

Thaw and Culture Details

Cell Line Name	JHU238i						
WiCell Lot Number	DB37055						
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 5 wells of a 6 well plate. WiCell recommends thawing using ROCK Inhibitor for best results.						
Culture Platform	Feeder Independent						
	Medium: E8						
	Matrix: Vitronectin						
Protocol	WiCell Feeder Independent E8 Medium Protocol						
Passage Number	p16 These cells were cultured for 16 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	22-March-2015						
Vial Label	P238 P16 3/22/15 1.0x10^6						
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

recurring recrimed by threen									
Test Description	Test Provider	Test Method	Test Specification	Result					
Karyotype by G-banding	WiCell	SOP-CH-003	Expected karyotype	See Report					
Post-Thaw Viable Cell Recovery	WiCell	SOP-CH-305	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.

- Embryoid bodies
- Infinium® Expanded Multi-Ethnic Genotyping Array (MEGAEX)



Approval Date	Quality Assurance Approval		
14-July-2016	2/14/2019 X JKG IKG Quality Assurance Signed by Gay, Jenna		



Chromosome Analysis Report: 074666

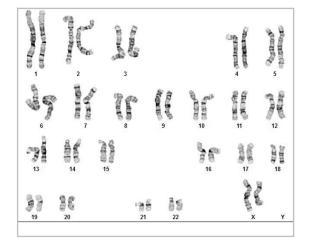
Date Reported: Tuesday, January 22, 2019 Cell Lin

Cell Line: JHU238i-DB37055 14232

Passage#: 17

Date of Sample: 1/15/2019 Specimen: Human IPS

Results: 46,XX



Cell Line Sex: Female

Reason for Testing: Lot Release Testing

Investigator: WiCell

Cell: 12

Slide: G03

Slide Type: Karyotype

Total Counted: 20
Total Analyzed: 8

Total Karyogrammed: 4
Band Resolution: 400 - 500

Interpretation:

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

Completed by: , CG(ASCP)

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



TRIP Laboratory (Molecular)

Short Tandem Repeat HISTOLOGY - IHC - MOLECULAR - IMAGING

WiCell Research Institute

Quality Assurance Department

Requestor:

Analysis



characterization@wicell.org

(608) 316-4145

Receive Date: 01/22/19 **Report Sent:** 01/29/19 **Assav Date:** 01/24/19

File Name: STR 190125 wmr

Report Date: 01/28/19

(608) 265-9168 **Sample Report:**

14232-STR

Sample Name on Tube: 14232-STR

https://research.pathology.wisc.edu/trip/

Department of Pathology and Laboratory Medicine

96.3 ng/ μ L, (A260/280=1.87)

Sample Type: Cells

Cell Count: ~2 million cells

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					
TPOX							
D8S1179	70 7-18						
vWA	vWA 10-22 nelogenin X,Y Penta_D 2.2, 3.2, 5, 7-17 CSF1PO 6-15						
Amelogenin							
Penta_D							
CSF1PO							
D16S539							
D7S820	6-14	Support.					
D13S317	7-15						
D5S818							
Penta_E	5-24						
D18S51							
D21S11							
TH01	4-9,9.3,10-11,13.3						
D3S1358	12-20						

Results: Based on the 14232-STR cells submitted by WiCell QA dated and received on 01/29/18, this sample (Label on Tube: 14232-STR) defines the STR profile of the human stem cell line JHU238i comprising 27 allelic polymorphisms across the 15 STR loci analyzed.

Interpretation: No STR polymorphisms other than those corresponding to the human JHU238i 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 14232-STR sample submitted corresponds to the JHU238i 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 $\sim 2-5\%$.

X RMB	Digitally Signed on	01/29/19	X WMR		Digitally Signed on	01/29/19
TRIP La	, BA boratory, Molecular		UWHC Mole	cular Diag	, Director / Co-Directors / Laboratory / UV	etor VSMPH TRIP Laboratory

Native Product Sterility Report



SAMPLE #:

19011133

DATE RECEIVED:

17-Jan-19

504 S Rosa Road, Rm 101

TEST INITIATED:

22-Jan-19

Madison, WI 53719

WiCell

TEST COMPLETED:

05-Feb-19

SAMPLE NAME / DESCRIPTION:

STAN039i-119-1 WB66980 14235

JHU152i DB36333 14236 JHU176i DB36383 14237 JHU183i DB36760 14238 JHU238i DB37055 14239 JHU006i-1 DB40948 14240

STAN065i-167-1 DB31085 14241 STAN066i-167-2 DB31097 14242 STAN069i-169-1 DB31068 14243 STAN070i-169-2 DB31078 14244

UNIQUE IDENTIFIER:

NA

TEST RESULTS:

	# Positives	
# Tested	(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

PD #:

000053

TEST METHODOLOGY:

USP - Direct Transfer

COMMENTS:

NA

REVIEWED BY

DATE OSFEBIT

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 Detection Assay Report Testing Performed by WiCell

Testing Performed by WiCell Lot Release Testing January 17, 2019

FORM SOP-QU-004.01 Version G Edition 02 Reported by: AP Reviewed by: JB Berthold Flash n' Glow 539

		Reading A		A	Read	ling B	В	Ratio		
#	Sample Name	RLU1	RLU2	Ave	RLU1	RLU2	Ave	B/A	Result	Comments/Suggestions
1	JHU238i-DB37055 14232	115	110	112.5	38	39	38.5	0.34	Negative	
2	Positive (+) Control	73	82	77.5	615	624	619.5	7.99	Positive	
3	Negative (-) Control	267	265	266	30	27	28.5	0.11	Negative	

