



Certificate of analysis

SFC865-03-07

Operator: Olga Perestenko

Date: 25/01/2017

Supervisor: Sally Cowley

Date: 25/02/2018

Signature:

A handwritten signature in black ink, appearing to read "SACowley", is written over a light grey rectangular background.

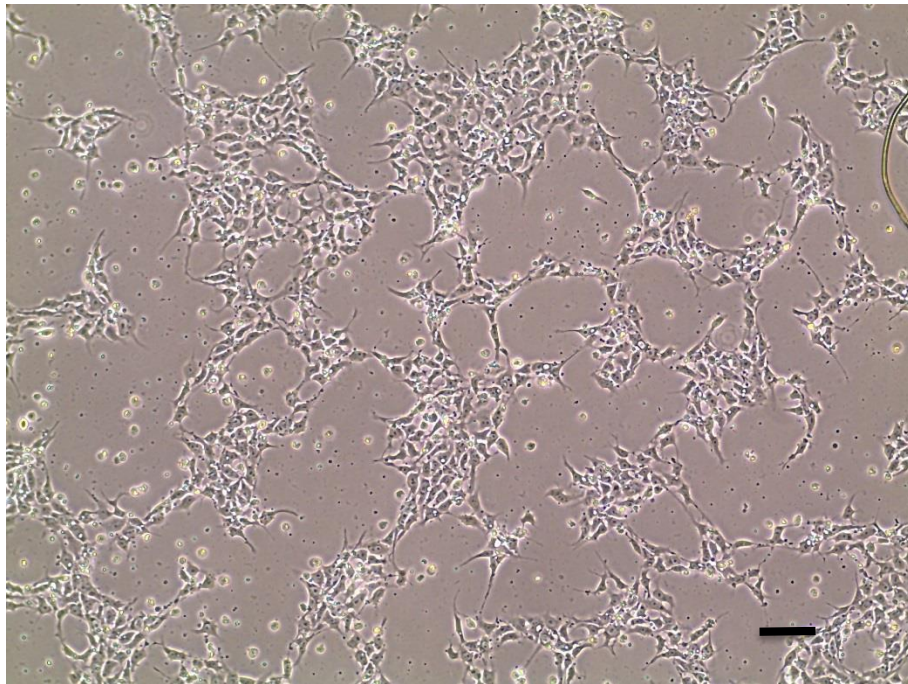
Source of fibroblasts and reprogramming information

- SF865 from UOXF
- Reprogrammed at UOXF-S
- Reprogrammed on 06/10/2016 at passage 4 OP
- Cytotune v2 WP3 SOP10

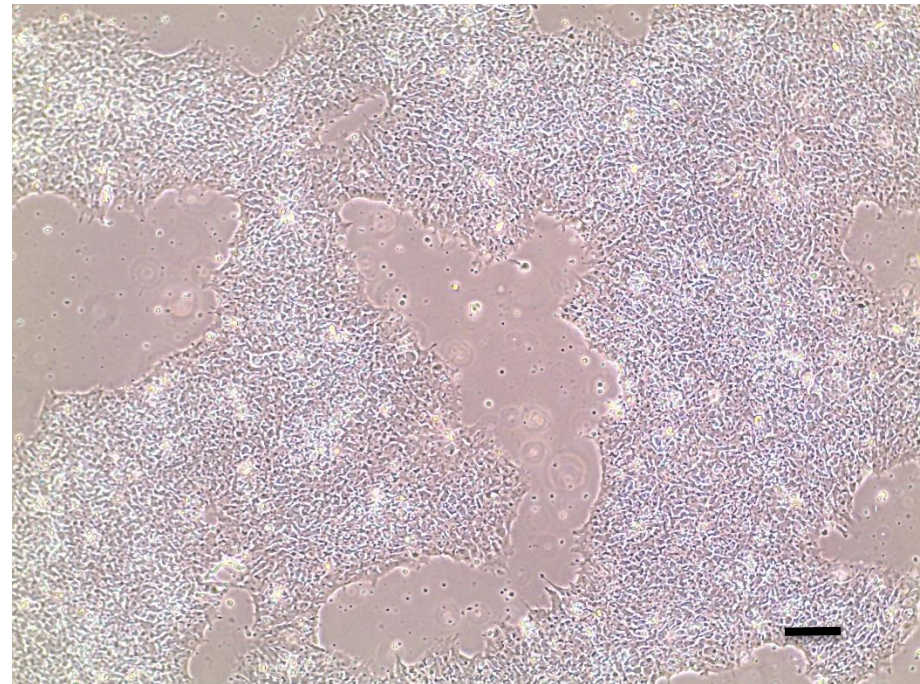
Viability post-thaw and Morphology according to SOP19 passage 10

- Cell count immediately post-thaw 1.25×10^6
- Viability immediately post-thaw 90.9%
- Photo at 24h and day 3 post-thaw (scale bar = $100\mu\text{m}$):

24h post-thaw 80% plated



Day 3 post-thaw 20% plated



Sendai clearance: according to WP3 SOP15

Faint residual myc virus detectable at passage 10 – recommend
passaging further and testing again at p13+

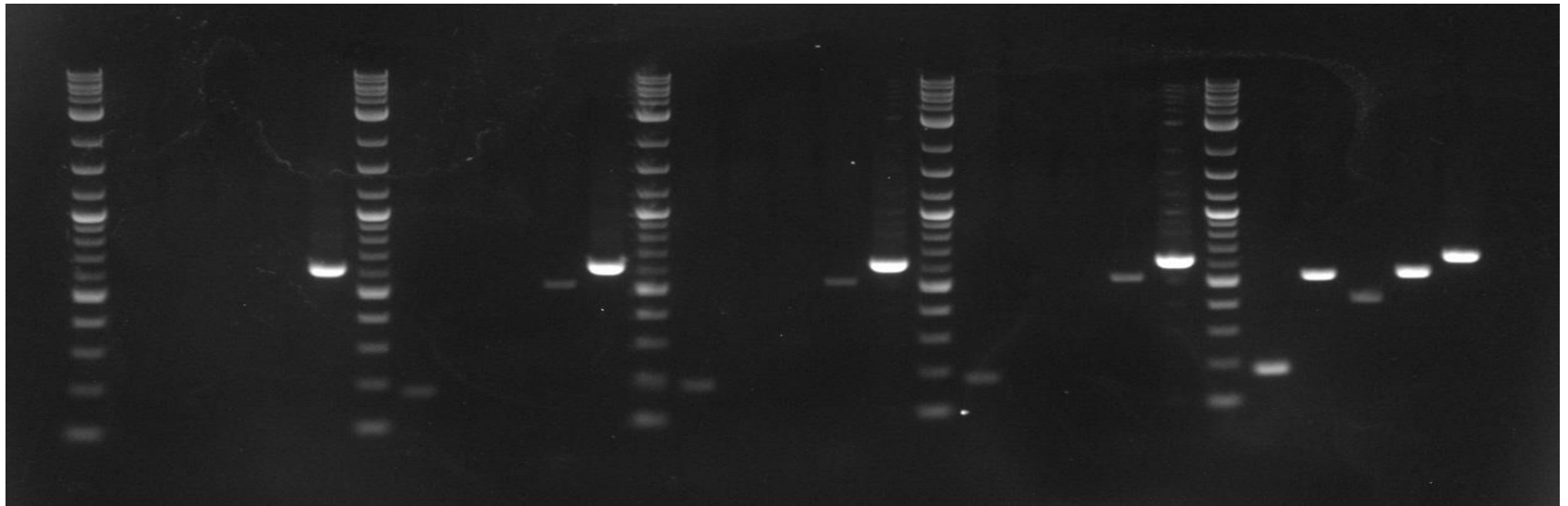
2Log ladder
Sendai virus
KOS
Klf4
C-Myc
Actin

2Log ladder
Sendai virus
KOS
Klf4
C-Myc
Actin

2Log ladder
Sendai virus
KOS
Klf4
C-Myc
Actin

2Log ladder
Sendai virus
KOS
Klf4
C-Myc
Actin

2Log ladder
Sendai virus
KOS
Klf4
C-Myc
Actin



SFCxxx-xx-xx

SFC865-03-07

SFCxxx-xx-xx

SFCxxx-xx-xx

+ control

Product sizes: SeV 181bp; KOS 528bp; SeV-Klf 410bp; SeV-Myc 532bp; Actin 623bp

Mycoplasma Test:

According to MycoAlert Lonza LT07-318 undetectable at passage 10

Sample	Clone	Passage number	Initial	Reading 1	Reading 2	Ratio/Status
+ve control				5.428	153.9	28.35
-ve control				3.384	0.385	0.11
	SFC865-03-07	p10	OP	0.974	0.463	0.48

Results mean

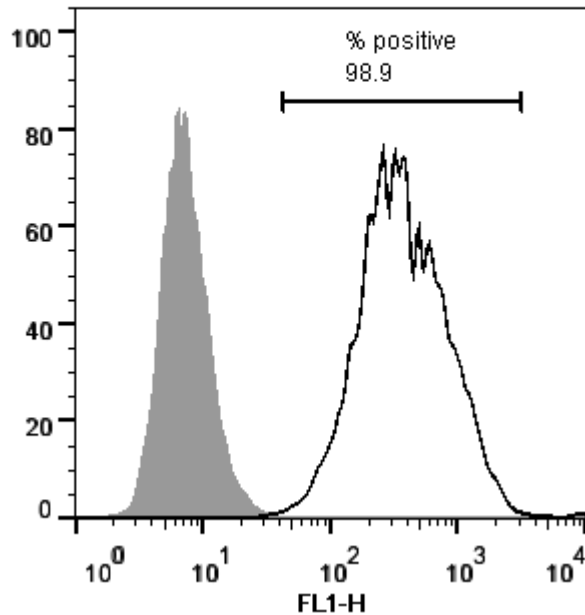
Ratio **0 - 0.999** negative for mycoplasma

Ratio **1 – 1.3** Borderline Result (retest required)

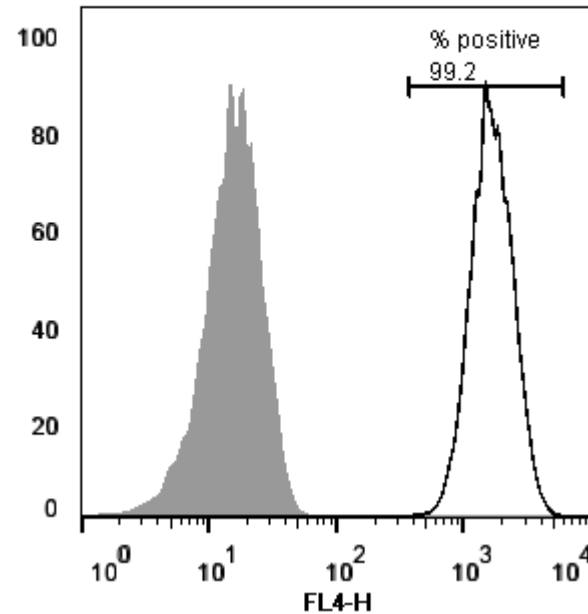
Ratio above **1.3** positive for mycoplasma

Flow cytometric analysis according to WP3 SOP 20 and 21 passage p11

Tra-1-60:



NANOG:



SNP analysis

according to WP3 SOP Preparation of DNA and RNA samples for Illumina arrays

- Passage 10
- Identity to parent fibroblasts confirmed
- Karyotype abnormalities: none detected
- For details and raw data see StemDB