



# Certificate of Analysis

## Thaw and Culture Details

Cell Line Name	<b>UCSD242i-LQT1-1</b>	
WiCell Lot Number	<b>DB68089</b>	
Provider/Client	University of California, San Diego – Dr. Kelly Frazer	
Banked By	University of California, San Diego – Dr. Kelly Frazer	
Thaw and Culture Recommendations	WiCell recommends thawing 1 vial into 2 well of a 6 well plate using mTeSR™ 1 and Cultrex®. WiCell recommends thawing using ROCK Inhibitor for best results.	
Protocol	WiCell Feeder Independent Pluripotent Stem Cell Protocol	
Culture Platform Prior to Freeze	mTeSR™ 1	Matrigel®
Passage Number	p17 Cells were cultured for 17 passages prior to freeze. Plated cells at thaw should be labeled passage 18.	
Date Vialied	25-August-2022	
Vial Label	LQT1_2_5 iPSC_P17 ADC_20220825	
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.	



# Certificate of Analysis

## Results

Test Description	Test Provider	Test Method	Test Specification	Result
Karyotype	WiCell	G-T-L Banding performed on 20 metaphase cells	Expected karyotype	See Report
	<b>Results:</b> 46,XX <b>Interpretation:</b> This is a normal karyotype; no clonal abnormalities were detected at the stated band level of resolution.			
Post-Thaw Viable Cell Recovery	WiCell	Thaw using specified Thaw & Culture Recommendations	Recoverable attachment after passage	Pass
Identity by STR	WiCell	PowerPlex 16 HS System by Promega™	Defines STR profile of deposited cell line	See Report
Mycoplasma	WiCell	PCR	Amplification of mycoplasma specific DNA detected with negative result	Pass
Sterility	Steris	Native Product Direct Transfer using FTM and TSB (ST/07)	Negative for growth following 14 days of culture	Pass

Approval Date	WiCell Quality Assurance Approval
30-November-2023	<div style="text-align: right; font-size: small;">11/30/2023</div> <div style="border: 1px solid black; padding: 2px; display: inline-block;"> X Hunter Hefti </div> <div style="font-size: x-small; margin-top: 2px;"> WiCell Quality Assurance  Signed by: Hefti, Hunter </div>

**Date Reported:** Tuesday, August 1, 2023

**Cell Line:** UCSD242i-LQT1-1-DB68089

**Submitted Passage #:** 19

**Date of Sample:** 7/24/2023

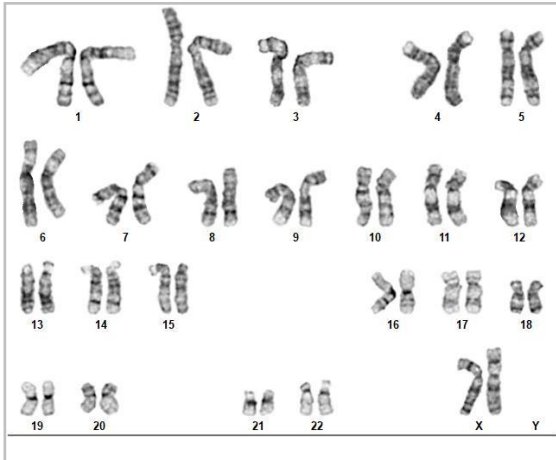
**Specimen:** Human Modified iPSC

**Results:** 46,XX

**Cell Line Sex:** Female

**Reason for Testing:** LOT\_RELEASE

**Investigator:** WiCell Stem Cell Bank, WiCell



**Cell:** 78

**Slide:** G05

**Slide Type:** Karyotype

**Total Counted:** 20

**Total Analyzed:** 8

**Total Karyogrammed:** 4

**Band Resolution:** 425 - 500

**Interpretation:**

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

**Completed by:** Erica Schutter, CG(ASCP)

**Reviewed and Interpreted by:** Vanessa Horner, PhD, FACMG

*For internal use only*

**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](http://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.*



# Short Tandem Repeat

Requestor: WiCell Stem Cell Bank, WiCell

Samples Received: 24Jul23

STR Amplification Date: 26Jul23

Form SOP-89.01

Version 10.0

Sample Name	UCSD243i-LQT3-1-DB68091 p17	UCSD242i-LQT1-1-DB68089 p19
WiCell CTR No. <sup>1</sup>	98010	98009
FGA	Identifying information has been redacted to protect donor confidentiality. If more information is required, please contact <a href="mailto:info@wicell.org">info@wicell.org</a>	
TPOX		
D8S1179		
vWA		
Amelogenin		
Penta_D		
CSF1PO		
D16S539		
D7S820		
D13S317		
D5S818		
Penta_E		
D18S51		
D21S11		
TH01		
D3S1358		
Allelic Polymorphisms	26	26
Matches*	See Matches Comment	See Matches Comment
Comments		

*\*Note: The STR profile of the following sample is a 100% match for the given sample/samples unless otherwise specified.*

<sup>1</sup> CTR No.: Characterization Test Request Number; also known as a laboratory accessioning number.



# Short Tandem Repeat

Requestor: WiCell Stem Cell Bank, WiCell

Samples Received: 24Jul23

STR Amplification Date: 26Jul23

Form SOP-89.01

Version 10.0

**Assay Description:** STR analysis is performed using the PowerPlex 16 HS System by Promega™. Results are reported as 13 CODIS STR markers, Amelogenin for gender determination and two low-stutter, highly discriminating pentanucleotide STR markers.

**Results:** The genotypic profiles comprise a range of 26 allelic polymorphisms across the 15 STR loci analyzed.

**Interpretation:** 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. These results suggests that the cells submitted correspond to the cell lines as named and were not contaminated with any other human 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 cell lines is ~2-4%.

**Matches:** Samples 98010 and 98009 are a 100% match to each other and to 97709, 97621, 73192, 72643, 72297, and 72296.

7/31/2023	8/2/2023	8/1/2023
<p><b>X</b> Justin Hobson</p> <hr/> <p>Tech #1 Characterization Signed by: Hobson, Justin</p>	<p><b>X</b> Amber Kuhn</p> <hr/> <p>Tech #2 Characterization Signed by: Kuhn, Amber</p>	<p><b>X</b> Ryen Smith</p> <hr/> <p>QA Review Quality Assurance Signed by: Smith, Ryen</p>

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# Mycoplasma Assay Report

PCR-based assay performed by WiCell  
WiCell Stem Cell Bank, WiCell  
18Jul23

Form SOP-83.01  
Version 5.0

Sample Name	Result	Interpretation
UCSD243i-LQT3-1-DB68091 p16 (97932)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
UCSD242i-LQT1-1-DB68089 p18 (97931)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
H1 MYH11-NLuc-tdTomato-WB68133 p32 (97828)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
Positive (+) Control	Positive	
Negative (-) Control	Negative	

Assay Description
Sample is tested for presence of mycoplasma using EZ-PCR™ Mycoplasma Detection Kit (Sartorius).

7/18/2023	7/18/2023	7/19/2023
<b>X</b> Justin Hobson <hr/> Tech #1 Characterization Signed by: Hobson, Justin	<b>X</b> Kaylie Petersen <hr/> Tech #2 Characterization Signed by: Petersen, Kaylie	<b>X</b> Dawn Graham <hr/> QA Review Quality Assurance Signed by: Graham, Dawn

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*A gel image is available upon request.*

# Native Product Sterility Report



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

SAMPLE #: 23040601  
DATE RECEIVED: 13-Apr-23  
TEST INITIATED: 14-Apr-23  
TEST COMPLETED: 28-Apr-23

SAMPLE NAME / DESCRIPTION: PACS2iPS01-C5-DB68076  
WA09-WB68097  
HES7-NLuc-2A-tdtomato-DB68082  
H9 T-2A-EGFP-PGK-DB68085  
H1-Fucci-DB68086  
H1 MYH11-NLuc-tdTomato-DB68087  
HES7-NLuc-2A-tdtomato-DB68090  
JHU255i-DB37129  
JHU256i-DB37132  
JHU257i-DB37136  
UCSD242i-LQT1-1-DB68089  
UCSD243i-LQT3-1-DB68091  
UCSD244i-LQT3-2-DB68092  
UCSD245i-CNTL-1-DB68093  
UCSD246i-CNTL-2-DB68094  
UCSD247i-LQT1-2-DB68095  
UCSD182i-3-2-DB68096  
JHU252i-DB37121  
JHU013i-2-DB40951  
JHU022i-DB40963

UNIQUE IDENTIFIER: N/A

## TEST RESULTS:

# Tested	# Positives (Growth)	- Control
20	1	2 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

# Native Product Sterility Report



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

PD #: 000053

TEST METHODOLOGY: USP - Direct Transfer

COMMENTS: Sample #23040601

Sample labeled H1myh11 c5 p28 5cm2 jab 28AUG17 is positive for both TSB and FTG media.

AUTHORIZED BY \_\_\_\_\_

A handwritten signature in blue ink, appearing to be "AD", written over a horizontal line.

DATE 09 MAY 2023

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.