Thaw and Culture Details

Cell Line Name	UCSD037i-26-2
WiCell Lot Number	WB65027
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™1 Protocol
Passage Number	p15 These cells were cultured for 14 passages prior to freeze and post reprogramming. WiCell adds +1 to the passage number to best represent the overall passage number of the cells at thaw.
Date Vialed	12-May-2017
Vial Label	UCSD037i-26-2 p15 WB65027
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	See Report
	Results: 46,XY,der(2)inv(2)(p23q31)t(2;8)(p23;q24.2),der(8)t(2;8)(p23;q24.2)[19] Nonclonal findings: 47,XY,+1,der(2)inv(2)(p23q31)t(2;8)(q31;q24.2),der(8)t(2;8)(p23;q24.2) *Interpretation:* This is an abnormal karyotype. There is an apparently balanced translocation between the long (q) arms of chromosomes 2 and 8 in twenty of twenty cells examined. The resulting derivative chromosome 2 also contains a pericentric inversion. No other clonal abnormalities were detected at the stated band level of resolution. There is a nonclonal finding, listed above, which contains the previously mentioned rearrangement and a chromosomal aberration (trisomy 1) recurrently acquired in cultures of this cell type. Nonclonal findings may result from technical artifact, but may be due to a developing clonal abnormality or to low-level mosaicism.			
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

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.

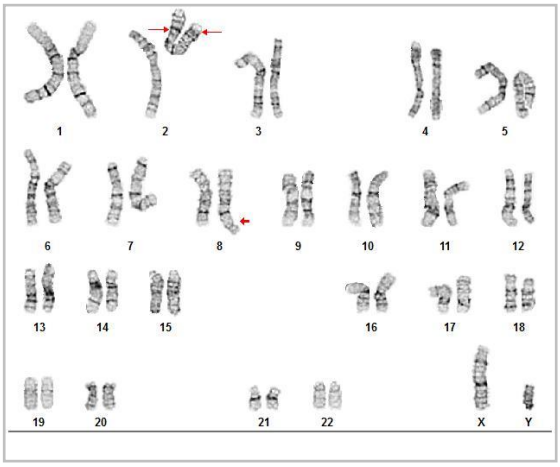
- Illumina® HumanCoreExome BeadChip Array
- RNA-Seq
- Flow Cytometry (SSEA-4, Tra 1-81)
- Infinium® Expanded Multi-Ethnic Genotyping Array (MEGA^{EX})

Approval Date	Quality Assurance Approval
14-June-2017	8/13/2018 X JKG JKG Quality Assurance Signed by: Gay, Jenna

Chromosome Analysis Report: 072411

Date Reported: Friday, July 13, 2018
Cell Line Sex: Male
Cell Line: UCSD037i-26-2-WB65027 13844
Reason for Testing: lot release testing
Passage#: 15
Date of Sample: 7/6/2018
Investigator: [redacted], WiCell
Specimen: Human IPS
Results: 46,XY,der(2)inv(2)(p23q31)t(2;8)(p23;q24.2),der(8)t(2;8)(p23;q24.2)[19]

Nonclonal findings: 47,XY,+1,der(2)inv(2)(p23q31)t(2;8)(q31;q24.2),der(8)t(2;8)(p23;q24.2)



Cell: 70
Slide: G03
Slide Type: Karyotype

Total Counted: 20
Total Analyzed: 8
Total Karyogrammed: 5
Band Resolution: 350 - 475

Interpretation:

This is an abnormal karyotype. There is an apparently balanced translocation between the long (q) arms of chromosomes 2 and 8 in twenty of twenty cells examined. The resulting derivative chromosome 2 also contains a pericentric inversion. No other clonal abnormalities were detected at the stated band level of resolution.

There is a nonclonal finding, listed above, which contains the previously mentioned rearrangement and a chromosomal aberration (trisomy 1) recurrently acquired in cultures of this cell type. 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".

TRIPath
HISTOLOGY - IHC - MOLECULAR - IMAGING
Department of Pathology and Laboratory Medicine
TRIP Laboratory (Molecular)
http://www.pathology.wisc.edu/research/trip

Short Tandem Repeat Analysis

WiCell®
info@wicell.org
(888) 204-1782

Sample Report:
13844-STR
Sample Name on Tube: 13844-STR
77.8 ng/µL, (A260/280=1.81)
Sample Type: Cells
Cell Count: ~2 million cells

Requestor:
WiCell Research Institute
Quality Department

Sample Date: N/A
Receive Date: 07/16/18
Assay Date: 07/17/18
File Name: STR 180718 wmr repeat
Report Date: 07/23/18

STR Locus	STR Genotype Repeat #	STR Genotype
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 been redacted to protect donor confidentiality. If more information is required, please, contact WiCell's Technical Support.
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 13844-STR cells submitted by WiCell QA dated and received on 07/16/18, this sample (Label on Tube: 13844-STR) defines the STR profile of the human stem cell line UCSD037i-26-2 comprising 29 allelic polymorphisms across the 15 STR loci analyzed.

Interpretation: No STR polymorphisms other than those corresponding to the human UCSD037i-26-2 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 13844-STR sample submitted corresponds to the UCSD037i-26-2 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 07/26/18**

[redacted] *BA*
TRIP Laboratory, Molecular

X *WMR* **Digitally Signed on 07/26/18**

[redacted], *PhD, Director / Co-Director*
UWHC Molecular Diagnostics Laboratory / UWSMPH TRIP Laboratory

Native Product Sterility Report

STERIS

SAMPLE #:	17121102
DATE RECEIVED:	14-Dec-17
TEST INITIATED:	14-Dec-17
TEST COMPLETED:	02-Jan-18

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

SAMPLE NAME / DESCRIPTION:
UCSD033i-41-2 WB54901 13153
UCSD037i-26-2 WB65027 13154
UCSD039i-14-3 WB57650 13155
UCSD040i-33-1 WB61158 13156
UCSD041i-33-2 WB60323 13157
UCSD043i-47-1 WB61824 13158
UCSD045i-49-1 WB62417 13159
UCSD046i-50-1 WB60581 13160
UCSD047i-51-1 WB54782 13161
UCSD049i-53-1 WB57867 13162
UCSD114i-69-1 WB55346 13163
UCSD150i-11-1 WB58932 13164
UCSD154i-90-1 WB58798 13165
UCSD164i-96-1 WB58713 13166
UCSD180i-27-2 WB60894 13167
UCSD204i-26-1 WB62522 13168
UCSD216i-114-1 WB65031 13169
UCSD220i-118-1 WB60019 13170
iPS (Foreskin)-4 WB66699 13171
WISC015i-SC7 DB66675 13172

UNIQUE IDENTIFIER: NA

PRODUCT REGISTRATION: Other: Human iPS cells

TEST RESULTS:

# Tested	# Positives (Growth)	- Control
20	0	4 Negatives

TEST SUMMARY:

# Samples	Media Type	Volume (mL)	Incubation Temperature (° C)	Incubation Duration (Days)
20	TSB	40	20-25	15
20	FTG	40	30-35	15

REFERENCE: Processed according to LAB-003: Sterility Test Procedure

METHOD VALIDATION / PD #: 000053

Native Product Sterility Report

STERIS

TEST METHODOLOGY: USP - Direct Transfer

COMMENTS: NA

REVIEWED BY _____ DATE 03JAN18

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

Lot Release Testing

July 5, 2018

FORM SOP-QU-004.01
Version G Edition 02
Reported by: AP
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	UCSD037i-26-2-WB65027 13844	229	221	225	99	92	95.5	0.42	Negative	
2	Positive (+) Control	390	381	385.5	61000	61163	61082	158.45	Positive	
3	Negative (-) Control	820	821	820.5	113	101	107	0.13	Negative