

Thaw and Culture Details

Cell Line Name	CREM025i-SS37-1							
WiCell Lot Number	DB48040							
Provider	Boston University – Laboratory of Dr. Martin Steinberg							
Banked By	Boston University - Laboratory of Dr. Gustavo Mostoslavsky							
Thaw and Culture Recommendations	WiCell recommends thawing 1 vial into 1 wells of a 6 well plate. WiCell recommends thawing using ROCK Inhibitor for best results.							
Culture Platform Feeder Dependent								
	Medium: hESC Medium (KOSR)							
	Matrix: MEF							
Protocol WiCell Feeder Dependent Protocol								
Passage Number p6 These cells were cultured for 6 passages after colony picking prior to freeze. Add +1 to the panumber to best represent the overall passage number of the cells at thaw.								
Date Vialed 14-April-2016								
Vial Label	SS37-1p6 hiPSC/KSR 4/14/16 SMP							
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

Total grant and any trace								
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.

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



Approval Date	Quality Assurance Approval			
05-December-2016	9/27/2018 X JKG IKG Quality Assurance Signed by Gay, Jenna			



Chromosome Analysis Report: 072438

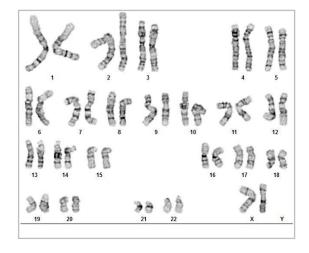
Date Reported: Tuesday, July 17, 2018

Cell Line: CREM025i-SS37-1-DB48040 13803

Passage#: 11

Date of Sample: 7/11/2018 Specimen: Human IPS

Results: 46,XX



Cell Line Sex: Female

Reason for Testing: Lot release testing

Investigator: WiCell

Cell: 37 Slide: G01

Slide Type: Karyotype

Total Counted: 20
Total Analyzed: 8

Total Karyogrammed: 4
Band Resolution: 500 - 550

Interpretation:

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

Completed by:

Reviewed and Interpreted by:

, CG(ASCP)

PhD, FACMG

A signed copy of this report is available upon request.

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 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:

13803-STR

Sample Name on Tube: 13803-STR

 $70.7 \text{ ng/}\mu\text{L}, (A260/280=1.75)$

Sample Type: Cells

Cell Count: ~2 million cells

Requestor:

WiCell Research Institute
Quality Department

Sample Date: N/A Receive Date: 07/09/18

Assay Date: 07/11/18

File Name: STR 180712 wmr

Report Date: 07/18/18

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	7-18 P								
D8S1179									
vWA									
Amelogenin	elogenin								
Penta_D									
CSF1PO	6-15	please, contact WiCell's Technical							
D16S539									
D7S820	820 6-14								
D13S317	7-15								
D5S818									
Penta_E	5-24								
D18S51	8-10, 10.2, 11-13, 13.2, 14-27								
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								
D3S1358	12-20								

<u>Results:</u> Based on the 13803-STR cells submitted by WiCell QA dated and received on 07/09/18, this sample (Label on Tube: 13803-STR) defines the STR profile of the human stem cell line CREM025i-SS37-1 comprising 27 allelic polymorphisms across the 15 STR loci analyzed.

<u>Interpretation:</u> No STR polymorphisms other than those corresponding to the human CREM025i-SS37-1 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 13803-STR sample submitted corresponds to the CREM025i-SS37-1 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 07/19/18

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Native Product Sterility Report



SAMPLE #: 17080685

DATE RECEIVED: 10-Aug-17

TEST INITIATED: 11-Aug-17

TEST COMPLETED: 25-Aug-17

SAMPLE NAME / DESCRIPTION: CREM032i-SS48-1 DB48070 12715

CREM019i-SS25-1 DB48022 12716
CREM020i-SS28-1 DB48025 12717
CREM021i-SS29-1 DB48028 12718
CREM022i-SS32-1 DB48031 12719
CREM024i-SS36-1 DB48037 12720
CREM025i-SS37-1 DB48040 12721
CREM026i-SS38-1 DB48043 12722
CREM027i-SS41-1 DB48055 12723
CREM028i-SS43-2 DB48058 12724

UNIQUE IDENTIFIER: NA

PRODUCT REGISTRATION: Other: Human iPS cells

TEST RESULTS:

WiCell

504 S Rosa Rd Rm 101

Madison, WI 53719

# Tested	# Positives (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

METHOD VALIDATION / PD #: 000053

TEST METHODOLOGY: USP - Filtration

COMMENTS: NA

REVIEWED BY _____

DATE 28 AU GUT

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 June 14, 2018

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

		Reading A		A	Read	ling B	В	Ratio		
#	Sample Name	RLU1	RLU2	Ave	RLU1	RLU2	Ave	B/A	Result	Comments/Suggestions
1	CREM025i-SS37-1-DB48040 13803	484	494	489	177	174	175.5	0.36	Negative	
2	Positive (+) Control	560	600	580	21715	21834	21775	37.54	Positive	
3	Negative (-) Control	910	902	560	91	87	89	0.16	Negative	

