence of nt703A and nt1096A, characteristic of B alleles. *RH: RHD* exons 4 and 7 are present. Negative for the inactivating *RHD* pseudogene. *RH*Cc* and *RH*ee*.

GenotypePredicted PhenotypeWA27 #10406: ABO^*BO^1 ; RH^*D , RH^*Cc , RH^*ee Group B; RhD+, C+E-c+e+

Manager, Genomics

Director of Immunohematology and Genomics

These *in vitro* diagnostic tests were developed and their performance characteristics established in the Molecular Analysis Laboratory. The tests have not been submitted to the Food and Drug Administration (FDA) for clearance or approval and; therefore, are not FDA-licensed tests. The Molecular Analysis Laboratory is certified under the Clinical Laboratory Improvement Amendment (CLIA) of 1988 as qualified to perform high complexity clinical testing. The New York Blood Center has been approved by the New York State Department of Health to perform these tests under its current Clinical Laboratory Permit.

These results are intended to predict a blood group antigen profile in a patient or donor, and are not intended for clinical diagnosis or as the sole means for patient management decisions. There are situations where testing DNA of a person may not reflect the red cell phenotype and not all performance characteristics have been determined. Nucleotide changes that inactivate gene expression or rare new variant alleles may not be identified in these assays. In addition, test results obtained from DNA isolated from leucocytes and other hematopoietic cells may differ from DNA isolated from other tissues in persons with a history of transplantation.

▲ NewYork BloodCenter

Immunohematology

Telephone: 718-752-4771

Genomics Telephone: 718-752-4637

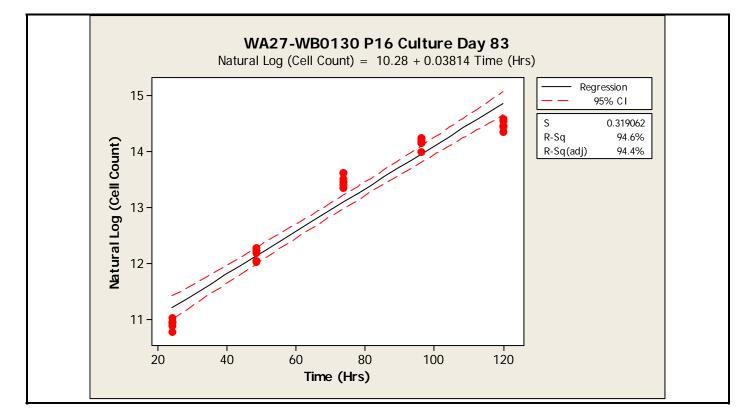
Sample: MA168-12; WA27 #10406

Test: ABO and RH - GF

Description/Molecular Testing	ABO/RH
Enzymatic digestion	X2
Separation by electrophoresis	x3
Amplification each nucleic acid seq	x3
Interpretation and report	X1
	-
	1
	Enzymatic digestion Separation by electrophoresis Amplification each nucleic acid seq Interpretation and report



Sample ID	Cell Line	Cell lot #	Passage	Culture Day		Medium	Matrix	Passag	ing Additive
10450	WA27	WB0130	16	83		E8 + PVA	rh-Vitronectin	Rho-kinase Inhibitor Y-27632	
Documentation of Growth Assay Data						ebook #	Page(s)	Date Initiated	
Doct	intentation	on of Growth Assay Data				148	73-81	0	2MAY12
Growth Assay Performed by		Report Prepared by		Date QA Review		QA Review	ed by	Date	
WiCell Deriv	ation Labor	atory	LAN			14AUG12	JKT	15Aug12	



Regression Analysis: Natural Log (Cell Count) versus Time (Hrs)								
The regressior	n equatio	on is Nati	ural Log (Cell Cou	nt) = 10	.3 + 0.0381 Time	(Hrs)	<u>Slope ± 95% C.I</u> 0.0381 ± 0.0035
Predictor	Coej	f SE	E Coef	т	Р			
Constant	10.284	2 0	.1371	75.00	0.000			Apparent Doubling Time (bours) + 05% C l
Time (Hrs)	0.038	3136 0	.001715	22.24	0.000		<u> </u>	Apparent Doubling Time (hours) ± 95% C.I.
								18.18 ± 2.05
S = 0.319062	R-Sq = 9	94.6% R	-Sq(adj) =	= 94.4%				
		Analy	vsis of Va	riance				Apparent Doubling Time (95% C.I.)
Source	DF	SS	MS		F	Р		16.64 hours – 20.02 hours
Regression	1	50.331	50.33	1 49	4.41	0.000		
Residual Error	· 28	2.850	0.10)2				
Total	29	53.181						



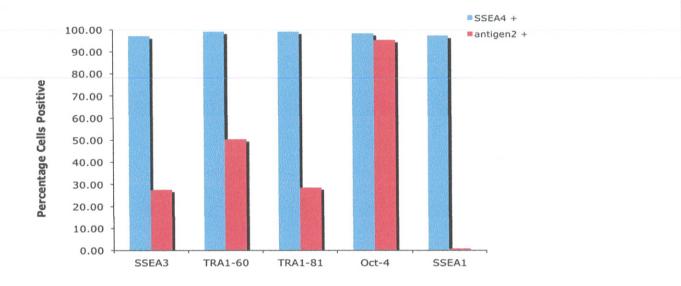
Procedure performed: Cell line: WA27

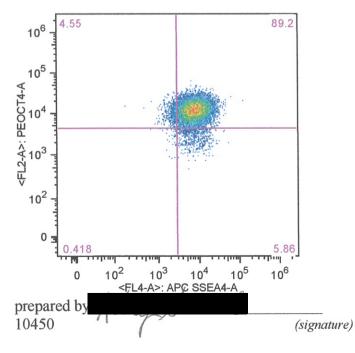
Passage P15 Sample ID: 10450 Date of: (05/05/12) acquisition: file creation: file submission:

			PERCEN	ГS		
	SSEA4 -	SSEA4 +	SSEA4 +	SSEA4 -	ALL	ALL
antigen2:	antigen2 +	antigen2 +	antigen2 -	antigen2 -	SSEA4 +	antigen2 +
SSEA3	0.37	10.30	78.80	10.50	89.10	10.67
TRA1-60	3.00	64.00	30.60	2.36	94.60	67.00
TRA1-81	2.26	44.00	50.00	3.79	94.00	46.26
Oct-4	4.57	89.2	5.85	0.42	95.05	93.77
SSEA1	0.31	3.08	89.10	7.51	92.18	3.39
Democrat and	lungable arrant					

Percent analyzable events: 27.9 #wells submitted: 6

Total # cells analyzed: 10.7×10^6





Sponsor: WiCell Research Institute Accession #: 2012-015912 **Diagnostic Summary Report** 20 Mar 2012 **Received: Approved:** 27 Mar 2012, 13:11 **Bill Method: Test Specimen:** Human ? PDG Sample Set Service (# Tested) Profile Tested + +/-Assay #1 Infectious Disease PCR (3) All Results Negative + = Positive, +/- = Equivocal, ? = Indeterminate, PDG = Pending Service Approvals Service **Approved By*** Date Infectious Disease PCR 27 Mar 2012, 13:11

To assure the SPF status of your research animal colonies, it is essential that you understand the sources, pathobiology, diagnosis and control of pathogens and other adventitious infectious agents that may cause research interference. We have summarized this important information in infectious agent **Technical Sheets**, which you can view by visiting http://www.criver.com/info/disease_sheets.

*This report has been electronically signed by laboratory personnel. The name of the individual who approved these results appears in the header of this service report. All services are performed in accordance with and subject to General Terms and Conditions of Sale found in the Charles River Laboratories-Research Models and Services catalogue and on the back of invoices.

Sponsor: WiCell Research Institute

Approved by

Product: Not Indicated

Test Specimen: Human

Accession #: 2012-015912

Received: 20 Mar 2012

Molecular Diagnostics Infectious Disease PCR Results Report

Department Review:

27 Mar 2012, 13:11*

Human Comprehensive Virus Panel

Sample #:	<u><u>1</u> WA25-WB0132</u>	<u>2</u> WA26-WB0131	<u>3</u> WA27-WB0130
Code :	10429	10430	10431
John Cunningham virus	-	-	-
BK virus	-	-	-
Herpesvirus type 6	-	-	-
Herpesvirus type 7	-	-	-
Herpesvirus type 8	-	-	-
Parvovirus B19	-	-	-
Epstein-Barr Virus	-	-	-
Hepatitis A virus	-	-	-
Hepatitis B virus	-	-	-
Hepatitis C virus	-	-	-
HPV-16	-	-	-
HPV-18	-	-	-
Human T-lymphotropic virus	-	-	-
Human cytomegalovirus	-	-	-
HIV-1	-	-	-
HIV-2	-	-	-
Adeno-associated virus	-	-	-
Human Foamy Virus	-	-	-
LCMV PCR	-	-	-
Hantavirus Hantaan PCR	-	-	-
Hantavirus Seoul PCR	-	-	-
Mycoplasma Genus PCR	-	-	-
DNA Spike	PASS	PASS	PASS
RNA Spike	PASS	PASS	PASS
NRC	PASS	PASS	PASS

Remarks: - = Negative; I = Inhibition, +/- = Equivocal; + = Positive.

Sample Suitability/Detection of PCR Inhibition:

Sample DNA or RNA is spiked with a low-copy number of a exogenous DNA or RNA template respectively. A spike template-specific PCR assay is used to test for the spike template for the purpose of determining the presence of PCR inhibitors. The RNA spike control is also used to evaluate the reverse-transcription of RNA. Amplification of spike template indicates that there is no detectable inhibition and the assay is valid.

NRC:

The nucleic acid recovery control (NRC) is used to evaluate the recovery of DNA/RNA from the nucleic acid isolation process. The test article is spiked with a low-copy number of DNA/RNA template prior to nucleic acid isolation. A template-specific PCR assay is used to detect the DNA/RNA spike.

^{*}This report has been electronically signed by laboratory personnel. The name of the individual who approved these results appears in the header of this service report.