

Certificate of Analysis

Thaw and Culture Details

Cell Line Name	STAN227i-516C5
WiCell Lot Number	DB35603
Provider	Stanford University – Laboratory of Dr. Thomas Quetermous
Banked By	Icahn School of Medicine at Mount Sinai Stem Cell Core
Thaw and Culture Recommendations	WiCell recommends thawing 1 vial into 2 wells of a 6 well plate using mTeSR™1 and Matrigel®.
Culture Platform	Feeder Independent
	Medium: mTeSR1 [™]
	Matrix: Matrigel [®]
Protocol	WiCell Feeder Independent mTeSR1 [™] Protocol
Passage Number	p11 These cells were cultured for 11 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	05-December-2015
Vial Label	ISMMS 516i C5 P11 MM 120515
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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Certificate of Analysis

Results

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Test Description	Test Provider Test Method		Test Specification	Result	
	WiCell	G-T-L Banding performed on 20 metaphase cells	Expected karyotype	See Report	
Karyotype	Results: 46,XX Interpretation: 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	

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.

- RNA-Seq
- Whole Genome Sequencing
- Infinium[®] Expanded Multi-Ethnic Genotyping Array (MEGA^{EX})

Approval Date	Quality Assurance Approval	
07-November-2016	4/90/2025 X HEB Guality Assurance Signed by: Hilfruner	

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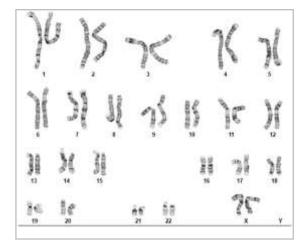
The material provided under this certificate has been subjected to the tests specified and the results and data described herein are accurate based on WiCell's reasonable knowledge and belief. Appropriate Biosafety Level practices and universal precautions should always be used with this material. For clarity, the foregoing is governed solely by WiCell's Terms and Conditions of Service, which can be found at http://www.wicell.org/privacyandterms.



Chromosome Analysis Report: 106844

Date Reported: April 17, 2025 Cell Line: STAN227i-516C5-DB35603 Submitted Passage #: 13 Date of Sample: 4/3/2025 Specimen: Human IPSC Results: 46,XX Cell Line Sex: Female Reason for Testing: LOT_RELEASE

Investigator: WiCell Stem Cell Bank, WiCell



Cell: 1 Slide: G02 Slide Type: Karyotype Total Counted: 20 Total Analyzed: 8 Total Karyogrammed: 4 Band Resolution: 450 - 500

Interpretation:

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

Completed by: Reviewed and Interpreted by: Jennifer Pecos, CG(ASCP) Justin Schleede, 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. 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 or effect.



Short Tandem Repeat

Requestor: WiCell Stem Cell Bank, WiCell Sample Receipt Date: 03Apr25, 04Apr25 STR Amplification Date: 15Apr25

STAN226i-516C3-STAN227i-516C5-STAN236i-551C1-Sample Name DB35603 p13 DB35595 p13 DB35687 p16 WiCell CTR No.1 106846 106845 106844 FGA TPOX D8S1179 Identifying vWA information has Amelogenin been redacted to protect donor Penta_D confidentiality. If more information CSF1PO is required, D16S539 please contact D7S820 D13S317 D5S818 Penta_E D18S51 D21S11 **TH01** D3S1358 **Allelic Polymorphisms** Matches² 106844 106715 106846 Comments

Form SOP-89.01 Version 15.0

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

² The STR profile of the sample(s) listed are a 100% match for the given sample unless otherwise specified.



Short Tandem Repeat Requestor: WiCell Stem Cell Bank, WiCell Sample Receipt Date: 03Apr25, 04Apr25 STR Amplification Date: 15Apr25 Form SOP-89.01 Version 15.0

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

<u>Results:</u> The genotypic profiles comprise a range of 25-28 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 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.

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

4/18/2025	4/21/2025	4/22/2025
X John Raff	X Anna Lisa Larson	X Dawn Graham
Tech #1	Tech #2	QA Review
Characterization	Characterization	Quality Assurance
Signed by: Raff, John	Signed by: Larson, Anna Lisa	Signed by: Graham Dawn

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Mycoplasma Assay Report

Form SOP-83.01 Version 7.0

PCR-based assay performed by WiCell WiCell Stem Cell Bank, WiCell 04Apr25

Sample Name	Result	Interpretation
STAN236i-551C1-DB35687 p16 (106845)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
STAN227i-516C5-DB35603 p13 (106844)	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).

4/4/2025	4/4/2025	4/9/2025
X Nina Montgomery	X John Raff	X Dawn Graham
Tech #1 Characterization Signed by: Montgomery, Nina	Tech #2 Characterization Signed by: Raff, John	QA Review Quality Assurance Signed by: Graham, Dawn

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A gel image is available upon request.

Native Product Sterility Report



		SAMPLE #:	25030859
Accounting@wicell.org; WiCell Research	DATE RECEIVED:	22-Mar-25	
504 S Rosa Road, Rm 101		TEST INITIATED:	28-Mar-25
Madison, WI 53719		TEST COMPLETED:	11-Apr-25
SAMPLE NAME / DESCRIPTION:	CBiPS-E12C1-PCBC-WB68767		
	STAN227i-516C5-DB35603		
	STAN237i-551C2-DB35694		
	STAN226i-516C3-DB35595		

STAN227i-516C5-DB35603
STAN237i-551C2-DB35694
STAN226i-516C3-DB35595
STAN236i-551C1-DB35687
JHU129i-DB41332
JHU139i-DB36275
JHU216i-DB36861
JHU123i-DB41320
JHU231i-DB37030
JHU220i-DB41414
JHU067i-DB36212
JHU159i-DB41368
JHU124i-DB41323
JHU205i-DB36820
JHU089i-DB41243
JHU200i-DB36800
JHU141i-DB41341
JHU111i-DB36250
JHU074i-DB41131

UNIQUE IDENTIFIER:

TEST RESULTS:		# Positives	
	# Tested	(Growth)	- Control
	20	0	2 Negatives
			1

TEST SUMMARY

Y:	# Samples	Media Type	Volume (mL)	Incubation Temperature (° C)	Incubation Duration (Days)	
	20	TSB	40	20-25	14	
	20	FTG	40	30-35	14	

REFERENCE:

PD #:

TEST METHODOLOGY:

Processed according to LAB-003: Sterility Test Procedure 000053 USP - Direct Transfer

STERIS 9303 West Broadway Ave Brooklyn Park, MN 55445

Native Product Sterility Report



COMMENTS: Sa

Sample #25030859

AUTHORIZED BY

DATE 11APR 2025

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.