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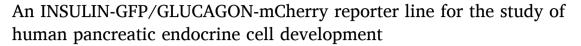
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Lab Resource: Single Cell Line



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Insulin expressing beta cells and glucagon expressing alpha cells are the two most abundant endocrine cell types of the human pancreatic islet. Both alpha and beta cells can be generated in vitro via the differentiation of pluripotent stem cells (PSCs), affording the opportunity to study their ontogeny and to examine their developmental inter-relationship. To aid these analyses, we have generated a PSC line in which insulin expression is reported by GFP and glucagon expression is reported by mCherry. This cell line enables viable isolation of cells expressing each hormone and optimisation of methods that lead to their generation.

### Resource Table

Cell Source

Unique stem cell line identifier ESIBIe003-A-10

https://hpscreg.eu/cell-line/ESIBIe003-A-10

Alternative name(s) of stem MCRIe004-A-10 cell line HES3 INS-GFP GCG-mCh

Institution Murdoch Children's Research Institute,

Melbourne, Australia

hES3 (ESIBIe003-A)

Contact information of Jacqueline Schiesser; jacqui.schiesser@mcri.edu.

distributor au
Type of cell line ESC
Origin Human
Additional origin info Age: n/a

Sex: Female Ethnicity if known: n/a

 $\begin{array}{ll} \mbox{Clonality} & \mbox{Clonal} \\ \mbox{Method of reprogramming} & \mbox{n/a} \end{array}$ 

Method of reprogramming n/a Genetic Modification YES

Type of Modification Reporter knock-in

Associated disease n/

Gene/locus Glucagon, 2q24.2

Method of modification TALEN

Name of transgene or mCherry

resistance

Inducible/constitutive system n/2

Inducible/constitutive system n/a
Date archived/stock date 25/6/2019
Cell line repository/bank N/A

Ethical approval RCH Human Research Ethics Committee 33001A

#### 1. Resource utility

This reporter line expresses GFP and mCherry fluorescent proteins under the control of the endogenous INSULIN (INS) and GLUCAGON (GCG) loci respectively. This facilitates the identification of endocrine cells in real time and enables isolation of the two most common hormone expressing cell types in the pancreas by flow cytometry (Table 1).

# 2. Resource details

We previously generated INS<sup>GFP/w</sup> embryonic stem cell (ESC) reporter lines (Micallef et al., 2012) that enable the isolation of putative pancreatic beta cells from differentiating ESC cultures. In order to further characterise endocrine cells generated during the differentiation process, we inserted sequences encoding a mCherry reporter gene into the GLUCAGON (GCG) locus of INS<sup>GFP/w</sup> hESCs, using gene targeting by homologous recombination. The targeting vector comprised two homology arms flanking the mCherry gene and an adjacent cassette conferring resistance to the antibiotic, hygromycin (Fig. 1A). Following identification of correctly targeted clones, the PGKHygroR was subsequently removed by CRE-mediated recombination, a modification confirmed by PCR analysis of sequences spanning the junctions between the targeting vector and flanking genomic sequences (Fig. 1B). The resultant cell line, here designated as INS<sup>GFP/w</sup>GCG<sup>mCh/w</sup> hESCs, appeared morphologically normal and expressed the pluripotency

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<sup>&</sup>lt;sup>1</sup> Edouard G. Stanley and Jacqueline V. Schiesser contributed equally to this study.

Table 1 Characterization and validation.

Classification	Test	Result	Data
Morphology Phenotype	Photography Qualitative analysis (Immunofluorescence)	Normal Positive staining for the pluripotency marker OCT4	Fig. 1C Fig. 1C
	Quantitative analysis (Flow Cytometry)	Positive staining for EPCAM, CD9 and SSEA4	Fig. 1C
Genotype	Karyotype (G-banding) and resolution	SNP array (resolution 0.50 Mb)	Submitted in archive with journal
Identity	Microsatellite PCR (mPCR) OR	Not performed	N/A
	STR analysis	SNPduo compartitive analysis of SNP arrays of parental and gene edited clone. Identical genotypes (>99.9%) of entire genome indicating the lines are from the same individual	Submitted in archive with journal
Mutation analysis (IF APPLICABLE)	PCR	Correct Junction fragments generated by integration dependent primer pairs	Fig. 1B
Microbiology and virology	Mycoplasma	Negative for Mycoplasma as determined by PCR	Submitted in archive with journal
Differentiation potential	Directed differentiation to derivatives of ectoderm, mesoderm and endoderm	Differentiation to ectoderm marked by Beta-III Tubulin, mesoderm marked by CD34 and CD43, and endoderm marked by INSULIN and GLUCAGON	Fig. 1D (ectoderm), E (mesoderm) and F (endoderm)
Donor screening (OPTIONAL)	HIV $1+2$ Hepatitis B, Hepatitis C	N/A	N/A
Genotype additional	Blood group genotyping	N/A	N/A
info (OPTIONAL)	HLA tissue typing	N/A	N/A

marker OCT4 by immunofluorescence, as well as the stem cell markers CD9, EPCAM and SSEA-4 by flow cytometry (Fig. 1C). DNA fingerprinting by STR analysis confirmed the relationship between the GCG-mCherry-targeted line to the parental INS-GFP line (submitted in archive with journal). The INS<sup>GFP/w</sup>GCG<sup>mCh/w</sup> hESC line had a normal molecular karyotype as determined by SNP analysis and tested negative for *Mycoplasma*.

We next tested whether  $INS^{GFP/w}GCG^{mCh/w}$  hESCs retained the potential to differentiate into derivatives of the three primary germ layers, ectoderm, mesoderm and endoderm.  $INS^{GFP/w}GCG^{mCh/w}$  hESCs differentiated toward the neural lineage, a derivative of ectoderm, formed cells that expressed the neural marker beta III-tubulin (Fig. 1D). To examine their capacity to form mesoderm lineages,  $INS^{GFP/w}GCG^{mCh/w}$  hESCs were subjected to a differentiation protocol that generates endothelial and haematopoietic cells. Under these conditions, early mesoderm cells were identified by the expression of CD13 at differentiation day 4 whilst day 8 cultures contained CD34 + endothelial cells and a small fraction of CD43 + blood cells (Fig. 1E).

The ability of INS<sup>GFP/w</sup>GCG<sup>mCh/w</sup> hESCs to form endoderm was

examined by differentiating them toward pancreatic endocrine lineages, marked by the expression of INSULIN-GFP and GLUCAGON-mCherry. Consistent with previously observations using the parental INSGFP/w hESC line (Micallef et al., 2012), INSGFP/WGCGmCh/w hESCs generated GFP+ (INS+) cells by differentiation day 28. Confocal microscopy and flow cytometry analysis revealed the presence of cells expressing either hormone alone or both INS and GCG (Fig. 1F, G). By differentiation day 38, most fluorescent cells expressed either GCG or INS but not both. At this stage, intracellular flow cytometry analysis was used to show that GCG expression was exclusively associated with mCherry + cells, confirming the fidelity of the reporter gene (Fig. 1H). When combined with the results of previous studies, these data indicate that  $INS^{GFP/w}GCG^{mCh/w}$ w hESCs enable the real time observation of cells expressing INS and GCG. As such, the INS<sup>GFP/w</sup>GCG<sup>mCh/w</sup> hESCs dual reporter line will facilitate research aimed at optimising endocrine differentiation protocols and for studying the developmental relationship between beta cells and alpha cells.

### 3. Materials and methods

# 3.1. Generation of targeted GCG-mCherry/w hESCs

The GLUCAGON-mCherry targeting construct comprised a 2 kb 5′ homology arm, mCherry coding sequences, a loxP flanked phosphoglycerol kinase (PGK)-promoter-hygromycin resistance cassette and a 2.9 kb 3′ homology arm. INSULIN<sup>GFP/w</sup> HES3 ESCs (Micallef et al., 2012) were electroporated with GLUCAGON specific TALENs and the targeting construct as described previously (Kao et al., 2016). Hygromycin resistant clones were screened for the presence of gene targeting events using primer P1, corresponding to genomic sequences 5′ of the 5′ end of the targeting vector, in conjunction with P2, a reverse primer located in the mCherry coding sequence. Correctly targeted clones were expanded and the PGK-hygromycin resistance cassette removed as previously described (Micallef et al., 2012), verified by PCR using primers P3 and P4. Single-cell cloning was performed using single-cell deposition as previously described (Micallef et al., 2012). A single clone, designated INS<sup>GFP/w</sup>GCG<sup>mCh/w</sup>, was used for subsequent analyses.

## 3.2. hESC culture and in vitro differentiation

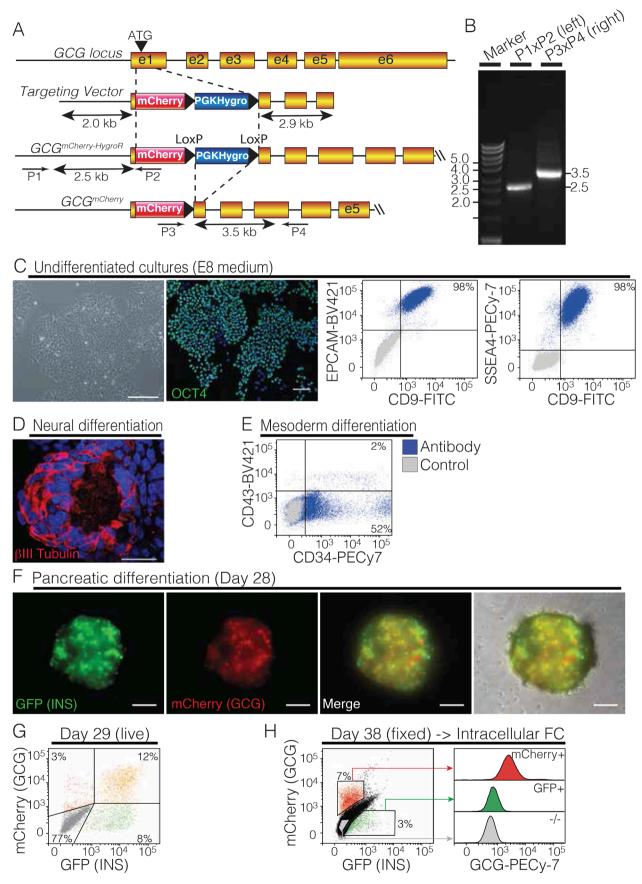
Cells were cultured and passaged enzymatically essentially as described previously (Costa et al., 2007). To assess ectodermal differentiation potential, hESCs were differentiated as described (Goulburn et al., 2011) and neural progenitors identified using immunofluorescence analysis. For mesoderm differentiation, hESCs were differentiated to CD34 + cells according to Ng et al. (2008) and analysed using flow cytometry. Endodermal differentiations were performed as described by Micallef et al. (2012).

#### 3.3. Extracellular and intracellular staining by flow cytometry

Cells were dissociated using TrypLE<sup>TM</sup> select (ThermoFisher) and analysed using a LSR Fortessa (Becton Dickinson, Bioscience). Expression of extracellular markers was evaluated in cells that were incubated with conjugated antibodies (Table 2) diluted in PBS containing 2% FCS for 20 min on ice. Dead cells were identified by propidium iodide (Sigma) staining. Expression of intracellular markers was evaluated in cells following fixation and permeablisation in Cytofix/Cytoperm (Becton Dickinson, Bioscience) as specified by the manufacturer.

## 3.4. Live cell sorting

Single cell cloning was performed by dissociating cultures using  $TrypLE^{TM}$  select (ThermoFisher) and single cells deposited into individuals wells of a 96 well plate using a BD Aria (Becton Dickinson, Bioscience).



 $\textbf{Fig. 1.} \ \ \text{Characterization and validation of an INSULIN}^{\text{GFP/w}} \ \ \text{GLUCAGON}^{\text{mCh/w}} \ \ \text{human embryonic stem cell line.}$ 

Table 2 Reagents details.

Antibodies used for	r immunocytochemistry/fl	ow-cytometr	y	
	Antibody	Dilution	Company Cat # and RRID	
Pluripotency Marker	Rabbit anti-OCT4	1:200	Cell Signaling Technology Cat# 2840 RRID:AB 2167691	
Pluripotency	FITC Mouse anti-	1:10	BD Biosciences	
Marker	human CD9		Cat#555371, RRID: AB_395773	
Pluripotency Marker	Brilliant Violet 421 Mouse anti-human CD326 (EpCAM)	1:50	Biolegend Cat#324220 RRID:AB_2563847	
Pluripotency Marker	PE/Cyanine 7 Mouse anti-human SSEA-4	1:50	Biolegend Cat#330420 RRID:AB 2629631	
Differentiation Marker	PE/Cyanine 7 Mouse anti-human CD34	1:100	Biolegend Cat#343516 RRID:AB 187725	
Differentiation Marker	Brilliant Violet Mouse anti-human CD43	1:25	BD Biosciences Cat#562916, RRID: AB 2737890	
Differentiation Marker	Mouse anti-Beta III Tubulin	1:200	Promega Cat#G712A, RRID:AB 430874	
Differentiation Marker	Mouse anti-human Glucagon	1:100	Sigma-Aldrich Cat#G2654, RRID: AB 259852	
Secondary Antibody	Alexa Fluor 594 Donkey anti-mouse IgG (H + L)	1:1000	Invitrogen Cat#A- 21203, RRID: AB_141633	
Secondary Antibody	Alexa Fluor 647 Donkey anti-mouse IgG (H + L)	1:1000	Invitrogen Cat#A- 21244, RRID: AB_2535812	
Secondary Antibody	PE/Cyanine 7 Goat anti-mouse IgG	1:100	Biolegend Cat#405315 RRID:AB_10662421	
Primers				
	Target	Forward/	Forward/Reverse primer (5'-3')	
Screening primer P1	GLUCAGON	ggttttctttggatgcatctgagt		
Screening primer P2	mCherry	tcgccctcgatctcgaactcgtg		
Screening primer P3	mCherry	cagtctcaatggatacaggctatc		
Screening primer P4	GLUCAGON	catcgtggaacagtacgaacgcg		

# 3.5. Immunofluorescence staining

Cells were fixed in 4% paraformaldehyde for 10 min at room temperature. Following this, cells were placed in blocking buffer (10% FCS  $\pm$  0.1% Triton-X in 1x PBS) for 1 h at room temperature. Primary antibodies were diluted in blocking buffer and incubated overnight at 4 °C. Secondary antibodies were diluted in blocking buffer and incubated at room temperature for 1 h (antibodies listed in Table 2). Nuclei were stained with DAPI at 1  $\mu g/ml$  (Sigma-Aldrich) and cells visualised using a LSM 780 confocal microscope running Zen Black software (Carl-Zeiss). Image analysis was performed using ImageJ software.

#### 3.6. SNP analysis

Karyotypes were analysed using Infinium CoreExome-24 SNP arrays. Data was compared to the human reference sequence hg19/CRCh37 (Feb 2009). The parental hES3 INSULIN-GFP/w was analysed using the SNP array and this data compared to the corresponding HES3 INSULIN-GFP/w GLUCAGON-mCh/w line using SNPduo comparative analysis (http://pevsnerlab.kennedykrieger.org/SNPduo). No differences were detected between the parental single-knock in and the derivative double knock-in line.

#### 3.7. Mycoplasma screening

hESCs were tested for Mycoplasma by PCR by the commercial service provider, Cerberus Sciences (Adelaide, Australia).

## **Declaration of Competing Interest**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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