



## Thaw and Culture Details

Cell Line Name	<b>AI06e-SOX2YFP</b>
WiCell Lot Number	<b>DB66691</b>
Provider	Allen Institute – Dr. Boaz Levi
Banked By	Allen Institute – Dr. Boaz Levi
Thaw and Culture Recommendations	WiCell recommends thawing 1 vial into 5 wells of a 6 well plate.
Culture Platform	Feeder Independent
	Medium: mTeSR™1
	Matrix: Matrigel®
Protocol	WiCell Feeder Independent mTeSR™1 Protocol
Passage Number	p65 These cells were cultured for 64 passages prior to freeze. Cells were modified at passage 40. The provider adds +1 to the passage number to best represent the overall passage number of the cells at thaw.
Date Viald	01-July-2014
Vial Label	SOX2 600 #6 H1 p65 July 1, 2014
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	Consistent with STR profile of deposited cell line	Pass
Sterility	Steris	ST/07	Negative	Pass
Mycoplasma	WiCell	SOP-QU-004	Negative	Pass
Expression of Reporter Proteins	WiCell	SOP-CH-032	Expression of reporter proteins reported	Pass

## Testing Reported by Provider

The provider has stated the following testing and results were performed for this cell line. A link to the relevant publication is provided on the cell line specific web page on the WiCell website.

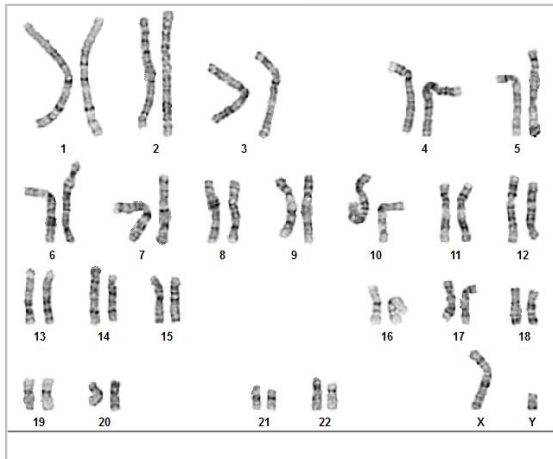
Test Description	Result
Karyotype by G-banding	Normal karyotype
Mycoplasma	Negative
Sterility	Negative
Expression of reporter protein	Pass



Approval Date	Quality Assurance Approval
25-January-2018	<p style="text-align: right; margin-right: 20px;"><small>5/27/2020</small></p> <p>X HEB</p> <p><small>HEB Quality Assurance Signed by: Bruner, Haley</small></p>

**Date Reported:** Monday, March 05, 2018  
**Cell Line:** AI06e-SOX2YFP-DB66691 13481  
**Passage#:** 65  
**Date of Sample:** 2/28/2018  
**Specimen:** Human ES  
**Results:** 46,XY

**Cell Line Gender:** Male  
**Reason for Testing:** lot release testing  
**Investigator:** [REDACTED], WiCell



**Cell:** 71  
**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:** [REDACTED], CG(ASCP)  
**Reviewed and Interpreted by:** [REDACTED], PhD, FACMG  
**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.*

*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.*

**Sample Report:**

13481-STR

**Sample Name on Tube:** 13481-STR

73.5 ng/μL, (A260/280=1.83)

**Sample Type:** Cells**Cell Count:** ~2 million cells**Requestor:**

WiCell Research Institute

Quality Department

**Sample Date:** N/A**Receive Date:** 03/12/18**Assay Date:** 03/13/18**File Name:** STR 180314 wmr**Report Date:** 03/16/18

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	20,24
TPOX	6-13	8,11
D8S1179	7-18	12,13
vWA	10-22	15,17
Amelogenin	X,Y	X,Y
Penta_D	2.2, 3.2, 5, 7-17	10,13
CSF1PO	6-15	12,13
D16S539	5, 8-15	9,13
D7S820	6-14	8,12
D13S317	7-15	8,11
D5S818	7-16	9,11
Penta_E	5-24	10,12
D18S51	8-10, 10.2, 11-13, 13.2, 14-27	17,18
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	28,32.2
TH01	4-9,9.3,10-11,13.3	9.3,9.3
D3S1358	12-20	15,15

**Results:** Based on the 13481-STR cells submitted by WiCell QA dated and received on 03/12/18, this sample (Label on Tube: 13481-STR) exactly matches the STR profile of the human stem cell line WA01 comprising 28 allelic polymorphisms across the 15 STR loci analyzed.

**Interpretation:** No STR polymorphisms other than those corresponding to the human WA01 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 13481-STR sample submitted corresponds to the WA01 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 ~2-5%.



Digitally Signed on 03/21/18

[Redacted], BA  
TRIP Laboratory, Molecular

Digitally Signed on 03/21/18

[Redacted], 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: <http://www.pathology.wisc.edu/research/trip/acknowledging>

TRIP agrees to maintain the confidentiality of any information provided to it in connection with its performance of this STR analysis on the same conditions as set forth in paragraph 2 of WiCell's Terms and Conditions of Service (<http://www.wicell.org/media.acux/1a429b84-2b54-44a4-8ad8-5c05db93dd8a>).

# Native Product Sterility Report



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

SAMPLE #: 18021403  
DATE RECEIVED: 22-Feb-18  
TEST INITIATED: 23-Feb-18  
TEST COMPLETED: 09-Mar-18

SAMPLE NAME / DESCRIPTION: AI03e-DCXYFP DB66690 13456  
AI06e-SOX2YFP DB66691 13457  
AI07e-Timothy DB66692 13458  
AI08e-PAX6YFP DB66693 13459  
AI09e-KCTD13a DB66694 13460  
AI10e-KCTD13b DB66695 13461  
AI11e-OTX2YFP DB66696 13462  
AI12e-HOPX-CIT+/- DB66697 13463  
AI13e-HOPX-CIT+/+ DB66698 13464  
CREM022i-SS32-1 WB66732 13466  
iPS(IMR90)-1 WB66731 13467  
STAN004i-147-1 DB31065 13468  
STAN005i-147-2 DB31088 13469  
STAN024i-29-1 DB30891 13470  
STAN025i-29-2 DB30897 13471  
WC034i-SOD1-D90A WB66734 13472  
WC035i-SOD1-D90D WB66733 13473  
WISC015i-SC7 WB66735 13474  
WC008i-C603-4 WB66741 13475  
WC034i-SOD1-D90A WB66740 13484

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

### TEST RESULTS:

# Tested	# Positives (Growth)	- Control
20	0	3 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

# Native Product Sterility Report



TEST METHODOLOGY: USP - Direct Transfer

COMMENTS: NA  
# 18021403

REVIEWED BY *Densad*

DATE 22 MAR 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 Detection Assay Report

Testing Performed by WiCell

Lot Release Testing

March 1, 2018

FORM SOP-QU-004.01

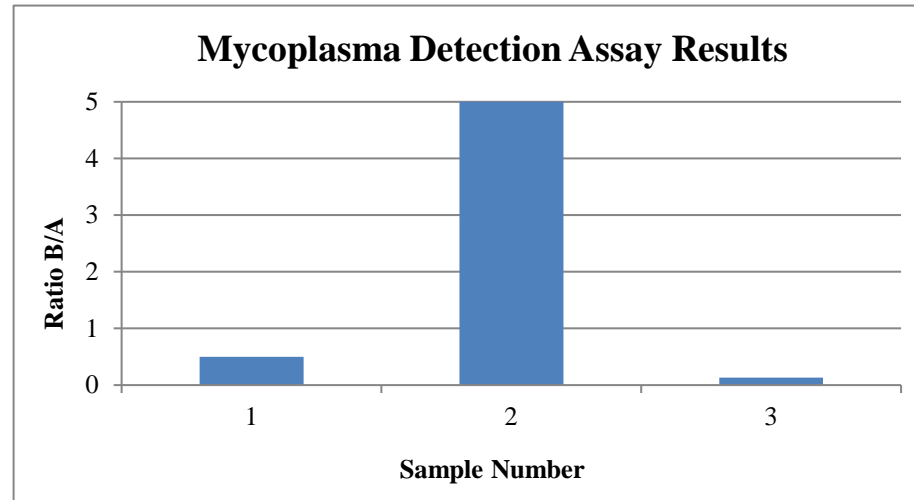
Version G Edition 02

Reported by: AP

Reviewed by: JB

BD Monolight 180

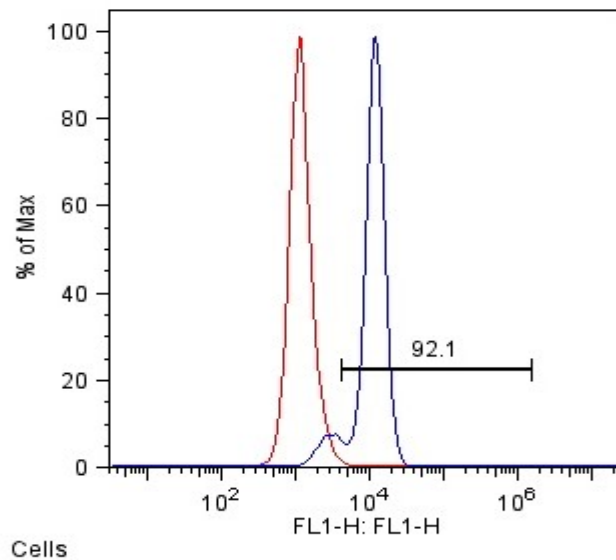
#	Sample Name	Reading A			Reading B			Ratio B/A	Result	Comments/Suggestions
		RLU1	RLU2	Ave	RLU1	RLU2	Ave			
1	A106e-SOX2YFP-DB66691 13481	255	264	259.5	130	128	129	0.50	Negative	
2	Positive (+) Control	388	400	394	34112	34379	34246	86.92	Positive	
3	Negative (-) Control	706	732	719	101	92	96.5	0.13	Negative	



Cell Line-Lot Number	AI06e-SOX2YFP-DB66691
Sample ID	13481
Passage Number	65
Reported By/Date	SM 13Mar18
QA Review By/Date	RK 26Mar18
Percent Positive for Reporter Protein	92.1
Deviations from Procedure	<input checked="" type="checkbox"/> N/A
Notes	<input checked="" type="checkbox"/> N/A

**Histogram Plot Indicating Positive Percentage of the Reporting Gene**

Red peak is negative control population. Blue peak is test population.



<span style="color: blue;">█</span>	A01 AI06e-SOX2YFP-DB66691.fcs	53.3
<span style="color: red;">█</span>	A01 WA01-DL-12-T1042530.fcs	76