

Murine pluripotent stem cells that escape differentiation inside teratomas maintain pluripotency

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Background. Pluripotent stem cells (PSCs) offer immense potential as a source for regenerative therapies. The teratoma assay is widely used in the field of stem cells and regenerative medicine, but there is limited understanding of the *in vivo* teratoma differentiation system.

Methods. We utilized PSCs expressing green fluorescent protein (GFP) under the control of the *Pou5f1* promoter to study the persistence of potential pluripotent cells during teratoma formation *in vivo*. OCT4-MES (mouse embryonic stem cells) were isolated from the blastocysts of 3.5-day OCT4-EGFP (enhanced green fluorescent protein) mice embryos, and TG iPS 1-7 (induced pluripotent stem cells) were generated from mouse embryonic fibroblasts (MEFs) from 13.5-day OCT4-EGFP mice embryos by infecting them with a virus carrying OCT4, SOX2, KLF4 and c-MYC. These pluripotent cells were characterized according to their morphology and expression of pluripotency markers. Their differentiation ability was studied with *in vivo* teratoma formation assays. Further differences between pluripotent cells were examined by real-time quantitative PCR (qPCR).

Results. The results showed that several OCT4-expressing PSCs escaped differentiation inside of teratomas, and these escaped cells (MES-FT, GFP-positive cells separated from OCT4-MES-derived teratomas; and iPS-FT, GFP-positive cells obtained from teratomas formed by TG iPS 1-7) retained their pluripotency. Interestingly, a small number of GFP-positive cells in teratomas formed by MES-FT and iPS-FT (MES-ST, GFP-positive cells isolated from MES-FT-derived teratomas; iPS-ST, GFP-positive cells obtained from teratomas formed by iPS-FT) were still pluripotent, as shown by alkaline phosphatase (AP) staining, immunofluorescent staining and PCR. MES-FT, iPS-FT, MES-ST and iPS-ST cells also expressed several markers associated with germ cell formation, such as *Dazl*, *Stella* and *Stra8*.

Conclusions. In summary, a small number of PSCs escaped differentiation inside of teratomas, and these cells maintained pluripotency and partially developed towards germ cells. Both escaped PSCs and germ cells present a risk of tumor formation. Therefore, medical workers must be careful in preventing tumor formation when stem cells are used to treat specific diseases. 

11 Abstract

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17 teratoma formation *in vivo*. OCT4-MES (mouse embryonic stem cells) were isolated from the
18 blastocysts of 3.5-day OCT4-EGFP mice (transgenic mice express EGFP cDNA under the control
19 of the *Pou5f1* promoter) embryos, and TG iPS 1-7 (induced pluripotent stem cells) were generated
20 from mouse embryonic fibroblasts (MEFs) from 13.5-day OCT4-EGFP mice embryos by infecting
21 them with a virus carrying OCT4, SOX2, KLF4 and c-MYC. These pluripotent cells were
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30 teratomas; iPS-ST, GFP-positive cells obtained from teratomas formed by iPS-FT) were still
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32 PCR. MES-FT, iPS-FT, MES-ST and iPS-ST cells also expressed several markers associated with
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39 **Introduction**

40 Pluripotent stem cells (PSCs), including embryonic stem cells (ESCs) and induced pluripotent
41 stem cells (iPSCs), have the potential to differentiate into all cell types of the body *in vitro* through
42 embryoid body formation or *in vivo* through teratoma formation. Due to these characteristics, stem
43 cells provide an option for treating a multitude of clinical problems, such as myocardium damage
44 after heart infarction, spinal cord damage after mechanical injury, brain damage after stroke, age-
45 related macular degeneration of the retina, liver damage, extensive skin burns, Parkinson's disease,
46 and diabetes (Abdelalim et al., 2014; Lodi et al., 2011; Orlic et al., 2001; Ratajczak et al., 2016).
47 When transplanted into immune-compromised mice, undifferentiated PSCs can form teratomas,
48 consisting of multiple tissue types derived from all three germ layers (Przyborski, 2005;
49 Takahashi and Yamanaka, 2006). As such, there have been many efforts to recapitulate an *in vivo*
50 developmental environment. For example, neural stem cells (NSCs) have been differentiated *in*
51 *vivo* through teratoma formation, and pure NSC populations exhibit properties similar to those of
52 brain-derived NSCs (Hong et al., 2016). Similarly, fully functional and engraftable hematopoietic
53 stem/progenitor cells (HSPCs), along with functional myeloid and lymphoid cells, have been
54 isolated from teratomas when human iPSCs were transplanted into immunodeficient mice
55 (Amabile et al., 2013; Suzuki et al., 2013). In addition, the teratoma assay can be applied to assess
56 the safety of human PSC-derived cell populations that are used for therapeutic application since a
57 small number of undifferentiated cells contaminating a given transplant material can be efficiently
58 detected by their multi-lineage differentiation ability (Stachelscheid et al., 2013).

59 However, the intrinsic self-renewal and pluripotency qualities of PSCs that make them
60 therapeutically promising are responsible for an equally fundamental tumorigenic risk (Lee et al.,
61 2013). Studies on teratomas will contribute to a better understanding of their stepwise development
62 processes and underlying molecular mechanisms and may provide helpful information for the
63 development of tissue engineering technologies (Aleckovic and Simon, 2008). These facts
64 prompted us to address the additional characteristics of teratoma growth and differentiation after
65 PSC injection.

66 In the present study, we aimed to isolate OCT4-expressing cells that escaped differentiation inside
67 of growing teratomas and to determine whether OCT4-expressing cells still possess self-renewal
68 and pluripotency abilities.

69 **Materials & methods**

70 All animal experiments were approved by the Animal Care and Use Committees of the State Key
71 Laboratories for Agrobiotechnology, College of Biological Sciences, China Agricultural
72 University (Approval number: SKLAB-2016-05-01). Briefly, mice were bred in a 12/12 h
73 light/dark period and sacrificed by cervical vertebra dislocation.

74 **Mouse strains**

75 OCT4-GFP transgenic mice (Model Animal Research Center of Nanjing University) express
76 EGFP (enhanced green fluorescence protein) cDNA under the control of the *Pou5f1* promoter,
77 which is active in pluripotent stem cells. This strain is useful for isolating pluripotent stem cells,
78 as they specifically express green fluorescent protein. These OCT4-GFP transgenic mice were the
79 source of the OCT4-MES and OG2 MEFs (mouse embryonic fibroblasts of 13.5-day OCT4-EGFP

80 mice embryos) used in this study.

81 **Derivation of MES and generation of iPSCs**

82 To obtain OCT4-MES, uteri containing E3.5 embryos were isolated from timed pregnancies and
83 transferred individually to the wells of a 24-well plate with irradiated MEF feeders. After 5 days
84 of incubation, embryo outgrowths were separated from trophectoderm, individually picked, and
85 expanded in MES medium (DMEM supplemented with 15% FBS, L-glutamine, nonessential
86 amino acids, β -mercaptoethanol, and 1,000 U/ml LIF).

87 OG2 MEFs were cultured in MEF medium (Dulbecco's modified Eagle's medium (DMEM)
88 supplemented with 10% FBS, L-glutamine and nonessential amino acids); infected with
89 retroviruses generated from pMX retroviral vectors encoding mouse *Pou5f1*, *Sox2*, *Klf4* and *c-*
90 *Myc*; and cultured on irradiated MEF feeder cells in MES medium. Subsequently, a single ESC-
91 like colony was individually picked and expanded on feeders to establish stable lines. Both OCT4-
92 MES and iPSCs originated from male embryos. Additional details can be found in our previous
93 study (Pei et al., 2015).

94 **Immunofluorescence**

95 Cells were fixed with 4% paraformaldehyde, permeabilized with 0.1% Triton X-100, and blocked
96 with 2% BSA. The cells were then stained with primary antibodies against OCT4 (Abcam,
97 ab19857, 1:500), SOX2 (Abcam, ab97959, 1:1000), NANOG (Abcam, ab80892, 1:500) and
98 SSEA1 (Abcam, ab16285, 1:200), followed by staining with the respective secondary antibodies
99 conjugated to Alexa Fluor (Invitrogen). Finally, cells were counterstained with DAPI (Sigma,
100 D9542).

101 RNA purification and cDNA preparation

102 Feeders were removed by plating ESCs on a gelatin-coated dish for 30 min, and unattached cells
103 were collected by centrifugation. Total RNA was extracted from pure PSCs using Trizol reagent
104 according to the manufacturer's instructions (Invitrogen). RNA was reverse-transcribed using
105 oligo-dT and M-MLV Reverse Transcriptase (Promega, Madison, WI).

106 Real-time quantitative PCR

107 qPCR was performed on a LightCycler 480 II Real-Time PCR System (Roche, Basel, Switzerland)
108 using the LightCycler 480 SYBR Green I Master Mix (Roche, Basel, Switzerland). The specific
109 calculation method for qPCR data is referred to the description about three separate RNA
110 preparations by Livak and Schmittgen (Livak and Schmittgen, 2001). And the most conservative
111 test, owing to its nonparametric nature, is the Wilcoxon two group test, which is distribution-
112 independent. So, we performed the SAS program developed for Wilcoxon two group test as shown
113 in methodology article of Joshua S Yuan and his colleagues (Yuan et al., 2006).

114 The primers used for qPCR and PCR and listed below.

Gapdh-F	AGGTCGGTGTGAACGGATTTG
Gapdh-R	TGTAGACCATGTAGTTGAGGTCA
β -tubulin-F	TGAGGCCTCCTCTCACAAGTA
β -tubulin-R	CCGCACGACATCTAGGACTG
EF1 α -F	GTGTTGTGAAAACCACCGCT
EF1 α -R	AGGAGCCCTTTCCCATCTCA
Pou5f1-F	GTTGGAGAAGGTGGAACCAA

Pou5f1-R	CTCCTTCTGCAGGGCTTTC
Sox2-F	AAGGGTTCTTGCTGGGTTTT
Sox2-R	AGACCACGAAAACGGTCTTG
klf4-F	CTCTGCTCCCGTCCTTCTC
klf4-R	AGAGTTCCTCACGCCAAC
Nanog-F	TTCTTGCTTACAAGGGTCTGC
Nanog-R	AGAGGAAGGGCGAGGAGA
Rex1-F	CAGTTCGTCCATCTAAAAAGGGAGG
Rex1-R	TCTTAGCTGCTTCCTTGAACAATGCC
Tbx3-F	ATCGCCGTTACTGCCTATCA
Tbx3-R	TGCAGTGTGAGCTGCTTTCT
Lin28a-F	GTCTTTGTGCACCAGAGCAAG
Lin28a-R	ATGGATTCCAGACCCTTGGC
Nr5a2-F	TAGGACCGGAAAGCGTCTGC
Nr5a2-R	GCTTCCGTCTCCACTTTGGG
Dazl-F	GCCCGCAAAGAAGTCTGTG
Dazl-R	ACCAACAACCCCTGAGATG
Stella-F	GAGAAGACTTGTTTCGGATTGAGC
Stella-R	CATCGTCGACAGCCAGGG
Stra8-F	CTCCTCCTCCACTCTGTTGC
Stra8-R	GCGGCAGAGACAATAGGAAG
Vasa-F	ACCAAGATCAGGGGACACAG
Vasa-R	TAACCACCTCGACCACTTCC

115 **Teratoma production and analysis**

116 Approximately 1×10^6 PSCs were suspended in 150 μ l of PBS (phosphate buffered solution) and
 117 injected into NOD/SCID mice to form teratomas. Three weeks after injection, the teratomas were

118 harvested, fixed overnight with 4% paraformaldehyde, embedded in paraffin, sectioned, HE
119 stained or immunostained (GFP, 1: 200, Cell Signaling Technology, Cat. #2956), and analyzed.

120 **Statistical analysis**

121 The results are presented as the mean \pm standard deviation (MS \pm SD). The significance of
122 differences was analyzed using Wilcoxon two group test.

123 **Results**

124 **Both OCT4-MES and TG iPS 1-7 are pluripotent**

125 OCT4-EGFP mice express green fluorescent protein under the control of the pluripotency-
126 associated *Pou5fl* promoter and are widely used to study the function of PSCs (Pei et al., 2015).
127 These mice were used to generate mouse embryonic stem cells (MES) and iPSCs. OCT4-MES
128 were isolated from the blastocysts of 3.5-day OCT4-EGFP mice embryos, while other mice were
129 selected to prepare MEFs after day 13.5. The isolated MEFs were used to generate iPSCs by
130 infecting them with a virus carrying OCT4, SOX2, KLF4 and c-MYC. Then, TG iPS 1-7 was
131 selected from the isolated iPSC clones.

132 Both OCT4-MES and TG iPS 1-7 were maintained on feeder cells in the presence of leukemia
133 inhibitory factor. They both exhibited typical MES-like morphologies (Figure 1a and b).
134 Immunofluorescent staining confirmed the expression of the three master transcription factors
135 (OCT4, NANOG and SOX2) as well as ESC-specific surface marker SSEA-1 in OCT4-MES and
136 TG iPS 1-7 (Figure 1c, d). The PCR results further demonstrated that these cells expressed
137 pluripotency marker genes, including *Pou5fl*, *Sox2*, *Nanog*, *Rex1*, *Tbx3*, *Nr5a2*, *Klf4* and *Lin28a*
138 (Figure 2). Next, *in vivo* teratoma formation assays were performed to further validate the

139 pluripotency of OCT4-MES and TG iPS 1-7. Approximately 1×10^6 PSCs were suspended in 150
140 μ l of PBS and injected into non-obese diabetic/severe combined immunodeficient (NOD/SCID)
141 mice to form teratomas. Three weeks after injection, OCT4-MES and TG iPS 1-7 formed
142 teratomas *in vivo*, and hematoxylin and eosin (H&E) staining confirmed the formation of all three
143 germ layers in each teratoma (Figure 3a, b). These results revealed that OCT4-MES and TG iPS
144 1-7 were pluripotent. Interestingly, we observed OCT4-positive cells growing in clusters in the
145 teratoma masses formed by OCT4-MES and TG iPS 1-7 (Figure 3c, d).

146 **OCT4-positive cells from OCT4-MES and TG iPS 1-7 teratomas have self-renewal and**
147 **pluripotency qualities**

148 To quantify the fraction of OCT4-positive pluripotent cells in teratomas generated by OCT4-MES
149 and TG iPS 1-7, we cut the teratomas into pieces and digested them with trypsin and then cultured
150 the cells in MEF medium. Three days later, we found that most of these cells separated from
151 OCT4-MES and that TG iPS 1-7-derived teratomas had the morphology of mouse embryonic cells,
152 but a small number of cells were round and expressed GFP (Figure 4a, c). After picking these cells
153 and culturing them in MES medium, we found that they had typical MES-like morphologies, and
154 they were AP-positive (Figure 4b, d). We named these cells MES-FT and iPS-FT, which were
155 derived from OCT4-MES and TG iPS 1-7, respectively. OCT4-expressing MES-FT and iPS-FT
156 cells were grown in the presence of leukemia inhibitory factor, and they expressed pluripotency
157 marker genes, including *Pou5f1*, *Sox2*, *Nanog*, *Rex1*, *Tbx3*, *Nr5a2*, *Klf4* and *Lin28a* (Figure 2).
158 The immunostaining results showed that these colonies were positive for OCT4, NANOG, SOX2
159 and SSEA-1 (Figure 5a, b). We performed *in vivo* teratoma formation assays to further validate

160 the pluripotency of MES-FT and iPS-FT. MES-FT and iPS-FT formed teratomas *in vivo*, and the
161 hematoxylin and eosin (H&E) staining results confirmed the formation of all three germ layers in
162 each teratoma (Figure 5c, d). As in the results described above, there were also OCT4-positive
163 pluripotent cells in the teratomas formed by MES-FT and iPS-FT (Figure 5e, f).

164 **OCT4-positive cells from MES-FT and iPS-FT teratomas are still pluripotent**

165 We discovered several round and bright cells expressing OCT4-GFP under a microscope in cells
166 separated from teratomas formed by MES-FT and iPS-FT cells (Figure 6a, c). These round and
167 bright cells formed AP positive clones (Figure 6b, d). We named these cells MES-ST and iPS-ST.
168 OCT4-expressing MES-ST and iPS-ST cells also expressed pluripotency marker genes, including
169 *Pou5f1*, *Sox2*, *Nanog*, *Rex1*, *Tbx3*, *Nr5a2*, *Klf4* and *Lin28a* (Figure 2), and the immunostaining
170 results demonstrated that they expressed the stemness regulators OCT4, NANOG, SOX2 and
171 SSEA-1 (Figure 5e, f).

172 The above results showed that OCT4-MES, TG iPS 1-7, MES-ST, iPS-ST, MES-ST and iPS-ST
173 had pluripotency characteristics. However, MES-FT, iPS-FT, MES-ST and iPS-ST were survivors
174 of the differentiation environment, so we wanted to know whether there were differences among
175 these cells. Thus, we next investigated their differences.

176 **OCT4-positive cells separated from teratomas highly express germ cell marker genes**

177 To explore the gene expression patterns of OCT4-MES, TG iPS 1-7, MES-FT, iPS-FT, MES-ST
178 and iPS-ST, cDNA was prepared from these cells without feeders for gene expression analysis.
179 First, we detected the expression of pluripotency genes. When normalized to the values for OCT4-
180 MES cells, the expression level of *Pou5f1* was higher in MES-FT, and that of *Lin28a* was higher

181 in both MES-FT and MES-ST cells, but there were no differences in the *Naong* expression levels
182 between these three cell lines (Figure 7a). When normalized to the values for TG iPS 1-7 cells,
183 iPS-FT and iPS-ST both highly expressed *Pou5fl* and *Nanog* (Figure 7b). However, there were no
184 differences in the expression level of *Lin28a* (Figure 7b). The expression of pluripotency marker
185 genes in these cells varied slightly, but they were all within reasonable levels. Thus, these cell
186 types were all pluripotent.

187 Previous results have shown that PSCs that escape from differentiation inside of embryonic bodies
188 express several markers associated with germ cell formation (Attia et al., 2014). As such, we
189 further assayed the differences between the expression levels of important germ cell-specific genes
190 (*Dazl*, *Stella*, *Stra8*, *Vasa*) in MES-FT, iPS-FT, MES-ST and iPS-ST (Figure 7c, d). When
191 normalized to the values for OCT4-MES, *Dazl* and *Stella* were more highly expressed in MES-FT
192 cells, and the expression level of *Stra8* was elevated nine-fold and ten-fold in MES-FT and MES-
193 ST, respectively.

194 Similarly, iPS-FT and iPS-ST highly expressed *Dazl*, *Stra8* and *Vasa* than TG iPS 1-7. iPS-FT
195 also highly expressed *Stella*. The above results show that OCT4-positive cells separated from
196 teratomas have elevated expression of several markers associated with germ cell formation, such
197 as *Dazl*, *Stella* and *Stra8*.

198 Discussion

199 ESCs and iPSCs are characterized by their ability to develop into any cell type of the adult
200 organism. As such, they can be widely applied to the treatment of many diseases. This is especially
201 true for iPSCs, as they do not present ethical issues.

202 A previous report demonstrated the presence of undifferentiated human ESCs expressing the
203 surface marker CD133 (Ritnerand Bernstein, 2010). However, no additional research has been
204 performed to investigate the characteristics of those undifferentiated cells in teratomas. Therefore,
205 in this study, we isolated OCT4-GFP positive cells, MES-FT and iPS-FT, from teratomas
206 generated by OCT4-MES and TG iPS 1-7, respectively. MES-FT and iPS-FT exhibit MES-like
207 morphologies, express pluripotency marker genes and proteins, and can generate all three germ
208 layers in an *in vivo* differentiation model. We discovered that there were still pluripotent cells in
209 the teratomas formed by MES-FT and iPS-FT, so we separated them from the teratoma mass and
210 named them MES-ST and iPS-ST. Further study confirmed that these isolated cells (MES-ST and
211 iPS-ST) retained pluripotency and were capable of differentiation. From these results, it can be
212 inferred that a subset of PSCs escape differentiation during *in vivo* differentiation, and the escaped
213 cells retain their PSC characteristics in the appropriate environment. Since the escaped PSCs
214 (MES-FT, iPS-FT, MES-ST and iPS-ST) still possessed PSC-like characteristics, these cells may
215 progress to tumor formation at an undefined later time point.

216 Bottai et al. reported that they used 5×10^5 undifferentiated murine ESCs to cure spinal cord injury.
217 However, some of the transplanted ESCs were found as dense aggregates in the tissue (Bottai et
218 al., 2010). This result supports our view that ESCs can be maintained *in vivo*. Another study
219 showed that transplantation of $1-2 \times 10^6$ MES cells into SV129 mice led to tumor formation in
220 100% of cases, whereas transplantation of 5×10^5 cells produced tumors in 2 of 6 mice and
221 transplantation of 1×10^5 ESCs gave rise to tumor formation in 1 of 6 transplanted mice within 100
222 days (Dressel et al., 2008). It can be deduced that there is likely a niche within teratomas that nurse

223 PSCs, and the number of cells determines the niche environment. The more PSCs used for
224 transplantation, the higher probability of tumor formation.

225 The escaped PSCs (MES-FT, iPS-FT, MES-ST and iPS-ST) showed slight similarities to
226 primordial germ cells (PGCs), as shown by the high expression of *Pou5f1*, *Dazl*, *Stella* and *Stra8*
227 in MES-FT, MES-ST, iPS-FT, and iPS-ST. *Pou5f1*, *Dazl*, *Stella*, *Stra8* and *Vasa* are well-known
228 germ cell markers, and they are also commonly expressed in ESCs (Cauffman et al., 2005; Kehler
229 et al., 2004; Tedesco et al., 2009; Toyooka et al., 2000; Wongtrakoongate et al., 2013). *Stra8* is
230 required for the chromosomal program of meiotic prophase (Soh et al., 2015). *Dazl*, an intrinsic
231 meiotic competence factor, is required for *Stra8*-mediated initiation of meiosis in germ cells (Lin
232 et al., 2008). Overexpression of *Stra8* and *Dazl* genes promotes the transdifferentiation of
233 mesenchymal stem cells and ESCs *in vitro* toward PGCs (Li et al., 2017; Shi et al., 2014). The
234 elevated expression of *Pou5f1*, *Dazl*, *Stella* and *Stra8* might indicate that the GFP positive cells
235 separated from teratomas partially develop towards germ cells. This suggests that it is possible to
236 isolate PGCs from teratoma differentiation models.

237 **Conclusions**

238 In summary, we found a small number of OCT4-expressing PSCs that escaped differentiation
239 inside teratomas. The escaped cells kept their unique properties of self-renewal and pluripotency
240 and were able to form teratomas *in vivo*. They also highly expressed several markers associated
241 with germ cell formation, such as *Pou5f1*, *Dazl*, *Stella* and *Stra8*, suggesting that these cells may
242 partially differentiate into germ cells. Therefore, this study serves as a warning that medical
243 workers using stem cells to treat specific diseases must pay careful attention to prevent tumor

244 formation because OCT4-expressing cells retain pluripotency, and it is feasible to isolate germ
245 cells from teratomas. This study of PSCs that remain undifferentiated within teratomas has
246 provided critical information for further investigation of the applications of stem cell therapy and
247 for obtaining germ cells from *in vivo* differentiation models.

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321

Figure 1

OCT4-MES and TG iPS 1-7 are pluripotent

(a and b) Phase contrast images of OCT4-MES (a) and TG iPS 1-7 (b). Both types of cells exhibit typical MES-like morphologies.

(c and d) Immunofluorescent staining showing that both OCT4-MES (c) and TG iPS 1-7 (d) express the pluripotency markers OCT4, NANOG, SSEA1 and SOX2.

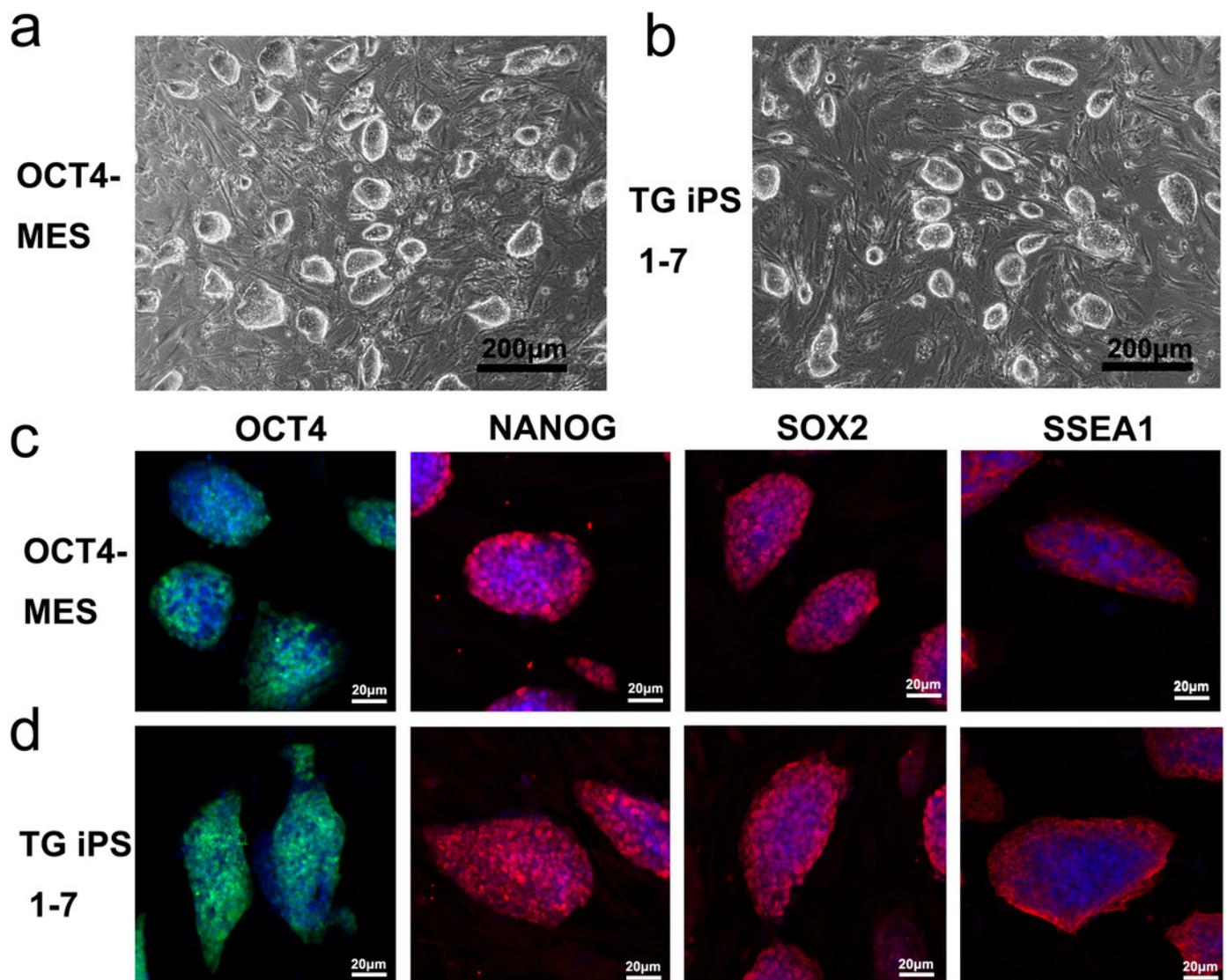


Figure 2

OCT4-MES, MES-FT, MES-ST, TG iPS 1-7, iPS-FT and iPS-ST express pluripotency genes.

Expression of pluripotency marker genes was evaluated by PCR, showing that OCT4-MES, MES-FT, MES-ST, TG iPS 1-7, iPS-FT and iPS-ST all express pluripotency marker genes, including *Pou5f1*, *Sox2*, *Nanog*, *Rex1*, *Tbx3*, *Nr5a2*, *Klf4* and *Lin28a*.

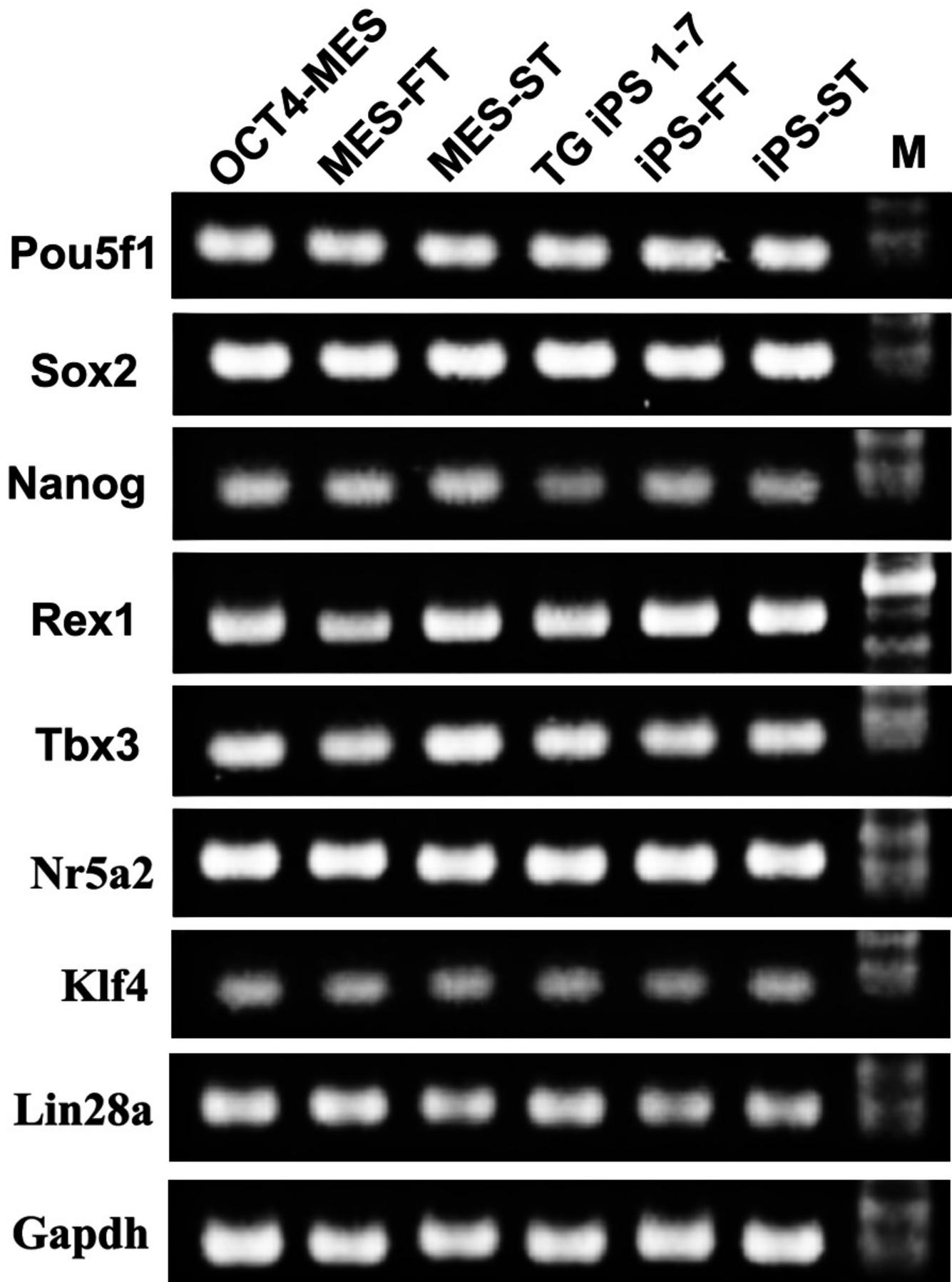


Figure 3

GFP-positive pluripotent cells are present in teratomas generated by OCT4-MES and TG iPS 1-7 .

(a and b) Hematoxylin and eosin staining of teratomas derived from OCT4-MES (a) and TG iPS 1-7 (b). Products of all three germ layers are seen in the image: Ectoderm: epidermis with keratin (left). Mesoderm: smooth muscle (middle). Endoderm: gastrointestinal lining cells/glands (right). Specified cells are indicated by arrows.

(c and d) Immunohistochemistry to detect the presence of GFP-positive pluripotent cells in teratomas generated by OCT4-MES (c) and TG iPS 1-7 (d). GFP-positive pluripotent cells were stained in brown with anti-GFP primary antibodies.

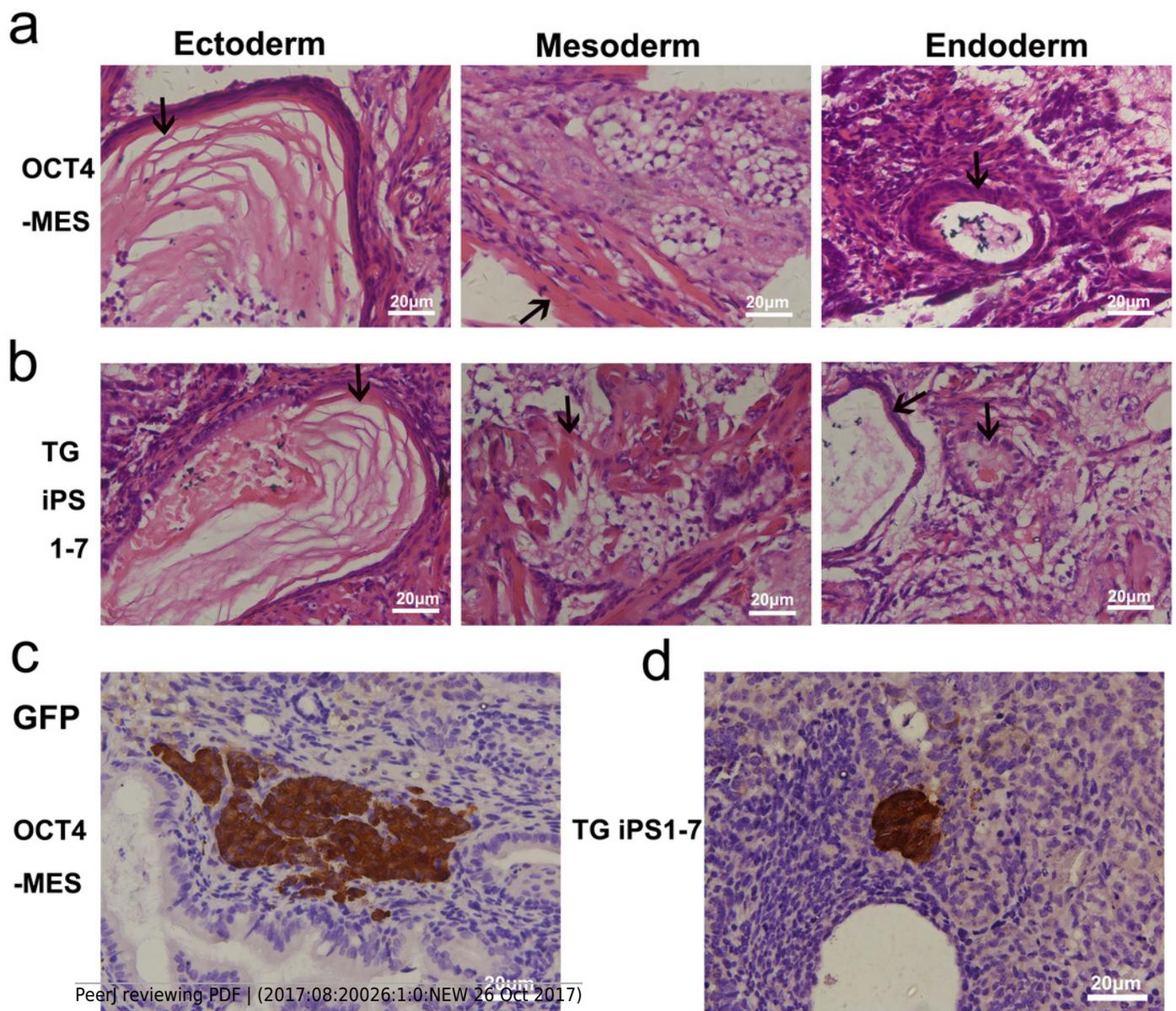


Figure 4

OCT4-positive pluripotent cells isolated from teratomas have typical mouse embryonic cell morphology .

(a and c) A small number of OCT4-GFP positive cells were found among teratoma cells generated by OCT4-MES (a) and TG iPS 1-7 (c) cultured in MEF medium.

(b and d) MES-FT (b) and iPS-FT (d) have typical mouse embryonic cell morphology and are AP-positive when cultured in MES medium.

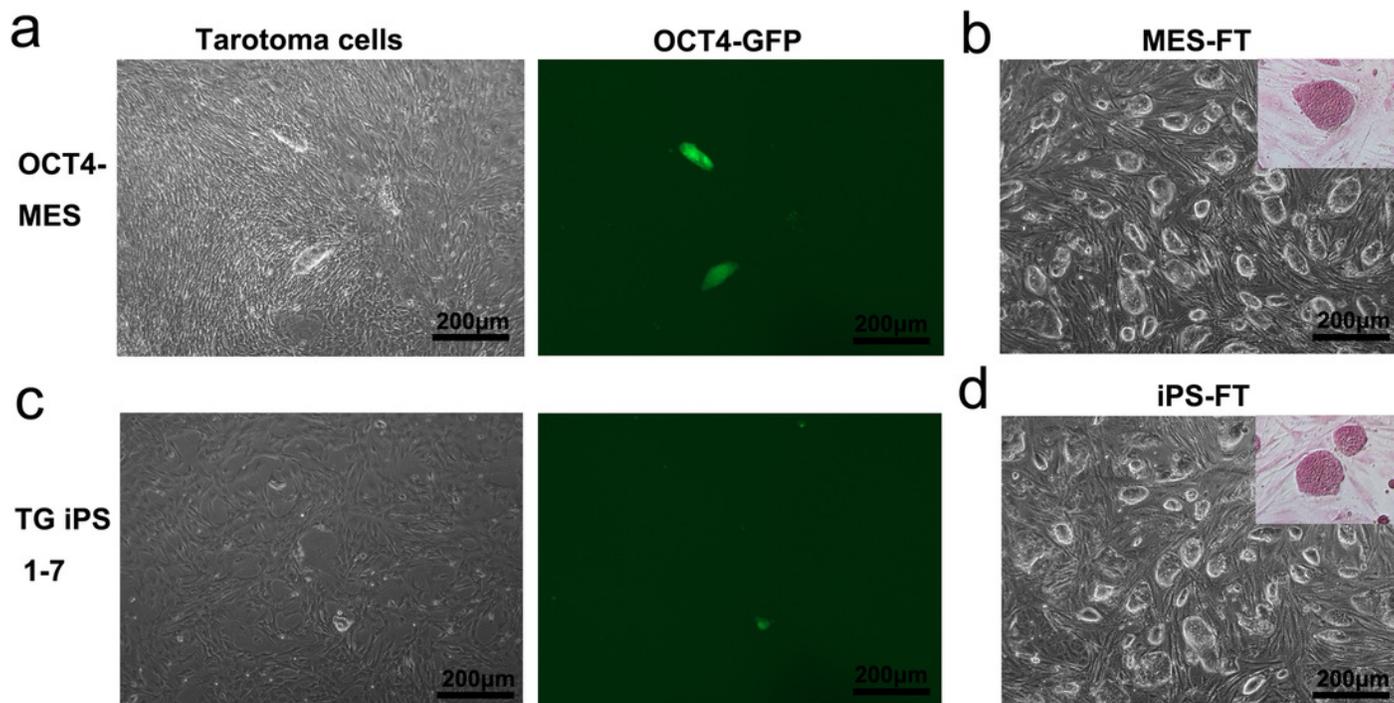


Figure 5

MES-FT and iPS-FT own self-renewal and pluripotency ability.

(a and b) Immunofluorescent staining of pluripotency markers OCT4, NANOG, SSEA1 and SOX2 in MES-FT (a) and iPS-FT (b). Both types of cells expressed all four markers.

(c and d) Hematoxylin and eosin staining of teratomas derived from MES-FT (c) and iPS-FT (d). Products of all three germ layers are seen in the image: Ectoderm: Neuronal (left of c) / epidermis with keratin (left of d). Mesoderm: smooth muscle (middle). Endoderm: gastrointestinal lining cells/glands (right). Specified cells are indicated by arrows.

(e and f) Immunohistochemistry to detect the presence of GFP-positive pluripotent cells in teratomas generated by MES-FT (e) and iPS-FT (f). GFP-positive pluripotent cells were stained in brown with anti-GFP primary antibodies.

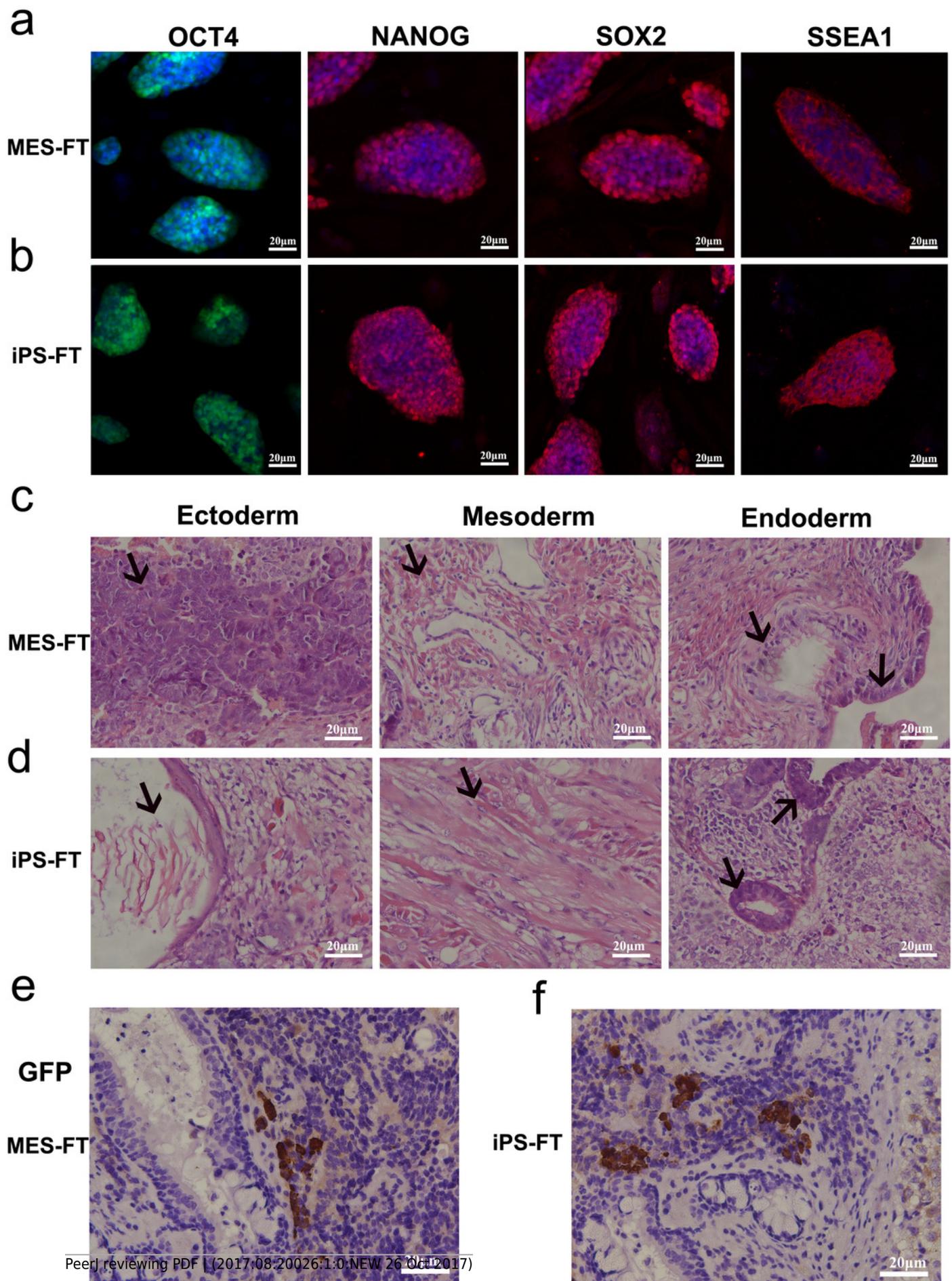


Figure 6

MES-ST and iPS-ST owned PSCs characteristic.

(a and c) A small number GFP-positive cells were found among teratoma cells generated by MES-FT (a) and iPS-FT (c) cultured in MEF medium.

(b and d) MES-ST (b) and iPS-ST (d) have typical mouse embryonic cell morphology and are AP-positive when cultured in MES medium.

(e and f) Immunofluorescent staining of pluripotency markers OCT4, NANOG, SSEA1 and SOX2 in MES-ST (e) and iPS-ST (f). Both MES-ST (e) and iPS-ST (f) expressed all four markers.

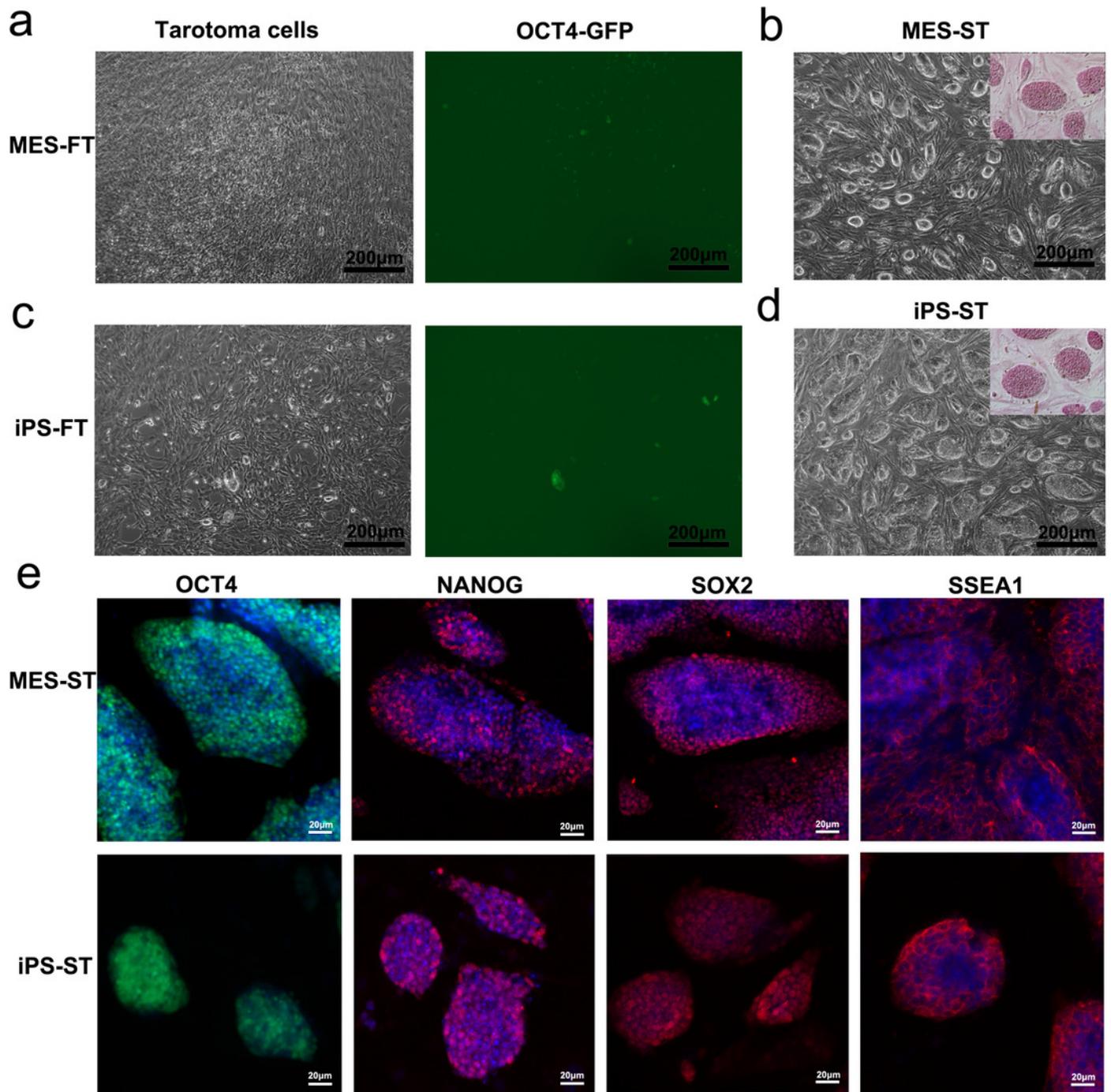


Figure 7

Oct4-MES, TG iPS 1-7, MES-FT, iPS-FT, MES-ST and iPS-ST express pluripotency genes; MES-FT, iPS-FT, MES-ST and iPS-ST more highly express several markers associated with germ cell formation.

(a) The expression levels of *Pou5f1*, *Nanog* and *Lin28a* in OCT4-MES, MES-FT and MES-ST were determined by qPCR. Both MES-FT and MES-ST highly expressed *Lin28a*, and MES-FT also highly expressed *Pou5f1*.

(b) The expression levels of *pou5f1*, *Nanog* and *Lin28a* in TG iPS 1-7, iPS-FT and iPS-ST were determined by qPCR. iPS-FT and iPS-ST highly expressed *pou5f1* and *Lin28a*.

(c) The expression levels of *Dazl*, *Stella*, *Stra8* and *Vasa* in Oct4-MES, MES-FT and MES-ST were determined by qPCR. Both MES-FT and MES-ST highly expressed *Stra8*. MES-ST also highly expressed *Dazl* and *Stella*.

(d) The expression levels of *Dazl*, *Stella*, *Stra8* and *Vasa* in TG iPS 1-7, iPS-FT and iPS-ST were determined by qPCR. iPS-FT and iPS-ST highly expressed *Dazl*, *Stra8* and *Vasa*. iPS-FT also highly expressed *Stella*. Relative expression (a and b) was quantified using the comparative threshold cycle (Ct) method ($2^{-\Delta\Delta Ct}$). n=3, Gapdh, EF1 α and β -tubulin were used as references. *p < 0.05.

