

Certificate of Analysis 2021

Invoice number: SCTC2021-00017

Name principal investigator: Sarah Weckhuysen

Cell line number: IPS17-00063

Project name: KCNQ2

Table 1: Information on the reprogrammed cell line

Information cell line:	
Product description	Trilineage differentiation
Parental cell line Parental cell type	HEP17-00110 PBMCs
Diagnosis Mutation	Benign neonatal familial epilepsy N/A*
Number of clones Passage (P) of iPSCs reported at delivery	1 P14

*N/A: Not Applicable

Table 2: Information on the characterization of the reprogrammed cell line

Test description:	Test method:	Test specification:	Result:
Activation of markers of undifferentiation	qPCR	Upregulation of <i>SOX2</i> , <i>LIN28</i> , <i>NANOG</i> , <i>OCT4</i> in iPSCs compared with PBMCs	N/A*
Expression of markers of undifferentiation	Immunocytochemistry	Expression of OCT4, NANOG, SSEA4, TRA-1-81	N/A*
Mycoplasma test	PCR	Negative	N/A*
Three lineage differentiation	Differentiation assay	Upregulation of germlayer-specific markers	Pass

Three germ layer differentiation

Clone 1 was differentiated into the endodermal, mesodermal and ectodermal germ layers. The RNA was isolated and the gene expression was checked by qPCR. Ct values are normalized with the housekeeping gene GUSB, set at 1. For each lineage two genes were assessed (Table 3). The differentiated cells were also stained for lineage-specific markers (Table 4).

Table 3: qPCR markers for three lineage differentiation

Lineage	Marker
Endoderm	FOXA2, SOX17
Mesoderm	Brachyury, HAND1
Ectoderm	PAX6, NCAM1

Table 4: ICC markers for three lineage differentiation

Lineage	Marker
Endoderm	SOX17
Mesoderm	NCAM1
Ectoderm	NESTIN

Endoderm

Upregulation of endodermal markers

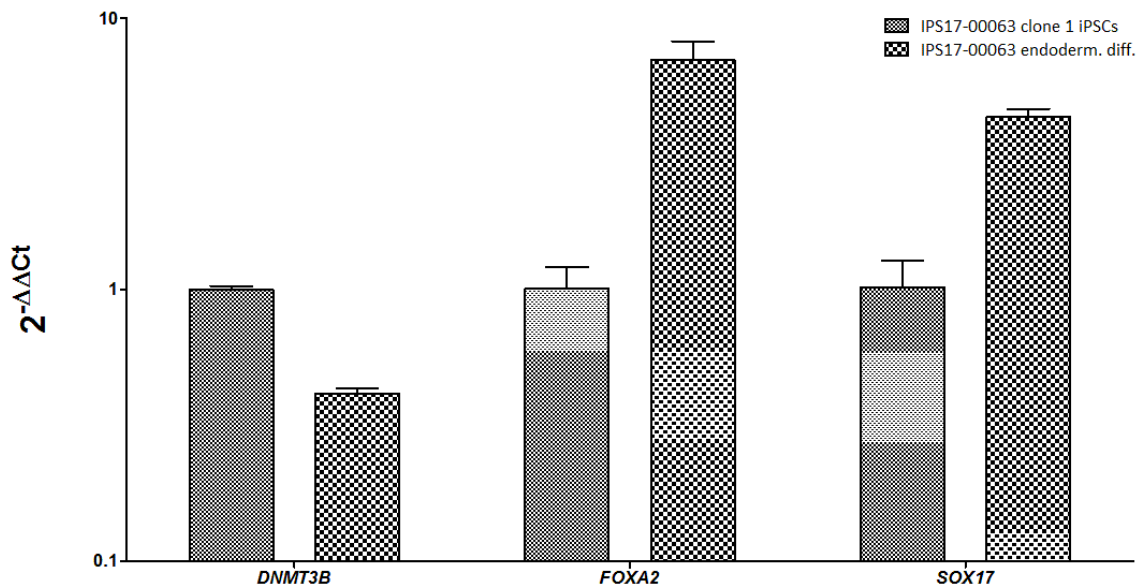


Figure 1: Expression fold difference of endoderm-specific genes in differentiated cells, compared with undifferentiated iPSCs. *DNMT3B* was used as a reference for undifferentiation.

Mesoderm

Upregulation of mesodermal markers

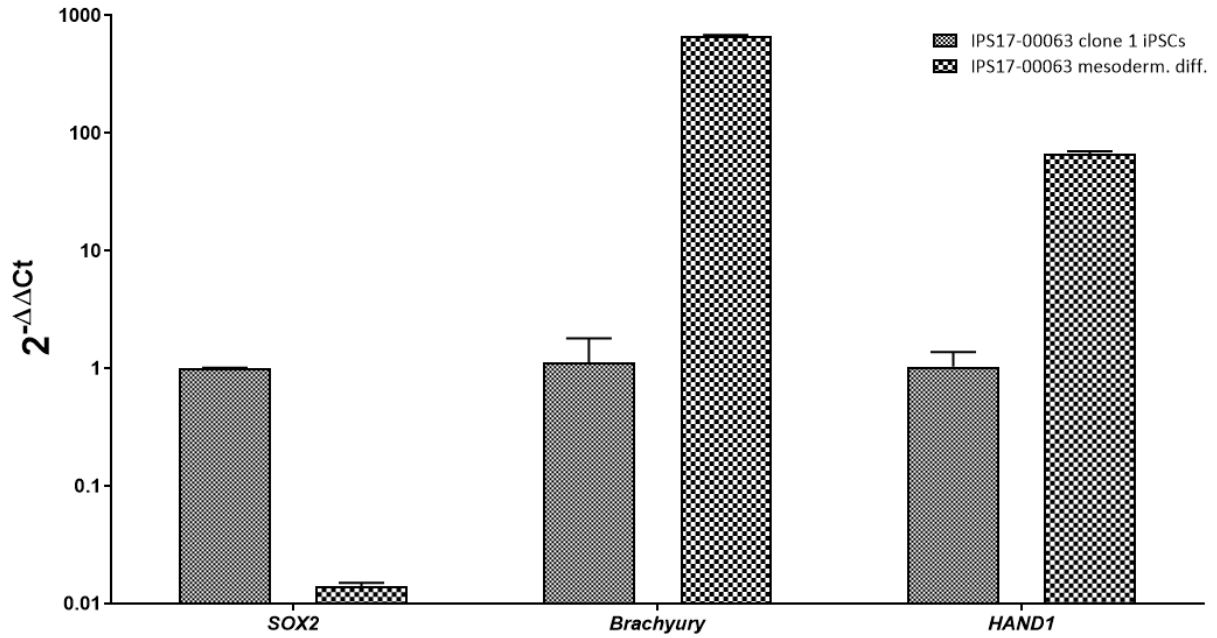


Figure 2: Expression fold difference of mesoderm-specific genes in differentiated cells, compared with undifferentiated iPSCs. *SOX2* was used as a reference for undifferentiation.

Ectoderm

Upregulation of ectodermal markers

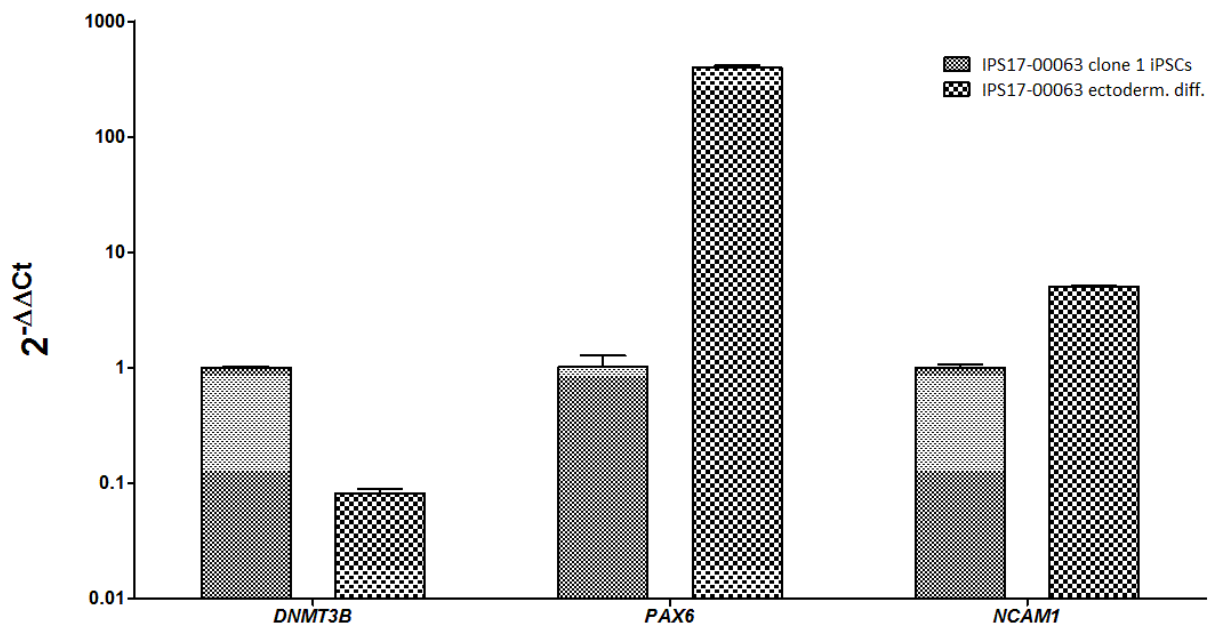


Figure 3: Expression fold difference of ectoderm-specific genes in differentiated cells, compared with undifferentiated iPSCs. *DNMT3B* was used as a reference for undifferentiation.

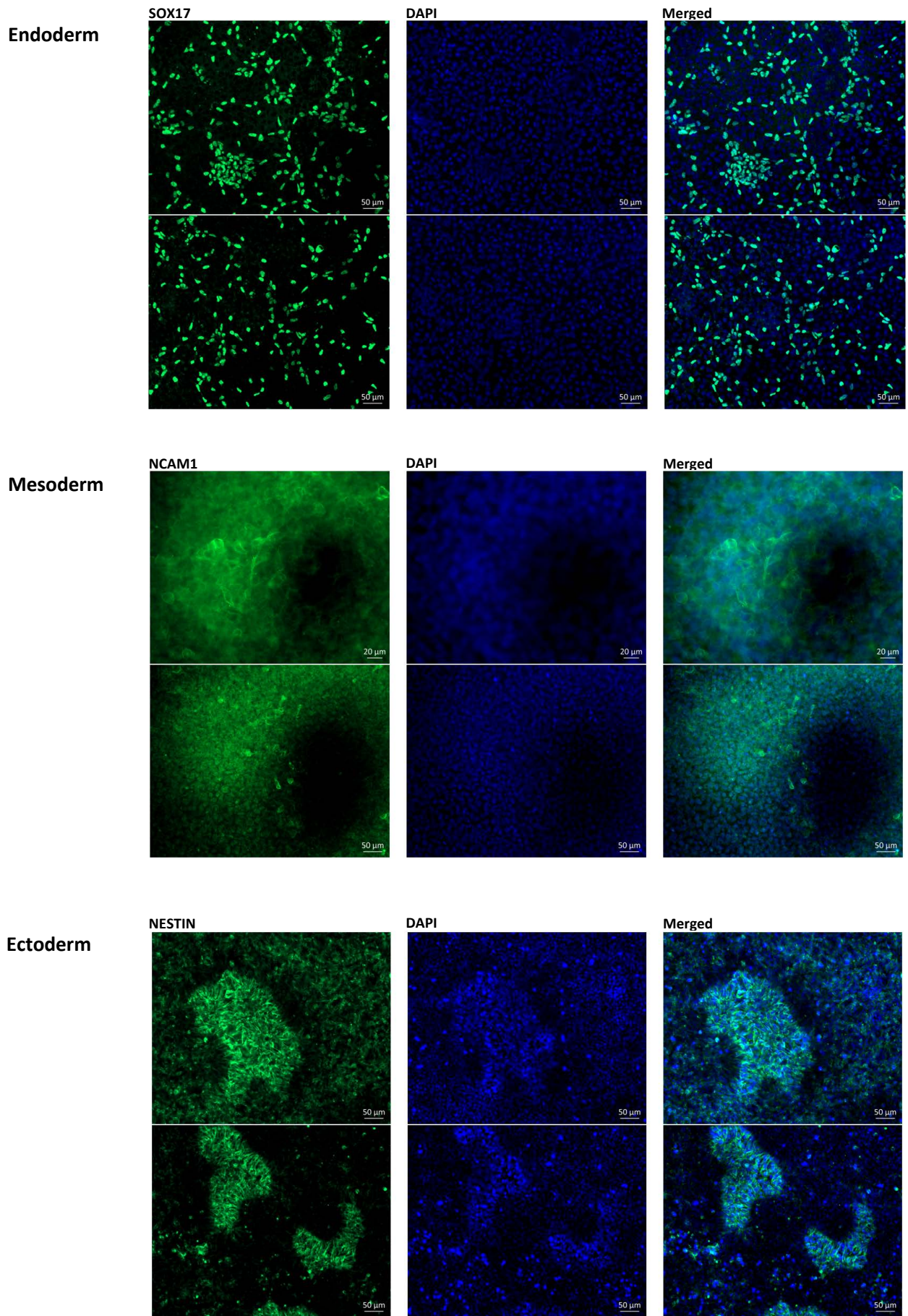


Figure 4: Immunofluorescence staining of differentiated cells showing a positive signal of germlayer-specific markers.

Pass

Fail

Other:

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Date