

Novel Function of Sprouty4 as a Regulator of Stemness and Differentiation of Embryonic Stem Cells

Jae-Young Lee¹, Sunghyun Park¹, Kwang-Soo Kim¹, Jeong-Jae Ko¹, Soohong Lee¹,
†Keun Pil Kim² and †Kyung-Soon Park¹

¹Dept. of Biomedical Science, College of Life Science, CHA University, Seoul 06135, Korea

²Dept. of Life Science, Chung-Ang University, Seoul 06975, Korea

ABSTRACT : Sprouty (Spry) genes encode inhibitors of the receptor tyrosine kinase signaling cascade, which plays important roles in stem cells. However, the role of Spry4 in the stemness of embryonic stem cells has not been fully elucidated. Here, we used mouse embryonic stem cells (mESCs) as a model system to investigate the role of Spry4 in the stem cells. Suppression of Spry4 expression results in the decreases of cell proliferation, EB formation and stemness marker expression, suggesting that Spry4 activity is associated with stemness of mESCs. Teratoma assay showed that the cartilage maturation was facilitated in Spry4 knocked down mESCs. Our results suggest that Spry4 is an important regulator of the stemness and differentiation of mESCs.

Key words : Sprouty4, Embryonic stem cell, Differentiation

INTRODUCTION

Sprouty (Spry) was first discovered as a common antagonist of fibroblast growth factor (FGF) and epidermal growth factor (EGF) signaling pathways in *Drosophila* (Hacohen et al., 1998; Kramer et al., 1999). In a search of the Expressed Sequence Tag database, three human homologs of the fly gene, designated *hSpry1-3*, were identified (Hacohen et al., 1998). The fourth mammalian homolog, hSpry4, was later discovered in mice and humans (De Maximy et al., 1999; Leeksa et al., 2002). Emerging evidence shows that Spry specifically inhibits mitogen-activated protein kinase (MAPK) signaling activated by a wide range of ligands,

including FGF, platelet-derived growth factor (Gross et al., 2001), vascular endothelial growth factor (Impagnatiello et al., 2001), nerve growth factor (Wong et al., 2002), brain-derived neurotrophic factor (Gross et al., 2007), and glial cell line-derived neurotrophic factor (Ishida et al., 2007), but does not affect MAPK signaling activated by EGF (Sasaki et al., 2001). Spry2 reportedly potentiates biological effects induced by EGF via inhibiting EGF receptor degradation (Wong et al., 2001; Egan et al., 2002; Rubin et al., 2003). These findings suggest that Spry isoforms regulate MAPK signaling in a negative or positive manner, possibly depending on the subcellular context.

Autocrine FGF-induced extracellular signal-regulated kinase

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† Corresponding Author : Kyung-Soon Park, Dept. of Biomedical Science, College of Life Science, CHA University, Seoul 06135, Korea. Tel. : +82-31-881-7144, E-mail : kspark@cha.ac.kr, Keun Pil Kim, Dept. of Life Science, Chung-Ang University, Seoul 06975, Korea. Tel. : +82-2-820-5792, E-mail: kpkim@cau.ac.kr

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(ERK) 1/2 signaling occurs in embryonic stem cells (ESCs) and performs controversial functions to maintain the pluripotency of mouse ESCs (mESCs). It primes mESCs to initiate their differentiation in response to external differentiation-inducing stimuli as well as inducing the expression of inhibitors of FGF signaling to generate a negative-feedback loop that would facilitate reversion back to the original unprimed state (Kunath et al., 2007; Stavridis et al., 2007; Stavridis et al., 2010). ERK signaling can determine whether mesenchymal stem cells commit to the osteogenic chondrogenic lineage. Inhibition of the ERK1/2 pathway completely abolishes chondrogenic differentiation and augments osteogenic commitment (Pelaez et al., 2012). *Spry2* and *Spry4* are the most highly expressed *Spry* isoforms in human ESCs. Knockdown of *Spry2* decreases the proliferation and increases the death of human ESCs, whereas knockdown of *Spry4* enhances the survival of the cells. This indicates that the *Spry* family regulates pathways involved in the proliferation and survival of ESCs (Felfly and Klein, 2013).

Although ERK signaling is a core signaling pathway to sustain the pluripotency of mESCs and to regulate the lineage commitment of mesenchymal stem cells, it is unknown whether *Spry4* sustains the differentiation potential of mESCs and adult stem cells functions as it does in human ESCs. Here, we used mESCs as models to investigate the role of *Spry4* in the stem cells. We report that *Spry4* contributes to the maintenance of stemness and balanced differentiation of mESCs. When *Spry4* was suppressed, stemness of mESCs was disrupted and the development of cartilage was facilitated interatomas formed by mESCs. Our results reveal that *Spry4* is an important regulator of the stemness and differentiation of mESCs.

MATERIALS AND METHODS

1. Reagents and cell culture

J1 mESCs (cat #SCRC-1010) were purchased from ATCC (www.atcc.org) and maintained as described previously (Jung et al., 2012). To form embryoid bodies (EBs), mESCs were trypsinized to form a single-cell suspension and subsequently cultured on uncoated Petri dishes in ESC medium lacking leukemia inhibitory factor (LIF).

2. Genetic modification of mESCs

shRNA plasmids targeting mouse *Spry4* were purchased (TL509780, Origene, MD, USA) to generate a stable *Spry4*-knockdown cell line. mESCs were grown to 80% confluence in 24-well plates, then transfected with shRNA plasmids using Lipofectamine 2000 (Invitrogen) following the manufacturer's instructions. Stable mESCs expressing shRNA for *Spry4* were selected in 1 µg/mL of puromycin-containing culture medium. Knockdown of *Spry4* expression was confirmed with real-time reverse transcription polymerase chain reaction (RT-PCR) analysis and immunoblot analysis.

3. RNA extraction and real-time RT-PCR

Total RNA was extracted using TRIzol (Invitrogen), and 2–5 µg of total RNA was reverse-transcribed using the SuperScriptII™ First-Strand Synthesis System (Invitrogen) according to the manufacturer's instructions. Real-time RT-PCR was performed using cDNA with the Quantitect SYBR Green PCR Kit (Qiagen, CA, USA). Reactions were performed in triplicate using an Exicycler™ 96 real-time quantitative thermal block (Bioneer). For quantification, target genes were normalized against the *glyceraldehyde 3-phosphate dehydrogenase* gene.

4. Immunoblotting

Whole-cell extracts were prepared, and 20–50 µg of protein was resolved by SDS-PAGE and blotted using antibodies against *Spry4* (ab59785, Abcam, MA, USA), phospho-Stat3 (Tyr-705; #9131, Cell Signaling Technology, Beverly, MA, USA), Stat3 (sc-482, Santa Cruz, CA, USA), cMyc

(ab32072, Abcam), Nanog (ab80892, Abcam), Sox2 (sc-20088, Santa Cruz), phospho-ERK (M9692, Sigma), ERK2 (sc-154, Santa Cruz), and β -actin (sc-47778, Santa Cruz). Immunoreactivity was detected by enhanced chemiluminescence (Amersham, Buckinghamshire, England).

5. Teratoma formation

For the teratoma formation assay, cells were trypsinized, and 5×10^5 cells were suspended in a DMEM/Matrigel solution (BD Biosciences Inc., Franklin Lakes, NJ, USA) (1:1 ratio (v/v)). The cell/Matrigel suspension was injected subcutaneously into NOD/SCID mice (Charles River Laboratories, Yokohama, Japan). Teratoma formation was examined for 6 weeks after injection. The experiments were reviewed and approved by the Institutional Animal Care and Use Committee of CHA University. All procedures were performed in accordance with the Guidelines for the Care and Use of Laboratory Animals published by the US National Institutes of Health (NIH publication no. 85-23, revised 1996).

6. Statistical analysis

Graphical data are presented as means \pm SD. Each experiment was performed at least three times and subjected to statistical analysis. Statistical significance between two groups was determined using Student's *t*-test. A *p* value below 0.05 was considered significant. Statistical analysis was performed using the SAS statistical package v.9.13 (SAS Inc., Cary, NC).

RESULTS AND DISCUSSION

Activation of the FGF-ERK signaling pathway is required for naïve mESCs to initiate lineage differentiation by exiting their self-renewal status. Therefore, FGF signaling should be tightly regulated at multiple levels in ESCs. We previously reported that Spry1, a member of the Spry family of proteins, is necessary to balance the self-renewal and pluripotency of

mESCs by inhibiting ERK1/2 signaling under self-renewal culture conditions (Jung et al., 2012). To increase our understanding of the regulatory mechanisms of ERK1/2 signaling in mESCs, we further investigated the role of Spry4 in mESCs.

We first examined whether Spry4 modulates phosphorylation of ERK1/2 in mESCs. When Spry4 expression was suppressed by transfection of shRNA-targeting *Spry4*, ERK phosphorylation was significantly increased, suggesting that Spry4 suppresses ERK1/2 signaling in mESCs (Fig. 1A). We next engineered mESCs expressing green fluorescent protein and *Spry4*-targeting shRNA to generate a stable Spry4-knockdown mESC line (Spry4 KD), and then examined the effect of Spry4 suppression on the morphology and proliferation of these cells. Although Spry4 KD showed strong phosphorylation of ERK, it maintained the undifferentiated round shape of mESCs and alkaline phosphatase activity (Fig. 1B, 1C). However, the proliferation of Spry4 KD was lower than that of control cells (Fig. 1D). To examine whether altered Spry4 expression affects mESC self-renewal, we examined the expression of undifferentiated ESC markers. Expression of Oct4 and Sox2 was decreased at the mRNA and protein levels, whereas expression of c-Myc, Nanog, and phosphorylated Stat3 was not affected by suppression of Spry4 expression (Fig. 1E). We examined the ability of Spry4 KD to maintain an undifferentiated state by analyzing the efficiency of secondary EB formation (Felfly & Klein, 2013). The ability of Spry4 KD to maintain an undifferentiated state was comparable to that of control mESCs, indicating that alteration of Spry4 did not affect the self-renewal of mESCs, despite the finding that suppression of Spry4 decreased cell proliferation and reduced expression of Sox2 and Oct4 (Fig. 1F). Our results are consistent with previous reports that Fgf4-null ESCs behave normally in the undifferentiated state in the presence of LIF, although they cannot exit the pluripotent state and initiate differentiation when LIF is removed (Kunath et al., 2007).

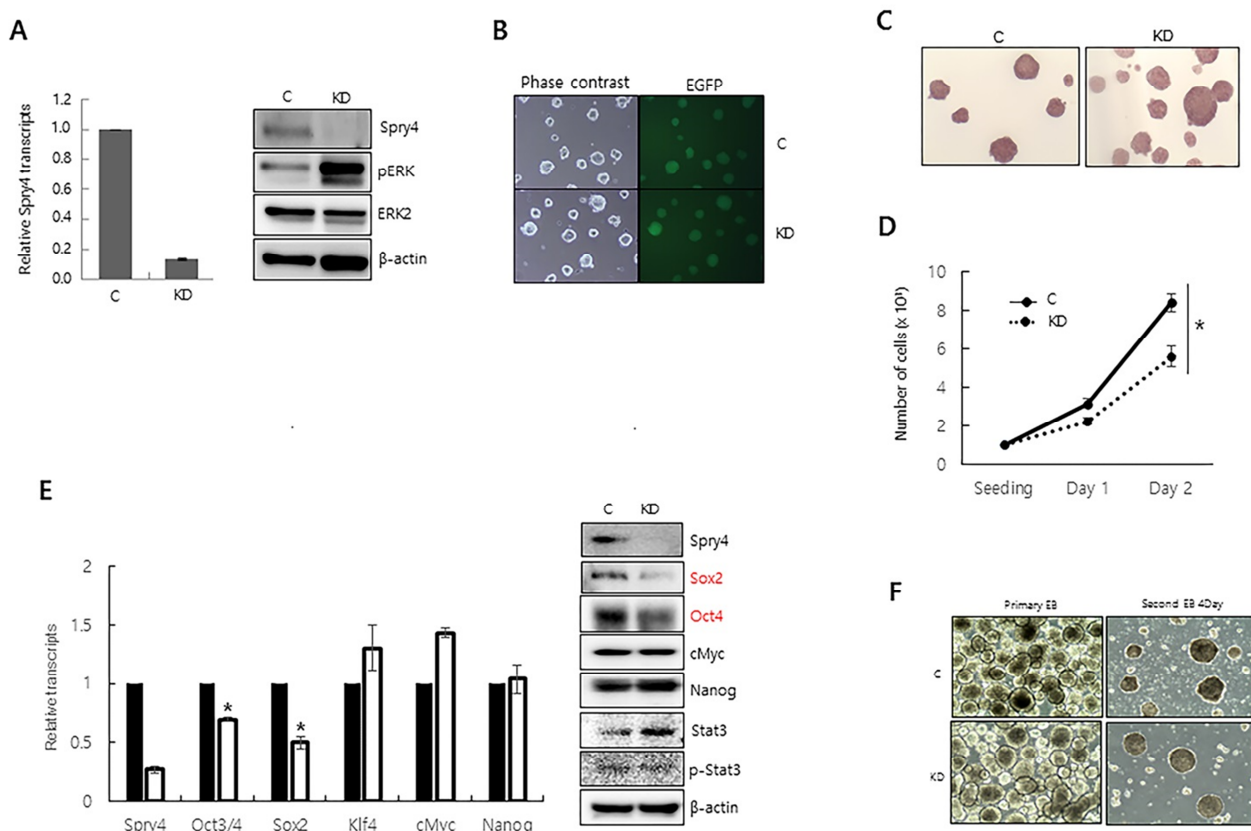


Fig. 1. The effect of Spry4 suppression on the self-renewal of mESCs. (A) Expression of Spry4 in Spry4-knockdown mESCs (KD) was quantitatively analyzed by real-time RT-PCR (left) and immunoblotting (right). The effect of Spry4 suppression on the level of phospho-ERK was analyzed by immunoblotting (right). (B) The morphologies of Spry4 KD and control mESCs were observed by phase contrast microscopy and fluorescence microscopy. (C) The growth of Spry4 KD and control mESCs were compared by counting cells at 24 h (Day 1) and 48 h (Day 2) after seeding. (D) Expression of self-renewal genes was analyzed by real-time RT-PCR (left) and immunoblotting (right). (E) Alkaline phosphatase staining of Spry4 KD and control mESCs. (F) Primary and secondary EB formation abilities of Spry4 KD and control mESCs. All values are means ± standard deviation of at least three independent experiments. * $P < 0.05$ based on the student's t -test. Abbreviations: C, mESCs with control shRNA integration.

Autocrine FGF/ERK signaling is a pro-differentiation signal (Kunath et al., 2007); therefore, we hypothesized that the differentiation potential of Spry4 KD was higher than that of control mESCs. To test this hypothesis, we analyzed the impact of *Spry4* suppression on the differentiation of mESCs in xenografts. As expected, Spry4KD gave rise to larger teratomas than control mESCs (Fig. 2A). We next analyzed differentiation of the three germ lineages in teratomas by hematoxylin-eosin (H&E) staining. The pro-

portions of mesoderm tissues, including cartilage, bone, neuroglia and primitive mesoderm, as well as endoderm tissues, such as glands, were significantly increased in 6-week-old teratomas formed by Spry4KD (Fig. 2B, 2C). On the other hand, the proportion of immature cartilage was significantly lower in teratomas formed by Spry4 KD than in control teratomas, suggesting that the balanced differentiation potential of mESCs was disturbed by suppression of *Spry4*. The expression level of *Spry4* is much lower than

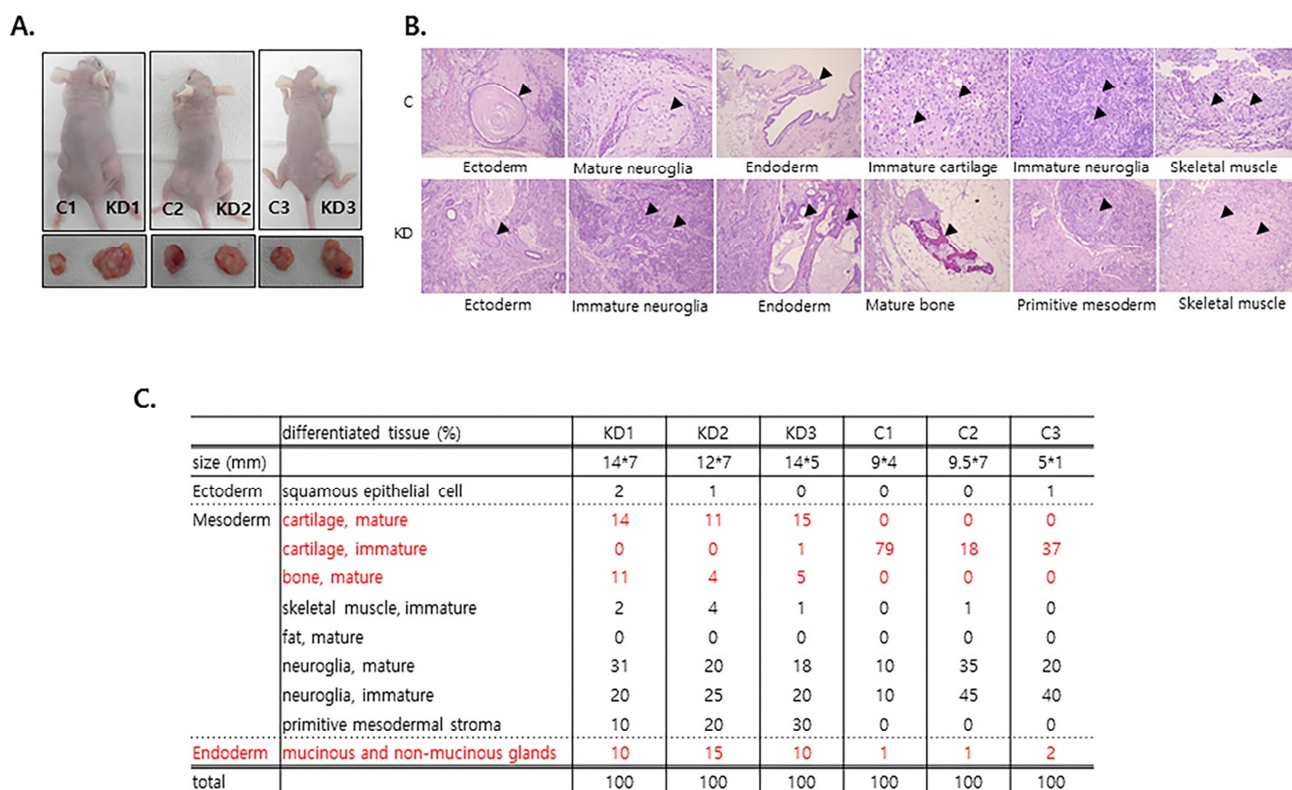


Fig. 2. Teratoma formation and differentiation of the three germ layers by Spry4 KD. (A) Spry4KD (KD) and control mESCs were injected into NOD/SCID mice and teratoma development was examined. The size of teratomas was examined 6 weeks after injection. (B) H&E staining and data analysis of teratoma sections were performed by a pathologist. Mature or immature germ layer differentiation is indicated by arrows. (C) Table of the relative percentages of differentiated tissues in teratomas formed by Spry4 KD and control mESCs. Abbreviations: C, mESCs with control shRNA integration.

that of *Spry1* in mESCs (Jung et al., 2012). Despite its slow expression level, our results show that *Spry4* plays a major role in control of the differentiation ability of mESCs.

In our analysis of the lineage differentiation of teratomas formed by mESCs, we noticed that mature cartilage was highly developed in teratomas formed by Spry4 KD in comparison to control teratomas (Fig. 2C). Cartilage loss is a typical degenerative disease because cartilage has a limited regeneration potential; however, effective therapy remains elusive. Therefore, further work to investigate the detailed mechanism by which Spry4 enhances cartilage differentiation will contribute to enhance the differentiation efficiency of stem cells.

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