

## **Derivation of ringed seal (*Phoca hispida*) tripotent induced pluripotent stem-like cells**

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## Abstract

Induced pluripotent stem (iPS) cells have been produced just for a few species among order Carnivora: snow leopard, Bengal tiger, serval, jaguar, cat, dog, ferret, and American mink. For the first time, we derived the ringed seal (*Phoca hispida*) iPS-like cells. We had shown the expression of pluripotency marker gene *Rex1*. Ringed seal iPS-like cells were able to differentiate into derivatives of endoderm (expression of *AFP*), mesoderm (adipocytes and osteocytes), and trophoctoderm (expression of *Cdx2*). To confirm pluripotency, we need to differentiate cells into ectoderm cell types, for instance into neurons.

## Keywords

pluripotency, reprogramming, ringed seal, iPS cells

## INTRODUCTION

Order Carnivora consists of two suborders: Caniformia and Feliformia. It includes domestic animals, that are used for disease modeling, fur-bearing animals, such as mink, marine mammals, and other species. Derivation of induced pluripotent stem (iPS) cells could allow insights into pluripotency and embryonic development of these species, as well as the development of new disease models. Currently, pluripotent cells were produced from several Carnivora species: dog (Luo et al., 2011; Baird et al., 2015; Lee et al., 2011; Vaags et al., 2009; Shimada et al., 2010; Whitworth et al., 2012; Koh et al., 2012; Tsukamoto et al., 2018), snow leopard (Verma et al., 2012), Bengal tiger, serval, jaguar, and cat (Verma et al., 2013; Dutton et al., 2019; Gómez et al., 2010), ferret (Gao et al., 2020), and American mink (Menzorov et al., 2015).

Marine mammals represent the basal Carnivora group. Their iPS cells would expand our knowledge of embryonic development and differentiation. We report the generation of ringed seal (*P. hispida*) iPS-like cells. These cells expressed pluripotency marker *Rex1* and were able to differentiate into endo-, meso-, and trophoctoderm.

## MATERIALS AND METHODS

### Production of ringed seal fibroblasts

Primary fibroblasts of the ringed seal were obtained from lung necropsy. Tissue samples from wild female were collected during aboriginal quota sealing in the coastal waters of the Bering Sea (Mechigmen bay, Chukotka Autonomous Okrug, Russia). To establish primary

fibroblast's cell culture, we used conventional technique (Stanyon, Galleni, 1991). The fibroblast culture medium consisted of a-MEM supplemented with 15 % fetal bovine serum (FBS), 1x MEM non-essential amino acids solution, 1x GlutaMAX supplement, and 1x penicillin-streptomycin (Thermo Fisher Scientific, USA).

### **Production of ringed seal iPS cell lines**

To produce iPS cells from the ringed seal fibroblasts we used lentiviral vectors LeGO (<http://www.lentigo-vectors.de/vectors.htm>) with EGFP and human reprogramming transcription factors: OCT4, SOX2, C-MYC and KLF4, courtesy of Dr. Sergei L. Kiselev, Moscow. We used previously the published protocol (Menzorov et al., 2015) with minor modifications. Lentiviruses were produced in the Phoenix cell line using Lipofectamine 3000 (Thermo Fisher Scientific, USA). The multiplicity of infection was estimated as 5.1 using EGFP lentiviral vector (Beklemisheva, Menzorov, 2018). Fibroblasts ( $3 \times 10^5$  cells,  $30 \times 10^3$  cells/cm<sup>2</sup>) plated the day before were transduced with viruses containing four reprogramming transcription factors: 50 % virus supernatant, 50 % fibroblast culture medium without antibiotics with heat-inactivated FBS, and 10 µg/ml Polybrene. Transduction was performed two consecutive days. Cells were passaged onto a 6 cm cell culture dish coated with 0.1 % gelatin on mouse strain CD-1 feeder cells on day 5. From day 6 we used iPS cell culture medium: a-MEM supplemented with 20 % ES cell qualified FBS, 1x MEM non-essential amino acids solution, 1x GlutaMAX supplement, 0.1 mM 2-mercaptoethanol, 1x penicillin-streptomycin, and 10 ng/ml bFGF recombinant human protein (Thermo Fisher Scientific, USA). From day 6 until day 12 the medium was changed once in two days with the addition of 1 mM valproic acid (Sigma-Aldrich, USA). On day 23 colonies were picked up and expanded on the feeder. The passage was performed with 0.25 % Trypsin-EDTA (Thermo Fisher Scientific, USA). All cell cultures were maintained at 37°C and 5 % CO<sub>2</sub>.

iPS cell derivation was performed at the Collective Center of ICG SB RAS "Collection of Pluripotent Human and Mammalian Cell Cultures for Biological and Biomedical Research" (<http://ckp.icgen.ru/cells/>; [http://www.biores.cytogen.ru/icg\\_sb\\_ras\\_cell/](http://www.biores.cytogen.ru/icg_sb_ras_cell/)).

All animal studies were undertaken with prior approval from the Interinstitutional Bioethical Committee of the Institute of Cytology and Genetics SB RAS and the Ethics Committee on Animal and Human Research of the Institute of Molecular and Cellular Biology SB RAS, Russia (order No. 32 of 5 May 2017).

### **Cytogenetic analysis**

Cytogenetic analysis for fibroblasts was carried out on passage 7 and for iPS-like cell line on passage 9. The preparation of metaphase chromosomes from fibroblasts was performed as previously described (Yang et al., 1999; Graphodatsky et al., 2000; Graphodatsky et al., 2001). GTG-banding of metaphase chromosomes was done according to a previously published protocol (Seabright, 1971). For each cell line, an average of 50 metaphase plates were analyzed using the VideoTest system (Saint Petersburg, Russia) with a CCD camera (Jenoptik, Germany) mounted on a Zeiss microscope Axioscope 2 (Zeiss, Germany). Metaphase spreads images were edited in Corel Paint Shop Pro Photo X2 (Corel, Canada). Chromosomes of the ringed seal (*P. hispida*) were arranged according to the nomenclature (Graphodatsky et al., 2020).

### **iPS cell differentiation**

Differentiation into embryoid bodies (EBs) was performed in EB differentiation medium, the same composition as iPS cell culture medium but with 20 % FBS instead of ES qualified FBS and without bFGF. Cells were passaged into 1 % agarose coated cell culture plates. The medium was changed every second day for 31 days. On day 5 part of EBs were plated onto 0.1 % gelatin-coated cell culture plates for osteocyte and adipocyte differentiation. Osteocyte differentiation was carried out in EB differentiation medium; adipocyte differentiation in a similar adipocyte differentiation medium supplemented with 10 % knockout serum replacement (KSR) (Thermo Fisher Scientific, USA) instead of FBS from day 6.

### **Cytochemical staining**

Cells were fixed by 4 % paraformaldehyde for 20 min and washed with PBS. Adipocytes were stained with 0.7 % Sudan black B in propylene glycol, washed twice with 0.85 % propylene glycol, and washed multiple times with PBS. Calcification was shown by staining with alizarin red (Sigma-Aldrich) according to the manufacturer's recommendations. Staining was analyzed on Zeiss Observer.Z1 fluorescent microscope with AxioCam HRm 3 CCD-camera (Zeiss, Germany). Digital images were analyzed using the ZEN 2 starter (Zeiss, Germany) software.

### **DNA isolation**

Genomic DNA was isolated from cells using a PCR buffer with nonionic detergents (PBND), which was adapted from a protocol from Perkin Elmer Cetus (Higuchi, 1989).

### **RNA isolation and cDNA synthesis**

RNA was isolated using Aurum Total RNA mini kit (Bio-Rad, USA). Genomic DNA was removed using *DNaseI* (Fermentas, USA), 0.4 micrograms of total RNA was used for cDNA synthesis by First Strand cDNA Synthesis kit (Thermo Fisher Scientific, USA).

### **Primer design and PCR**

We used Primer-BLAST software (Ye et al., 2012) to design primers for mink *Rex1* and canine *Cdx2*: qNvRex1F 5'-AAA GCG TTT TCC ACA CCC CT-3', qNvRex1R 5'-CTC CTT GTC CAT GGT CCT CG-3', CfCdx2E1F 5'-GGA ACC TGT GCG AGT GGA TG-3', and CfCdx2E3R 5'-TTC CTT TCC TTG GCT CTG CG-3'.

We used previously published primer sequences for human-specific *KLF4* transgene (Mathew et al., 2010), mycoplasma detection (Choppa et al., 1998), mink *Hprt1* (Rouvinen-Watt et al., 2012), mink *Oct4* (Menzorov et al., 2015), and human *AFP* (Huangfu et al., 2008).

PCR was performed using BioMaster HS-Taq PCR-Color (2x) (Biolabmix, Russia) in 10  $\mu$ L reaction volume.

## **RESULTS**

### **iPS cell derivation and differentiation**

Experiment outline is shown in Figure 1, A. To produce iPS cells from primary ringed seal fibroblasts we used the following human reprogramming transcription factors: OCT4, SOX2, C-MYC, and KLF4. On day 26 after transduction, there were colonies with different morphology. Two colonies were picked up and expanded. One of them, iPHIS1, gave rise to cells with epithelial-like morphology (Fig. 1, B) that grew in a monolayer. Overgrown culture formed “bubbles” (Fig. 1, C) that later detached and floated as cyst-like structures resembling canine embryos (Hayes et al., 2008).

We attempted to differentiate iPHIS1 cells into derivatives of the three germ layers. Solid and cyst-like EBs were formed after passage to non-adhesive culture plates (Fig. 1, E). EB gene expression analysis is described below. Adipo- and osteogenic differentiation was successfully performed in the adipo- and EB differentiation media, respectively (Fig. 1, H, I). Interestingly, there was some calcification in the adipocyte differentiation medium (data not shown).

### **Cytogenetic analysis**

We performed the cytogenetic analysis of the ringed seal primary fibroblasts and iPHIS1 cells. Fibroblasts had 32 chromosomes, XX (n = 50) (Fig. 1, D) with 15.7 % polyploid cells. iPHIS1

cells (n = 50) had 32 (n = 47), 33 (n = 2), and 35 (n = 1) chromosomes, with 5 % polyploid cells. We had not revealed a difference in the pattern of GTG-banding between karyotypes of the primary fibroblasts and iPHIS1 cells. We conclude that the iPHIS1 karyotype is stable.

### **iPS cell and EB gene expression**

We were able to show expression of one of the key pluripotency markers *Rex1* (*Zfp42*) in iPHIS1 cells (Fig. 1, F), as well as *Oct4*. There also was a weak expression of *Nanog* (data not shown). We were not able to determine its expression level as qualitative PCR was performed, though we previously showed a very low level of *Nanog* expression in American mink pluripotent stem cells (Menzorov et al., 2015). Importantly, we were not able to amplify human *KLF4* transgene (Fig. 1, F), it suggests transgene silencing.

Analysis of EBs revealed the expression of *AFP*, endoderm marker, and *Cdx2*, trophoblast marker (Fig. 1, G). Ectoderm marker, *Tubb3*, was expressed in both iPHIS1 cells and embryoid bodies (data not shown).

Ringed seal fibroblasts and iPHIS1 cells were negative for *Mycoplasma* contamination.

### **DISCUSSION**

Transcription factors Oct4, Sox2, c-Myc, and Klf4 were used to generate iPS cells from a variety of Carnivora species. We decided to produce iPS cells from the ringed seal (*P. hispida*), the basal representative of the suborder Caniformia. Only one cell line expressed pluripotency marker genes *Rex1* and *Oct4*. Its morphology differed from mouse, human, and mink iPS cells (Fig. 1, B). Similar to American mink iPS cells, overgrown cell culture formed “bubbles” (Fig. 1, C) that later unfastened and floated as cysts. We differentiated iPHIS1 cells into EBs to analyze differentiation potential (Fig. 1, E). Cyst-like EBs resembled canine embryos (Hayes et al., 2008), expressed endoderm marker *AFP* and trophoblast marker *Cdx2* (Fig. 1, G). EBs seated on gelatin differentiated into mesoderm derivatives: adipo- and osteocytes. We were not able to show ectoderm differentiation, as *Tubb3* was expressed in both iPHIS1 and EBs. The presence of *Tubb3* transcript in undifferentiated pluripotent stem cells is in accordance with our previous data on mouse ES cells (Menzorov et al., 2019). Quantitative PCR or another ectoderm marker and/or different differentiation conditions should solve this problem.

Another potential sign of the iPHIS1 pluripotency is human *KLF4* transgene silencing (Fig. 1, F). Viral transgene silencing in pluripotent stem cells is a well-known phenomenon (Maherali et al., 2007; Wernig et al., 2007; Okita et al., 2007).

One of the interesting properties of iPHIS1 cell line is its ability to differentiate into trophoblast, as shown by *Cdx2* expression. Trophoblast differentiation was shown for human and mouse primed pluripotent stem cells (Xu et al., 2002; Kojima et al., 2014) and canine iPS cells (Luo et al., 2011; Wilcox et al., 2008). Differentiation into trophoblast suggests a primed pluripotency state of iPHIS1.

We used bFGF to derive and propagate the iPHIS1 cell line. Colonies with iPS-like morphology were not formed in the media supplemented with LIF or combination of LIF and bFGF. Canine pluripotent stem cells were produced with LIF (Luo et al., 2011; Baird et al., 2015; Lee et al., 2011; Vaags et al., 2009) or LIF and bFGF (Shimada et al., 2010; Whitworth et al., 2012; Koh et al., 2012; Tsukamoto et al., 2018). Also, cells were later cultured in the presence of LIF only (Whitworth et al., 2012). Other Carnivora species pluripotent cells include snow leopard (Verma et al., 2012), Bengal tiger, serval, jaguar, and cat (Verma et al., 2013; Dutton et al., 2019) cultured with LIF; cat cultured with LIF and bFGF (Gómez et al., 2010), and ferret cultured with bFGF (Gao et al., 2020). Interestingly, cat iPS cells required species-specific feline bFGF (Dutton et al., 2019). We generated American mink iPS cells without LIF or bFGF supplementation (Menzorov et al., 2015), though inactivated mouse embryonic fibroblasts secrete both growth factors. Different requirements of the iPS cell culture of various species indicate that different signaling pathways are activated. It leads to different pluripotency states, naïve, primed, or other, as additional distinct pluripotency states had been described recently. More high-quality transcriptome data may facilitate the distinction between various pluripotency states in different species.

Pluripotent stem cells of different pluripotency status also have distinctive morphology. American mink pluripotent stem cells form colonies unlike mouse and human naïve and primed cells (Menzorov et al., 2015). Ringed seal iPHIS1 cell line is at least multipotent and also has different morphology.

We produced only one ringed seal iPS-like cell line, other experiments were not successful. Suboptimal culture conditions might be the main reason. ES cell qualified FBS batch was only tested for mouse ES cell growth support. Also, growth factor origin may be important. We used human bFGF, and for cat iPS cells only species-specific feline LIF was able to support pluripotency (Dutton et al., 2019). Also, growth factor concentration is an important factor. Ferret iPS cells were derived with human bFGF, but in 10x concentration compared to human iPS cells and this study (Gao et al., 2020). Different researchers used small molecules to enhance iPS cell derivation efficiency. In our experience, 2i inhibitors (PD0325901 and

CHIR99021) and TGH- $\beta$  antagonist A83-01 caused substantial fibroblast death, thus we were not able to use them. Another way to increase efficiency is to use different transgenes and/or delivery vectors. We have recently shown that Sendai virus-based vector transgene expression level is superior to lentiviruses (Beklemisheva, Menzorov, 2018). Also, another set of reprogramming factors was successfully used to reprogram not only human, but ferret fibroblasts to pluripotency, OCT4, SOX2, KLF4, L-MYC, LIN28A, mp53DD, and 160 oriP/EBNA-1 (Gao et al., 2020).

## **CONCLUSIONS**

We produced tri-potent ringed seal iPS-like cells, capable of differentiation into endoderm, mesoderm, and trophoblast. We need to achieve ectoderm differentiation to prove pluripotency.

## **ACKNOWLEDGMENT**

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## **CONFLICT OF INTERESTS**

The authors declare that there is no conflict of interest.

## **AUTHOR CONTRIBUTIONS**

VRB obtained ringed seal fibroblasts and performed cytogenetic analysis. AGM produced and differentiated iPS cells, performed gene expression analysis. AGM carried out interpretations of the data and project coordination; AGM did most of the writing with contributions from VRB. All authors read and approved the final manuscript.

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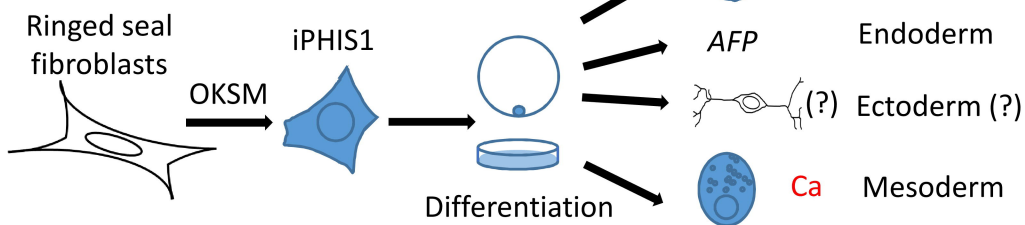
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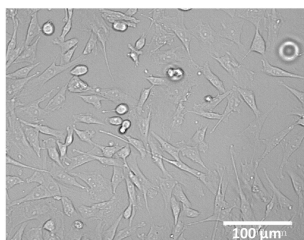
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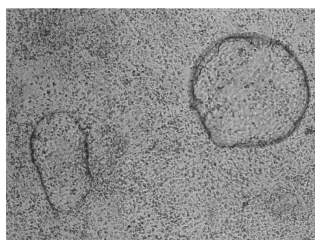
### A Experiment outline



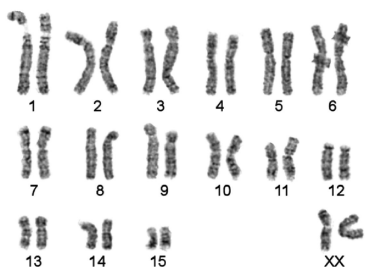
### B iPHIS1 on gelatin without feeder



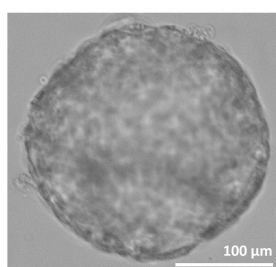
### C Overgrown iPHIS1 forms “bubbles”, similar to mink iPS cells



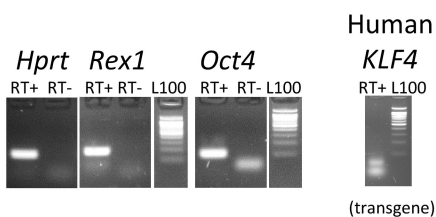
### D iPHIS1 karyotype



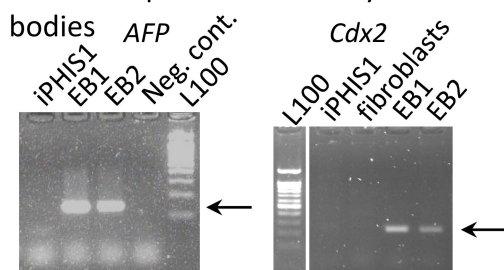
### E iPHIS1 embryoid body on day 17



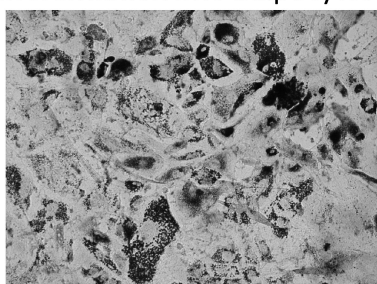
### F iPHIS1 gene expression



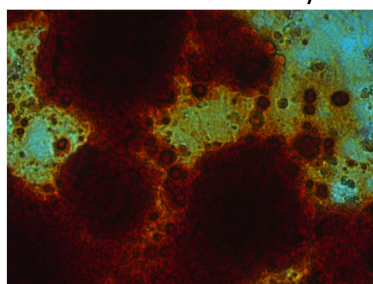
### G Gene expression in embryoid bodies



### H Differentiation into adipocytes



### I Differentiation into osteocytes



**Figure 1.** Derivation and characterization of ringed seal tripotent induced pluripotent stem-like cells