

Thaw and Culture Details

Cell Line Name	JHU053i			
WiCell Lot Number	DB36209			
Provider	Johns Hopkins University – Laboratory of Dr. Lewis Becker			
Banked By	Johns Hopkins University – Laboratory of Dr. Lewis Becker			
Thaw and Culture Recommendations	WiCell recommends thawing 1 vial into 3 wells of a 6 well plate using TeSR [™] -E8 [™] and Recombinant Human Vitronectin. WiCell recommends thawing using ROCK Inhibitor for best results.			
Protocol	WiCell Feeder Independent Pluripotent Stem Cell Protocol			
Culture Platform Prior to Freeze	Feeder Independent			
	Medium: E8			
	Matrix: Vitronectin			
Passage Number	p5 These cells were cultured for 5 passages post reprogramming prior to freeze. Add +1 to the passage number to best represent the overall passage number of the cells at thaw.			
Date Vialed	23-February-2016			
Vial Label	P053 P5 1x10^6 2/23/16			
Biosafety and Use Information	This cell line is of human origin. 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	
	WiCell	SOP-49	Expected karyotype	See Report	
Karyotype by G-banding	Results: 46,XY,del(18)(q21.3)[6]/47,XY,+mar[4]/46,XY[10] Interpretation: This is an abnormal karyotype. There are two unrelated abnormal clones. The cells in the predominant clone (six of twenty cells examined; representative image on the right) contain a terminal deletion of the long (q) arm of chromosome 18. Loss of chromosome 18q is recurrently acquired in pluripotent stem cell cultures. The cells in the secondary clone (four of twenty cells examined; representative image on the left) contain an additional structurally abnormal chromosome (marker, "mar") that cannot be identified by G-banded karyotyping. Additional testing is recommended to aid in identification of the marker chromosome. No other clonal abnormalities were detected at the stated band level of resolution.				
Post-Thaw Viable Cell Recovery	WiCell	SOP-99	Recoverable attachment after passage	Pass	
Identity by STR	WiCell	PowerPlex 16 HS System by Promega	Defines STR profile of deposited cell line	Pass	
Sterility	Steris	ST/07	Negative	Pass	
Mycoplasma	WiCell	SOP-79	Negative	Pass	

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The material provided under this certificate has been subjected to the tests specified and the results and data described herein are accurate based on WiCell's reasonable knowledge and belief. Appropriate Biosafety Level practices and universal precautions should always be used with this material. For clarity, the foregoing is governed solely by WiCell's Terms and Conditions of Service, which can be found at http://www.wicell.org/privacyandterms.



Testing Reported by Provider

The Provider stated that some or all of the additional analyses listed below may have been performed for this cell line. For more information, publication and dbGaP links, where available, are provided on the cell line specific web page on the WiCell website.

- Embryoid bodies
- Infinium[®] Expanded Multi-Ethnic Genotyping Array (MEGA^{EX})

Approval Date	Quality Assurance Approval
08-July-2016	2/11/2021 Xing Min Quality Assurance Signed by Gay, Janna

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Date Reported: Friday, January 15, 2021 Cell Line: JHU053i-DB36209 Submitted Passage #: 6

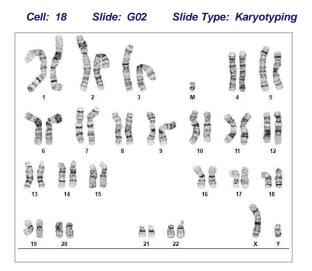
Date of Sample: 1/11/2021

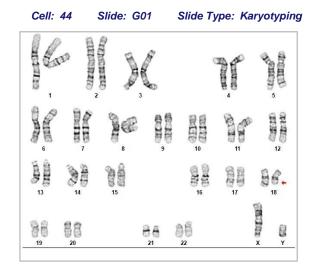
Cell Line Sex: Male Reason for Testing: LOT_RELEASE

Investigator: WiCell Stem Cell Bank, WiCell

Specimen: Human IPSC

Results: 46,XY,del(18)(q21.3)[6]/47,XY,+mar[4]/46,XY[10]





Total Counted: 20 Total Analyzed: 9 Total Karyogrammed: 5 Band Resolution: 425 - 525

Interpretation:

This is an abnormal karyotype. There are two unrelated abnormal clones.

The cells in the predominant clone (six of twenty cells examined; representative image on the right) contain a terminal deletion of the long (q) arm of chromosome 18. Loss of chromosome 18q is recurrently acquired in pluripotent stem cell cultures.

The cells in the secondary clone (four of twenty cells examined; representative image on the left) contain an additional structurally abnormal chromosome (marker, "mar") that cannot be identified by G-banded karyotyping. Additional testing is recommended to aid in identification of the marker chromosome.

No other clonal abnormalities were detected at the stated band level of resolution.

Completed by: Reviewed and Interpreted by:		, PhD, FACMG	
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

Case #: 084425

Cell Line: JHU053i-DB36209

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 of this assay are for research use only. Unless otherwise mutually agreed in writing, the services provided to you hereunder by WiCell Research Institute, Inc. ("WiCell") are governed solely by WiCell's Terms and Conditions of Service, found at www.wicell.org/privacyandterms. Any terms you may attach to a purchase order or other document that are inconsistent, add to, or conflict with WiCell's Terms and Conditions of Service or effect.



Form SOP-89.01 Version 3.0

Requestor: WiCell Stem Cell Bank, WiCell Samples Received: 11Jan21, 14Jan21, 19Jan21, 21Jan21, 25Jan21, 26Jan21 STR Amplification Date: 25Jan21, 28Jan21

Sample Name	<mark>JHU053i-</mark> DB36209 p6	JHU157i- DB36352 p16	EMe-TPint5GCA5- DB67600 p40	EMe-TPint5GCC1- DB67601 p40	MIN28i-35833.A- WB67616 p14	MIN29i-35833.B- WB67612 p12	WA09-WB67614 p26
Label on tube	84425	84426	84447	84448	84468	84469	84476
FGA							
ΤΡΟΧ			1-1				
D8S1179				entifying formation has			
vWA			be	en redacted to			
Amelogenin				otect donor nfidentiality. If			
Penta_D				pre information			
CSF1PO				required,			
D16S539				ease contact			
D7\$820							
D13S317							
D5\$818							
Penta_E							
D18S51							
D21S11							
TH01							
D3S1358							
Allelic Polymorphisms	27	26	24	24	25	25	24
Matches*							See Matches Comment
Comments							

*Note: The STR profile of the following sample is an exact match for the given sample/samples.



Form SOP-89.01 Version 3.0

Requestor: WiCell Stem Cell Bank, WiCell Samples Received: 11Jan21, 14Jan21, 19Jan21, 21Jan21, 25Jan21, 26Jan21 STR Amplification Date: 25Jan21, 28Jan21

Sample Name	WA09-WB67615 p26	MIN30i- 33109.2G- WB67613 p35	MIN27i-35326.K- WB67617 p10	MIN26i-35326.I- WB67609 p26	BWHi009- WB66301 p180	MIN25i-35613.SF- 1-WB67607 p17	EMe-TPint5GC42- DB67599 p39
Label on tube	84477	84496	84531	84534	84550	84551	84552
FGA							
ТРОХ							
D8S1179							
vWA				Identifying			
Amelogenin				Identifying information has			
Penta_D				been redacted to protect donor			
CSF1PO				confidentiality. If			
D16S539				more information is required,			
D7S820				please contact			
D13S317				info@wicell.org			
D5S818							
Penta_E							
D18551							
D21S11							
TH01							
D3S1358							
Allelic Polymorphisms	24	26	29	34	28	26	24
Matches*	See Matches Comment						
Comments				¹ See Triploid Genotype Comment			

*Note: The STR profile of the following sample is an exact match for the given sample/samples.



Requestor: WiCell Stem Cell Bank, WiCell Samples Received: 11Jan21, 14Jan21, 19Jan21, 21Jan21, 25Jan21, 26Jan21 STR Amplification Date: 25Jan21, 28Jan21 Form SOP-89.01 Version 3.0

Sample Name	EMe-TPint5GC23- DB67598 p40
Label on tube	84553
FGA	
ΤΡΟΧ	Identifying
D8S1179	information has
vWA	been redacted to
Amelogenin	protect donor confidentiality. If
Penta_D	more information
CSF1PO	is required, please contact
D16S539	info@wicell.org
D7S820	
D13S317	
D5S818	
Penta_E	
D18S51	
D21S11	
TH01	
D3S1358	
Allelic Polymorphisms	24
Matches*	
Comments	

*Note: The STR profile of the following sample is an exact match for the given sample/samples.



Requestor: WiCell Stem Cell Bank, WiCell Samples Received: 11Jan21, 14Jan21, 19Jan21, 21Jan21, 25Jan21, 26Jan21 STR Amplification Date: 25Jan21, 28Jan21 Form SOP-89.01 Version 3.0

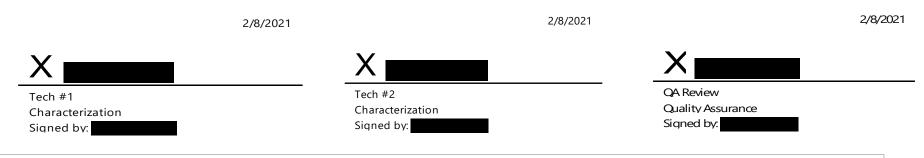
<u>Results:</u> The genotypic profiles comprise a range of 24-34 allelic polymorphisms across the 15 STR loci analyzed.

Interpretation: 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 suggests that the cells submitted correspond to the cell lines as named and were not contaminated with any other human 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 cell lines is ~2-5%.

¹Triploid Genotype: A triploid genotype was detected at the vWA, Penta_D, D16S539, Penta_E, and D18S51 loci. This observation could be the result of chromosomal gain, loss, and/or amplification in this cell line.

Matches: Samples 84476 and 84477 are exact matches to each other and to 14630, 74319, 74844, 74924, 74925, 83593, 84032, 84034, and 84095.



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Raw data is available upon request.

Native Product Sterility Report



WiCell 504 S Rosa Road, Rm 101 Madison, WI 53719		SAMPLE #: DATE RECEIVED: TEST INITIATED: TEST COMPLETED:	21010718 14-Jan-21 20-Jan-21 03-Feb-21
SAMPLE NAME / DESCRIPTION:	BWHi009-WB66301 PENN038i-366-6-DB36313 JHU042i-DB41048 JHU133i-DB41335 JHU053i-DB36209 JHU157i-DB36352		
UNIQUE IDENTIFIER:	JHU233i-DB37038 JHU214i-WB67611 WA09-WB67614 WA09-WB67615 N/A		

TEST RESULTS:	# Tested	# Positives (Growth)	- Control		
	10	0	2 Negatives		
TEST SUMMARY:	# Samples	Media Type	Volume (mL)	Incubation Temperature (° C)	Incubation Duration (Days)
	10	TSB	40	20-25	14
	10	FTG	40	30-35	14
REFERENCE: PD #: TEST METHODOLO	GY:	Processed accord 000053 USP - Direct Tran		erility Test Procedu	Ire

COMMENTS: NA

REVIEWED BY

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DATE OY FEB 2021

Specific test results may not be indicative of the characteristics of any other samples from the same lot or similar lots. This test report shall not be reproduced, except in full, without prior written approval. Liability is limited to the costs of the tests. Results applied to samples as received.



Mycoplasma Assay Report

PCR-based assay performed by WiCell WiCell 13Jan21

Sample Name	Result	Interpretation
MIN27i-35326.K-DB67585 (84386)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
MIN28i-35833.A-DB67586 (84387)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
MIN29i-35833.B-DB67587 (84388)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
MIN30i-33109.2G-DB67588 (84389)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
JHU042i-DB41048 (84391)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
JHU053i-DB36209 (84392)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
JHU133i-DB41335 (84393)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
JHU157i-DB36352 (84394)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
JHU233i-DB37038 (84395)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
STAN245i-601C4-WB67605 (84403)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
STAN366i-282C2-WB67606 (84404)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
STAN274i-729C2-WB67604 (84411)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
MIN24i-35613.B-WB67610 (84414)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
EMe-TPint5GCC1-DB67601 (84416)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
EMe-TPint5GCA5-DB67600 (84417)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
EMe-TPR208X12-DB67602 (84418)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
EMe-TPR208X25-DB67603 (84419)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
Positive (+) Control	Positive	
Negative (-) Control	Negative	

Reported by: Cell C Reviewed by: Senior Cel

, Cell Culture Specialist , Senior Cell Culture Specialist

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A gel image is available upon request.