

Certificate of Analysis 2021

Invoice number: SCTC2021-00004

Name principal investigator: Anneke den Hollander Cell line number: IPS19-00051 CRISPR clone H6 Project name: VICI

Table 1: Information on the reprogrammed cell line

| Information cell line: | |
|---|----------------------------|
| Product description | Trilineage differentiation |
| Parental cell line | IPS19-00051 |
| Parental cell type | N/A* |
| Diagnosis | N/A* |
| Mutation | N/A* |
| Number of clones | 1 |
| Passage (P) of iPSCs reported at delivery | P25 |

*N/A: Not Applicable

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

| Test description: | Test method: | Test specification: | Result: |
|----------------------------------|-----------------------|--|-----------------------------|
| Activation of stem cell markers | qPCR | Upregulation of <i>SOX2, LIN28, NANOG, OCT4</i> in iPSCs compared with PBMCs | N/A* |
| Expression of stem cell markers | Immunocytochemistry | Expression of OCT4, NANOG, SSEA4, TRA-1-81 | N/A* |
| Mycoplasma test | PCR | Negative | N/A* |
| Three lineage differentiation | Differentiation assay | Upregulaton of germlayer-specific markers | No upregulation of HAND1 |

Three germ layer differentiation

IPS19-00051 CRISPR clone H6 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).

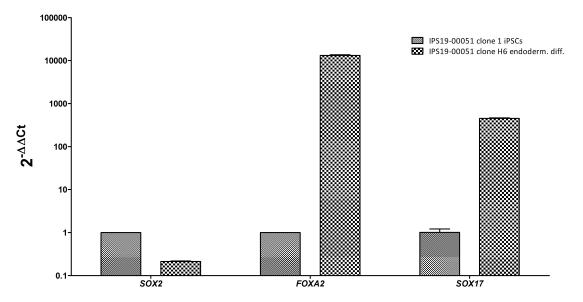
Table 3: qPCR markers for three lineage differentiation

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

Table 4: ICC markers for three lineage differentiation

| Lineage | Marker |
|----------|--------|
| Endoderm | SOX17 |
| Mesoderm | DESMIN |
| Ectoderm | NESTIN |

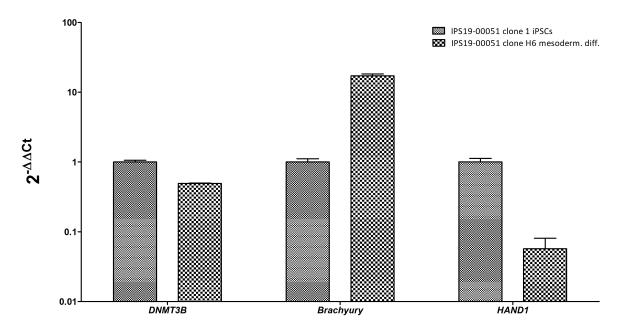
Endoderm



Upregulation of endodermal markers

Figure 1: Expression fold difference of endoderm-specific genes in differentiated cells, compared with undifferentiated iPSCs. *SOX2* was used as a reference for pluripotency.

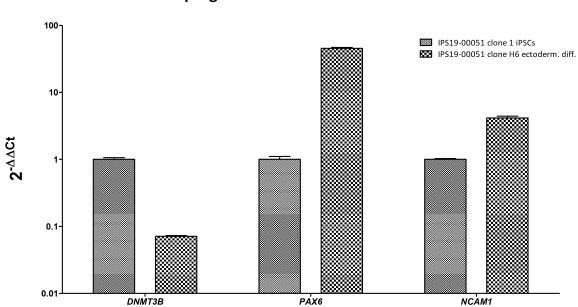
Mesoderm



Upregulation of mesodermal markers

Figure 2: Expression fold difference of mesoderm-specific genes in differentiated cells, compared with undifferentiated iPSCs. *DNMT3B* was used as a reference for pluripotency.

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 pluripotency.

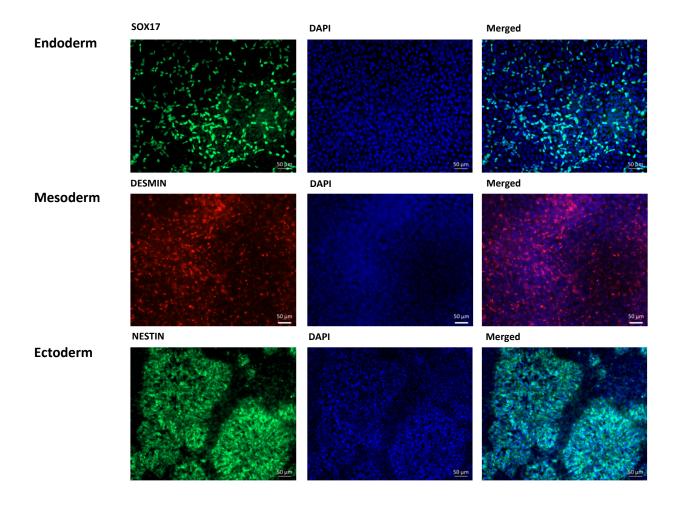


Figure 4: Immunofluorescence staining of differentiated cells showing a positive signal of germlayer-specific markers.

Pass Fail Other:

> Silvia Albert, PhD Manager, Radboud Stem Cell Technology Center Date