



UNIVERSIDADE DE LISBOA
Faculdade de Medicina Veterinária

EVALUATION OF DIFFERENT EXTENDERS FOR COLD STORAGE OF MEAGRE
(*ARGYROSOMUS REGIUS*) SEMEN

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CONSTITUIÇÃO DO JÚRI

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DISSERTAÇÃO DE MESTRADO INTEGRADO EM MEDICINA VETERINÁRIA

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ABSTRACT

“EVALUATION OF DIFFERENT EXTENDERS FOR COLD STORAGE OF MEAGRE (*ARGYRO SOMUS REGIUS*) SEMEN”

Semen refrigeration is usually recommended as a cheap and simple procedure that facilitates artificial reproduction techniques. The main objective of this experiment was to develop a semen refrigeration protocol for meagre that is considered a potential candidate for aquaculture diversification in Southern Europe. This thesis also contributes for the understanding of the causes of the fish sperm quality degradation during refrigeration.

Three extenders (non-activating medium (Fauvel et al., 1998); NaCl 0.9% and; NaCl 0.9% with glycine and glucose) and three different dilutions (1:4, 1:9 and 1:19, sperm:extender) were tested in a full factorial design. The following quality parameters were assessed along the storage time: sperm motility parameters; percentage of viable sperm; adenosine triphosphate (ATP); lipid peroxidation in the form of malondialdehyde (MDA); and bacteriology.

The 0.9% NaCl and the 0.9% NaCl with glycine and glucose extenders and the dilutions 1:4 and 1:9 kept a higher percentage of motile cells for longer, as well as higher sperm velocity. Sperm viability and ATP had better results with 0.9% NaCl and the 0.9% NaCl with glycine and glucose extenders. The MDA values were lower in treatments with dilution 1:4 when compared to those with dilution 1:9. In the CFU/ml values, no differences were found between extenders and dilutions. Motility parameters were strongly correlated with viability, whereas no or weak correlations existed with the remaining parameters. Thus, motility and viability seem to have the most impact in the loss of semen quality. According to the results, meagre semen could be kept refrigerated using 0.9% NaCl, in dilution 1:4, for up to 10 days.

Key-words: Semen; Fish; *Argyrosomus regius*; Cold storage; Extenders

RESUMO

“AVALIAÇÃO DE DIFERENTES DILUIDORES PARA A REFRIGERAÇÃO DE SÉMEN DE CORVINA (*ARGYROSOMUS REGIUS*)”

A refrigeração de sémen é tipicamente recomendada como um procedimento barato e simples que facilita as técnicas de reprodução artificial. O principal objectivo desta experiência foi desenvolver um protocolo de refrigeração de sémen para a corvina, que é considerada uma potencial candidata para a diversificação de aquacultura no sul da Europa. Esta tese contribui também para a compreensão das causas da degradação da qualidade de sémen de peixe durante a refrigeração.

Três diluidores (*non-activating medium* (Fauvel et al., 1998); NaCl 0,9%; e NaCl 0,9% com glicina e glucose) em três diferentes diluições (1:4, 1:9 e 1:19, sémen:diluidor) foram testados num plano factorial completo. Os seguintes parâmetros de qualidade espermática foram avaliados ao longo do tempo de armazenamento: parâmetros de mobilidade do sémen; percentagem de espermatozoides viáveis; adenosina trifosfato (ATP); peroxidação lipídica na forma de malondialdeído (MDA); e bacteriologia.

Os diluidores NaCl 0,9% e NaCl 0,9% com glicina e glucose e as diluições 1:4 e 1:9 mantiveram uma percentagem de células móveis mais elevada por mais tempo, bem como maior velocidade dos espermatozóides. A viabilidade e o ATP tiveram melhores resultados com NaCl 0,9% e NaCl 0,9% com glicina e glucose. Os valores de MDA foram mais baixos em tratamentos com as diluições 1:4, quando comparados com aqueles com as diluições 1:9. Nos valores de bacteriologia não foram encontradas diferenças entre diluidores e diluições testadas. Os parâmetros de mobilidade correlacionaram-se fortemente com a viabilidade, enquanto inexistentes ou fracas correlações foram encontradas entre os restantes parâmetros. Por conseguinte, mobilidade e viabilidade parecem ter o maior impacto na perda de qualidade do sémen. De acordo com os resultados, o sémen de corvina pode ser mantido refrigerado usando o diluidor NaCl 0,9% na diluição 1:4 até 10 dias.

Palavras-chave: Sémen; Peixe; *Argyrosomus regius*; Refrigeração; Diluidores

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LIST OF ABBREVIATIONS, INITIALS AND ACRONYMS

AEC	Adenylate energy charge
ATP	Adenosine triphosphate
BSA	Bovine serum albumen
C	Control/undiluted
CASA	Computer-assisted sperm analysis
CFU	Colony-forming units
FAO	Food and Agriculture Organization
GnRH _a	Gonadotropin-releasing hormone agonist
GzLM	Generalized Linear Model
HBSS	Hanks' balanced salt solution
HPLC	High-performance liquid chromatography
IPMA Atmosphere)	Instituto Português do Mar e Atmosfera (Portuguese Institute of Sea and
LIN	Linearity
MDA	Malondialdehyde
MOT	Percentage of motile cells
NAM	Extender NAM
NAM4	Treatment with extender NAM in dilution 1:4
NAM9	Treatment with extender NAM in dilution 1:9
NAM19	Treatment with extender NAM in dilution 1:19
NC	Extender NC
NC4	Treatment with extender NC in dilution 1:4
NC9	Treatment with extender NC in dilution 1:9
NC19	Treatment with extender NC in dilution 1:19
NCG	Extender NCG
NCG4	Treatment with extender NCG in dilution 1:4
NCG9	Treatment with extender NCG in dilution 1:9
NCG19	Treatment with extender NCG in dilution 1:19
PI	Propidium iodide
PROG	Percentage of progressive spermatozoa
PS	Pseudomonas CN Agar
STR	Straightness
TBA	Thiobarbituric acid
TBARS	Thiobarbituric acid reactive substances
TSA	Trypticase soya agar
VAP	Average path velocity
VCL	Curvilinear velocity
VIAB	Viability
VSL	Straight line velocity

ACTIVITIES DEVELOPED IN THE SCOPE OF THE MASTER'S DEGREE

1. Internship

Faculty of Veterinary Medicine – University of Lisbon (FMV-UL)

For three weeks, in July of 2014, some work was done in the facilities of the Faculty of Veterinary Medicine dedicated to research concerning Fish and Aquaculture. The work consisted in a preliminary approach to a possible research concerning the behavior of a species of shrimp, *Palaemonetes varians*. The preferences of this species regarding color of the substrate was assessed. During this period, aquaria were prepared for the individuals, and their behavior was observed and registered. A visit to an aquaculture was also done.

During these three weeks it was possible to acquire some experience in the planning and implementation of a preliminary project.

Oceanário de Lisboa

The first segment of the internship took place from January 5th to March 27th of 2015, in Oceanário de Lisboa (Lisbon Oceanarium). The team of veterinarians was accompanied during the entirety of the internship. The species that were observed and handled during the internship consisted of: mostly fish, both bony and cartilaginous, from different ecosystems; some species of amphibians; aquatic birds; a species of reptile (*Caretta caretta*); and one species of mammal (*Enhydra lutris*). Some of the activities which I accompanied included observation and diagnosis of individuals, routine analysis, reproductive ecology and necropsies. Due to an agreement with the institution, this section won't be further elaborated. This internship allowed to obtain knowledge and a better understanding of aquatic animal medicine, mostly of fish, including some experience in reproduction of different species.

Aquaculture Research Station of the Portuguese Institute for the Ocean and Atmosphere (EPPO – IPMA)

The second segment of the internship took place from the 7th of April to the 26th of June of 2015, with two weeks of additional work in September, in the Aquaculture Research Station of the Portuguese Institute for the Ocean and Atmosphere, in Olhão. A portion of work took place in the facilities of the Center of Marine Sciences of the University of Algarve, in Faro. During this internship took place the research project addressed by this thesis. The activities done in the scope of this project were done mostly by myself and my supervisor, but a team of biologists, as well as aquaculture and research technicians provided us with valuable support.

Experience was obtained in induction of handling of fish, as well as induction of anesthesia and collection of semen by stripping. I prepared the extenders that were tested and used them to dilute the semen. The semen quality parameters that were evaluated also represent areas of acquired expertise. I learned to use a computer-assisted sperm analysis in order to evaluate concentration, motility and viability. Nonetheless, the latter was mostly evaluated by myself through observation and counting in a fluorescence microscope. I prepared solutions and did ATP extraction and quantification, which included manipulation of a microplate reader. The MDA assay procedure was observed by myself. For bacteriology, I prepared growth medium plates and performed inoculations, as well as counting of colonies. I also obtained experience in the isolation, propagation and identification of genera of bacteria with biochemical tests. Lastly, in general, time planning and teamwork skills were developed.

2. Formation and Publications

A poster was displayed at the Aquaculture Europe 2014 congress, organized by the European Aquaculture Society: “Studies on behavior and activity of the shrimp *Palaemonetes varians*”, by Santos, M. and Afonso, F. (2014).

In July of 2015 I attended the seminar “Meagre production – science and practice”, organized by the EPPO-IPMA. During this seminar the results of the several works of the AQUACOR project were presented, regarding different aspects of the production of meagre, such as: performance of different production systems; feed; reproduction; larvae and fry production.

In October of 2015 I attended the XV National Spanish Congress and I Iberian Congress of Aquaculture in Huelva (Spain), organized by the Spanish Aquaculture Society. The communications I attended were related to different subjects such as: reproduction in captivity of Senegalese sole (*Solea senegalensis*) and other species; diversification of aquaculture with species such as meagre (*Argyrosomus regius*) and common octopus (*Octopus vulgaris*); production and sanitary quality of mollusks and crustaceans; sustainability and environmental impact in the scope of aquaculture. A poster was displayed: “Refrigeração do sêmen de corvina (*Argyrosomus regius*): desenvolvimento de um protocolo e avaliação de parâmetros de qualidade” (Refrigeration of meagre semen (*Argyrosomus regius*): development of a protocol and evaluation of quality parameters), by Santos, M., Soares, F., Moreira, M., Cabrita, E. and Beirão, J. (2015).

BIBLIOGRAPHIC REVISION

1. Aquaculture in the world, and in Portugal

The human population is expected to increase to more than 11 billion by the end of the century (Lutz, Sanderson & Scherbov, 2001), leading to a bigger consumption of resources such as food (Pimentel, Huang, Cordova & Pimentel, 1997). This will result in an increase in fish consumption, a protein-rich food recommended as part of the human diet (Sargent & Tacon, 1999). Under these circumstances, and since the vast majority of the wild fish stocks are already overexploited or have reached their maximum productive capacity, further demand of fish should be met by aquaculture instead of capture (Grafton, Kompas & Hilborn, 2007).

The first written records regarding aquaculture date back to 2 500 years ago in China (Rabanal, 1988). Nowadays, it is the fastest growing animal production industry (Gjedrem, Robinson & Rye, 2012). The portion of global fish production corresponding to aquaculture, in 2015, was forecast to be 46% (Food and Agriculture Organization of the United Nations [FAO], 2015), while in 2003 it was only 31% (FAO, 2015). Quoting the words of the inventor of modern management and business visionary Peter Drucker, “Aquaculture, not the Internet, represents the most promising investment opportunity of the 21st century”.

The fish consumption per capita in Portugal is the highest in the European Union, and it is estimated to remain so, with a predicted consumption of 58 kg per capita per year, in 2020 (Failler, 2007). To support this high demand the country relies on a large volume of fish commodity imports, which in 2009 represented more than half of the national supply (Teixeira, Rodrigues, Cavadas & Neto, 2013). At the same time, only 5% of fish produced in Portugal comes from aquaculture, whereas the remaining 95% comes from fisheries (FAO, 2014). Thus, the development of aquaculture would decrease Portugal’s dependence on imports.

There are several factors in favour of the development of aquaculture in Portugal. As previously stated, the consumption of fish is very high. There is funding from the European Commission towards the development of aquaculture (Associação Portuguesa de Aquacultura [APA], 2014). Furthermore, it is one of the warmest European countries, due to the Mediterranean climate (Kalkan & Canuyrt, 2012). Finally, it has a vast coastline and a large area covered with lagoons, which can be used for aquaculture (Kalkan & Canuyrt, 2012). Nonetheless, the lack of economic power and the bureaucracy involved (licensing procedures take several years) have hindered the investment in aquaculture (APA, 2014).

Aquaculture production in Portugal, over the last 30 years, has fluctuated between 5 and 11 thousand tonnes, approximately (FAO, 2014). Of the total volume produced, 93% refers to brackish or marine aquaculture, and fish comprises 55% of its volume (Instituto Nacional de Estatística [INE], 2016). The main species produced in 2014 were grooved carpet shell (*Ruditapes decussatus*), turbot (*Scophthalmus maxima*), mussels (*Mytilus sp.*) and gilthead seabream (*Sparus aurata*) (INE, 2016). When it comes to marine fishes, the markets are currently saturated (Couto et al., 2016), especially because of competition with countries like Spain or Greece. An alternative to prevent this saturation is the development of technology for the production of new species. This is the aim of projects such as the AQUACOR project of IPMA, regarding meagre (*Argyrosomus regius*) and its production.

2. Biology of the meagre

The meagre, *Argyrosomus regius* (Asso, 1801), is part of the *Sciaenidae* family, which belongs to the Perciformes order. A common characteristic of this family, that gave its members the name of croakers and drums, are the sounds they produce by beating specific muscles against the swim bladder, making it resonate (Ramcharitar, Gannon & Popper, 2006). The *Argyrosomus* name comes from the Greek: *argyros*, meaning silver, and *soma*, body. Indeed, this species has a silver to bronze coloured elongated body, with a fairly large head and a yellow mouth cavity (Stipa & Angelini, 2005). In the wild, they reach up to 2 m in length and 50 kg in weight (Stipa & Angelini, 2005). The meagre's geographic distribution includes the Eastern Atlantic coast and the Mediterranean and Black Seas (Haffray et al., 2012; Lagardère & Mariani, 2006). They are carnivorous and, although the diet varies with their size, they feed mostly on small crustaceans when juveniles (Jiménez et al., 2005), and fish when adults (Quéro & Vayne, 1985; Stipa & Angelini, 2005).

Meagre are gonochorists (Gil, Grau, Basilone, Ferreri & Palmer, 2013; Schiavone, Zilli, Storelli & Vilella, 2012), meaning each individual has one constant gender. The reproductive season occurs from April to June (Gil et al., 2013), with aggregation near estuaries and salt marshes (Quéro, 1989; Stipa & Angelini, 2005). During this season the males emit the characteristic *croaks* (Lagardère & Mariani, 2006), whilst spawning occurs. The females lay eggs that are fertilized externally and hatch in 48 hours (Haffray et al., 2012). The larvae leave the nursing area as juveniles, when the summer ends, and migrate back to spawning areas upon reaching maturity (González-Quirós et al., 2011). During the winter, adults migrate to deeper waters, and reduce feeding activity. Water temperature seems to be a

determining factor in this species' biology, affecting namely its feeding requirements, trophic migration and reproduction (Stipa & Angelini, 2005).

Meagre appears to be a resilient species, due to its high fecundity (Haffray et al., 2012) and growth rate (Monfort, 2010). However, their aggregation and emission of spawning sounds makes meagre a species vulnerable to capture (Catalán, Jiménez, Alconchel, Prieto & Muñoz, 2006; Sadovy & Cheung, 2003).

3. Meagre as an advantageous species for aquaculture

Meagre has been farmed in Europe since the late nineties, first in France and then Italy. Since then, other countries like Spain, Portugal and Greece have started to show some interest in this species. In 2000, the total global production was of only 33 tonnes, but the number has been growing and in 2010 over 10 000 tonnes were produced in aquaculture (Monfort, 2010). This species has several strengths that support its production. It has a fast growth rate, reaching about 1 kg in the first year and 2.5 kg in the second (Duncan et al., 2013) and a low feed conversion rate, between 0.9 and 1.2 (Monfort, 2010). Meagre responds well to the hormonal induction of spawning and the eggs and larvae have high quality (Mylonas et al., 2013a; Mylonas, Mitrizakis, Sigelaki & Papadaki, 2011). A study by Roo, Hernández-Cruz, Borrero, Schuchardt & Fernández-Palacios (2010) obtained promising results for larval production at an industrial rate. As a euryhaline species, meagre tolerates diverse environments, such as land based cultivation with brackish water (Jiménez et al., 2005; Ribeiro, Soares, Quental-Ferreira, Gonçalves & Pousão-Ferreira, 2013), while also withstanding tank captivity fairly well (Pastor, Grau, Massutí & Sánchez-Madrid, 2002). It is also resistant to handling and there are few publications reporting disease problems, indicating that meagre might be somewhat resistant to infectious diseases (Stipa & Angelini, 2005). Additionally, it has a good filleting yield of 42% (Grigorakis, Fountoulaki, Vasilaki, Mittakos & Nathanailides, 2011) and its fillets received high sensory acceptability scores (Giogios, Grigorakis & Kalogeropoulos, 2013). The intramuscular fat content is low, around 1% (Grigorakis et al., 2013), when compared to species such as gilthead seabream (Özogul, Özogul & Alagoz, 2007) and European sea bass (*Dicentrarchus labrax*) (Lanari et al., 1999). At the same time, the fat has a high nutritional value (Giogios, Grigorakis & Kalogeropoulos, 2013) and its low content allows refrigeration for a longer period of time (Poli, et al., 2003). Altogether, these characteristics make meagre a very attractive species for the aquaculture diversification (Monfort, 2010; Poli et al., 2003), particularly in southern Europe.

As a new species adapted to aquaculture, important advances are expected to occur in meagre production, which should lead to a reduction of the juveniles' price (Stipa & Angelini, 2005). In this context, most of the research is focused on the improvement of meagre production techniques and the increase of its efficiency and profitability (Duncan et al., 2013). As an example, research in the field of reproduction will certainly result in regular production of high quality spawns.

4. Overview of meagre reproduction

Reproduction of meagre, as in fish in general, depends on the brain-hypothalamus-pituitary-gonad axis. Environmental stimuli like photoperiod and temperature are responsible for the control of this axis and have a crucial role in the control of gamete maturation and release (Cardinaletti et al., 2010). In the end of the axis lie the gonads, which are the reproductive organs in which gametes' production occurs, termed gametogenesis (Wootton & Smith, 2014). Gametes originated from germ cells after several mitotic divisions and meiosis (Okutsu, Suzuki, Takeuchi, Takeuchi & Yoshizaki, 2006; Stickney, 2000). In males, the gametogenesis occurs in the testes and is called spermatogenesis, producing spermatozoa. In females, the gametogenesis occurs in the ovaries and is called oogenesis, producing oocytes (Patiño & Sullivan, 2002). During oogenesis takes place the vitellogenesis that consists in the accumulation of yolk proteins within the oocyte (Patiño & Sullivan, 2002), essential for the formation of eggs. Mylonas, Mitrizakis, Papadaki & Sigelaki (2013) evaluated female and male gamete development of meagre over the reproductive period, in captivity, and observed that both spermatogenesis and vitellogenesis took place between April and June, in accordance with their natural reproductive season.

Meagre reach maturity at the age of 2.7 years for males and 3.5 years for females (Gil et al., 2013) and the length at maturity ranged 26.8-61.6 cm for males and 35.8-110 cm for females (González-Quirós et al., 2011; Schiavone et al., 2012; Soares et al., 2015). Additionally, Gil et al. (2013) concluded that the annual fecundity increases with length, weight and age, ranging from 0.9 to 4.2 million oocytes in reared females.

The reproduction of meagre in captivity is one of the bottlenecks of its production, similar to most species new to aquaculture (Schiavone et al., 2012). One example is the fry cost, which is still very high (Schiavone et al., 2012), in part because of unpredictability of its production. Nonetheless, industrial fish farming requires broodstocks that provide an adequate supply of fertilized eggs (Duncan et al., 2012) and juveniles (Stickney, 2000). However, some advances

in meagre reproduction are taking place. As an example, different research groups recently reported spontaneous spawning of meagre in captivity (Mylonas et al., 2013b; Pastor et al., 2013; Soares et al., 2015). Nowadays, fertilized eggs can either be obtained with wild-caught fish acclimated to captivity (Duncan et al., 2012; Soares et al., 2015) or with first-generation fish born in captivity (Soares et al., 2015). Indeed, when adequate nutrition is provided, both types of broodstocks can spawn good eggs and produce high larval quality, suitable for a hatchery production (Soares et al., 2015). Oviparous females in particular, like the meagre, have a higher and more diverse nutritional demand due to oogenesis (Stickney, 2000). A commonly observed problem in meagre reproduction is the failure to undergo oocyte formation and spawning. This also occurs in other species, due to a deficiency in the feed (Cabrita, Robles & Herráez, 2008).

Additionally, there have been some developments in hormonal induction protocols for meagre. These consist in the usage of a gonadotropin-releasing hormone agonist (GnRHa), given via injections or slow-release implants, to both males and females (Duncan et al., 2008; Duncan et al., 2012; Mylonas et al., 2013a; Pastor et al., 2013). Biglino (2015) observed a reduction in semen volume and sperm concentration when using GnRHa implants in males, but no apparent effect over motility or viability. At the same time, Duncan et al. (2008) and Pastor et al. (2013) obtained high egg fecundity after the administration of GnRHa to the females. Indeed, a single injection or a slow-release implant of GnRHa led to the production of fertilized eggs in sufficient number and quality for a commercially viable hatchery production (Duncan et al., 2012).

The current and future research should lead to new knowledge on meagre reproductive biology, which will be crucial to improve the broodstocks' reproductive performance (Mylonas et al., 2013b).

5. Artificial reproduction

Adequate management of broodstock and gametes requires certain environmental conditions and procedures, including artificial reproduction techniques. Artificial reproduction is the reproduction in a non-natural way, with the interference and manipulation of humans. It ensures the availability of fertilized eggs over time or when necessary, and maximizes the number of larvae produced (Suquet, Dreanno, Fauvel, Cosson & Billard, 2000). Furthermore, it grants some protection from the environment and pathogens, leading to a bigger survival rate and growth. In addition, species that cannot spawn naturally or reproduce in captivity can be propagated through these methods, using for example *in vitro* fertilization. Some

technologies require artificial reproduction. Polyploidy production, already done in other sciaenid species, is one example (Ballarin et al., 2004). It leads to the formation of fish that, being sterile, have less incidence of diseases or higher growth rate (Piferrer et al., 2009). Lastly, these techniques allow selective breeding, which consists in crossing the genetic material of specific individuals (Dupont-Nivet, Vandeputte, Haffray & Chevassus, 2006). Meagre gametes may be collected for artificial reproduction (Poli et al, 2003), particularly when spawning is not occurring naturally (Marino et al., 2000). Semen can be collected by stripping, which might not be effective in every individual (Mylonas et al., 2013b), whereas eggs can also be collected by stripping after hormonal treatment, as seen in Figure 1 (DIVERSIFY, n.d.).

Figure 1: The process of stripping eggs from a female meagre (DIVERSIFY, n.d.).



Sperm and eggs are usually mixed in a certain ratio that maximizes the gamete usage. In general, to ensure genetic variability when doing artificial reproduction, a pool of semen from several males is used (Mylonas, Fostier & Zanuy, 2010). Nevertheless, for selective breeding, the use of only one specific male is also possible.

6. Gamete storage

When conducting artificial reproduction male and female gametes are not always available at the same time. In these cases gametes have to be stored in advance. In this context, gamete storage is an essential tool for artificial fertilization and breeding programs for several reasons. Fertilization can be synchronized according to gametes' availability (Cabrita et al., 2010; Suquet, 2000). Furthermore, it also allows for the gametes' transportation between locations such as collection site and hatchery, making easier and cheaper genetic exchanges (Cabrita et al., 2010; Kopeika et al., 2007; Suquet, 2000). The total volume of the gametes can

be efficiently used, which is particularly important when low volumes are available (Cabrita et al., 2010). Finally, in the case of semen cryopreservation, genetic material of valuable males can be theoretically preserved indefinitely (Cabrita et al., 2010; Suquet, 2000).

After collection, gametes start to suffer degradation processes. Their quality decreases, causing a reduced fertilizing ability. In some species, gametes may be stored unaltered from hours to days, at room temperature (Mylonas et al., 2010). Alternatively, the storage conditions can be manipulated to delay the degradation processes (Mishra, Patra, Dash, Verma & Routray, 2016; Yasui et al., 2015). Some examples of parameters that can be manipulated are temperature, oxygen supply, gas exchange, the storage medium (extender) and its dilution ratio, as well as the replenishment of the extender during storage (Kowalski et al., 2014; Mishra et al., 2016). For both female and male gametes of several fish species there are different storage protocols, which involve the manipulation of one or more of these factors.

Regarding storage temperatures, refrigeration is done above 0°C (Yasui et al., 2015). This can be achieved in a simple way, with the use of a refrigerator or a cooler with ice packs for both oocytes and semen, either undiluted or diluted (Komrakova & Holtz, 2009; Wayman, Tiersch & Thomas, 1998). Depending on the species, the gametes may be refrigerated for several days while keeping some quality.

Alternatively, gametes can be stored at temperatures below 0°C. In fishes, only semen can be stored at subzero temperatures (Asturiano, Cabrita & Horváth, 2016). The larger size of the oocytes, the yolk content and the different permeability of the oocytes' membranes causes some difficulties in their storage, namely the penetration of cryoprotectants and lack of uniformity in the cooling process (Asturiano et al., 2016; Chao & Liao, 2001). On the other hand, it is relatively easier to preserve semen of most fish species, either through short term cooling or long term cryopreservation in liquid nitrogen, at -196°C (Agarwal, 2011; Gwo, Chen & Cheng, 2002; Stoss, 1983). Although cryopreservation allows the cells' preservation for a longer period of time, and several cryopreservation protocols have been developed for different fish species, there are not many cases of practical applications outside of the research environment (Cabrita et al., 2010). Indeed, from the aquaculture perspective semen refrigeration remains the best alternative (Viveiros, Isaú, Figueiredo, Leite & Maria, 2010a).

Extenders are the substances in which gametes are diluted and preserved. Oocytes are generally stored in ovarian fluid (Bellard, 1988), since the usage of dilution media has shown no advantages (Stoss, 1983). In the case of semen storage the usage of extenders is more common. Although it is only essential for cryopreservation, for the incorporation of

cryoprotectants (Agarwal, 2011; Kopeika, Kopeika & Zhang, 2007), they are an important complement for refrigeration as well, extending the spermatozoa's viability (Lahnsteiner, Berger, Weismann & Patzner, 1996; Maria, 2014). An ideal extender, according to Mann (1964), should not activate semen motility, should be isotonic and have a good buffering capacity and should include nutrients, antioxidants, antibacterial substances and stabilizing colloids. Keeping these characteristics in consideration, the extender's composition may vary. A simple and effective extender is a solution of NaCl (Fabbrocini et al., 2000; Maria, Viveiros, Freitas & Oliveira, 2006; Oliveira, Viveiros, Maria, Freitas, & Izaú, 2007). The addition of other substances to this extender may be proven beneficial, such as glucose (Viveiros, Orfão, Maria & Allaman, 2009). He and Woods III (2003) also proved that the addition of some aminoacids to a NaCl solution, such as glycine, improved the quality of stored semen. There are also extenders with more complex compositions, usually mimicking the seminal plasma composition. As an example, for the European sea bass it is usually recommended a non-activating medium, with different salts, glucose and BSA (Fauvel, Suquet, Dreanno, Zonno, & Menu, 1998). Additionally, there are commercially available extenders that may be used, such as BTS™ and ACP® (Nascimento, Maria, Pessoa, Carvalho, & Viveiros, 2010; Viveiros, Maria, Orfão, Carvalho, & Nunes, 2008).

The atmosphere with which the gametes are stored, excluding cryopreservation, affects their quality. Generally, normal air is used, but oxygen can be added (Billard, 1981). During storage, gametes consume oxygen (Robitaille, Mumford & Brown, 1987), and thus aerobic conditions should be provided. For this reason, addition of oxygen to the containers results in a longer storage time (Bellard, 1988; Komrakova & Holtz, 2011) for both oocytes (Billard, 1981) and semen (Bellard, 1988). Nevertheless, the presence of oxygen can also be the cause of lipids' peroxidation, resulting in damage to the cells with deterioration of quality (Chen et al., 2010). Some factors must also be taken in consideration, such as frequency of renewal of atmosphere (Marques & Godinho, 2004), which depends on the amount of oxygen consumed. This varies with the species, semen dilution and ratio of volume of semen to atmosphere. Diluting semen or stocking it with more air allows a reduction in this frequency (Bellard, 1988).

Other aspect to take into consideration in the refrigeration protocols is, in the case of semen, the sedimentation over time, which can be avoided by slightly shaking the containers periodically (Stoss, Büyükhatoğlu & Holtz, 1978). In the case of oocytes their storage is affected by the weight of eggs and the number of layers (Komrakova & Holtz, 2009). In most

cases, these protocols' optimization is done empirically, testing different parameters and evaluating the gametes' quality over time.

Semen storage protocols are far more diverse and more frequently used than oocytes' storage protocols. Artificial reproduction techniques normally involve semen storage until collection of oocytes (Rurangwa, Kime, Ollevier & Nash, 2004; Stoss, 1983). In the case of the Sciaenidae family, there are already semen storage protocols for different species, testing different extenders, osmolalities and dilution ratios (Leclercq et al., 2014; Wayman et al. 1998; Wayman, Thomas & Tiersch, 1997). More recently, research done under the DIVERSIFY-EU project, in 2014, found that in meagre sperm quality decreases rapidly after collection, if not diluted in an extender.

Meagre is a species in the early phases of adaptation to aquaculture, which will benefit from the improvement of artificial reproduction techniques that could assist in the management of broodstocks. Preliminary results regarding the refrigeration of meagre semen, done under the DIVERSIFY-EU project, gave promising results for the development of artificial reproduction. The usage of European sea bass extenders efficiently sustained the spermatozoa's motility for more than 24 hours. However, there are no protocols for semen storage in this species.

7. General considerations on semen and spermatozoa

The ejaculate semen is composed of spermatozoa and seminal plasma. Spermatozoa are the cells responsible to deliver the male genetic information to the egg, and have the capacity to become motile (Miller, Brinkworth & Iles, 2010). The seminal plasma is an enriched medium responsible for maintaining spermatozoa in a quiescent state until ejaculation and providing energy resources (Alavi & Cosson, 2006).

In teleost fish, the spermatozoa structure is divided in head, mid-piece and flagellum (Ginzburg, 1968; Islam & Akhter, 2011). The head is occupied in the most part by the nucleus, containing the genetic material (Ginzburg, 1968; Islam & Akhter, 2011). In the case of the teleosts, the head does not contain an acrosome (Dzyuba & Cosson, 2014), and in meagre it is oval-shaped (Schiaivone et al., 2012). The mid-piece is linked to the head (Ginzburg, 1968; Islam & Akhter, 2011) and contains the mitochondria (Berois et al., 2011; Ginzburg, 1968). The flagellum is the apparatus responsible for the movement (Islam & Akhter, 2011), and originates from the posterior part of the nucleus (Cosson et al., 2008). It produces waves that propagate from proximal to distal tip (Boryshpolets, 2011), resulting in cell motility. ATP is the major source of energy for the flagellar beating, produced by the

mitochondria present in the mid-piece (Boryshpolets, 2011; Dzyuba et al., 2016). Nevertheless, during the fast motility period there is a very high ATP consumption that is not compensated by the mitochondria production (Ingermann, 2008). As a result, the energy stores are depleted during this time leading to the spermatozoa immobilization in a short period (Cosson, 2010; Dreanno, Seguin, Cosson, Suquet & Billard, 1999b). This means that motility depends on the endogenous ATP and other energy molecules' stores present before activation (Cabrita et al., 2010). In most external fertilizing fishes spermatozoa are immotile when inside the testes (Cosson et al., 2008; Ginzburg, 1968). After ejaculation motility is triggered by changes in the osmotic or ionic environment, which in the case of most marine species is caused by seawater (Cosson et al., 2008).

8. Sperm quality and its evaluation

Sperm quality can be defined as the ability to successfully fertilize an egg and develop into a normal embryo (Bobe & Labbé, 2010; Rurangwa et al., 2004). The measurement of different sperm quality parameters has several applications in aquaculture. They can be used as predictors of the fertilizing ability (Bobe & Labbé, 2010; Rurangwa et al., 2004), and thus their evaluation is important to increase artificial reproduction efficiency (Zilli, Schiavone, Zonno, Storelli & Vilella, 2004). When there is a reduced amount of eggs (Rurangwa et al., 2004), the risk of a low fertilization rate and loss of egg batches is prevented (Bobe & Labbé, 2010). Furthermore, cryopreservation should be only done to sperm of proven high quality (Zilli et al., 2004). Sperm quality evaluation also helps to determine the optimal time for semen collection, when its quality is variable along the season (Babiak, Ottesen, Rudolfsen & Johnsen, 2006; Fabbrocini et al., 2000). The occurrence of contamination during collection (Dreanno et al., 1998) and the effect of repeated stripping (Aas, Refstie & Gierde, 1991) may also be assessed. Finally, it helps to improve handling and storage protocols (Fabbrocini et al., 2000; Linhart et al., 2004), as is the case of the present thesis.

Fertilization success is the most integrative parameter and the best indicator of sperm quality (Bobe & Labbé, 2010). Nonetheless, for practical reasons, there is the need to use other parameters, which do not require the usage of eggs. The most commonly used are motility parameters, such as percentage of motile cells and motility duration or spermatozoa's swimming speed (Beirão et al., 2015, 2011b; Schiavone et al., 2012). Other parameters less frequently used are sperm concentration, sperm cell membrane integrity or viability (Beirão, Pérez-Cerezales, Martínez-Páramo & Herráez, 2010; Cabrita et al., 2011), ATP content (Montgomery, Brown, Gendelman, Ota & Clotfelter, 2014), lipid peroxidation (Martínez-

Páramo, Diogo, Beirão, Dinis & Cabrita, 2012) or bacteriology (Jenkins & Tiersch, 1997; Viveiros et al., 2010b), to mention a few.

In most cases it is difficult to correlate one specific parameter with fertilization rate, since they evaluate specific cellular functions, and fertilization depends on many (Bobe & Labbé, 2010). Furthermore, they show the mean value for the whole spermatozoa population, whereas only a small number of spermatozoa is needed to ensure fertilization (Bobe & Labbé, 2010). Sperm quality parameters are, then, partial descriptors of fertilization ability (Bobe & Labbé, 2010). In this context, sperm quality schemes should be built based on several parameters at the same time. These parameters should be investigated and improved, to increase the number of spermatozoa which meet the requirements for fertilization to occur (Bobe & Labbé, 2010).

According to Bobe & Labbé (2010) there are several factors that can influence the sperm quality, most of them related to broodstock management (Billard, Cosson, Perchec & Linhart, 1995). One example is stress, which can be caused by fish handling or environmental conditions. Nutrition is another factor with impact in sperm quality, as feed should have an adequate formula, especially regarding lipid composition and vitamins (Asturiano et al., 2001; Dabrowski & Ciereszko, 1996). Sperm handling during and after collection can also affect its quality, such as adequate storage conditions as temperature, medium and duration (Babiak et al., 2006; Peñaranda, Pérez, Fakriadis, Mylonas & Asturiano, 2008; Suquet et al., 1998).

In the case of meagre, the sperm quality has already been evaluated for concentration, for aspects of motility such as percentage of progressive motile cells and duration of progressive forward movement and morphology (Mylonas et al., 2013b; Mylonas et al., 2016; Schiavone et al., 2012). While Mylonas et al. (2013b) state that sperm quality does not significantly change during the reproductive season., Schiavone et al. (2012) verified a variation during the reproductive season, which goes in accordance to what occurs in other species, and may be due to the aging of spermatozoa (Babiak et al., 2006; Peñaranda et al., 2008).

9. Sperm quality parameters

9.1. Concentration

Sperm concentration is usually presented as the number of cells per milliliter (Junior et al., 2008; Wirtz & Steinmann, 2006). In teleost semen the cell concentration is highly variable, due to factors such as species, individual and time in the reproductive season (Butts, Litvak & Trippel, 2010; Piironen & Hyvärinen, 1983). In several species studied by Piironen and

Hyvärinen (1983) and Poole and Diilane (1998), the concentration ranged from 2.2 to 127.4 x 10⁹ ml⁻¹. Results obtained by Mylonas et al. (2013b) in meagre semen obtained a mean cell concentration between 19 and 32 x 10⁹ ml⁻¹. This parameter can be correlated, in some instances, with other parameters such as fertilization rate and motility (Ciereszko & Dabrowski, 1994; Hwang & Idler, 1969). Nevertheless, its main importance involves the optimization of the usage of semen when doing artificial reproduction, and the sperm to egg ratio needs to be adjusted (Bombardelli, Mörschbacher, Campagnolo & Syperreck, 2006; Sanches et al., 2011).

Several methods can be used to assess sperm concentration of fish semen. The most common involves counting the number of spermatozoa with a microscope and a cell counting chamber, such as a Neubauer chamber (Junior et al., 2008; Wirtz & Steinmann, 2006). These methods, despite cheaper, are time consuming (Fauvel, Suquet & Cosson, 2010). Spermocrit determination, with hematocrit capillary tubes, is faster and simpler, giving a result in percentage equivalent to sperm concentration (Ciereszko & Dabrowski, 1993). However, this technique is not as effective in marine fish because spermatozoa do not sediment efficiently, which may be due to their density being similar to that of seminal plasma (Fauvel et al., 2010). Alternatively, the value of absorbance, obtained by spectrophotometry, can be used if a correlation with concentration has been previously established (Ciereszko & Dabrowski, 1993). Furthermore, some authors also suggest the use of flow cytometry or coulter counter, which are highly precise but expensive methods (Fauvel et al., 2010). Lastly, a modern image analysis software connected to a microscope is also frequently used to determine concentration fast and efficiently (Fauvel et al., 2010). Depending on the species, the correlation coefficient between these three methods may be high (Ciereszko & Dabrowski, 1993).

9.2. Motility

Sperm motility is an integrative parameter that combines different cellular structures (Bobe & Labbé, 2010). It is the most commonly used parameter to compare different experimental conditions such as sperm collection procedures, semen extenders, or sperm storage conditions (Bobe & Labbé, 2010).

In general, the initiation, duration and the patterns of motility differ between species (Rurangwa et al., 2004). In teleosts with external fertilization, such as meagre, the motility tends to be brief (Coward, Bromage, Hibbitt & Parrington, 2002) and spermatozoa usually swim at high speed and frequency, in rectilinear motion (Cosson et al., 2008; Cosson, 2010).

In meagre the percentage of motile spermatozoa varies between 53% and 74%, with a duration of 34 to 80 seconds (Schiavone et al., 2012). In most species, fertilization usually occurs between 5 and 20 seconds after spermatozoa's activation, thus the most useful motility data is obtained during this period (Kime et al., 2001).

In several species a correlation between motility and ability to fertilize the eggs has been found (e.g. Cosson et al., 2008; Ottesen, Babiak & Dahle, 2009), including refrigerated and cryopreserved sperm (e.g. Beirão et al., 2011a; Ciereszko & Dabrowski, 1994). Some examples are the correlations between sperm motility parameters and fertilization or hatching rate, in cryopreserved semen in African catfish (*Clarias glariepinus*) (Rurangwa et al. 2001), gilthead seabream (Beirão et al. 2011a), common carp (*Cyprinus carpio*) (Leveroni, Calvi, Zoccarato, Gasco & Andrione, 1993; Leveroni, Calvi, Zoccarato, Gasco & Andrione, 1994) and *Alburnus alburnus* (Lahnsteiner et al., 1996), or in refrigerated rainbow trout (*Oncorhynchus mykiss*) semen (Ciereszko & Dabrowski, 1994).

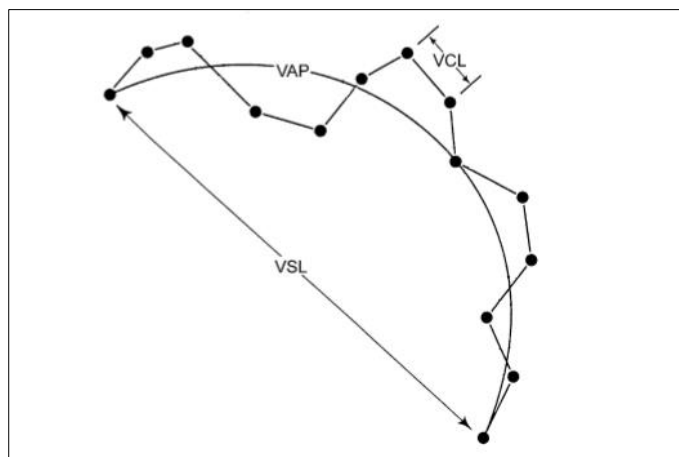
Motility can be assessed subjectively, by direct observation on a microscope (Kime et al., 2001). In these cases, the researcher measures the motility duration and estimates the percentage of motile sperm (Fauvel et al., 2010). To increase the accuracy of these measurements, sperm motility may be videotaped and posteriorly analyzed by different researchers (Kime et al., 2001). Alternatively, nowadays, more laboratories are using computer-assisted sperm analysis (CASA) systems (Fauvel et al., 2010; Kime et al., 2001) to objectively assess motility. The CASA systems analyze sperm movement of every single spermatozoa in a short video file, which is recorded with a camera attached to a microscope and connected to a computer (Rurangwa et al., 2004). This system measures several motility and velocity parameters (Fauvel et al., 2010). The parameters most commonly used for fish sperm analysis are described in Table 1. The differences between the three velocity parameters are represented in Figure 2. The large data sets obtained can be used to compare different sperm samples and thus reveal differences in experimental protocols, individuals and species, to mention a few (Linhart, Rodina, Gela, Kocour & Vandeputte, 2005; Rurangwa et al., 2001).

Table 1: Sperm motility parameters, their abbreviation, unit, and description (Rurangwa et al., 2004).

PARAMETER	DESCRIPTION
(unit)	
Duration of Motility	
(s)	time interval in which the cells remain motile
MOT • Percentage of Motile Cells	
(%)	percentage of cells that exhibit some motility

PROG • Percentage of Progressive Spermatozoa	(%)	percentage of cells with forward progressive movement
MOC • Concentration of Motile Cells	(spz/ml)	concentration of cells that exhibit some motility
VCL • Curvilinear Velocity	($\mu\text{m/s}$)	considering the entire distance travelled
VSL • Straight Line Velocity	($\mu\text{m/s}$)	considering a straight line between initial and final point
VAP • Average Path Velocity	($\mu\text{m/s}$)	considering the distance travelled as a smoothed path
LIN • Linearity	(%)	obtained by VSL/VAP • a measure of path curvature
STR • Straightness	(%)	obtained by VAP/VCL • a measure of side to side movement

Figure 2: Schematic representation of some motility patterns measured by a CASA system. VCL, curvilinear velocity, in the entire distance travelled; VSL, velocity in a straight line; VAP, velocity of an average path. Black circles represent the position of the cell in each frame. Adapted from Rurangwa et al. (2004).



9.3. Viability

Sperm viability refers to the percentage of live spermatozoa (Garner & Johnson, 1995). Although viability should be attributed to the capacity of sperm to fertilize an egg, this parameter evaluates the percentage of spermatozoa with an integral plasma membrane (Fauvel et al., 2010; Rurangwa et al., 2004).

Different authors have found a correlation between sperm viability and other semen quality parameters, such as sperm velocity parameters in common carp semen and hatching rate in African catfish (Linhart et al., 2005; Rurangwa et al., 2001).

Viability tests are based on staining protocols, which confer different coloration to viable and dead cells (Rurangwa et al., 2004). Each stain has a different cell permeation ability,

depending on the plasma membrane's integrity (Fauvel et al., 2010). Protocols may only use one dye, such as trypan blue (Lubzens et al., 1997; Rurangwa, Biegniewska, Slominska, Skorkowski & Ollevier, 2002), or a combination of two, such as eosin and nigrosine (Maria et al., 2006). Alternatively, fluorescent dyes can also be used, to facilitate cell distinction. A commonly used combination of these is propidium iodide (PI) with SYBR-14, for dead and viable cells respectively (Rurangwa et al., 2004), but other combinations have been proposed, such as PI with YO-PRO 1 (Beirão et al., 2010). These stains usually target nucleic acids of the spermatozoa. In the case of the PI/SYBR-14 combination, PI cannot cross the intact plasma membrane, whereas SYBR-14 leaks out when the plasma membrane is deteriorated (Nagy, Jansen, Topper & Gadella, 2003; Rurangwa et al., 2004). The percentage of viable cells can be assessed under a microscope, by direct counting or using an image analysis system (De Baulny, Labbé & Maise, 1999). Other ways to do so are through flow cytometry (Beirão et al., 2010), with an automatic counter (Fauvel et al., 2010) or a fluorometer (McNiven, Gallant & Richardson, 1993).

9.4. ATP

ATP, together with adenosine diphosphate (ADP) and adenosine monophosphate (AMP), compose the Adenylate Energy Charge (AEC). The AEC can be considered the intracellular content of available energy, for it is composed of high energy nucleotides. Since the values of ATP and AEC are highly correlated, it is reasonable to evaluate just one of these parameters (Fauvel et al., 2010), such as ATP, which has simple and precise protocols (Perchec, Jeulin, Cosson, Andre & Billard, 1995). As previously mentioned, the ATP present in the spermatozoa is the major energy source for flagellar beating and, as demonstrated by Dreanno et al. (1999b) and Cabrita et al. (2005), it can be used as a sperm quality estimator. Indeed, a positive correlation between ATP content and percentage of motile cells was found in some species such as turbot (De Baulny et al., 1996) and steelhead trout (*Oncorhynchus mykiss*) (Bencic, Krisfalusi, Cloud & Ingermann, 1999b). Furthermore, Bencic et al. (1999b) also observed a positive correlation between ATP and fertility rate in steelhead trout.

ATP determination can be done with a bioluminescence technique that uses firefly luciferin-luciferase, to which there are commercial assay kits (Fauvel et al., 2010; Perchec et al., 2005). The reaction between ATP, luciferin and luciferase leads to the production of luminescence, which can be detected with a luminometer or scintillation counter (Rieger, 1997; Yang, Ho, Chen & Hu, 2002). There are also methods which enable the quantification of not only ATP but also the other components of AEC, such as high pressure liquid chromatography

(Dziewulska, Rzemieniecki & Domagala, 2010; Mukai & Okuno, 2004) and nuclear magnetic resonance spectroscopy (van den Thillart, van Waarde, Muller, Erkelens & Lugtenburg, 1990).

9.5. Lipid Peroxidation

During the aerobic metabolic processes, cells produce reactive oxygen species (ROS). These have cytotoxic properties, causing damages to the desoxyribonucleic acid (Cabrita et al., 2011; Potts, Notarianni & Jefferies, 2000), ATP depletion (Aramli, Kalbassi, Nazari & Aramli, 2013), abnormalities in morphology (Ball, 2008), and lipid peroxidation (Zhou et al., 2006), which leads to a decrease in motility and cell viability (Aramli et al., 2013; Ball, 2008), and a decrease in sperm quality (Apel & Hirt, 2004). The cell membrane's high content in polyunsaturated fatty acids (PUFAs) makes fish spermatozoa quite susceptible to lipid peroxidation (Labbe et al., 1995; Zhou et al., 2006). To counteract the ROS effects semen has antioxidant mechanisms (Lahnsteiner & Mansour, 2010), which include different antioxidant substances present in the seminal plasma (Cabrita et al., 2011; Potts et al., 2000). However, *in vitro* storage of the semen increases ROS formation (Ball, 2008; Zhou et al., 2006) namely because dilution in the extenders reduces the antioxidants' concentration of seminal plasma (Cabrita et al., 2011; Martínez-Páramo, Martínez-Pastor, Martínez-Rodríguez, Herráez & Cabrita, 2009). Nevertheless, antioxidants could be added to stored semen to prevent cellular injuries caused by ROS (Aramli et al., 2013).

During lipid peroxidation, several by-products are formed. Their measure is frequently used to quantify lipid peroxidation (Devasagayam, Bolor & Ramasama, 2003). The thiobarbituric acid (TBA) assay is the most used to evaluate lipid peroxidation *in vitro* (Buege & Aust, 1978; Devasagayam et al., 2003). It measures the by-products which react with thiobarbituric acid – thiobarbituric acid reactive substances (TBARS) -, such as malondialdehyde (MDA) (Potts et al., 2000; Devasagayam et al., 2003; Yahyavi, Kaykhaii & Hashemi, 2016). This assay is sensitive but non-specific (Devasagayam et al., 2003; Zhou et al., 2006), since TBA might react with other organic components. Alternatively, MDA can be measured using commercially available kits (Beirão et al., 2015; Martínez-Páramo et al., 2012).

9.6. Bacteriology

Bacterial growth can be the cause of sperm quality decrease especially when semen is stored in refrigeration. Bacteria may be present in semen as part of the normal flora, due to contamination or with origin in the extender (Bartha, 2009). They can reduce sperm quality

due to the oxygen consumption, and the production of extracellular enzymes and metabolic by-products that affect spermatozoa (Jenkins & Tiersch, 1997). The growth of bacteria has been associated to decreased fertility (Saad, Billard, Theron & Hollebecq, 1988; Stoss & Refstie, 1983), low motility (Christensen & Tiersch, 1996; Isaú, 2014) and viability (Stoss et al., 1978), as well as increased morphological changes (Jenkins & Tiersch, 1997). Indeed, Jenkins & Tiersch (1997) observed that the usage of a sterile extender, as opposed to a non-sterile one, extended the number of days in which there was sperm motility. Moreover, some authors have suggested the use of antibiotics to inhibit bacterial growth and improve sperm survival and fertility (Billard et al., 2004; Viveiros et al., 2010b).

Bacteriological evaluation can be done quantitatively or qualitatively. The quantitative analysis is the estimation of the number of bacteria per unit of volume by counting the number of colony-forming units (CFUs) (Elain et al., 2015; Jenkins & Tiersch, 1997). The qualitative analysis involves determining the presence of bacteria and their identification (Plusquellec, Beucher, Le Lay, Le Gal & Cleret, 1991) by means of various techniques that evaluate different phenotypical characteristics of the bacteria. Examples of these are observation of morphology and motility, Gram staining, catalase or oxidase test, and glucose fermentation (Salanitro, Blake & Muirhed, 1977).

9.7. Fertilization success

Fertilization success is the most integrative estimator of sperm quality, since it requires the interaction of all sperm cellular functions, rather than one or some of them (Bobe & Labbé, 2010). In fact, in numerous occasions there is no correlation between other sperm quality parameters, such as motility parameters or viability, and fertilization (Bobe & Labbé, 2010; Cabrita et al., 2010). It should also be kept in mind that only a small fraction of spermatozoa are needed to fertilize the eggs, whereas most of the sperm quality parameters take into account the average of overall sperm population (Bobe & Labbé, 2010).

In general, to evaluate sperm fertilization success, sperm and eggs are mixed at a certain ratio to observe the percentage of ova fertilized per amount of sperm cells (Beirão et al. 2011a). Usually, only the amount of fertilized eggs is evaluated, without contemplating further developmental success (Shields, Brown & Bromage, 1997). This is easy to observe in species with transparent eggs, such as meagre (Gamsiz & Neke, 2008). Nonetheless, the hatching rate might also be assessed, incubating the eggs (Cabrita et al., 2010). It should be taken in consideration that there may be significant differences between fertility and hatching rate

(Cabrita et al., 2009; Pérez-Cereales et al., 2010). Furthermore, the impact of egg quality must also be taken in consideration when evaluating the results (Bobe & Labbé, 2010).

EXPERIMENTAL STUDY

1. Objectives

The primary objective of this study was to develop a protocol for the refrigeration of meagre semen that facilitates artificial reproduction techniques both for research and for meagre production. The secondary objective was to contribute for the understanding of the causes of the fish sperm quality degradation during refrigeration. For this, meagre sperm was refrigerated using three extenders in three different dilutions, taking in consideration the differences in the semen quality parameters along the time. Semen quality was analyzed in terms of sperm motility, viability, ATP content, lipid peroxidation and bacteriology.

2. Material and methods

2.1. Experimental animals

The meagre broodstock used in this experiment was kept in two 18 m³ outdoor in the Aquaculture Research Station of the Portuguese Institute for the Ocean and Atmosphere (EPPO-IPMA). Each tank had 9 individuals born in the research station, with a mean weight of $7\ 315 \pm 750$ g and length of 91.8 ± 4.4 cm. Most of the individuals were not previously sexed, but an initial screening showed that the ratio was of two males per female. The tanks were covered with shade nets and exposed to natural photoperiod and temperature conditions. Tanks were supplied with filtered seawater from the nearby estuary, pumped at a continuous rate of 1 L/s. The water salinity changed according to the natural tidal cycles. The mean water temperature was 21.9°C (maximum of 26.9°C and minimum of 15.6°C), and the mean dissolved oxygen concentration was 6.3 mg/L (maximum of 9.3 and minimum of 2.2). Fish were fed *ad libitum* with commercial pellets twice a week.

2.2. Semen collection

Semen was collected in seven different occasions during meagre reproductive season, between April and June. Sampling took place between 10 am and 12 pm, once every 6 to 8 days. One tank was used in each day of sampling, alternating between them. This way fish in each tank had at least 12 days to recover from sampling.

In the sampling days, the water level of the tank was lowered to about 30 cm. Before semen collection, fish were mildly sedated with 50 ppm of 2-phenoxyethanol directly added to the tank and a net was placed to separate fish already sampled, as shown in Figure 3.

Figure 3: Image of one tank during semen collection, showing the lowered water level and the net separating fish already sampled.



Fish were caught individually and placed onto a 200 liters tank shown in Figure 4, with 300 ppm of 2-phenoxyethanol (Neiffer and Stamper, 2009). Meagre were kept there until they showed signs of anesthesia.

Figure 4: Two meagre on the 200 liters tank with anesthesia, showing loss of equilibrium, by being upside down.



Besides the loss of equilibrium, which causes fish to turn upside down (Fig. 4), fish were also tested for a lack of response to stimuli, grasping the base of the tail (Neiffer and Stamper, 2009). Before removing the fish from the anesthesia tank, they were gently pressed near the

genital papilla to observe whether they were spermiating or not. Males that were not spermiating were returned to the main tank. Spermiating males were enveloped in a wet towel, and placed in dorsal decubitus in an acrylic cylinder half with some water on its bottom (Fig. 5).

Figure 5: Moment of semen collection, with the meagre upside down and enveloped in a wet towel, in the acrylic cylinder half. Abdominal pressure was being applied and semen was collected with a syringe.



Since the sperm sampling procedure took less than 2 minutes, it could be performed out of the water, without the need of an artificial ventilation system (Harms and Bakal, 1995). The semen was obtained by applying pressure abdominally in a craniocaudal motion. Semen was collected with a 5 ml sterile syringe and placed in 1.5 ml microtubes kept in a styrofoam box with ice packs. The genital papilla was frequently wiped with tissue paper to avoid contamination, particularly before applying abdominal pressure and after the discharge of urine or feces. In the event of contamination the syringe and its contents were discarded.

2.3. Refrigeration and samples' processing

Samples with no signs of contamination and a percentage of motile spermatozoa over 70% were pooled in a sterilized 15 ml falcon. Each pool had semen of 3 to 6 males. A total of 7 pools were used in this study.

Subsamples of 1.5 to 2.5 ml of each pool were placed in sterilized and labelled 15 ml falcons, and combined with one of three different extenders, in one of three different dilutions, in a full factorial design. A control sample, with 1.5 to 2.5 ml of undiluted sperm, was also kept. A total of 10 treatments were tested (3 extenders \times 3 dilutions + 1 control). Dilutions were

selected after a preliminary test. The dilutions used were (in ratio of volume semen:extender) 1:4, 1:9 and 1:19.

The extenders' composition was the following:

- Extender NAM, Non-activating medium: NaCl 7.5 mg/mL, KCl 0.11 mg/mL, MgCl₂ 1.23 mg/mL, CaCl₂ 0.39 mg/mL, NaHCO₃ 1.68 mg/mL, Glucose 0.08 mg/mL, BSA 10 mg/mL, diluted in distilled water (Fauvel et al., 1998);
- Extender NC: 0.9% NaCl;
- Extender NCG: 0.9% NaCl, Glycine 0.0038 mg/mL, Glucose 0.001 mg/mL (He and Woods III, 2003).

The osmolality was adjusted to 300 mOsm, according to the observed value of meagre's seminal plasma in a preliminary experiment. The pH of the extenders was adjusted according to the pH of the NAM, which was 7.7 (Fauvel et al., 1998). NAM was chosen because it is the extender used for European sea bass sperm (Fauvel et al., 1998), and there are some resemblances between European sea bass and meagre sperm. NC was designed to be a simple and inexpensive extender. Finally, the extender NCG was based on NC, with the addition of glycine that, according to the results obtained by He and Woods III (2003), improves the quality of striped bass' (*Morone saxatilis*) stored semen. All the extenders were autoclaved before mixing with the semen samples. In the case of NAM, BSA was only added after this procedure, to avoid protein denaturation.

The falcons were kept in a refrigerator at 4°C (Fig. 6). They were opened daily for gas exchanges and gently shaken to avoid formation of cell deposit. There was a temperature sensor in the refrigerator to control the daily maximum and minimum temperature it reached (Annex A, Table 13).

Figure 6: The refrigerator shelf. In the front, the falcons with the various samples from three separated pools. On the left, bottles with the extenders and reagents for the ATP protocol. On the right, a falcon filled with water and with the temperature sensor.



All the samples were analyzed for concentration. Due to the limited samples' volume, not all analysis were conducted in all pools. The number of pools evaluated, as well as the frequency of evaluation of each quality parameter, is shown in table 2. The first 5 pools were analyzed for sperm motility and viability within one hour after collection (considered 0 h data point) in the control sample. Motility and viability were evaluated in all the treatments at 12, 24, 36, 48 h, and every 24 h after that. The motility was measured until the percentage of motile cells was zero, or until 15 days. The viability was measured until 7 days or after observing zero percent of motility. Pools 1 to 4 were also analyzed for ATP within one hour after collection (considered 0 h data point) in the control sample, and in all treatments at 12 and 24 h, and every 48 h, for as long as viability was evaluated. MDA was evaluated in the last two pools (6 and 7). It was measured on day 0 in the control and at days 3, 6 and 9 in all the treatments. Bacteriology was analyzed in 3 pools (5, 6 and 7). In day 0 only the control was inoculated, whereas the control, the dilutions 1:4 from each extender, and the three extenders as negative control, were inoculated at days 5 and 10. It was not logistically feasible to inoculate the plates for the evaluation of all the 10 treatments. Therefore, only the 1:4 dilution treatments were assayed for bacteriology, as representatives of each extender, these had the highest semen concentration, and thus the highest concentration of bacteria from semen. To allow for comparisons, sperm motility and viability was measured on the same days as MDA and inoculations.

Table 2: Number of pools and frequency of evaluation of each sperm quality parameter.

	MOTILITY	VIABILITY	ATP	MDA	BACTERIOLOGY
NUMBER OF POOLS EVALUATED	5	5	5	2	3
FREQUENCY OF EVALUATION	Daily	Daily	Days 0, 0.5, 1, 3, 5	Days 0, 3, 6, 9	Days 0, 5, 10

2.4. Concentration and motility

Sperm concentration and motility were evaluated using the CASA software ISAS® by PROiSER R+D and a UOP UB202i microscope by Prymus© with a 10x lens, attached to a Basler© camera connected to a laptop with the software.

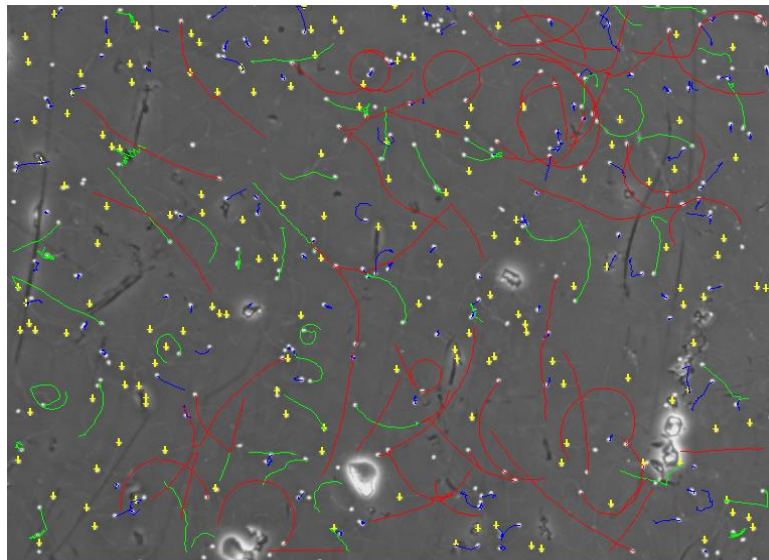
For concentration, the samples were diluted between 10 to 100 times in 0.9% NaCl dependent on the treatment dilution. This was repeated six times for each treatment of each pool.

For motility evaluation, the sperm was activated by adding 1.5 μl of the semen (volume adjusted according to treatment dilution) in a Makler chamber with 20 μl of the research station seawater. Motility was recorded 15 seconds after activation.

The following parameters were analyzed: MOT, PROG, VCL, VAP, VSL, LIN and STR. The software settings were adjusted to meagre sperm: 50 frames captured per second; scale was calibrated at 10x; particle area (μm^2) between 1 and 70; connectivity, 14; Progressivity: 80% of the STR and; minimum number of images to calculate ALH: 10. An example of an image obtained during this evaluation is in Figure 7.

Each reading was repeated three times.

Figure 7: Image obtained with the CASA software. Different colours represent spermatozoa with different motility speeds. Yellow dots represent immotile spermatozoa, whereas, blue, green and red represent slow, medium and fast spermatozoa respectively, and their paths.



2.6. Viability

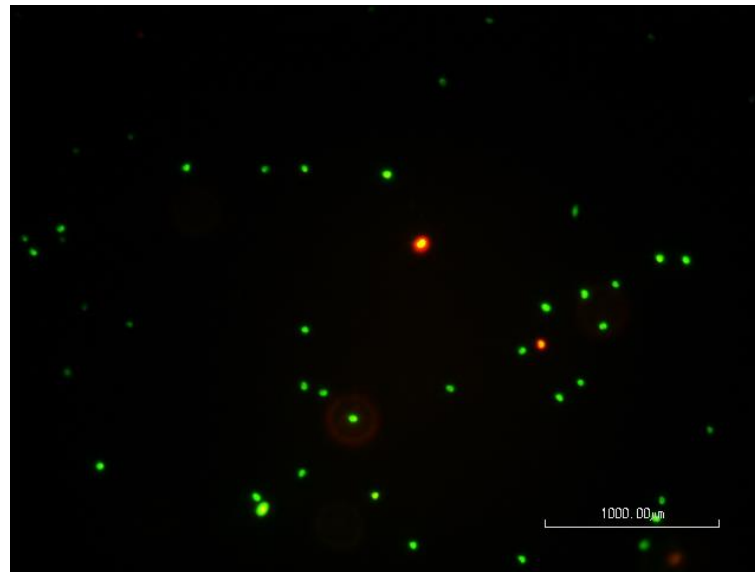
Sperm viability was measured with the double stain PI and SYBR-14 from the kit Live/Dead® Sperm Viability Kit from Invitrogen™. The cells were observed under a Nikon® Eclipse Ci fluorescence microscope with 10x magnification lens with a Nikon® LH-M100C-1 100w Mercury Lamphouse.

One hundred μl of 0.9% NaCl were added to different volumes of sperm according to the different treatments dilution (control and 1:4 dilutions, 0.5 μl of sperm; 1:9 dilutions, 1 μl of sperm and; 1:19 dilutions, 1.5 μl of sperm). The cell suspensions were then stained with 0.5 μl pre-diluted SYBR-14 (1:9 in 0.9% NaCl) and 0.75 μl of PI (stock solution). After 10 minutes of incubation in the dark and on ice, the suspension was slightly mixed, and a droplet

placed between a slide and cover slip. The preparation was immediately observed under the fluorescence microscope in a dimly lit room. Different fields were observed and at least 200 cells were counted. Cells were differentiated between viable cells (marked green with SYBR-14) and damaged or dead cells (marked red with PI) as observed in Figure 8, and the percentage of viable cells was recorded.

This process was repeated three times per sample.

Figure 8: Image taken in the fluorescence microscope of a sample observed after incubation, showing the difference between cells coloured green (viable cells) and red (dead cells).



2.7. ATP

The following solutions were prepared in advance of the ATP extractions: Sorensen stock solutions A and B, and ethylenediaminetetraacetic acid (EDTA) solution (composition in Annex B, Table 14). The three solutions were autoclaved and stored refrigerated for 1 month maximum. Sorensen buffer and the 2% TCA solution (composition in Annex B, Table 14) were freshly prepared every day.

For the ATP extraction, 10 μl of each treatment were mixed with 150 μl of 2% TCA solution in microtubes kept on ice, and left for 15 min. Subsequently, the microtubes were vortexed for some seconds, and centrifuged at 4°C, 10 000 g, for 10 min. One hundred μl of the resulting supernatant was diluted in 500 μl of Sorensen buffer and stored at -80°C until further analysis. ATP extracts were analyzed with the Adenosine 5'-triphosphate (ATP) Bioluminescent Assay Kit from Sigma-Aldrich®, following the manufacturer's instructions with some adjustments to sperm samples. The kit ATP mix, buffer and ATP standard were divided in aliquots and frozen at -80°C, in advance. The ATP mix was kept from the light, wrapped in aluminum foil. These aliquots were thawed before usage and discarded afterwards.

ATP extracts and reactives were kept on ice during all the procedure. For the ATP curve, serial dilutions of 1 $\mu\text{mol/ml}$ ATP standard were done with HPLC grade H_2O (Table 3).

The curve was kept for a maximum of 8 hours. A working ATP mix solution was prepared by mixing 1.2 ml of ATP buffer with 50 μl of ATP mix.

Table 3: Curve points used in the ATP protocol.

CURVE POINTS	1	2	3	4	5	6	7
ATP (nmol/ml)	25	6.25	1.25	0.25	0.0625	0.0125	0.0025

The luminescence was measured in a BioTek's Synergy 4TM multi-mode microplate reader, using black 96-well microwell plates. For each plate reading a new curve and two blanks (HPLC grade H_2O) were prepared. One hundred μl of the working ATP mix solution was added to the plate wells and left to rest for 5 minutes at room temperature in the dark. After this time, 100 μl of the curve points, blanks and extracts were added to the respective wells. The plate was automatically shaken for 3 minutes and read for luminescence. Because strong luminescence can affect the results in adjacent wells, one well of interval was left between all wells with a sample. Each sample was analyzed in duplicate in different plates.

2.8. MDA

To measure the MDA a spectrophotometric assay kit, BIOXYTECH® MDA-586TM (OxisResearchTM), was used following the protocol developed for fish sperm by Martínez-Páramo et al. (2012). From each treatment 50 μl were used, that were kept in ice all the time. In the case of the control, this volume was diluted 1:1 with 0.9% NaCl. For the cell suspension from treatments NAM and NCG the extender was replaced by 0.9% NaCl after centrifugation at 13 000 g and 4°C for 10 min. All the cell suspensions were briefly vortexed, to resuspend the sperm cells, after which 450 μl of 0.9% NaCl were added to all of them, followed by 5 μl of a solution with 200 μM of sodium ascorbate and 40 μM of iron sulfide. This step fragments the existing lipid peroxides and generates MDA, which will be measured (Annerén, & Epstein, 1987). Then, the cell suspensions were slightly shaken and incubated in a dry bath for 30 minutes at 37°C, in the dark. Meanwhile, a calibration curve was prepared (Table 4), by diluting MDA standard solution (20 μM) with bidistilled water.

Table 4: Curve points used in the MDA protocol.

CURVE POINTS	1	2	3	4	5	6
MDA (μM)	10	8	4	2	1	0

After the incubation time, 100 μl of each cell suspension and curve points were transferred to another microtube, each with 320 μl of R1 solution (N-methyl-2-phenylindole, in acetonitrile). In addition, a control of turbidity was prepared with 100 μl from a previously prepared cell suspension. To prevent further lipid peroxidation in this control 320 μl of acetonitrile solution were added (Botsoglou, 1994). Five μl of Probucol were then added to all the microtubes (cell suspensions, control and curve points). This minimizes the reaction with 4-hydroxyalkenals, although they do not produce significant colour under the conditions of the assay (*OxisResearch*TM, 2001). Subsequently, all microtubes were briefly vortexed and 75 μl of R2 reagent (hydrochloric acid) were added in each one, and then incubated at 45°C in a dry bath, in the dark, for 60 minutes. Afterwards, the microtubes were centrifuged at 10 000 g and 4°C for 10 minutes. Two hundred μl of supernatant from each microtube were placed in well of a 96-well microplate, in duplicate for each cell suspension, curve point, blank and control. Absorbance was read at 586nm in a BioTek's Synergy 4TM multi-mode microplate reader immediately, as acetonitrile may damage the bottom of the multiwell plate.

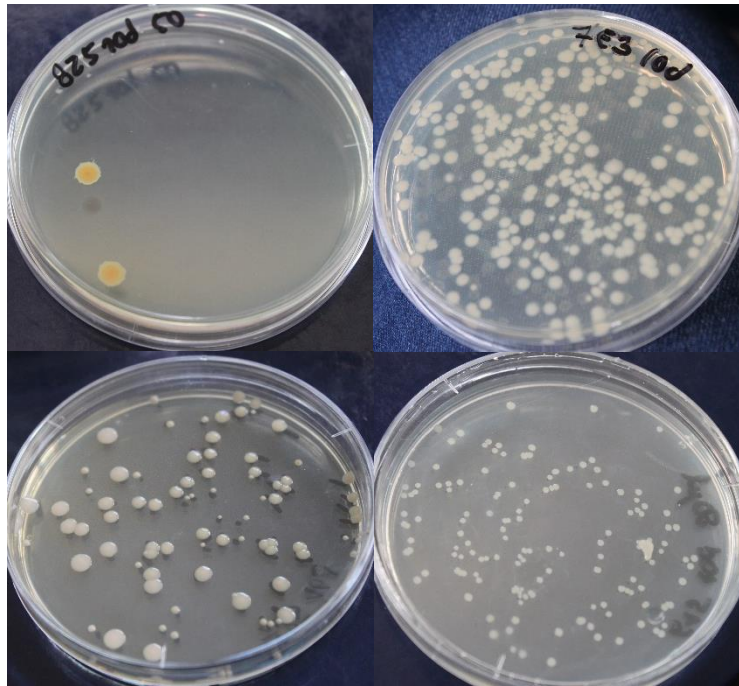
2.9. Bacteriology

For each sample, three plates with Trypticase Soya Agar (TSA), from Sigma-Aldrich®, and three with Pseudomonas CN Agar (PS), from Merck Millipore®, were inoculated by spread plate technique. TSA was chosen as a general nutrient medium for the growth of bacteria, also used by Viveiros et al. (2010s) and Jenkins and Tiersch (1997). PS, a medium selective for *Pseudomonas aeruginosa*, was chosen because *Pseudomonas* sp. are prevalent not only in semen but also in the gastrointestinal tract, water and soil (Boonthai et al., 2016; Jenkins & Tiersch, 1997; Nayak, 2010; Sader & Jones, 2005). Fifty μl of the corresponding treatment were used per plate. In the case of the control, 100 μl were used per plate after pre-dilution 1:9 in sterilized 0.9% NaCl. The incubations were conducted at 24°C. Incubators were weekly disinfected with 70% ethanol to prevent cross contamination.

A potential fungal presence suspected by macroscopic observation was confirmed microscopically by wet mount of a portion of colony. The number of colony forming units (CFUs) of bacteria (Fig. 9) were counted on the 2nd and 7th days of incubation. The last count was the one used for data analysis.

Representative colonies of plates from the same medium, pool, treatment and day of inoculation, were selected based on phenotypical differences. They were photographed when observed and isolated on the 7th day of incubation of their respective plate. Isolation was done with an inoculating loop, touching the colony (preferably isolated from others, and not old) and streaking onto a plate with the TSA medium, which was incubated again for 2 to 7 days at 24°C, in order to purify the colonies.

Figure 9: Plates with apparent bacterial growth after 7 days of incubation.



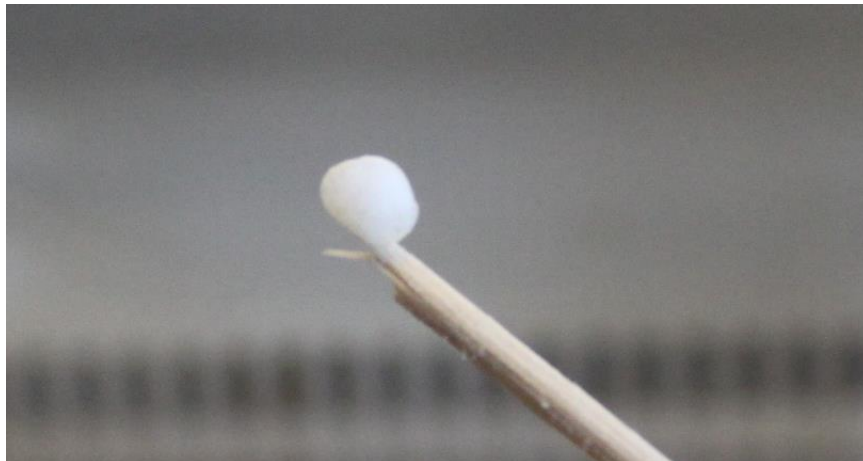
So as to identify the genus of the bacteria, several biochemical identification tests were done in fresh colonies, with 2 days, as described by Holt (1994) and Buller (2004). The following tests were made:

- Morphology and mobility assessment (Bryant and Small, 1956): a portion of colony was mixed with 0.9 ml of sterilized 0.9% NaCl. A drop of this wet mount was placed between slide and cover slip, and observed in a Nikon® Eclipse Ci microscope with a 100x oil-immersion lens. Bacteria were evaluated as to morphology and motility.
- Gram test with potassium hydroxide (KOH) (Buck, 1982): A drop of 3% KOH was placed on a sterile surface. The colony was touched with a sterile toothpick, and dipped repeatedly in 3% KOH. If a viscous string was formed between the drop and toothpick, the bacteria was Gram negative. If not, the bacteria was Gram positive.
- Oxidase test (Ishikawa, Nakajima, Yanagi, Yamamoto & Yamasato, 2003): A portion of the colony was touched with a sterile toothpick and streaked in an oxidase test strip

from bioMérieux ©. A color change to purple or deep blue indicated that the colony was oxidase positive.

- Catalase test (Chang, Kim & Shin, 1997): A portion of the colony was touched with a sterile toothpick and in it was placed a drop of 3% hydrogen peroxide solution. The formation of gas bubbles (Fig. 10) indicated a positive result, meaning the colony had the enzyme catalase.

Figure 10: Positive result of a catalase test, with the formation of gas bubbles.



- Oxidative/fermentation glucose test (Hugh and Leifson, 1953): The colony was mixed in two tubes with a liquid medium of Oxidative Fermentative Test Nutrient Agar of Sigma-Aldrich®. One tube was covered by paraffin, to differentiate growth on anaerobic and aerobic conditions. Both tubes were incubated for 2 days. If the anaerobic or both tubes turned yellow, the result was fermentation of glucose. If only the aerobic tube turned yellow, the result was oxidation of glucose. If the aerobic tube turned blue, then it did not metabolize glucose, and oxidized peptones.
- Growth in mediums with different percentages of salinity: Each bacteria was incubated in non-selective mediums with 0, 3, 5 and 10% of salinity. After 5 days it was determined if the bacteria grew or not in each percentage of salinity.

2.10. Data analysis

The statistical analysis was conducted with the IBM SPSS® Statistics 20 for the correlation analysis, generalized linear models (GzLM) and multiple comparisons and with the R software (version 3.3.1) and the packages Rcmdr (Fox, 2005) and RcmdrPlugin.survival (Fox & Sá Carvalho, 2012) for the survival analysis.

The concentration values of each pool were determined from control sample. The mean and standard error of the mean were calculated for the total number of pools ($n = 7$).

The values of the triplicates done for sperm motility were averaged by pool, treatment and day. In time zero only the control was tested, and the value obtained was assumed to be the time zero value of the other treatments from the same pool. No value of VCL, VAP, VSL, LIN or STR was attributed when motility was 0%. The values of each parameter (MOT, PROG, VCL, VAP, VSL, LIN and STR) were tested for normality with a Shapiro-Wilk test (Annex C, Table 15). Since the data was not normally distributed, they were tested for correlations with a Spearman's rank correlation coefficient. Some parameters were discarded from further analysis when they had similar biological meaning and were significantly correlated. Significant differences between treatments for MOT, VCL and LIN, from the five initial pools for the first seven days were detected with a GzLM because the data was not normally distributed (Annex C, Table 16). Both extender and dilution were considered as factors. MOT, VCL and LIN were tested for multiple comparisons with a Mann-Whitney U, performed for the different combinations of pairs of each of the factors. The average time it takes for the percentage of motile cells to reach zero was determined with two survival analysis. One to compare extenders and another to compare dilutions.

Regarding viability, the triplicates done for each day, treatment and pool were averaged. In time zero only the control was tested, and assumed to be the value at time zero for the other treatments. The Shapiro-Wilk tests showed that for some treatments the values were not normally distributed (Annex C, Table 17). Therefore, significant differences between treatments were investigated with a GzLM, followed by a Mann-Whitney U test for the pairs of values of the different extenders. The average time it takes for a population to present only 50% of viable cells was detected with survival analysis. Due to the fluctuation verified in viability values over time, the 50% value was obtained with linear regression equations (Annex C, Table 18) on each of the five initial pools. Two different survival analysis were done. One to compare extenders and another to compare dilutions.

In the ATP luminescence readings the mean of the two blanks was subtracted to all readings in the respective plate. The ATP value in $\mu\text{mol/ml}$ of each extract was calculated from the curve equation. The values which gave less or equal to zero were considered to be zero. After this, the values of ATP were converted from $\mu\text{mol/ml}$ to $\mu\text{mol}/10^6$ cells, dividing each value

of ATP for the respective concentration value. For samples with a repeated ATP analysis, the two values obtained for the same sample were averaged. When the highest value for a data point was more than 15 times superior to the second highest value, it was considered to be procedural error and it was removed. In time zero only the control was tested. This was assumed to be the value of the other treatments in time zero. To study the significant differences between treatments only the values obtained at 12 h and days 1 and 3 were considered since after that the ATP values were below the detection limit. The Shapiro-Wilk test showed that the data was not normally distributed (Annex C, Table 18), and so significant differences were tested with a GzLM, with extender and dilution as factors.

For MDA, the values were corrected with respective cell concentration and dilution, in order to obtain MDA in $\mu\text{M}/10^6$ cells. In time zero only the control was tested. This was assumed to be the initial value for the other treatments. The Shapiro-Wilk test showed that the data was not normally distributed (Annex C, Table 19). A GzLM was performed to detect significant differences between the values, with extender and dilution as factors, followed by a Mann-Whitney U test for the pairs of values of the different dilutions.

Finally, for the bacteriology results, the number of colonies counted in each plate at the seventh day of incubation was divided by the inoculation volume, to obtain the CFU/ml value of that inoculation. Counts under 30 were considered to be 0. When the highest value of a triplicate was above one hundred and more than 3 times superior to the second highest value, it was removed. The CFU/ml values of the three plates for each pool, treatment, day and medium were averaged. According to the Shapiro-Wilk test of normality the values were not normally distributed (Annex C, Table 21). Thus, significant differences between treatments were tested with a Kruskal-Wallis, performed for the values of day 5, and another for the values of day 10.

Correlations between the results obtained in each different assay were tested with the Spearman's coefficient test, since the data was not normally distributed (Annex C, Tables 22, 22, 23 and 24). Pairs of values with zeros were discarded. When the sample size was three or less the correlation was not performed.

3. Results

3.1. Concentration

The sperm concentration values obtained for the control sample of each pool ranged from 26.30 to 62.47 x 10⁹ spz/ml (Annex D, Table 23). The average concentration was 43.45 ± 18.17 x 10⁹ spz/ml.

3.2. Motility

Most sperm motility parameters were highly correlated (Table 5) with each other. MOT was selected for further analysis because is the most used motility parameter, whereas progressive movement, PROG, was discarded as it was correlated ($\rho = 0.942$, $n = 379$, $p < 0.01$) with MOT. VCL was also selected for analysis, this is the velocity parameter that corresponds to the spermatozoa's actual path and had a high and significant correlation with the other two velocity parameters, VSL ($\rho = 0.931$, $n = 379$, $p < 0.01$) and VAP ($\rho = 0.992$, $n = 379$, $p < 0.01$). Lastly, LIN also correlated to STR ($\rho = 0.804$, $n = 379$, $p > 0.01$), and was selected to be further used because it is more widely reported than STR.

Table 5: Spearman's rho test of correlation between the values obtained for all the sperm motility parameters measured with CASA.

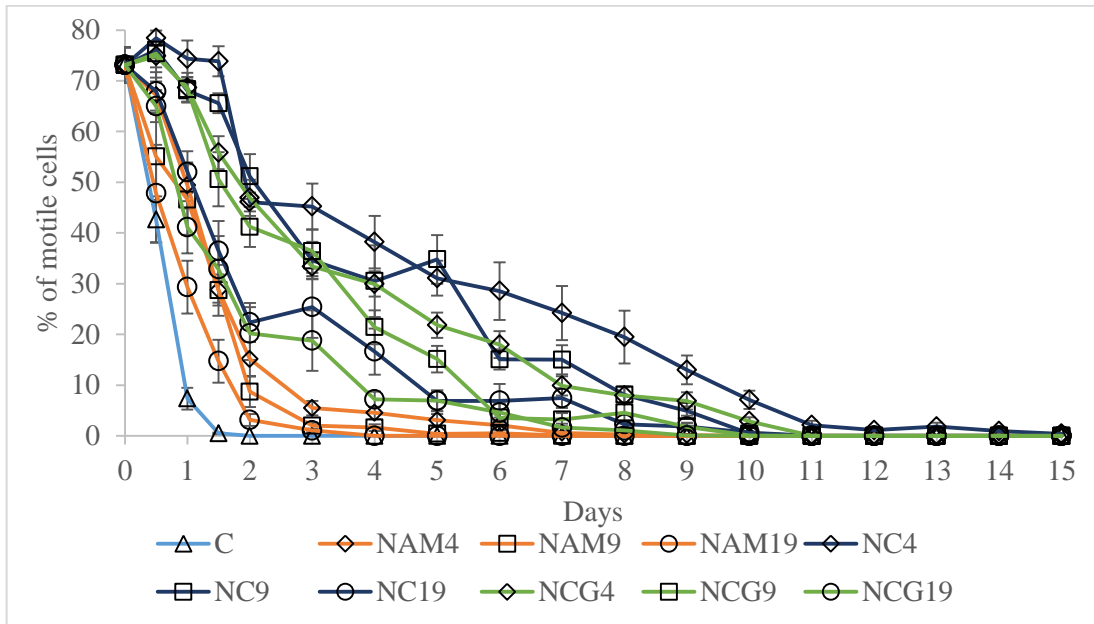
	PROG	VCL	VSL	VAP	LIN	STR
MOT	0.942*	0.794*	0.646*	0.758*	0.017	-0.203*
PROG		0.827*	0.756*	0.804*	0.213*	-0.042
VCL			0.931*	0.992*	0.220*	-0.136*
VSL				0.953*	0.485*	0.108
VAP					0.279*	-0.114
LIN						0.804*

Note: MOT, percentage of motile cells; PROG, percentage of progressive motile cells; VCL, curvilinear velocity; VSL, straight line velocity; VAP, average path velocity; LIN, linearity; and STR, straightness. The sample size refers to the seven pools, and all the days during which motility was assessed. N = 379. *Correlation is significant at the 0.01 level (2-tailed).

The mean percentage of motility was 73.12 ± 3.48% at day 0 (Annex E, Table 27). This value decreased along the time for all treatments. All the treatments had a higher percentage of motile cells than the control at all times (Graph 1; Annex E, Table 27). The control reached 0% motility on average at one and a half days of storage (Graph 2), which was less than half of the average of the extenders and dilutions tested.

There were significant differences between the extenders for the MOT parameter (Wald $\chi^2(2) = 67.186$, $p < 0.001$). The post-hoc analysis (Annex E, Table 28 for a more complete overview of the Mann-Whitney U test) revealed significant differences between all the pairs of extenders. The treatments with the extenders NC and NCG, when compared with those with NAM, kept higher percentage of motile cells during a longer period.

Graph 1: Mean percentage of motile spermatozoa (MOT) of each treatment for each day of assessment.

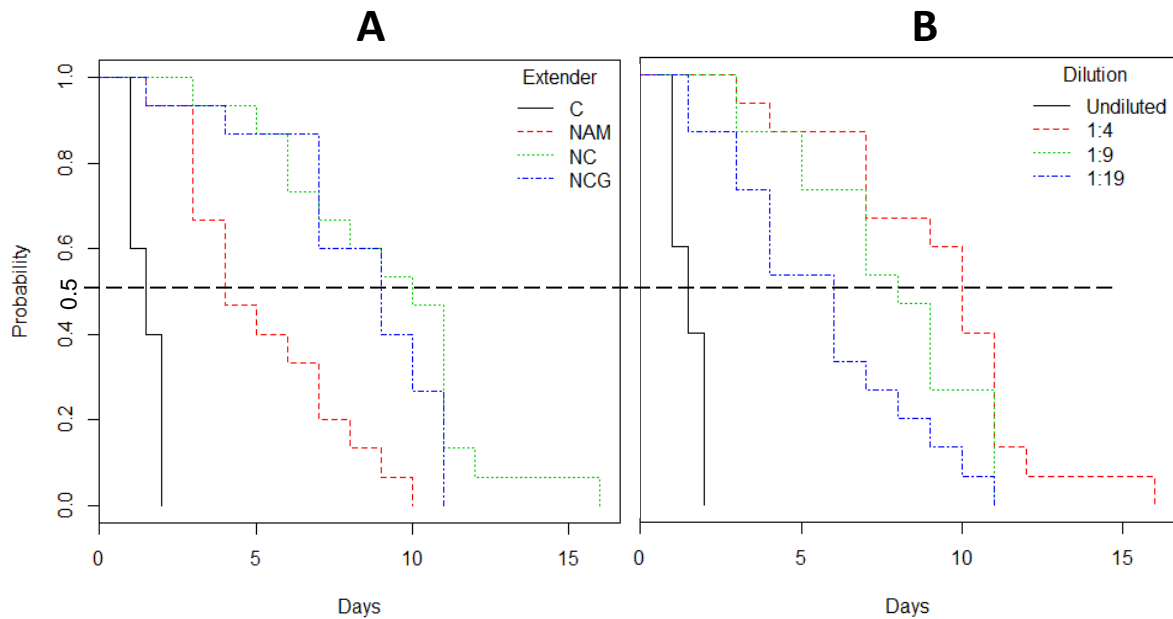


Note: Control in light blue, NAM in orange, NC in dark blue and NCG in green. 1:4 dilution is represented with \diamond , 1:9 dilution with \square and 1:19 dilution with \circ . The value at day zero, for all the treatments, was considered to be equal to that of the control. Motility was assessed every 12 h on the first two days and every 24 h after that. Mean values \pm standard error of the mean. The sample size refers to the first five pools in which sperm motility was assessed, with CASA. N = 5.

In the first day of refrigerated storage the percentage of motile cells of the control treatment dropped to less than 10% and to 0% in the second day. In the treatments with NC the percentage of motile cells was kept above 50% in the first day. Whereas, in the treatments with NCG and NAM it was kept above 40% and 25% respectively. After one and half days, the percentage of motile cells in all treatments with the extender NAM were below the values of the treatments with NC and NCG. At the fourth day, the treatments with NAM had less than 5% motile cells, whereas, the treatments with extenders NC and NCG had more than 15% and 5% motile cells respectively. As seen in Graph 2, the samples with the extenders NC or NCG took longer, on average, to reach 0% motile cells, when compared to the samples with the extender NAM (11, 9 and 4 days, respectively).

There were also significant differences between the dilutions for the MOT parameter (Wald $\chi^2(2) = 32.811, p < 0.001$) (Table 6; Annex E, Table 28 for a more complete overview of the Mann-Whitney U test). The post-hoc analysis revealed that there were significant differences between all the pairs with the exception of 1:4 with 1:9.

Graph 2: Survival analysis until the percentage of motility reaches zero, comparing the values obtained in five pools for: A, the different combinations of extenders (NAM, NC, NCG, C); B, the



different combinations of dilutions (1:4, 1:9, 1:19, Control).

Note: Probability 0.5 is indicated with a dashed line. $N_{NAM, NC, NCG, 1:4, 1:9, 1:19} = 15$. $N_C = 5$.

Table 6: Mann-Whitney U test for the values of MOT, VCL and LIN in pairs of extenders and pairs of dilutions.

	Pair		N	Significantly higher element		
	1	2		MOT	VCL	LIN
Extender	C	NAM	166	2	2	2
	C	NC	180	2	2	2
	C	NCG	180	2	2	2
	NAM	NC	256	2	2	2
	NAM	NCG	256	2	2	2
	NC	NCG	270	1	none	none
Dilution	C	1:4	180	2	2	2
	C	1:9	173	2	2	2
	C	1:19	170	2	2	2
	1:4	1:9	263	none	none	none
	1:4	1:19	260	1	1	1

1:9	1:19	253	1	1	1
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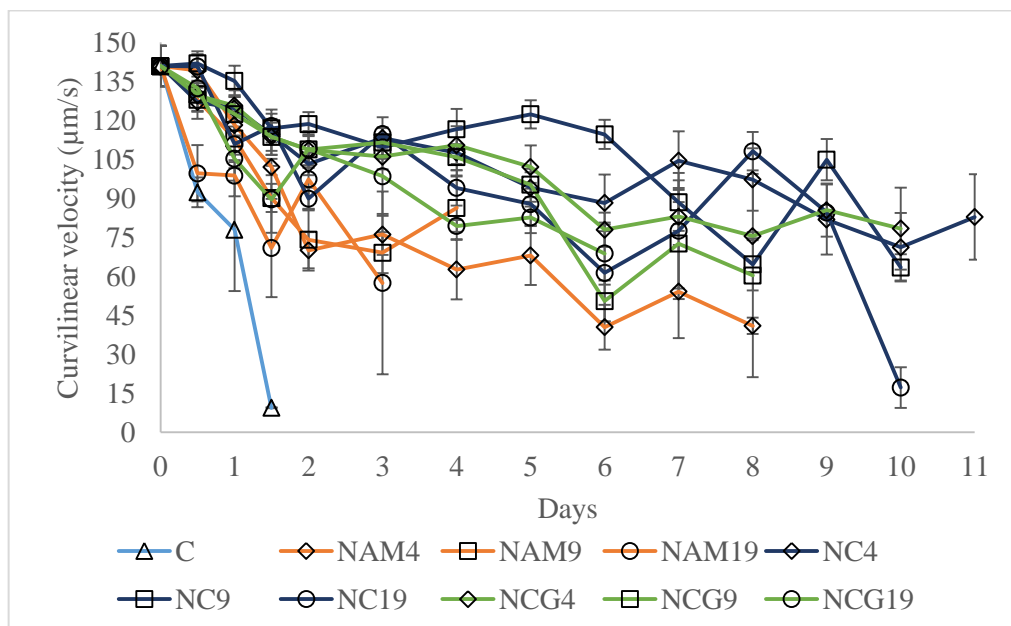
Note: The “Significantly higher element” indicates: if there was no significant difference between the elements of the pair (none); or that there was significant difference between elements of the pair (1 or 2 corresponding to the element with the significantly higher value). Values obtained from the seven pools, from all the days in which motility was assessed for the pair of treatments. The N column corresponds to the sample size value.

Although the dilutions 1:4 and 1:9 did not differ, in the survival analysis (Graph 2) the dilution 1:4 retained the percentage of motile cells above 0% for the longer time, on average ten days. The dilution 1:9 had an average of 8 days, and 1:19 had an average of 6 days. Along the days of evaluation the control had always lower MOT values. When comparing the three dilutions of the same extender, the 1:19 dilution always had lower MOT values, whereas the values of dilution 1:4 were generally higher.

The last recorded day of refrigerated storage was the fifteenth day, for the treatment NC dilution 1:4, with values barely above the 0% of motile cells. This treatment was the only one with a percentage of motile cells above zero after day 11.

The VCL was on average $140.90 \pm 7.75 \mu\text{m/s}$ at day 0 (Annex E, Table 29). This value decreased along the time for all treatments and this decrease was faster in the control treatment (Graph 3). There were significant differences between the extenders (Wald $\chi^2(2) = 82.962, p < 0.001$).

Graph 3: Mean curvilinear velocity (VCL) of each treatment for each day of assessment and treatment.



Note: Control in light blue, NAM in orange, NC in dark blue and NCG in green. 1:4 dilution is represented with ◊, 1:9 dilution with ◻ and 1:19 dilution with ◉. The value at day zero, for all the treatments, was considered to be equal to that of the control. Motility was assessed every 12 h on the first two days and every 24 h after that. Mean values \pm standard error of the mean. The sample size refers to the first five pools in which motility was assessed, with CASA. N = 5.

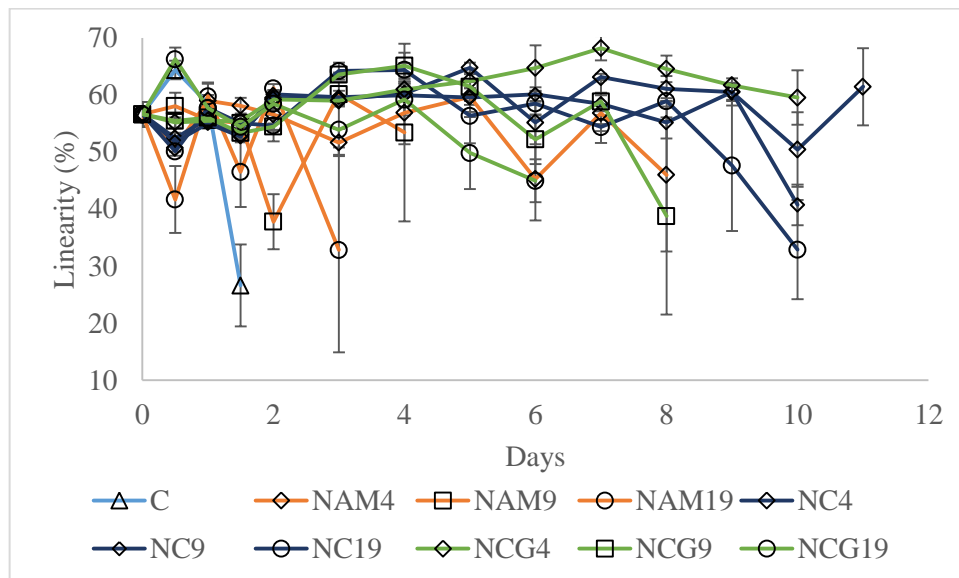
The post-hoc analysis (Annex E, Table 30 for a more complete overview of the Mann-Whitney U test) revealed significant differences between all the pairs of extenders, with the exception of NC with NCG. All the extenders had significantly higher values than the control. Indeed, in every time point the VCL of the control was lower than that of the other treatments tested. In the first day of refrigerated storage the VCL of the control was below 80 $\mu\text{m/s}$, while it was above 95 $\mu\text{m/s}$ in all the samples tested. The VCL values of the extender NAM were significantly lower than those of the extenders NC and NCG. After two days of refrigerated storage, VCL of the treatments with NAM was between 70 and 97 $\mu\text{m/s}$, while it was between 90 and 119 $\mu\text{m/s}$ in those with NC and NCG. At the fourth day of refrigerated storage, the treatments with NAM had an average VCL of less than 86 $\mu\text{m/s}$, nonetheless this value only refers to motile cells and for this treatment means less of 5% of the cells. In treatments with NCG and NC the VCL was above 79 $\mu\text{m/s}$ and 94 $\mu\text{m/s}$ respectively.

Also for VCL, there were significant differences between the dilutions (Wald $\chi^2(2) = 37.054$, $p < 0.001$). The post-hoc analysis (Table 6; Annex E, Table 30 for a more complete overview of the Mann-Whitney U test) revealed that there were significant differences between all the pairs with the exception of 1:4 with 1:9. The dilution 1:19 had the lowest VCL. In fact, the general trend for the different dilutions from the same extender (Annex E, Table 29), at the same time point, was that the dilution 1:19 had the lowest value. Indeed, at the fourth day of storage VCL of the dilution 1:19 of the extender NAM there were no motile cells. Whereas, at day four, the dilution 1:19 of the extender NC had a VCL of 75 $\mu\text{m/s}$, while it was above 100 $\mu\text{m/s}$ in the other two dilutions. In day 7 the dilution 1:19 had an average VCL below 20 $\mu\text{m/s}$, while the dilutions 1:4 and 1:9 had values above 70 $\mu\text{m/s}$. A similar trend was observed for the extender NCG.

The mean LIN was 56.61 ± 2.16 % at day 0 (Annex E, Table 31). For the control LIN was the second highest at 12 h of storage, but from then on had the lowest values. There were significant differences between the extenders (Wald $\chi^2(2) = 71.353$, $p < 0.001$). The results of the post-hoc analysis (Annex E, Table 32 for a more complete overview of the Mann-Whitney U test) are similar to those from the VCL parameter. The control had lower values than the

extenders tested. From these three, NAM had the lowest percentage of LIN. There were no differences between the values from the extenders NC and NCG. At the sixth day of storage, samples from the extender NAM had less than 45% LIN, while those from NC and NCG had more than 45% LIN (Annex E, Table 31).

Graph 4: Mean percentage of linearity (LIN) of each treatment for each day of assessment and treatment.



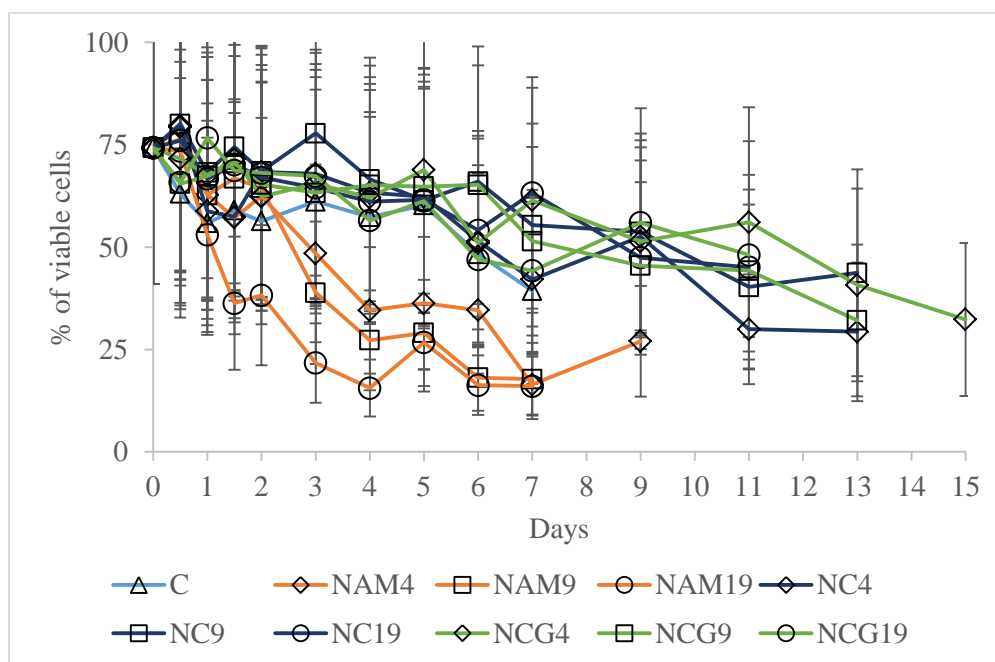
Note: Control in light blue, NAM in orange, NC in dark blue and NCG in green. 1:4 dilution is represented with \diamond , 1:9 dilution with \square and 1:19 dilution with \circ . The value at day zero, for all the treatments, was considered to be equal to that of the control. Motility was assessed every 12 h on the first two days and every 24 h after that. Mean values \pm standard error of the mean. The sample size refers to the first five pools in which motility was assessed, with CASA. N = 5.

There were also significant differences between the dilutions used (Wald $\chi^2(2) = 47.142$, $p < 0.001$). The post-hoc analysis (Table 6; Annex E, Table 32 for a more complete overview of the Mann-Whitney U test) revealed that, similar to what was verified for MOT and VCL, there were differences between the values of all the pairs with the exception of 1:4 with 1:9. The control presented the lowest values, and of the three dilutions tested, 1:19 had the lowest LIN. For the different dilutions of the same extender (Annex E, Table 31), at the same time point, in general the dilution 1:19 had the lowest LIN. For example, at the fourth day, the LIN of the dilution 1:19 of NCG was 36%, while it was 61% and 65% in dilutions 1:4 and 1:9, respectively. The difference was smaller in NC, for which dilutions 1:4 and 1:9 had 60% LIN, and dilution 1:19 had 51%. At the seventh day of storage the LIN of the dilution 1:19 was less than half than that of 1:4 and 1:9, in the case of the extenders NC and NCG.

3.3. Viability

The average percentage of viability was $74.12 \pm 33.20\%$ at day 0 (Annex F, Table 33). The values decreased along the storage time in all the samples. There were significant differences between the extenders used (Wald $\chi^2(2) = 114.887$, $p < 0.001$), but not between the dilutions (Wald $\chi^2(2) = 6.285$, $p = 0.099$). The post-hoc analysis (Table 7; Annex F, Table 34 for a more complete overview of the Mann-Whitney U test) revealed significant differences between most pairs of extenders.

Graph 5: Mean values obtained for each treatment, along the days of evaluation.



Note: Control in light blue, NAM in orange, NC in dark blue and NCG in green. 1:4 dilution is represented with \diamond , 1:9 dilution with \square and 1:19 dilution with \circ . The value at day zero, for all the treatments, was considered to be equal to that of the control. Error bars with the standard error of the mean. Samples were only measured for viability when the MOT value was higher than 0. Therefore in some instances the date points were obtained with less than five pools. No mean value was attributed when it had values from less than three pools. N value ranged from 5 to 3.

Table 7: Mann-Whitney U test done to the pairs of extenders.

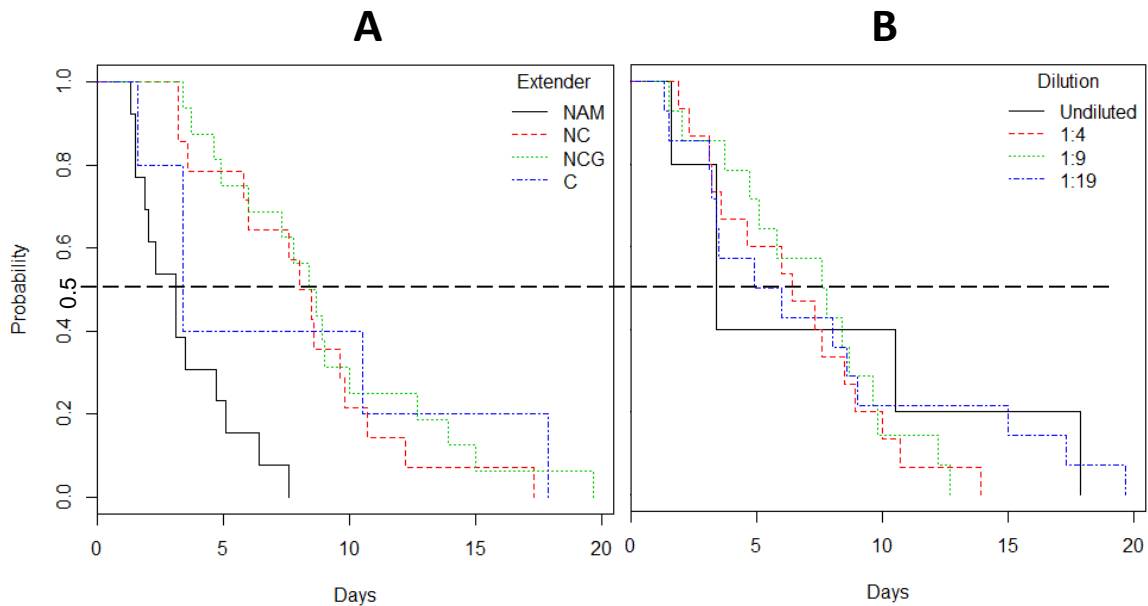
	Pair		Significantly higher element
	1	2	
Extender	C	NAM	1
	C	NC	2
	C	NCG	2
	NAM	NC	2
	NAM	NCG	2

NC	NCG	none
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Note: The “Significantly higher element” indicates: if there was no significant difference between the elements of the pair, with “none”; or that there was significant difference ($p < 0.05$) between elements of the pair, indicating the number (previously attributed in column “Pair”) corresponding to the element with the significantly higher VIAB. N corresponds to the values obtained from the seven pools, from all the days in which viability was assessed. $N_C = 42$, $N_{NAM} = 133$, $N_{NC} = 130$ and $N_{NCG} = 127$.

From the third day on, the extenders NC, NCG, and the control had viability values higher than the treatments with the extender NAM. At the fourth day of storage the viability of the samples with the extender NAM ranged 16 and 35%, while those from extender NC, NCG and control ranged from 57 to 67%. At the seventh day of storage the viability of extender NAM was lower than 18%, while the control was 39% and the values from the extenders NC and NCG were above 42%.

Graph 6: Survival analysis until the viability reaches 50%, comparing the values obtained in five pools for: A, the different combinations of extenders (NAM, NC, NCG, C); B, the different combinations of dilutions (1:4, 1:9, 1:19, Control).



combinations of dilutions (1:4, 1:9, 1:19, Control).

Note: Probability 0.5 is indicated with a dashed line. $N_{NAM, NC, NCG, 1:4, 1:9, 1:19} = 15$. $N_C = 5$.

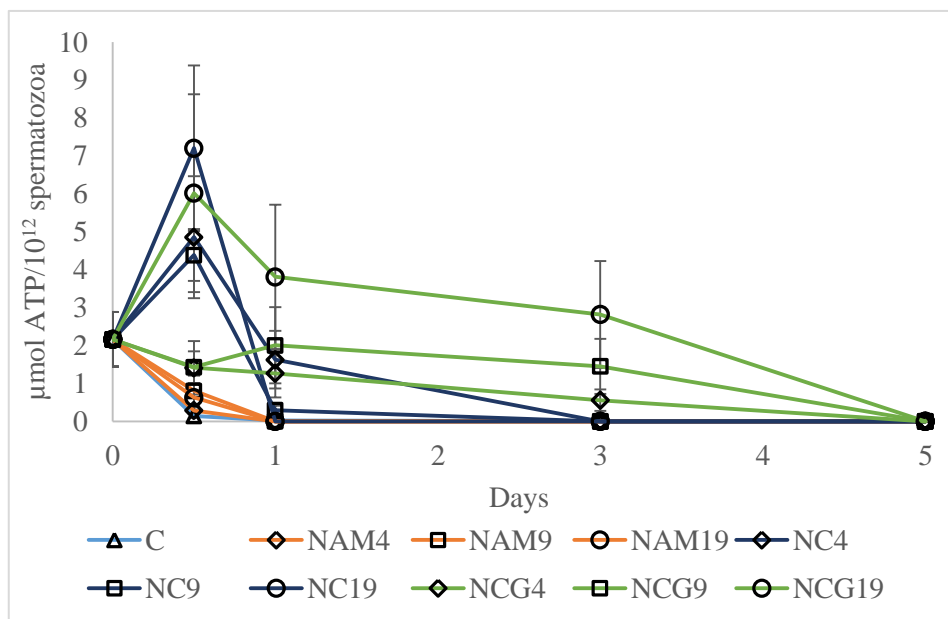
Samples incubated in the extenders NC and NCG had in most cases higher viability than the control. The values from these two extenders did not differ between themselves.

As seen in Graph 5, which represents the average time to reach 50% viability, the samples with the extenders NC or NCG took, on average, eight and a half days to reach 50% viability, while the control and the samples with the extender NAM took on average less than four days. Although there were no significant differences between dilutions, the samples from dilution 1:19 reached 50% viability on average at around 6 days of storage, the dilution 1:4 reached the same percentage on average at six and a half days, and the dilution 1:9 at around 8 days (Graph 6).

3.4. ATP

The average value of ATP was $2.16 \pm 0.72 \mu\text{mol ATP per } 10^{12}$ spermatozoa at day zero (Annex G, Table 35) and at the fifth day of storage these values were below the detection limit for all the treatments (Graph 7).

Graph 7: Mean ATP values obtained for each treatment, from five pools, along the days of viability



evaluation.

Note: Control in light blue, NAM in orange, NC in dark blue and NCG in green. 1:4 dilution is represented with ◊, 1:9 dilution with ◻ and 1:19 dilution with ◉. Mean value at time zero was the same for all the samples, based on the mean from the control. Error bars with the standard error of the mean. N = 4.

Table 8: Mann-Whitney U test done to the pairs of extenders.

	Pair		Significantly higher element
	1	2	
Extender	C	NAM	none
	C	NC	none
	C	NCG	none
	NAM	NC	2
	NAM	NCG	2
	NC	NCG	none

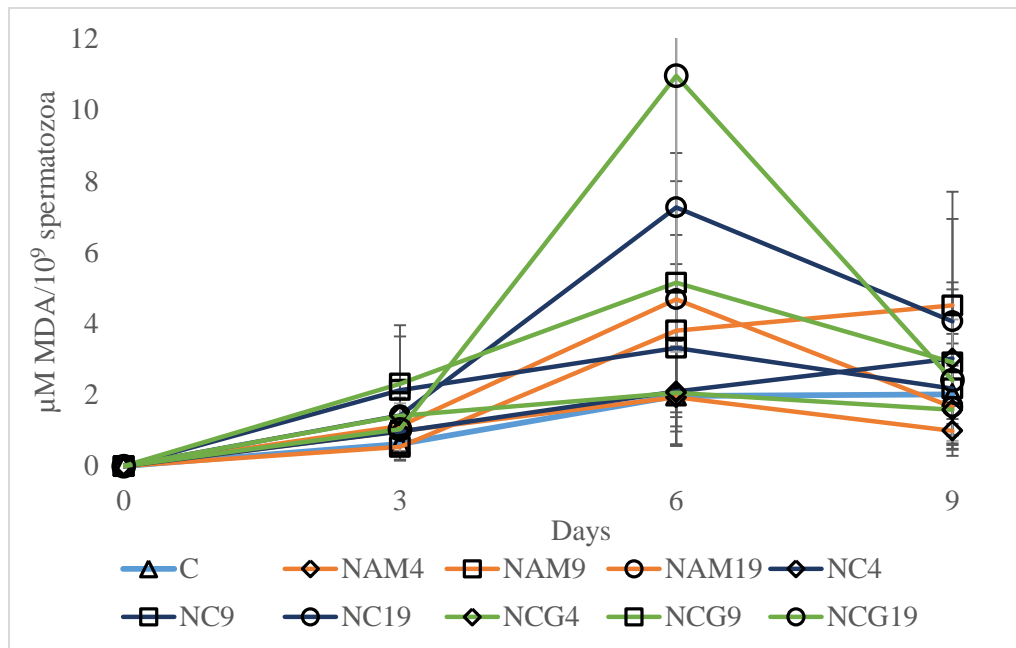
Note: The “Significantly higher element” indicates: if there was no significant difference between the elements of the pair, with “none”; or that there was significant difference ($p < 0.05$) between elements of the pair, indicating the number (previously attributed in column “Pair”) corresponding to the element with the significantly higher ATP. N corresponds to the values obtained from the seven pools, from all the days in which viability was assessed. $N_C = 12$, $N_{NAM} = 34$, $N_{NC} = N_{NCG} = 35$.

There were significant differences between the extenders (Wald $\chi^2(2) = 6.645$, $p < 0.05$), but not between the dilutions (Wald $\chi^2(2) = 4.364$, $p = 0.113$). The post-hoc analysis (Table 8; Annex G, Table 36 for a more complete overview of the Mann-Whitney U test) revealed significant differences between the pairs NAM and NC, and NAM and NCG, but none with the control treatment. The extender NAM had lower values than NC and NCG. In fact, the ATP values of the samples with the extender NAM were lower than the other two extenders at all times. Samples kept in the extender NAM only kept the ATP value above the detection limit until the first day of storage. At 12 hours of refrigerated storage all the treatments with the extenders NC or NCG had more than $1.40 \mu\text{mol ATP per } 10^{12}$ spermatozoa, while the ATP content of the samples with NAM was below $0.85 \mu\text{mol ATP per } 10^{12}$ spermatozoa.

3.5. MDA

The value of MDA on time zero was 0 μM MDA per 10^6 spermatozoa (Annex H, Table 37). The values increased in all the treatments until the sixth day of storage, and decreased between days 6 and 9 in most treatments (Graph 8). Furthermore, the values from day 9 were all higher than those at day 3, from the corresponding treatment.

Graph 8: Mean and standard error of the mean of the values of MDA at days 0, 3, 6 and 9, in which it was quantified.



Note: Control in light blue, NAM in orange, NC in dark blue and NCG in green. 1:4 dilution is represented with \diamond , 1:9 dilution with \square and 1:19 dilution with \circ . Mean value of the treatments at time zero was considered to be equal to the mean of the control. Pools 6 and 7 were used in this evaluation. N = 2.

There were no significant differences in the values between the groups of extenders (Wald $\chi^2(2) = 1.765$, $p = 0.414$), but there were significant differences between dilutions (Wald $\chi^2(2) = 7.216$, $p < 0.05$). The post-hoc analysis (Table 9; Annex H, Table 38 for a more complete overview of the Mann-Whitney U test) revealed significant differences between only one pair of dilutions. The MDA content of dilution 1:9 was higher than that of dilution 1:4. As seen in Table 37, from Annex H, the mean values from dilution 1:9 were higher than those from dilution 1:4 from the same extender, with a couple of exceptions.

Table 9: Mann-Whitney U test done to the MDA values in pairs of different dilutions.

	Pair (1 x 2)	Significantly higher element
Dilution	C 1:4	none
	C 1:9	none
	C 1:19	none

1:4	1:9	2
1:4	1:19	none
1:9	1:19	none

Note: The “Significantly higher element” indicates: if there was no significant difference between the elements of the pair, with “none”; or that there was significant difference ($p < 0.05$) between elements of the pair, indicating the number (previously attributed in column “Pair”) corresponding to the element with the significantly higher MDA. N corresponds to the values obtained from the two pools, from all the days in which MDA was assessed. $N_{C \times 1:4} = N_{C \times 1:9} = N_{C \times 1:19} = 24$. $N_{1:4 \times 1:9} = N_{1:4 \times 1:19} = N_{1:9 \times 1:19} = 36$.

3.6. Bacteriology

The CFU/ml obtained from the extenders was discarded as a contamination source, since its values were mostly zero.

Table 10: Mean \pm standard error of the mean of the CFU/ml obtained for each treatment (C, NAM, NC and NCG) and day, in TSA medium. Values from three pools. $N = 3$.

Treatment	Day		
	0	5	10
C	0	0	242200.00 ± 39984.00
NAM		15689.00 ± 2643.34	18513.00 ± 2860.36
NC		7528.90 ± 1271.05	84.44 ± 28.15
NCG		0	23255.56 ± 3516.67

Table 11: Mean \pm standard error of the mean of the CFU/ml obtained for each treatment (C, NAM, NC and NCG) and day, in PS medium. Values from three pools. $N = 3$.

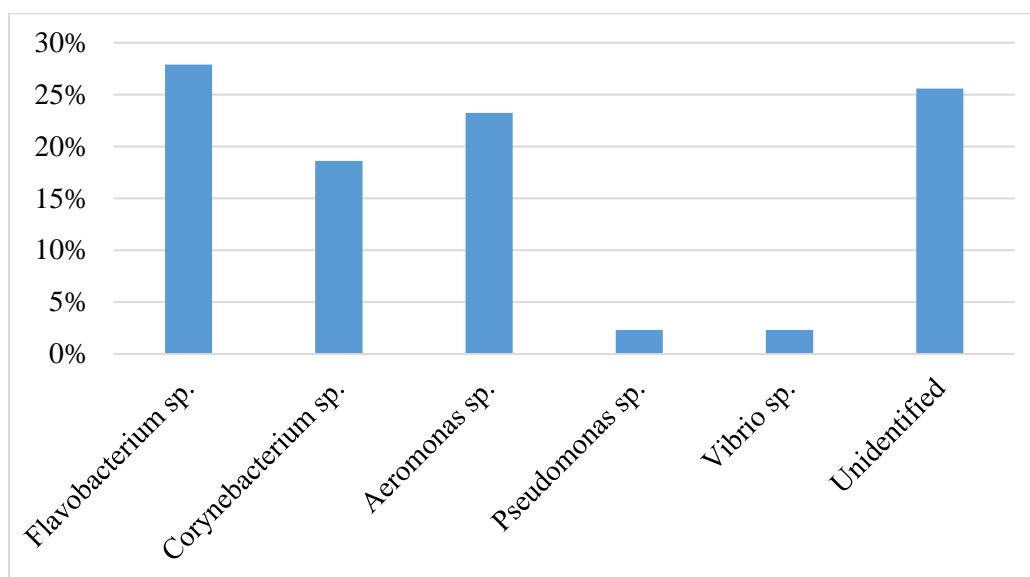
Treatment	Day		
	0	5	10
C	333.33 ± 111.11	1222.20 ± 407.41	0
NAM		120.00 ± 40.00	2268.89 \pm 296.61
NC		0	0
NCG		0	413.33 ± 74.77

The value of CFU/ml on time zero was 0 for TSA (Table 10). At day 5 and at day 10 of incubation there were no significant differences between CFU/ml of the treatments tested (χ^2 (3) = 2.212, $p = 0.530$ and χ^2 (3) = 1.528, $p > 0.676$ respectively).

The value of CFU/ml on time zero was 333.33 ± 111.11 for PS mediums (Table 11). The CFU/ml was not significantly different between the treatments tested at day 5 ($\chi^2(3) = 4.992$, $p = 0.172$), nor at day 10 ($\chi^2(3) = 2.212$, $p = 0.530$).

Of all the isolated colonies, 25% were not possible to identify (Graph 9 and Table 39 from Annex I). The most identified genus was *Flavobacterium* sp., comprising 28% of all the colonies, followed by *Aeromonas* sp. and *Corynebacterium* sp., 23% and 19%, respectively. Lastly, *Pseudomonas* sp. and *Vibrio* sp. represent around 2% each.

Graph 9: Relative frequency of the genera of the bacteria that were isolated at the seventh day of incubation in the TSA and PS mediums from the three pools. $N = 43$.



3.7. Correlations

Viability was positively correlated with MOT ($\rho = 0.323$, $n = 293$, $p < 0.01$) and VCL ($\rho = 0.223$, $n = 293$, $p < 0.01$) and negatively correlated with LIN ($\rho = -0.231$, $n = 293$, $p < 0.01$), as seen in Table 12. ATP content was correlated with MOT ($\rho = 0.168$, $n = 35$, $p < 0.05$) (Table 12).

Table 12: Spearman's rank correlation coefficient between the different parameters tested.

	MOT	VCL	LIN	VIAB
VIAB	0.323**	0.223**	-0.231**	
ATP	0.350*	-0.024	-0.148	0.261
MDA	0.071	0.198	0.266	0.222
CFU/ml TSA	***	***	***	-0.100
CFU/ml PS	***	***	***	0.500

-
- * Correlation is significant at the 0.05 level (2-tailed).
 - ** Correlation is significant at the 0.01 level (2-tailed).
 - *** Correlation not performed due to the sample size ($N \leq 3$).

Note: MOT, percentage of motile cells; VCL, curvilinear velocity; LIN, linearity; VIAB, percentage of viable cells; ATP, adenosine triphosphate; MDA, malondialdehyde; TSA, Trypticase Soya Agar medium; PS, CN Pseudomonas Agar medium. Values equal to zero were excluded. For VIAB x MOT, VCL and LIN, $N = 293$. Values from seven pools and all the days of evaluation. For ATP x MOT, VCL and LIN, $N = 35$. For ATP x VIAB, $N = 32$. Values from five pools, obtained between 0 and 3 days of storage. For MDA with MOT, VCL and LIN, $N = 33$. For MDA x VIAB, $N = 57$. Values from two pools, obtained between 3 and 9 days of storage. For CFU/ml TSA x VIAB, $N = 5$. For CFU/ml PS x VIAB, $N = 4$. Values from three pools, obtained between 0 and 10 of storage.

There was no correlation between the MDA values and the other measured parameters ($\rho \geq 0.266$, $n \geq 32$ and $p > 0.05$) (Table 12). The correlation tests between CFU/ml of the TSA and PS medium and the remaining parameters were considered not biologically meaningful because of the reduced sample size ($n \leq 3$).

4. Discussion

The reproduction of meagre is one of the areas that requires research and improvement, for the production of high quality spawns at a lower price. In this work a protocol for semen refrigeration in meagre was developed that will facilitate the artificial reproduction techniques. To determine the best extender and dilution ratio for the protocol, semen quality was evaluated during several days based on the following quality parameters: sperm motility, sperm viability, energy content (ATP), lipid peroxidation (MDA) and bacteriology.

4.1. Concentration

The cell concentration values obtained in the present thesis were generally higher compared to the values reported for meagre by Mylonas et al. (2013b). The maximum concentration obtained on this experiment was 63×10^9 spz/ml versus 32×10^9 spz/ml obtained by Mylonas et al. (2013b). Nonetheless, these values are within the range reported for several teleost species, by Piironen and Hyvärinen (1983) and Poole and Diilane (1998).

The differences between cell concentration values in this work and those obtained by Mylonas et al. (2013b) could be due to normal variability between individuals and different experimental conditions, but also caused by different measurement methods. While in this study the concentration was assessed with a CASA system, Mylonas et al. (2013b) used a Neubauer chamber. This last method, hemacytometry, is the gold standard, while CASA systems require the determination of a compensation factor to ensure accuracy (Kuster, 2005).

Nonetheless, different factors could have affect cell concentration such as different feeding regimes, the size of the fish (in this study fish were larger, average of 7.3 kg versus average of 4.6 kg in Mylonas et al. (2013b)), or the rearing conditions (outdoor tanks in this study versus indoor tanks in the study by Mylonas et al. (2013b)).

4.2. Motility

The percentage of motile spermatozoa of fresh semen immediately after collection in this study ($73 \pm 3.5\%$) was similar to the one obtained in the other two studies in meagre. Schiavone et al. (2012) obtained between 53 and 74% (depending on the time during the spawning season), and Mylonas et al. (2013b) had values that ranged between 44% and 80%. In all these studies, the motility was evaluated with a CASA system. Samples with this high percentage of motile cells should be able to achieve high fertilization rates, given that the quality of eggs is good and enough volume of sperm is used (e.g. Casselman et al. 2006; Ottesen et al. 2009). In relation to the sperm velocity immediately after collection, the values of $141 \pm 8 \mu\text{m/s}$ observed for meagre in this study were slightly higher than the values described for other Perciformes. As an example, the Lahnsteiner & Patzner (1998) described for the white seabream (*Diplodus sargus*) $127 \pm 23 \mu\text{m/s}$ VCL, whereas, the European sea bass semen, which has some similarities to meagre semen had a curvilinear velocity lower than $80 \mu\text{m/s}$ at 15 seconds after activation (Fauvel et al., 1998). On the other hand, the linearity of meagre's fresh semen immediately after collection of $57 \pm 2\%$ appears to be lower, when compared to other teleosts. While zebrafish (*Danio rerio*) semen have an average linearity of 84% (Wilson-Leedy & Ingermann, 2007) the bluegill (*Lepomis macrochirus*) has an average linearity of 76%.

As expected, during refrigeration, the values of percentage of motility, curvilinear velocity and linearity obtained with the undiluted sample were lower than those from any of the diluted treatments. Also, in another sciaenid, the Atlantic croaker (*Micropogonias undulatus*), it was verified that the undiluted sample reached zero motility earlier than the diluted treatments, in semen refrigerated at 4°C (Leclercq et al., 2012). In effect, as stated by Lahnsteiner et al. (1996) and Maria et al. (2014), diluting semen with an extender contributes to its preservation, which translates in better values of quality parameters throughout time.

The highest percentages of motility along the time were obtained with dilutions 1:4 and 1:9, and with the extenders 0.9% NaCl and 0.9% NaCl with glucose and glycine. No differences were found between the results obtained with dilutions 1:4 and 1:9. This was also verified in Atlantic cod (*Gadus morhua*) (DeGraaf & Berlinsky, 2004) and black sea bass (*Centropristis*

striata) (DeGraaf, King, Benton & Berlinsky, 2004), when comparing refrigerated semen, diluted with 1:4 and 1:9 modified Mounib's extender. Nonetheless, and although not significantly, the dilution 1:4 kept motility for longer in black sea bass semen, as also observed in this experiment.

Comparing with other Sciaenids, the results obtained for motility along storage time appear similar or better. Atlantic croaker semen diluted in 1:2 or 1:9 HBSS retained motility for less than 4 days (Leclercq et al., 2014), whereas in the present work, at 4 days, it had on average 38% and 31% motile cells when diluted 1:4 or 1:9 in 0.9% NaCl, respectively. Semen of black drum (*Pogonias cromis*) diluted in 202 or 286 mOsm/kg HBSS retained 8% of motile cells at the eighth day of refrigerated storage (Wayman et al., 1997), while in this experiment it retained 13% of motility until the ninth day, in semen diluted 1:4 in 0.9% NaCl. In the present experiment the motility was kept for a maximum of 15 days, with the 1:4 dilution of 0.9% NaCl. These good results could be due to the meagre semen being more resistant to refrigeration or because of a positive effect of some of the treatments tested. Additionally, in this thesis only extenders with 300 mOsm/kg were tested, chosen according to the osmolality of meagre's seminal plasma. Whereas, in red drum (*Sciaenops ocellatus*) semen refrigerated at 1°C and diluted in HBSS or NaCl with 200 mOsm/kg kept some motility until the thirteenth day, while the same extenders with 300 or 400 mOsm/kg kept motility for only 11 days (Wayman et al., 1998). The storage temperature of 4°C in this study was selected, for practical reasons of usage in a fish farm facility. In this sense, the Wayman et al. (1998) results could indicate that a lower refrigeration temperature together with a slightly decrease of the extenders' osmolality might extend meagre sperm usage.

From the different semen quality parameters, motility has been shown to be the parameter that more frequently correlates with fertilization rate. See examples in species such as European sea bass (Fauvel, Savoye, Dreanno, Cosson, & Suquet, 1999) and gilthead seabream (Beirão et al., 2011a). Therefore, when selecting for a semen refrigeration protocol special emphasis should be placed in the sperm motility evaluation. According to the sperm motility values, semen should be refrigerated with 0.9% NaCl and 0.9% NaCl with glucose and glycine in dilutions 1:4 and 1:9.

4.3. Viability

The percentage of viability of fresh semen immediately after collection was $74 \pm 33\%$. This value is in a range normally observed in other species. For example, in the literature the values for other euryhaline species vary between 94% in Mangrove red snapper (*Lutjanus*

argentimaculatus) (Vuthiphandchai, Chomphuthawach & Nimrat, 2009) to 83% for brook trout (*Salvelinus fontinalis*) (Nynca & Ciereszko, 2009).

In the present work, while there was no difference between the dilutions tested (1:4, 1:9 or 1:19), the extender had a significant effect in the percentage of viable cells during refrigeration. Whereas the semen viability was not prolonged by NAM, compared with the undiluted sample, both the samples diluted in 0.9% NaCl and 0.9% NaCl with glycine and glucose had more than 50% viability for 10 days. Different authors have already described that the dilution of semen prolonged its viability during refrigeration (e.g. Maria et al. 2006; McNiven et al. 1993). Thus, the negative effects of the extender NAM in the sperm viability were unexpected. Even more so, the undiluted sample had lower values of MOT and VCL when compared to NAM, and these two parameters are correlated with VIAB. This unexpected result may be due to the NAM composition, for example the ionic composition and the addition of the BSA could be unappropriate, as this extender was developed for the cryopreservation of sea bass (Fauvel et al., 1998). However, the cause for the lower viability values with the NAM extender are not clear.

In general, the decrease in the viability with time helped explained the decrease in the sperm motility parameters. In effect, and similar to the observation made by other authors in different teleosts, there was a positive correlation between sperm viability and the percentage of motile cells (Beirão et al. 2011b; Horváth et al. 2008; Lanes et al. 2008), as well as sperm velocity (Linhart, Rodina, Flajshans, Gela & Kocour, 2005). Linearity was also correlated with viability. Despite the absence of publications correlating both parameters, this goes in accordance with the correlation found between the other two motility parameters.

4.4. ATP

The ATP content of fresh meagre semen, assessed after its collection, was relatively low (2.16 μmol of ATP per 10^{12} spermatozoa) compared with other marine teleosts with a similar motility period and velocity. As an example, European sea bass semen had 115 μmol of ATP per 10^{12} spermatozoa (Dreanno et al., 1999a), and turbot semen had between 1450 and 1080 μmol of ATP per 10^{12} spermatozoa (Suquet et al., 1998). Expectedly, the ATP content decreased over time, by more than half between 12 h and day 1, with the exception of the samples stored in the extender NCG. When compared with other species, the decrease verified in the present study was more abrupt. In refrigerated beluga (*Huso huso*) and steelhead trout semen the ATP decreased by half after three to four days (Aramli, 2014; Bencic et al., 1999b).

In the present work the values fell to undetectable levels in all treatments within five days, even though some still had motility values above 30%. Despite the fact that the samples diluted in 0.9% NaCl with glycine and glucose were the only ones with measurable values above the detection limit in the third day of refrigeration, they were not statistically different from the control. In contrast, Dziewulska et al. (2010) reported that the content of ATP in Atlantic salmon semen was significantly higher when it was diluted, when compared to undiluted. Nevertheless, the present work analysis refers to the entire period of refrigeration, whereas the observations by Dziewulska et al. (2010) refers only to one time point (10 h). An analysis focused in specific time points of the meagre sperm refrigeration, is likely to find similar differences between undiluted sperm and the two best extenders (0.9% NaCl and 0.9% NaCl with glycine and glucose).

As expected, the ATP content, associated to the energy and thus movement of the spermatozoa, was positively correlated with MOT. A similar correlation has been found also in some freshwater teleosts, such as rainbow trout (De Baulny et al., 1996; Lahnsteiner et al., 1998) and bluegill (*Lepomis macrochirus*) semen (Burness et al., 2005), but not in bleak (*Alburnus alburnus*) semen (Lahnsteiner et al., 1996). Therefore, the depletion of the energy reserves in the form of ATP may partially explain the loss of the motility of the refrigerated sperm. Nonetheless, contrary to the correlation found by Burness et al. (2005) in bluegill, in this work there was no significant correlation between ATP and swimming velocity, neither between ATP and linearity. Characteristics other than ATP content may influence the swimming parameters.

4.5. MDA

As expected, the MDA values increased in all treatments with the refrigeration time, peaking in the 6th day, with the highest value being 109.63 ± 77.52 from the extender 0.9% NaCl with glucose and glycine in the dilution 1:19. Likewise, in another teleost, the Persian sturgeon (*Acipenser persicus*), the levels of TBARS, also indicative of lipid peroxidation, increased significantly with the storage time (Aramli et al., 2013). In meagre, however, a decrease was observed between days 6 and 9, in most treatments. Since this was verified in both pools, assessed in different occasions, it is assumed that this was not due to an error in the thiobarbituric acid assay. The production of MDA requires oxygen and lipids. A depletion of these would lead to a lower production of MDA, which was not the case. The quantity of oxygen did not differ between the several days, as the falcons were opened regularly. Furthermore, as lipids are the main component of the spermatozoa's membranes, it appears

that these would not be a limiting factor. However, there are no publications regarding the half-life of MDA. Thus, the cause for this decrease is not clear.

There was no difference between the extenders used, only between dilutions 1:4 and 1:9, with the latter having higher MDA values. Either the samples are not differently affected by the dilutions and extenders tested, or this is due to the small sample size ($n = 2$). The MDA values had to be measured in the same day in which the samples were collected and it was not logistically feasible to evaluate it in more than two pools.

Finally, although lipid peroxidation leads to degradation of spermatozoa, there was no correlation between the MDA values and viability or motility parameters. Aramli et al. (2013) and Ball (2008) also did not find correlations between them. Thus, despite not being a parameter with a greater influence, MDA appears to somewhat impact semen quality.

The results evidence that lipid peroxidation occurs during the refrigerated storage of meagre semen. Taking this in consideration, antioxidants should be added to the extenders design to prevent cell injury caused by oxidative stress (Lahnsteiner & Mansour, 2010; Shaliutina-Kolešová et al., 2013).

4.6. Bacteriology

The number of CFU/ml increased with the refrigeration time in most treatments, as expected, and also observed by Viveiros et al. (2010a) in piracanjuba. No differences were found between the CFU/ml of the different treatments at days 5 and 10, in TSA and in the PS mediums. Nonetheless, the mean CFU/ml in the TSA medium was more than ten times superior for the undiluted treatment at the last day of incubation than for the diluted treatments tested. This could indicate that undiluted semen promotes the growth of more bacteria when compared to the diluted sperm. The number of pools tested might not have been sufficient to obtain statistically significant results. Additionally, and despite care taken with disinfection during the plate's incubation, the possibility of cross contamination can not be totally discarded.

Most authors associate the presence and growth of bacteria with the degradation of semen quality, with negative effects in fertilization, motility and viability (Nimrat & Vuthiphandchai, 2008). The sample size for correlation of CFU/ml from either TSA or PS mediums with the parameters was small, meaning that the results of the statistical test were inconclusive. Statistics with a larger sample size could lead to the verification of additional correlations.

During the qualitative analysis five bacteria genera were identified. The most prevalent genus was *Flavobacterium* sp. Although it is normally found in the water of freshwater fish aquaculture (Kubilay, Altun & Savas, 2009; Welker, Shoemaker, Arias & Klesius, 2005), it has also been identified in the intestinal tract of marine fish (Nayak, 2010). Additionally, Boonthai et al. (2016) found it in semen collected without cleaning the urogenital area. Despite the care taken in the present work to clean the urogenital papilla from water, urine and feces, there is the possibility of some contamination during the sampling procedure, as there was no disinfection. Pathogenic species of *Flavobacterium* sp. have been isolated in fish semen (Kumagai & Nawata, 2011; Madsen & Dalsgaard, 2008), but Oplinger and Wagner (2015) do not consider semen as a relevant infectious medium for this genus. Therefore, the main concern regarding the presence of *Flavobacterium* sp. is associated to the possible effect in semen degradation.

Concerning *Aeromonas* sp., it is found in both freshwater and marine fish (Nimrat & Vuthiphandchai, 2008). It has been found in hatchery water of rainbow trout (Kubilay et al., 2009), skin of freshwater fish (Austin, 2006), digestive tract of freshwater fish (Nayak, 2010; Nimrat & Vuthiphandchai, 2008) as well as semen of silver barb (*Barbodes gonionotus*) (Boonthai et al., 2016). In fact, in rainbow trout, *Aeromonas* sp. was isolated in semen collected with and without cleaning the urogenital area (Boonthai et al., 2016). Thus, its presence in the present experiment could be due to contamination or with origin in the sperm. Regarding *Corynebacterium* sp. and *Vibrio*, these genera have been isolated from the skin (Austin, 2006; Austin & Al- Zahrani, 1988; Gillespie & Macrae, 1975; Gilmour, McCallum & Allan, 1976) and intestinal tract (Austin & Al- Zahrani, 1988; Nayak, 2010; Trust & Sparrow, 1974) of marine and freshwater fish, as well as euryhaline in the case of *Corynebacterium* sp. This genus is the most prevalent in hatchery water of rainbow trout (Kubilay et al., 2009). The presence of these two genera is most likely related with contamination.

Finally, bacteria from the genus *Pseudomonas* sp. are commonly found in soil and water (Sader & Jones, 2005), but are also present in fish, as part of the microbiota of the semen (Boonthai et al., 2016; Jenkins & Tiersch, 1997) and gastrointestinal tract (Nayak, 2010) of both marine and freshwater fish. Boonthai et al (2016) isolated it in channel catfish (*Ictalurus punctatus*), even when collecting semen with a catheter and cleaning of urogenital pore, although in a smaller number, meaning that it may be present in the semen of that species. Therefore, the detection of *Pseudomonas* sp. in the present study could have been isolated due to contamination or presence in the semen. Jenkins and Tiersch (1997) consider *Pseudomonas*

sp. to be responsible for the shortening of the semen storage period. Since in this experiment the genus was only identified once its effects were not adequately assessed.

The usage of antibiotics such as gentamycin (Viveiros et al., 2010a) is considered essential by some authors (DeGraaf & Berlinsky, 2004; Segovia, Jenkins, Paniagua-Chavez & Tiersch, 2000). However, as observed by Oplinger and Wagner (2015) in production-size trials, the usage of antibiotics in the extenders causes a decrease in egg fertilization rate. In the present work we discarded the hypothesis of using antibiotics, since one of the objectives was to develop a protocol that could be used by fish farmers.

5. Conclusions

The sperm samples in the extenders 0.9% NaCl and 0.9% NaCl with glycine and glucose in the 1:4 and 1:9 (sperm:extender) dilutions kept better semen quality, namely in motility, viability and bacteriology, when compared to the undiluted treatment and the other treatments tested.

Both the decrease in sperm viability, the ATP depletion, the increase in lipid peroxidation and the increase in bacterial counts seem to contribute to the degradation of the sperm quality during refrigerated storage. Thus, the addition of antioxidants to the extender may increase its quality.

Since extenders with different osmolalities gave better motility results in another Sciaenidae, this may also be tested futurely.

Overall, the best option according to the results for the refrigeration of meagre semen was the dilution 1:4 (sperm:extender) in the extender 0.9% NaCl. At the fourth day of storage, it had more than 40% of motile cells, and at the 10th day of storage it had approximately 10%. This means that if enough volume of sperm is available, good fertilization rates could still be obtained even after 10 days of refrigeration, in these conditions.

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ANNEX

Annex A: Temperature of refrigeration.

Table 13: Mean maximum and minimum values of temperature of semen storage, as registered daily in the temperature sensor, during the days in which the seven pools were stored..

	Temperature (°C)
Mean Minimum	2.6
Mean Maximum	7.7

Annex B: Composition of solutions used in the ATP protocol.

Table 14: Composition of the solutions that were used during the procedure of extraction and quantification of ATP from meagre semen.

Solution	Composition
Sorensen stock solution A	0.1 M KH_2PO_4 in distilled water
Sorensen stock solution B	0.1 M $\text{Na}_2\text{HPO}_4 \cdot 2 \text{H}_2\text{O}$ in distilled water
Sorensen buffer (pH \approx 7.8)	10 parts solution A : 95 parts solution B
Ethylenediaminetetraacetic acid (EDTA) solution	2mM $\text{C}_{10}\text{H}_{16}\text{N}_2\text{O}_8$ in distilled water (dissolved at pH=8 with the help of Borax)
2% TCA solution	2% trichloroacetic acid in EDTA solution

Annex C: Intermediate tests of the data analysis.

Table 15: Shapiro-Wilk normality test for each motility parameter.

	Statistic
MOT	0.904
PROG	0.924
VCL	0.986
VSL	0.983
VAP	0.989*
LIN	0.898
STR	0.865

Note: MOT, percentage of motile cells; PROG, percentage of cells with progressive movement; VCL, curvilinear velocity; VSL, straight line velocity; The values used were the ones obtained in the seven pools, from all the treatments, in all the days in which motility was assessed. For each test, N = 379. Degrees of freedom = 379. *Normality is statistically significant, $p > 0.05$.

Table 16: Values for the Statistic of the Shapiro-Wilk normality test, done for each combination of treatment.

	MOT	VCL	LIN
C	0.409	0.473	0.524
NAM4	0.727	0.915*	0.796
NAM9	0.658	0.870	0.801
NAM19	0.533	0.650	0.649
NC4	0.949*	0.960*	0.719
NC9	0.936	0.846	0.654
NC19	0.845	0.907	0.799
NCG4	0.953*	0.969*	0.756
NCG9	0.898	0.960*	0.775
NCG19	0.809	0.917	0.789

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19. MOT, percentage of motile cells; VCL, curvilinear velocity; and LIN, linearity. The values used were obtained during the initial seven days of evaluation, in five pools. For each test, N = 38. Degrees of freedom = 38. *Normality is statistically significant, $p > 0.05$.

Table 17: Shapiro-Wilk normality test done to the percentage of viability values of each treatment.

	N	Statistic
C	42	0.990*
NAM4	45	0.954*
NAM9	44	0.949*
NAM19	44	0.929
NC4	42	0.983*
NC9	43	0.853
NC19	45	0.912
NCG4	38	0.979*
NCG9	44	0.983*
NCG19	45	0.969*

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19. Values obtained in five pools, in the first seven days, grouped according to combinations of extender with dilution. N corresponds to the sample size. For all the groups, degrees of freedom = 38. *Normality is statistically significant, $p > 0.05$.

Table 18: Linear regression equations, and r^2 value, for the viability values along the time of each treatment from each pool.

	Pool				
	1	2	3	4	5
C	$y = -0,2129x + 72,496$ $r^2 = 0,8146$	$y = -0,1533x + 55,904$ $r^2 = 0,5621$	$y = -0,1864x + 65,138$ $r^2 = 0,7945$	$y = -0,0861x + 87,051$ $r^2 = 0,7169$	$y = -0,0976x + 74,632$ $r^2 = 0,7688$
NAM4	$y = -0,3328x + 74,919$ $r^2 = 0,8468$	$y = -0,4319x + 70,177$ $r^2 = 0,8684$	$y = -0,4206x + 72,97$ $r^2 = 0,9118$	$y = -0,1582x + 74,141$ $r^2 = 0,7516$	$y = -0,1604x + 79,141$ $r^2 = 0,7686$
NAM9	$y = -0,2927x + 48,009$ $r^2 = 0,7736$	$y = -0,4126x + 69,586$ $r^2 = 0,7743$	$y = -0,4183x + 65,494$ $r^2 = 0,8668$	$y = -0,3441x + 89,152$ $r^2 = 0,9069$	$y = -0,2965x + 86,559$ $r^2 = 0,8793$
NAM19	$y = -0,3342x + 39,696$ $r^2 = 0,8168$	$y = -0,4314x + 63,377$ $r^2 = 0,7294$	$y = -0,4634x + 66,17$ $r^2 = 0,8237$	$y = -0,3408x + 75,506$ $r^2 = 0,7189$	$y = -0,3821x + 82,486$ $r^2 = 0,8373$
NC4	$y = -0,1297x + 76,569$ $r^2 = 0,7044$	$y = -0,1611x + 62,456$ $r^2 = 0,7571$	$y = -0,2155x + 68,683$ $r^2 = 0,7109$	$y = -0,133x + 84,3$ $r^2 = 0,5621$	$y = -0,2484x + 85,481$ $r^2 = 0,7259$
NC9	$y = -0,1542x + 78,223$ $r^2 = 0,8373$	$y = -0,1729x + 90,042$ $r^2 = 0,7181$	$y = -0,2778x + 88,393$ $r^2 = 0,8665$	$y = -0,1289x + 80,172$ $r^2 = 0,7409$	$y = -0,144x + 92,122$ $r^2 = 0,8871$
NC19	$y = -0,1498x + 78,651$ $r^2 = 0,5524$	$y = -0,1454x + 79,964$ $r^2 = 0,811$	$y = -0,2317x + 67,591$ $r^2 = 0,761$	$y = -0,0612x + 75,346$ $r^2 = 0,8332$	$y = -0,1074x + 88,709$ $r^2 = 0,806$
NCG4	$y = -0,1677x + 85,707$ $r^2 = 0,7395$	$y = -0,1175x + 70,463$ $r^2 = 0,6474$	$y = -0,1943x + 71,371$ $r^2 = 0,6895$	$y = -0,098x + 82,725$ $r^2 = 0,6255$	$y = -0,1017x + 74,289$ $r^2 = 0,716$
NCG9	$y = -0,1103x + 73,096$ $r^2 = 0,7754$	$y = -0,1563x + 79,183$ $r^2 = 0,858$	$y = -0,1952x + 69,946$ $r^2 = 0,7588$	$y = -0,1572x + 81,742$ $r^2 = 0,7726$	$y = -0,1234x + 87,504$ $r^2 = 0,6107$
NCG19	$y = -0,2204x + 68,156$ $r^2 = 0,8367$	$y = -0,1166x + 66,914$ $r^2 = 0,6423$	$y = -0,213x + 74,932$ $r^2 = 0,7449$	$y = -0,0679x + 82,073$ $r^2 = 0,6309$	$y = -0,1734x + 87,275$ $r^2 = 0,8177$

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19.

Table 19: Shapiro-Wilk normality test done to the ATP values of each treatment.

	N	Statistic
C	12	0.557
NAM4	12	0.507
NAM9	11	0.345
NAM19	11	0.366
NC4	11	0.644
NC9	12	0.709
NC19	12	0.512
NCG4	12	0.401
NCG9	12	0.702
NCG19	11	0.630

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and

NCG19, treatment NCG in dilution 1:19. Values obtained at 12 h, 1 and 3 days, in the four pools, grouped according to the different combinations of independent variables (extender and dilution). A significance (Sig.) value below 0.05 indicates the absence of a normal distribution. The N column corresponds to the sample size value. Degrees of freedom = 11. *Normality is statistically significant, $p > 0.05$.

Table 20: Shapiro-Wilk normality test done to the MDA values obtained in each treatment.

	Statistic
C	0.954*
NAM4	0.778
NAM9	0.822*
NAM19	0.856*
NC4	0.931*
NC9	0.948*
NC19	0.837*
NCG4	0.987*
NCG9	0.751
NCG19	0.825*

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and

NCG19, treatment NCG in dilution 1:19. Values obtained in days 3, 6 and 9, and in both pools. Significance (Sig.) greater than 0.05 indicates the existence of normal distribution. N = 6. Degrees of freedom = 6.

*Normality is statistically significant, $p > 0.05$.

Table 21: Shapiro-Wilk normality test done to the CFU/ml values obtained in each treatment, medium and day.

		Day 5	Day 10
		Statistic	
TSA	C	0.750	0.862
	NAM	0.500	0.790
	NC	0.500	0.750
	NCG	0.500	0.750
PS	C	0.750	0.500
	NAM	0.750	0.947*
	NC	0.500	0.500
	NCG	0.500	0.750

Note: C, control; NAM, treatment with extender NAM; NC, treatment with extender NC; NCG, treatment with extender NCG; TSA, Trypticase Soya Agar medium; PS, CN Pseudomonas Agar medium. Values from three pools. The N column corresponds to the sample size value. Degrees of freedom = 3. * Normality is statistically significant, $p > 0.05$.

Table 22: Shapiro-Wilk normality test done to the values obtained in all the pools for VIAB, and the respective values of MOT, VCL and LIN.

	Statistic
VIAB	0.968
MOT	0.931
VCL	0.994*
LIN	0.967

Note: VIAB, percentage of viable cells; MOT, percentage of motile cells; VCL, curvilinear velocity; LIN, linearity. Values of all the treatments and days of evaluation, grouped according to parameter. Values of zero were excluded. N = 293. Degrees of freedom = 293. *Normality is statistically significant, $p > 0.05$.

Table 23: Shapiro-Wilk normality test done between ATP and the respective values of MOT, VCL and LIN, as well as ATP and VIAB.

	ATP x MOT, VCL, LIN	ATP x VIAB
	Statistic	Statistic
ATP	0.836	0.833
MOT	0.874	
VCL	0.462	
LIN	0.716	
VIAB		0.772

Note: ATP, adenosine triphosphate; MOT, percentage of motile cells; VCL, curvilinear velocity; LIN, linearity; VIAB, percentage of viable cells. Values above zero obtained between 0 and 3 days of storage, in five pools. A significance (Sig.) value below 0.05 indicates the absence of a normal distribution. For ATP x MOT, VCL and LIN, N = 35 and degrees of freedom = 35. For ATP x VIAB, N = 32 and degrees of freedom = 32. *Normality is statistically significant, $p > 0.05$.

Table 24: Shapiro-Wilk normality test done between MDA and the respective values of MOT, VCL and LIN, as well as MDA and VIAB.

	MDA x MOT, VCL, LIN	MDA x VIAB
	Statistic	Statistic
MDA	0.707	0.707
MOT	0.809	
VCL	0.957	
LIN	0.891	
VIAB		0.866

Note: MDA, malondialdehyde; MOT, percentage of motile cells; VCL, curvilinear velocity; LIN, linearity; VIAB, percentage of viable cells. For MDA x MOT, VCL, LIN, N = 33 and degrees of freedom = 33. For MDA x VIAB, N = 57 and degrees of freedom = 57. *Normality is statistically significant, $p > 0.05$.

Table 25: Statistic values of the Shapiro-Wilk normality test for the different assessments of correlation, as indicated in the first row.

	CFU/ml TSA x MOT, VCL, LIN	CFU/ml TSA x VIAB	CFU/ml PS x MOT, VCL, LIN	CFU/ml PS x VIAB
TSA	0.999	0.839*		
PS			0.750*	0.820
MOT	0.896		0.790	
VCL	0.997		0.984	
LIN	0.996		0.968	
VIAB		0.861		0.908

Note: TSA, Trypticase Soya Agar medium; PS, CN Pseudomonas Agar medium; MOT, percentage of motile cells; VCL, curvilinear velocity; LIN, linearity; VIAB, percentage of viable cells. The values corresponding to the same days of inoculation and of all the pools, treatments and days were used for each test. For CFU/ml TSA x MOT, VCL and LIN, N = degrees of freedom = 3. For CFU/ml TSA x VIAB, N = degrees of freedom = 10. For CFU/ml PS x MOT, VCL and LIN, N = degrees of freedom = 3. For CFU/ml PS x VIAB, N = degrees of freedom = 6. *Normality is statistically significant, $p < 0.05$.

Annex D: Concentration values and data treatment.

Table 26: Concentration values obtained for the control sample, in each and in all of the pools tested.

Pool	Mean \pm S.E.M.
1	26.30 \pm 2.54
2	46.40 \pm 1.74
3	46.87 \pm 4.14
4	23.83 \pm 3.68
5	35.87 \pm 8.37
6	62.47 \pm 5.89
7	62.40 \pm 3.29
Total	43.45 \pm 2.80

Note: Values shown as mean \pm standard error of the mean. Values expressed in 10^9 spz/ml. For each pool, N = 6.

Total N = 42.

Annex E: Motility values and data treatment.

Table 27: Mean \pm standard error of the mean of the percentage of motile cells for each treatment and day.

	C	NAM 4	NAM 9	NAM 19	NC 4	NC 9	NC 19	NCG 4	NCG 9	NCG 19
0	73.12 \pm 3.48									
0.5	42.63 \pm 4.58	67.13 \pm 3.47	55.05 \pm 8.54	47.79 \pm 9.60	78.40 \pm 1.52	76.07 \pm 1.66	67.91 \pm 3.78	74.87 \pm 2.24	75.37 \pm 5.48	64.99 \pm 3.16
1	7.34 \pm 2.15	49.36 \pm 4.55	46.56 \pm 6.95	29.31 \pm 5.20	74.32 \pm 3.62	68.07 \pm 2.20	52.00 \pm 4.09	68.64 \pm 2.92	68.32 \pm 2.45	41.12 \pm 5.14
1.5	0.48 \pm 0.17	28.76 \pm 3.05	28.71 \pm 5.01	14.71 \pm 4.21	73.83 \pm 2.96	65.57 \pm 1.95	36.48 \pm 5.87	55.86 \pm 3.25	50.59 \pm 5.31	32.83 \pm 6.59
2	0	15.14 \pm 3.33	8.69 \pm 2.97	3.15 \pm 1.18	46.11 \pm 1.84	51.13 \pm 4.44	22.35 \pm 3.85	46.92 \pm 3.56	41.19 \pm 3.93	20.18 \pm 5.19
3	0	5.47 \pm 1.44	2.00 \pm 0.59	1.05 \pm 0.36	45.22 \pm 4.54	34.56 \pm 3.68	25.39 \pm 6.09	33.38 \pm 2.41	36.39 \pm 4.29	18.76 \pm 5.96
4	0	4.52 \pm 1.24	1.62 \pm 0.67	0	38.20 \pm 5.18	30.52 \pm 7.07	16.61 \pm 4.57	29.95 \pm 2.47	21.40 \pm 3.32	7.19 \pm 1.51
5	0	3.09 \pm 1.18	0.39 \pm 0.19	0	31.09 \pm 3.45	34.80 \pm 4.81	6.89 \pm 2.02	21.82 \pm 2.51	15.14 \pm 2.62	6.92 \pm 2.06
6	0	2.09 \pm 0.87	0.54 \pm 0.55	0	28.54 \pm 5.70	15.06 \pm 1.99	6.91 \pm 3.29	17.99 \pm 2.65	3.57 \pm 0.91	4.59 \pm 1.83

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19. Values from five pools, ten treatments, and all the days of motility assessment. N = 5.

Table 27 (cont.): Mean \pm standard error of the mean of the percentage of motile cells for each treatment and day.

	C	NAM 4	NAM 9	NAM 19	NC 4	NC 9	NC 19	NCG 4	NCG 9	NCG 19
7	0	0.60 \pm 0.19	0.09 \pm 0.04	0	24.22 \pm 5.35	15.01 \pm 2.87	7.45 \pm 1.99	9.86 \pm 1.88	3.22 \pm 1.29	1.64 \pm 0.73
8	0	0.25 \pm 0.07	0	0	19.48 \pm 5.18	8.11 \pm 1.59	2.29 \pm 0.64	7.94 \pm 1.63	4.57 \pm 2.00	1.06 \pm 0.47
9	0	0.13 \pm 0.06	0	0	13.00 \pm 2.85	4.90 \pm 0.93	1.81 \pm 0.68	6.88 \pm 1.72	1.78 \pm 0.80	0.05 \pm 0.02
10	0	0	0	0	7.10 \pm 1.80	0.43 \pm 0.09	0.65 \pm 0.18	2.86 \pm 0.79	0.05 \pm 0.02	0
11	0	0	0	0	2.11 \pm 0.88	0	0	0	0	0
12	0	0	0	0	1.13 \pm 0.51	0	0	0	0	0
13	0	0	0	0	1.81 \pm 0.81	0	0	0	0	0
14	0	0	0	0	0.95 \pm 0.42	0	0	0	0	0
15	0	0	0	0	0.41 \pm 0.18	0	0	0	0	0

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19. Values from five pools, ten treatments, and all the days of motility assessment. N = 5.

Table 28: Results of the Mann-Whitney U of all the pairs tested of the various combinations of extenders, and of dilutions, for percentage of motility (MOT) values.

	Pairs (1 x 2)	Mann-Whitney U	Mean rank (1)	Mean rank (2)
Extender	C x NAM	1689.000*	60.53	92.04
	C x NC	716.500*	38.92	107.69
	C x NCG	873.000*	42.40	106.53
	NAM x NC	3802.500*	92.43	160.83
	NAM x NCG	4568.500*	98.76	155.16
	NC x NCG	7775.500*	145.40	125.60
Dilution	C x 1:4	751.000*	39.69	107.44
	C x 1:9	907.500*	43.17	102.41
	C x 1:19	1620.000*	59.00	96.84
	1:4 x 1:9	7890.000	137.56	126.14
	1:4 x 1:19	5330.000*	156.52	104.14
	1:9 x 1:19	5815.000*	147.07	109.93

Note: The mean rank of each element of the pair is shown, and highlighted in bold when the mean rank corresponds to an element of the pair with statistically significantly higher MOT values. Values from seven pools. $N_C = 45$, $N_{NAM} = 121$, $N_{NC} = 135$, $N_{NCG} = 135$, $N_{1:4} = 135$, $N_{1:9} = 128$ and $N_{1:19} = 128$. * Difference is statistically significant, $p < 0.05$.

Table 29: Mean \pm standard error of the mean of the curvilinear velocity for each treatment and day.

	C	NAM4	NAM9	NAM19	NC4	NC9	NC19	NCG4	NCG9	NG19
0	140.90 \pm 7.75									
0.5	92.34 \pm 5.76	139.17 \pm 5.87	127.91 \pm 7.31	99.62 \pm 10.84	127.06 \pm 3.45	141.91 \pm 3.54	140.73 \pm 5.83	129.56 \pm 2.97	130.05 \pm 6.80	132.36 \pm 1.93
1	77.95 \pm 23.63	118.19 \pm 3.98	112.98 \pm 8.27	98.78 \pm 8.04	124.65 \pm 5.07	135.15 \pm 5.88	110.83 \pm 6.85	125.68 \pm 5.88	122.41 \pm 6.51	105.24 \pm 8.39
1.5	9.40 \pm 0.13	102.05 \pm 3.19	90.13 \pm 5.45	70.90 \pm 18.94	114.59 \pm 7.93	116.95 \pm 3.09	117.80 \pm 6.33	113.72 \pm 5.42	113.68 \pm 6.90	89.56 \pm 12.89
2		69.98 \pm 6.94	74.06 \pm 11.92	97.35 \pm 11.96	103.21 \pm 4.29	118.56 \pm 4.67	89.90 \pm 4.21	108.60 \pm 5.89	108.78 \pm 7.10	109.00 \pm 6.35
3		76.09 \pm 7.90	69.07 \pm 7.89	57.44 \pm 35.15	113.24 \pm 3.79	109.60 \pm 4.41	114.68 \pm 6.47	106.03 \pm 4.87	111.53 \pm 5.67	98.48 \pm 15.19
4		62.57 \pm 11.44	86.33 \pm 12.07		107.56 \pm 8.83	116.60 \pm 7.86	94.00 \pm 6.85	110.47 \pm 7.30	105.80 \pm 5.10	79.35 \pm 2.29
5		68.03 \pm 11.34			94.10 \pm 7.31	122.33 \pm 5.40	87.77 \pm 8.12	102.10 \pm 8.32	95.22 \pm 4.85	82.60 \pm 6.04
6		40.41 \pm 8.63			88.25 \pm 10.90	114.58 \pm 5.61	61.28 \pm 18.61	77.91 \pm 6.58	50.48 \pm 7.87	68.73 \pm 11.99
7		54.12 \pm 17.93			104.50 \pm 11.27	88.48 \pm 8.41	77.52 \pm 22.22	83.00 \pm 5.53	72.59 \pm 21.36	
8		40.93 \pm 3.14			97.22 \pm 12.04	64.58 \pm 10.07	108.12 \pm 7.42	75.45 \pm 1.55	60.29 \pm 39.15	
9					81.84 \pm 13.45	104.88 \pm 7.94	84.58 \pm 0.86	85.47 \pm 10.26		

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19. Values from five pools, ten treatments, and all the days of motility assessment. N = 5.

Table 29 (cont.): Mean \pm standard error of the mean of curvilinear velocity for each treatment and day.

	C	NAM4	NAM9	NAM19	NC4	NC9	NC19	NCG4	NCG9	NG19
10					71.18 \pm 13.14	63.40 \pm 5.06	17.17 \pm 7.85		78.35 \pm 15.76	
11					82.82 \pm 16.42					

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19. Values from five pools, ten treatments, and all the days of motility assessment. N = 5.

Table 30: Results of the Mann-Whitney U of all the pairs tested of the various combinations of extenders, and of dilutions, for curvilinear velocity (VCL) values.

	Pairs (1 x 2)	Mann-Whitney U	Mean rank (1)	Mean rank (2)
Extender	C x NAM	1607.000*	58.71	92.72
	C x NC	677.500*	38.06	107.98
	C x NCG	789.000*	40.53	107.16
	NAM x NC	4250.000*	96.12	157.52
	NAM x NCG	4806.000*	100.72	153.40
	NC x NCG	8304.000	141.49	129.51
Dilution	C x 1:4	710.000*	38.78	107.74
	C x 1:9	835.500*	41.57	102.97
	C x 1:19	1528.000*	56.96	97.56
	1:4 x 1:9	8402.000	133.76	130.14
	1:4 x 1:19	5975.000*	151.74	111.18
	1:9 x 1:19	5985.000*	145.74	111.26

Note: The mean rank of each element of the pair is shown, and highlighted in bold when the mean rank corresponds to an element of the pair with statistically significantly higher VCL values. Values from seven pools. $N_C = 45$, $N_{NAM} = 133$, $N_{NC} = 130$, $N_{NCG} = 127$, $N_{1:4} = 135$, $N_{1:9} = 128$, $N_{1:19} = 128$. *Difference is statistically significant, $p < 0.05$.

Table 31: Mean \pm standard error of the mean of the percentage of linearity for each treatment and day.

	C	NAM4	NAM9	NAM19	NC4	NC9	NC19	NCG4	NCG9	NCG19
0	56.61 \pm 2.16									
0.5	64.37 \pm 1.62	50.71 \pm 0.62	58.04 \pm 2.32	41.67 \pm 5.88	52.41 \pm 0.97	51.77 \pm 0.75	50.10 \pm 0.77	55.17 \pm 0.92	55.45 \pm 1.29	66.25 \pm 2.06
1	58.45 \pm 3.49	58.99 \pm 1.04	55.61 \pm 0.78	59.77 \pm 2.42	55.86 \pm 1.06	55.02 \pm 0.91	57.69 \pm 0.68	55.47 \pm 0.59	56.03 \pm 0.79	57.70 \pm 0.61
1.5	26.60 \pm 7.18	58.11 \pm 1.37	55.08 \pm 1.23	46.52 \pm 6.16	52.99 \pm 1.28	52.70 \pm 0.64	55.04 \pm 1.38	55.39 \pm 1.15	53.32 \pm 0.90	54.39 \pm 0.83
2		56.58 \pm 2.85	37.78 \pm 4.80	61.15 \pm 0.72	59.47 \pm 1.00	60.11 \pm 0.75	54.62 \pm 2.77	59.20 \pm 1.64	54.44 \pm 0.54	58.45 \pm 1.96
3		51.65 \pm 2.39	60.16 \pm 2.19	32.83 \pm 17.94	59.20 \pm 0.77	59.63 \pm 1.08	64.18 \pm 1.48	59.00 \pm 0.76	63.56 \pm 0.39	53.90 \pm 4.41
4		56.88 \pm 5.54	53.41 \pm 15.57		60.20 \pm 1.33	59.96 \pm 2.04	64.39 \pm 1.49	60.97 \pm 1.35	65.10 \pm 2.33	59.18 \pm 2.76
5		59.66 \pm 2.63			64.78 \pm 0.87	59.57 \pm 0.66	56.34 \pm 4.82	62.41 \pm 1.23	61.44 \pm 3.20	49.83 \pm 6.32
6		45.27 \pm 7.27			55.15 \pm 3.81	60.08 \pm 1.27	89.25 \pm 17.82	64.71 \pm 3.96	52.21 \pm 4.35	44.95 \pm 3.75
7		56.82 \pm 3.14			63.13 \pm 0.55	58.44 \pm 1.97	54.36 \pm 2.77	68.20 \pm 2.13	58.86 \pm 3.97	
8		45.98 \pm 13.42			61.08 \pm 2.19	55.16 \pm 2.79	58.92 \pm 0.03	64.56 \pm 2.32	38.79 \pm 17.28	
9					60.53 \pm 1.61	60.52 \pm 2.36	47.64 \pm 11.48	61.68 \pm 0.28		

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and

NCG19, treatment NCG in dilution 1:19. Values from five pools, ten treatments, and all the days of motility assessment. N = 5.

Table 31 (cont.): Mean \pm standard error of the mean of the percentage of linearity for each treatment and day. Values from five pools, ten treatments, and all the days of motility assessment. N = 5.

	C	NAM4	NAM9	NAM19	NC4	NC9	NC19	NCG4	NCG9	NCG19
10					50.42 \pm 6.53	40.74 \pm 3.55	32.88 \pm 8.69	59.54 \pm 4.78		
11					61.43 \pm 6.78					

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19. Values from five pools, ten treatments, and all the days of motility assessment. N = 5.

Table 32: Results of the Mann-Whitney U of all the pairs tested of the various combinations of extenders, and of dilutions, for the percentage of linearity (LIN) values.

	Pairs (1 x 2)	Mann-Whitney U	Mean rank (1)	Mean rank (2)
Extender	C x NAM	1745.000*	61.78	91.58
	C x NC	1019.500*	45.66	105.45
	C x NCG	1068.000*	46.73	105.09
	NAM x NC	5061.000*	102.83	151.51
	NAM x NCG	5162.000*	103.66	150.76
	NC x NCG	9107.000	135.54	135.46
Dilution	C x 1:4	982.000*	44.82	105.73
	C x 1:9	1126.500*	48.03	100.70
	C x 1:19	1724.000*	61.31	96.03
	1:4 x 1:9	7540.000	140.15	123.41
	1:4 x 1:19	5763.000*	153.31	109.52
	1:9 x 1:19	6380.000*	142.66	114.34

Note: C, control. The mean rank of each element of the pair is shown, and highlighted in bold when the mean rank corresponds to an element of the pair with statistically significantly higher LIN values. Values from seven

pools. $N_C = 45$, $N_{NAM} = 121$, $N_{NC} = 135$, $N_{NCG} = 135$, $N_{1:4} = 135$, $N_{1:9} = 128$, $N_{1:19} = 128$. *Difference is statistically significant, $p < 0.05$.

Annex F: Viability values and data treatment.

Table 33: Mean \pm standard error of the mean of the percentage of viability for each treatment and day.

	C	NAM 4	NAM 9	NAM 19	NC 4	NC 9	NC 19	NCG 4	NCG 9	NCG 19
0	74.24 \pm 33.20									
0.5	63.06 \pm 28.20	79.65 \pm 35.62	76.31 \pm 34.13	73.46 \pm 32.85	79.29 \pm 35.46	80.08 \pm 35.81	76.04 \pm 34.01	71.61 \pm 35.80	65.51 \pm 32.76	65.80 \pm 29.43
	55.88 \pm 24.99	62.81 \pm 28.09	62.72 \pm 28.05	53.02 \pm 23.71	58.78 \pm 26.29	68.24 \pm 30.52	66.65 \pm 29.81	67.56 \pm 39.01	67.37 \pm 30.13	76.69 \pm 34.30
1.5	59.00 \pm 26.38	57.20 \pm 25.58	66.79 \pm 29.87	36.30 \pm 16.24	57.40 \pm 28.70	74.49 \pm 33.31	70.16 \pm 31.38	71.51 \pm 31.98	70.49 \pm 31.53	68.69 \pm 30.72
	56.35 \pm 25.20	62.46 \pm 27.93	64.49 \pm 28.84	38.26 \pm 17.11	68.39 \pm 30.58	68.51 \pm 30.64	67.04 \pm 29.98	62.32 \pm 27.87	65.31 \pm 29.21	68.10 \pm 30.45
3	61.14 \pm \pm 27.34	48.46 \pm 21.67	38.87 \pm 17.38	21.70 \pm 9.70	67.89 \pm 30.36	77.79 \pm 34.79	64.48 \pm 28.83	65.47 \pm 29.28	63.26 \pm 28.29	67.34 \pm 30.11
	57.36 \pm 25.65	34.57 \pm 15.46	27.25 \pm 12.19	15.59 \pm 6.97	63.25 \pm 28.29	66.53 \pm 29.75	61.08 \pm 27.32	62.11 \pm 27.77	65.17 \pm 29.15	56.51 \pm 25.27
5	60.26 \pm 30.13	36.28 \pm 16.22	29.05 \pm 12.99	26.72 \pm 11.95	62.35 \pm 31.17	61.41 \pm 30.70	61.60 \pm 27.55	68.86 \pm 48.69	64.81 \pm 28.98	61.28 \pm 27.40
	48.35 \pm 21.62	34.60 \pm 15.47	18.10 \pm 8.10	16.27 \pm 7.28	51.45 \pm 25.73	66.00 \pm 33.00	54.13 \pm 24.21	50.99 \pm 25.49	65.24 \pm 29.17	47.02 \pm 21.03
7	39.33 \pm 22.71	16.42 \pm 7.34	17.76 \pm 8.88	16.06 \pm 8.03	42.11 \pm 18.83	55.37 \pm 24.76	63.23 \pm 28.28	61.42 \pm 27.47	51.46 \pm 23.01	44.21 \pm 19.77

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and

NCG19, treatment NCG in dilution 1:19. Values from five pools, ten treatments, in all the days of viability evaluation. N = 5.

Table 33 (cont.): Mean \pm standard error of the mean of the percentage of viability for each treatment and day.

	C	NAM 4	NAM 9	NAM 19	NC 4	NC 9	NC 19	NCG 4	NCG 9	NCG 19
9		27.05 \pm 13.52			52.58 \pm 23.51	53.73 \pm 24.03	47.44 \pm 23.81	51.39 \pm 22.98	45.49 \pm 20.34	55.95 \pm 28.98
11					29.95 \pm 13.39	40.29 \pm 20.14	45.09 \pm 22.54	56.08 \pm 28.04	44.27 \pm 19.80	48.10 \pm 27.77
13					29.33 \pm 16.93	43.73 \pm 25.25		40.75 \pm 23.53	32.10 \pm 18.53	
15								32.34 \pm 18.67		

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19. Values from five pools, ten treatments, in all the days of viability evaluation. N = 5.

Table 34: Results of the Mann-Whitney U of all the pairs tested of the various combinations of extenders, for percentage of viability (VIAB) values.

	Pairs (1 x 2)	Mann-Whitney U	Mean rank (1)	Mean rank (2)
Extender	C x NAM	1873.500*	4615.50	10784.50
	C x NC	1908.500*	2811.50	12066.50
	C x NCG	1996.000*	2899.00	11466.00
	NAM x NC	4263.000*	13174.00	21542.00
	NAM x NCG	4425.500*	13336.50	20593.50
	NC x NCG	7540.500	17484.50	15668.50

Note: The mean rank of each element of the pair is shown, and highlighted in bold when the mean rank corresponds to an element of the pair with statistically significantly higher VIAB values. The N column

corresponds to the sample size value. $N_C = 42$, $N_{NAM} = 133$, $N_{NC} = 130$ and $N_{NCG} = 127$. *Difference is statistically significant, $p < 0.05$.

Annex G: ATP values and data treatment.

Table 35: Mean \pm standard error of the mean of the ATP value for each treatment and day.

	C	NAM 4	NAM 9	NAM 19	NC 4	NC 9	NC 19	NCG 4	NCG 9	NCG 19
0	2.16 \pm 0.72									
0.5	0.14 \pm 0.05	0.29 \pm 0.08	0.81 \pm 0.40	0.63 \pm 0.34	4.86 \pm 1.61	4.38 \pm 0.68	7.20 \pm 2.18	1.41 \pm 0.71	1.43 \pm 0.42	6.01 \pm 2.61
1	0.02 \pm 0.01	0	0	0	1.63 \pm 0.76	0.29 \pm 0.11	0	1.26 \pm 0.63	2.01 \pm 1.00	3.81 \pm 1.90
3	0.01 \pm 0.00	0	0	0	0	0	0	0.56 \pm 0.28	1.45 \pm 0.73	2.82 \pm 1.41

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19. Values from five pools, in $\mu\text{mol ATP per } 10^{12}$ spermatozoa. N = 4.

Table 36: Results of the Mann-Whitney U of all the pairs tested of the various combinations of extenders, for ATP values.

	Pairs (1 x 2)	Mann-Whitney U	Mean rank (1)	Mean rank (2)
Extender	C x NAM	173.000	26.08	22.59
	C x NC	186.000	22.00	24.69
	C x NCG	187.000	22.08	24.66
	NAM x NC	464.500*	31.16	38.73
	NAM x NCG	469.500*	31.31	38.59
	NC x NCG	601.500	35.81	35.19

Note: The mean rank of each element of the pair is shown, and highlighted in bold when the mean rank corresponds to an element of the pair with statistically significantly higher ATP values. $N_C = 12$, $N_{NAM} = 34$, $N_{NC} = N_{NCG} = 35$. * Difference is statistically significant, $p < 0.05$.

Annex H: MDA values and data treatment.

Table 37: Mean \pm standard error of the mean of the MDA value for each treatment and day.

	C	NAM 4	NAM 9	NAM 19	NC 4	NC 9	NC 19	NCG 4	NCG 9	NCG 19
0	0.00 \pm 0.00									
3	6.17 \pm 4.36	9.83 \pm 6.95	5.36 \pm 3.79	11.26 \pm 7.96	9.74 \pm 6.89	21.32 \pm 15.08	14.22 \pm 10.05	14.15 \pm 10.01	23.15 \pm 16.37	10.34 \pm 7.31
6	19.70 \pm 13.93	19.31 \pm 13.65	38.06 \pm 26.91	46.89 \pm 33.16	20.93 \pm 14.80	33.21 \pm 51.40	72.68 \pm 51.40	20.62 \pm 14.58	51.49 \pm 36.41	109.63 \pm 77.52
9	20.18 \pm 14.27	9.93 \pm 7.02	45.13 \pm 91.91	16.56 \pm 11.71	30.21 \pm 21.36	21.77 \pm 28.77	40.69 \pm 28.77	15.81 \pm 11.18	29.10 \pm 20.57	24.14 \pm 17.07

Note: C, control; NAM4, treatment NAM in dilution 1:4; NAM9, treatment NAM in dilution 1:9; NAM19, treatment NAM in dilution 1:19; NC4, treatment NC in dilution 1:4; NC9, treatment NC in dilution 1:9; NC19, treatment NC in dilution 1:19; NCG4, treatment NCG in dilution 1:4; NCG9, treatment NCG in dilution 1:9; and NCG19, treatment NCG in dilution 1:19. Values from two pools, in $\mu\text{M MDA}/10^{10}$ cells. N = 2.

Table 38: Results of the Mann-Whitney U of all the pairs tested of the various combinations of dilutions, for MDA values.

	Pairs (1 x 2)	Mann-Whitney U	Mean rank (1)	Mean rank (2)
Dilution	C x 1:4	50.000	11.83	12.72
	C x 1:9	26.000	7.83	14.06
	C x 1:19	34.000	9.17	13.61
	1:4 x 1:9	83.000*	14.11	22.89
	1:4 x 1:19	110.000	15.61	21.39
	1:9 x 1:19	159.000	18.67	18.33

Note: The mean rank of each element of the pair is shown, and highlighted in bold when the mean rank corresponds to an element of the pair with statistically significantly higher VCL values. Values from two pools and from days 3, 6 and 9. $N_C = 6$, $N_{1:4} = 18$, $N_{1:9} = 18$, $N_{1:19} = 18$. *Difference is statistically significant, $p < 0.05$.

Annex I: Bacteriology values and data treatment.

Table 39: Results from the bacterial identification procedures, showing the percentage and number of isolates for each of the species that were found, including the ones that were not identified.

Species	Percentage	Number of isolates
Flavobacterium sp.	27.91%	12
Corynebacterium sp.	18.60%	8
Aeromonas sp.	23.26%	10
Pseudomonas sp.	2.33%	1
Vibrio sp.	2.33%	1
Unidentified	25.58%	11

Note: Three pools and TSA and PS plates were used, and isolates done at the seventh day of incubation. N = 43.

