



Thaw and Culture Details

Cell Line Name	UCSD015i-3-3
WiCell Lot Number	WB60296
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	p21 These cells were cultured for 20 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 Vialied	28-February-2017
Vial Label	UCSD015i-3-3 p21 WB60296
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

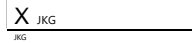
Test Description	Test Provider	Test Method	Test Specification	Result
Karyotype by G-banding	WiCell	SOP-CH-003	Expected karyotype	Pass
	Results: 46,XX,der(10)t(X;10)(p11.2;p15)[20] Interpretation: This is an abnormal karyotype. There is an unbalanced translocation between the short (p) arm of chromosome 10 and the p arm of an X chromosome in twenty of twenty cells examined. This abnormality results in three copies of Xp from Xp11.2 and a cryptic deletion of 10p15. No other clonal abnormalities were found. Karyotyping of the parental culture may help to determine the origin of this abnormality.			
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
23-March-2017	<div style="text-align: right;">12/1/2017</div> <div style="text-align: center;"> JKG Quality Assurance Signed by: Gay, Jenna</div>



Chromosome Analysis Report: 069213

Date Reported: Tuesday, November 21, 2017

Cell Line: UCSD015i-3-3-WB60296 13062

Passage#: 21

Date of Sample: 11/13/2017

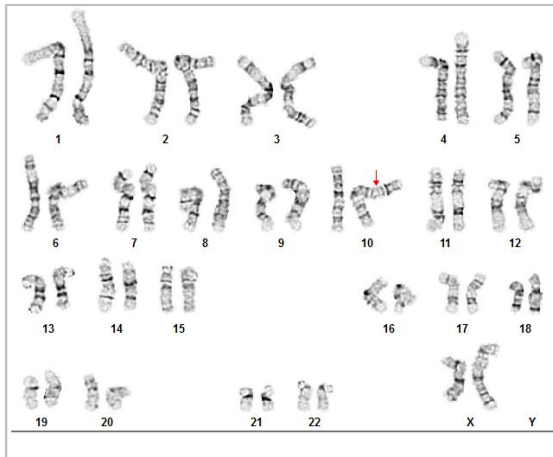
Specimen: Human IPS

Results: 46,XX,der(10)t(X;10)(p11.2;p15)[20]

Cell Line Gender: Female

Reason for Testing: lot release testing

Investigator: [REDACTED], WiCell CDM



Cell: 124

Slide: G02

Slide Type: Karyotype

Total Counted: 20

Total Analyzed: 8

Total Karyogrammed: 4

Band Resolution: 475 - 550

Interpretation:

This is an abnormal karyotype. There is an unbalanced translocation between the short (p) arm of chromosome 10 and the p arm of an X chromosome in twenty of twenty cells examined. This abnormality results in three copies of Xp from Xp11.2 and a cryptic deletion of 10p15. No other clonal abnormalities were found.

Karyotyping of the parental culture may help to determine the origin of this abnormality.

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.

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 are null and void and of no legal force or effect.

Sample Report:

13062-STR

Sample Name on Tube: 13062-STR

124.0 ng/μL, (A260/280=1.85)

Sample Type: Cells**Cell Count:** ~2 million cells**Requestor:**

WiCell Research Institute

Quality Department

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	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 13062-STR cells submitted by WiCell QA dated and received on 11/20/17, this sample (Label on Tube: 13062-STR) defines the STR profile of the human stem cell line UCSD015i-3-3 comprising 27 allelic polymorphisms across the 15 STR loci analyzed.

Interpretation: No STR polymorphisms other than those corresponding to the human UCSD015i-3-3 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 13062-STR sample submitted corresponds to the UCSD015i-3-3 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%.



Digitally Signed on 12/01/17

BA
TRIP Laboratory, Molecular

Digitally Signed on 12/01/17

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

Native Product Sterility Report



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

SAMPLE #: 17041445
DATE RECEIVED: 19-Apr-17
TEST INITIATED: 21-Apr-17
TEST COMPLETED: 05-May-17

SAMPLE NAME / DESCRIPTION: JFMD3 WB62418 12473
JFRBi2 WB62419 12474
JFMD1 WB62435 12475
WISCI004-A-1 WB62846 12476
WISCI004-A-2 WB62848 12477
WISCI004-A-3 WB62903 12478
WISCI004-A-4 WB62825 12479
UCSD017i-3-5 WB54903 12480
UCSD019i-3-7 WB62523 12481
UCSD069-19-3 WB55070 12482
CREM010i-SS9-1 DB47994 12463
CREM011i-SS12-1 DB51676 12464
CREM012i-SS13-1 DB48001 12465
CREM013i-SS14-1 DB48004 12466
CREM014i-SS15-1 DB48007 12467
CREM015i-SS16-1 DB48010 12468
CREM016i-SS18-1 DB48013 12469
CREM017i-SS19-1 DB48016 12470
CREM023i-SS35-1 DB48034 12471
UCSD015i-3-3 WB60296 12472

UNIQUE IDENTIFIER: NA
PRODUCT REGISTRATION: Human iPS cells

TEST RESULTS:

# Tested	# Positives (Growth)	- Control
20	0	3 Negatives

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 #: 000053

Native Product Sterility Report



TEST METHODOLOGY: USP - Direct Transfer

REVIEWED BY 

DATE 08MAY17

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

November 15th, 2017

FORM SOP-QU-004.01

Version G Edition 02

Reported by: SM

Reviewed by: JB

Berthold Flash n' Glo 539

#	Sample Name	Reading A			Reading B			Ratio B/A	Result	Comments/Suggestions
		RLU1	RLU2	Ave	RLU1	RLU2	Ave			
1	UCSD015i-3-3-WB60296 13062	84	90	87	36	36	36	0.41	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	

