



## Thaw and Culture Details

Cell Line Name	JFHZ3
WiCell Lot Number	DB29774
Provider	Jain Foundation
Banked By	Cellular Dynamics International
Thaw and Culture Recommendations	WiCell recommends thawing 1 vial into 1 well of a 6 well plate.
Culture Platform	Feeder Independent
	Medium: E8 Medium
	Matrix: Matrigel®
Protocol	WiCell Feeder Independent E8 Medium Protocol
Passage Number	p11 These cells were cultured for 11 passages prior to freeze and post reprogramming. Add +1 to the passage number to best represent the overall passage number of the cells at thaw.
Date Vialied	20-August-2014
Vial Label	MyCell® Products Cat #: iPSC Lot #: 01460.103.11 Passage #: 11 Storage Temp Liquid Nitrogen
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.

## Testing Performed by WiCell

Test Description	Test Provider	Test Method	Test Specification	Result
Karyotype by G-banding	WiCell	SOP-CH-003	Expected karyotype	Fail
	<b>Result from report: This is an abnormal karyotype. There is an isochromosome ("i") of the long (q) arm of chromosome 20 in two of twenty cells examined. This imbalance results in trisomy for 20q and monosomy for 20p. i20q is a recurrent acquired abnormality in human pluripotent stem cell cultures. No other clonal abnormalities were found.</b>			
Post-Thaw Viable Cell Recovery	WiCell	SOP-CH-305	Recoverable attachment after passage	Pass
Identity by STR	UW Translational Research Initiatives in Pathology Laboratory	PowerPlex 16 HS System by Promega	Defines profile	Pass
Sterility	Biotest Laboratories	ST/07	Negative	Pass
Mycoplasma	WiCell	SOP-QU-004	Negative	Pass



## Testing Reported by Provider

**This testing was performed prior to banking unless otherwise specified.**

Test Description	Method	Result
Genetic Analysis	Karyotype by G-Banding	Normal Karyotype
Pluripotency	Multiplex RT-PCR to quantify endogenous expression of 7 genes. Scores generated from the analysis predict probability samples are iPSC-like.	Passing sample score $\geq 0.9$
Mycoplasma	Commercially available mycoplasma detection kit.	Negative
Human Virus Testing	HIV I/II CPT Code 87389; detects both antigen and antibodies for HIV I and HIV II. HBV CPT Code 87340; detects Hepatitis B surface antigen. HCV CPT Code 86803; Immunoassay detects Hepatitis C antibody.	Donor samples tested negative for the following human viruses. HIV I HIV II HBV HCV
Identity	Multiplex STR analysis of 9 commonly used alleles.	Match of iPS cell line to incoming donor material.

Approval Date	Quality Assurance Approval
21-December-2015	<div>8/9/2017</div> <div>X AMK</div> <div>AMK Quality Assurance Signed by Klade, Anjelica</div>



## Chromosome Analysis Report: 060428

**Date Reported:** Monday, March 06, 2017

**Cell Line:** JFHZ3-DB29774 12295

**Passage#:** 13

**Date of Sample:** 2/28/2017

**Specimen:** iPSC

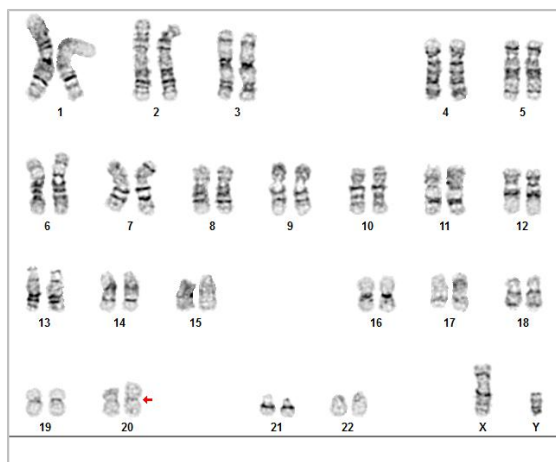
**Results:** 46,XY,i(20)(q10)[2]/46,XY[17]

**Cell Line Gender:** Male

**Reason for Testing:** lot release testing

**Investigator:** [REDACTED], WiCell CDM

**Nonclonal findings:** 47,XY,+12



**Cell:** 5

**Slide:** 1

**Slide Type:** Karyotype

**Total Counted:** 20

**Total Analyzed:** 9

**Total Karyogrammed:** 5

**Band Resolution:** 400 - 450

### Interpretation:

**This is an abnormal karyotype. There is an isochromosome ("i") of the long (q) arm of chromosome 20 in two of twenty cells examined. This imbalance results in trisomy for 20q and monosomy for 20p. i20q is a recurrent acquired abnormality in human pluripotent stem cell cultures. No other clonal abnormalities were found.**

**There is one nonclonal finding, listed above. Nonclonal findings may result from technical artifact, but may be due to a developing clonal abnormality or to low-level mosaicism.**

**Completed by:** [REDACTED], CG(ASCP)

**Reviewed and Interpreted by:** [REDACTED], PhD, FACMG

**A signed copy of this report is available upon request.**

**Date:** \_\_\_\_\_ **Sent By:** \_\_\_\_\_ **Sent To:** \_\_\_\_\_ **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.

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# Short Tandem Repeat Analysis

Department of Pathology and Laboratory Medicine  
TRIP Laboratory (Molecular)  
<http://www.pathology.wisc.edu/research/trip>

**Sample Report:**

12295-STR  
**Sample Name on Tube:** 12295-STR  
61.2 ng/μL, (A260/280=1.79)  
**Sample Type:** Cells  
**Cell Count:** ~2 million cells

**Requestor:**

WiCell Research Institute  
Quality Department

**Sample Date:** N/A

**Receive Date:** 03/06/17  
**Assay Date:** 03/07/17  
**File Name:** 170308 STR TCS  
**Report Date:** 03/10/17

STR Locus	STR Genotype Repeat #	Identifying information has been redacted to protect donor confidentiality. If more information is required, please, contact <a href="#">WiCell's Technical Support</a> .
FGA	16-18,18.2,19,19.2,20,20.2,21,21.2,22, 22.2, 23, 23.2, 24, 24.2, 25, 25.2, 26-30, 31.2, 43.2, 44.2,45.2, 46.2	
TPOX	6-13	
D8S1179	7-18	
vWA	10-22	
Amelogenin	X,Y	
Penta_D	2.2, 3.2, 5, 7-17	
CSF1PO	6-15	
D16S539	5, 8-15	
D7S820	6-14	
D13S317	7-15	
D5S818	7-16	
Penta_E	5-24	
D18S51	8-10, 10.2, 11-13, 13.2, 14-27	
D21S11	24,24.2,25,25.2,26-28,28.2,29,29.2, 30, 30.2,31, 31.2,32,32.2,33,33.2, 34,34.2,35,35.2,36-38	
TH01	4-9,9.3,10-11,13.3	
D3S1358	12-20	

**Results:** Based on the 12295-STR cells submitted by WiCell QA dated and received on 03/06/17, this sample (Label on Tube: 12295-STR) defines the STR profile of the human stem cell line JFHZ3 comprising 23 allelic polymorphisms across the 15 STR loci analyzed.

**Interpretation:** No STR polymorphisms other than those corresponding to the human JFHZ3 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 12295-STR sample submitted corresponds to the JFHZ3 stem cell line and was not contaminated with any other human stem cells or a significant amount of mouse feeder layer cells.

**Sensitivity:** Sensitivity limits for detection of STR polymorphisms unique to either this or other human stem cell lines is ~2-5%.

X *RMB*

Digitally Signed on 03/10/17

TRIP Laboratory, Molecular

X *WMR*

Digitally Signed on 03/10/17

PhD, Director / Co-Director  
UWHC Molecular Diagnostics Laboratory / UWSMPH TRIP Laboratory

# Sterility Report

Biotest Laboratories, Inc.

Making life-saving products possible

WiCell Research Institute, Inc.  
WiCell Quality Assurance  
504 South Rosa Road, Room 101  
Madison, WI 53719

BIOTEST SAMPLE # 16020409

VALIDATION # NG

TEST PURPOSE NG

PRODUCT MIN19i-33811.D-WB20032 11514  
MIN20i-34363.A-WB20384 11515  
MIN21i-34363.B-WB20385 11516  
MIN15i-33363.D-WB20945 11517  
MIN14i-33363.C-WB20811 11518  
MIN17i-33808.B-WB20714 11519  
MIN16i-33808.A-WB20715 11520  
DF19-9-7T-WB0136 11521  
JFHZ3-DB29774 11522  
JFHZ2-DB29769 11523

PRODUCT LOT NA

STERILE LOT NA

BI LOT NA

STERILIZATION LOT NA

BI EXPIRATION DATE NA

STERILIZATION DATE NA

DATE RECEIVED 2016-02-02

STERILIZATION METHOD NA

TEST INITIATED 2016-02-05

SAMPLING BLDG / ROOM NA

TEST COMPLETED 2016-02-19

REFERENCE Processed according to LAB-003: Sterility Test Procedure

Ten (10) products were each divided between 40 mL TSB and 40 mL FTG. The samples were then cultured at 20-25 C and 30-35 C respectively and were monitored for a minimum of 14 days.

- ☒ USP  
☐ BI Manufacturers Specifications  
☐ Other

RESULTS  
Sterile

# POSITIVES  
0

# TESTED  
10

POSITIVE CONTROL  
NA

NEGATIVE CONTROL  
2 Negatives

COMMENTS NA

REVIEWED BY



DATE

22 FEB 16

Specific test results may not be indicative of the characteristics of any other samples from the same lot or similar lots. Liability is limited to the costs of the tests.

Biotest Laboratories • 9303 West Broadway Ave. • Brooklyn Park, MN 55445 • USA • (763) 315-1200

A subsidiary of STERIS Corporation





# Mycoplasma Detection Assay Report

Testing Performed by WiCell

Lot Release

February 21, 2017

FORM SOP-QU-004.01

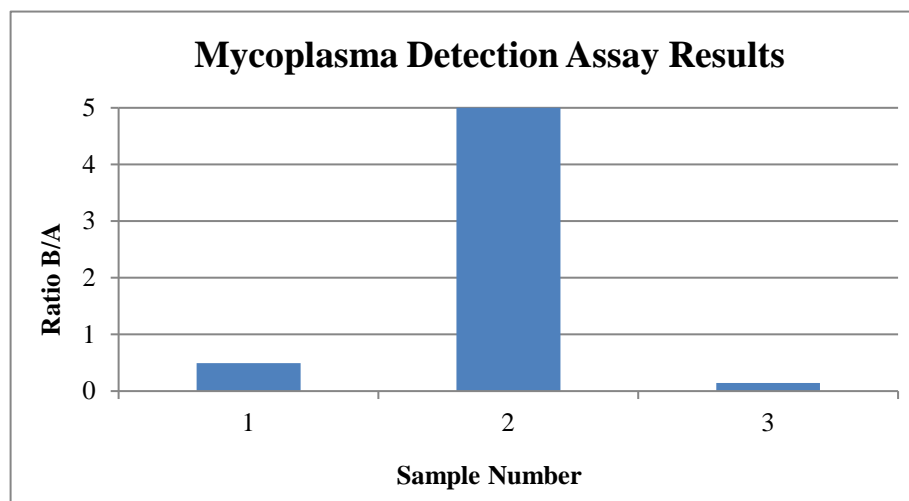
Version F Edition 02

Reported by: OG

Reviewed by: JB

BD Monolight 180

#	Sample Name	Reading A		A Ave	Reading B		B Ave	Ratio B/A	Result	Comments/Suggestions
		RLU1	RLU2		RLU1	RLU2				
1	JFHZ3-DB29774 12295	248	242	245	114	129	121.5	0.50	Negative	
2	Positive (+) Control	331	332	331.5	46049	46466	46258	139.54	Positive	
3	Negative (-) Control	538	563	550.5	76	79	77.5	0.14	Negative	





# Testing Reported by Provider

The testing reports following this placeholder are described on the certificate of analysis found in the beginning of this packet.



## Chromosome Analysis Report: 014915

**Date Reported:** Wednesday, August 13, 2014

**Cell Line:** VIH0036

**Passage#:** 8

**Date of Sample:** 8/5/2014

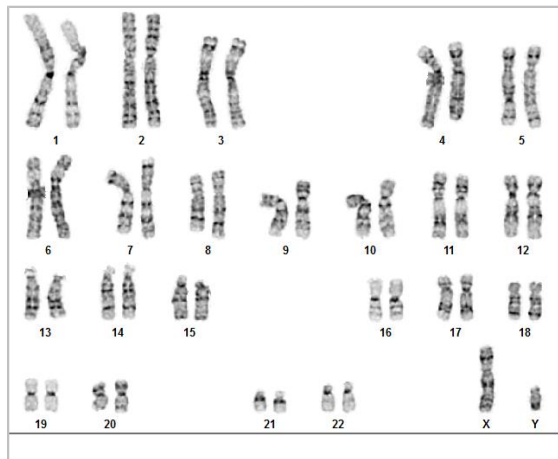
**Specimen:** iPSC

**Results:** 46,XY

**Cell Line Gender:** Male

**Reason for Testing:** Routine testing

**Investigator:** [REDACTED], CDI



**Cell:** 46

**Slide:** 4

**Slide Type:** Karyotype

**Total Counted:** 20

**Total Analyzed:** 8

**Total Karyotyped:** 4

**Band Resolution:** 450 - 525

### Interpretation:

**This is a normal karyotype. No clonal abnormalities were detected at the stated band level of resolution.**

**Completed by:** [REDACTED] MS, CG(ASCP)

**Reviewed and Interpreted by:** [REDACTED], PhD, FACMG

**A signed copy of this report is available upon request.**

**Date:** \_\_\_\_\_ **Sent By:** \_\_\_\_\_ **Sent To:** \_\_\_\_\_ **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".

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