Metabolic dynamics of human Sertoli cells are differentially modulated by physiological and pharmacological concentrations of GLP-1

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ABSTRACT
Obesity incidence has pandemic proportions and is expected to increase even further. Glucagon-like peptide-1 (GLP-1) based therapies are well-established pharmacological resources for obesity treatment. GLP-1 regulates energy and glucose homeostasis, which are also crucial for spermatogenesis. Herein, we studied the GLP-1 effects in human Sertoli cells (hSCs) metabolism and mitochondrial function. hSCs were cultured in absence or exposed to increasing doses of GLP-1 mimicking physiological post-prandial (0.01 nM) levels or equivalent to pharmacological levels (1 and 100 nM) used for obesity treatment. We identified GLP-1 receptor in hSCs. Consumption/production of extracellular metabolites were assessed, as well as protein levels or activities of glycolysis-related enzymes and transporters. Mitochondrial membrane potential and oxidative damage were evaluated. Glucose consumption decreased, while lactate production increased in hSCs exposed to 0.01 and 1 nM GLP-1. Though lactate dehydrogenase (LDH) protein decreased after exposure to 100 nM GLP-1 its activity increased in hSCs exposed to the same concentration of GLP-1. Mitochondrial membrane potential decreased in hSCs exposed to 100 nM of GLP-1, while formation of carbonyl groups was decreased in those cells. Those effects were followed by an increase in p-mammalian target of rapamycin (mTOR) Ser(2448). Overall, the lowest concentrations of GLP-1 increased the efficiency of glucose conversion to lactate, while GLP-1 concentration of 100 nM induces mTOR phosphorylation, decreases mitochondrial membrane potential and oxidative damage. GLP-1 regulates testicular energy homeostasis and pharmacological use of GLP-1 analogues could be valuable to counteract the negative impact of obesity in male reproductive function.

1. Introduction
Obesity has emerged as a major healthcare problem increasing the risk of cardiovascular diseases and decreases lifespan (Fontaine et al., 2003). Reduced fertility is a silent obesity-related complication, which among other co-morbidities is becoming a matter of concern (Alves et al., 2016). Dysregulation in the hormonal control of energy homeostasis promoted by the balance between food intake and energy expenditure has been hypothesized to interfere not only in body metabolism, but also with fertility though the underlying mechanisms remain unknown.

Glucagon like peptide-1 (GLP-1) is a 29-amino acid amidated...
peptide hormone, produced by posttranslational cleavage of the proglucagon precursor protein (Bell et al., 1983). This peptide is mainly produced by intestinal L-cells and secreted post-prandially (Guedes et al., 2015). GLP-1 is known for its role in glucose homeostasis predominantly mediated by the incretin effect. It potentiates insulin secretion after oral glucose ingestion, as compared to intra-venous glucose administration. GLP-1 also acts in the central nervous system to reduce the appetite and delays gastric emptying. Glucose equilibrium is crucial for energy homeostasis and the proper functioning of physiological functions, including male fertility. However, due to in vivo fast-proteolytic digestion mainly by dipeptidyl peptidase 4 (DPP4) and renal clearance, GLP-1 has a short half-life (Ruiz-Grande et al., 1993) that limits the use of the native peptide for pharmacological purposes. Thus, several GLP-1 analogues are available, being options for treatment of diabetes and obesity, as these promote weight loss, while having a low risk of causing hypoglycaemia (Crane and McGowan, 2016).

GLP-1 exerts its activity via GLP-1 receptor (GLP-1R), which belongs to a G-protein coupled receptor family (Bullock et al., 1996). GLP-1R expression within the male reproductve system has been detected in mice Sertoli cells (SCs) (Zhang et al., 2015), evidencing that GLP-1 should have an impact on male fertility. In fact, GLP-1R knockout mice have reduced seminal vesicle and gonadal weights, despite depicting normal testicular sex steroids levels and retaining fertility (MacLusky et al., 2000).

While it is known that GLP-1 and its analogues are key regulators of metabolism, the molecular means by which they impact cell metabolism, remain unknown. It has been postulated that mitochondria are the main metabolic targets for this hormone (for review Alves et al., 2016). For instance, GLP-1 improves mitochondrial membrane potential (Ogata et al., 2014) and also mitochondrial mass in a pancreatic cell line (Kang et al., 2015). Indeed, both GLP-1 analogues and DPP4 inhibitors, which increase endogenous GLP-1 levels by preventing its proteolytic digestion mainly by dipeptidyl peptidase 4 (DPP4) and renal clearance, GLP-1 has a short half-life (Ruiz-Grande et al., 1993) that limits the use of the native peptide for pharmacological purposes. Thus, several GLP-1 analogues are available, being options for treatment of diabetes and obesity, as these promote weight loss, while having a low risk of causing hypoglycaemia (Crane and McGowan, 2016).

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While it is known that GLP-1 and its analogues are key regulators of metabolism, the molecular means by which they impact cell metabolism, remain unknown. It has been postulated that mitochondria are the main metabolic targets for this hormone (for review Alves et al., 2016). For instance, GLP-1 improves mitochondrial membrane potential (Ogata et al., 2014) and also mitochondrial mass in a pancreatic cell line (Kang et al., 2015). Indeed, both GLP-1 analogues and DPP4 inhibitors, which increase endogenous GLP-1 levels by preventing its degradation, modulate mitochondrial structure and functioning (for review Ranganath, 2008). These metabolic processes depend on signalling pathways network. The mammalian target of rapamycin (mTOR) kinase emerged as pivotal for these processes (for review Oliveira et al., 2017), controlling mitochondrial bioenergetics and glucose metabolism. GLP-1 interferes with mTOR pathway (Park et al., 2015) and recently it was suggested that mTOR has an essential role in male reproduction (Jesus et al., 2017). In addition, mTOR controls glucose consumption and redox balance of human SCs (hSCs) (Jesus et al., 2016). The SCs are responsible for the nutritional support of spermatogenesis and the major hormonal target inside the testis (Martins et al., 2015; Martins et al., 2016). Herein, we studied the GLP-1 effects on hSCs metabolism and mitochondrial function. We hypothesized that GLP-1 could interfere in hSC function with potential implications for the nutritional support of spermatogenesis.

2. Methods

2.1. Chemicals

All chemicals were obtained from Sigma-Aldrich (St. Louis, MO, USA) unless specified otherwise.

2.2. Patient’s selection and testicular tissue preparation

Testicular tissue was obtained from testicular biopsies (n = 6) performed to men with conserved spermatogenesis, suffering from anejaculation due to previous vasectomy or traumatic section of the vas deferens, with the ultimate aim of recovering gametes for medically assisted procreation. After informed written consent, hSCs were isolated from the testicular tissue left in culture plate once gamete retrieval was completed. Testicular biopsies and handling of testicular tissue was done at the Centre for Reproductive Genetics Professor Alberto Barros (Porto, Portugal) in accordance with the Guidelines of Local, National and European Ethical Committees and performed in agreement with the Declaration of Helsinki.

2.3. Sertoli cells culture

Human SCs were isolated following the protocol optimized by Oliveira and collaborators (Oliveira et al., 2009). Anti-Mullerian hormone and vimentin, were used, as specific protein markers, to assess the purity of the hSCs cultures (Steger et al., 1996). Only cultures with a purity above 95% were used, SC cultures purity was determined by phase contrast microscopy using an ExtrAvidin Peroxidase Staining Kit.

To establish the various experimental groups, cells were washed with PBS, and placed in culture serum-free medium (DMEM: Ham’s F12, pH7.4) with insulin-transferrin-sodium selenite medium (final concentration of 10 mg/L; 5.5 mg/L; 6.7 μg/L, respectively) to which GLP-1 (Bachem AG, Bubendorf, Switzerland) was added (or not). One group of cells was not exposed to GLP-1 (0 nM) (no GLP-1) and three other groups were treated with increasing concentrations of GLP-1 (0.01, 1 and 100 nM). The concentration of 0.01 nM was chosen taking into consideration the postprandial GLP-1 levels found in healthy individuals (Vilsboll et al., 2001; Balestrieri et al., 2015). The other GLP-1 concentrations were chosen to mimic the highest plasmatic concentrations attained after administration of a GLP-1 analogue at the maximum therapeutic dosage recommended for obese individuals (3 mg/daily for liraglutide) either after a single administration (1 nM) or at steady state after a daily administration for 5 weeks (100 nM) (Elbround et al., 2002; Jiang et al., 2011; Danne et al., 2017).

After 6 h of treatment, cell culture medium was also collected. Afterwards, the cells were washed and exposed to GLP-1 for 6 h. We chose this time period for the exposure of cells to GLP-1 based on GLP-1 half-life (Ruiz-Grande et al., 1993). The cells were then washed with PBS and detached using a Trypsin-EDTA solution (0.05% / 0.02% (w/v)), counted using a Neubauer chamber and collected for protein, DNA and RNA extraction. Only groups with viability averaging 85–90%, evaluated by Trypan Blue Exclusion test, were considered for analysis.

2.4. RNA and DNA extraction

Total RNA (RNA) was extracted using the E.Z.N.A.* RNA commercial kit (Omega Bio-Tek, Norcross, USA) and DNA was extracted using the E.Z.N.A.* Tissue DNA Kit Commercial Kit (Omega Bio-Tek, Norcross, USA), as indicated by the manufacturer. The amount of DNA and RNA was determined using a NanoDrop 1000 Spectrophotometer (Thermo Fisher Scientific, Wilmington, USA). The 260/280 nm of the samples were used to assess the purity of DNA and RNA. DNA presented a ratio of ≈1.8 and RNA presented a ratio of ≈2 in the extracted samples. To assess the integrity of RNA and DNA, an aliquot of each sample was run on a denaturing agarose gel stained with GreenSafe (NZYTech, Lisboa, Portugal).

2.5. cDNA synthesis and Reverse Transcriptase Polymerase Chain Reaction (RT-PCR)

RNA was reversely transcribed as previously described (Martins et al., 2015). The resultant cDNA was used with exon-exon spanning primer sets designed to amplify specific cDNA fragments (Table 1). PCR was performed with a primer’s optimal annealing temperature, using standard methods (Martins et al., 2015) (Table 1). The samples were visualized using the software Image Lab (BioRad, Hercules, CA, USA) coupled to an image acquisition system BioRad FX-Pro-plus (BioRad, Hemel Hempstead, UK). The product size was compared to a DNA ladder (NZYTech, Lisboa, Portugal). Human heart and human liver RNAs (AMS Biotechnology, Abingdon, UK) were used as positive control and cDNA-free sample was used as negative control.

The mRNA expression levels of GLP-1R were evaluated in the different experimental groups. qPCR experiments were carried out in
encoded beta-2-microglobulin (β2M) gene were used to quantify mtDNA copy number relative to the following model proposed by Pfaffl: $2^{-\Delta\Delta CT}$, the fold variation the target gene expression was calculated.

### 2.6. Determination of mtDNA copy number

A qPCR analysis was performed to study the mtDNA copy number as described (Fuke et al., 2011) with small modifications. The efficiency of the amplification was determined by serial dilutions of cDNA (Martins et al., 2015). Amplification conditions were followed as previously described (Martins et al., 2016). β-2-microglobulin (β2M) transcript levels were used to normalize the mRNA expression of the target genes. The target genes, sequences and annealing temperatures of the primers are described in Table 1. Following the mathematical model proposed by Pfaffl in the formula: $2^{-\Delta\Delta CT}$, the fold variation the target gene expression was calculated.

### 2.7. Cytotoxicity assay

A sulforhodamine B (SRB) colorimetric assay was performed to test the cytotoxicity of GLP-1 to hSCs (Skehan et al., 1990). The cells were seed and treated with the selected concentrations of GLP-1. After 6 h, the assay was performed as previously described (Martins et al., 2015). No cytotoxicity was observed for the GLP-1 concentrations used (data not shown).

### 2.8. Western blot

Total proteins isolated from hSCs were extracted using the M-PER Mammalian Protein Extraction Reagent (Thermo Scientific, Rockford, USA). Western blot was performed as previously described (Alves et al., 2014). The membranes were incubated overnight at 4 °C with primary and secondary antibodies (Table 2). Mouse β-tubulin was used as protein loading control. ECF detection system was used and the membranes were read in the BioRad FX-Pro-plus (BioRad, Hemel Hempstead, UK). The densities from each band were quantified according to standard methods using Image Lab (BioRad, Hemel Hempstead, UK). Whenever possible the membranes were stripped with Restore Western Blot Stripping Buffer (Thermo Scientific, Rockford, USA) following the manufacturer's instructions, blocked and marked with other primary and consequently secondary antibody.

### 2.9. Lactate dehydrogenase (LDH) activity

The commercial Pierce LDH Kit (Thermo Scientific, Rockford, USA) was used to measure LDH activity. The obtained values of activity were calculated using the molar absorptivity of formazan ($e = 19,900 \text{ M}^{-1} \text{cm}^{-1}$).

### 2.10. Measurement of oxidative damage

The oxidative damage to proteins and lipids was evaluated by Slot-blot. We determined the protein carbonyl, nitro-tyrosine (NT) and 4-hydroxynonenal (4-HNE) group levels as previously described (Martins et al., 2015). The membranes were incubated with primary and secondary antibodies (Table 2). ECF detection system was used, the membranes were read in the BioRad FX-Pro-plus and the densities from each band were quantified using Image Lab (BioRad, Hemel Hempstead, UK).

### 3. Results

#### 3.1. GLP-1 receptor is expressed in hSCs

GLP-1R was previously identified in the mice testis, only by immunohistochemistry (Zhang et al., 2015). We investigated the presence of GLP-1R in hSCs and detected a 300 bp amplicon, corresponding to the presence of GLP-1R mRNA (Fig. 1A). Our results also show that, when exposed to the different concentrations of GLP-1, hSCs do not present alteration in the transcript levels of GLP-1R (Fig. 1B).
acetate (Alves et al., 2012), which is exported to the intratubular fluid by monocarboxylate transporters (MCTs), as is lactate. Our results showed that lactate production increased to 11.49 ± 2.06 and 1.34 ± 0.31 pmol/cell of acetate (Supplementary Fig. 2C). On the other hand, acetate production by hSCs exposed to 1 nM of GLP-1 increased to 2.49 ± 0.38 pmol/cell when compared to non-exposed cells that produced 1.34 ± 0.31 pmol/cell of acetate (Supplementary Fig. 1D).

3.3. Exposure to GLP-1 increased lactate production by hSCs

Once inside the cell, glucose is metabolized to pyruvate. Pyruvate is consumed by hSCs exposed to GLP-1 as their major source of energy. Our data showed that mitochondrial membrane potential decreased to 0.78 ± 0.11-fold variation relative to non-exposed cells when compared to hSCs exposed to 100 nM of GLP-1, which, presented an activity of 0.78 ± 0.11-fold variation relative to non-exposed cells (Fig. 2D).

3.4. Human SCs exposed to 100 nM of GLP-1 presented decreased mitochondrial membrane potential

Our data showed that mitochondrial membrane potential decreased in SCs exposed to 100 nM (ratio of 0.61 ± 0.02), when compared to non-exposed or those exposed to 0.01 and 1 nM of GLP-1 (ratio of 1 ± 0.06; 1.15 ± 0.07 and 1.11 ± 0.10, respectively) (Fig. 3A). Still, when we analysed the relative quantity of mitochondrial DNA no alteration was observed in hSCs exposed to GPL-1 when compared to non-exposed cells (Fig. 3B).

3.5. Carbonyl group levels decreased in hSCs exposed to 1 and 100 nM when compared to 0.01 nM of GLP-1

The high glycolytic flux, followed by the conversion of pyruvate to lactate, as happens in SCs, enhances a pro-oxidative environment. The products of protein carbonylation (dinitrophenol (DNP)), nitration (NT) and lipid peroxidation (4-HNE) are biomarkers for oxidative stress.

Table 2

| Antibody | Source | Molecular Weight (kDa) | Dilution | Vendor | Catalog # |
|----------|--------|------------------------|----------|--------|-----------|
| GLUT1    | Rabbit | 55                     | 1:500    | Santa Cruz Biotechnology, Heidelberg, Germany | sc-7903   |
| GLUT2    | Rabbit | 61                     | 1:5000   | Santa Cruz Biotechnology, Heidelberg, Germany | sc-9117   |
| GLUT3    | Goat   | 48                     | 1:500    | Santa Cruz Biotechnology, Heidelberg, Germany | sc-7582   |
| MCT4     | Rabbit | 43                     | 1:1000   | Santa Cruz Biotechnology, Heidelberg, Germany | sc-50529  |
| LDH      | Rabbit | 37                     | 1:10000  | Abcam, Cambridge, MA, USA | ab52488   |
| Catalase | Mouse  | 60                     | 1:4000   | Sigma-Aldrich, Roedermark, Germany | C0979     |
| P-mTOR   | Rabbit | 289                    | 1:500    | Cell Signaling Technology, Danvers, MA, USA | 2971S     |
| β-Tubulin| Mouse  | 50                     | 1:5000   | Thermo Fisher Scientific, Rockford, USA | MAS-16308 |
| DNP      | Rabbit | —                      | 1:5000   | Sigma-Aldrich, Roedermark, Germany | D9656     |
| 4-HNE    | Goat   | —                      | 1:5000   | EMD Millipore, Temecula, CA, USA | AB5605    |
| Nitro-Tirosine | Rabbit | —                      | 1:2500   | Cell Signaling Technology, Danvers, MA, USA | 9691S     |
| Mouse    | Goat   | —                      | 1:5000   | Sigma-Aldrich, Roedermark, Germany | A3562     |
| Rabbit   | Goat   | —                      | 1:5000   | Sigma-Aldrich, Roedermark, Germany | A3687     |
| Goat     | Goat   | —                      | 1:5000   | Sigma-Aldrich, Roedermark, Germany | A4187     |

Abbreviations: 4-HNE- 4 Hydroxynonenal; DNP- Dinitrophenol; GLUT1- Glucose transporter 1; GLUT2- Glucose transporter 2; GLUT3- Glucose transporter 3; LDH- Lactate dehydrogenase; MCT4- Monocarboxylate transporter 4; P-mTOR- Phospho-mTOR

**Fig. 1.** Identification and effects of exposure to glucagon like peptide-1 (GLP-1) in its receptor (GLP-1R) in human Sertoli cells. Panel A - Expression of GLP-1R in cultured human Sertoli cells identified by reverse transcriptase polymerase chain reaction. Panel B - Expression of mRNA transcript levels of GLP-1R in human Sertoli cells exposed to increasing concentrations of GLP-1. Panels B shows pooled data of independent experiments, indicating the mRNA transcript levels of GLP-1R. hSCs: human Sertoli cells; hH: human heart; Neg: Negative control; Neg(RT): Negative without reverse transcriptase. Results are expressed as mean ± SEM (n = 6 for each condition).

**Fig. 2.** Glucose consumption is decreased in hSCs exposed to GLP-1

Glucose is essential for SCs metabolism and spermatogenesis. This metabolite enters SCs mostly through glucose transporters (GLUTs), GLUT1–3 (Martins et al., 2015; Martins et al., 2016). After treatment with GLP-1, no alterations were detected in the protein levels of those GLUTs in hSCs (Supplementary Fig. 1A to C). Nevertheless, when hSCs were exposed to 0.01 and 1 nM of GLP-1, glucose consumption decreased to 45.35 ± 18.40 and 25.66 ± 15.60 pmol/cell, respectively, while cells exposed to 100 nM of GLP-1 consumed 75.82 ± 28.40 pmol of glucose/cell and those not exposed to GLP-1 consumed 122.30 ± 30.70 pmol of glucose/cell (Fig. 2A). SCs also produce acetate (Alves et al., 2012), which is exported to the intratubular fluid by monocarboxylate transporters (MCTs), as is lactate. Our results show that hSCs exposed to all GLP-1 concentrations did not alter the protein levels of MCT4 (Supplementary Fig.1D). On the other hand, acetate production by hSCs exposed to 1 nM of GLP-1 increased to 2.49 ± 0.38 pmol/cell when compared to non-exposed cells that produced 1.34 ± 0.31 pmol/cell of acetate (Supplementary Fig. 2C).
Carbonyl group levels did not alter in any of the hSCs exposed groups to GLP-1 when compared with the no GLP-1 group. However, hSCs exposed to 1 and 100 nM of GLP-1 presented lower levels of carbonyl groups (0.78 ± 0.07 and 0.86 ± 0.05-fold variation to non-exposed cells (no GLP-1), respectively (Fig. 4A)) when compared with hSCs exposed to 0.01 nM of GLP-1 (1.10 ± 0.02-fold variation to non-exposed cells (no GLP-1)). Yet, the levels of 4-HNE and NT groups were not altered in cells exposed to GLP-1 when compared with non-exposed cells (Fig. 4B and C, respectively). Additionally, we determined the protein levels of antioxidant enzymes such as catalase and the total/reduced glutathione ratio. Our results show that cells exposed to all GLP-1 concentrations did not present alterations in the protein levels of catalase or in the total/reduced glutathione ratio when compared with the levels detected in non-exposed cells (Supplementary Fig. 3A and 3B, respectively).

Fig. 2. Effects of glucagon like peptide-1 (GLP-1) in glucose metabolism in human Sertoli cells. Glucose consumption (Panel A), lactate production (Panel B), expression of lactate dehydrogenase (LDH) (Panel C) and LDH activity (Panel D) in human Sertoli cells after exposure to increasing concentrations of GLP-1 are represented. Panels A and B show pooled data of independent experiments, indicating the glucose consumption and lactate production, respectively. Panels C and D show pooled data of independent experiments, indicating the protein levels and activity of LDH, respectively. Panels C also show representative Western Blot experiments. Results are expressed as mean ± SEM (n = 6 for each condition). Significantly different results (p < .05) are as indicated: * relative to non-exposed cells (no GLP-1); § relative to 1 nM.

Fig. 3. Effects of glucagon like peptide-1 (GLP-1) in mitochondria function of human Sertoli cells. JC-1 ratio (Panel A) and mitochondrial relative copies (Panel B) in human Sertoli cells exposed to increasing concentrations of GLP-1 are represented. Panel A shows JC-1 ratio found in human Sertoli cells cultured with increasing concentrations of GLP-1 or without GLP-1. Panel B shows pooled data of independent experiments indicating mitochondrial DNA (mtDNA) quantity by quantitative PCR. Results are expressed as mean ± SEM (n = 6 for each condition). Significantly different results (p < .05) are as indicated: * relative to non-exposed cells (no GLP-1); # relative to 0.01 nM; § relative to 1 nM.
3.6. Phosphorylated mTOR is increased in hSCs exposed to 100 nM of GLP-1

mTOR has a crucial role in coordinating cellular homeostasis and energy status (Chiang and Abraham, 2005). Recent studies demonstrate that mTOR signalling modulates the glycolytic and oxidative profile in hSCs (Jesus et al., 2016). GLP-1 had no effects regarding mTOR phosphorylation in hSCs of any of the exposed groups when compared with expression in hSCs of no GLP-1 group. Still, hSCs exposed to 100 nM of GLP-1 presented an increase in the protein levels of phosphorylated mTOR (Ser2448), to 1.41 ± 0.28-fold variation to non-exposed cells (no GLP-1) when compared to hSCs exposed to 0.1 nM of GLP-1 (0.77 ± 0.12-fold variation relative non-exposed cells (no GLP-1)) (Fig. 5).

4. Discussion

GLP-1 is a peptide hormone with an active role in the regulation of circulating glucose levels (Brubaker, 2006). Glucose equilibrium is crucial for energy homeostasis and the proper functioning of physiological functions, including male fertility (Dunning et al., 2005). Excessive fat accumulation is associated with dysregulation of energy homeostasis signalling systems, and with a concurrent impairment of GLP-1 mediated functions (Dirksen et al., 2013). Several GLP-1 analogues are now used as pharmacological agents to promote weight loss (Blundell et al., 2017). Recent studies have consistently associated the increase in obesity rates with decreased male fertility (Rato et al., 2014a). In 2030, it is estimated that over 50% of the world population will be overweight/obese (Smith and Smith, 2016), which will aggravate male infertility trends. Alterations in SC metabolism were hypothesized to contribute for reduced fertility in obese males (Winters et al., 2006). In fact, the SCs are responsible for the nutritional support of spermatogenesis.

There is evidence that GLP-1 may influence male reproductive function (Jeibmann et al., 2005), although few studies have been focused on the molecular mechanisms. Since SCs are major hormonal targets within the testis (Alves et al., 2014; Martins et al., 2015; Martins et al., 2016), we hypothesized that GLP-1 could directly modulate the metabolic functions of these cells and has a possible effect on male fertility. For that, we exposed hSCs to increasing doses of GLP-1 to evaluate: (1) the impact of the postprandial levels of GLP-1; (2) the impact of GLP-1 when present at the levels reached by its analogues (liraglutide) in the plasma of healthy/obese individuals. The maximum plasma concentrations found in overweight and obese subjects treated with GLP-1 analogues is similar to the one observed in healthy individuals (Onge et al., 2016). GLP-1 actions are mediated by the GLP-1R. So far GLP-1R has only been identified in mice SCs by immunohistochemistry of the testicular tissue (Zhang et al., 2015). Our results allowed us to observe for the first time the expression of GLP-1R in isolated hSCs. Interestingly, GLP-1 exposure did not change GLP-1R expression in cultured hSCs, illustrating that the mRNA expression of this receptor in these cells is not rapidly responsive to exposure to increasing GLP-1 concentrations.

We then evaluated the impact of GLP-1 postprandial levels in hSCs metabolism and bioenergetics. These cells uptake glucose from the interstitial fluid by the action of GLUTs (GLUT1–3). Glucose is then used to produce essential metabolites (mainly lactate) that will serve as energy sources for developing germ cells (Martins et al., 2015; Martins et al., 2016). Although GLP-1 exposure did not alter GLUTs expression, hSCs decreased the consumption of glucose after exposure to postprandial levels of this hormone, as compared to non-exposed cells. Concurrent results had been reported in other mammalian cell lines, with GLP-1 exposure decreasing deoxyglucose uptake (Morales et al., 2014). Our results suggest that this hormone is capable of eliciting an alteration of glucose uptake, likely due to an alteration on its
metabolism. Still, GLP-1 postprandial levels were capable to stimulate lactate production by hSCs. Previous studies, show that lactate is essential for spermatogenesis (Oliveira et al., 2011) since it is used as metabolic fuel by developing germ cells (Boussour and Benahmed, 2004) and has an anti-apoptotic effect on those cells (Rato et al., 2014b). The postprandial levels of GLP-1 increased the production of lactate, while glucose consumption was decreased, as compared to non-exposed cells, illustrating the metabolic commitment of these cells to an efficient production of lactate. No effect was detected on mitochondria functionality in these cells. Neither mitochondrial membrane potential nor DNA content were affected by exposure to these levels of GLP-1. Moreover, the oxidative stress markers (protein carbonyl and nitrotyrosine groups, and lipid peroxides) were not altered, which correlates with a normal functioning of the mitochondria. Hence, at postprandial levels, GLP-1 seems to be vital for eliciting the production of lactate by hSCs, which in turn consume lower amounts of glucose. These cells have adaptive mechanisms, and show a metabolic plasticity (Alves et al., 2014; Martins et al., 2015; Martins et al., 2016; Meneses et al., 2016) that is very important to sustain the metabolic support of spermatogenesis. They utilize a wide sort of metabolic sources to produce lactate, namely from triglycerides that accumulate as lipid droplets (Gorga et al., 2017). Depending on the substrate availability and on stimuli, SCs can oxidize these cytoplasmic lipidic droplets to support their metabolic requirements (Jutte et al., 1985). It has been described that GLP-1 may promote lipid droplet remodelling and lipolysis in human adipocytes, when present at concentrations above 10^{-11}M (Villanueva-Penacarrillo et al., 2001). A similar event may be occurring in hSCs exposed to the postprandial levels of GLP-1 (Vilsbøll et al., 2001; Balestrieri et al., 2015).

Similar results were obtained when the hSCs were exposed to GLP-1 concentrations that mimic the levels of GLP-1 analogues observed in the plasma after a single administration of liraglutide at the therapeutic dosage recommended for obese individuals (Elboudou et al., 2002). Both glucose consumption and lactate production were maintained, when compared to cells exposed to postprandial levels of GLP-1. The SC is responsible for producing lactate favouring the glycolytic flux (Oliveira et al., 2015), but it also uses mitochondrial oxidative phosphorylation to sustain its own energetic needs. Human SC exposed to 1 nM of GLP-1 presented an increase in acetate production, however this did not lead to an increase in mitochondrial membrane potential, as observed in other cell lines (Moraes et al., 2014). No changes were observed in the pro-oxidant environment, nitration and lipid peroxidation, nor in the levels of antioxidant defences, but we found less oxidative damage in proteins of cells exposed to this concentration of GLP-1. This protective effect mediated by GLP-1 may be essential to counteract the pro-oxidant environment promoted by an increased metabolic activity.

Contrasting results were obtained with cells exposed to GLP-1 concentrations that mimic the levels of GLP-1 analogues observed in the plasma after a prolonged administration of liraglutide at the therapeutic dosage recommended for obese individuals (Jiang et al., 2011; Danne et al., 2017). In this case, hSCs appear to consume higher amounts of glucose, producing the same amounts of lactate, as compared to cells exposed to postprandial GLP-1 levels. This increased glucose consumption was associated with LDH activity stimulation, which suggests an adaptation to sustain lactate production. In fact, hSCs exposed to 100 nM of GLP-1, presented a decrease in mitochondrial membrane potential. As said, SCs have a distinct metabolic behaviour when exposed to GLP-1, presented an increase in acetate production, however this did not lead to an increase in mitochondrial membrane potential, as observed in other cell lines (Moraes et al., 2014). No changes were observed in the pro-oxidant environment, nitration and lipid peroxidation, nor in the levels of antioxidant defences, but we found less oxidative damage in proteins of cells exposed to this concentration of GLP-1. This protective effect mediated by GLP-1 may be essential to counteract the pro-oxidant environment promoted by an increased metabolic activity.

Conflict of interest

The authors declare that they have no conflict of interest.

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Appendix A. Supplementary data

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