Tbx3 and Nr5 α 2 Play Important Roles in Pig Pluripotent Stem Cells

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Abstract Pigs are valuable animal models in pre-clinical research due to their anatomical and similarity to human-beings. Little is known about porcine embryonic development and porcine pluripotent stem cells. Recently, porcine-induced pluripotent stem cells (piPSCs) have been generated with Oct4 (Pou5f1), Sox2, Klf4 and c-Mvc (termed OSKM, 4 F). Here, we found two other factors (*Tbx3* and *Nr5\alpha2*, termed TN), with important roles in piPSCs induction. They could improve the generation of piPSCs by supplementing these two factors on the basis of OSKM (OSKMTN, 6 F) orientated to mouse ESCs-like. Surprisingly, $Nr5\alpha 2$ alone could induce piPSCs formation in the presence or absence of *c-Myc*. These results suggested that *Tbx3* and *Nr5\alpha2* may have vital roles in Sus scrofa and proposed new insights into pig pluripotent stem cells.

Keywords piPSCs \cdot Nr5 α 2 \cdot Tbx3 \cdot Pluripotency \cdot Efficiency

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Introduction

Pigs are ideal animal models for a variety of diseases. However, porcine embryonic stem cells (pESCs) have not so far been successfully established, thus impeding the gene target research. Induced pluripotent stem cells were established by infecting mouse embryonic fibroblasts with retroviral vectors carrying four pluripotent genes Oct4, Sox2, c-Myc, and Klf4 [1]. Subsequently, a number of groups confirmed the reliability of this finding, and improved the quality of the mouse-induced pluripotent stem cells (miPSCs) [2, 3]. In 2009, pig-induced pluripotent stem cells (piPSCs) were established simultaneously by three groups with Yamanaka's classic four-factor OSKM approach, and these cells have been proved as having pluripotency [4–6]. As far as we know, there exist many defects in piPSCs. For example, the pluripotency maintenance of piPSCs needs the expression of exogenous genes. Some groups claimed that piPSCs-derived chimeric offspring have been obtained; however, this still needs to be further confirmed [7, 8]. The developmental efficiency of piPSCs nuclear transfer (NT)-derived embryos was lower than that of porcine embryonic fibroblasts (PEFs) NTderived ones both in vitro and in vivo, but this was not the case in mouse-induced pluripotent stem cells (miPSCs) [9, 10]. So, this suggested that the existed piPSCs may not be the true pluripotent stem cells. Therefore, factors and the medium which is used in piPSCs need to be further optimized.

The liver receptor homolog-1(*Lrh-1*, $Nr5\alpha 2$) is an orphan nuclear receptor family member [11] which could replace *Oct4* in the process of miPSCs induction when using the classic OSKM approach [12], and mESCs like humaninduced pluripotent stem cells (hiPSCs) could be induced

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by the addition of $Nr5\alpha 2$ and retinoic acid receptor gamma (RAR gamma) to OSKM [13]. Mouse epiblast stem cells (mEpiSCs) could be reset to the naïve state by ectopic expression of $Nr5\alpha 2$ [14], while T-box transcription factor, Tbx3, could improve the germline transmission of miPSC chimeras [15]. Therefore, in this study, $Nr5\alpha 2$ and Tbx3 have been chosen for induction of piPSCs and to determine whether these two transcription factors play similar roles in porcine stem cells.

We optimized the medium by modified the LBX medium which had roles in pluripotent states conversion and the improvements of hiPSCs [16]. This modified medium was named MX. In our study, TN was supplemented in the piPSCs induction system on the basis of OSKM. The results showed that TN increased the number of alkaline phosphatase-positive colonies. In addition, we observed the morphology of piPSCs, and some conventional pluripotency characteristics in 6 F group were very similar to mESCs.

Materials and Methods

Reagents and Media

The reagents and media were purchased from Life Technologies, R&D, Millipore and BD (all USA). The porcine embryonic fibroblast cells (PEF) were derived from Duroc and Landrace pigs which were provided by Harbin Sanyuan pig farm. ICR mice for mouse embryonic stem cells (MEF) were purchased from Beijing Vital River.

Cell Culture

The KOSR medium consisted of 76 % knockout Dulbecco's modified eagle medium (DMEM), 20 % knockout serum replacement, 8 ng/ml bFGF, 2 mmol/L L-glutamine, 0.1 mmol /L β-mercaptoethanol, 1 % MEM nonessential amino acids, and 1 % penicillin-streptomycin. MX medium is mixture of 1:1 MX-1 and MX-2 medium, but the concentrations of bFGF and LIF were not changed. The MX-1 medium consisted of 76 % knockout DMEM, 20 % knockout serum replacement, 8 ng/ml bFGF, 2 mmol/L L-glutamine, 0.1 mmol /L β-mercaptoethanol, 1 % MEM nonessential amino acids, and 1 % penicillin-streptomycin. MX-2 medium consisted of 48 % DMEM/F12, 48 % Neurobasal, 1,000U/ml LIF (human), 0.5 % N2, 1 % B27, 0.5 mg/mL BSA and 1 % penicillin-streptomycin. FBS medium consisted of 86 % high glucose DMEM, 10 % FBS, 1 % NEAA, 2 mmol/L L- glutamine, and 1 % penicillin-streptomycin. Freezing medium consisted of 90 % FBS and 10 % dimethyl Sulfoxide (DMSO). Pig embryonic fibroblasts (PEF) were obtained from 33.5-dayold pig fetus via collagenase IV and DnaseIdigestion followed by culturing in FBS medium and then passaged by 0.25 % Trypsin-EDTA enzyme. Mouse embryonic fibroblasts (MEFs) were acquired from 13.5-day-old mouse embryo and were treated with 10 µg/mL mitomycin C as feeder layer. MEF and feeder layer were cultured in FBS medium. piPSCs were cultured in MX medium and dissociated with TrypleTM. All the porcine and mouse cells were cultured in an incubator under the conditions of 5 % CO₂ and 37 °C. These cells were frozen according to a program freezing method. In brief, after digestion, the cells were suspended with 500 µl freezing medium and subsequently transferred into a -80 °C freezer for storage.

Retroviral Transduction

pMX plasmids containing mouse *Oct4*, *Sox2*, *Klf4*, *c-Myc*, *Nr5* α 2 and *Tbx3*, and VSV-G and Gag-Pol were purchased from Addgene. A total of 293 T cells were transfected with 42 µl LTX and 21 µl of PLUS regent. The ratio of VSV-G, Gag-Pol, and pMX was 5:10:16. Supernatants were collected at 24 and 48 h after transfecting, and filtered through a 0.45-µm filter. Finally, 10⁴ PEF per well were infected with concentrated virus.

Immunofluorescence Analysis and Alkaline Phosphatase

piPSCs were fixed with 4 % paraformaldehyde for 30 min at room temperature. After three times washing in PBS, they were permeabilized with 0.5 % triton for 1 h at 37 °C, followed by blocking with 2 % bovine serum albumin (BSA). Subsequently, these cells were incubated with primary antibodies to Oct4 (Santa Cruz, Sc-8628, 1:200), Sox2 (Santa Cruz, sc-17320, 1:200) and Nanog (Abcam) overnight at 4 °C. The following day, the cells were washed three times with PBS and incubated with secondary antibodies for 1 h at 37 °C. Finally, nuclei were stained with Hochest for 8 min. A Nikon 80i was used for the analysis of these cells. Alkaline phosphatase (AP) staining was performed with BCIP/NBT Alkaline Phosphatase Color Development Kit (Beyotime) following the manufacturer's instructions.

Karyotype Analysis

When piPSCs reached 70~80 % confluence, these cells were treated with 0.4 μ g/ml colchicine for 2.5 h. Subsequently, these cells were harvested, plated into a gelatin-coated dish for 10 min, and further cultured in a hypotonic solution at 37 °C for 15 min. Later, piPSCs were fixed in fixative (3:1 methanol and glacial acetic acid) at room temperature for 5 min. The cells were collected by centrifugation at 800 rpm for 10 min, followed by cold fixative with 6 ml at 4 °C for 20 min, and the process repeated once again.

Table 1Primer sequences forPCR

| Gene name | Forward primers | Reverse primers |
|-----------|------------------------------|---------------------------|
| pOct4 | CAAACTGAGGTGCCTGCCCTTC | ATTGAACTTCACCTTCCCTCCAACC |
| pSox2 | CATCAACGGTACACTGCCTCTC | ACTCTCCTCCCATTTCCCTCTTT |
| pNanog | AATGATCGTCACATATCTTCAGGCTGTA | GTTCCATGGGCTCAGTGGTCAAG |
| pRex1 | ATCCAAGACCACCACCACTG | GTTCACAGCAACATTCAGGTAGA |
| mOct4 | GAGGCCCTTGGAAGCTTAGCC | CCCAGTGTGGTGGTACGGGAAATC |
| mSox2 | TTGACGCGGTCCGGGCTGTTCT | CCCAGTGTGGTGGTACGGGAAATC |
| mKlf4 | TAGTCGGGGCACCTGCTGGACGC | CCCAGTGTGGTGGTACGGGAAATC |
| mc-Myc | CGCTCTGCTGTTGCTGGTGAT | CCCAGTGTGGTGGTACGGGAAATC |
| Fgfr1 | ACTGCTGGAGTTAATACCACCG | GCAGAGTGATGGGAGAGTCC |
| Fgfr2 | TGATGATGAGAGACTGTTGGCATGC | TCCAAGTAGTCCTCATTGGTCGTG |
| Lif | CACTGGAAACACGGGGGCA | AGGGCGGGAAGTTGGTCA |
| Bmp4 | CGTCATCCCAGATTACAT | GAGTCGAAGCTCTGCGGAT |
| Smad4 | GGCTTCAGGTGGCTGGTCGGA | ACCTGATGGAGCATTACT |
| Lifr | CTCATCCCAGTGGCAGTG | CCAGAACCTCAACATTAT |
| bFGF | GCGACCCTCACATCAAACT | CAGTGCCACATACCAACT |

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Then, 2 ml of cold fixative was added and centrifuged. Finally, spreads were made by "huffing" on slides, holding at 45 °C, and dropping one drop of cells onto the top of the slide. After drying for 5 min, the sides were observed under a Nikon 80i microscope. Thirty samples were selected for chromosome number analysis.

Embryoid Body Formation

piPSCs were digested into single cells, transferred into gelatin-coated plates, and incubated in FBS medium for 10 min. Subsequently, the suspensions were harvested and transferred to a dish with FBS medium. The medium was changed every other day and, 6 days later, the EBs formed.

Teratoma Formation

Before harvesting, the piPSCs were transferred to gelatincoated plates. Then, piPSCs were transferred into 1.5-ml eppendorf tube with 600 μ l DPBS at 4 °C. Next, a 300 μ l suspension containing 1×10⁷ piPSCs was injected under the skin of non-obese diabetic/severe combined immunedeficient (NOD/SCID) mice. The majority of piPSCs lines at 15–20 passages can form teratomas. These teratomas were harvested for histological analysis at 4 weeks.

RT-PCR and Quantitative PCR

Total RNA was extracted using TRIzol reagent. RT-PCR was performed using High-Capacity cDNA Reverse Transcription Kits (Applied Biosystems, 4368814) according to the manufacturer's instructions. Electrophoresis was performed on a 2 % agarose gel. QPCR was performed using SYBRPremix Ex TaqTM (TaKaRa) and the 7300

Real-Time PCR System (Applied Biosystems). Primers are listed in Table 1.

Results

Tbx3 and Nr5α2 Could Improve pig iPSCs Generation

Porcine embryonic fibroblast cells (PEF N and PEF-9) were infected by the retroviral factors OSKM (4 F), OSKMN, OSKMT, and OSKMNT (6 F). The infected cells were cultured in two kinds of culture media, KOSR medium (hESCs culturing) and MX medium. AP staining was carried out in these groups and positive colonies were counted (Fig. 1b). The number of AP positive colonies in the 6 F group were significantly increased compared with the other groups (Fig. 1c). On day 7 after infection, the majority of the colonies induced by 4 F were flattened similar to the hESCs colonies. However, 6 F-induced colonies were domed, similar to mESCs colonies. The colonies were dissociated mechanically and were then passaged using type III (6 F groups)or type IV collagenase (4 F groups)(Fig. 1a). Two and 12 pig iPSCs cell lines were established from 4 F and 6 F induction separately. The 6 F-induced pig iPSCs lines were morphologically similar to mESCs and could be digested into single cells, while the 4 F-induced cell lines were like hESCs whose single cell colony formation ability was poor (Fig. 1a). Attempts to culture piPSCs have been made in different feeder-free systems, in which the plates were treated with matrigel (BD), FBS, or gelatin, or in a matrix-free culture. The results showed that none of these systems is suitable for stable culture of 4 F-induced piPSCs, while the matrigel and matrix-free culturing system could produce 6 F-induced piPSCs which survived. In the latter



Fig. 1 Tbx3 and Nr5 α 2 can improve piPSCs generation. a The PEF was infected by 6 F and 4 F, respectively. *D1* shows infected PEF 1 day later. After 7 days, some colonies came out and picked for establishing cell lines. 6 F piPSCs were passaged routinely using TrypleTM and appeared domed. 4 F piPSCs were passaged routinely using 1 mg/mL type IV collagenase and appeared flattened. **b** AP staining for counting the positive colonies of diverse factors groups in MX medium. *OSKM* represents *Oct4*, *Sox2*, *Klf4* and *c-Myc*; *OSKMN* represents *Oct4*, *Sox2*,

subculturing, piPSCs could be cultured for more than 30 passages in the matrix-free systemwhereas, when cultured on matrigel, these cells readily formed EBs (Fig. 3e).

Characterization of piPSCs

The piPSCs had a normal karyotype of 38 chromosomes both in 4 F and 6 F groups (Fig. 2b and Supplementary

Klf4, c-Myc and *Nr5* α 2; *OSKMT* represents *Oct4, Sox2, Klf4, c-Myc* and *Tbx3*; *OSKM* represents *Oct4, Sox2, Klf4, c-Myc, Nr5\alpha2* and *Tbx3*. Each had three replicates. **c** Number of AP-positive colonies of the four groups in KOSR and MX medium. The numbers of each group were counted from 3 wells, repectively, and each well was counted for three times. Values with different *superscripts* are significantly different on the same histogram figure by one-way ANOVA. *P*<0.05

Fig. 1B). We characterized the pluripotency of these pig iPSCs by using immunofluorescence staining and RT-PCR. The results showed that the classic pluripotent markers (OCT4, SOX2 and NANOG) were expressed in both 4 F and 6 F groups (Fig. 2a and Supplementary Fig. 1A). RT-PCR showed that the exogenous genes were not silenced and the endogenous pluripotent genes including *Oct4*, *Sox2* and *Nanog* were expressed in both the 4 F and 6 F groups.

Fig. 2 Characterization of piPSCs. a Immunostaining for pluripotency markers of 6 F piPSCs. Positive OCT4 (green), SOX2 (green) and NANOG (purple) were observed. DNA was stained with Hoechst 33342 (blue) and propidium iodide (PI, red). b Karyotype analyses of 6 F piPSCs. More than 80 % of the cells showed normal pig karyotype of 38 chromosomes. c RT-PCR for transgenes (mOct4, mSox2, mKlf4, mc-Myc, mNr5 α 2 and *mTbx3*) and endogenous (pOct4, pSox2, pNanog and pRex1) expression. The first column is the 6 F piPSCs and the second and third columns are 4 F piPSCs and H₂O, respectively. d Embryoid body (EB) formation of 6 F piPSCs. Scale bars 200 µm. e Histological examination of stained teratomas derived from 6 F piPSCs. Nervous tissues represent the ectoderm; cartilage tissue represents the mesoderm: hepatocvte represents the endoderm (arrows)



However, *rex1* was only expressed in the 6 F group (Fig. 2c). Both the 4 F and 6 F-induced piPSCs could form embryoid bodies (EBs) which indicated the differentiation ability of piPSCs in vitro (Fig. 2d and Supplementary Fig. 1C). The in vivo differentiation ability of 6 F-induced piPSCs was checked by injection of these cells into severe combined immune-deficient (SCID) mice and detecting teratoma formation. Teratomas were observed in the 6 F group by 3 weeks, while in the 4 F group, teratomas appeared a little later. Histological examination showed that the

teratomas contained three germ layers including hepatocyte (endoderm), cartilage (mesoderm), and nervous tissue (ectoderm) as shown by the arrows in Fig. 2e.

Tbx3 and Nr5 α 2 Generated piPSCs That are Morphologically Similar to Mouse ESCs

As the 6 F-induced piPSCs satisfied the conventional pluripotency, their characteristics were further detected. Morphologically, 6 F-induced piPSCs colonies were similar to the



Fig. 3 6 F piPSCs were similar to mESCs. **a** The morphology of 6 F and 4 F piPSCs. **b** Proliferation detection of the 6 F piPSCs. Double time: TD=txlog2/(logNtlogN0)=18.5 h. **c** Single-cell cloning efficiency of piPSCs. 6 F and 4 F piPSCs were cultured in MX medium. **d** Quantitative RT-PCR for *Fgfr1*, *Fgfr2*, *Lifr*, *Lifr*, *Bmp4*, and *Smad4*.

mESCs, while the 4 F-induced piPSCs colonies were similar to hESCs (Fig. 3a). Then, we assayed the growth of piPSCs by measuring the cell cumulative increase. The 6 F-induced The expressions of these genes were relative to the expression of β actin. e 6 F piPSCs could be cultured in feeder-free system. The 4 F and 6 F piPSCs were cultured in three matrix (matrigel, FBS, gelatin) and without matrix. *Scale bars* 200 µm

piPSCs grew faster whose doubling time was 18.5 h similar to mESCs (Fig. 3b). To test the piPSCs viability, single-cell colony efficiency was performed and the results showed that Fig. 4 piPSCs generated with MN or N. a piPSCs generated with MN. D10 the morphology of the colonies on the tenth day after PEFs were infected. P5 piPSCs morphology at the fifth passage. AP piPSCs were positive for AP activity. b piPSCs generated with N. D10 the morphology of the colonies on the tenth day after PEFs were infected. P5 piPSCs morphology at the fifth passage. AP piPSCs were positive for AP activity. c RT-PCR for detection of transgenes



the colony efficiency was decreased in the 4 F-induced piPSCs compared with the 6 F-induced piPSCs. Meanwhile, we also found that the single-cell cloning efficiency was decreased in piPSCs when cultured in KOSR medium compared with that in MX medium in both the 4 F and 6 F groups (Fig. 3c). To determine which signaling pathway involved in the regulation of single-cell colony efficiency of piPSCs, key downstream effector molecules of LIF and bFGF pathways were examined by using real-time PCR. The results showed that the expression of *Lifr*, *Smad4*, and *Bmp4* were higher in the 6 F than in the 4 F group, and, reversely, the expression of *Fgfr1*, *Fgfr2*, and *bFGF* were higher in the 4 F than that in the 6 F piPSCs (Fig. 3d).

PiPSCs Could be Induced by two Factors ($Nr5\alpha 2$ and c-Myc, MN) or a Single Factor ($Nr5\alpha 2$, N)

So far, we could generate piPSCs by 6 F efficiently. In order to use fewer transcription factors to induce piPSCs formation, we found that two factors, $Nr5\alpha 2$ together with *c*-Myc, could generate piPSCs in place of *Oct4*. MN-induced piPSCs had normal proliferative ability and could be passaged stably (Fig. 4a). Surprisingly, we found $Nr5\alpha 2$ alone could induce piPSCs generation but *Oct4* alone never got piPSCs. Single-factor $Nr5\alpha 2$ -induced piPSCs could be cultured routinely and AP staining was positive (Fig. 4b). The expression of $Nr5\alpha 2$ and *c*-*Myc* were detected in MN and N-induced piPSC by RT-PCR, and no other transgenes were found to be involved in this process (Fig. 4b). Further detection will be carried out in the future experiments

Discussion

Morphologically, 6 F-derived piPSCs were similar to mESCs but different from those derived by the 4 classical factors. The 6 F-induced piPSCs colonies were domed. The cells in these colonies were closely connected with each other. 6 F-induced piPSCs can be digested into single cells and cultured in a feeder-free system. The differences

between 6 F-induced piPSCs and 4 F-induced piPSCs indicated that Nr5a2 and Tbx3 played an important role in establishing mESCs-like piPSCs. The 6 F-induced piPSCs possess better viability and more rapid proliferative ability than the 4 F-induced iPSCs. $Nr5\alpha 2$ can bind both the proximal enhancer and proximal promoter regions of Oct4 and regulate Oct4 expression in the epiblast stage of mouse embryonic development [17]. And it can replace Oct4 in miPSCs induction [12]. The combination of $Nr5\alpha 2$ and RAR gamma can increase the miPSCs generation due to the activation of endogenous Oct4 and can generate mESCslike hiPSCs by maintaining Oct4 expression [13]. Therefore, in the induction of piPSCs, $Nr5\alpha 2$ may also play a similar role. It can enhance the expression of endogenous Oct4, and activate the LIF signal pathway (Fig. 3d). Tbx3 can maintain ESC self-renewal in an undifferentiated state by its persistent expression in the absence of LIF [18, 19] and can improve the germ-line competency of miPSCs [12]. Practically, 6 F piPSCs in this study had a dominant position in culture. Therefore, in this study, the existence of $Nr5\alpha 2$ and Tbx3 has promoted piPSCs to grow more like mESCs which were distinct from the 4 F piPSCs.

6 F-induced piPSCs' mESCs-like pluripotent state can be proved through the signal genes analysis. *Lifr* and *Smad4* expressions were higher in 6 F piPSCs compared to 4 F piPSCs (Fig. 3d). In mESCs, *Lifr* had a higher expression and *Smad4* can maintain the pluripotent state through taking part in inhibiting ERK activity [20, 21]. These findings suggested that *Nr5a2* and *Tbx3* may have some functional roles in pig. Single cell colony formation ability can be used for transgenic porcine production. 6 F piPSCs can be cultured and passaged in a feeder-free system and this can accelerate the piPSCs application for transgenic research and screening for some drugs.

In this study, piPSCs have been successfully induced by two factors (MN) or a single factor (N). It has been reported that piPSCs have been obtained by other two factors (*Oct4* and *Klf4*) [22, 23] and miPSCs can be obtained by *Oct4* alone [24]. In murine, $Nr5\alpha 2$ can mediate pluripotent genes expression [25]. *Nanog* was able to rescue $Nr5\alpha 2$ knockdown but not *Oct4* knockdown in ESCs [12]. So far, there has been no report on the generation of miPSCs by a single factor or two factors induction without *Oct4* being involved. The MN and N piPSCs have not been detected for more pluripotent markers, but the proliferation ability of these cells indicated that $Nr5\alpha 2$ may had an important function in porcine stem cell which is similar to the role of *Oct4* in mESCs.

Researchers have tried to establish porcine embryo stem cells (pESCs) for about 20 years, but it has proved to be very difficult [26, 27]. It seems that these pluripotent genes which are important in mouse and human ESCs, such as *Oct4* and *Nanog*, do not have crucial roles in porcine ESCs, but some

other pluripotent genes may affect the establishment of pESCs. In this study, we proved that $Nr5\alpha^2$ and Tbx3 play important roles in piPSCs induction. Tracking the expression of $Nr5\alpha^2$ and Tbx3 in the early porcine embryo and the analysis of $Nr5a^2$ - and Tbx3-related signaling pathways may contribute to the future establishment of pESCs.

In summary, we found that the genes $Nr5\alpha^2$ and Tbx3 improved the generation of piPSCs, and the generated cells exhibited outstanding cell viability. The features of the 6 F piPSCs imply that they could be a good source for future domestic applications as well as for the basic research that utilizes porcine pluripotent cells.

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Conflict of Interest The authors declare no potential conflicts of interest.

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