

## Generation and characterization of induced pluripotent stem cells in domestic Asian water buffalo (*Bubalus bubalis*)

Research Article

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### Abstract

Induced pluripotent stem cells (iPSCs) have numerous applications in livestock to improve production traits, disease resistance, biopharming, conservation of germplasm, disease modeling, regenerative medicine etc. Here, we have described the derivation of iPSCs from buffalo fetal fibroblasts (BuiPSCs) by lentivirus based transduction of mouse derived pluripotency marker genes Oct4, Sox2, Klf4 and c-Myc. The BuiPSCs showed typical buffalo embryonic stem cells like colony morphology which were alkaline phosphatase (AP) positive and expressed pluripotency markers Oct4, Nanog, SOX2, KLF4, FoxD3 and SSEA1, TRA-1-60, TRA-1-81. The cells were carrying normal karyotype and were able to differentiate into cell of all three germ layers *in vitro*. The BuiPSCs could be propagated beyond 20<sup>th</sup> passages. To the best of our knowledge, this is the first report on generation of buiPSCs in domestic water buffalo (*Bubalus bubalis*) using mouse derived transcription factors and the reprogrammed cells could self renew more than 20<sup>th</sup> passage.

**Keywords:** Buffalo; Fetal Fibroblast; iPSC Cells; Lenti Virus Vector; Mouse Derived Transcription Factors.

### Introduction

Since the discovery of embryonic stem cells (ESCs) in mouse [1], stem cells technology has been envisioned to revolutionize livestock industry. Because ESCs are pluripotent, has remarkable plasticity and a single ESC can divide to form unlimited number of stem cells, they can be manipulated in Petri dish for wide range of applications including production of transgenic and cloned animal, generation of gametes, drug testing etc. However, in spite of considerable efforts, derivation of proven ESCs line in farm animals has so far been elusive [2, 3]. The induced pluripotent stem cell (iPSCs) technology first described in 2006 [4] has revolutionized the stem cell research arena by providing another source of pluripotent stem cells even from each individual surpassing the constraint of availability of embryos and other issues. Surprisingly it was observed that the iPSCs resembled ESCs in cell morphology, expression of surface markers, telomerase activity, teratoma formation, gene expression, DNA methylation, formation of chimera etc [4] and thus iPSCs have been designated an alterna-

tive to ESCs. The success of this technology has been proved by generation of iPSCs in human [5], rat [6, 7], non-human primates [8], pig [9], canine [10], cattle [11], sheep [12], hircine [13], equine [14] etc. Tremendous progress in iPSC technology within a very short period of time has emphasized its promising applications in regenerative medicine, “disease-in-a-dish” modeling, drug and toxicity screening, custom tailored therapeutics, transgenesis and for studying the developmental biology. Recently, preclinical studies has been successfully done for treating retina injury in pigs using pig iPSCs [15], cardiovascular disease in dog using dog iPSCs [16] and myocardial infarction in pigs using human iPSCs [17].

India is home to more than 50% of the buffalo (*Bubalus bubalis*) population of the world. It is also found in other south East Asian countries and commonly used for milk, meat and draught purpose. Buffalo iPSCs can be an alternative to buffalo ESC and can play a role in generation of improvement of milk production, growth rate, carcass composition, diseases resistance, reproductive performance, generation of gametes, conservation of germ

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plasm etc. Research conducted so far indicates that derivation of pluripotent ESCs in buffalo either from early stage embryos or from blastocyst are limited and the success rate has been very low [18, 19]. Therefore, iPSCs technology can be used for deriving pluripotent stem cells in this species. Recently, attempt has been made to derive iPSCs in buffalo using proteins of buffalo defined transcription factors, however, the cells could not be propagated beyond 10<sup>th</sup> passage [20]. In the present study, we have generated buffalo iPSCs using mouse derived transcription factors (OKSM) which could be propagated beyond 20<sup>th</sup> passages. The buffalo fetal skin fibroblast cells were transfected with lentivirus particle carrying the pluripotent genes and the colonies developed were mechanically propagated, characterized for expression of pluripotent markers, chromosomal integrity and ability to form three germ layer specific cells.

## Material and Methods

### Ethics statement

The entire procedure of collection of buffalo fetus from local large animal slaughter house and its processing to obtain skin fibroblast cells in lab, was approved by the Animal Ethics Committee of Indian Veterinary Research Institute, India.

### Preparation buffalo fetal skin fibroblast cells (BFF)

Buffalo fetuses of about 80 days of gestation were collected from the local large animal slaughter house, Bareilly and were transported in ice to the laboratory. The fetuses were washed thoroughly with PBS and all the appendages including head were removed, followed by washing again with PBS. The fetal skin was gently removed and minced into small pieces using BP blade and scissors. The skin explants were collected in fibroblast media (FM) containing knock out DMEM (Invitrogen), 10% FBS (Gibco), 1mM L-glutamine (Gibco), 1% Non essential amino acids (Gibco) and 0.5% Penicillin- Streptomycin (Gibco) as a suspension, centrifuged twice at 1000 rpm for 5 minutes and the pellet containing explants was cultured in FM for harvesting fibroblast cells. The culture plates were treated with 0.05% trypsin (Gibco) and the dispersed cells were subsequently passaged in FM.

### Plasmid preparation and production of Lentivirus

pLentG-KOSM plasmid vector (Cell Biolabs) in which the mouse transcription factors (OKSM) are in-frame fused into a single open reading frame (ORF) via self-cleaving 2A peptides followed by IRES-GFP as a reporter for viral transduction was procured commercially in bacterial glycerol stock. The bacteria were grown in LB broth and plasmid DNA was isolated from the bacteria using mini and maxi prep culture kit (MDI) as per the manufacturer's instruction. The 293 T cells were plated in 35 mm dishes at a concentration of  $1 \times 10^5$  and incubated at 37°C and 5% CO<sub>2</sub> for overnight. About  $1 \times 10^5$  number of 293 T cells were plated in 35 mm tissue culture dishes and incubated at 37°C and 5% CO<sub>2</sub> for overnight. The cells were then transfected with a mixture of vectors containing 2 µg of pLentG KOSM, 0.7 µg each of pRSV-Rev packaging vector, pCMV-VSV-G envelope vector as well as pCgpV packaging vector (Cell Biolabs) using Lipofectamine™ Plus (Invitrogen) as a transfection reagent following the manufacturer's instruction. Briefly, The viral supernatant was

harvested after every 12 hours from 36 to 72 hours post transfection and stored at 4°C over the collection period. The lentivirus supernatant were pooled, and centrifuged at 1500 rpm for 5 minutes to remove the cell debris, filtered in 0.22 µm filter and stored at -80°C in aliquots.

### Estimation of lentiviral titer

To ensure that the pseudoviral medium is viable and to control the number of copies of integrated viral constructs per target cell, the lenti viral titer was estimated by Quick Titer™ Lentivirus Titer Kit (Cell Biolabs, Vat No VPK-107). The viral titer estimates were  $8 \times 10^7$  lentiviral particles/ml. The stocks were frozen stored and used for transductions of fibroblast cells after thawing.

### Transduction of BFF

One day before transduction, BFF of 3rd passage were seeded in a six well plate at a concentration of  $2 \times 10^4$  cells/well in FM. Lentivirus supernatant (2 ml) with ViraDuctin™ Lentivirus Transduction Reagent (Cell Biolabs) were added in each well as per the manufacturer's instruction. This was equivalent to approximately  $16 \times 10^7$  viral particles per 20000 cells. After 48 hrs of viral transduction, FM was replaced with buffalo ESC media containing knock out DMEM, 20% FBS, 1mM L-glutamine, 0.1mM β mercaptoethanol, 1% non essential amino acids, 0.5% penicillin-streptomycin, 10ng/ml bFGF (Millipore) and 4ng/ml human LIF (Millipore). The transduced cells were monitored daily and ESC media was changed regularly. The cells were subcultured in 90mm tissue culture dish following 90% confluence. The putative iPSC colonies were picked up and subcultured mechanically on BFF monolayer which was inactivated using mitomycin C as described earlier [21]. The same process was repeated for further propagation of buiPSC colonies. The detail time line for reprogramming of BFF has been presented in Figure 1& 2.

### Alkaline phosphatase (AP) staining

The buiPSC like colonies were stained for alkaline phosphatase (AP) as per the earlier described method [22]. Briefly, the culture plates were rinsed twice in 1X PBS (pH 7.4) and fixed in 4% paraformaldehyde in 1X PBS for 15 min at room temperature. The fixed cells were washed twice with distilled water, stained with 0.5 mg/ml naphthol AS-MX phosphate (Sigma) and 1 mg/ml FAST RED TR salt (Sigma) in 100 mM tris buffer (pH 8.2) for 4 h at room temperature. The staining was stopped by washing the cells in distilled water and the cells were observed under microscope for AP activity.

### In vitro differentiation

The differentiation ability of buiPSC cells were examined by hanging drop culture. Briefly, a single cell suspension of iPSCs ( $2.5 \times 10^4$  cells/ml) were prepared in differentiation media consisting of IMDM (Gibco) supplemented with 20% fetal calf serum, 1% non-essential amino acids, 2 mmol/l L-glutamine, and 100 µmol/l β-ME. From this cell suspension, 20 µl was pipetted on to the lid of a tissue culture dish and then inverted over its bottom dish containing 5 ml phosphate-buffered saline. On day 3, the embryo bodies (EBs) was transferred on gelatin coated coverslip in 6 well tissue culture dishes and grown up to day 11, with

Figure 1. Time line for reprogramming of buffalo fetal skin fibroblast.

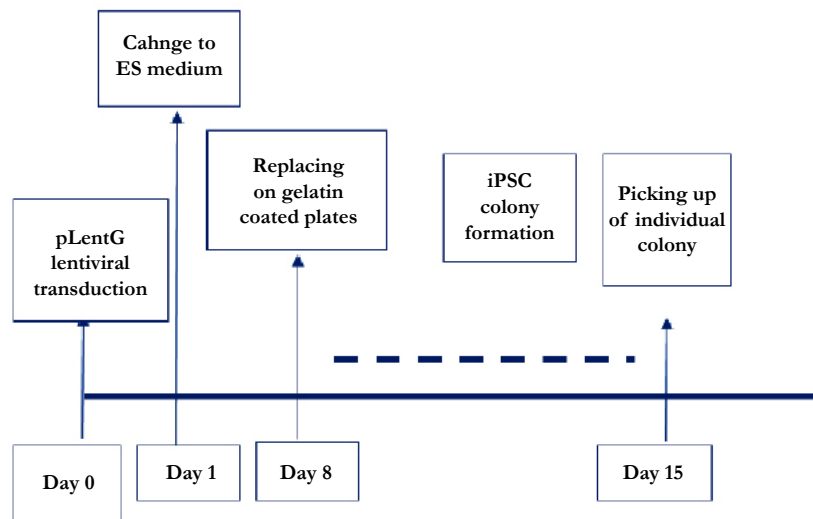
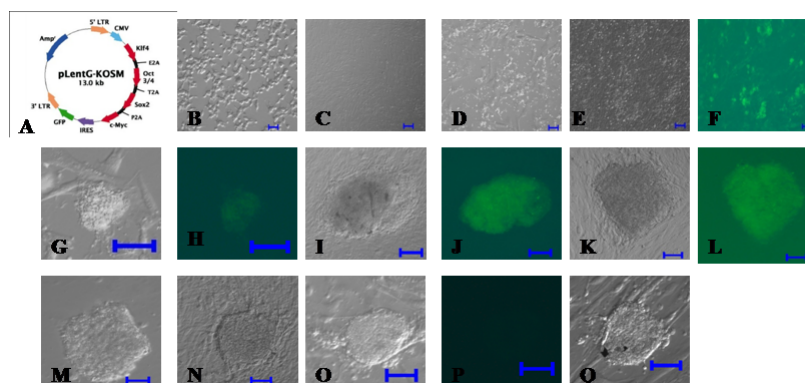


Figure 2. Generation of BuiPSC. (A) A 13.0 kb polycystronic lentivirus with mouse defined pluripotent genes (OSKM) was used to reprogram buffalo fetal skin fibroblast, (B) 293T cells were used for lentivirus production, (C) buffalo fetal skin fibroblast cells, (D) buffalo fibroblast with lentivirus particle having pLentG KOSM, (E) Transduced cells after 48 hrs, (F) Transduced cells after 48 hrs showing GFP expression, (G-H) Small colony after day 4 of transduction, (I-J) iPSC like colonies after day 20th of transduction, (K,L) BuiPSC colony after first passage (P1), (M) BuiPSC colonies at passage Six (P6), (N). BuiPSC colonies at passage 12, (O) BuiPSC in 16th passage (P) No expression of transgene as indicated by GFP expression at P16<sup>th</sup>. (Q) P24th Passage. Scale bar 100µm



periodical changing of the media. For studying the expression of genes indicating the formation of cell of three germ layers, 8-10 numbers of EBs were collected on different days (EB-D0, EB-D3, EB-D7, EB-D9, EB-D11) for isolation of total RNA. On day 9, the EBs on cover slip was also processed for immunocytochemistry using ectoderm (G-FAP), endoderm (AFP) and mesodermal (alpha SMA) cells specific antibody.

### Immunocytochemistry

Immunocytochemistry was done as per the methods of National Institutes of Health, resource for stem cell research, USA. Briefly, cells were fixed with 4% paraformaldehyde, blocked with 5% BSA in PBS, then incubated with the following primary antibodies overnight at 4°C: mouse monoclonal anti-TRA-1-60 and TRA-1-81, Oct4, Nanog, SSEA1 (Santa Cruz Biotechnology), G-FAP (Sigma), AFP (Sigma) and alpha SMA (Sigma). The cells were then immunostained with appropriate Alexa FluorR594 conjugated secondary antibodies (Life Technology) for 2 h. The negative control was performed following the entire procedure, except adding primary antibodies.

### Molecular characterization - RNA extraction and real time-polymerase chain reaction

For gene expression analysis, 5-6 number of BuiPSCs colonies of different passages (Po, P5, P10) as well as 8-10 numbers of EBs of each time interval of differentiation were used. The total RNA was isolated from all the samples using Quick-RNA™ MicroPrep kit (Zymo Research Kit) as per manufacturer's protocol and the cDNA was synthesized using fixed amount of isolated RNA by using iScript select cDNA Synthesis kit (Biorad). Amplification of genes was performed using gene-specific primers (Table 1) and Eva Green supermix (Biorad, USA) as per manufacturer's instructions. The PCR reaction was under denaturation at 95°C for 3 min followed by 40 cycles of denaturation at 95°C for 5s, annealing at 54°C (Klf4), 55°C (Oct4), 56°C (Foxd3), 60°C (β-actin, Nanog, Sox 2, C-myc, BMP4 and CDX2) and 65°C (Nestin) for 15s. To ensure the specificity of target amplification, melting curves and agarose (2%) gel electrophoresis of the PCR reactions were performed. The relative mRNA expression of genes was calculated using the  $2^{-\Delta\Delta CT}$  method (Livak and Schmittgen, 2001)

Table 1. Real Time PCR was performed with gene specific primers as mentioned below:

S.N	Gene	Primer	Product (bp)	Reference
1	$\beta$ -actin	F 5'-tcataccatcggaatg-3' R 5'-ccaatccacacggagtac-3'	286	AY141970
2	Oct-4	F 5'-caattgccaagctcctaaag-3' R 5'-gcctctcactcggttctc-3'	288	NM_174580
3	Nanog	F 5'-gtcccggtaagaacaaaa-3' R 5'-tgcatttgctggagactgag-3'	107	[51]
4	Sox 2	F 5'-cggcaaccagaagaacag-3' R 5'-cggcagtggtacttatcc-3'	292	NM_001105463
5	Klf4	F 5'-gaactgaccaggcactac-3' R 5'-ctgattatcattcacaagttg-3'	271	NM_001105385
6	Foxd3	F 5'-agcaagccaagaatagc-3' R 5'-tccagggtccagtagttg-3'	257	XM_610898
7	C-myc	F 5'-tttgctctctgaccag-3' R 5'-tgcccaaagtccatttg-3'	400	Z68501.1
8	Nestin	F-5'-aacgctgagtcattgagaac-3' R-5'-cactgctcctggtcttc-3'	276	NM_001206591
9	BMP4	F-caggctaccaggcattctac-3' R-accacctgtcactcatcc-3'	190	BC105344
10	CDX2	F-5'-aacctgtgcagtgatg-3' R-5'-ctgtgctgctgctgttg-3'	276	NM_001206299

and the results were expressed as a fold change in expression over the control. The samples of at least three experiments were analyzed for gene expression studies.

## Results

The chronology of BuiPSC generation has been presented in Figure 1 and 2. The Lentivirus virus particles having polycistronic vector (Figure 2A) was produced in 293T cell (Figure 2B). The BFF cells ( $2 \times 10^4$  cells/well) (Figure 2C) in a six well culture plate were transduced with pLentG lentiviral vector on day 0 (Figure 2D) and the GFP expression was visible within 2-3 days post transduction (Figure. 2E,F), which indicated the successful integration of transgenes into the host genome. Small clumps of cells were observed on day 4-8 (Figure. 2G, H); however, the confluent transduced cells were further replated in 90 mm dishes to avoid the interference in the formation of colonies due to high number of cells. On day 15, morphologically uniform compact, flattened AP positive colonies had emerged (Figure. 2I, J). From day 15 to day 25 post transduction, numerous well defined colonies resembling buffalo ESC colony morphology were formed. Out of these, 18 colonies were picked up and sub cultured by mechanical passaging on inactivated BFF feeder cells. They grew rapidly and within 3-4 days, well defined secondary colonies had appeared (Figure. 2K, L) which were subsequently passaged following the same procedure (Figure. 2 M, N, O, P, Q). The newly formed colonies on subsequent passages maintained buffalo ESC morphology which were propagated beyond 20<sup>th</sup> passages (Data not shown). An average 24 numbers of well defined colonies were formed out of 20,000 transduced cells with a reprogramming efficiency of 0.12%. Initially, the GFP expression in colonies were strong but started declining at passage 6 to 7 and no expression was observed at 12<sup>th</sup> passage indicating probable silencing of transgene (Figure. 3 A-E). The colonies had well de-

finer borders and were positive for alkaline phosphatase activity (Figure 4A). Using RTPCR, we could detect robust expression of pluripotency marker genes Oct4, Sox2, Nanog, C-Myc, KLF4 and FoxD3 (Figure 4B) in passage P0, P5 and P10. Further, the colonies were strongly immunopositive for Oct4, Nanog, Sox2, SSEA1, TRA-1-60 and TRA-1-81 (Figure. 5A). When analyzed at passage 5, during *in vitro* propagation, the colonies exhibited normal chromosome numbers (Figure. 5B). Further, the colonies readily formed EBs following *in vitro* differentiation (Figure. 6A, B). RTPCR and immunocytochemistry analysis revealed the expression of lineage-specific genes representing the three germ layers: endoderm-cdx2 ( $\alpha$ -feto-protein), ectoderm-Nestin and mesoderm-BMP4 (Figure. 6CDEF). After proper characterization, the colonies generated have been named BuiPSCs.

## Discussion

In the present study, we have generated BuiPSCs by reprogramming fetal skin fibroblast using mouse derived pluripotency factors. The cell colonies aroused from transduced fibroblast had well defined morphology resembling buffalo ESC [18, 19] and expressed various pluripotency markers qualifying them as pluripotent stem cells. The lentivirus transfected cells showed expression of GFP within 48-72h which indicated successful integration of transgenes into fibroblast genome. Small colonies appeared as early as on day 4-8, but it took about 15-25 days to get alkaline phosphatase positive numerous compact large colonies resembling buffalo ESC morphology. In three replicates, an average of 24 colonies were observed providing a comparatively higher reprogramming efficiency. In most of the species, similar trend has been observed and about 10-25 days were the usual time of post transfection when the colonies attained the size for propagation. AP positive compact colonies of incorporate sheep iPSCs were observed after about 13 days post transduction of fibroblast with



Figure 3. Silencing of transgene in BuiPSC. (A, B) representing GFP expression in passage 3, (C) Reducing expression of GFP in passage 7 and no expression was observed at passage 12. Scale bar 100µm.

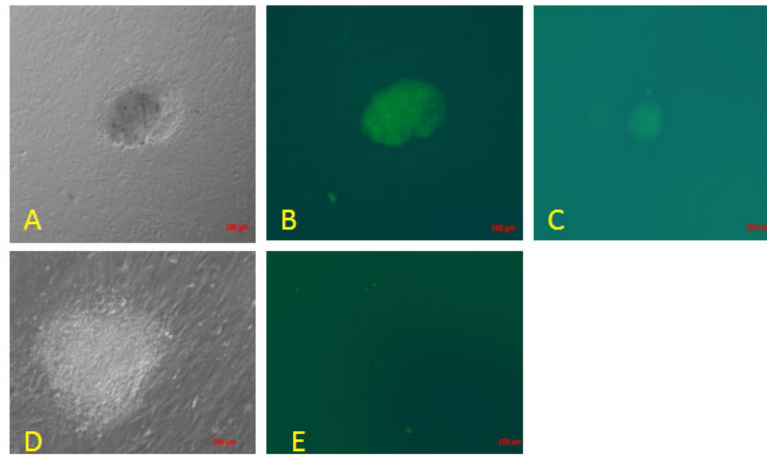


Figure 4. Characterisation of BuiPSC. (A) AP staining, (B) Expression of pluripotent markers genes in different passages of BuiPSC.

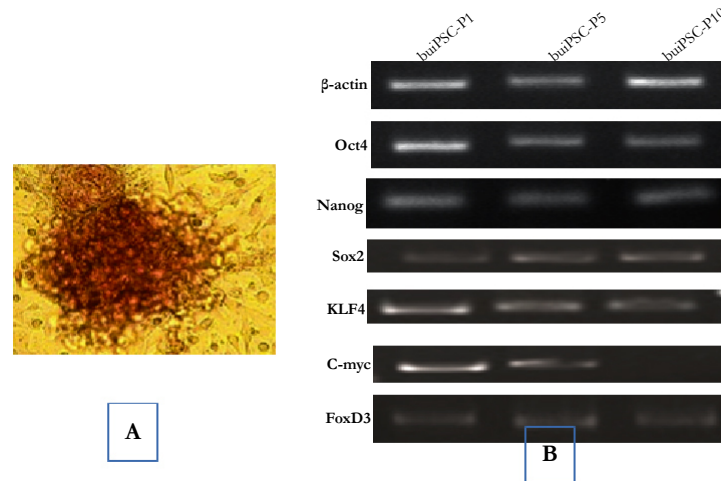
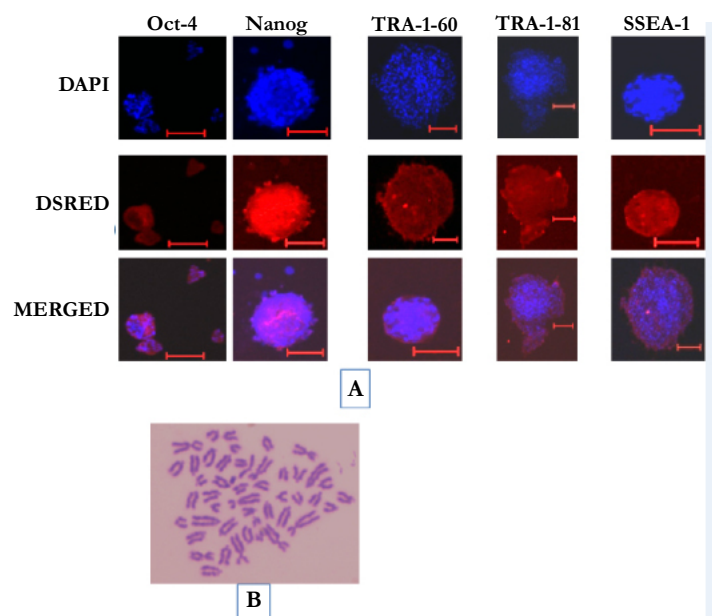
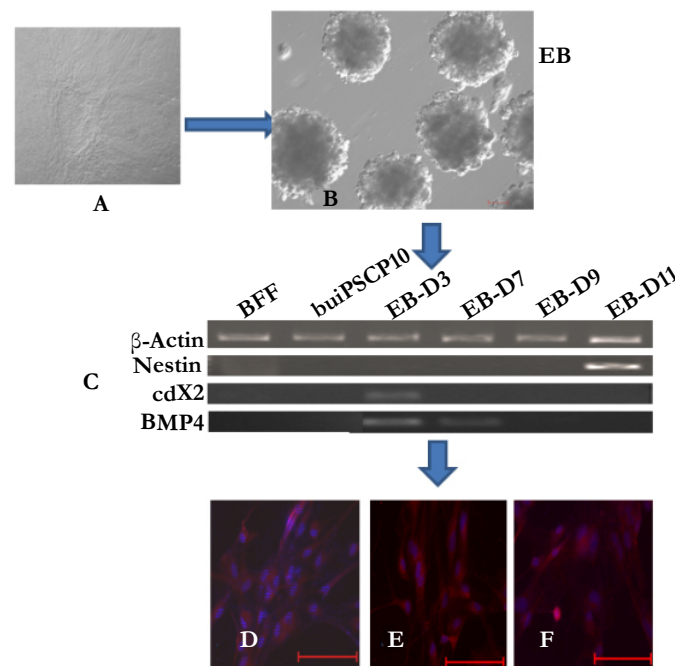


Figure 5. Characterisation of BuiPSC. (A) Immunofluorescence staining showing expression of pluripotent markers in BuiPSC (Scale bar 100µm). B. BuiPSC showing normal Karyotype.



**Figure 6.** *In vitro* differentiation of BuiPSC. (A) BuiPSC colony (B) Embryobodies formed after 72 hrs in differentiation medium © Expression of lineage specific gene expression in BuiPSC derived embryobodies, buffalo fibroblast and BuiPSC: Nestin, (ectoderm), cdX2 (Endoderm) and BMP4 (mesoderm). D, E and F represents the neuronal cells (G-fap), Endodermal cells (AFP) and Mesodermal cells (alpha SMA) (Scale bar 100µm).



very low reprogramming efficiency (0.001875%) [3]. In case of bovine iPSCs, small colonies were formed on day 4 of transduction and colonies were ready for picking up as early as on day 7 [23]. Again the reprogramming efficiency recorded was only 0.0002%. Further, they reported that repeated transfection increased the reprogramming efficiency (0.0007%). In equine iPSCs generation, it took 17-18 days post-transfection, when many clearly defined colonies were formed [14] with a reprogramming efficiency of 0.028% based on the transfected (GFP+) cells which was comparable to those obtained when reprogramming human and mouse fibroblasts with the same transfection system [24]. In the present study, the BFF was transfected only once for 48 h which may be the reason for taking longer days for development of well defined colonies. However, the reprogramming efficiency was 0.048% which was comparable with equine iPSC generation [14]. However, it has been reported that the reprogramming efficiency can be enhanced by using small molecules like valproic acid, suberoylanilide hydroxamic acid (SAHA) and trichostatin A (TSA) [25], p53 siRNA [26] and vitamin C [27]. It is assumed that transcription factor of same species would be better for reprogramming, but in the present study, mouse defined transcription factor was able to generate buiPSCs. In several studies, mouse defined factors have already been shown to reprogram fibroblasts of pig [7, 9], sheep [28], horse [11] etc. These results indicated that defined factors related to iPSCs are highly conserved among species. Silencing of transgenes is important for potential clinical applications of iPSCs. Although transgenes introduced by lentiviral vectors are generally not silenced or poorly silenced [29], integrated transgene of pLentG KOSM was silenced 12 days after transduction and no GFP could be observed through-out subsequent culturing. In the present study, GFP expression was not detectable at 12<sup>th</sup> passage [30]. Whereas, it is reported that the EGFP expression in bovine iPSCs was fully absent only after 20<sup>th</sup> passage [23]. *In vitro* embryo body formation and *in vivo* teratoma formations are the two key

factors of successful generation of pluripotent stem cells. In the present study, the BuiPSCs readily formed well defined embryobodies which on differentiation, formed cells of three germ layers. The iPSCs derived from mouse, human, sheep, bovine, equine etc have all been shown to form embryo bodies and teratoma expressing the cells of ectoderm, endoderm and mesoderm [4, 14, 28, 31, 32]. In the present study, only four transcription factors viz oct4, Sox2, KLF4 and c-myc were able to generate buiPSC. In earlier studies involving different species, iPSCs have been generated by different combinations of transcription factors. For example, only Nanog and Lin28 were able to generate iPSCs from the IMR90 fetal fibroblast line and post-natal fibroblasts from human newborn foreskin [31]. Other experiments which resulted in generation of iPSCs in different species have used combination of Oct4, Sox2, Lin28 and Nanog [33], Oct4 with either Klf4 or c-MYC [34], Oct4, Sox2, Nanog [35], Oct4 and Sox2 [25, 36], Oct4 and Klf4 [37], or even by using only Oct4 [38]. In cattle it has been shown that in addition to Yamanaka factors, nanog is necessary for generating iPSC [39]. We assumed that nanog in addition of KOSM may improve the reprogramming efficiency of buffalo fibroblast cells. We have used polycistronic lentivirus vector produced in the laboratory for transduction of buffalo fibroblast cells. Lentiviruses are a subclass of retroviruses capable of transducing a wide range of both dividing and non-dividing cells. Various reports suggest that cellular reprogramming is best achieved when lentiviral transduction is done [31, 40]. However, successful reprogramming have been achieved by using virus backbone free plasmids carrying the reprogramming factors, piggyBac (PB) transposon gene-delivery system, non-DNA approach using small molecules or chemicals, delivering proteins of transcription factors and miRNA etc [15, 41-44]. Recently reprogramming of fetal fibroblast of buffalo and goat has been attempted using egg extract of poultry [45, 46]. Similarly reprogramming of adult cells have been tried using oocyte extracts in an attempt to

generate iPSC without foreign DNA [47]. In the present study, a single polycistronic vector was used for reprogramming of fibroblast cells. In several studies, multiple individual viral vectors have been used for delivering each of the transcription factors that resulted in high genomic integration. However, the number of viruses necessary to reprogram somatic cells had been reduced by delivering reprogramming factors in a single virus using 2A “self-cleaving” peptides, which support efficient polycistronic expression from a single promoter, thus increasing reprogramming efficiency [48, 49].

Pluripotent stem cell generation in domestic ungulates has so far been very difficult [50]. As stated earlier, derivation of pluripotent ESCs in buffalo either from early stage embryos or from blastocyst are limited and the success rate has been very low [18, 19]. Therefore, iPSCs technology can be used for deriving pluripotent stem cells in this species. Unfortunately, the initial attempt to produce buffalo iPSCs has been partially successful as the reprogrammed cells could be propagated only upto 10<sup>th</sup> passage [21]. Therefore, our study is important as we could propagate the cells beyond the numbers of passages achieved by Deng group. This better progression may be due to use of poly cistronic vectors consisting of pluripotent genes and delivered through lentivirus vector.

Application of stem cell technology in improving animal production is a promising area. As stated earlier, despite more than two decades of effort, establishment of pluripotent embryonic stem cells from ungulates has remained an elusive goal. Recently derivation of buffalo ESCs has been reported but the success rate has been very low. Therefore, buffalo iPSCs would be the alternative to ESC. The advantage of this technology is that, quality pluripotent cells can be generated from individual genetically superior animal and these cells not only be used for animal cloning and transgenesis, but gametes can also be produced *in vitro* ensuring propagation of the superior germ plasm. Further, if a high producing buffalo male or female fails to produce gametes due to toxicity, radiation exposure or failure of ejaculation due to permanent disability, iPSCs can be generated from such quality animal for further propagation. Apart from these, such cells would also open up possibilities to use the buffalo as an excellent pre-clinical model for future human stem cell-based therapies. However, virus back bone free iPSCs need to be generated for their large scale application in future.

## Conclusion

In the present study, we report on successful derivation of iPSC cells from buffalo fetal fibroblasts. The BuiPSCs displayed pluripotent characteristics as evidenced by expression of key pluripotency markers, alkaline phosphatase activity, and the formation of cells representing all the three embryonic germ layers.

## Acknowledgements

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