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Advancing Male Reproductive Toxicology with Innovative 3D hiPSCs-based Models of Testicular Steroidogenesis

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BACKGROUND



Approximately **1 out of 6 couple** experience **INFERTILITY** in their lifetime ¹ World Health Organization

 \checkmark In male, androgens (e.g. testosterone) play a

Endocrine disruptors (e.g. drugs, pollutants)
can disrupt testicular steroidogenesis
(converting cholesterol into androgens) →
severe consequences for human health



AIM OF THE STUDY

To develop an in vitro model for testicular steroidogenesis for both toxicological purposes and biomedicine research, characterized by:

- Based on human induced pluripotent stem cells (hiPSCs)
- Emphasis on mimicking the natural development of Leydig cells (LCs), focusing on critical developmental windows
- Replication of physiological conditions through **3D cell culture**

pivotal role not only in reproduction health but also in development, behaviour, gender identity and many other processes. Implementation of hormonally controlled conditions, facilitating the study of men's individual hormone ratios, daily variations, and age-related changes.

CURRENT IN VITRO MODELS

Conducted a comprehensive review of current in vitro models for testicular steroidogenesis (**Figure 1**), revealing limitations:

- One OECD validated test employs H295R cell line, derived from adrenal carcinoma (female patient) → possibly lacking gonadal relevance.
- Majority of models rely on rodent (74%), cancer (75%) cell lines → Species differences and lack of key enzymes → lacking human relevance.
- Predominantly cultured in 2D (94%) → ignoring the physiological microenvironment.



METHODOLOGY



2D DIFFERENETIATION PROTOCOL ACCORDING TO ROBINSON ET AL (2023)²



Figure 1: Overview of characteristics of currently utilized in vitro testicular steroidogenesis models. F, female; LCs, Leydig cells, M, male.

RESULTS 2D CONDITION

Successful adaptation of differentiation protocol:

The presence of lipid droplets typical for LCs present in LCs-like cells (**Figure 2**).

- Decrease of pluripotency markers (Nanog).
- Expression of NR5A1 and LHCGR driving testicular steroidogenesis.



Figure

Expression of steroidogenic enzymes (STAR, HSD3B2, HSD3B1, CYP17A1, CYP11A1).

SLCs-like cells (D0) exhibit mitotic activity while showing markers indicating a transition toward LCs differentiation (**Figure 3 & 4**) \rightarrow optimal phase for transfer to 3D conditions.







Figure 2: Accumulation of neutral lipid-containing lipid droplets, blue: nucleus by DAPI; green: lipid droplets by AdiporedTM.



RESULTS 3D CONDITION

- Mouse cell line **TM3 was used for the optimization** process.
- Encapsulation into Matrigel® Matrix unsuitable for dynamic culture \rightarrow hydrogel breakdown.
- Successful optimization of encapsulation of TM3 cells into alginate and long-term culture in dynamic conditions in RWV.



Figure 6: TM3 cells encapsulated in Matrigel® Matrix under static culture



Figure 4: qPCR results, fold change relative to the level in hiPSCs.



Figure 5: TM3 cells encapsulated in alginate, culture under dynamic cultivation in RWV

REFERENCE

[1] World Health Organization. (Licence: CC BY-NC-SA 3.0 IGO., 2023)

HSD17B3, and NC (negative control) in SLCs-like cells (D0). Red:

3: Immunostaining analysis of HSD3B1, LHCGR,

[2] Robinson, M. et al. Adv Biol 7, (2023)

Target protein; Blue: DAPI for nuclear staining.

ACKNOWLEDGMENT

Research is supported by the Czech Science Foundation project No. GA22-30004S. Images were created in BioRender. Brno Ph.D. Talent Scholarship Holder- Funded by the Brno City Municipality.





Research group: Cell and Tissue Toxicology, RECETOX

