

## **Thaw and Culture Details**

Cell Line Name	UCSD178i-17-3						
WiCell Lot Number	WB61149						
Provider	University of California, San Diego – Dr. Kelly Frazer						
Banked By	WiCell						
Thaw and Culture Recommendations	WiCell recommends thawing 1 vial into 3 wells of a 6 well plate.						
Culture Platform	Feeder Independent						
	Medium: mTeSR™1						
	Matrix: Matrigel®						
Protocol WiCell Feeder Independent mTeSR <sup>™</sup> 1 Protocol							
Passage Number p18 These cells were cultured for 17 passages prior to freeze and post reprogramming. WiC the passage number to best represent the overall passage number of the cells at thaw.							
Date Vialed 13-March-2017							
Vial Label	UCSD178i-17-3 p18 WB61149						
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>Results:</b> 46,XY Nonclonal findings: 47,XY,+12 47,XY,+20 <b>Interpretation:</b> This is a normal karyotype. No clonal abnormalities were detected at the stated band level of resolution. There are two nonclonal findings, listed above. Nonclonal findings may result from technical artifact, but may be due to a developing clonal abnormality or to low-level mosaicism. Standard analysis requires that chromosomes are counted in twenty cells. Twenty additional cells were examined with no further evidence of this nonclonal aberration.						
Post-Thaw Viable Cell Recovery	WiCell	SOP-CH-305	≥ 15 Undifferentiated Colonies, ≤ 30% Differentiation and 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	Steris	ST/07	Negative	Pass			
Mycoplasma	WiCell	SOP-QU-004	Negative	Pass			

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# **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.

- Illumina® HumanCoreExome BeadChip Array
- RNA-Seq
- Flow Cytometry (SSEA-4, Tra 1-81)
  Infinium<sup>®</sup> Expanded Multi-Ethnic Genotyping Array (MEGA<sup>EX</sup>)

Approval Date	Quality Assurance Approval		
17-April-2017	1/2/2018 XG XG Quality Assurance Signed by Gay, Anna		

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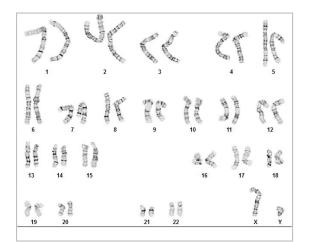


Date Reported: Monday, November 20, 2017 Cell Line: UCSD178i-17-3-WB61149 13063 Passage#: 18 Date of Sample: 11/13/2017 Specimen: Human IPS Results: 46,XY Cell Line Gender: Male Reason for Testing: lot release testing

Investigator:

, WiCell CDM

Nonclonal findings: 47,XY,+12 47,XY,+20



Cell: 7 Slide: G02 Slide Type: Karyotype

Total Counted: 40 Total Analyzed: 8 Total Karyogrammed: 4 Band Resolution: 450 - 550

### Interpretation:

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

There are two nonclonal findings, listed above. Nonclonal findings may result from technical artifact, but may be due to a developing clonal abnormality or to low-level mosaicism. Standard analysis requires that chromosomes are counted in twenty cells. Twenty additional cells were examined with no further evidence of this nonclonal aberration.

Completed by: Reviewed and Interpreted by: A signed copy of this report is a		CG(ASCP) , PhD, FACMG equest.	
Date:	Sent By:	Sent To:	QC Review By:
Limitations: This assay allows for microscopic	visualization of numerica	al and structural chromosome abnormal	ities. The size of structural abnormality that can be detected

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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#### HISTOLOGY - IHC - MOLECULAR - IMAGING

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

Sample Report: 13063-STR Sample Name on Tube: 13063-STR 165.6 ng/µL, (A260/280=1.89) Sample Type: Cells Cell Count: ~2 million cells **Requestor:** WiCell Research Institute Quality Department

Short Tandem Repeat

Analysis

WiCell<sup>®</sup> info@wicell.org (888) 204-1782

Sample Date: N/A Receive Date: 11/20/17 Assay Date: 11/28/17 File Name: STR 171129 wmr Report Date: 11/30/17

STR Locus	ocus STR Genotype Repeat # STR C			
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	Identifying information has		
ТРОХ	6-13	been redacted to		
D8S1179	7-18	protect donor		
vWA	10-22	confidentiality. If		
Amelogenin	X,Y	more information		
Penta D	2.2, 3.2, 5, 7-17	is required,		
CSF1PO	6-15	please, contact WiCell's Technical		
D16S539	5, 8-15	Support.		
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			

<u>Results:</u> Based on the 13063-STR cells submitted by WiCell QA dated and received on 11/20/17, this sample (Label on Tube: 13063-STR) defines the STR profile of the human stem cell line UCSD178i-7-3 comprising 25 allelic polymorphisms across the 15 STR loci analyzed.

<u>Interpretation:</u> No STR polymorphisms other than those corresponding to the human UCSD178i-7-3 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 13063-STR sample submitted corresponds to the UCSD178i-7-3 stem cell line and was not contaminated with any other human stem cells or a significant amount of mouse feeder layer cells.

<u>Sensitivity</u>: 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 12/01/17	X WMR Digitally Signed on 12/01/17
BA	PhD, Director / Co-Director
TRIP Laboratory, Molecular	UWHC Molecular Diagnostics Laboratory / UWSMPH TRIP Laboratory

Testing 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. Acknowledge TRIP in your publications, posters & presentations. For details, see: http://www.pathology.wisc.edu/research/trip/acknowledging TRIP agrees to maintain the confidentiality of any information provided to it in connection with its performance of this STR analysis on the same conditions as set forth in paragraph 2 of WiCell's Terms and Conditions of Service (http://www.wicell.org/media.acux/1a429b84-2b54-44a4-8ad8-5c05db93dd8a).

# Native Product Sterility Report

WiCell 504 S Rosa Rd, Rm 101 Madison, WI 53719

# **STERIS**

SAMPLE #: 17111201 DATE RECEIVED: 16-Nov-17 TEST INITIATED: 20-Nov-17 TEST COMPLETED: 04-Dec-17

### SAMPLE NAME / DESCRIPTION:

iPS(Foreskin)-1-WB666667 13067 UCSD234i-SAD2-3-WB666668 13068 UCSD193i-106-1-WB57372 13069 UCSD178i-17-3-WB61149 13071 UCSD165i-97-1-WB64665 13072 WISC013i-SCID-DB66578 13073 WISC012i-SCA-DB55679 13074 UCSD067i-19-1-WB64878 13075 UCSD166i-98-1-WB59911 13076 UCSD210i-112-1-WB63447 13077 UCSD208i-111-1-WB58973 13079 UCSD160i-92-1-WB61150 13080 UCSD189i-28-1-WB60070 13081 UCSD190i-28-2-WB58714 13082 UCSD191i-13-1-WB65029 13083 UCSD196i-30-1-WB57099 13084 UCSD197i-30-2-WB54408 13085 UCSD202i-108-1-WB57850 13086 UCSD215i-113-1-WB59923 13087 STAN054i-149-2-WB666669 13088

## UNIQUE IDENTIFIER: PRODUCT REGISTRATION:

NA Human iPS Cells

TEST RESULTS:

	# Positives	
# Tested	(Growth)	- Control
20	2	2 Negative

### TEST SUMMARY:

# Samples	Media Type	Volume (mL)	Incubation Temperature (° C)	Incubation Duration (Days)
20	TSB	40	20-25	14
20	FTG	40	30-35	14

## **REFERENCE:**

Processed according to LAB-003: Sterility Test Procedure





METHOD VALIDATION / PD #: TEST METHODOLOGY:

000053 USP - Direct Transfer

COMMENTS:

Sample labeled UCSD208i-111-1-WB58973 13079 was positive in TSB and FTG. Sample #17111201

REVIEWED BY

DATE OGDECIT

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.



## Mycoplasma Detection Assay Report Testing Performed by WiCell

Testing Performed by WiCell Lot Release Testing November 15th, 2017 FORM SOP-QU-004.01 Version G Edition 02 Reported by: SM Reviewed by: JB Berthold Flash n' Glo 539

		Reading A A		Read	ing B	В	Ratio			
#	Sample Name	RLU1	RLU2	Ave	RLU1	RLU2	Ave	B/A	Result	<b>Comments/Suggestions</b>
1	UCSD178i-17-3-WB61149 13063	86	88	87	35	35	35	0.40	Negative	
2	Positive (+) Control	150	136	143	6886	6897	6892	48.19	Positive	
3	Negative (-) Control	246	231	238.5	25	24	24.5	0.10	Negative	

