

## Molecular Mechanism to Maintain Stem Cell Renewal of ES Cells

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**ABSTRACT.** Embryonic stem (ES) cells are pluripotent cells directly derived from early stage embryos that retain the ability to differentiate into all cell types. This unique feature is the basis of various applications of ES cell technology such as *in vitro* models of mammalian development, germline transgenesis to make knockout mice, and a generic source for cell therapy in regenerative medicine. To achieve success in these applications, the pluripotency of ES cells has to be kept stable during long-term culture *in vitro*, leading to the necessity of determining the molecular basis for maintaining ES self-renewal. This paper summarizes the recent progress in this area, focusing mainly on the LIF signaling pathway and the transcription factor Oct-3/4. Although it is still unclear how these components work together, a model is presented here that provides a plan to solve this problem.

**Key words:** ES cell/pluripotency/LIF/Stat3/Oct-3/4

Cellular pluripotency can be defined as the ability of a cell to differentiate into various types of cells and belongs to all definitive tissues: ectoderm, mesoderm, and endoderm. In mammalian development, only particular subsets of cells in the early stage embryos transiently possess pluripotency. Therefore, it is difficult to characterize pluripotency *in vivo* because of inaccessibility due to small embryo size and in utero development after implantation. Embryonic stem (ES) cells are directly derived from such pluripotent cell populations and can maintain pluripotency under particular culture conditions *in vitro* (reviewed by Smith, 1992). Thus, they can provide a good model system to analyze molecular mechanism to maintain cellular pluripotency (reviewed by Pesce *et al.*, 1999). Recent progress on the study of signal transduction and transcriptional regulation supporting stem cell renewal of ES cells is reviewed in this report.

### Exogenous signal to maintain ES self-renewal

Leukemia inhibitory factor (LIF) is a cytokine belonging to the IL-6 family which was initially identified by its activity to induce differentiation of M1 leukemia cells (Tomida *et al.*, 1984; Gearing *et al.*, 1987). However, it was re-discovered as an activity to inhibit differentiation of mouse ES cells (Smith *et al.*, 1988; Williams *et al.*, 1988). Addition of LIF is sufficient to establish and maintain ES cells without feeder cells in the presence of fetal calf serum (Nichols *et al.*, 1990), indicating that this is an unique extrinsic factor specifically required for ES self-renewal. LIF function is mainly limited to prevent differentiation, and a previous report indicated that stimulation of proliferation could be separated from LIF action (Raz *et al.*, 1999). Since the analysis of the LIF signal transduction pathway was conducted in M1 cells (reviewed by Hirano *et al.*, 1997), studies in ES cells followed that sought to identify the difference that mediates the opposite cellular responses, which inhibits differentiation in ES cells while inducing differentiation in M1 cells.

### Signal transduction of LIF

The LIF receptor consists of the LIF-specific receptor subunit LIFR $\beta$  and the common signal transducer gp130, which is shared between the members of the IL-6 cytokine family (reviewed by Taga and Kishimoto, 1997). Since

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Abbreviations: ES, embryonic stem; LIF, leukemia inhibitory factor; GCSF, granulocyte colony stimulating factor; ESRF, ES renewal factor; ICM, inner cell mass; EC, embryonal carcinoma; PEC, primitive ectoderm; FGF, fibroblast growth factor; PDGF, platelet-derived growth factor; hCG, human chorionic gonadotropin; ELA, E1A-like activity; EG, embryonic germ.

LIFR $\beta$  possess an intracellular domain homologous to that of gp130, the roles of these subunits on signal transduction were investigated. Using the chimeric molecules consisting of an extracellular domain of human granulocyte colony stimulating factor (GCSF) and an intracellular domain of either LIFR $\beta$  or gp130, it was clearly shown that LIFR $\beta$  is not sufficient to mediate the signal to maintain ES self-renewal whereas gp130 is (Starr *et al.*, 1997; Niwa *et al.*, 1998). Indeed, activation of gp130 by the combination of IL-6 with soluble IL-6 receptor can support establishment of ES cell lines and maintain them (Nichols *et al.*, 1994; Yoshida *et al.*, 1994).

There are two major pathways of intracellular signal transduction downstream of gp130 in M1 leukemia cells (reviewed by Hirano *et al.*, 2000), the Jak-Stat pathway and the Shp2-Erk pathway. GCSF-gp130 chimeric receptors, which carry various deletions in the intracellular domain, were assayed for their ability to support ES self-renewal (Matsuda *et al.*, 1999; Niwa *et al.*, 1998). Since Jak and Shp2 interact with separate subdomains of the intracellular domain of gp130, variant chimeric receptors which can recruit only one of them can be generated, and the results clearly showed that activation of Jak, but not Shp2, is sufficient. The role of Stat3, which mainly acts in the LIF signaling pathway in M1 cells, was then tested using a dominant-negative mutant of Stat3 (Stat3F) (Minami *et al.*, 1996; Nakajima *et al.*, 1996). Overproduction of Stat3F in ES cells resulted in induction of differentiation in the presence of LIF, the morphological change of which is similar to that induced by withdrawal of LIF, indicating that activation of Stat3 is essential to the LIF signaling pathway (Niwa *et al.*, 1998; Starr *et al.*, 1997). Moreover, activation of Stat3 without LIF is sufficient for it, which was elegantly proved by Matsuda *et al.* (Matsuda *et al.*, 1999). They made a chimeric protein consisting of Stat3 and the variant ligand binding domain of estrogen receptor and introduced it into ES cells. This mutant Stat3 (Stat3ER) can be dimerized by addition of estrogen-derivative 4-hydroxy tamoxifen (4-HT), and the ES cells expressing them can be maintained by 4-HT without LIF while maintaining germline competency.

Shp2 signal pathway is not essential for ES self-renewal, but can still act as a modifier of the LIF signal. Shp2 mutant ES cells exhibited a lower magnitude of LIF dependency than wild-type ES cells (Qu and Feng, 1998; Saxton *et al.*, 1997), and the addition of the Erk kinase inhibitor PD98059 in the medium resulted in a similar phenotype (Burdon *et al.*, 1999a). Thus, the LIF signal is mainly transmitted to the nuclei by the Jak-Stat pathway, and the Shp2-Erk pathway does not contribute directly to stem cell self-renewal (reviewed by Burdon *et al.*, 1999b).

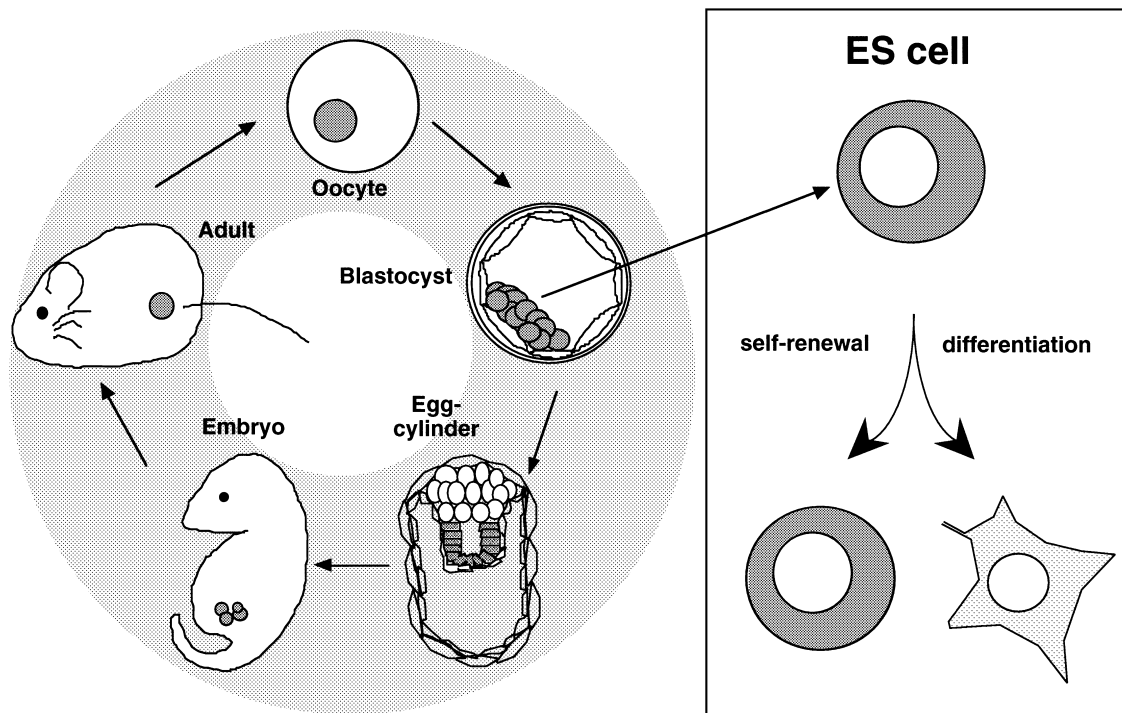
### ***LIF-independent signal: ESRF***

Although activation of Stat3 via gp130 by LIF-related

cytokines is sufficient for derivation and maintenance of ES cells (Nichols *et al.*, 1994; Yoshida *et al.*, 1994), there is no direct evidence indicating the importance of this cascade for keeping pluripotent cell population *in vivo*. It was reported that all of *Lif* (Stewart *et al.*, 1992), *Lifr $\beta$*  (Li *et al.*, 1995; Ware *et al.*, 1995), *gp130* (Yoshida *et al.*, 1996), and *Stat3* (Takeda *et al.*, 1997) knockout mice can develop beyond the egg cylinder stage. Why is such a discrepancy evident in LIF function *in vivo* and *in vitro*? One possible hypothesis is there is an unknown cascade which is functionally redundant to LIF signaling. Indeed, Dani *et al.* reported that ES cells lacking both endogenous LIF alleles still produce activity to support the undifferentiated state of ES cells in the absence of exogenous LIF after differentiation to parietal endoderm-like cells. They defined this activity as ES renewal factor (ESRF) (Dani *et al.*, 1998). ESRF can maintain the pluripotency of ES cells during a week without LIF. Interestingly, ESRF does not activate Stat3, indicating the presence of an alternative intracellular signaling pathway to maintain stem cell renewal.

### ***Function of Oct-3/4 in ES cells***

Stat3 acts as a transcription factor to activate target genes, which should include essential genes to maintain pluripotent cell phenotype. To date, there are only a few genes identified whose functions are essential to establish or maintain pluripotent cell population in pre- and early postimplantation embryos (reviewed by Pesce *et al.*, 1999). Moreover, in many cases, the phenotype may reflect their housekeeping function in proliferating cells rather than a specific function on pluripotent cell phenotype. For example, outgrowth of the inner cell mass (ICM) was perturbed in *B-myb* (Tanaka *et al.*, 1999) or *Chk1* (Takai *et al.*, 2000) mutant blastocyst, but it looks more like growth arrest of ICM rather than the disruption of the molecular machinery to maintain pluripotent phenotype because of their broad expression during embryogenesis and proposed general function. There is only one gene whose specific function in pluripotent cell population is confirmed, that is the POU-family transcription factor *Oct-3/4* (also known as *Oct-3* or *Oct-4*, encoded by *Pou5f1*). *Oct-3/4* was initially identified as a POU family member expressed in embryonal carcinoma (EC) cells (Rosner *et al.*, 1990; Schöler *et al.*, 1990a) or a transcriptional regulator binding to the retrotransposon enhancer which showed undifferentiated state-specific activity in EC cells (Okamoto *et al.*, 1990). *Oct-3/4* expression is tightly restricted in totipotent and pluripotent cells in mouse life cycle (Fig. 1) (reviewed in Pesce *et al.*, 1998a). Expression was observed in oocytes, early cleavage stage embryos, the ICM of the blastocyst, primitive ectoderm (PEC) in egg cylinder stage embryos, and primordial germ cells (Palmieri *et al.*, 1994; Schöler *et al.*, 1990b), suggesting its important role in maintaining cellular pluripotency. One exceptional loss of *Oct-3/4* in



**Fig. 1.** Stem cell-specific expression of Oct-3/4. Oct-3/4 expression is seen in fertilized eggs, inner cell mass of blastocyst, primitive ectoderm in egg cylinder, and primordial germ cells in later stage embryo and germ cells in adult. It also expresses in a stem cell-specific manner in mouse ES cells cultured *in vitro*.

pluripotent cell population can be found during oogenesis and spermatogenesis coincident with entry into meiosis (Pesce *et al.*, 1998b). Furthermore, the up-regulation in oocyte at the completion of prophase I of meiotic division is evident in ovary whereas Oct-3/4 expresses only in type A spermatogonia in testis, indicating specific role of Oct-3/4 in oocyte growth (Pesce *et al.*, 1998b).

The essential role of Oct-3/4 in mouse development has been revealed by targeting gene deletion (Nichols *et al.*, 1998). Oct-3/4 deficient embryos fail to initiate fetal development because the prospective founder cells of the ICM do not acquire pluripotency and become diverted into the trophoblast lineage, indicating Oct-3/4 is essential to establish pluripotent cell population in preimplantation development. Further investigation via conditional repression/expression in ES cells has revealed that the precise level of Oct-3/4 governs three different cell fates (Niwa *et al.*, 2000). ES cells require a critical level of Oct-3/4 to maintain stem cell renewal, and a less than twofold increase causes differentiation into endoderm and mesoderm whereas reduction to less than 50% of the normal expression level triggers dedifferentiation into trophoblast (Fig. 2) (Niwa *et al.*, 2000). In the presence of fibroblast growth factor (FGF)-4 and feeders, it is even possible to isolate trophoblast stem (TS) cells (Tanaka *et al.*, 1998) from ES cells by repression of Oct-3/4 expression (Niwa *et al.*, 2000). These observations not only accord

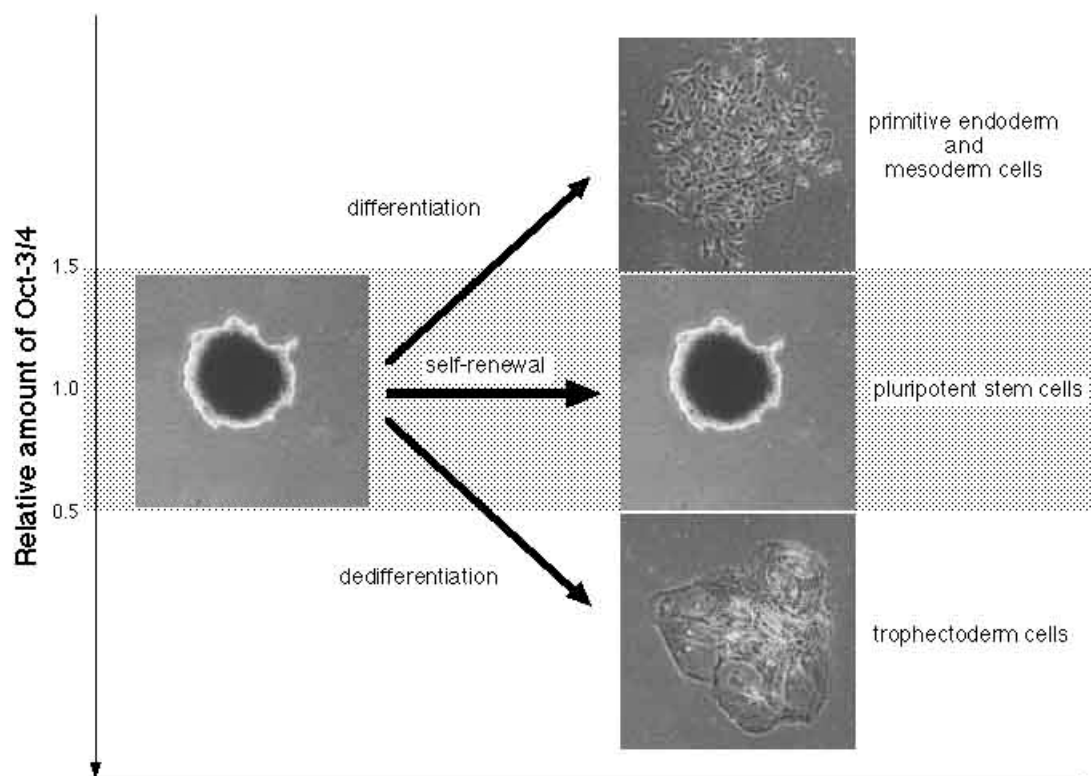
with the expression pattern in preimplantation embryos, in which Oct-3/4 is up-regulated in primitive endoderm and down-regulated in trophoblast (Palmieri *et al.*, 1994), but also fit the phenotype of Oct-3/4-null embryos (Nichols *et al.*, 1998). Therefore, Oct-3/4 can be regarded as a candidate for master regulator of initiation, maintenance and differentiation of pluripotent cells.

Does Stat3 keep cellular pluripotency via activation of Oct-3/4? This simple hypothesis is now ruled out because keeping the expression of Oct-3/4 at appropriate levels cannot prevent differentiation of ES cells induced by withdrawal of LIF (Niwa *et al.*, 2000). However, the up-regulation of Oct-3/4 induces a differentiation event which is quite similar to that induced by withdrawal of LIF (Niwa *et al.*, 2000). To investigate the relationship between these differentiation events, it is necessary to inquire into the molecular mechanism of Oct-3/4 function in detail.

### How Oct-3/4 works in ES cells

#### Target genes

A few target genes of Oct-3/4 have been identified to date, and the Oct-3/4 binding sites confirmed in their regulatory elements are listed in Fig. 3 (Nishimoto *et al.*, 1999; Ben-Shushan *et al.*, 1998; Botquin *et al.*, 1998; Kraft *et al.*, 1996; Liu and Roberts, 1996; Saijoh *et al.*, 1996;



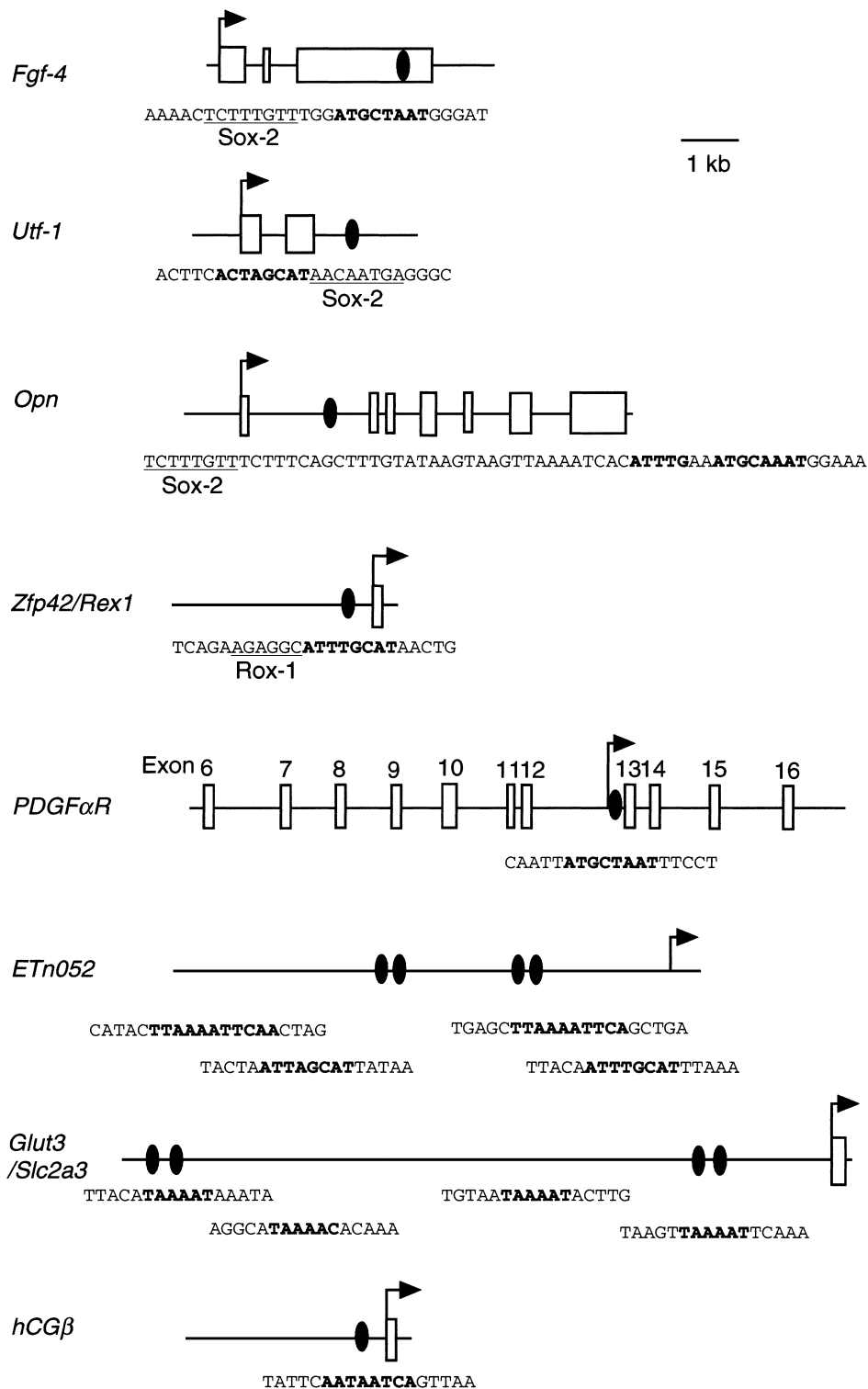
**Fig. 2.** Relationship between Oct-3/4 expression and stem cell fate. To maintain the undifferentiated phenotype, Oct-3/4 expression must remain within plus or minus 50% of normal diploid expression. If Oct-3/4 expression is increased beyond the upper threshold level, differentiation into primitive endoderm and mesoderm is triggered. If Oct-3/4 expression is decreased below the lower threshold level, stem cells are dedifferentiated into trophectoderm lineage.

Yuan *et al.*, 1995; Okamoto *et al.*, 1990). It was shown that Oct-3/4 can bind a variety of sequences including the consensus octamer motif (ATGCAAAT) and the AT-rich sequence (Saijoh *et al.*, 1996; Okamoto *et al.*, 1990), hence the high-affinity binding sites would be determined by association with appropriate co-factors which also have sequence-specific DNA-binding abilities. Oct-3/4 adopts various monomer configurations on DNA and can form homo- and heterodimers on the palindromic Oct factor recognition element and its derivatives (Tomilin *et al.*, 2000; Botquin *et al.*, 1998).

The genes encoding *Fgf-4* (Yuan *et al.*, 1995), the transcriptional co-factor *Utf-1* (Nishimoto *et al.*, 1999), the zinc-finger protein *Zfp42/Rex-1* (Ben-Shushan *et al.*, 1998) and the platelet-derived growth factor  $\alpha$  receptor (*PDGFR*) (Kraft *et al.*, 1996) were identified as targets of Oct-3/4 by their stem cell-specific expression. Regulation of *osteopontin* (*Opn*) by Oct-3/4 was determined by immunoprecipitation of the first intron of *Opn* from covalently-fixed chromatin of EC cells by Oct-3/4-specific antibodies (Botquin *et al.*, 1998). Saijoh *et al.* systematically isolated several candidates of target genes by their elegant cDNA subtraction screening method (Saijoh *et al.*, 1996), and we confirmed Oct-3/4-dependent expression in ES cells for some of them, *Otx-2*, *Lefty-1/Ebaf*, *uridine*

*phosphorylase* (*Upp*)/383 and *Tera*/226 (Niwa *et al.*, 2000). All of these defined and putative target genes showed stem cell-specific expression in ES cells, but they exhibit differential expression pattern in pre- and early postimplantation embryos (Table I). Although all these genes express in the ICM of blastocyst, expressions of *Lefty-1/Ebaf*, *Opn*, *Zfp42/Rex-1* and *Upp*/383 were not observed in PEC in egg cylinder stage embryos that possess pluripotency and express Oct-3/4. Oct-3/4 has been identified as a repressor on the expression of *human chorionic gonadotropin*  $\beta$  subunit (*hCG* $\beta$ ) in choriocarcinoma cells (Liu and Roberts, 1996), and our previous data suggested that Oct-3/4 repress directly or indirectly the caudal-related homeobox transcription factor *Cdx-2* and the basic helix-loop-helix transcription factor *Hand-1* (also known as *eHand* or *Hxt*) (Niwa *et al.*, 2000), both of which express in the trophectoderm lineage in the early stage embryos (Beck *et al.*, 1995; Cross *et al.*, 1995).

Little is known about the function of these target genes. *Fgf-4* is the best analyzed one, the function of which is essential for peri-implantation development (Feldman *et al.*, 1995). However, it has been revealed that the function of *Fgf-4* is not essential for stem cell renewal of ES cells and might be important as a paracrine growth factor for polar trophectoderm and primitive endoderm (Tanaka *et al.*,



**Fig. 3.** Target genes of Oct-3/4. Position and sequence of the Oct-3/4 binding elements of *Fgf-4* (Yuan *et al.*, 1995), *Utf1* (Okuda *et al.*, 1998), *Opn* (Botquin *et al.*, 1998), *Zfp42/Rex-1* (Ben-Shushan *et al.*, 1998), *PDGFαR* (Kraft *et al.*, 1996), *ETn052* (Okamoto *et al.*, 1990), *Slc2a3/Glut3* (Saijoh *et al.*, 1996), and *hCGβ* (Liu and Roberts, 1996) are highlighted. Bold letters show the binding sequence of Oct-3/4, and the underlined letters indicate the binding site of co-factors. The task is to find the various positions of the binding sites in these target genes, which may correlate with different dependency of cofactors.

**Table I.** EXPRESSION PATTERN OF OCT-3/4 TARGET GENES IN PLURIPOTENT CELLS

Gene	ICM	embryos			sqelching <sup>m</sup>
		PEC	ES <sup>m</sup>		
<i>Fgf-4</i>	+ <sup>a</sup>	+ <sup>a</sup>	+		—
<i>Utf-1</i>	+ <sup>b</sup>	+ <sup>b</sup>	+ <sup>n</sup>		— <sup>n</sup>
<i>Tera/226</i>	+ <sup>c</sup>	+ <sup>d</sup>	+		+
<i>Lefty-1</i>	+ <sup>e</sup>	— <sup>f</sup>	+		—
<i>Otx-2</i>	+ <sup>e</sup>	+ <sup>g</sup>	+		—
<i>Opn</i>	+ <sup>h</sup>	— <sup>h</sup>	+ <sup>h</sup>		nd
<i>Zfp42/Rex-1</i>	+ <sup>i</sup>	— <sup>i</sup>	+		+
<i>Upp/383</i>	+ <sup>d</sup>	— <sup>d</sup>	+		+
<i>Slc2a3/Glut-3</i>	+ <sup>d</sup>	+ <sup>d</sup>	+		—
<i>Sox-2</i>	+ <sup>e</sup>	+ <sup>j</sup>	+		
ELA	+ <sup>k,l</sup>	— <sup>k,l</sup>	+ <sup>o</sup>		

All genes except Sox-2 and ELA are candidates as Oct-3/4 targets. References for their expression in pluripotent cells are listed below; <sup>a</sup>(Rappolee *et al.*, 1994); <sup>b</sup>(Okuda *et al.*, 1998); <sup>c</sup>hit to EST of blastocysts; <sup>d</sup>(Saijoh *et al.*, 1996); <sup>e</sup>(Matsui, H. & Niwa, H., unpublished); <sup>f</sup>(Meno *et al.*, 1997); <sup>g</sup>(Simeone *et al.*, 1993); <sup>h</sup>(Botquin *et al.*, 1998); <sup>i</sup>(Rogers *et al.*, 1991); <sup>j</sup>(Collignon *et al.*, 1996); <sup>k</sup>(Suemori *et al.*, 1988); <sup>l</sup>(Dooley *et al.*, 1989); <sup>m</sup>(Niwa *et al.*, 2000); <sup>n</sup>Niwa H., unpublished; <sup>o</sup>(Suemori *et al.*, 1988).

1998; Wilder *et al.*, 1997). Functional analysis of *Otx-2* (Matsuo *et al.*, 1995), *Lefty-1/Ebaf* (Meno *et al.*, 1998) and *Opn* (Hynes, 1996) using knockout mice revealed that functions of these genes were not essential to establish the pluripotent cell population in preimplantation-stage embryos which affected in *Oct-3/4* knockout embryos. In contrast, targeted inactivation of *Cdx-2* resulted in embryonic lethality between 3.5–5.5 day postcoitum (Chawengsaksophak *et al.*, 1997) and *Hand-1* mutant embryos exhibit defect in differentiation of secondary giant cells in placenta (Firulli *et al.*, 1998; Riley *et al.*, 1998), indicating the important role of these genes in placental development.

### Co-factors

It is known that POU family transcription factors can act as both transcriptional activators and repressors by cooperating with various co-factors (Xu *et al.*, 1998; Gstaiger *et al.*, 1995; Zwilling *et al.*, 1995; Lai *et al.*, 1992), and several different co-factors of Oct-3/4 have been reported. The first co-factor of Oct-3/4 identified was the adenovirus E1A (Schöler *et al.*, 1991), which was regarded as a substitute of the so-called E1A-like activity (ELA) observed in undifferentiated cells (La Thangue and Rigby, 1987). For activation of the artificial promoter in heterologous cells, E1A served as a bridging factor between Oct-3/4 and the basic transcription machinery. The second

one was the Sry-related factor Sox-2, which was initially identified as a co-factor to activate *Fgf-4* enhancer (Yuan *et al.*, 1995) and then found to activate the *Utf-1* enhancer (Nishimoto *et al.*, 1999). In contrast, it was reported that Sox-2 prevents activation of the *Opn* enhancer by homodimer of Oct-3/4 (Botquin *et al.*, 1998). In the case of the *Zfp42/Rex-1* promoter, the unidentified factor Rox-1 was hypothesized as a new co-factor of Oct-3/4 (Ben-Shushan *et al.*, 1998), and screening of the phage display library revealed interaction of several factors with Oct-3/4, which included HMG-1 (Butteroni *et al.*, 2000). It was known that the allosteric effect results in the recruitment of a corepressor complex on the POU factor Pit-1 which usually interacts with a coactivator complex (Scully *et al.*, 2000), so a similar effect might provide the repressor function on Oct-3/4.

One interesting observation in the cooperation between Oct-3/4 and co-factors was the sqelching phenomenon (Schöler *et al.*, 1991). In transactivation of the artificial promoter element by Oct-3/4 and E1A, the quantitative balance between these factors is important for proper activation by a ternary complex that consists of Oct-3/4, E1A and the basal transcription machinery. Excess amounts of Oct-3/4 or E1A prevent formation of active complex by occupation of binding site or surface saturation, respectively, resulting in prevention of activating the target promoter. A similar phenomenon was observed in the activation of the *Zfp42/Rex-1* promoter (Ben-Shushan *et al.*, 1998), which was repressed by excess amounts of Oct-3/4 in EC cells, but not in the *Fgf-4* and *Utf-1* enhancers activated by Oct-3/4 and Sox-2 (Nishimoto *et al.*, 1999; Yuan *et al.*, 1995).

Another point of interest is that these co-factors of Oct-3/4 exhibit different expression patterns in pluripotent cells. Sox-2 expresses in both ICM and PEC (Botquin *et al.*, 1998)(Collignon *et al.*, 1996), but ELA was observed only in ICM (Dooley *et al.*, 1989; Suemori *et al.*, 1988) (Table I). Therefore, the differential expression of target genes might be due to their different dependency of co-factors. An alternative explanation is that the expression of these target genes is differentially regulated by factors other than Oct-3/4, which mostly happens on the two separate enhancers of *Oct-3/4* itself (Yeom *et al.*, 1996) (see below).

### Functional domains

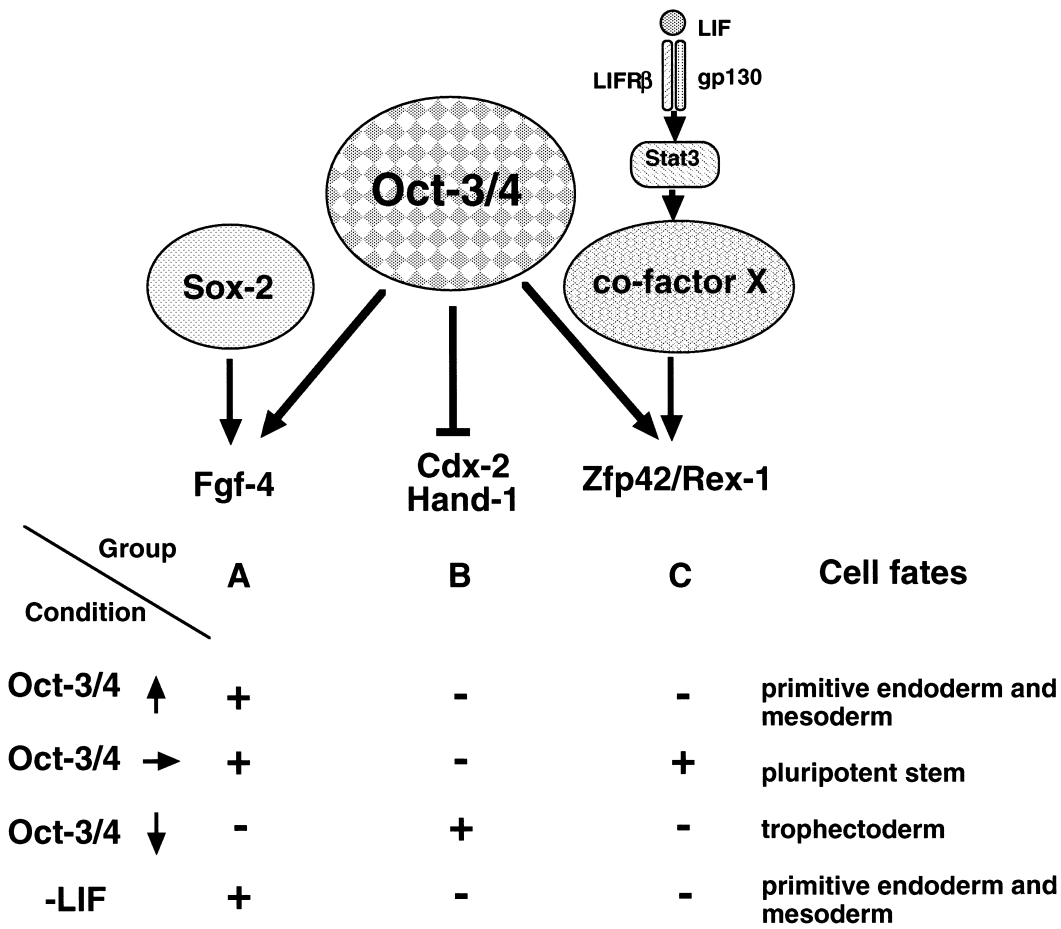
Oct-3/4 consists of a bipartite DNA-binding POU domain and both amino-terminal and carboxy-terminal transactivation domains (Okamoto *et al.*, 1990; Rosner *et al.*, 1990; Schöler *et al.*, 1990a). The functional difference between these separate transactivation domains was reported (Ambrosetti *et al.*, 2000; Brehm *et al.*, 1997; Viganó and Staudt, 1996; Imagawa *et al.*, 1991), but our recent finding clearly indicated that they share redundant functions on ES self-renewal, and that the combination of one of them with the proper POU domain is sufficient to substitute for the function of Oct-3/4 in ES cells (Niwa *et al.*, submitted for

publication). Interestingly, the DNA-binding ability of Oct-3/4 is essential to maintain ES cell phenotype but not to induce differentiation by overproduction, suggesting that the titration out of relevant partner(s) (Schöler *et al.*, 1991) is achieved by protein-protein interactions that are independent of DNA-binding (Niwa *et al.*, submitted for publication).

**Model for molecular mechanism governs ES cell phenotype**

A model for Oct-3/4 and Stat3 cooperative function is shown in Fig. 4 as proposed in our previous report (Niwa *et al.*, 2000). The categorization of Oct-3/4 target genes into three groups A to C is based on (1) the function of Oct-3/4 on these targets (activation or repression) and (2) the expression pattern in ES cells with excess amounts of Oct-3/4 (repressed by squelching or not). As shown above, up-regulation of Oct-3/4 results in differentiation to primitive endoderm and mesoderm, which is different from

the phenotype generated by its down-regulation. The squelching phenomenon can give the explanation that excess amounts of Oct-3/4 result in its loss-of-function on activation of target genes, but repression by overexpression of Oct-3/4 occurs only in a part of target genes such as *Zfp42/Rex-1* and *Upp/383*. Therefore, a hypothesis was introduced in a key position of this model, in which the target genes activated by Oct-3/4 can be divided into two groups, one is repressed by squelching mechanism and the other is not, and the difference between them is based on their different co-factor dependency. This hypothesis can be supported by the correlation between differential expression patterns of co-factors and target genes in pluripotent cells (Table I) and the different co-factor dependencies of known target genes. Indeed, the co-factor *Sox-2* and two *Sox-2*-dependent Oct-3/4 target *Fgf-4* and *Utf-1* co-express in both ICM and PEC (Table I), and neither *Fgf-4* nor *Utf-1* was repressed by overproduction of Oct-3/4 in ES cells (Niwa *et al.*, 2000, and unpublished result), as there is no evidence suggesting that the squelching occurs between Oct-3/4 and



**Fig. 4.** Model for Oct-3/4 and Stat3 co-operative function. Putative and defined Oct-3/4 target genes are divided into three groups and the typical members of genes in each group are shown. By introduction of two hypotheses in the model, we can explain all phenomenon observed by the change of Oct-3/4 expression and/or withdrawal of LIF. See the text for details.

Sox-2. In contrast, the two target genes repressed by overexpression of Oct-3/4, *Zfp42/Rex-1* and *Upp/383*, express in ICM but not in PEC, and the putative co-factor ELA exhibits the same expression pattern as these genes (Table I), which substitute E1A to mediate the squelching phenomenon. In Fig. 4, Sox-2 represents a co-factor that does not mediate squelching, while co-factor X represents a co-factor that mediates squelching. Group A target genes are activated by Oct-3/4 and the Sox-2-class co-factors and not repressed by squelching, while group B ones are repressed by Oct-3/4, and group C ones are activated by Oct-3/4 and co-factor X and repressed by squelching. When the Oct-3/4 expression is maintained within the appropriate level, the expression pattern of these groups [A,B,C] is [ON,OFF,ON], resulting in stem cell renewal. By down-regulation of Oct-3/4, the expression pattern of these groups is inverted to [OFF,ON,OFF], resulting in differentiation to trophectoderm. Since the genes involved in placental development belongs to group B, it would seem that a component of Oct-3/4 functions as a gatekeeper that prevents differentiation into the trophectoderm lineage and thereby locks pluripotent capacity. Up-regulation of Oct-3/4 results in shutting off group C and provides the expression pattern [ON, OFF, OFF] for the expression of the groups [A,B,C], which is different from the pattern shown above and results in differentiation to primitive endoderm and mesoderm. Such choice of cell fate might be achieved by down-regulation of group C, which may be required for maintaining stem cell renewal, while keeping repressed the group B genes involved in commitment to trophectoderm lineage. A similar situation obtains by repression of co-factor X, and it may also happen in PEC as observed in ELA (Dooley *et al.*, 1989; Suemori *et al.*, 1988) to allow their differentiation to embryonic tissues via gastrulation.

To explain the relationship between the LIF/Stat3 pathway and Oct-3/4, a second hypothesis is incorporated in this model, in which Stat3 activates the expression of co-factor X. This hypothesis is based on the evidence that overexpression of Oct-3/4 and withdrawal of LIF induce a similar differentiation and allows explanation of this result because both of these events give the same expression pattern [ON,OFF,OFF] of the groups [A,B,C]. It also fits the observation that maintenance of Oct-3/4 expression is not sufficient to keep stem cell phenotype in the absence of LIF (Niwa *et al.*, 2000).

To confirm this model, it is necessary to prove these two hypotheses. The common component of them is co-factor X, which may correspond to the hypothetical factor Rox-1 or the unidentified ELA. For this purpose, the systematic identification of the proteins that interact with Oct-3/4 is essential in future.

### Regulation of Oct-3/4 expression

If the change of Oct-3/4 expression level is a primary

event to determine cell fates in preimplantation development, how is it achieved? In ES cells, the transcription level of Oct-3/4 should be kept in the narrow range to maintain pluripotent cell phenotype. For such tight regulation of transcription, a negative feedback mechanism should be present in pluripotent cells expressing Oct-3/4. Indeed, in the model system in *Escherichia coli*, the presence of a simple negative feedback loop dramatically keeps the expression level within the narrow range, within 5% of the average level (Becskei and Serrano, 2000). However, there is no experimental evidence which suggests the presence of such a simple negative feedback loop in the regulation of Oct-3/4 expression. Analysis of the regulatory element of Oct-3/4 revealed that there are two separate enhancer elements upstream the Oct-3/4 promoter, a distal enhancer (DE) and a proximal enhancer (PE) (Yeom *et al.*, 1996). DE is active in ICM and PGC but not in PEC, whereas PE is active only in PEC. Among the pluripotent cell lines, DE is active in ES and embryonic germ (EG) cells whereas PE is active in EC cells. Only PE has been analyzed in detail and it was revealed that transcription factors that belong to the nuclear receptor superfamily bind to the core element of PE (Sylvester and Schöler, 1994). COUP-TFs act as a negative regulator on this element (Ben-Shushan *et al.*, 1995) and GCNF exhibits a pattern of expression that suggests it could be involved in regulating the activity of Oct-3/4 promoter (Chen *et al.*, 1994), so they can be regarded as candidate components of a negative feedback loop.

### Evolutional location of LIF/Stat3 and Oct-3/4

In mouse ES cells, both the LIF/Stat3 pathway and the transcription factor Oct-3/4 have a pivotal role to keep cellular pluripotency. Such an important mechanism tends to be evolutionally conserved, but there are lines of evidence to prevent generalization of this model to other species. For example, all mouse ES cell lines have dependency to LIF, although there are variations in degree; for instance, the recently established rhesus and human ES cells do not exhibit clear dependency to LIF (Thomson *et al.*, 1998; Thomson *et al.*, 1995). Medaka ES cell lines also undergo self-renewal without LIF (Hong *et al.*, 1996), but ES cell lines of chicken (Pain *et al.*, 1996), rat (Lannaccone *et al.*, 1994), and EG cell lines of mouse (Matsui *et al.*, 1992) and human (Shamblott *et al.*, 1998) undergo stem cell renewal in LIF-dependent manner. Interestingly, most of mouse EC cell lines can grow up without LIF and overexpression of Stat3F cannot prevent it (Niwa *et al.*, 1998), which rules out the hypothesis that an alternative pathway activates Stat3 in these EC cells to keep the expression of co-factor X without LIF. Therefore, it is still unclear how the pluripotency is maintained in LIF-independent ES and EC cell lines. One possible explanation is the variation in balance between the dependency to the



LIF/Stat3 pathway and the ESRF pathway. To test this hypothesis, identification of ESRF should be done first.

Oct-3/4 function looks like well conserved among the mammals. The human ES cells express Oct-3/4 in stem cell-specific manner (Reubinoff *et al.*, 2000), and expression of Oct-3/4 is tightly regulated in human ICM cells whereas it is repressed in trophectoderm (Hansis *et al.*, 2000). In domestic animals, Oct-3/4 expression can be detected in both ICM and trophectoderm (Kirchhof *et al.*, 2000; van Eijk *et al.*, 1999), but the decrease of Oct-3/4 expression to induce differentiation, which is only a 50% reduction in mouse ES cells (Niwa *et al.*, 2000), may happen in the trophectoderm although the precise estimation of expression level has not been done. Overexpression of Oct-3/4 in mouse EC cells can induce differentiation (Niwa, H., unpublished results), indicating the same function in LIF-independent pluripotent cells as in mouse ES cells. However, to date, Oct-3/4 homologue has been found only in mammals such as marsupial brushtail possum (Frankenberg *et al.*, 2001), mouse (Okamoto *et al.*, 1990; Rosner *et al.*, 1990; Schöler *et al.*, 1990a), bovine (van Eijk *et al.*, 1999) and human (Takeda *et al.*, 1992). Oct-3/4 belongs to the class V of the POU family (Rosenfeld, 1991), and the members of this class were identified in Zebrafish and *Xenopus* but not in *Caenorhabditis elegans* and *Drosophila melanogaster* genomes, and systematic search of chicken genome failed to identify any homologous gene (Soodeen-Karamath and Verrinder Gibbins, 2001). These data indicate that the class V POU factor is evolutionally new in this family because members in the other class can be observed in *Caenorhabditis elegans* and *Drosophila melanogaster* (reviewed by Ryan and Rosenfeld, 1997), and the conserved mechanism to keep pluripotency other than Oct-3/4 should be present because ES cell lines were established from the species in which Oct-3/4 homolog have not been identified such as chicken (Soodeen-Karamath and Verrinder Gibbins, 2001; Pain *et al.*, 1996). Since the precise level of Oct-3/4 governs three distinct fates of mouse ES cells, it can be regarded as a switch to separate the placental and embryonic lineages. Therefore, this would lead us to speculate that Oct-3/4 is an optional switch to generate the placenta in the evolution of mammals, in which the evolutionally conserved mechanism to keep pluripotency is under the control of Oct-3/4 as the group C target genes.

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