



## Thaw and Culture Details

Cell Line Name	<b>STAN099i-108C2</b>
WiCell Lot Number	<b>DB44602</b>
Provider	Stanford University – Laboratory of Dr. Thomas Quettermous
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. WiCell recommends thawing using ROCK Inhibitor for best results.
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 Vialied	15-October-2015
Vial Label	ISMMS 108i C2P11 AP 101515
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	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 STR profile of deposited cell line	Pass
Sterility	Steris	ST/07	Negative	Pass
Mycoplasma	WiCell	SOP-CH-044	Negative	Pass




## Testing Reported by Provider

Test Description	Method	Result
Mycoplasma	Lonza MycoAlert kit	Negative

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<sup>EX</sup>)

Approval Date	Quality Assurance Approval
27-October-2016	<div style="text-align: right;">1/16/2020</div> <div style="text-align: center;"> JKG Quality Assurance Signed by Gay, Jenna</div>

**Date Reported:** Monday, November 19, 2018

**Cell Line:** STAN099i-108C2-DB44602 14102

**Passage#:** 13

**Date of Sample:** 11/12/2018

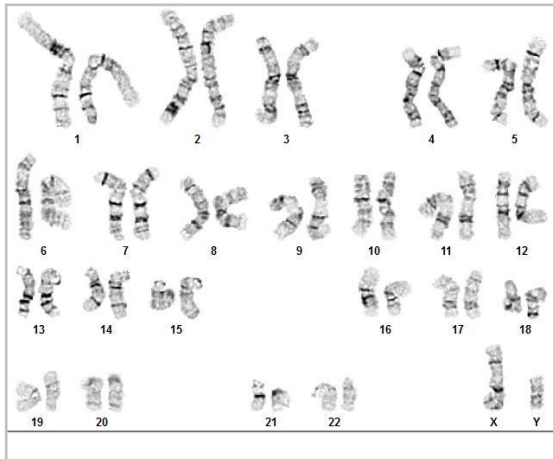
**Specimen:** Human IPS

**Results:** 46,XY

**Cell Line Sex:** Male

**Reason for Testing:** Lot Release Testing

**Investigator:** [REDACTED], WiCell



**Cell:** 3

**Slide:** G01

**Slide Type:** Karyotype

**Total Counted:** 20

**Total Analyzed:** 8

**Total Karyogrammed:** 4

**Band Resolution:** 475 - 550

### Interpretation:

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

**Completed by:** [REDACTED]

**Reviewed and Interpreted by:** [REDACTED], PhD, FACMG

**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](http://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.*



HISTOLOGY - IHC - MOLECULAR - IMAGING

Department of Pathology and Laboratory Medicine  
TRIP Laboratory (Molecular)  
<https://research.pathology.wisc.edu/trip-home/>  
(608) 265-9168

# Short Tandem Repeat Analysis



Your Lab Partner

characterization@wicell.org  
(608) 316-4145

**Sample Report:**

15200-STR

**Sample Name on Tube:** 15200-STR

98.1 ng/μL, (A260/280=1.75)

**Sample Type:** Cells

**Cell Count:** ~2 million cells

**Requestor:**

WiCell Research Institute

Quality Assurance Department

**Receive Date:** 12/19/19

**Report Sent:** 01/09/20

**Assay Date:** 01/07/20

**File Name:** STR 1200108 wmr

**Report Date:** 01/09/20

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 been redacted to protect donor confidentiality. If more information is required, please, contact <a href="#">WiCell's Technical Support</a> .
TPOX	6-13	
D8S1179	7-18	
vWA	10-22	
Amelogenin	X,Y	
Penta_D	2.2, 3.2, 5, 7-17	
CSF1PO	6-15	
D16S539	5, 8-15	
D7S820	6-14	
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 15200-STR cells submitted by WiCell QA dated and received on 12/19/19, this sample (Label on Tube: 15200-STR) defines the STR profile of the human cell line STAN099i-108C2 comprising 26 allelic polymorphisms across the 15 STR loci analyzed.

**Interpretation:** No STR polymorphisms other than those corresponding to the human STAN099i-108C2 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 15200-STR sample submitted corresponds to the STAN099i-108C2 cell line and was 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 ~2-5%.

X *RMB*

Digitally Signed on 01/09/20

X *WMR*

Digitally Signed on 01/09/20

██████████, BA  
TRIP Laboratory, Molecular

██████████, PhD, Director / Co-Director  
UWHC Molecular Diagnostics Laboratory / UWSPH TRIP Laboratory

Testing was accomplished by analysis of human genetic polymorphisms at STR loci. This methodology has not yet been approved by the FDA and is for investigational use only.

**Acknowledge TRIP in your publications, posters & presentations. For details, see: <https://research.pathology.wisc.edu/acknowledging-trip/>**

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# Native Product Sterility Report



WiCell  
504 S Rosa Rd , Rm 101  
Madison, WI 53719

SAMPLE #: 18111110  
DATE RECEIVED: 15-Nov-18  
TEST INITIATED: 26-Nov-18  
TEST COMPLETED: 10-Dec-18

SAMPLE NAME / DESCRIPTION: LUEL8357i-3 WB66939 14103  
LUEL8679i-4 WB66940 14104  
STAN100i-108C4 DB44605 14105  
STAN099i-108C2 DB44602 14106  
STAN207i-459C2 DB35961 14107  
STAN206i-459C1 DB35958 14108  
STAN216i-496C1 DB35535 14109  
LUEL7159i-7 WB66914 14110  
EFNB2-tdTomato/EPHB4-EGFP DB66613 14116  
JHU012i-2 DB36196 14117

UNIQUE IDENTIFIER: NA  
PRODUCT REGISTRATION: Other: Human iPS cells

TEST RESULTS:

# Tested	# Positives (Growth)	- Control
10	1	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 - Direct Transfer

COMMENTS: Sample labeled LUEL7159i-7 WB66914 14110 is positive in TSB and FTG.

REVIEWED BY

DATE

20 DEC 18

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 Assay Report

PCR-based assay performed by WiCell

WiCell

12Dec19

FORM SOP-CH-048.01

Version B Edition 01

Sample Name	Result	Comments/Suggestions
WC068i-310-17-2-36-DB67343 15198 (79400)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
WC065i-247-1-2-32-DB67338 15197 (79401)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
WC071i-335-1-2-35-DB67346 15199 (79402)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
STAN099i-108C2-DB44602 15200 (79403)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
MCW085i-40002118-WB67193 15109 (79405)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
MCW082i-U2052-WB67222 15108 (79406)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
MCW095i-U2311-WB67185 15160 (79407)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
MCW103i-40000237-WB67208 15158 (79408)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
MCW096i-40000169-WB67199 15161 (79409)	Negative	Band was not seen at 270bp, indicating the absence of mycoplasma.
Positive (+) Control	Positive	
Negative (-) Control	Negative	

**Reported by: Alex Paguirigan, Assistant Cell Culture Specialist**

**Reviewed by: Katie Remondini, Cell Culture Specialist**

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*A gel image is available upon request.*