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16

17 **Abstract**

18 Leukemia inhibitory factor (LIF) is a cytokine which is essential for oocyte and

1 embryo development, embryonic stem cell, and induced pluripotent stem cell
2 maintenance. Leukemia inhibitory factor improves the maturation of oocytes in the
3 human and the mouse. However, feline LIF (fLIF) cloning and effects on oocytes during
4 IVM have not been reported. Thus, we cloned complete cDNA of feline LIF and
5 examined its biological activity and effects on oocytes during IVM in the domestic cat.
6 The amino-acid sequence of fLIF revealed a homology of 81% or 92% with that of
7 mouse or human. The fLIF produced by pCold[®] TF DNA in *Escherichia coli* was readily
8 soluble and after purification showed bioactivity in maintaining the undifferentiated
9 state of mouse embryonic stem cells and enhancing the proliferation of human
10 erythrocyte leukemia cells. Furthermore, 10- and 100- ng/mL fLIF induced cumulus
11 expansion with or without FSH and EGF ($P < 0.05$). The rate of metaphase II oocytes
12 was also improved with 100 ng/mL fLIF ($P < 0.05$). We therefore confirmed the
13 successful production for the first time of biologically active fLIF and revealed its
14 effects on oocytes during IVM in the domestic cat. Feline LIF will further improve
15 reproduction and stem cell research in the feline family.

16

17 *Keywords:* Recombinant protein; Feline; IVM; Leukemia inhibitory factor; Cumulus
18 expansion; Nuclear maturation

1 **1. Introduction**

2

3 Leukemia inhibitory factor (LIF), belonging to the interleukin-6 (IL-6) family, was
4 discovered as a cytokine that induced the differentiation of mouse myeloid leukemia
5 cells [1]. Subsequent studies have reported various biological effects of LIF on a large
6 range of cell types, including monocytes, megakaryocytes, neuronal cells, osteoblasts,
7 oocytes, hepatocytes, adipocytes, embryonic stem cells (ESCs), and induced pluripotent
8 stem cells (iPSCs), which have receptors for LIF [2-7]. Most of effects exerted by LIF
9 are similar to those of IL-6 [8].

10 Leukemia inhibitory factor also has further effects, not shared with IL-6. LIF
11 supplementation during IVM of human and mouse oocytes induced cumulus expansion
12 and in mouse enhanced cleavage rate at two-cell stage as well as birth delivery [9].
13 Furthermore, LIF prevented naive ESCs and iPSCs from differentiating, which have
14 higher abilities to proliferate and differentiate [10]. These specific effects of LIF help us
15 to improve reproductive techniques and generate naive ESCs and iPSCs, which could be
16 used in the preservation of wild animals [11-13], regenerative medicine [14, 15] and
17 treatment of transmissible diseases [16, 17].

18 Recombinant human and mouse LIF have been produced by utilizing *Esheria coli*,

1 yeast and COS cells [18-20]. Protein expression in *E coli* is superior to that with yeast
2 or COS cells in ease of manufacturing and quantity of production [21]. Some drawbacks
3 remain, such as the dependence of the efficiency of collecting protein on its behavior
4 and size, because of solubility and inclusion bodies, and loss of activity caused by few
5 posttranslational modifications [22-24]. The protein expression vector pCold[®] TF DNA
6 possesses a trigger factor which is known to improve the solubility of synthetic proteins
7 [25]. In previous studies, LIF was produced by utilizing *E coli*, which showed that LIF
8 may still be biologically active without posttranslational modifications [26].

9 During IVC of bovine oocytes, human LIF (hLIF) reduced the blastocyst rate, though
10 the sequence of bovine LIF revealed a homology of 89.1% with that of hLIF [27].
11 Human and mouse LIF, whose aminoacid homology with that of porcine was 88% and
12 79%, respectively, failed to generate porcine ESCs and iPSCs, however, the cultivation
13 with porcine LIF could establish porcine iPSCs [28, 29]. Although, ESCs of domestic
14 cats were generated with mouse LIF (mLIF), they could not be maintained in an
15 undifferentiated state for a prolonged period or differentiate into three germ layers [30,
16 31]. The iPSCs of endangered felids were also established with mLIF, whose transgene
17 expressions were sustained [32, 33]. These studies suggest that heterospecific LIF
18 should be carefully used instead of homospecific LIF. Biologically active LIF has been

1 produced only in the human [19], the mouse [1], the rat [34], the possum [35] and the
2 chicken [36], but not in the domestic cat which is a good research model for wild felids
3 faced with extinction. Thus, the purpose of this study was to clone the complete cDNA
4 of feline LIF (fLIF), produce biologically active fLIF in *E coli* with effortless
5 purification at a higher efficiency, and examine its activity in relation to mouse ESCs,
6 human erythroleukemia cells, and domestic cat oocytes.

7

8 **2. Materials and methods**

9

10 2.1. Culture of feline embryonic fibroblasts

11 We collected domestic short-haired cat fetuses from a cat that underwent
12 ovariectomy at a local veterinary clinic at day 30 of pregnancy. The cat was
13 privately owned, and the owner's consent was obtained before the collection of the
14 fetuses. No cats were bred and operated upon specifically for this study. The fetuses
15 were washed with PBS, and their heads and livers were removed. The fetuses were then
16 cut into small pieces and cultured in Dulbecco's modified Eagle medium (Life
17 Technologies, San Diego, CA, USA) supplemented with 10% (v:v) fetal bovine serum
18 (FBS; PAA Laboratories, Pasching, Australia), 100-IU/mL penicillin and 100- μ g/mL

1 streptomycin (Sigma-Aldrich, St. Louis, MO, USA), and 2-mM L-glutamine
2 (Sigma-Aldrich) at 37°C under 5% CO₂ in humidified air. The adherent cells were
3 passaged with 0.25% trypsin (Sigma-Aldrich). After three passages, the spindle cells
4 were used as feline embryonic fibroblasts for RNA isolation.

5 deoxyribonucleotides

6 2.2. Cloning of fLIF

7 Total RNA was extracted from cat embryonic fibroblasts with RNeasy Mini
8 (QIAGEN, Hilden, Germany) according to the manufacturer's protocol. Total RNA was
9 used in a reverse transcription (RT) reaction with a reactive solution (20 µL) containing
10 0.5-mM deoxyribonucleotides mixture (TOYOBO, Osaka, Japan), 2.5-µM random
11 mer (TOYOBO) and 5-U/µL ReverTra Ace (TOYOBO) in the appropriate RT buffer.
12 The RT solution was incubated at 30°C for 10 min, at 42°C for 50 min, then at 99°C for
13 5 min.

14 Following the comparison of the nucleotide sequences of LIF in human
15 (NM_002309) and rat (NM_022196), primers for cloning were designed with reference
16 to common parts between both of them. Polymerase chain reaction (PCR) was carried
17 out in 20-µL KOD buffer containing 0.5-µL RT products, 0.2-mM dNTPs, 0.02-U/µL
18 KOD plus (TOYOBO), 1- mM MgSO₄ (TOYOBO), and 0.5 µM of a pair of primers

1 (Table 1). The thermocycling protocol comprised preincubation at 94°C for 2 min and
2 35 cycles at 94°C for 15 seconds, 60°C for 30 seconds, 68°C for 1 minutes, and finally
3 68°C for 2 minutes. The chain reaction product was inserted into pGEM[®]-T Easy Vector
4 (Promega, Madison, WI, USA) and sequenced by Sigma-Aldrich, Japan, using a
5 BigDye[®] Terminator version 3.1 Cycle Sequencing Kit (Life Technologies) with a pair
6 of primers (Table 1) and an auto sequencer (ABI 3130xl, Applied Biosystems). The
7 nucleotide sequence of fLIF cDNA was compared with sequences of hLIF
8 (NM_002309) and mLIF (NM_008501), and the coding region of mature protein of
9 fLIF was predicted. This was amplified with a forward primer containing a BamHI site
10 and a reverse primer containing HindIII site (Table 1) for subcloning and inserted into
11 pCold[®] TF DNA (Takara, Shiga, Japan).

12

13 2.3. Recombinant synthetic protein expression and purification

14 BL21 (DE3 strain; Nippon Gene) was transformed with reconstructed pCold[®] TF
15 DNA and positive clones were selected on LB agar plate with antibiotics (50-μg/mL
16 ampicillin; TOYOBO). The selected clone was incubated in LB medium with
17 antibiotics at 37°C until OD₆₀₀ reached 0.5. For recombinant synthetic protein
18 expression, isopropyl-β-D-thiogalactopyranoside (Wako, Osaka, Japan) was added to

1 cultures to a final concentration of 1 mM, and after stationary incubation at 15°C for 30
2 min, cultures were incubated at 15°C for 24 hours. Induced cells were centrifuged at
3 4°C for 10 min at 800 x g in an ice-cold tube, and resulting pellets were resuspended
4 with PBS. Resuspended cells were centrifuged at 4°C for 10 min at 800 x g, and pellets
5 were frozen at -80°C prior to protein purification.

6 Ice-cold PBS was added to the frozen pellets, and the mixture was sonicated on ice.
7 This sample was centrifuged, and supernatant was harvested and used for purification.
8 The supernatant was run through Ni-NTA Sepharose resin (Wako) which had been
9 equilibrated with PBS. Then, to remove non-specifically bound protein, Ni-NTA
10 Sepharose was washed with 10 column volumes of PBS. Bound proteins were harvested
11 with 10 column volumes of PBS for which the imidazole (Nacalai Tesque, Kyoto,
12 Japan) concentration was increasing in the series. The flow-through PBS were collected
13 and the protein concentration was determined by the Bradford method. Thereafter,
14 samples were digested using a Thrombin kit (Merck, Darmstadt, Germany) at 4°C for
15 24 h to remove His-tag and Trigger Factor and were analyzed with SDS-PAGE.

16

17 2.4. SDS-PAGE analysis

18 Samples of the elution fractions were vortexed with 2 µL of 5× loading buffer (Wako),

1 and then heated at 97°C for 3 minutes. These samples were analyzed by silver staining
2 after SDS-PAGE with 10% or 12% separating gel.

3

4 2.5. Biological activity assay of recombinant protein

5

6 *2.5.1 Ability to maintain mouse ESCs (TT2) in an undifferentiated state*

7 Mouse ESC line, TT2 (RIKEN Bioresource Center, Japan) was cultured on gelatin-
8 coated dishes and inactivated feeder cells in mouse ES medium comprising Dulbecco's
9 modified Eagle medium Nutrient Mixture F-12 HAM (Sigma-Aldrich) supplemented
10 with 20% (v:v) FBS (lot for ES culture; Life Technologies), 100-IU/mL penicillin and
11 100-μg/mL streptomycin, 2-mM L-glutamine, 0.1-mM MEM Non-Essential Amino
12 Acids (Life Technologies), 0.1-mM 2-mercaptoethanol (Sigma-Aldrich), 1-mM sodium
13 pyruvate, 0.075% (wt/vol) sodium bicarbonate (Life Technologies) and 1000-U/mL
14 mLIF (ESGRO; Millipore, Billerica, MA, USA) at 37°C under 5% CO₂ in humidified
15 air. Mouse embryonic fibroblasts were collected from fetuses at day 12 of pregnancy,
16 inactivated with mitomycin C (Kyowa Hakko Kirin, Tokyo, Japan) and used as feeder
17 cells. The protocol was approved by the animal experiment committee of the Graduate
18 School of Life and Environmental Sciences of Osaka Prefecture University. TT2 was

1 passaged with 0.25% trypsin every 2-3 days.

2 To assay the biological activity of fLIF produced in *E coli*, TT2 was cultured in
3 mouse ES medium with a series of increasing concentrations of mLIF, commercial hLIF
4 (Millipore), or fLIF (0.2 – 10 ng/ml). After five passages, the undifferentiated state of
5 TT2 colonies was analyzed by the total number of colonies and the rate of colonies
6 positive to alkaline phosphatase (ALP) stained with Alkaline Phosphatase Staining Kit
7 II (Stemgent, Cambridge, MA, USA) [37]. Colonies were classified as ALP positive
8 when the entire cells in the colony stained red. The number of colonies was counted in
9 three high-power fields at each experiment, and the experiment was repeated three
10 times.

11

12 *2.5.2 Ability to enhance proliferation of human erythroleukemia cells (TF-1)*

13 TF-1 cells (Health Protection Agency, London, UK) were maintained in RPMI 1640
14 medium (Nacalai Tesque) supplemented with 10% (v:v) FBS, 100-IU/mL penicillin and
15 100- μ g/mL streptomycin, 2-mM L-glutamine and 1-mM sodium pyruvate
16 (Sigma-Aldrich) at 37°C under 5% CO₂ in humidified air. Human
17 granulocyte-macrophage colony stimulating factor (GM-CSF; Millipore) was added
18 into the medium at a final concentration of 4 ng/mL every 48 h. TF-1 cells were

1 incubated without GM-CSF for 24 h before addition of a series of increasing dilution of
2 mLIF, hLIF or fLIF (twofold serial dilution) and reseeded into a 96-well plate at a
3 density of 2×10^4 cells/well. After 48 h incubation at 37°C under 5% CO₂ in humidified
4 air, the proliferation rate of TF-1 cells was quantified by
5 2-(2-methoxy-4-nitrophenyl)-3-(4-nitrophenyl)-5-(2,4-disulfophenyl)-2H tetrazolium in
6 a WST-8 colorimetric assay using a Cell Counting Kit-8 (DOJINDO) according to the
7 manufacturer's protocol [38, 39]. In summary, WST-8 solution was supplemented with
8 the medium at a dilution of 1:10. After 4-hour incubation at 37°C under 5% CO₂ in
9 humidified air, the absorbance was measured at 450 nm. The proliferation rate was
10 calculated as $[(\text{absorbance}_{450} \text{ sample} - \text{absorbance}_{450} \text{ control}) / (\text{absorbance}_{450} \text{ Max} -$
11 $\text{absorbance}_{450} \text{ control})] \times 100(\%)$.

12

13 2.6. Effect of fLIF on domestic cat oocytes matured *in vitro*

14 Ovaries from domestic short-haired cats were collected during ovariohysterectomy at
15 a local veterinary clinic. The cats were privately owned, and owner's consent was
16 obtained before ovaries collection. Cats were not specifically ovariohysterectomized for
17 this study. Collected ovaries were sliced with scissors in M-199 medium (Life
18 Technologies), and collected cumulus-oocyte complexes (COCs) were rinsed three

1 times in IVM medium consisting of M-199 medium supplemented with 0.4% (wt/vol)
2 BSA (Sigma), 10-IU/mL 17 β -estradiol (Sigma), 100- μ g/mL gentamycin (Sigma) and
3 137- μ g/mL sodium pyruvate. Immature oocytes with dark cytoplasm and two or more
4 layers of cumulus cells were randomly divided into each experimental group [40]. Then,
5 we added fLIF at each concentration (0 - 100 ng/mL) into the IVM medium to evaluate
6 the activity of fLIF on COCs. After 28-hour incubation of COCs in IVM medium at
7 38.5°C under 5% CO₂ in humidified air, the cumulus layer sizes were measured with a
8 micrometer. Furthermore, we incubated COCs in IVM medium with 0.02-IU/mL
9 Follistim[®] (human recombinant FSH, MSD, Tokyo, Japan) and 25 ng/mL human
10 recombinant EGF (Sigma) at 38.5°C under 5% CO₂ in humidified air for 28 hours to
11 examine whether the effects of LIF on cumulus expansion are shared with those of FSH
12 or EGF and measured the cumulus layers with a micrometer. The COCs incubated in the
13 IVM medium with FSH and EGF were pipetted gently many times to remove the
14 cumulus layers. Nude oocytes were collected, rinsed with PBS twice, and fixed in PBS
15 containing 3.7% paraformaldehyde (Merck) and 1% Triton-X100 (Merck) for 15
16 minutes and in PBS containing 0.3% polyvinylpyrrolidone (Sigma) for 15 minutes.
17 Fixed oocytes were placed on slides and mounted with 90% glycerol including 10-
18 μ g/mL bis-benzimide (Hoechst 33342, Sigma). The oocytes were examined with

1 fluorescence microscope and categorized into germinal vesicle (GV), germinal vesicle
2 breakdown (GVBD), metaphase I/anaphase/telophase I (MI), metaphase II (MII) stage,
3 parthenogenetic activation (PA), or unclassified.

4

5 2.7. Statistical Analyses

6

7 All experiments were repeated independently at least three times. The results
8 expressed as percentages were subjected to arcsine transformation and analyzed with
9 ANOVA, followed by Fisher's protected least significant difference post hoc analysis
10 (Stat View; Hulinks Inc., Tokyo, Japan). A value of $P < 0.05$ was taken as significant.

11

12 **3. Results**

13

14 3.1. Reconstruction of plasmid vectors

15 The cDNA of fLIF was composed of 609 base pairs (AB853322), whose nucleotide
16 sequence revealed homology of 99% with predicted fLIF mRNA (XM_003994784) (Fig.
17 1). Comparison of fLIF with mLIF or hLIF revealed a homology of 81% or 92% of the
18 aminoacid sequence of mature protein (Fig. 2). The sequence of mature protein of fLIF

1 was inserted into pCold[®] TF DNA.

2

3 3.2. Expression and purification of recombinant synthetic protein

4 After purification with Ni-NTA Sepharose, digestion by thrombin, and second
5 purification with Ni-NTA Sepharose, the extracted proteins were analyzed with
6 SDS-PAGE and silver staining (Fig. 3). The band seen around 70 kDa was divided into
7 two bands around 50 kDa and 20 kDa. The second purification of sample digested by
8 thrombin with Ni-NTA Sepharose led to the isolation of the band around 20 kDa. The
9 final yield of fLIF without the tag sequence was about 1.8 mg/L of fermentation broth.

10

11 3.3. Biological activity assay of recombinant protein

12

13 *3.3.1 Ability to maintain TT2 in an undifferentiated state*

14 The total number of colonies of TT2 was less following incubation without LIF than
15 with 0.2-, 1-, or 10-ng/mL LIF ($P < 0.01$). No differences were detected between species
16 (Fig. 4). These were fewer ALP-positive colonies of TT2 upon incubation without LIF
17 than with 1- or 10-ng/mL LIF ($P < 0.01$). These were fewer upon incubation with 0.2
18 ng/mL LIF than with 1 ng/mL or 10 ng/mL LIF ($P < 0.01$). No differences were detected

1 between species (Fig. 5).

2

3 *3.3.2 Ability to enhance TF-1 proliferation*

4 Proliferation of TF-1 was not seen with mLIF. However, hLIF and fLIF did enhance
5 the proliferation of TF-1 according to the LIF concentration (Fig. 6).

6

7 *3.4 Effect of fLIF on domestic cat oocytes matured in vitro*

8 A total of 117 COCs was incubated in IVM medium with fLIF. Cumulus expansion
9 was induced after incubation with 10- or 100-ng/mL fLIF ($P < 0.05$) (Fig. 7A).
10 Furthermore, a total of 414 COCs was cultured with FSH, EGF and fLIF. Cumulus
11 expansion was improved at higher concentration than 10-ng/mL fLIF ($P < 0.01$; Fig.
12 7B). The percentage of oocytes at MII stage were greater upon incubation with 100-
13 ng/mL fLIF than control group ($P < 0.05$; Table 2).

14

15 **4. Discussion**

16 We cloned here for the first time fLIF cDNA. Although the cloned fLIF cDNA
17 possesses a different aminoacid with predicted fLIF (XM_006938581), we considered
18 the difference as individual variability. The comparison of the aminoacid sequence of

1 fLIF with mLIF or hLIF revealed a homology of 81% or 92%, respectively.
2 Cross-species receptor affinity of LIF depends on aminoacids sequence and the
3 secondary structure [41]. The most important six aminoacids of hLIF that are used in
4 binding the hLIF receptor correspond to only five in the cat [42]. The secondary
5 structure of the fLIF receptor is necessary to determine the hLIF affinity for fLIF
6 receptor accurately. We predict, however, that hLIF does not bind the fLIF receptor
7 effectively [30-33].

8 The 70 kDa protein, which matches the predicted size of synthetic protein, was
9 isolated at a higher efficiency with Ni-NTA Sepharose. It was digested with thrombin
10 into a 50- and a 20-kDa protein, which again matches the predicted size of the tag
11 sequence and fLIF. This suggests that fLIF was indeed produced in *E coli* and that
12 isolation was carried out accurately. Although inclusion body formation occasionally
13 makes it difficult to isolate synthetic proteins [24], no such problems occurred with 1-
14 mM isopropyl- β -D-thiogalactopuranoside and reconstructed pCold[®]-TF DNA, because
15 of the Trigger Factor. Finally, the yield of fLIF was about 1.8 mg/L of fermentation
16 broth, demonstrating that soluble fLIF was collected and purified at a higher efficiency
17 [19, 24].

18 It has been reported that mouse and human LIF influenced the undifferentiated state

1 and proliferation of mouse ESCs by activating the signal transducer and activator of
2 transactivation (STAT) 3 signaling pathway [3, 8, 43]. TT2 is one of major mouse ESC
3 lines used in previous studies [44-46]; therefore, we utilized TT2 to assay the biological
4 activity of fLIF. It has been reported that there are no differences between mLIF and
5 hLIF affinity to mLIF receptor [41, 42]. Similarly, fLIF produced in this study
6 influenced the undifferentiated state and the proliferation of TT2, as indicated by
7 affinity to mLIF receptor. Although mouse ESCs are usually cultured with 10-ng/ml LIF
8 [44-46], the total number of colonies of TT2 during incubation with 0.2-ng/mL LIF did
9 not change. The number of ALP-positive colonies was significantly reduced; however,
10 we assume that these results arise because of lack of the desired LIF concentration
11 leading to loss of the undifferentiated state of TT2, which eventually caused TT2 to
12 slow the proliferation activity.

13 TF-1 was established from a human erythroleukemia patient; it is well known for its
14 characteristic of proliferation stimulated by various cytokines, such as GM-CSF or LIF,
15 following their concentration, and has been utilized in many experiments [21, 23]. In
16 the previous study, hybrid LIF of the human and mouse revealed that mLIF has no
17 affinity to hLIF receptor and at least six amino-acids are important for hLIF to bind
18 hLIF receptor [41]. Moreover, the secondary structure models of LIF and its receptor

1 were made to explain the binding interaction of human and mouse LIF [42]. In the
2 present study, although hLIF and fLIF demonstrated the significant effects of
3 enhancement of TF1 proliferation, mLIF did not possess any such activity. These results
4 therefore indicate that fLIF possesses the similar characteristics to hLIF and mLIF and
5 will help us to establish naive ESCs and iPSCs, which have higher abilities to
6 proliferate and differentiate.

7 The addition of fLIF significantly induced oocyte cumulus expansion regardless of
8 FSH and EGF. The effects of LIF on the cumulus expansion have been reported in
9 porcine [47], mouse, and human [9]. Previous studies have revealed that both FSH and
10 EGF induce cumulus expansion through mitogen-activated protein kinases (MAPK)
11 signaling pathway in the mouse [48]. On the other hand, LIF has effects on not only
12 MAPK but also STAT signaling pathways [8]. The activation of STAT signaling
13 pathway also induces the cumulus expansion in the porcine [47]. These studies suggest
14 that the supplementation of fLIF into IVM medium activates MAPK and STAT
15 signaling pathways in cumulus cells and enhances cumulus expansion regardless of the
16 addition of FSH and EGF.

17 The effects of LIF on nuclear maturation of oocytes have been reported in the mouse
18 [49], the porcine [47], and the bovine [50]. In the present study, the addition of fLIF

1 significantly increased the rate of oocytes at MII stage with FSH and EGF, which agrees
2 with previous studies. This result may come from two sources, cumulus expansion and
3 direct effects of fLIF on oocytes. Previous studies have revealed that cumulus expansion
4 is correlated with loss of gap junctions between cumulus cells, and leads to nuclear
5 maturation of oocytes [51, 52]. On the other hand, the activation of STAT signaling
6 pathways directly improves nuclear maturation in bovine oocytes [50]. These studies
7 indicate that fLIF has direct and indirect effects on IVM of oocytes in the domestic cat.
8 To evaluate if the beneficial effect of feline LIF on nuclear maturation and cumulus
9 expansion is reflected on embryonic developmental competence, additional studies are
10 required.

11 In the present study, we have cloned fLIF cDNA and also produced fLIF in *E coli*
12 which is biologically active as LIF in relation to TT2, TF-1 and domestic cat oocytes.
13 These results will improve reproduction and stem cell research in the feline family.

14

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3

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1 Table 1

2 Primers for feline leukemia inhibitory factor (LIF).

Primer	Sequence (5'-3')	Appliance
Human LIF36-56	CAGCCCATAATGAAGGTCTTG	Cloning
Human LIF653-633	CTAGAAGGCCTGGGCCAACAC	
Cat-LIF-for-BamHI	AGGGATCCAGCCCCCTTCCTATCACCCC	Subcloning
Cat-LIF-rev-HindIII	ACAAGCTTGAAGGCCTGGGCCAACACAG	
Universal primer T7	TAATACGACTCACTATAGG	Sequencing
Universal primer SP6	TATTTAGGTGACACTATAG	pGEM [®] T Easy
pCold-TF-F1	CCACTTTCAACGAGCTGATG	Sequencing
pCold-TF-R	GGCAGGGATCTTAGATTCTG	pCold [®] TF

3 Primers for subcloning were added BamHI or HindIII site (underlined).

1 Table 2
 2 The effect of feline leukemia inhibitory factor (fLIF) on nuclear maturation of oocytes
 3 in the domestic cat.

		Percentage of oocytes at the stage of					
fLIF (ng/mL)	No. of oocytes examined	GV	GVBD	MI	MII	PA	unclassified
0	108	6.6 ± 2.4	14.1 ± 3.4	27.0 ± 6.3	48.7 ± 4.2 ^a	0.7 ± 0.7	2.9 ± 2.0
1	107	3.3 ± 1.6	18.6 ± 4.0	18.8 ± 4.4	54.4 ± 5.9 ^{ab}	0.0 ± 0.0	4.9 ± 1.9
10	95	3.1 ± 1.6	15.8 ± 5.0	16.9 ± 3.9	64.1 ± 5.9 ^{ab}	0.0 ± 0.0	0.0 ± 0.0
100	104	2.6 ± 1.7	10.8 ± 2.4	14.4 ± 4.5	72.2 ± 4.9 ^b	0.0 ± 0.0	0.0 ± 0.0

4 The oocytes were categorized into germinal vesicle (GV), germinal vesicle breakdown
 5 (GVBD), metaphase I/anaphase/telophase I (MI), metaphase II (MII), parthenogenetic
 6 activation (PA) or unclassified. Values present the mean ± SEM of eight independent
 7 experiments and more than eight cumulus-oocyte complexes were used in each
 8 experiment. Different superscripts indicate significant differences (P < 0.05).
 9

1 **Figure legends**

2

3 Fig. 1. Nucleotide and aminoacid sequence of feline leukemia inhibitory factor (fLIF)
4 cDNA. Signal sequence is underlined and the termination codon is marked with an
5 asterisk.

6

7 Fig. 2. Aminoacid sequence of mature protein of feline leukemia inhibitory factor (fLIF).
8 A comparison of fLIF with human LIF (hLIF) and mouse LIF (mLIF) revealed that they
9 have a so homologous sequence of amino-acids of mature protein (81% and 92%).
10 However, most important aminoacids of hLIF to bind the hLIF receptor (*) have
11 difference. The predicted fLIF possesses not L but V as 176th aminoacid.

12

13 Fig. 3. SDS-PAGE analysis of proteins. The synthetic protein (about 70 kDa) was
14 digested with thrombin and feline leukemia inhibitory factor (fLIF) was isolated by the
15 second purification with Ni-NTA Sepharose. BT, synthetics protein before thrombin
16 digestion; M, molecular weight marker; SP, protein after the second purification by
17 Ni-NTA Sepharose

18

1 Fig. 4. The alkaline phosphatase-positive colonies of TT2 with leukemia inhibitory
2 factor (LIF). (A) TT2 cultured without LIF. TT2 cultured with 10- ng/mL LIF of human
3 (B), mouse (C) or cat (D). Mouse embryonic stem cells are recommended to culture
4 with 10-ng/mL mouse LIF. Scale bars = 100 μ m

5

6 Fig. 5. The effect of leukemia inhibitory factor (LIF) on maintenance of undifferentiated
7 state of TT2. (A) A comparison of proliferation of mouse embryonic stem cells (ESCs)
8 with different doses of human, mouse and feline LIF. The total colony number was
9 counted after 5 passages. (B) A comparison of alkaline phosphatase (ALP)-positive rate
10 of mouse ESCs with different doses of human, mouse and feline LIF. ALP-positive
11 colonies were detected by ALP Staining Kit II. Values present the mean \pm SEM of three
12 independent experiments. Values with different letters differ by $P < 0.01$.

13

14 Fig. 6. The effect of leukemia inhibitory factor (LIF) on proliferation of TF-1. A
15 comparison of growth rate of TF-1 cells with different doses of human, mouse and
16 feline LIF after 48-hour incubation. Values present the mean \pm SEM of three
17 independent experiments.

18

1 Fig. 7. The effect of feline leukemia inhibitory factor (fLIF) on cumulus expansion in
2 the domestic cat. (A) A comparison of cumulus expansion with different doses of fLIF
3 without FSH and EGF. Four independent experiments were done and more than three
4 cumulus-oocyte complexes (COCs) were used in each experiment. (B) Cumulus
5 expansion incubated with FSH, EGF and fLIF. Eight independent experiments were
6 done and more than eight COCs were used in each experiment. Values present the mean
7 \pm SEM Values with different letters differ by $P < 0.01$.

ATGAAGGTCTTGGCGGCAGGAGTCGTGCCCTGCTGCTGGTTCTGCACTGGAAACATGGG
M K V L A A G V V P L L L V L H W K H G

GCGGGGAGCCCCCTTCCTATCACCCCTGTCAACGCCACCTGTGCCACACGCCACCCATGT
A G S P L P I T P V N A T C A T R H P C

CACAGCAACCTCATGAACCAGATCAGGAACCAACTGGCGCAGCTCAATGGCAGTGCCAAT
H S N L M N Q I R N Q L A Q L N G S A N

GCCCTCTTTATTCTCTATTACACGGCCCAGGGGAGCCGTTCCCCAACAACTGGACAAA
A L F I L Y Y T A Q G E P F P N N L D K

CTGTGCGGCCCAACGTGACGGACTTCCCGCCATTCCATGCCAACGGCACAGAGAAGACC
L C G P N V T D F P P F H A N G T E K T

CGGTTAGTGGAGCTGTACCGCATCATCGCTTACCTTGGTGCTCCCTGGGCAACATCACC
R L V E L Y R I I A Y L G A S L G N I T

CGGGACCAGAAGGTCTCAATCCCAATGCCCTCAGCCTCCACAGCAAACCTGAACGCCACT
R D Q K V L N P N A L S L H S K L N A T

GCAGACATCATGCGGGCCTCCTCAGCAACGTGCTTTGCCGCTGTGTAACAAGTATCAC
A D I M R G L L S N V L C R L C N K Y H

GTGGCCACGTGGACGTGGCCTATGGCCCTGACACCTCAGGCAAGGACGTCTTTCAGAAG
V A H V D V A Y G P D T S G K D V F Q K

AAGAAGCTGGGCTGTCAGCTCCTGGGGAAGTATAAACAGGTCATTGCTGTGTTGGCCCAG
K K L G C Q L L G K Y K Q V I A V L A Q

GCCTTCTAG
A F *

Fig. 1.

fLIF	SPLPITPVNATCATRHPCHS	NLMNQIRNQLAQLNGSANAL	FILYYTAQGEFPNNDKLC	*
hLIF		I N S		
mLIF		I G K	S VE	
fLIF	GPNVTDFPPFHANGTEKTRL	VELYRI IAYLGASLGNITRD	QKVLNPNALSLHSKLNATAD	* **
hLIF		AK VV T	I S	
mLIF	A M S G	K MV S T	T V QV I	
fLIF	IMRGLLSNVLCRLCNKYHVA	HVDVAYGPDTSQKDVFKKK	LGCQLLGKYKQVIAVLAQAF	* *
hLIF	L	S G T	I	
mLIF	V	R G PPV H D EA R	T S VV	

Fig. 2.

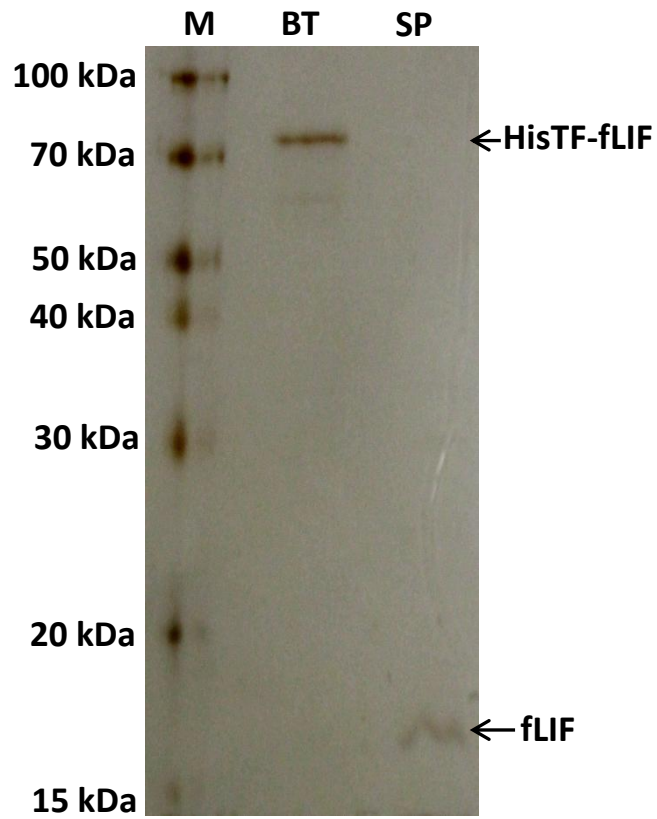


Fig. 3.

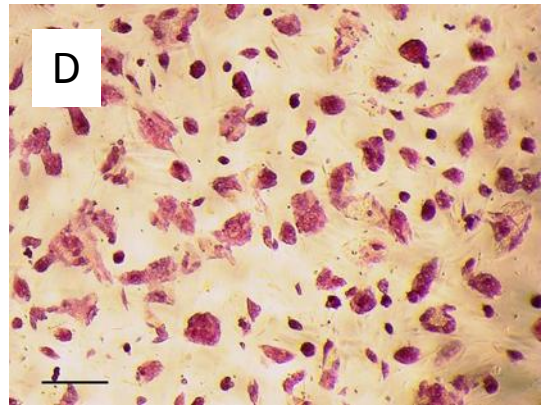
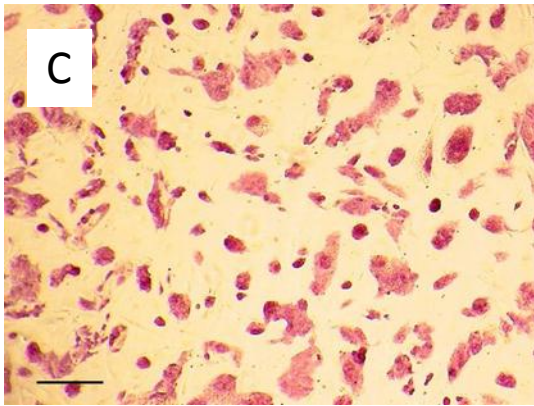
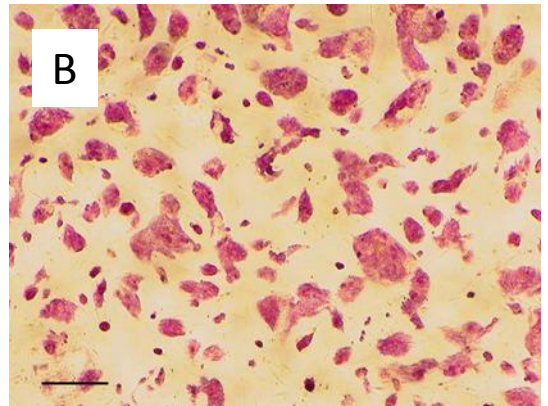
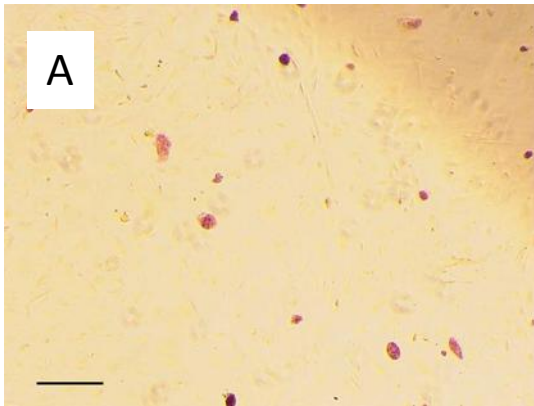


Fig. 4.

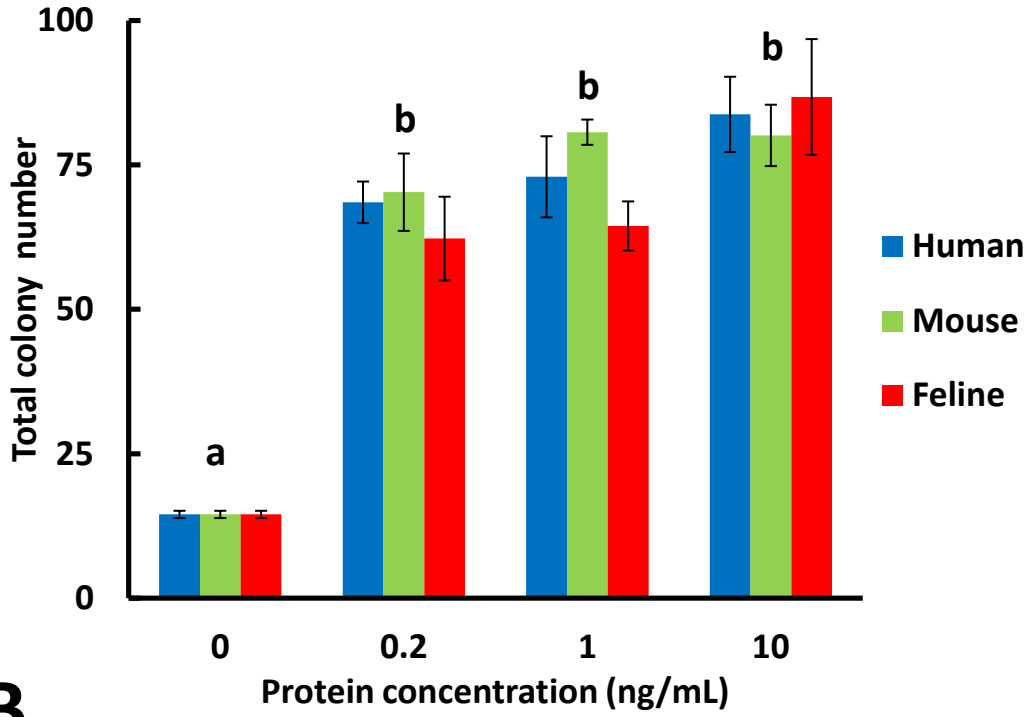
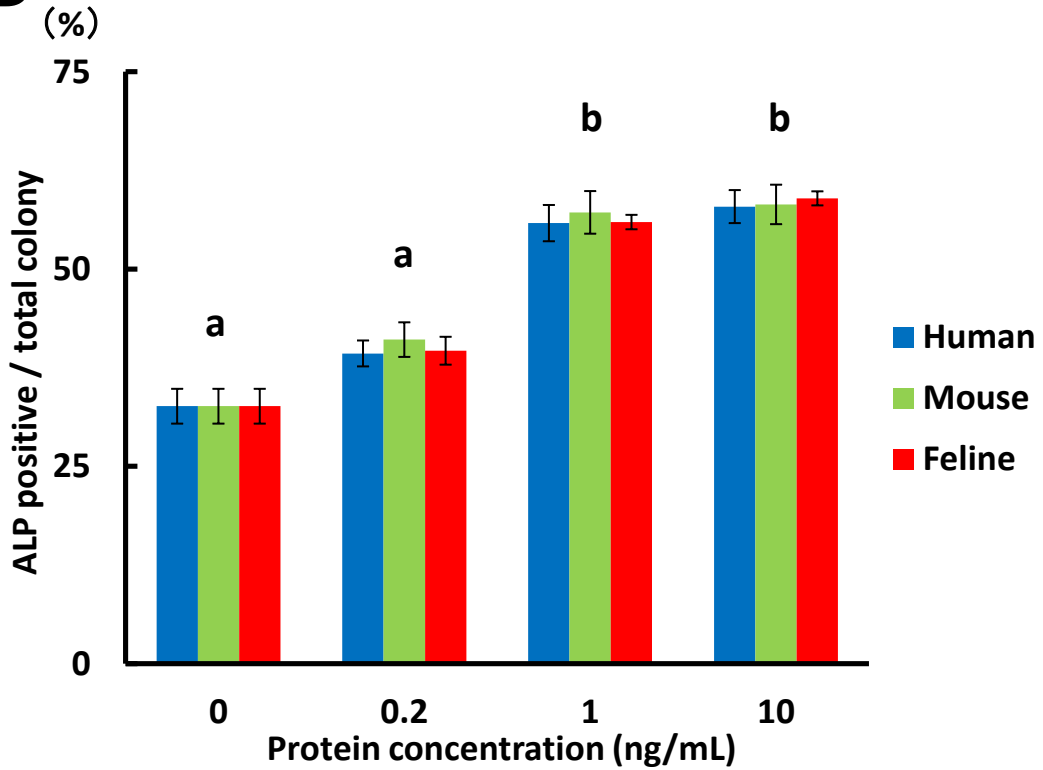
A**B**

Fig. 5.

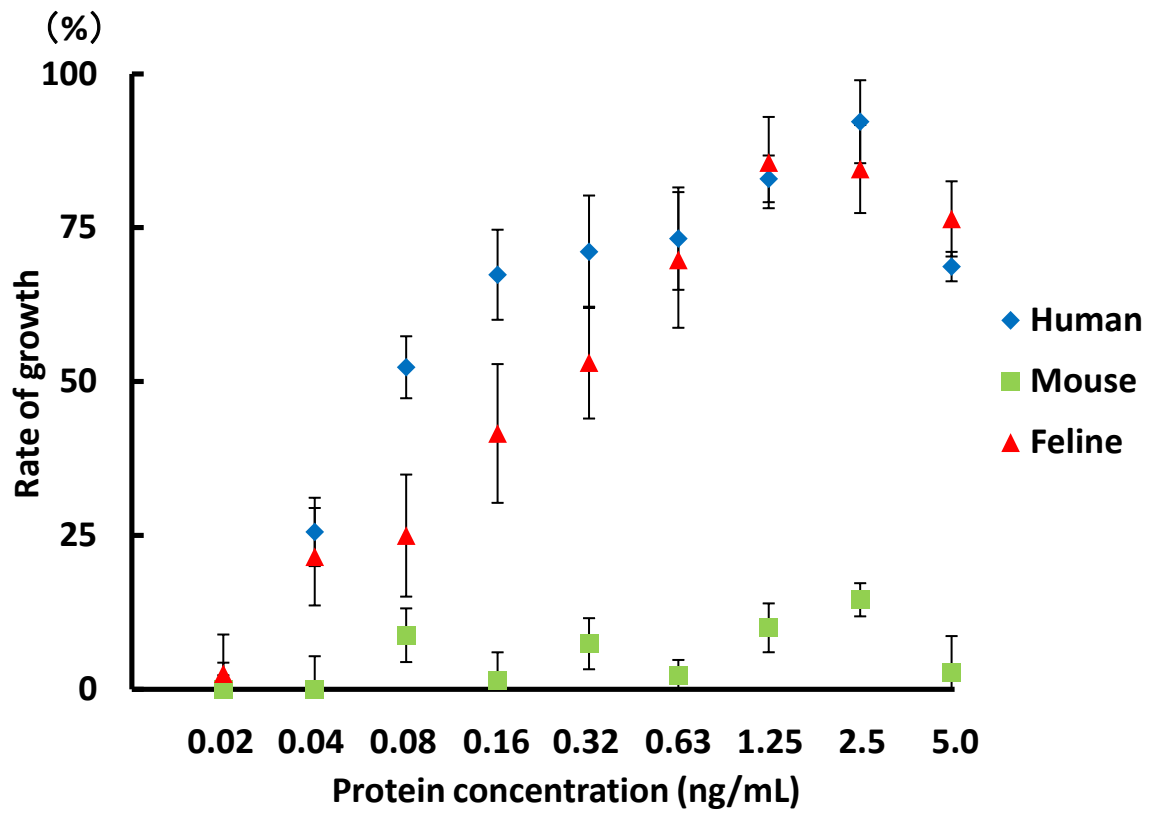


Fig.
6.

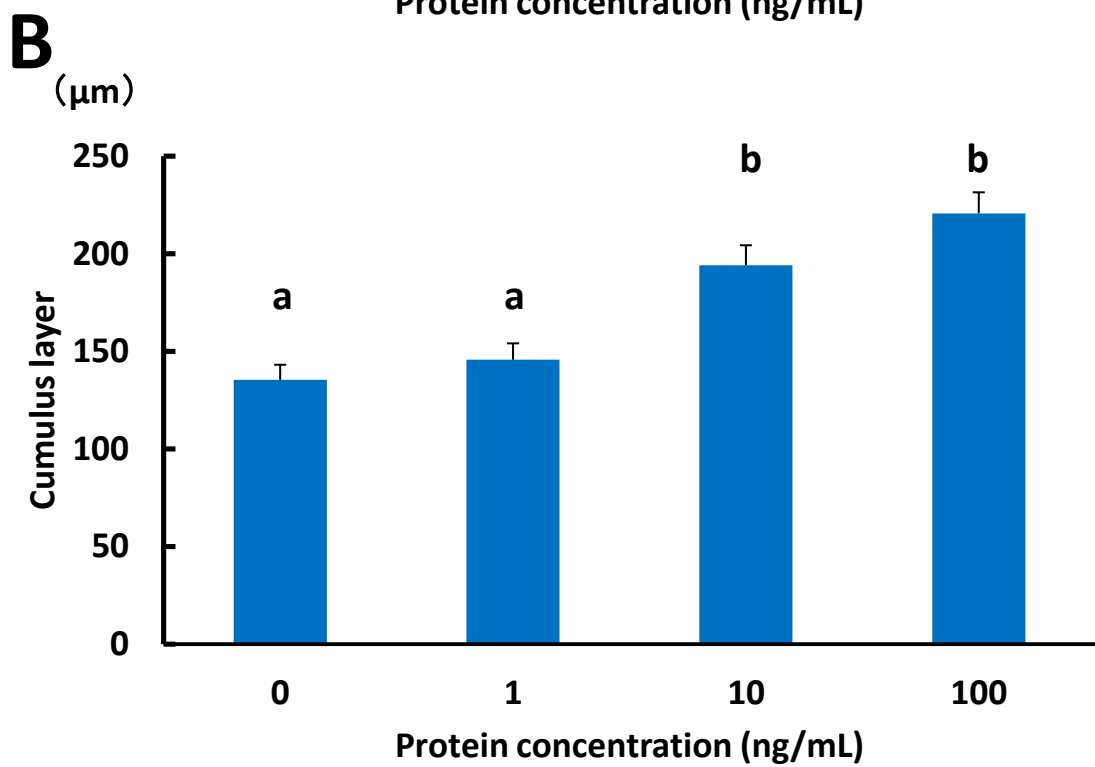
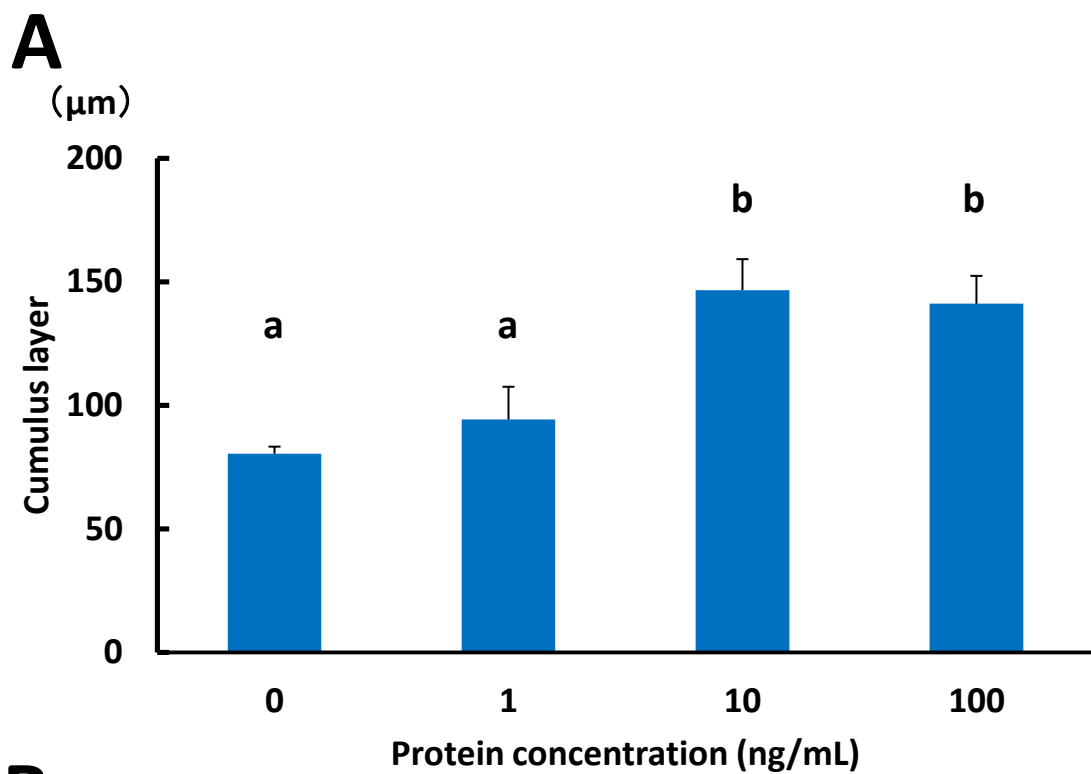


Fig. 7.