

Thaw and Culture Details

Cell Line Name	CREM032i-SS48-1					
WiCell Lot Number	WB66711					
Parent Material	CREM032i-SS48-1- DB48070					
Provider	Boston University – Laboratory of Dr. Martin Steinberg					
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	p11 These cells were cultured for 10 passages after colony picking prior to freeze. WiCell adds +1 to the passage number at freeze to best represent what the overall passage number of the cells at thaw. Plated cells at thaw should be labeled passage 11.					
Date Vialed	14-December-2017					
Vial Label	CREM032i-SS48-1 p11 WB66711					
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 Specification	Result					
	WiCell	SOP-CH-003	Expected karyotype	See Report				
Karyotype by G-banding	Interpretation: This is a stated band level of res analysis requires that cluvere examined with no	terpretation: This is a normal karyotype. No clonal abnormalities were detected at the ated band level of resolution. There is one nonclonal finding, listed above. Standard halysis requires that chromosomes are counted in twenty cells. Twenty additional cells are examined with no further evidence of this nonclonal aberration. Nonclonal findings ely result from technical artifact, but may be due to a developing clonal abnormality or to welevel mosaicism.						
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.

- Digital Genome Sequencing
- Infinium® Expanded Multi-Ethnic Genotyping Array (MEGAEX)

Approval Date	Quality Assurance Approval			
11-February-2018	7/8/2018 X JKG JKG Quality Assurance Signed by Gay, Jenna			



Chromosome Analysis Report: 070203

Date Reported: Friday, January 26, 2018

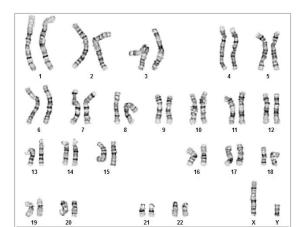
Cell Line: CREM032i-SS48-1-WB66711 13235

Passage#: 11

Date of Sample: 1/23/2018 Specimen: Human IPSC

Results: 46,XY

Nonclonal findings: 47,XY,+8



Cell Line Gender: Male

Reason for Testing: lot release testing

Investigator: , WiCell CDM

Cell: 52 Slide: G03

Slide Type: Karyotype

Total Counted: 40
Total Analyzed: 8
Total Karyogrammed: 4
Band Resolution: 450 - 525

Interpretation:

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

There is one nonclonal finding, listed above. Standard analysis requires that chromosomes are counted in twenty cells. Twenty additional cells were examined with no further evidence of this nonclonal aberration. Nonclonal findings likely result from technical artifact, but may be due to a developing clonal abnormality or to low-level mosaicism.

Completed By: Reviewed and Interpreted By: , CG(ASCP) , PhD, FACMG

A signed copy of this report is available upon request.

Date:	Sent By:	Sent To:	QC Review By:
Dutc	ocin by	OCH 10	quincinelli 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.



Short Tandem Repeat Analysis

HISTOLOGY - IHC - MOLECULAR - IMAGING

Department of Pathology and Laboratory Medicine TRIP Laboratory (Molecular)

http://www.pathology.wisc.edu/research/trip

WiCell® info@wicell.org (888) 204-1782

Sample Report:

13235-STR

Sample Name on Tube: 13235-STR

 $40.0 \text{ ng/}\mu\text{L}, (A260/280=1.80)$

Sample Type: Cells

Cell Count: ~2 million cells

Requestor:

WiCell Research Institute Quality Department Sample Date: N/A Receive Date: 01/29/18 Assay Date: 01/30/18

File Name: STR 180131 wmr

Report Date: 02/08/18

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	NV 6 13							
D8S1179	7-18	been redacted to protect donor						
vWA	10-22	confidentiality. If						
Amelogenin	X,Y	more information						
Penta_D	2.2, 3.2, 5, 7-17	is required,						
CSF1PO	6-15	please, contact						
D16S539	5, 8-15	WiCell's Technical Support.						
D7S820	6-14							
D13S317	7-15							
D5S818	D5S818 7-16 Penta_E 5-24 D18S51 8-10, 10.2, 11-13, 13.2, 14-27							
Penta_E								
D18S51								
D21S11	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							
TH01								
D3S1358	D3S1358 12-20							

<u>Results:</u> Based on the 13235-STR cells submitted by WiCell QA dated and received on 01/29/18, this sample (Label on Tube: 13235-STR) defines the STR profile of the human stem cell line CREM032i-SS48-1 comprising 26 allelic polymorphisms across the 15 STR loci analyzed.

<u>Interpretation:</u> No STR polymorphisms other than those corresponding to the human CREM032i-SS48-1 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 13235-STR sample submitted corresponds to the CREM032i-SS48-1 stem cell line and was not contaminated with any other human stem 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 stem cell lines is ~2-5%.

X RMB Digitally Signed on 02/08/18

X WMR Digitally Signed on 02/08/18

BA
TRIP Laboratory, Molecular

Digitally Signed on 02/08/18

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

Native Product Sterility Report



SAMPLE #:

17121502

WiCell

DATE RECEIVED:

21-Dec-17

504 S. Rosa Rd., Rm 101 Madison, WI 53719 TEST INITIATED:

26-Dec-17

TEST COMPLETED:

09-Jan-18

SAMPLE NAME / DESCRIPTION:

UCSD050i-54-1 WB54411 13186 UCSD051i-55-1 WB54717 13187 UCSD052i-56-1 WB57717 13188 UCSD053i-57-1 WB55067 13189 UCSD054i-58-1 WB55461 13190 UCSD055i-59-1 WB54168 13191 UCSD056i-60-1 WB57571 13192 UCSD057i-61-1 WB55674 13193 UCSD058i-62-1 WB57057 13194 UCSD059i-63-1 WB63472 13195 UCSD060i-64-1 WB57102 13196 UCSD063i-20-1 WB62421 13197 WISCO15i-SC7 WB66708 13198 UCSD235i-SAD2-4 WB66703 13199

STAN053i-149-1 WB66707 13200 HVRDi002-A WB66709 13201 WISCO14i-SC1 WB66706 13202 CREM032i-SS48-1 WB66711 13203 UCSD207i-31-2 WB66716 13204

UCSD065i-20-3 WB60829 13205

UNIQUE IDENTIFIER:

NA

PRODUCT REGISTRATION:

Other: Human iPS cells

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

METHOD VALIDATION / PD #:

000053

STERIS Laboratories, Inc. 9303 West Broadway Ave Brooklyn Park, MN 55445

LAB-003 rev 30 Form 5 Effective: 2017-08-29 Page 1 of 2

Native Product Sterility Report



1	FQT	MET	HOD	\cap	OGY:	
ı	E01		пор	UL	UGT.	

USP - Direct Transfer

COMMENTS:

Sample # 17121502

REVIEWED BY Wessel

DATE 10JANI8

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

Testing Performed by WiCell Lot Release Testing January 25, 2018

FORM SOP-QU-004.01 Version G Edition 02 Reported by: AP Reviewed by: JB BD Monolight 180

		Read	ing A	A Reading B		В	Ratio			
#	Sample Name	RLU1	RLU2	Ave	RLU1	RLU2	Ave	B/A	Result	Comments/Suggestions
1	CREM032i-SS48-1-WB66711 13235	124	126	125	51	48	49.5	0.40	Negative	
2	Positive (+) Control	377	389	383	13283	13495	13389	34.96	Positive	
3	Negative (-) Control	607	619	613	64	63	63.5	0.10	Negative	

