

Thaw and Culture Details

Cell Line Name	UCSD082i-40-1							
WiCell Lot Number	WB60394							
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.							
	Feeder Independent							
	Medium: mTeSR™1							
	Matrix: Matrigel®							
Protocol WiCell Feeder Independent mTeSR™1 Protocol								
Passage Number p18 These cells were cultured for 17 passages prior to freeze and post reprogramming. WiCell ad the passage number to best represent the overall passage number of the cells at thaw.								
Date Vialed	01-March-2017							
Vial Label	UCSD082i-40-1 p18 WB60394							
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

recurring remaining and remain								
Test Description	Test Provider	Test Method	Test Specification	Result				
Karyotype by G-banding	WiCell	SOP-CH-003	Expected karyotype	Pass				
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	Sterility Steris		Negative	Pass				
Mycoplasma	WiCell	SOP-QU-004	Negative	Pass				

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® Expanded Multi-Ethnic Genotyping Array (MEGAEX)



Approval Date	Quality Assurance Approval			
17-April-2017	9/25/2017 X RK RK Qualify Assurance Signed by Kremers, Erik			



Chromosome Analysis Report: 067663

Date Reported: Thursday, September 07,

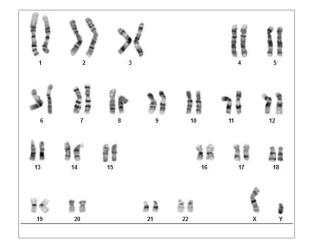
2017

Cell Line: UCSD082i-40-1-WB60394 12776

Passage#: 18

Date of Sample: 8/25/2017 Specimen: Human IPS

Results: 46,XY



Cell Line Gender: Male

Reason for Testing: lot release testing

Investigator:

Cell: 17 Slide: G02

Slide Type: Karyotype

Total Counted: 20
Total Analyzed: 9

Total Karyogrammed: 4
Band Resolution: 425 - 525

Interpretation:

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

Completed by: Reviewed and Interpreted by:



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

info@wicell.org (888) 204-1782

Department of Pathology and Laboratory Medicine TRIP Laboratory (Molecular)

http://www.pathology.wisc.edu/research/trip

Sample Report: 12776-STR

Sample Name on Tube: 12776-STR

 $43.6 \text{ ng/\mu L}, (A260/280=2.26)$

Sample Type: Cells

Cell Count: ~2 million cells

Requestor:

WiCell Research Institute **Ouality Department**

Sample Date: N/A **Receive Date:** 09/05/17

Assav Date: 09/12/17

File Name: 170913 STR WMR

Report Date: 09/15/17

STR Locus	Locus STR Genotype Repeat #							
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						
TPOX	6-13	been redacted to						
D8S1179	7-18	protect donor						
vWA	10-22							
Amelogenin								
Penta D	2.2, 3.2, 5, 7-17	is required,						
CSF1PO	6-15	please, contact						
D16S539	5, 8-15	WiCell's Technical Support.						
D7S820	6-14							
D13S317	7-15							
D5S818								
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 12776-STR cells submitted by WiCell QA dated and received on 09/05/17, this sample (Label on Tube: 12776-STR) defines the STR profile of the human stem cell line UCSD082i-40-1 comprising 29 allelic polymorphisms across the 15 STR loci analyzed.

Interpretation: No STR polymorphisms other than those corresponding to the human UCSD082i-40-1 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 12776-STR sample submitted corresponds to the UCSD082i-40-1 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 $\sim 2-5\%$.

 \mathbf{X} RMB \mathbf{X} WMR **Digitally Signed on Digitally Signed on** PhD. Director / Co-Director TRIP Laboratory, Molecular UWHC Molecular Diagnostics Laboratory / UWSMPH TRIP Laboratory

Native Product Sterility Report



SAMPLE #:

17081954

DATE RECEIVED:

31-Aug-17

TEST INITIATED:

06-Sep-17

TEST COMPLETED:

20-Sep-17

SAMPLE NAME / DESCRIPTION:

WC027i-5807-5-WB66542 12785

WC028i-5807-6-WB66555 12786 WC029i-5907-1-WB66543 12787 WC030i-5907-2-WB66544 12788 WC031i-5907-6-WB66556 12789 UCSD082i-40-1-WB60394 12790 UCSD092i-1-10-WB63301 12791

UCSD093i-1-11-WB64617 12792 MCW109i-40001470-WB66547 12793

MCW064i-40001159-WB66546 12794

UNIQUE IDENTIFIER:

NΑ

PRODUCT REGISTRATION:

Human iPS cells

TEST RESULTS:

WiCell

504 S Rosa Rd, Rm 101

Madison, WI 53719

# 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:

Processed according to LAB-003: Sterility Test Procedure

METHOD VALIDATION / PD #:

000053

TEST METHODOLOGY:

USP - Direct Transfer

COMMENTS:

NA

REVIEWED BY

DATE 218 EPI7

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 August 28, 2017

FORM SOP-QU-004.01 Version F Edition 02 Reported by: KR Reviewed by: JB Berthold Flash n' Glo 539

		Reading A		A	Read	ling B	В	Ratio		
#	Sample Name	RLU1	RLU2	Ave	RLU1	RLU2	Ave	B/A	Result	Comments/Suggestions
1	UCSD082i-40-1-WB60394 12776	95	91	93	51	52	51.5	0.55	Negative	
2	Positive (+) Control	165	166	165.5	17611	17643	17627	106.51	Positive	
3	Negative (-) Control	257	263	260	36	30	33	0.13	Negative	

