



## Certificate of Analysis - Amended

Product Description	iPS IMR90-1	
Cell Line Provider	University of Wisconsin – Laboratory of Dr. James Thomson	
Lot Number	iPS(IMR90)-1-MCB-01	
Date Viald	01-July-2008	
Passage Number	p33	
Culture Platform	Feeder Independent	
	Media: mTeSR™1	Matrix: Matrigel®

The following testing specifications have been met for the specified product 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 1.2 System by Promega	Consistent with known profile	Pass
Sterility - Direct transfer method	Apptec	30744	Negative	Pass
Mycoplasma	Apptec	30055	No contamination detected	Pass
Karyotype by G-banding	WiCell	SOP-CH-003	Normal karyotype	Pass

<sup>1</sup> These cells were cultured for 32 passages post reprogramming, at least 5 of them in mTeSR™1/Matrigel®. WiCell adds +1 to the passage number to best represent the overall passage number of cells at thaw. Fibroblasts were reprogrammed at p18.

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.

Date of Lot Release	Quality Assurance Approval
10-September-2008	<div>3/12/2018</div> <div>X HEB</div> <div>HEB Quality Assurance Signed by: Bruner, Haley</div>

## Short Tandem Repeat Analysis\*

Sample Report: WiCell 4825

UW HLA#: 59289

Sample Date: 08/11/08

iPS(IMR90)-1-MCB-1  
clone 1

Received Date: 08/11/08

Requestor: WiCell Research Institute

Test Date: 08/11/08

File Name: 080812

Report Date: 08/20/08

Sample Name: (label on tube) DNA 194  
4825-STR

Description: DNA Extracted by WiCell

252.35ug/mL; 260/280 > 1.9

Locus	Repeat #	STR Genotype
D16S539	5, 8-15	Identifying information has been redacted to protect donor confidentiality. If more information is required, please, contact <a href="#">WiCell's Technical Support</a> .
D7S820	6-14	
D13S317	7-15	
D5S818	7-15	
CSF1PO	6-15	
TPOX	6-13	
Amelogenin	NA	
TH01	5-11	
vWA	11, 13-21	

**Comments:** Based on the DNA 194 4825 STR submitted dated and received on 08/11/08 from WiCell, this sample (UW HLA# 59289) matches exactly the STR profile of the human stem cell line iPS(IMR90) comprising 16 allelic polymorphisms across the 8 STR loci analyzed. No STR polymorphisms other than those corresponding to the human iPS(IMR90) 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. These results suggest that the WiCell 4825 DNA sample submitted corresponds to the iPS(IMR90) stem cell line and it 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%. A preliminary copy of this report was issued via electronic mail to the WI Cell Research Institute on Friday, August 22, 2008.

HLA/Molecular Diagnostics Laboratory

HLA/Molecular Diagnostics Laboratory

Date

\* 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.

Report Number

**783236****Page 3 of 5**

WiCell Research Institute

August 11, 2008

P.O. #: [REDACTED]**STERILITY TEST REPORT****Sample Information:** 2: iPS(IMR90)-1-MCB-1 Lot 1, iPS cells**Date Received:** July 23, 2008**Date in Test:** July 25, 2008**Date Completed:** August 08, 2008

**Test Information:** Test Codes: 30744, 30744A  
Immersion, USP / 21 CFR 610.12  
Procedure #: BS210WCR.201

TEST PARAMETERS	PRODUCT	
Approximate Volume Tested	0.5 mL	0.5 mL
Number Tested	2	2
Type of Media	SCD	FTM
Media Volume	400 mL	400 mL
Incubation Period	14 Days	14 Days
Incubation Temperature	20 °C to 25 °C	30 °C to 35 °C
RESULTS	2 NEGATIVE	2 NEGATIVE

QA Reviewed:                     

Page 1 Signed

Reviewed:                     

Page 1 Signed



APPENDIX I

Document #: DCF3008A  
Edition #: 06  
Effective date: 9/17/2003  
Title: DNA FLUOROCHROME ASSAY RESULTS

**DNA-FLUOROCHROME ASSAY RESULTS**

Procedures 3008, 3009, 3011

Sample ID # 54002 M-250 Date Rec'd: 08/05/2008 P.O. #

Indicator Cells Inoculated: Date/Initials: 8/7/08 / JA  
Fixation: Date/Initials: 8/11/08 / JA  
Staining: Date/Initials: 8/11/08 / JA

TEST/CONTROL ARTICLE:

IPS(IMR90) 1-MCB-1

LOT# NA

Wicell QA  
WiCell Research Institute

**DNA FLUOROCHROME ASSAY RESULTS:**

X **NEGATIVE:** A reaction with staining limited to the nuclear region, which indicates no mycoplasmal contamination.

       **POSITIVE:** A significant amount of extranuclear staining which strongly suggests mycoplasmal contamination.

       **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:

Date: 8/11/08 Results Read by: JA Date of Review: 8/11/08 Reviewed by: SSA



**Report Date:** September 10, 2008

## Case Details:

**Cell Line:** iPS(IMR90)-I-MCB-1 (2982-KAR)

**Passage #:** 18+35(8)

**Date Completed:** 9/9/2008

**Cell Line Gender:** Female

**Investigator:** MW

**Specimen:** iPS cells on Matrigel

**Date of Sample:** 9/3/2008

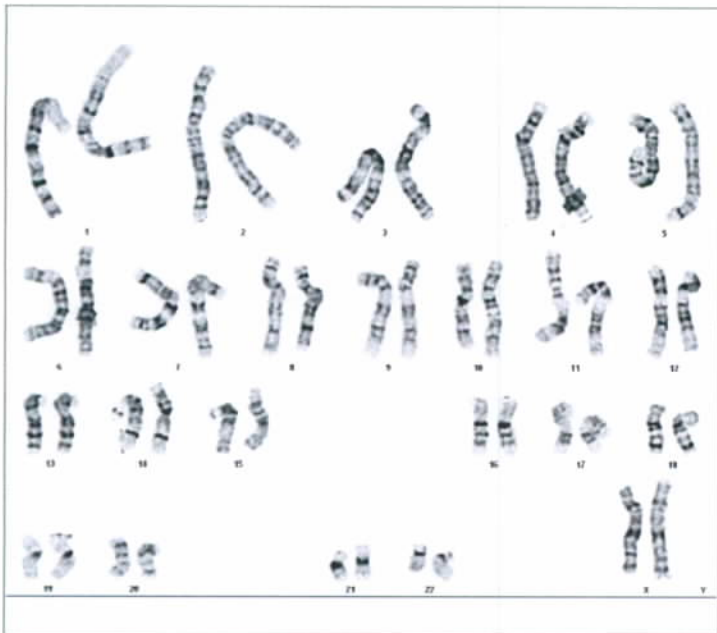
**Tests, Reason for:** Lot release

**Results:** 46,XX

Completed by ST, CLSp(CG), on 9/9/2008

Reviewed and interpreted by KDM, PhD, FACMG, on 9/9/2008

**Interpretation:** No clonal abnormalities were detected at the stated band level of resolution.



**Cell:** S01-01

**Slide:** A

**Slide Type:** Karyotyping

**Cell Results:** Karyotype: 46,XX

**# of Cells Counted:** 40

**# of Cells Karyotyped:** 4

**# of Cells Analyzed:** 8

**Band Level:** 500-550

**Results Transmitted by Fax / Email / Post**  
**Sent By:** \_\_\_\_\_

**Date:** \_\_\_\_\_  
**Sent To:** \_\_\_\_\_