

Functional Assessment of the Induced Pluripotent Stem Cells (iPSC)-derived β -like Cells

Manon Jaffredo, Nicole Krentz, Benoite Champon, Fernando Abaitua, Antoine Pirog, Nicola Beer, Anne Clark, Yannick Bornat, Sylvie Renaud, Matthieu Raoux, et al.

▶ To cite this version:

Manon Jaffredo, Nicole Krentz, Benoite Champon, Fernando Abaitua, Antoine Pirog, et al.. Functional Assessment of the Induced Pluripotent Stem Cells (iPSC)-derived β -like Cells. EASD Islet Study Group, Apr 2019, Oxford, United Kingdom. hal-02518480

HAL Id: hal-02518480 https://hal.science/hal-02518480v1

Submitted on 25 Mar 2020

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

Functional Assessment of the Induced Pluripotent Stem Cells (iPSC)-derived β-like Cells

Manon Jaffredo¹, Nicole Krentz², Benoite Champon^{2,3}, Fernando Abaitua², Antoine Pirog⁴, Nicola Beer², Anne Clark³, Yannick Bornat⁴, Sylvie Renaud⁴, Matthieu Raoux¹, Jochen Lang¹, Anna Louise Gloyn^{2,3,5}, Benoit Hastoy^{3,5}

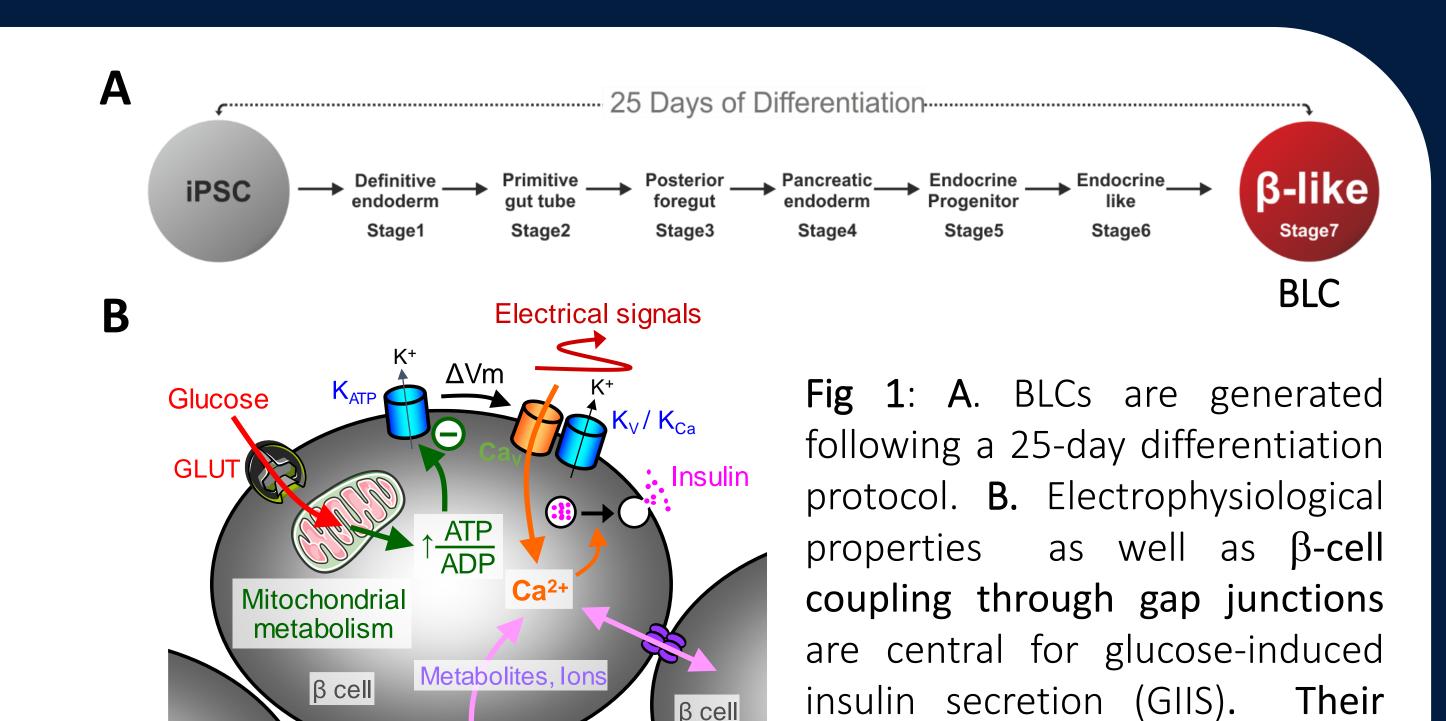
¹CBMN, CNRS UMR 5248, University of Bordeaux, France ²Wellcome Centre for Human Genetics, University of Oxford, UK ⁴IMS, CNRS UMR 5218, Bordeaux INP, University of Bordeaux, France ⁵Oxford NIHR Biomedical Research Centre, Churchill Hospital, Oxford, UK

Background and Aim

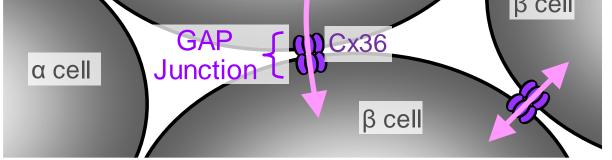
Human β-like cells (BLCs) derived from induced Pluripotent Stem are an emerging tool to study human β -cell Cells (iPSC) development and function.

established a **functional assessment** of each of our We differentiation on the following properties:

- **Secretory response** (insulin secretion assay)
- **Exocytosis kinetics** (capacitance measurement)

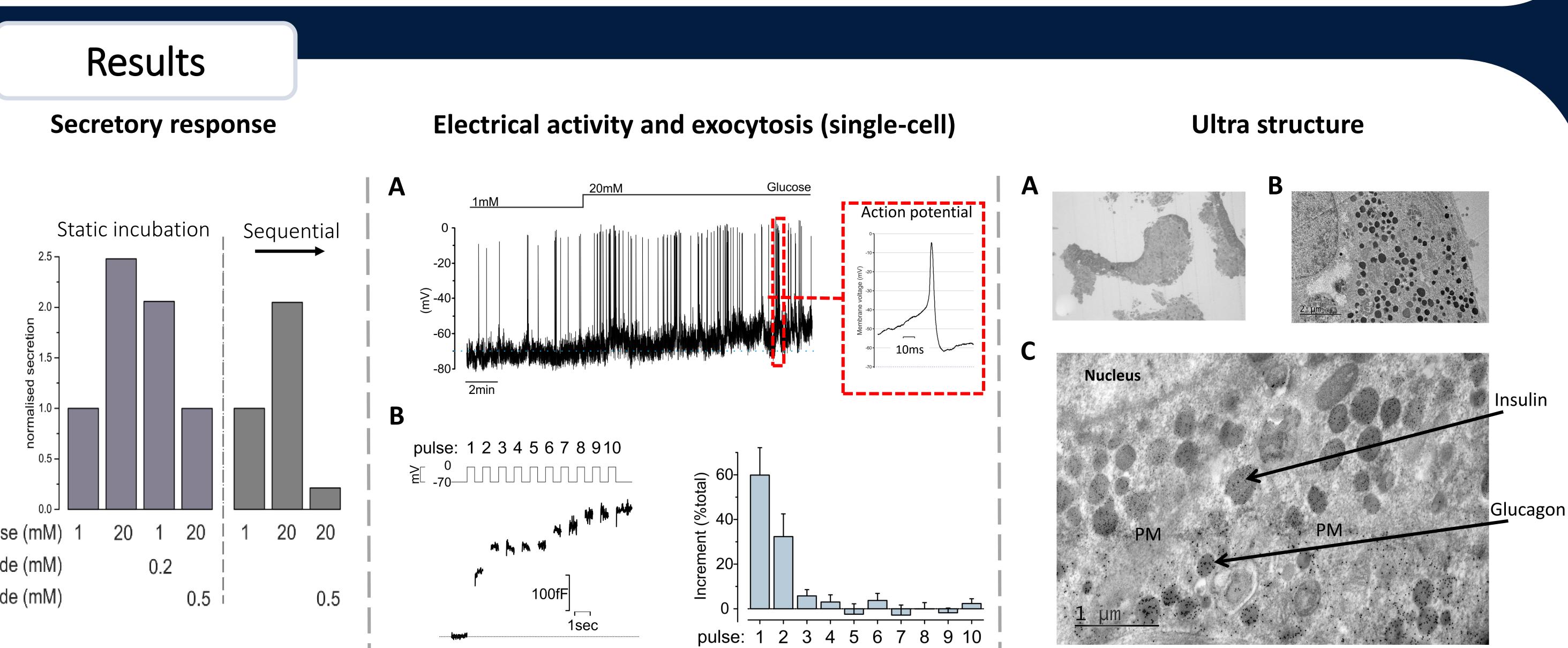


- **Ultrastructure** (Electron microscopy)
- **β-cells electrical coupling** (Multi-Electrode Array, MEA)



Cx36 = Connexin 36

assessment in BLC provides a functional quality control of the differentiation.



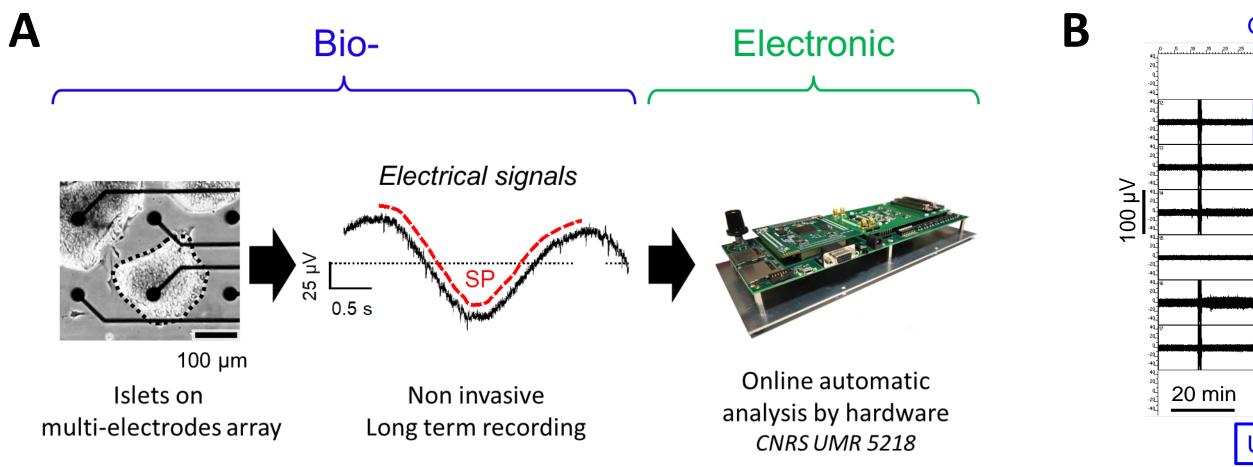
Glucose (mM)	1 2	20	1	20	1	20	20
Tolbutamide (mM)			0.2	ļ			
Diazoxide (mM)				0.5			0.5

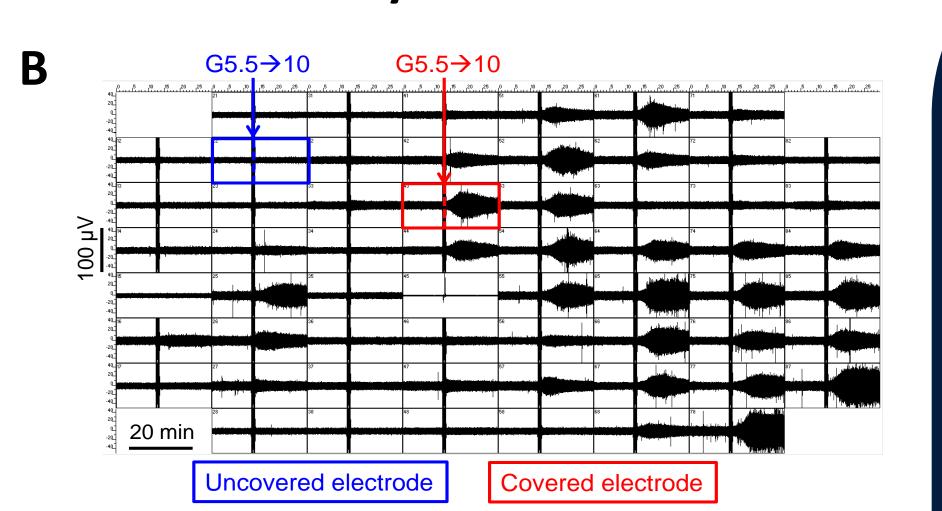
Fig 2: Representative GIIS induced by 1 and 20mM glucose and modulators of K_{ATP} channels activity (Tolbutamide/Diazoxide). The stimulations are either parallel static incubations (left), or sequential on the same BLC preparation (right).

Fig 3: A. Glucose stimulation triggers action potential firing (n=3 cells). Action potential kinetics is similar to human primary β -cells (10ms duration). **B.** Exocytosis triggered by a train of 10 depolarisations to 0mV is biphasic as in human primary β -cells (Raw trace left, quantification right, n=7 cells).

Fig 4: A. Semi thin section showing cell heterogeneity. B. Ultra structure of BLC. Scale bar: 2µm C. Immunogold labelling of insulin (BLC, 10nm gold particles, top) and glucagon (15nm gold particle, alpha-like cell, bottom). PM: Plasma Membrane. Scale bar: 1µm.

Multicellular activity assessed by MEA and its automatic analysis in hard real-time





β-like cells derived from **iPSC** exhibit critical human β -cell features:

Conclusions

- Glucose-induced insulin secretion
- Electrical excitability and exocytosis
- Electrical coupling

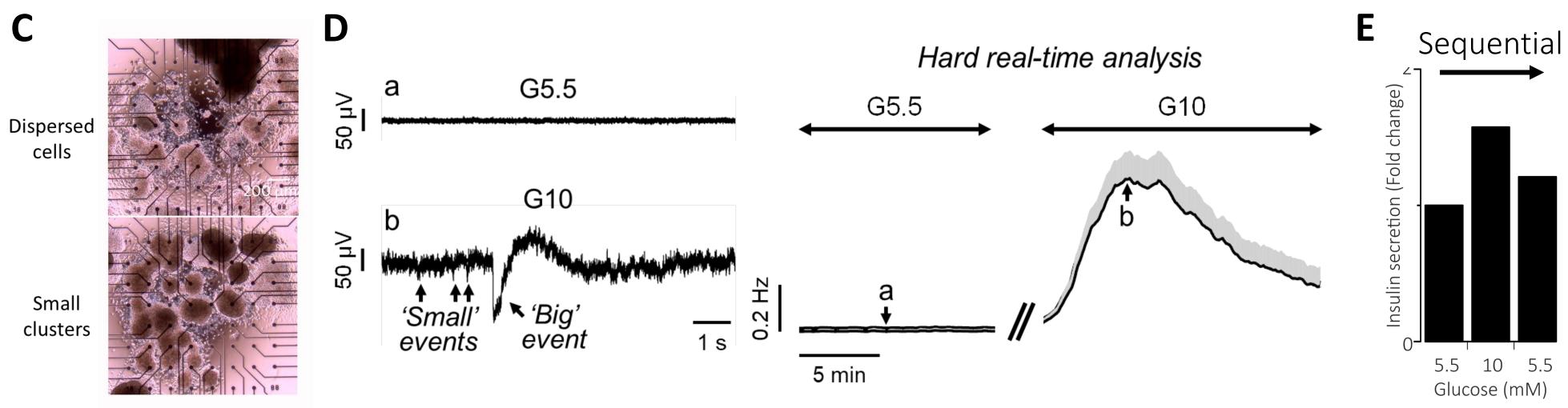


Fig 5: A. Principle of the approach: islets on MEA generate slow potentials (SP) analysed by a microelectronic circuit. SPs are β cell-specific coupling signals. **B.** 60 recording channels showing β -like cells activity. C. Images of preparations. D. Representative traces at 5.5 and 10mM glucose and realtime analysis of SP frequencies (means+SEM, n= 29 electrodes). E. Corresponding insulin secretions.

BLCs allow functional studies on human β -cell differentiation. The emergence of differentiation protocols should new their homogeneity and improve ultrastructure. This pipeline will the support characterization of T2D-risk alleles on human β -cell function.









