



Lab resource: Stem Cell Line

## Induced pluripotent stem cell line, ICAGi001-A, derived from human skin fibroblasts of a patient with 2p25.3 deletion and 2p25.3-p23.3 inverted duplication



Khabarova A.A.<sup>a,\*</sup>, Pristyazhnyuk I.E.<sup>a</sup>, Nikitina T.V.<sup>b</sup>, Gayner T.A.<sup>c,d</sup>, Torkhova N.B.<sup>b</sup>, Skryabin N.A.<sup>b</sup>, Kashevarova A.A.<sup>b</sup>, Babushkina N.P.<sup>b</sup>, Markova Zh.G.<sup>e</sup>, Minzhenkova M.E.<sup>e</sup>, Nazarenko L.P.<sup>b</sup>, Shilova N.V.<sup>e</sup>, Shorina A.R.<sup>f</sup>, Lebedev I.N.<sup>b</sup>, Serov O.L.<sup>a</sup>

<sup>a</sup> Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>b</sup> Research Institute of Medical Genetics, Tomsk NRMС, Tomsk, Russia

<sup>c</sup> Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia

<sup>d</sup> Group of companies "Center of new medical technologies", Novosibirsk, Russia

<sup>e</sup> Research Centre for Medical Genetics, Moscow, Russia

<sup>f</sup> Clinical City Hospital №1, Novosibirsk, Russia

### A B S T R A C T

Skin fibroblasts from a patient with developmental delay and chromosome 2p25.3 deletion syndrome were reprogrammed into induced pluripotent stem cells (iPSCs) and the clonal stem cell line ICAGi001-A (iTAF9-11) was established. ICAGi001-A pluripotency was demonstrated *in vitro* by three germ layer differentiation capacity. This line is a good model for studying of the developmental delay and brain disorder.

| Resource table                        |  | arr[GRCh38] 2p25.3(42444_2684871)x1 dn, 2p23.3-p25.3(2771354_24258056)x3 dn  |
|---------------------------------------|--|--|
| Unique stem cell line identifier      | ICAGi001-A   | Method of modification<br>N/A  |
| Alternative name(s) of stem cell line | iTAF9-11   | Name of transgene or resistance<br>N/A   |
| Institution                           | Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia<br>Research Institute of Medical Genetics, Tomsk NRMС, Tomsk, Russia | Inducible/constitutive system<br>N/A   |
| Contact information of distributor    | Anna Khabarova, <a href="mailto:anya.khabarova@gmail.com">anya.khabarova@gmail.com</a>   | Date archived/stock date<br>May 2018   |
| Type of cell line                     | iPSC   | Cell line repository/bank<br>Collective Center of ICG SB RAS "Collection of Pluripotent Human and Mammalian Cell Cultures for Biological and Biomedical Research"; Bioresource collection of the Research Institute of Medical Genetics, Tomsk NRMС, "Biobank of the population of Northern Eurasia" |
| Origin                                | Human  | Ethical approval<br>Written informed consent was obtained from the patient's parents. The study was approved by the Scientific Ethics Committee of Research Institute of Medical Genetics, Tomsk NRMС (protocol number 106/2017)   |
| Additional origin info                | Age: 3 y<br>Sex: Female<br>Ethnicity: Caucasian  |  |
| Cell source                           | Human skin fibroblasts Taf9  |  |
| Clonality                             | Clonal   |  |
| Method of reprogramming               | Lentiviral reprogramming with four transcription factors (OCT4, SOX2, KLF4, C-MYC)   | <b>Resource utility</b>  |
| Genetic modification                  | Yes  | The combination of 2p25.3 deletion and 2p25.3-p23.3 inverted duplication is a rarely described cytogenetic aberration in patients with intellectual disability. The use of hiPSC-derived neurons opens prospects in patient-specific studies to model human brain development <i>in</i>              |
| Type of modification                  | spontaneous ( <i>de novo</i> )   |  |
| Associated disease                    | Developmental delay, autosomal dominant 39; MRD39; OMIM 616521 (chromosome 2p25.3 deletion syndrome)                                 |  |
| Gene/locus                            |  |  |

\* Corresponding author.

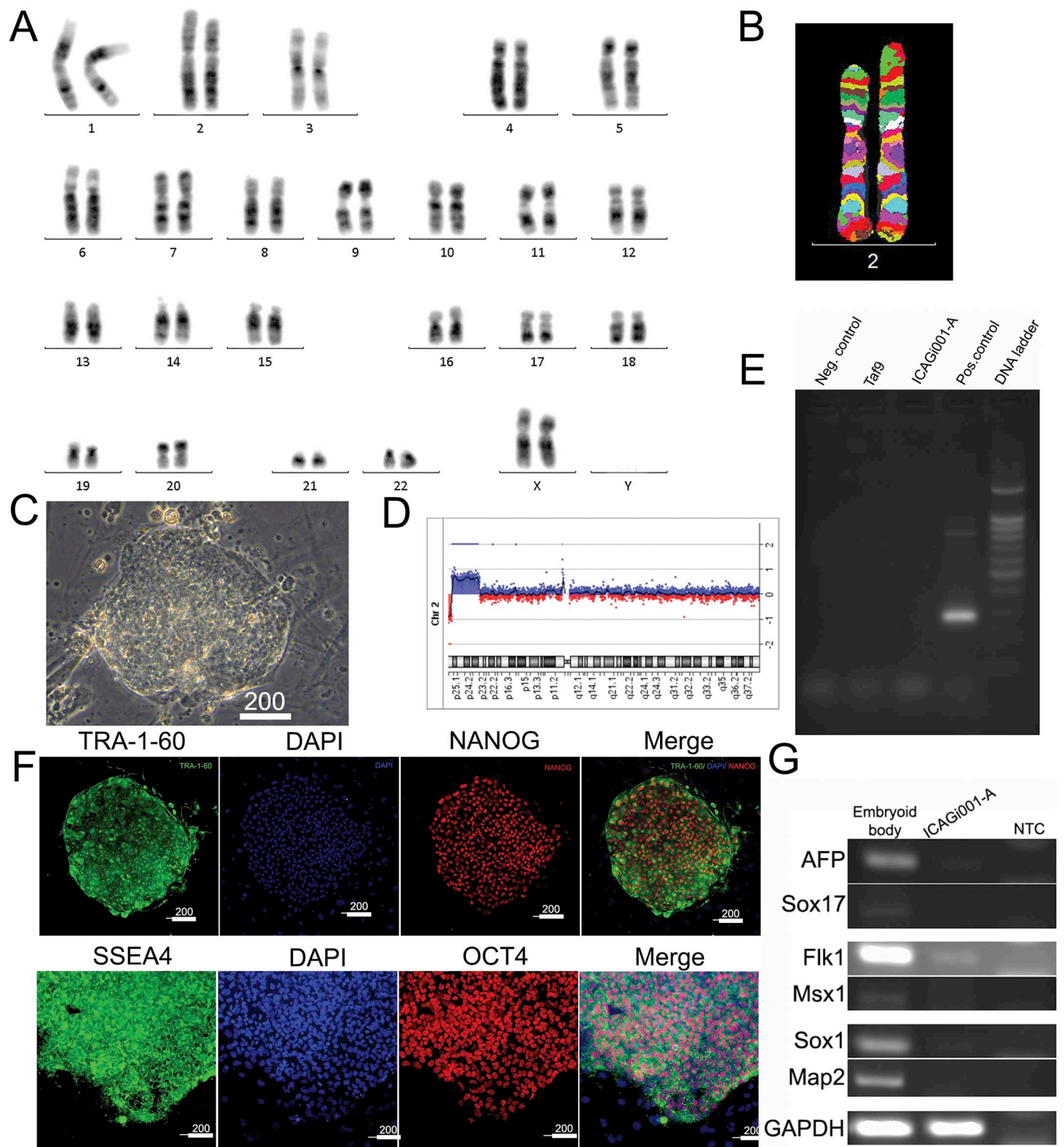
E-mail addresses: [anya.khabarova@gmail.com](mailto:anya.khabarova@gmail.com), [khabarova@bionet.nsc.ru](mailto:khabarova@bionet.nsc.ru) (A.A. Khabarova).

<https://doi.org/10.1016/j.scr.2018.101377>

Received 19 November 2018; Received in revised form 13 December 2018; Accepted 17 December 2018

Available online 18 December 2018

1873-5061/ © 2018 The Authors. Published by Elsevier B.V. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).



**Fig. 1.** Characterization of ICAGi001-A line. (A) Karyotypes. (B) The multicolor banding (MCB) analysis of chromosome 2. (C) Morphology of the iPSC colonies. (D) a CGH analysis. (E) Mycoplasma contamination test. (F) Immunofluorescence staining for the pluripotency markers NANOG, OCT4, SSEA4, TRA-1-60. (G) Expression of the endoderm (AFP, SOX17), mesoderm (MSX1, FLK1) and ectoderm (SOX1, MAP2) markers in the embryoid bodies and in ICAGi001-A.

*in vitro*. ICAGi001-A hiPSC line is a good model for studying of the developmental delay *in vitro*.

**Resource details**

Human skin fibroblasts TAF9 were derived from a 3-year-old female with developmental delay and chromosome 2p25.3 deletion syndrome.

The chromosomal rearrangement was investigated using aCGH, which detected the 2p25.3 deletion and 2p23.3-p25.3 duplication (Fig. 1D). The multicolor banding (MCB) analysis was applied to investigate the structural aberration of the der (2). The result showed an inverted duplication of distal 2p on the der (2) – dup (2) (p25.3p23.3) (Fig. 1B). Fibroblasts were reprogrammed into iPSCs through lentiviral delivery of four reprogramming factors (OCT4, SOX2, KLF4, and C-MYC)

**Table 1**  
Characterization and validation.

| Classification                      | Test   | Result   | Data   |
|-------------------------------------|--|--|--|
| Morphology                          | Photography  | Normal   | Fig. 1 panel C   |
| Phenotype                           | Qualitative analysis   | Positive for pluripotency markers: OCT4, NANOG, SSEA4 and TRA-1-60           | Fig. 1 panel F   |
|                                     | Immunocytochemistry  |  |  |
| Genotype                            | Quantitative analysis Immunocytochemistry counting (300 cells counted) | % of positive cells<br>OCT4: 98.7% TRA1-60: 97.4% SSEA-4: 96.1% NANOG 98.55% | Fig. 1 panel F   |
|                                     | Karyotype (G-banding) and resolution                                   | 46,XX,der (2) resolution 450   | Fig. 1 panel A   |
| Identity                            | STR analysis   | 20 sites tested, and 20/20 matched   | available with the authors   |
| Mutation analysis (IF APPLICABLE)   | Sequencing   | n/a  | n/a  |
|                                     | aCGH and MCB analysis  | arr[GRCh38] 2p25.3(42444_2684871)x1 dn, 2p23.3-p25.3(2771354_24258056)x3 dn  | Fig. 1 Panel B and D   |
| Microbiology and virology           | Mycoplasma   | Mycoplasma testing by PCR: Negative  | Fig. 1 panel E   |
|                                     | Differentiation potential  | Embryoid body formation  | Positive for expression of genes of the three germ layers in embryoid bodies ( <i>AFP</i> and <i>SOX17</i> for endoderm; <i>MSX1</i> and <i>FLK1</i> for mesoderm; <i>SOX1</i> and <i>MAP2</i> for ectoderm) |
| Donor screening (OPTIONAL)          | HIV 1 + 2 Hepatitis B, Hepatitis C                                     | n/a  | n/a  |
| Genotype additional info (OPTIONAL) | Blood group genotyping   | n/a  | n/a  |
|                                     | HLA tissue typing  | n/a  | n/a  |

(Gridina et al., 2018) (Table 1). ICAGi001-A line display typical morphology of human pluripotent stem cells under feeder-dependent conditions in phase contrast microscopy (Fig. 1C), and expressed OCT4 and NANOG pluripotency markers (red) in the nucleus and SSEA4 and TRA-1-60 surface markers (green), as detected by immunofluorescence staining. Nucleus stained by DAPI (blue) (Fig. 1F). Immunocytochemistry counting allowed quantifying the percentage of ICAGi001-A cells positive for OCT4, NANOG, SSEA4 and TRA-1-60 as 98.7%, 98.55%, 96.1% and 97.4%, respectively. ICAGi001-A cell line had 46,XX,der(2p) karyotype (Fig. 1A). The STR profile of the ICAGi001-A cell line fully matched with that of the parental TAF9 fibroblasts (loci analyzed: *DIS1656*, *D2S441*, *D3S1358*, *D5S818*, *D7S820*, *D8S1179*, *D10S1248*, *D12S391*, *D13S317*, *D16S539*, *D18S51*, *D21S11*, *D22S1045*, *AMEL*, *CSF1PO*, *FGA*, *SE33*, *TH01*, *TPOX*, and *vWA*). The ICAGi001-A cell line was negative for *Mycoplasma* contamination (Fig. 1E). The ability of the ICAGi001-A cell line to differentiate into cells of the three germ layers following embryoid body formation was assessed by RT-PCR for endodermal (*AFP*, *SOX17*), mesodermal (*MSX1*, *FLK1*) and ectodermal (*SOX1*, *MAP2*) genes (Fig. 1G). These results clearly demonstrate that the ICAGi001-A cells are pluripotent.

## Materials and methods

### Cell culture

TAF9 human fibroblasts were derived from a patient with 2p25.3 - p23.3 inverted duplication and 2p25.3 deletion and cultured in growth media (DMEM/F12 (Gibco) supplemented with 10% fetal bovine serum (FBS) (HyClone), 1% Pen Strep (Gibco), 1% MEM Non-essential Amino Acid solution, 1% MEM Vitamin solution, 2 mM L-glutamine (all from Sigma)) at 37 °C in 5% CO<sub>2</sub>.

### Generation of iPSCs from patient's fibroblasts

To produce iPSC from the patient's fibroblasts we used LeGO lentiviral vectors (MOI = 19) containing the human reprogramming transcription factors OCT4, SOX2, C-MYC, and KLF4 (kindly provided to us by Dr. Sergei L. Kiselev (Moscow, Russia)). The lentiviruses were produced in the Phoenix cell line using Lipofectamine 3000 (Invitrogen) according to the manufacturer's recommendations. Fibroblasts plated on the previous day were transduced with viruses containing the four reprogramming transcription factors for two days (on the second day, the C-MYC lentivirus was omitted). From day 7 to 16 the culture medium was changed daily with addition of 1 mM valproic acid (Sigma) and Selleck Human iPSC Enhancer Kit (Selleckchem,

K2010). On day 5, the transduced cells were seeded onto 10-cm culture dishes ( $2 \times 10^3$  cells per cm<sup>2</sup>) containing mitomycin C-treated CD-1 mouse embryonic fibroblast feeder cells in iPSC medium (DMEM/F12 medium supplemented with 20% KnockOut Serum Replacement, 1% GlutaMAX™-I, 1% MEM NEAA, 1% Pen Strep (all from Gibco), 0.1 mM 2-mercaptoethanol, and 10 ng/ml bFGF (Invitrogen)). On day 16, colonies (near 46 iPSC colonies per  $15 \times 10^4$  transformed fibroblasts) with iPSC morphology were picked up and expanded. All cell cultures were maintained at 37 °C in an atmosphere of 5% CO<sub>2</sub>. iPSCs maintained on feeder cells were expanded mechanically with ratio of split 1:3.

### Immunocytochemistry and immunocytochemistry counting

The iPSCs fixed with 3% formaldehyde for 20 min at room temperature. After fixation, the cells were permeabilized with 0.1% Triton X-100 in PBS for 3 min at room temperature, blocked with 3% bovine serum albumin in PBS for 25 min and incubated overnight at 4 °C with primary antibodies. Secondary antibodies were incubated for 1 h at room temperature (Table 2) and further counterstained with DAPI. All antibodies were diluted in PBS with 1.5% BSA. Immunofluorescence and immunofluorescence counting (300 cells counted) were examined with a fluorescence microscope AxioObserver Z1 (Zeiss) using ZEN software in collective Microscopic Center of ICG SB RAS, Novosibirsk and Fiji soft (ImageJ).

### In vitro differentiation and RT-PCR

Embryoid bodies were produced according to previously published protocol (Bock et al., 2011). RNA were isolated by TRI Reagent (Sigma), cDNA were obtained by RevertAid RT kit (ThermoFisher) and RT-PCR were performed with HP-Taq DNA polymerase in following conditions: 15 s–95 °C, 15 s–62 °C and 30 s–72, 34 cycles (BioRad T100 Thermal Cycler).

### Karyotyping

Preparation of metaphase chromosomes from iPSC cells was performed on passages 10–12 as previously described with minor modifications (Prokhorovich et al., 2007). Fifty eight metaphase spreads were analyzed using a Carl Zeiss Axioplan 2 imaging microscope, digital images were analyzed using ISIS 3 (*In Situ* Imaging System, MetaSystems GmbH) software at the Center for Microscopy of the Institute of Cytology and Genetics. Multicolor banding (MCB) was carried out using XCyte 2 mBAND probe (MetaSystems GmbH).

**Table 2**  
Reagents details.

| Antibodies used for immunocytochemistry/flow-citometry |                                       |  |   |
|--|---------------------------------------|--|---|
|  | Antibody                              | Dilution   | Company Cat # and RRID  |
| Pluripotency Markers                                   | Rabbit anti-NANOG                     | 1:100  | Abcam Cat# 21624, RRID: <a href="#">AB_446437</a>               |
| Pluripotency Markers                                   | Rabbit anti-OCT4                      | 1:200  | Abcam Cat# 19857, RRID: <a href="#">AB_445175</a>               |
| Pluripotency Markers                                   | Mouse anti-SSEA4                      | 1:600  | Abcam Cat# 16287, RRID: <a href="#">AB_778073</a>               |
| Pluripotency Markers                                   | Mouse anti-TRA-1-60                   | 1:600  | Abcam Cat# 16288, RRID: <a href="#">AB_778563</a>               |
| Secondary antibodies                                   | Alexa Fluor 546 Goat Anti- Rabbit IgG | 1:400  | Life technologies Cat# A-11010, RRID: <a href="#">AB_143156</a> |
| Secondary antibodies                                   | Alexa Fluor 488 Goat Anti-Mouse IgG   | 1:400  | Life technologies Cat# A-11029, RRID: <a href="#">AB_138404</a> |
| Primers  |                                       |  |   |
|  | Target                                | Forward/Reverse primer (5'-3')   |   |
| House-Keeping Genes                                    | <i>GAPDH</i>                          | GTGGACCTGACCTGCCGTCT/GGAGGAGTGGGTGTCGCTGT<br>Expected product size: 153 bp                 |   |
| Differentiation Markers                                | <i>AFP</i>                            | AAATGCGTTTCTCGTTGCTT/GCCACAGGCCAATAGTTTGT<br>Expected product size: 136 bp                 |   |
| Differentiation Markers                                | <i>SOX1</i>                           | CACAACTCGGAGATCAGCAA/GGTACTTGTAAATCCGGGTGC<br>Expected product size: 133 bp                |   |
| Differentiation Markers                                | <i>MAP2</i>                           | CAGGTGGCGGACGTGTGAAAATTGAGAGTG/CACGCTGGATCTGCCTGGGGACTGTG<br>Expected product size: 212 bp |   |
| Differentiation Markers                                | <i>SOX17</i>                          | CTCTGCCTCCTCCAGAA/CAGAATCCAGACCTGCACAA<br>Expected product size: 102 bp                    |   |
| Differentiation Markers                                | <i>MSX1</i>                           | CGAGAGGACCCCGTGGATGCAGAG/GCGGGCCATCTTCAGCTTCTCCAG<br>Expected product size: 307 bp         |   |
| Differentiation Markers                                | <i>FLK1</i>                           | TGATCGGAAATGACACTGGA/CACGACTCCATGTTGGTCAC<br>Expected product size: 131 bp                 |   |

### *Mycoplasma* contamination detection

The absence of *Mycoplasma* contamination was confirmed by PCR using primers from [Choppa et al., 1998](#).

### STR analysis

Parent TAF9 fibroblasts and their derivative iPSC line ICAGi001-A (iTAF9-11) were authenticated by STR analysis by Gordiz (<http://gordiz.ru/>).

### Acknowledgments

This study was supported by the Russian Science Foundation (grant № -16-15-10231). We are grateful to the family for participation in the study.

### References

- Bock, C., Kiskinis, E., Verstappen, G., Gu, H., Boulting, G., Smith, Z.D., Ziller, M., Croft, G.F., Amoroso, M.W., Oakley, D.H., Gnirke, A., Eggan, K., Meissner, A., 2011. Reference maps of human ES and iPSC cell variation enable high-throughput characterization of pluripotent cell lines. *Cell* 144, 439–452. <https://doi.org/10.1016/j.cell.2010.12.032>.
- Choppa, P.C., Vojdani, A., Tagle, C., Andrin, R., Magtoto, L., 1998. Multiplex PCR for the detection of *Mycoplasma fermentans*, *M. hominis* and *M. penetrans* in cell cultures and blood samples of patients with chronic fatigue syndrome. *Mol. Cell. Probes* 12, 301–308. <https://doi.org/10.1006/mcpr.1998.0186>.
- Gridina, M.M., Matveeva, N.M., Fishman, V.S., Menzorov, A.G., Kizilova, H.A., Beregovoy, N.A., Kovrigin, I.I., Pristiyazhnyuk, I.E., Oskorbin, I.P., Filipenko, M.L., Kashevarova, A.A., Skryabin, N.A., Nikitina, T.V., Sazhenova, E.A., Nazarenko, L.P., Lebedev, I.N., Serov, O.L., 2018. Allele-specific biased expression of the CNTN6 gene in iPSC cell-derived neurons from a patient with intellectual disability and 3p26.3 microduplication involving the CNTN6 gene. *Mol. Neurobiol.* 55, 6533–6546. <https://doi.org/10.1007/s12035-017-0851-5>.
- Prokhorovich, M.A., Lagarkova, M.A., Shilov, A.G., Karamysheva, T.V., Kiselyov, S.L., Rubtsov, N.B., 2007. Cultures of hESM human embryonic stem cells: chromosomal aberrations and karyotype stability. *Bull. Exp. Biol. Med.* 144 (1), 126–129. <https://doi.org/10.1007/s10517-007-0271-z>.