

## **Thaw and Culture Details**

| Cell Line Name                      | CREM001i-bBU1C2   |  |
|-------------------------------------|---|--|
| WiCell Lot Number                   | DB47959   |  |
| Provider                            | Boston University – Laboratory of Dr. Martin Steinberg  |  |
| Banked By                           | Boston University - Laboratory of Dr. Gustavo Mostoslavsky  |  |
| Thaw and Culture<br>Recommendations | WiCell recommends thawing 1 vial into 2 wells of a 6 well plate.  |  |
| Culture Platform                    | Feeder Independent  |  |
|                                     | Medium: mTeSR™1   |  |
|                                     | Matrix: Matrigel®   |  |
| Protocol                            | WiCell Feeder Independent mTeSR1 Protocol   |  |
| Passage Number                      | p30 These cells were cultured for 30 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                         | 11-March-2015   |  |
| Vial Label                          | hiPSC<br>bBU1C2<br>P30/mTeSR<br>3-11-15 AS  |  |
| 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 Method                             | Test Specification                   | Result |
|-----------------------------------|---|---|--------------------------------------|--------|
| Karyotype by G-banding            | WiCell  | SOP-CH-003                              | Expected karyotype                   | Pass   |
| Post-Thaw Viable Cell<br>Recovery | WiCell  | SOP-CH-305                              | Recoverable attachment after passage | Pass   |
| Identity by STR                   | UW Translational<br>Research Initiatives in<br>Pathology Laboratory | PowerPlex 16 HS<br>System by<br>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                                    |
|------------------|---|
| 02-December-2016 | 11/30/2017  X JKG  RG  Quality Assurance Signed by Gay, Jenna |



### Chromosome Analysis Report: 069184

| Date Repo | orted: Thursd  | ay, November 16    | Cell Li  | ne Gender: | Male |
|-----------|----------------|--------------------|----------|------------|------|
| Date Nept | orteu. Triursu | ay, itovellibel io | , Och Li | ne dender. | maic |

2017

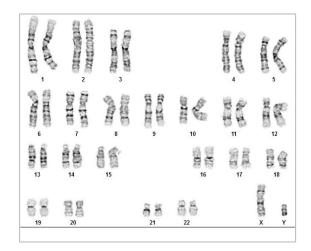
Cell Line: CREM001i-bBU1C2-DB47959

13025

Passage#: 32

Date of Sample: 11/9/2017 Specimen: Human IPSC

Results: 46,XY



Reason for Testing: lot release testing

Investigator: WiCell CDM

> Cell: 41 Slide: G01

Slide Type: Karyotype

Total Counted: 20 Total Analyzed: 8 Total Karyogrammed: 4 Band Resolution: 450 - 550

#### Interpretation:

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

| Completed by:                | CG(ASCP)     |
|------------------------------|--------------|
| Reviewed and Interpreted by: | , PhD, FACMO |

A signed copy of this report is available upon request.

| 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 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.

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# **Short Tandem Repeat Analysis**

**HISTOLOGY - IHC - MOLECULAR - IMAGING** 

Department of Pathology and Laboratory Medicine TRIP Laboratory (Molecular)

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

info@wicell.org (888) 204-1782

**Sample Report:** 

13025-STR

Sample Name on Tube: 13025-STR

71.2 ng/ $\mu$ L, (A260/280=2.07)

Sample Type: Cells

Cell Count: ~2 million cells

**Requestor:** 

WiCell Research Institute

Quality Department

Sample Date: N/A **Receive Date:** 11/13/17 **Assay Date:** 11/16/17

File Name: STR 171120 wmr

**Report Date:** 11/21/17

| 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                          |
| TPOX       | 6-13  | information has                      |
| D8S1179    | 7-18  | been redacted to                     |
| vWA        | 10-22   | protect donor                        |
| Amelogenin | X,Y   | confidentiality. If 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                   |
| D7S820     | 6-14  | Support.                             |
| D13S317    | 7-15  |                                      |
| D5S818     | 7-16  |                                      |
| 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   |                                      |

Results: Based on the 13025-STR cells submitted by WiCell QA dated and received on 11/13/17, this sample (Label on Tube: 13025-STR) defines the STR profile of the human stem cell line CREM001i-bBU1C2 comprising 26 allelic polymorphisms across the 15 STR loci analyzed.

Interpretation: No STR polymorphisms other than those corresponding to the human CREM001i-bBU1C2 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 13025-STR sample submitted corresponds to the CREM001i-bBU1C2 stem cell line and was not contaminated with any other human stem 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 stem cell lines is  $\sim 2-5\%$ .

X WMR  $\mathbf{X}$  RMB **Digitally Signed on** 11/22/17 **Digitally Signed on** 11/22/17 PhD, Director / Co-Director TRIP Laboratory, Molecular UWHC Molecular Diagnostics Laboratory / UWSMPH TRIP Laboratory

## Native Product Sterility Report



SAMPLE #:

17050657

DATE RECEIVED:

04-May-17

TEST INITIATED:

08-May-17

TEST COMPLETED:

22-May-17

SAMPLE NAME / DESCRIPTION:

H9-SOX2-GFP DB47477 12513

CREM001i-bBU1C2 DB47959 12514
CREM002i-BU2C10 DB47962 12515
CREM003i-BU3C2 DB47974 12516
CREM004i-SS2-1 DB47977 12517
CREM006i-SS4-1 DB47980 12518
CREM007i-SS5-1 DB47983 12519
CREM008i-SS6-1 DB47988 12520
CREM009i-SS8-2 DB47991 12521
CREM018i-SS24-1 DB48019 12522

**UNIQUE IDENTIFIER:** 

NA

PRODUCT REGISTRATION:

Human iPS cells

**TEST RESULTS:** 

WiCell

504 S Rosa Rd, Rm 101

Madison, WI 53719

| # Tested | # Positives<br>(Growth) | - Control   |
|----------|-------------------------|-------------|
| 10       | 0                       | 2 Negatives |

TEST SUMMARY:

| # Samples | Media Type | Volume (mL) | Incubation<br>Temperature<br>(° C) | Incubation<br>Duration<br>(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 - Direct Transfer

COMMENTS:

NA

# Native Product Sterility Report



|             | $\mathcal{C}$ |              |
|-------------|---------------|--------------|
| REVIEWED BY |               | DATE 24MAYI) |

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.

STERIS Laboratories, Inc. 9303 West Broadway Ave Brooklyn Park, MN 55445 LAB-003 rev 29 Form 5 Effective: 2017-04-20 Page 2 of 2



## Mycoplasma Assay Report

PCR-based assay performed by WiCell WiCell Stem Cell Bank, WiCell 20Oct23

Form SOP-83.01 Version 5.0

| Sample Name                         | Result   | Interpretation  |
|-------------------------------------|----------|---|
| CREM001i-bBU1C2-DB47959 p34 (99198) | 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<sup>TM</sup> Mycoplasma Detection Kit (Sartorius).

|  | 10/20/2023 | 10/25/2023   | 10/25/2023   |
|--|------------|--|--|
| X Justin Hobson  |            | X Kaylie Petersen  | X Andy Arntz   |
| Tech #1<br>Characterization<br>Signed by: Hobson, Justin |            | Tech #2<br>Characterization<br>Signed by: Petersen, Kaylie | QA Review<br>Quality Assurance<br>Signed by: Arntz, Andy |

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A gel image is available upon request.