

### Original Reference:

Susaimanickam PJ, Maddileti S, Pulimamidi VK, Boyinpally SR, Naik RR, Naik MN, Reddy GB, Sangwan VS, **Mariappan I** (2017). Generating minicorneal organoids from human induced pluripotent stem cells. *Development*. 144(13): 2338-2351.

### Figure S1. Derivation and characterization of a human induced pluripotent stem cell line.

(A) Growing colony morphology of BJNhem20 cells (i-ii) and hiPSC-F2-3F1 cells at passage p0 on irradiated MEF feeders (iii) and at passage p20 on Matrigel coated plates (iv). (B) Immunocytochemistry of hiPSC-F2-3F1 cells at p20 on Matrigel coated chamber slides showing a homogenous expression of OCT4 (i), NANOG (ii), SSEA4 (iii) and alkaline phosphatase (iv). (C) A teratoma of about 8X8 mm, formed by hiPSC-F2-3F1 cells at p25, transplanted in the subcutaneous space of nude mice (i). H&E staining of tissue sections reveal the development of ectoderm derived RPE-like pigmented cell patches (arrow, ii), mesoderm derived adipose, cartilage and muscle tissues (arrow heads, iii) and endoderm derived gut epithelium like structures (\*; asterisk, iv). (D) G-band karyotype of hiPSC-F2-3F1 cells at p20, confirmed a normal chromosomal pattern for this female line. (E) Genomic PCR profiles of transgene specific amplicons confirmed the genomic integration of human *OCT4*, *SOX2* and *KLF4* but not the *cMYC* transgenes (i). RT-PCR profiles of transgene-specific amplicons confirmed the genomic integration and expression of all transgenes except *cMYC*, at p20 (ii). RT-PCR profiles of endogenous gene specific amplicons confirmed the expression of all four transcripts, at levels comparable to that of BJNhem20, hESCs (iii). Plasmid DNA and no-RT samples were used as PCR controls. Genomic DNA and cDNA samples were normalized using eukaryotic elongation factor (*EEF1 $\alpha$* ) as the loading control. Scale bars, 100  $\mu$ m or as specified.

