

# **Certificate of Analysis**

### **Thaw and Culture Details**

Cell Line Name	SCRP8305i		
WiCell Lot Number	DB43120		
Provider	The Scripps Research Institute – Laboratory of Dr. Eric Topol		
Banked By	Gladstone Institutes – Laboratory of Dr. Sheng Ding		
Thaw and Culture Recommendations	WiCell recommends thawing 1 vial into 4 wells of a 6 well plate using using mTeSR™1 and Matrigel®. WiCell recommends thawing using ROCK Inhibitor for best results.		
Culture Platform	Feeder Independent		
	Medium: mTeSR™1		
	Matrix: Matrigel®		
Protocol	WiCell Feeder Independent Pluripotent Stem Cell Protocol		
Passage Number	p10 These cells were cultured for 10 passages after colony picking prior to freeze. Add +1 to the passage number to best represent the overall passage number of the cells at thaw.		
Date Vialed	20-May-2016		
Vial Label	HE00812, Passage 10, May-20-2016		
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.		



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### **Results**

Test Description	Test Provider	Test Method	Test Specification	Result	
	WiCell G-T-L Banding performed on 20 metaphase cells		Expected karyotype	See Report	
	Results: 45,X[20	]			
Karyotype	Interpretation: This is an abnormal karyotype. There is only one sex chromosome, an X, in all cells examined. This analysis cannot determine the cell line sex. No other clonal abnormalities were detected at the stated band level of resolution.				
	Comparison of this karyotype with the karyotype of the source (parental) specimen may be informative regarding the significance and origin of the chromosomal abnormality.				
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 <sup>™</sup>	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	



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## **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.

- HumanCore Exome Kit
- Methylation
- Tra1-60 marker expression via flow cytometry
- Infinium® Expanded Multi-Ethnic Genotyping Array (MEGAEX)

Approval Date	Quality Assurance Approval	
12-September-2016	7/30/2025  X HEB  HEB  Quality Assurance Signed by Brune, Haley	



#### Chromosome Analysis Report: 106449

Date Reported: March 29, 2025 Cell Line: SCRP8305i-DB43120

Submitted Passage #: 13

Date of Sample: 3/17/2025

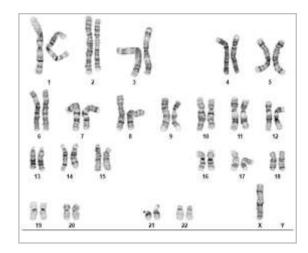
Specimen: Human IPSC

Results: 45,X[20]

Cell Line Sex: Unknown

Reason for Testing: LOT\_RELEASE

Investigator: WiCell Stem Cell Bank, WiCell



Cell: 20 Slide: G02

Slide Type: Karyotype

Total Counted: 20
Total Analyzed: 8

Total Karyogrammed: 4
Band Resolution: 425 - 525

#### Interpretation:

This is an abnormal karyotype. There is only one sex chromosome, an X, in all cells examined. This analysis cannot determine the cell line sex. No other clonal abnormalities were detected at the stated band level of resolution.

Comparison of this karyotype with the karyotype of the source (parental) specimen may be informative regarding the significance and origin of the chromosomal abnormality.

Completed by: Kate Bird, CG(ASCP)

Reviewed and Interpreted by: Xianggiang Shao, PhD, FACMG

For internal use only			
Date:	Sent By:	Sent To:	QC Review By:
Limitations: This assay allows for microscopic visu	ialization of numerical a	nd structural chromosome abnormalities	The size of structural abnormality that can be detected

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.



## **Short Tandem Repeat**

Requestor: WiCell Stem Cell Bank, WiCell Sample Receipt Date: 17Mar25 STR Amplification Date: 24Mar25

Sample Name	SCRP8305i- DB43120 p13	SCRP6101i- DB42990 p12	
WiCell CTR No.1	106449	106448	
FGA			
TPOX			
D8S1179	- Identifyin	a	
vWA	informati	on has	
Amelogenin	been red		
Penta_D	protect donor confidentiality. If more information is required,		
CSF1PO			
D16S539	please co	ontact	
D7S820	info@wicell.org		
D13S317			
D5S818	_		
Penta_E	_		
D18S51	_		
D21S11	_		
TH01	_		
D3S1358	_		
Allelic Polymorphisms			
Matches <sup>2</sup>			
Comments			

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

<sup>&</sup>lt;sup>2</sup> The STR profile of the sample(s) listed are a 100% match for the given sample unless otherwise specified.



### **Mycoplasma Assay Report**

Form SOP-83.01 Version 7.0

PCR-based assay performed by WiCell WiCell Stem Cell Bank, WiCell 26Mar25

Sample Name	Result	Interpretation
SCRP8305i-DB43120 p15 (106579)	<b>Negative</b>	Band was not seen at 270bp, indicating the absence of mycoplasma.
SCRP6101i-DB42990 p14 (106578)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
SCRP5603i-DB42976 p16 (106577)	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).

	3/26/2025	3/27/2025	3/28/2025
X John Raff	X	Michael Mussar	X Dawn Graham
Tech #1 Characterization Signed by: Raff, John	Ch	ch #2 aracterization ned by: Mussar, Michael	QA Review Quality Assurance Signed by: Graham, Dawn

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A gel image is available upon request.



### **Short Tandem Repeat**

Form SOP-89.01 Version 14.0

Requestor: WiCell Stem Cell Bank, WiCell Sample Receipt Date: 17Mar25 STR Amplification Date: 24Mar25

<u>Assay Description:</u> Short Tandem Repeat (STR) analysis is performed using the PowerPlex® 16 HS System by Promega<sup>TM</sup>. Results are reported as 13 CODIS STR markers, Amelogenin for sex determination and two low-stutter, highly discriminating pentanucleotide STR markers.

**Results:** The genotypic profiles comprise a range of 26-27 allelic polymorphisms across the 15 STR loci analyzed.

<u>Interpretation:</u> 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 suggest 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.

<u>Sensitivity:</u> Sensitivity limits for detection of STR polymorphisms unique to either this or other human cell lines is ~2-4%.

3/27/2025	3/31/2025	4/1/2025
X John Raff  Tech #1 Characterization	X Amber Kuhn  Tech #2 Characterization	Dawn Graham      QA Review     Quality Assurance
Signed by: Raff, John	Signed by: Kuhn, Amber	Signed by: Graham, Dawn

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## Native Product Sterility Report



SAMPLE #:

21070812

DATE RECEIVED:

14-Jul-21

TEST INITIATED:

27-Jul-21

TEST COMPLETED:

10-Aug-21

SAMPLE NAME / DESCRIPTION:

504 S Rosa Road, Rm 101

Madison, WI 53719

WiCell

CREM017i-SS19-1-WB67673 PENN042i-258-12-WB67671 UCSD239i-APP2-1-WB67672 STAN151i-303C3-DB35736 STAN248i-617C1-DB35488 STAN249i-617C2-DB35491

WA01-WB67657 WA01-WB67656

STAN366i-282C2-WB67655

SCRP5803i-DB42982 SCRP6101i-DB42990 SCRP6904i-DB43007 SCRP7301i-DB43010 HVRDi001-A-WB67674 SCRP8105i-DB43117 SCRP8305i-DB43120 SCRP8503i-DB43126 SCRP8601i-DB43129 SCRP8717i-DB43132 SCRP8901i-DB43135

UNIQUE IDENTIFIER:

N/A

**TEST RESULTS:** 

# Tested	# Positives (Growth)	- Control
20	0	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

REFERENCE:

Processed according to LAB-003: Sterility Test Procedure

PD #:

000053

TEST METHODOLOGY:

**USP** - Direct Transfer

STERIS 9303 West Broadway Ave Brooklyn Park, MN 55445

LAB-003 rev 35 Form 5 Effective: APR 06, 2021 Page 1 of 2

## Native Product Sterility Report



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Sample # 21070812

REVIEWED BY DATE 13 pug 2021

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.