

1 *Brief Report*

2 3 **Lipid Signature of Motile Human Sperm: Characterization of Sphingomyelin,** 4 **Ceramide and Phospholipids with a Focus on Very Long Polyunsaturated Fatty Acid** 5

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18 **Abstract:**

19 Sperm membrane lipids play a crucial role in male fertility, influencing sperm motility,
20 viability, and functional competence. This study provides a comprehensive characteriza-
21 tion of the phospholipid and sphingolipid composition in highly motile human sperma-
22 tozoa obtained through the swim-up method, a widely used technique in assisted repro-
23 ductive technology (ART). Using two-dimensional thin-layer chromatography and phos-
24 phorus analysis, we identified choline glycerophospholipids (CGP, 45%), ethanolamine
25 glycerophospholipids (EGP, 26%), and sphingomyelin (SM, 17%) as predominant phos-
26 pholipids, with minor components including cardiolipin, lysophospholipids, phosphati-
27 dylinositol, phosphatidylserine, phosphatidic acid, and neutral lipids. Gas chromatog-
28 raphy analysis of glycerophospholipids (GPL) revealed a high long chain (C20-C22) pol-
29 yunsaturated fatty acid (PUFA) content (46.3%), particularly docosahexaenoic acid (DHA,
30 22:6n-3), which was more abundant in CGP (46%) than EGP (26%). Sphingolipid analysis
31 indicated that ceramide (Cer) and SM shared similar fatty acid profiles due to their meta-
32 bolic relationship, with very long-chain (VLC) PUFA (\geq C26) being more prevalent in SM
33 (10%) than in Cer (6%). Additionally, argentation chromatography allowed the identifica-
34 tion of highly unsaturated VLCPUFA species in Cer, including 28:3n-6, 28:4n-6, and 30:4n-
35 6, which had not been previously quantified in motile human spermatozoa. Given the
36 essential function of sphingolipid metabolism in spermatogenesis, capacitation, and acro-
37 somal exocytosis, our findings suggest that the balance of VLCPUFA-containing SM and
38 Cer emerges as a key factor in sperm performance and fertilization potential. This study
39 provides novel insights into the lipid signature of human sperm and highlights the rele-
40 vance of membrane lipid remodeling for male fertility and ART outcomes.

41
42 **Keywords:** sphingolipids; ceramide; sphingomyelin; phospholipids; VLPUFA; human
43 sperm
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1. Introduction

The importance of the lipid composition of the sperm plasma membrane lies in its unique characteristics, which are closely related to the ability of these fully differentiated cells to successfully fertilize the oocyte [1]. In both murine and large mammal spermatozoa, the high content of C20-C22 polyunsaturated fatty acids in their glycerophospholipids (GPL) [2-4] and the presence of sphingolipids (SLs) with very-long-chain polyunsaturated fatty acids \geq C26 (VLCPUFA) stand out [5, 6]. It is worth noting that these VLCPUFA are exclusively bound to sperm sphingomyelins (SM) and not to other phosphoglycerolipids [5]. Notably, SM with VLCPUFA has been found to be exclusively located in the heads of bull and rat spermatozoa [6, 7]. The VLCPUFAs are produced through the elongation and desaturation of n-3 and n-6 dietary C18-C22 PUFA by a series of elongase and desaturase enzymes that involve the sequential action of ELOVL2, ELOVL5, and especially ELOVL4. The latter is expressed in a limited number of tissues, including the skin, brain, retina, Meibomian glands, and testes [8, 9].

In the rat testicular environment, Santiago Valtierra et al. (2018) demonstrated that the expression and activity of ELOVL4 correlate with the abundance of VLCPUFA. These unusual molecular species are incorporated into SLs compounds through the enzymatic action of ceramide synthase 3 (CerS3) in mice testis [10]. When CerS3 is conditionally deleted, it results in an almost-complete absence of SL products harboring VLCPUFA, increased apoptosis during meiosis, and spermatogenic arrest that culminates in infertility [11].

The requirement of VLCPUFA in human fertility is underscored by observations correlating diminished levels of these fatty acids in sperm with reduced quantity and quality of spermatozoa, as reported by Craig et al. in 2019 [12]. Although the majority of these investigations have focused on the total semen samples from various mammalian species, including human, our study aims to elucidate the fatty acid composition of main GPL, SM, and Cer, within the membranes of human sperm isolated by swim up commonly used for in vitro fertilization treatment. This issue has not yet been explored. Our findings have prompted us to investigate the relationship between lipids and the human sperm functions we previously studied. Maintaining lipid homeostasis, which is crucial for cell health, significantly impacts sperm viability, development, and performance during fertilization.

2. Results

2.1. Phospholipid composition and fatty acid of total glycerophospholipids (GPL) in non-capacitated motile human sperm

To assess the qualitative and quantitative phospholipid profile of motile non-capacitated human sperm we first resolved the lipids in a two-dimensional TLC (Figure 1A). After the exposure of the plates to iodine vapors, we transferred the silica gel containing each lipid class to different tubes. Then, we quantified the phospholipids by phosphorus analysis. The results showed in Figure 1A and B revealed the lipid composition of sperm

membranes, which are choline glycerophospholipids (CGP), diphosphatidylglycerol, commonly known as “cardiolipin” (DPG), ethanolamine glycerophospholipids (EGP), free fatty acids (FFA), lysophosphatidylcholine (LPC), lysophosphoethanolamine (LPE), phosphatidylinositol (PI), phosphatidylserine (PS), sphingomyelin (SM), phosphatidic acid (PA), seminolipid (SL), neutral lipids (NL, e.g. cholesterol and their esters) and some unknown lipids indicated with a question mark.

As expected, we found that major PL were CGP, EGP, and SM that accounted on average 45%, 26% and 17%, respectively, and minor percentages (3% or less) of lysoPLs (LPC and LPE), PI, PS, DPG followed of negligible amounts of PA (Figure 1C). The CGP/EGP and CGP/SM were 1,73 and 2,64, respectively.

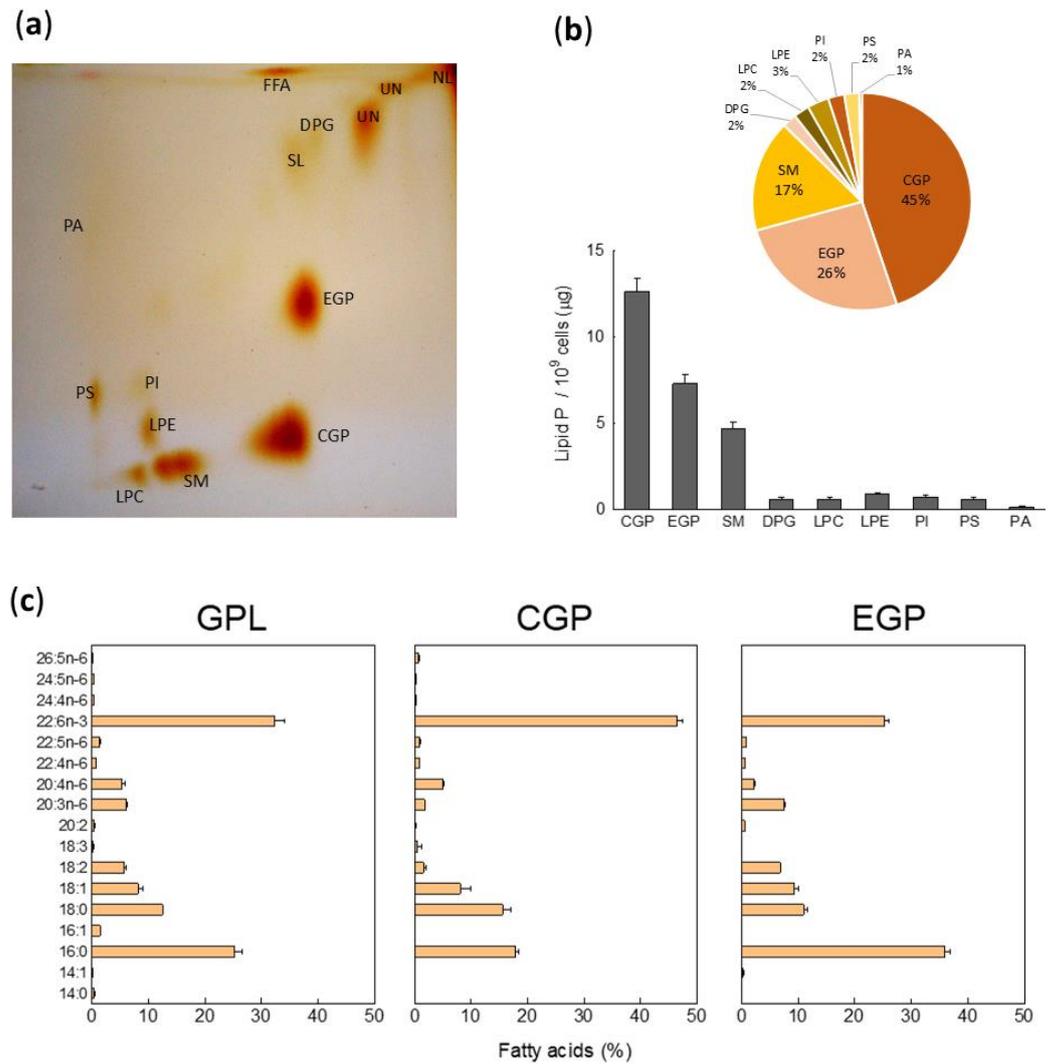


Figure 1. Qualitative phospholipid and fatty acid profile of human spermatozoa obtained by swim-up. (a) shows a representative TLC to resolve phospholipids. To perform this two-dimensional TLC two solvents were used, chloroform/methanol/ammonia (65:25:5) and chloroform/acetone/methanol/acetic acid/water (30:40:10:10:5 by vol). The lipids were located by exposure of the plates to iodine vapors. (b) Phospholipid composition (upper panel) and amounts per cell estimated from lipid phosphorus (bottom panel) of each phospholipid constituents of human spermatozoa. (c) Fatty acid composition of total glycerophospholipids (GPL), choline and ethanolamine glycerophospholipids (CGP and EGP, respectively). Other abbreviations: DPG, diphosphatidylglycerol (cardiolipin); FFA, free fatty acids; LPC, lysophosphatidylcholine; LPE, lysophosphoethanolamine; PI, phosphatidylinositol; PS, phosphatidylserine; SM, sphingomyelin; PA, phosphatidic acid ; SL, Seminolipid; UN, Unknown lipids.

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Then, aliquots of isolated total PL from human sperm membranes were analyzed by GC to determine the fatty acid composition of the GPL. As shown in Figure 1C, saturated FA represented 37.9 ± 1.5 %, being the 16:0 and 18:0 the most prevalent in human sperm samples. Further, we found MFA (9.5 ± 1.1 %, mainly 18:1) and DFA (5.9 ± 0.4 %, mainly 18:2). Strikingly, PUFA constituted 46.3 % of the total sperm GPL fatty acids and as observed in the Figure 1C, 22:6n-3 is the most abundant PUFA (32%), followed by minor amounts of 20:3n-6 and 20:4n-6. In addition, we found that only 0.5% of the total FA are C24-26 VLCPUFA in human sperm GPL. This fatty acid composition pattern, with some differences, was similar in major GPL, CGP and EGP. Notably, the percentage of 22:6n-3 in CGP reached 46%, while in EGP was 26% (Figure 1C). The latter was characterized by high amounts of molecular species of 16:0, about 36% of total fatty acids (Figure 1C).

2.2. Fatty acid composition in sphingomyelin (SM) and ceramide (Cer) from human motile spermatozoa

Our investigation was particularly directed towards the fatty acid composition of the SM and specially Cer molecular species isolated from the membranes of human motile spermatozoa obtained via swim up. In Figure 2, representative chromatograms of Cer and SM are shown, demonstrating that both lipids had the same fatty acids, which is expected due to their metabolic relationship. The total content of SM per unit of lipid phosphorus is 6.5 times greater than that of Cer (Figure 2B), and the latter exhibited the same molecular species of VLCPUFAs with \geq C26 carbon atoms as SM. Furthermore, when comparing the proportions of fatty acid groups, no differences were observed between the two lipids, with saturated (S), monoenoic (M), and dienoic (D) fatty acids, ranging from 56-58%, 25-30%, and 6-8% respectively. However, VLCPUFAs (V) were proportionally more abundant in SM than in Cer, at 10% compared to 6%, respectively ($p < 0.05$).

To concentrate individual molecular species of \geq C26 VLCPUFAs from the minor Cer, we resolved the fatty acid methyl esters (FAME) using argentation TLC. This method allowed us to separate total FAME into discrete bands containing saturated, monoenoic, and dienoic fatty acids, as well as those containing trienoic, tetraenoic, and pentaenoic VLCPUFA series. Each molecular species was then identified by GC (Figure 2C). Notably, 28:3n-6, 28:4n-6, and 30:4n-6 were the most abundant VLCPUFAs found in human sperm Cer (Figure 2C and Figure 3). Additionally, the longest VLCPUFA detected (Figure 2C, lower panel) and quantified (Figure 3) was 32:4n-6.

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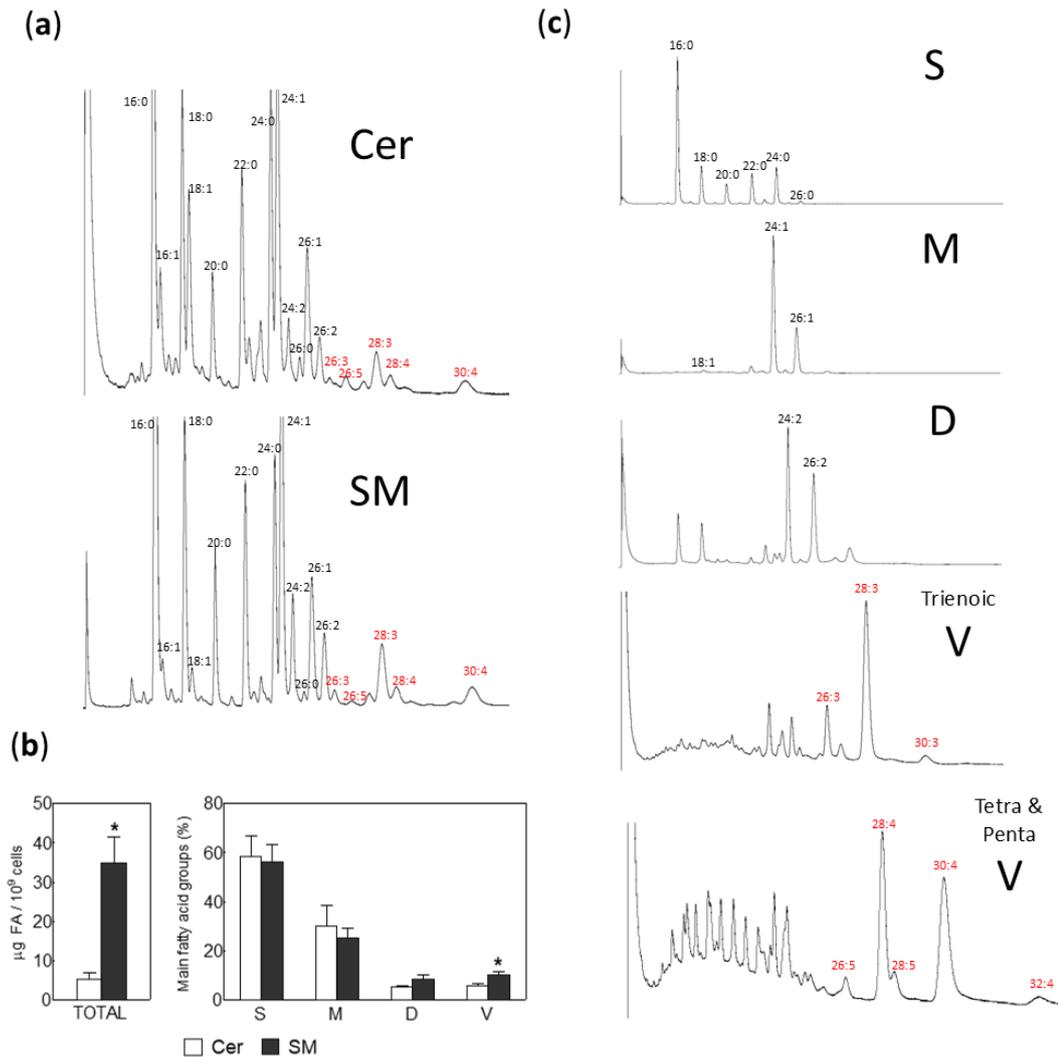


Figure 2. Ceramide (Cer) and sphingomyelin (SM) fatty acid profile of human spermatozoa obtained by swim-up. (A) shows a representative chromatograms showing the qualitative presence of $\geq\text{C}26$ VLCPUFA. (B) Content of Cer and SM. To compare them, the lipids were quantified by their fatty acids and expressed as μg per cells. Left panel: amounts of total fatty acids; right panel, percentages of main fatty acids, grouped into saturated (S), monoenoic (M), dienoic (D), and total VLCPUFA (V) (* $p < 0.01$). (C) Enrichment of different groups of fatty acids from sperm Cer, each chromatograms represent different fraction of fatty acids separated using AgNO_3 -TLC, VLCPUFA fractions were composed by tri-, tetra- and pentaenoic fatty acids.

In the quantitative comparison of fatty acid profiles of Cer and SM in human male gametes, we observed a predominance of saturated and monounsaturated fatty acids, ranging from 16 to 26 carbon atoms (Figure 2C and Figure 3). Notably, palmitic acid (16:0) was the most abundant, accounting for 18.0% of Cer and 32.5% of SM fatty acids. As shown in Figure 3, the proportion of 16:0 in SM is nearly double that of Cer. In contrast, Cer exhibited approximately double the proportions of the longer fatty acids 22:0 and 24:0 compared to SM, with 9.5% vs. 4.7% and 13.1% vs. 5.4%, respectively.

Within the individual molecular species of $\geq\text{C}26$ VLCPUFAs, no qualitative differences were found, but quantitative differences were observed. In both lipids, the predominant molecular species of ceramides (Cer) and sphingomyelins (SM) were 28:3n-6, 28:4n-

6, and 30:4n-6, although the proportions were slightly different: 4.4/1.6/2.3 in SM, while they were 2.5/1.0/1.4 in Cer (Figure 3).

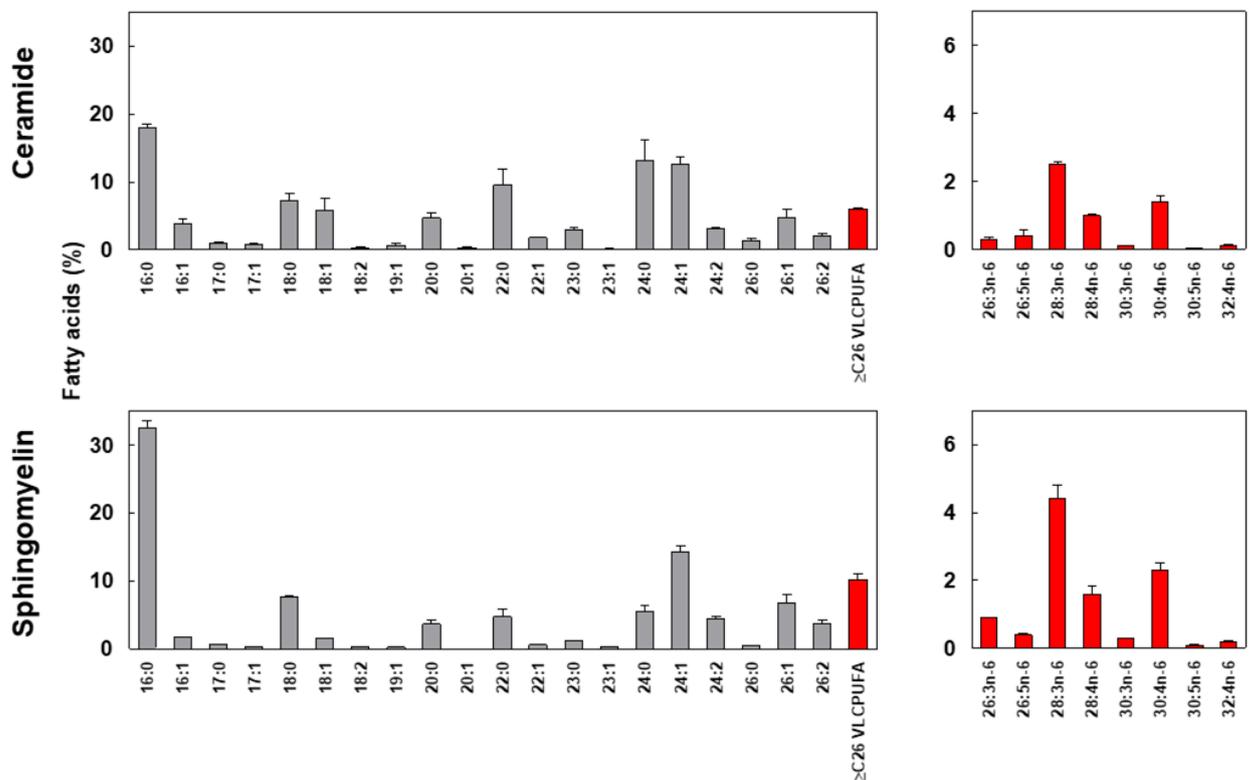


Figure 3. Cer and SM main individual fatty acids. Left panels: percentages of individual saturated, monoenoic and dienoic fatty acids in comparison with total \geq C26 VLCPUFA. Right panels: percentages of individual \geq C26 VLCPUFA are showed. Notably, in both Cer and SM from human sperm 28:3n-6, 28:4n-6, and 30:4n-6 are the main VLCPUFA molecular species.

3. Discussion

This study aimed to elucidate the comprehensive lipid composition of human sperm membranes of motile spermatozoa before the capacitation. Maintaining lipid homeostasis is crucial for sperm health, influencing the survival and functionality of male gametes during fertilization [13]. Thus, we characterized membrane lipids in human spermatozoa with high motility obtained by swim-up that are commonly used for in vitro fertilization assays, with a special focus on molecular species of GPL and the major sphingolipids SM and Cer.

Here, we showed that alongside the abundant GPL rich in 22:6n-3 and the “common” SM and Cer with saturated and monounsaturated fatty acids, molecular species of SM and Cer incorporating VLCPUFA are quantitatively critical constituents of human sperm membranes. In addition, this is the first report showing the quantification of endogenous molecular species of Cer pool in motile human sperm cells obtained by the same method used in assisted reproductive technology procedures. Interestingly, we found that SM from motile spermatozoa had higher percentage of VLCPUFA (about 10% of total fatty acid) in comparison with previous reports on total sperm cells, where VLCPUFA containing SM comprised 0 to 6.1% of the overall SM pool (mean 2.1%) [12, 14]. This is likely attributable to our analysis being conducted on a selected sperm population with the highest motility. Our findings stress the importance of the VLCPUFA concentration in SM for sperm functionality.

239 This prospect prompts intriguing inquiries into the characteristics and significance
240 of VLCPUFA within SL in human sperm membranes. Recognizing the presence of signifi-
241 cant amounts of SM and Cer containing "unusual" fatty acids in spermatozoa could facil-
242 itate exploration of their role in male reproductive physiology, potentially serving as key
243 players in fertilization-related processes. Indeed, Craig and colleagues [12] investigated
244 this issue in the human sperm total cell pool, noting that lower levels of VLCPUFA con-
245 taining SM were strongly associated with reduced sperm count and total motility.

246 Sphingoglycolipids and the sphingophospholipid, SM, constitute the main SLs
247 within eukaryotic cells, playing crucial roles in shaping membrane structure and func-
248 tionality [15]. Over the past decade, there has been a lot of interest in these lipids due to
249 certain metabolites they generate, such as sphingoid bases, phosphorylated sphingoid ba-
250 ses, and Cer, which are now essential messengers in cell signaling [15-20]. SM and Cer
251 alterations in their molar ratio impact membrane physical properties, influencing micro-
252 domain formation, vesicular trafficking, membrane fusion, as well as, different processes
253 involving membrane dynamics [21, 22]. Specific sphingomyelinases catalyze the hydroly-
254 sis of human sperm SM [20] generating most of the ceramide present in the male gamete.
255 In rodents and boar sperm, the enzyme activity rises just before fertilization hydrolyzing
256 SM, to produce ceramides during the acrosome reaction, with a consistent SM drop [23-
257 26]. Consequently, the steady-state levels of Cer in cells can be modulated by a set of en-
258 zymes, which remove or modify the SM or Cer. Cer is a minor lipid class in most animal
259 tissues and cells but is essential for human sperm function and fertilizing capability [18,
260 20, 27, 28].

261 We hypothesize that the fatty acid composition of these SLs plays a crucial role in
262 their biological function in membrane fusion. In secretory cells, SM is primarily regarded
263 as a structural lipid, yet it can also generate metabolites with highly bioactive properties
264 like Cer, ceramide 1-phosphate (C1P), or sphingosine 1-phosphate (S1P) [29, 30]. Further-
265 more, it remains unclear whether the incorporation of a VLCPUFA into the molecular
266 structure of the distinctive sperm ceramides affects their bioactivity, and if so, in what
267 manner.

268 Some authors have demonstrated the accumulation of VLCPUFA-rich SM, Cer and,
269 even glycosphingolipids (e.g. fucosylated GSL) in normal spermatogenic cells across dif-
270 ferent species [6, 10, 11, 31-33]. This finding underscores the incorporation of these sphin-
271 golipids into the spermatozoa where they play crucial physiological roles. The importance
272 of different sphingolipids in human male reproductive physiology has been demon-
273 strated. Our laboratory has elucidated the signaling cascades mediated by different sphin-
274 golipids during sperm acrosomal exocytosis [17, 18, 20]. Some enzymes involved in SL
275 metabolism, such as neutral-sphingomyelinase, ceramidases, ceramide synthase [20],
276 sphingosine kinase 1[17], and, ceramide kinase [18], are present in fully differentiated and
277 terminal cells like the human sperm. We determined how some of these enzymes and SL
278 regulate exocytosis. Cer induces the acrosome reaction and enhances the gamete response
279 to progesterone. Furthermore, we outlined the signaling sequence linking ceramide to the
280 internal and external mobilization of calcium during acrosome secretion [20], demonst-
281 rating that some enzymes involved in sphingolipid metabolism are active. For example, the
282 ceramide kinase is able to synthesize C1P from sperm ceramide under a calcium increase
283 during spermatozoa physiological function [18]. In addition, the ceramide effect is mainly
284 due to C1P synthesis. On the other hand, progesterone needs ceramide kinase activity to
285 rise intracellular calcium and induce the acrosome release. S1P, a bioactive sphingolipid,

286 initiates the acrosome reaction by binding to Gi-coupled receptors, thereby activating ex-
287 tracellular calcium influx and calcium efflux from intracellular reserves. [17].

288 Numerous studies in sperm and testis of different animal species stress the im-
289 portance of VLCPUFA in the reproductive physiology. The C20-C24 PUFA and \geq C26
290 VLCPUFA in spermatogenic cells originate from fatty acids derived from the diet, includ-
291 ing those from the n-6 series derived from linoleic acid, and the n-3 series derived from
292 linolenic acid (Figure 4). The PUFA elongases Elovl5 and Elovl2 are recognized for syn-
293 thesizing C18-C22 and C20-C24 PUFAs of the n-3 and n-6 series, respectively, operating
294 both sequentially and in conjunction with position-specific fatty acid desaturases. These
295 enzymes are essential for the major PUFAs (C20 and C22) found in the glycerophospho-
296 lipids of mammalian cells [34, 35]. Previously, it has been demonstrated that Lysophos-
297 phatidic acid acyltransferase 3 (LPAAT3) is essential for incorporating 22:6n-3 into the
298 membrane GPLs of differentiating germ cells. In LPAAT3-KO mice, there is a drastic and
299 specific decrease in DHA-containing phospholipids, leading to male infertility due to a
300 failure in spermiogenesis [36]. In addition, PUFA-rich GPLs, especially those containing
301 DHA, facilitate membrane reshaping and flexibility [37], a crucial characteristic for sper-
302 matogenic cells to progress through spermiogenesis and generate spermatozoa. The ma-
303 ture gamete retains these GPL molecular species until their final destination in the ovi-
304 duct. This suggests that maintaining this membrane deformability is important even up
305 to the moment of fertilization.

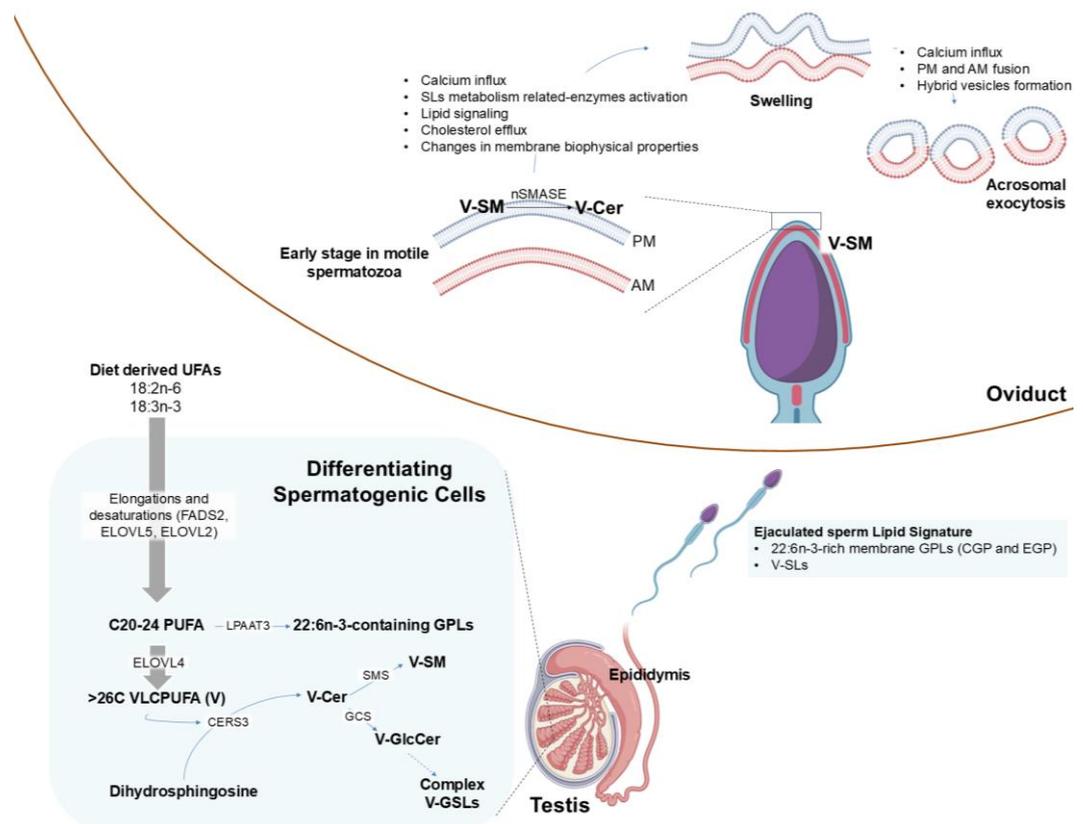
306 It has been shown that this enzyme group can elongate C20-C24 PUFAs up to C32 n-
307 V, a process mediated by the elongase Elovl4 [38], in rat seminiferous tubules [39] and,
308 more recently, in isolated germ cells [38]. The critical role of Elovl2 in PUFA and
309 VLCPUFA synthesis in germ cell lipids is highlighted by the fact that Elovl2 $-/-$ mice are
310 sterile, with their testes containing only spermatogonia and primary spermatocytes [40].

311 The production of sphingolipids with VLCPUFA in spermatogenic cells relies on the
312 Cers3 gene [10]. When Cers3 is specifically deleted in the germ cells of mice, leading to
313 the lack of VLCPUFA-containing Cer, SM, GlcCer, and complex GSL, it causes a halt in
314 spermatogenesis due enhanced apoptosis during meiosis and formation of multinuclear
315 giant cells [11]. Consequently, in rodents, both simple and complex sphingolipids contain-
316 ing VLCPUFA are crucial for normal spermatogenesis and male fertility. Interestingly,
317 CerS3-mRNA and SLs with VLCPUFA were absent in infertile human patients with Ser-
318 toli cell-only syndrome, and normal adult human testes contain the same molecular spe-
319 cies of VLCPUFA-containing Cer and SM that we observed in motile spermatozoa [11].
320 This demonstrates that these lipids, as well as VLCPUFA themselves, play a role early in
321 testis physiology and are conserved during epididymal transit to function in fully differ-
322 entiated, fertilization-ready spermatozoa.

323 Conversely, changes in the SM/Cer ratio (including those molecular species with
324 VLCPUFA) induce changes in rat sperm membrane stability [41]. Our results show that
325 human sperm Cer share the same fatty acids as sphingomyelins (SM), suggesting that Cer
326 may originate from the action of sphingomyelinase on pre-existing SM. In systems con-
327 taining SM where the Cer ratio is progressively increased by sphingomyelinase activity,
328 the formation of Cer leads to changes in lipid bilayer properties. Cer molecules spontane-
329 ously associate to form Cer-enriched microdomains that fuse into large Cer-rich mem-
330 brane platforms as Cer concentrations increase [42]. These Cer structures segregate from
331 the bulk liquid-crystalline fluid phase and exhibit gel-like properties. These properties
332 depend not only on the amount but also on the acyl chain length and unsaturation of the

333 generated Cer [43]. In agreement varied levels of Cer with VLCPUFA induced changes in
 334 cell membrane biophysical properties [23, 41], which could promote, along with other en-
 335 zymatic and signaling processes, the development of motility and/or acrosomal reaction
 336 (Figure 4).

337 In summary, the constituents of sperm membranes play a crucial role in development
 338 and sperm function. Recent lipidomics studies have highlighted the importance of
 339 membrane lipids—such as sulfogalactosylglycerolipid (SGG, seminolipid), cholesterol
 340 sulfate, and GPL with PUFAs—as key predictors of semen quality [44]. Notably, SLs with
 341 VLCPUFA have emerged as particularly significant.



363 **Figure 4.** The scheme summarizes the biosynthetic pathways of long (C20-C22) and very long-chain
 364 ($\geq C26$) polyunsaturated fatty acids (PUFA and VLCPUFA, respectively) and their incorporation into
 365 differentiating spermatogenic cell GPLs and SLs in the testis. After leaving the testis and passing
 366 through epididymal transit, ejaculated spermatozoa experience lipid remodeling but conserve
 367 22:6n-3-rich GPLs (CGP and EGP) and VLCPUFA-containing SM. Upon ejaculation, sperm cells
 368 carry these molecular species of lipids into the female reproductive tract, where additional lipid
 369 remodeling occurs in the oviduct, including the V-SM \rightarrow V-Cer reaction. These final modifications
 370 facilitate membrane fluidity and the ability of spermatozoa to undergo capacitation and the acro-
 371 some reaction, both of which are essential for fertilization. Abbreviations: AM, acrosomal mem-
 372 brane; Cer, ceramide; CERS3, ceramide synthase 3; CGP and EGP, choline and ethanolamine glyce-
 373 rophospholipids, respectively; ELOVL, elongation of very-long-chain fatty acid protein; FADS2,
 374 $\Delta 6$ desaturase; GPL, glycerophospholipids; GlcCer, glucosylceramide; GCS, GlcCer synthase; GSLs,
 375 glycosphingolipids; LPAAT3, lysophosphatidic acid acyltransferase 3; nSMASE, neutral sphingo-
 376 myelinase; PM, plasma membrane; SL, sphingolipids; SMS, SM synthase; SM, sphingomyelin;
 377 UFAs, unsaturated fatty acids; V, VLCPUFA. The scheme was created with BioRender.com.

4. Materials and Methods

4.1. Ethics statement and human sperm preparation.

Twelve healthy male donors provided ejaculates by masturbation, after at least 48 h of sexual abstinence. We used only semen samples that accomplished the World Health Organization (WHO, 2021) specifications, for the experiments shown here. Data collection adheres to the guidelines established in Argentina (ANMAT 5330/97) and the International Declaration of Helsinki. All donors signed an informed consent according to supply semen samples. The protocol for semen manipulation was accepted by the Ethics Committee of the School of Medicine, National University of Cuyo. After semen liquefaction (30–60 min at 37° C) and highly motile sperm were recovered after a swim-up separation for 1 h in HTF (5.94 g/liter NaCl, 0.35 g/liter KCl, 0.05 g/liter MgSO₄·7H₂O, 0.05 g/liter KH₂PO₄, 0.3 g/liter CaCl₂·2H₂O, 2.1 g/liter NaHCO₃, 0.51 g/liter glucose, 0.036 g/liter sodium pyruvate, 2.39 g/liter sodium lactate, 0.06 g/liter penicillin, 0.05 g/liter streptomycin, 0.01 g/liter phenol red supplemented with 5 mg/ml of BSA) at 37 °C in an atmosphere of 5% CO₂, 95% air. Cell concentration was then adjusted with HTF to 10 ×10⁶ sperm/ml. We developed the protocol as described in our publications[17, 18, 20, 45-47].

4.2. Lipid separation and analysis

The bulk of sperm suspensions obtained were then subjected to a gentle centrifugation (5 min at 400 x g). After collecting cells, lipid extracts were prepared and partitioned according to Bligh and Dyer [48]. Aliquots from these extracts were taken to determine the total phospholipid (PL) content in the samples by measuring the amount of lipid phosphorus[49].

For preparative isolation of lipid classes, most of the lipid extracts were spotted on TLC plates (500 µm, silica gel G) under N₂, along with commercial standards (Sigma Chemical Co, MO, USA). The polar lipids remained at the origin of the plates, and the neutral lipids were resolved in two steps. Chloroform: methanol: aqueous ammonia (90:10:2 by vol) was run up to the middle of the plates to separate the ceramides. Then, these solvents were evaporated and the plates were developed again by running n-hexane:diethyl ether (80:20, by vol.) up to the top of the plates to resolve minor neutral lipids (e.g. cholesterol esters and/or triacylglycerols).

The total phospholipid fraction was subjected to further separations and analyses. Aliquots were set aside to study the total GPL fatty acid composition. Most of the rest was used for preparative isolation of SM and major GPL using chloroform: methanol:acetic acid:0.15 mol/L NaCl (50:25:8:2.5, by vol). The phospholipids were resolved into classes by two-dimensional TLC [50]. The spots containing each phospholipid class were scraped from the plates followed by elution and quantification by phosphorus analysis in eluate aliquots. After drying, other aliquots of choline glycerophospholipids (CGP) and ethanolamine glycerophospholipids (EGP) were subjected to study fatty acid composition.

A mild alkali treatment was performed on the SM and Cer samples in order to remove any potential lipid contaminant containing ester-bound fatty acids. Both lipids were taken to dryness and treated (under N₂) with 0.5 N NaOH in anhydrous methanol at 50°C for 10 min. After this alkaline treatment, SM and Cer were recovered again by TLC. This procedure, involving alkaline methanolysis, was also used to obtain, as methyl esters, the fatty acids ester-bound to total GPL (thus excluding the fatty acids amide-bound to SM from this group). All solvents used in this study were HPLC-grade (JT Baker, NJ, USA; UVE, Dorwill, Argentina), and most procedures were carried out under N₂.

425 With the exception of the data shown in Fig. 1, in which lipid classes were located
426 with iodine vapors, lipids were located under UV light after spraying the plates with di-
427 chlorofluorescein, scraped from the plates and collected into tubes for elution. This was
428 achieved by three successive extractions of the silica support by vigorously mixing it with
429 chloroform:methanol:water (5:5:1, by vol.), centrifuging, collecting the solvents, and par-
430 titioning the resulting solvent mixtures with four volumes of water to recover the lipids
431 in the organic phases.

432 After elution, fatty acid analysis of Cer and SM was performed after converting the
433 lipids to fatty acid methyl esters (FAME), followed by gas-chromatography (GC), using
434 the conditions and instrumentation described in previous work [51]. Briefly, after adding
435 appropriate internal standards, transesterification was performed with 0.5N H₂SO₄ in N₂-
436 saturated anhydrous methanol by keeping the samples overnight at 45°C under N₂ in
437 screw-capped tubes. The resulting FAME were routinely purified by TLC on pre-cleaned
438 silica Gel G plates using hexane:ether (95:5, by vol) and then injected in the GC instrument.

439 To resolve according unsaturation, aliquots of FAME from Cer were subjected to ar-
440 gentation thin-layer chromatography (TLC) (Silica Gel G:AgNO₃, 80:20 by weight, and
441 chloroform:methanol, 90:10 by vol), which separated the esters in different bands contain-
442 ing saturates and different unsaturated fractions. After elution, these fractions were ana-
443 lyzed by GC/FID to identify the VLCPUFA linked to human sperm Cer.

444 4.3. Statistical analysis

445 Data are presented as mean values \pm SD from at least three samples pool, each ob-
446 tained from independent cell preparations. Statistical analysis was performed using the
447 Graph Pad Prism software, version 5.0 © (San Diego, CA). The Student t-test was used to
448 compare differences between two data. P values <0.05 were considered significant.

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474 **Abbreviations**

475 The following abbreviations are used in this manuscript:

476	ART	Assisted reproductive technology
477	Cer	Ceramide
478	CERS3	Ceramide synthase 3
479	CGP	Choline glycerophospholipids
480	DHA	Docosahexaenoic acid
481	EGP	Ethanolamine glycerophospholipids
482	ELOVL	Elongases
483	FADS2	Δ 6 desaturase
484	GCS	Glucosylceramide synthase
485	GluCer	Glucosylceramide
486	GPL	Glycerophospholipids
487	GSLs	Glycosphingolipids
488	LPAAT3	Lysophosphatidic acid acyltransferase
489	LPC	Lysophosphatidylcholine
490	LPE	Lysophosphoethanolamine
491	NL	Neutral lipids
492	nSMASE	Neutral sphingomyelinase
493	PA	Phosphatidic acid
494	PI	Phosphatidylinositol
495	PS	Phosphatidylserine
496	PUFA	Polyunsaturated fatty acid
497	SL	Seminolipid
498	SLs	Sphingolipids
499	SM	Sphingomyelin
500	SMS	Sphingomyelin synthase
501	UFAs	Unsaturated fatty acids
502	VLCPUFA	Very long-chain polyunsaturated fatty acid
503		

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