

The *Pou5f1* distal enhancer is sufficient to drive *Pou5f1* promoter-EGFP expression in embryonic stem cells

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ABSTRACT The POU5F1 transcription factor is the gatekeeper of the pluripotent state in mammals. It is essential for epigenetic reprogramming events and also for germ cell viability. *Pou5f1* gene expression is tightly controlled during embryogenesis, but its regulatory regions are not fully deciphered. The GOF18 \triangle PE-EGFP transgene, harboring the enhanced green fluorescence protein reporter gene inserted into a 17- kilobase long mouse *Pou5f1* genomic sequence, has been widely used to visualize pluripotent embryonic cells and primordial germ cells in the mouse and other mammalian species. This construct includes the *Pou5f1* promoter under the control of the distal enhancer and also includes the *Pou5f1* gene body and flanking sequences. In search of the essential regulatory regions of *Pou5f1*, we generated four shorter forms of this construct. We found that the shortest form, containing the *Pou5f1* promoter and distal enhancer but lacking the gene body and upstream flanking sequences, correctly expressed EGFP in transiently transformed undifferentiated ES cells, correctly switched it off upon ES cell differentiation, and correctly kept it silenced in differentiated Hep3B cells. Similarly to the original GOF18 \triangle PE-EGFP, this shortest form was expressed in the fetal mouse gonad. Our data suggest that the *Pou5f1* distal enhancer and proximal promoter may be sufficient to specify transgene expression in pluripotent cells.

KEY WORDS: Pou5f1, Oct4, EGFP transgene, pluripotency

Introduction

The Pou5f1 gene (also termed Oct3, Oct4 or Oct3/4) encodes a member of the Pic-1, Oct1,2, Unc-86 (POU) transcription factor family (Yeom et al., 1996). Pou5f1 protein is present in the totipotent zygote as a maternal factor. The Pou5f1 gene is activated during cleavage stages and remains active in the inner cell mass (ICM) and epiblast. After gastrulation, Pou5f1 is exclusively expressed in the developing germ line. POU5F1 transcription factor is essential for the pluripotency of ICM cells in vivo (Nichols et al., 1998) and for the viability of primordial germ cells (PGC) (Kehler et al., 2004). Introduction of POU5F1 and few other transcription factors can reprogram somatic cells into an induced pluripotency state or even can induce trans-differentiation, altering cell fate (Sterneckert et al., 2012). Therefore, POU5F1 is not only a gatekeeper in the early mammalian development, but also a gatekeeper for reprogramming expressway (Pesce and Scholer, 2001, Sterneckert et al., 2012). Because of its strict regulation during development, reporter genes under the control of the mouse Pou5f1 regulatory elements provide suitable tools for identifying pluripotent cell types (Yeom

et al., 1996). A reporter transgenic construct, GOF18APE-EGFP, consisting of a 7.5 kb promoter and distal enhancer region upstream of the Pou5f1 gene, an enhanced green fluorescent protein (EGFP) gene, and the five exons of Pou5f1, has been used as a powerful tool to visualize pluripotent cells in mouse development (Yoshimizu et al., 1999). The GOF18 APE-EGFP transgene provided an EGFP expression pattern that was faithful to the endogenous Pou5f1 expression in other mammals including pigs (Kirchhof et al., 2000, Nowak-Imialek et al., 2011). Transgenic mouse strains made with GOF18APE-EGFP, such as the TgOG2 line (Szabó et al., 2002, Yoshimizu et al., 1999) have been very useful for isolating PGCs and fetal germ cells using flow cytometry based on EGFP expression. We decided to find the essential components of the GOF18∆PE-EGFP transgene that are sufficient to specify reporter expression in pluripotent embryonic stem (ES) cells and in fetal germ cells and specify its silencing in differentiated cell types. We

Abbreviations used in this paper: DE, distal enhancer; EGFP, enhanced green fluorescent protein; PGC, primordial germ cell; Pou5f1, Pic-1, Oct1,2, Unc-86 transcription factor 1; PP, proximal promoter; RFP, red fluorescent protein.

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constructed four shorter forms of GOF18 Δ PE-EGFP and analyzed their expression patterns in cultured cells and mouse gonads. To this end, we found that the shortest form consisting of the *Pou5f1* distal enhancer and promoter is sufficient to drive EGFP expression in undifferentiated ES cells and in the 14.5 days post coitum (dpc) fetal gonad and is also sufficient to be silenced in differentiated Hep3B cells.

Results

$Considerations \, for \, shortening \, the \, {\it GOF18} \Delta PE{\it -}EGFP \, construct$

The Schöler laboratory has characterized the regulatory regions of the Pou5f1 gene in great detail using LacZ reporter transgenic constructs (Fig. 1.) From these analyses we concluded that in order to keep the specific expression pattern of the GOF18∆PE-EGFP in the shortened construct, we must keep at least two essential regions, the proximal promoter (PP) and the distal enhancer (DE) together with the EGFP reporter. The 230 bp long PP is essential for gene activity in pluripotent cells, because the promoterless GOF18ΔPP-LacZ construct is completely silent in embryonic stem (ES) cells (Fig. 1). The Pou5f1 PP is also essential for restricted germ cell-specific expression after gastrulation. GCNF orphan nuclear receptor binds and represses the Pou5f1 PP upon differentiation, restricting its activity to germ cells (Fuhrmann et al., 2001). The DE is required for activating the PP during preimplantation development and in ES cells, as well as in the germ cells (compare GOF 9 and GOF6; compare GOF18∆PE-LacZ and GOF18ADE-LacZ). We hypothesized that we could shorten the GOF18APE-EGFP construct from the 3' end: it might be possible to remove the downstream sequences and the gene body without compromising specificity. The downstream part was not essential for expression in pluripotent cells (compare GOF18 and GOF13). The gene body may not be required, but this deletion hasn't been tested before. Additionally, we considered that the upstream region 5' to the DE might also be shortened, because GOF9, that lacks this region is expressed in pluripotent cells.

Four shorter forms of GOF18 Δ PE-EGFP were expressed in mouse ES cells

We generated four shorter forms GOF18∆PE-EGFP, each containing the EGFP gene (Fig. 2) fused to variable lengths of

TABLE 1

POU5F1 REGULATORY SEQUENCES USED IN THE DIFFERENT CONSTRUCTS

	Pou5f1 regulatory sequence length (kb)					
Construct	Upstream	Downstream	Total			
GOF18∆PE-EGFP	7.5	9	16.5			
GOF18∆PE-EGFP S1	7.5	2	9.5			
GOF18∆PE-EGFP S2	5.5	2	7.5			
GOF18∆PE-EGFP S3	4.5	9	13.5			
GOF18∆PE-EGFP S4	4	0	4			

TABLE 2

DOWNREGULATION OF POU5F1-EGFP IN THE ABSENCE OF CONDITIONED MEDIUM

	Percent GFP+ cells in FACS							
	AVERAGE (n=3) STDEV((n=3)		Change			
Construct	LIF+	LIF-	LIF+	LIF-	Total	%	TTEST	
Pgk-EGFP control	28.51	25.15	2.01	0.93	-3.36	-11.77	0.01847	
GOF18∆PE-EGFP	1.06	0.46	0.16	0.04	-0.60	-56.78	0.01592	
GOF18∆PE-EGFP S4	3.25	1.88	0.20	0.36	-1.38	-42.32	0.00348	

Pou5f1 sequences (Table 1). GOF18 Δ PE-EGFP S1 retained the 7.5 kb of promoter/enhancer region and the first exon of *Pou5f1* gene. GOF18 Δ PE-EGFP S2 contained 5.5 kb of the promoter/ enhancer region and the first exon. GOF18 Δ PE-EGFP S3 contained 4.5 kb of the *Pou5f1* enhancer/promoter region and five exons. GOF18 Δ PE-EGFP S4, the shortest form, only harbored the 3.5 kb DE-PP region to drive EGFP expression. To analyze if these shorter versions of GOF18 Δ PE-EGFP retain the expression specificity of the original transgene, we transfected them into mouse ES cells (Fig. 3). Each of the four shorter (S1-S4) constructs drove EGFP expression in ES cells similarly to the original GOF18 Δ PE-EGFP. Only a subset of cells expressed EGFP. This was expected, because the efficiency of transient transfection is never 100%.

In addition, we found that the shortest form, GOF18 Δ PE-EGFP S4, carried the signal for repression in response to differentiation. We transfected the ES cells with the GOF18 Δ PE-EGFP, GOF18 Δ PE-EGFP S4 and positive control Pgk promoter-EGFP

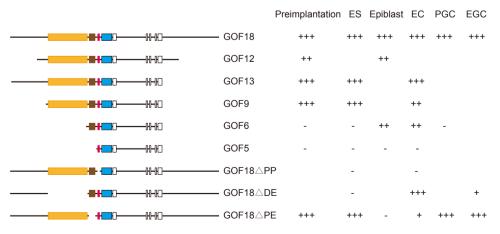
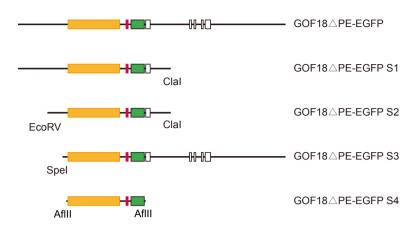


Fig.1. Considerations for shortening the GOF18ΔPE-EGFP construct. Summary of transgenic experiments done by the Schöler group (Yeom et al., 1996). The different constructs are depicted with their components: LacZ reporter with SV40 polyA termination, blue box; distal enhancer (DE), orange box; proximal enhancer (PE), brown box; Pou5f1 proximal promoter (PP), red box; Pou5f1 exons, open boxes. Expression of the reporter gene is tabulated to the right: +++, high expression; ++, reduced expression;

+, greatly reduced expression, compared to GF18; -, lack of expression at the different developmental stages: during preimplantation, in the epiblast, in primordial germ cells (PGC) and the cultured cell types: ES (embryonic stem) EC (embryonic carcinoma) EG (embryonic germ) cells.



plasmids in triplicates. 24 hours later we trypsinized the transfected plates and plated the ES cells on two culture dishes each. One contained ES-conditioned medium whereas the other one contained regular medium. This latter plate, therefore, had no lymphocyte inhibitory factor (LIF) to suppress the differentiation of ES cells. We trypsinized the plates three days later and subjected the cells to FACS analysis (Table 2). We found that the percent of GFP positive cells was greatly and significantly reduced (-42%, p=0.00348) in the absence of LIF in the plates transfected with the GOF18 Δ PE-EGFP S4 construct, similarly to the plate transfected

with the parental construct (-57%, p=0.01592). This suggested that the GOF constructs have started to shut down in the absence of LIF. As cell division times may also be affected in the different culture conditions, a decrease in EGFP protein or RNA levels may simply indicate the differential loss of plasmid content or episome inactivation upon cell division in cell culture and not necessarily the downregulation of the *Pou5f1* promoter upon differentiation. This may also occur. However, this doesn't account for the great level of reduction observed for the GOF constructs, because the control, Pgk promoter-EGFP, construct showed only a modest (-11%, p=0.01847) reduction in percent GFP positive cells.

Four shorter forms of GOF18∆PE-EGFP were not expressed in Hep3B cells

To test if these shorter forms could be correctly silenced in differentiated cells, we transfected them into Hep3B cells. We found that similarly to the parent GOF18 Δ PE-EGFP transgene, none of the shorter (S1-S4) forms drove EGFP expression in Hep3B cells (Fig. 4). The transfection control, red fluorescent protein (RFP), driven by the CMV promoter was expressed in each sample. Taken together, these results suggested that these four (S1-S4) shorter forms retained their restricted expression pattern specific to undifferentiated ES cells.

The shortest vector gives EGFP expression in 14.5 dpc mouse testis

We next investigated if the DE-PP regulatory sequences of *Pou511* could drive the EGFP expression in fetal mouse germ cells. We employed a simple ap**Fig. 2. Shortening of GOF18ΔPE-EGFP.** Four shortened forms (S1-S4) of the 17 kb long GOF18ΔPE-EGFP were generated by restriction endonuclease digestions: EGFP reporter with SV40 polyA termination, green box; distal enhancer (DE), orange box; Pou5f1 proximal promoter (PP), redbox; Pou5f1 exons, open boxes.

proach to accomplish this goal through electroporation of this construct into cultured mouse embryonic gonads (Nakamura *et al.*, 2002). We injected and electroporated the GOF18ΔPE-EGFP S4 into 14.5 dpc male gonads. Green fluorescence signal was found in the testicular cords where the germ cells reside. Based on this EGFP pattern it is most likely that EGFP was expressed in the fetal germ cells (Fig. 5).

Discussion

Pou5f1 expression is tightly controlled during embryogenesis. Even though the *Pou5f1* gene's transcriptional regulation is not yet completely understood, the *Pou5f1* transgene has been a very useful tool to drive reporter gene expression in pluripotent embryonic cells and allowed the isolation of primordial germ cells from mixed cell population (Szabó *et al.*, 2002, Yoshimizu *et al.*, 1999). To find the minimal regulatory regions, essential and sufficient for

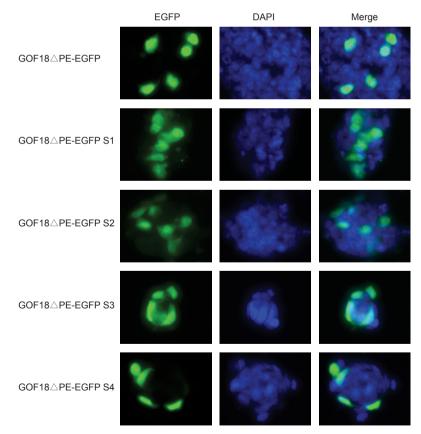


Fig. 3. Four shorter forms of GOF18 Δ PE-EGFP were expressed in mouse ES cells. Each plasmid DNA was transfected into mouse ES cells. Microscopy showed that all four shorter forms (S1-S4) of GOF18 Δ PE-EGFP expressed green fluorescent protein in ES cells, similarly to the original GOF18 Δ PE-EGFP.

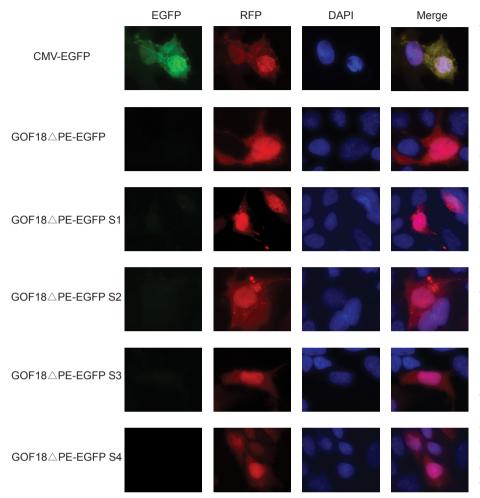


Fig. 4. Four shorter forms of GOF18∆PE-EGFP were not expressed in Hep3B cells. The expression constructs indicated to the left were transfected into Hep3B cells together with the CMV-dsRed plasmid as transfection control. Microscopy examination showed that similarly to the original GOF18ΔPE-EGFP, none of the shorter forms drove EGFP expression in differentiated Hep3B cells. However, the CMV promoter drove the control dsRed expression in each sample.

gonad at different time points. However, the suboptimal efficiency of transient transfection of ES cells and the normal downregulation of episomes in culture would pose technical limitations to these experiments and would make the interpretation difficult. Previous experiments in other species have shown that the entire Pou5f1 genomic sequences are needed for specifying the correct expression pattern of the GOF18∆PE-EGFP transgene (Kirchhof et al., 2000). Our results suggest that the sequences that are essential and sufficient for proper Pou5f1 gene regulation in the mouse may be concentrated on a shorter genomic region. To fully answer these questions one will need to develop a transgenic mouse line and test EGFP expression in embryos during development at different stages. Generation of a transgenic mouse with the shortest transgenic vector, GOF18ΔPE-EGFP S4, is needed to confirm whether the distal enhancer is sufficient to drive Pou5f1 expression exclusively in pluripotent cells in the early embryo and in

primordial germ cells after gastrulation. Nevertheless, we found that the most widely used Pou5f1 reporter construct can be short-

specifying Pou5f1 expression, we generated four shorter forms of GOF18ΔPE-EGFP and analyzed their expression patterns in pluripotent mouse ES cells and a differentiated cell line, Hep3B cells. We found that all four shorter forms were expressed in ES cell but not in Hep3B cells, similarly to the original GOF18∆PE-EGFP form. The shortest vector contained the DE and PP regions in front of the EGFP. This suggested that the shortest form harbors the essential regulatory elements in the DE that activate the promoter in pluripotent ES cells and also the essential elements that repress the promoter in differentiated Hep3B cells. We then used microinjection and gonad electroporation to introduce the shortest form into cultured fetal gonads and found that EGFP expression was localized in small foci in the testicular cords where germ cells reside. This suggests that the distal enhancer is sufficient to specify Pou5f1 expression in fetal germ cells at least in cultured gonads. There may be differences in the level of expression between the constructs, but these may not hinder visualizing pluripotent cells as long as the EGFP signal is detectable in the right place and the right time. It remains to be proven that the GFP-positive cells within the gonad are genuine gonocytes. It would be tempting to differentiate the transfected ES cells to embryoid bodies and ask whether the EGFP transgene expression becomes completely downregulated upon differentiation. One could transplant S4-transfected ES cells into blastocysts, and then perform embryo transfers to assess the EGFP expression pattern in the embryo at 6.5-7.5 dpc and in the

GOF18 APE-EGFP S4 EGFP Fig. 5. Distal enhancer of Pou5f1 drove EGFP expression in 14.5 dpc Merge mouse fetal gonad. Male gonads from 14.5 dpc mouse fetuses were injected (red arrow) with GOF18∆PE-EGFP S4 and cultured for 48 hours after electroporation. GFP positive cells were observed DIC inside the testicular cords, where the germ cells

reside.

ened to its one third without losing its specific expression pattern, at least in cultured cell types and in cultured gonad. We feel that this is useful information for future studies where visualization or genetic manipulation of pluripotent cells is desired.

Materials and Methods

Construction of shorter forms of GOF18 PE-EGFP

Four shorter forms of GOF18 Δ PE-EGFP were generated by restriction endonuclease digestion (Fig. 2). GOF18 Δ PE-EGFP S1 was made by Clal digestion, removing 7 kb of the *Pou5f1* gene sequence from the 3' end. GOF18 Δ PE-EGFP S2 was made by further digestion of S1 with EcoRV to remove 2 kb from the 5'end. GOF18 Δ PE-EGFP S3 was made by subcloning a 14 kb long Spel/NotI fragment (lacking 3 kb from the 5'end) into pBlueScript vector. GOF18 Δ PE-EGFP S4 was made by subcloning a 4.7 kb-long AfIII fragment, in which only *Pou5f1* distal enhancer and the promoter was retained, into pSL1180 vector. Pgk-EGFP plasmid was generated by replacing the CMV promoter of pEGFP-N1 (Clontech, USA) with 0.5kb of Pgk promoter.

Cell lines and transient transfection

Mouse A2 ES cells (129S1), provided by Jeffrey Mann, were grown in Dulbecco's modified Eagle's medium (DMEM, Gibco-BRL, USA) supplemented with 12% fetal bovine serum, 10^{-4} M β -mercaptoethanol, nonessential amino acids, L-glutamine, and antibiotics at standard concentration on a layer of mitomycin-inactivated LIF-producing STOC feeder cells. One day before transfection, coverslips were coated with 0.1% gelatin for 1 hour at 37°C in 12-well plates. ES cells were trypsinized and plated on a 10 cm plate for 30 minutes twice to remove feeder cells by differential attachment. Suspended cells were counted and plated into 12-well plate at the density of 5x10⁵ per well and were subsequently grown in ES-conditioned medium but without feeders. The following day, 2 µg of plasmid DNA was used for transfection with 5µl of LipoFectamine 2000 according to the standard procedure provided by the manufacturer (Invitrogene, USA). Microscopy was done 48 hours after transfection when cells were washed with phosphatebuffered saline (PBS), and fixed with 4% formaldehyde for 10 minutes at room temperature. After three time of PBS wash, cells were mounted with Prolong Gold antifade regent with DAPI (Invitrogene, USA) overnight. Slides were sealed and kept at 4°C before microscopy examination using an inverted fluorescence microscope. Alternatively, cells were trypsinized 24 hours after transfection and plated on duplicate 35 mm dishes with LIF (as supplied by freshly prepared conditioned medium) or without LIF (using regular medium). Three days later, cells were trypsinized, washed in PBS and fixed using 4% formaldehyde and analyzed by FACS for GFP positive cell content using a FACScalibur (BD Biosciences, USA) sorter.

Hep3B cells (ATCC) were grown in DMEM supplemented with 6% fetal bovine serum, nonessential amino acids, L-glutamine, and antibiotics, at standard concentrations. The transfection procedure was the same as for ES cells except the cell density was $3x10^5$ in 12-well plate, and DNA:LipoFectamine 2000 ratio used was 1 μ g : 2.5 μ l. For internal control, one tenth of CMV-dsRed plasmid DNA (0.1 μ g) was cotransfected with each shorter construct DNA.

Gonad injection

Plasmid DNA was prepared using Qiagen endotoxin free maxiprep kit (Qiagen, USA), precipitated by ethanol, and dissolved in TE buffer at a concentration of 5 μ g/µl. As described by Nakamura *et al.*, (Nakamura *et al.*, 2002), gonads were isolated from 14.5 dpc CF1 mouse embryos and

kept in cold DMEM supplemented with 10% fetal bovine serum, nonessential amino acids, L-glutamine, and antibiotics at standard concentrations. A single gonad was washed with PBS and placed between a pair of electrodes (0.2 mm diameter, 15 mm length, 1 mm distance between electrodes; Nepa Gene Co., Ltd, Chiba, Japan) on glass dish with a small volume of PBS. The anterior-posterior axis of the gonad was parallel to the electrodes. Under microscope, approximately 0.3 μ l of DNA solution was hand injected by a glass capillary with mouthpiece. Right after injection, a set of electric pulses (50V, 50-ms, 100 ms intervals, 10 times) was given by an electroporator (CUY21; Nepa Gene Co., Ltd, Japan) to the injected gonad to induce uptake of DNA by gonadal cells. Injected and electroporated gonads were cultured at 37°C with 5% CO₂ for 48 hours on a culture insert membrane (Falcon, USA) in a 24-well dish with culture medium and examined using a Zeiss fluorescent dissecting microscope.

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