

## Characterization of hPSC

<b>Cell Line Name</b>	NCRM5AS-iCAGcGFP.c9		
<b>Type of Cell Line</b>	hiPSC		
<b>Depositor (Institution)</b>	NIH Center for Regenerative Medicine (US)		
<b>Passage #</b>	p73*		
<b>Day of Cell Freezing</b>	20181004		
Analysis	Result	Passage #	Day of analysis
Cell viability	Pass (67.33±0.05 %)	p74	20181017
Authentication (STR)	pass	p74	20181030
Mycoplasma test (PCR)	Pass	p74	20181031
Cell attachment and colony morphology	Pass	p70	20180911
Karyotype (G-banding)	46,XY	p73	20181016
Stem Cell Marker Expression			
· AP staining	Pass	p71	20181018
· ICC	Pass	p71	20181018
· qRT-PCR	Pass	p71	20181011
Differentiation Marker Expression			
· EB formation, GFP expression	pass	p73	20181004
· qRT-PCR	pass	p74	20181026

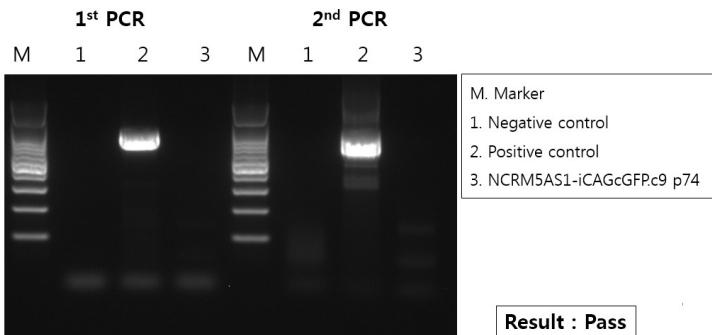
### Cell Culture Condition

- Feeder/matrix Vitronectin (Gibco, A14700)
  - Media TeSR-E8 (Stem Cell Technol, ST05940)
  - Passage method EZPassage(StemPro, 23181-010)

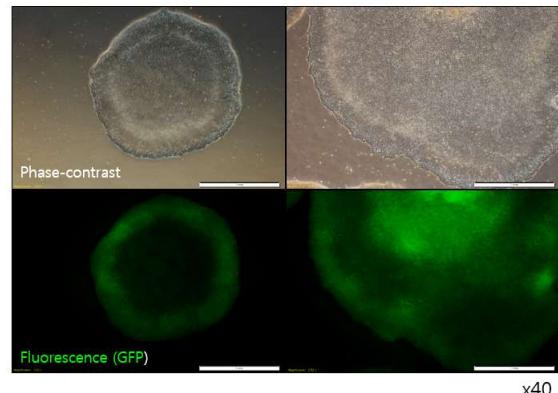
## Description by depositor

- |                        |   |
|------------------------|---|
| - Parental Cell        | NCRM-5 (NCRM Cat#CR0000013)<br>(CD34+ cord blood ; episomal plasmid)  |
| - Genetic modification | GFP-report tagged; TALEN<br><br>Luo et al., 2014. Stem Cell Translational Med (3) 821-835.  |
| - Reference            | Stable enhanced green fluorescent protein expression after differentiation and transplantation of reporter human induced pluripotent stem cells generated by AAVS1 transcription activator-like effector nucleases. |

## Mycoplasma contamination test



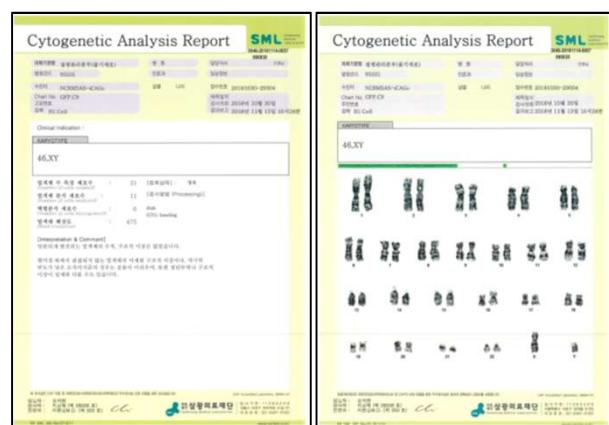
## Cell morphology



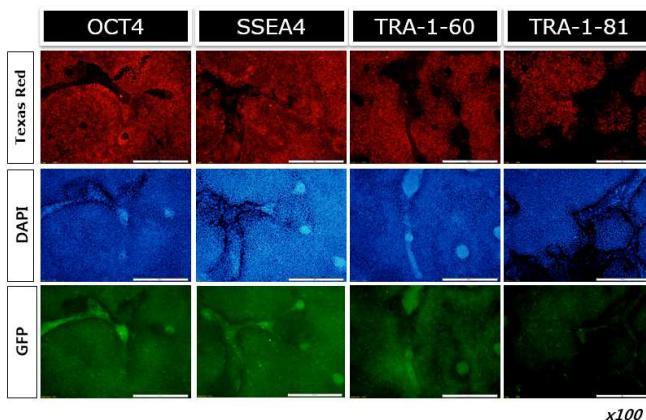
## Microbial contamination test



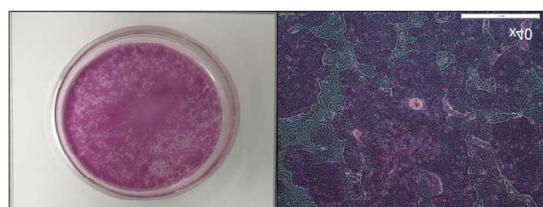
## Karyotype



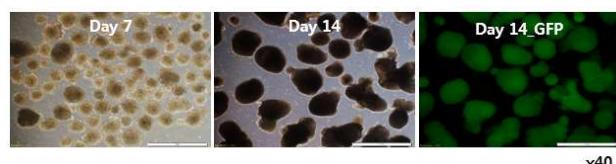
## Stem cell marker gene and GFP expression



## AP staining



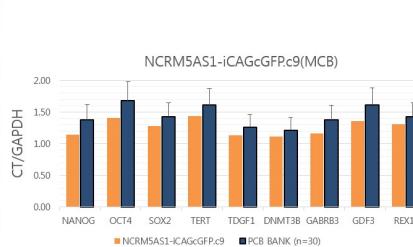
## EB formation



## Differentiation marker gene expression (EB14d)

### <qRT-PCR>

Gene	Ct Mean
NANOG	21.47
OCT4	26.31
SOX2	23.91
TERT	26.96
TDGF1	21.23
DNMT3B	20.89
GABRB3	21.70
GDF3	25.53
REX1	24.51
GAPDH	18.73



Gene	Ct Mean
PAX6	25.29
SOX1	31.75
HNF3B	23.75
AFP	22.54
T	27.92
MYOG	32.83
ACTB	18.33
GAPDH	19.30

### <qRT-PCR>

