

# Characterization of hPSC

<b>Cell Line Name</b>	<b>NU01-EiPS07</b>		
<b>Type of Cell Line</b>	<b>hiPSC</b>		
<b>Depositor (Institution)</b>	<b>Korea National Institute of Health</b>		
<b>Passage #</b>	<b>p15*</b>		
<b>Day of Cell Freezing</b>	<b>20190910*</b>		
<b>Analysis</b>	<b>Result</b>	<b>Passage #</b>	<b>Day of analysis</b>
Authentication (STR)	Pass	p12	20190920
Mycoplasma test (PCR)	Pass	p11	20190920
Cell attachment and colony morphology	Pass	p11	20190805
Microbial contamination test (Virus, Fungi, bacteria)	Pass	p12	20191002
Karyotype (G-banding)	46,XY	p7	20190730
HLA genotype	HLA-A *11:01:01G *31:01:02G HLA-B *39:01:01G *54:01:01 HLA-DRB1 *04:05:01G *08:03:02G	p12	20190927
ABO genotype	OO	p12	20191001
CNV analysis (CNV_Chip)	CNV calls(2) - Gain : 16p13.13 - Loss: 15q25.3	p12	20191127
Stem Cell Marker Expression			
· AP staining	Pass	p11	20190905
· ICC	Pass	p12	20190923
· qRT-PCR	Pass	p11	20191021
Differentiation Marker Expression			
· EB formation	Pass (EB14d)	p12	20190910
· qRT-PCR	Pass	p12	20191021

\* Freezing media : Stem Cell Banker (Amsbio, 11897)

## Cell Culture Condition

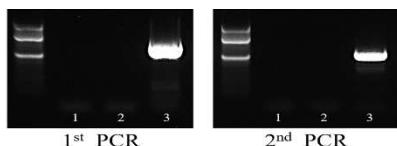
- Feeder /matrix · Vitronectin (Gibco, A14700)
- Media · TeSR-E8 Basal Medium(Stem Cell Technol, #05990)
- Passage (Cell dissociation) · EDTA/Gentle Cell Dissociation Reagent (Stem cell Technol, 07174)

## Description of the hPSC

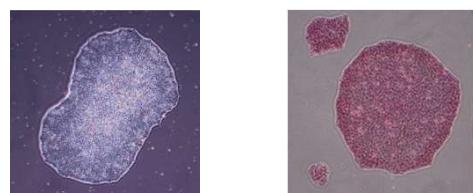
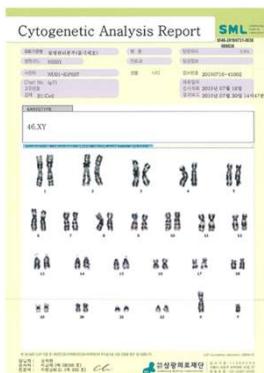
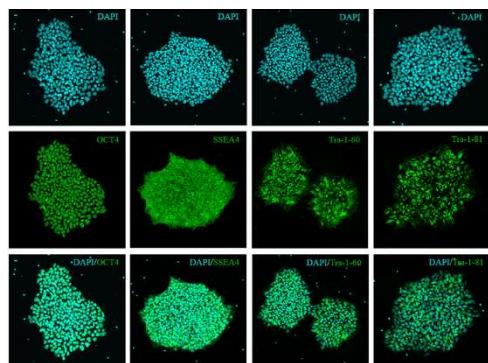
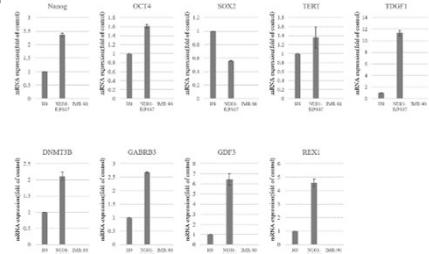
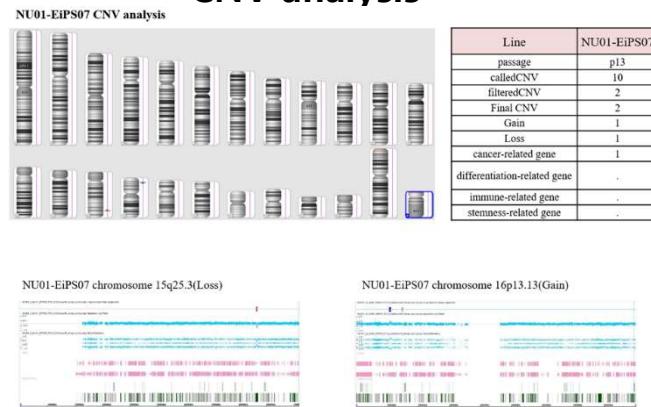
- Parental Cell · Urine derived cell (NU01)
- Reprogram · Episomal Vector (SOP#202.1)
- Epi5 Episomal iPSC Reprogramming Kit (Thermo Fisher, A15960)
- Oct4, Sox2, Klf4, c-Myc, Lin28, p53DD

## Reference

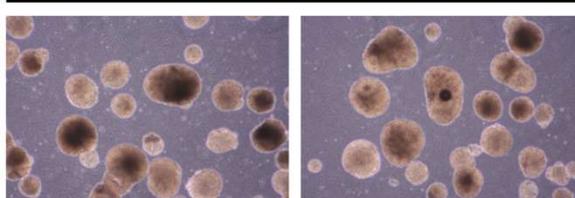
Im YS et al. Generation of integration-free induced pluripotent stem cell line (KSCBi012-A) from urinary epithelial cells of a healthy male individual. Stem Cell Res. 2022 Aug;63:102841.

**Mycoplasma contamination test**

1. Negative control  
2. NU01-EiPS07(TeSR-E8 + Vitronectin, p12)  
3. Positive control

**Cell Morphology AP staining****Microbial contamination test****Karyotype****Stem cell marker gene expression <ICC>****<qRT-PCR>****CNV analysis****Differentiation marker gene expression****<EB formation>**

EB Day 14

**<qRT-PCR, EB 14d>**

Gene	Ct mean
ACTB	24.2
PAX6	29.1
SOX1	34.9
HNF3B	not detected
AFP	35.4
T	35.3
MYOG	33.75

