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Proteins of seminal fluid and spermatozoa in the trout (*Oncorhynchus mykiss*): Partial characterization and variations

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Abstract

The protein composition of seminal fluid, blood serum, sperm plasma membrane and flagellum of rainbow trout were analysed by SDS-polyacrylamide gel electrophoresis. Immunological identity between proteins of the 2 fluids and sperm components was studied using crossed immunoelectrophoresis, rocket immunoelectrophoresis and immunoblotting. Results indicate that many seminal proteins are antigenically-related to serum proteins, proteins of sperm origin are present in seminal fluid in varying amounts, depending on the animals and sampling time, and several serum-like seminal proteins are bound to spermatozoa.

Lipoproteins were isolated from seminal fluid (mean level: 33 $\mu\text{g}/\text{ml}$) and characterized. They were identified as being HDL-like lipoproteins. A possible physiological role is proposed for these seminal lipoproteins.

Introduction

In rainbow trout, spermatogenesis is completed about 2 months before spawning starts and spermatozoa are then in the lobular lumen. During the spawning period, spermatozoa are stored for variable periods in the sperm ducts. Although it has been shown in sea bass (Billard *et al.* 1977) that sperm fertility decreases throughout the reproductive period as a consequence of the sperm aging process, it has been demonstrated in trout that the capacity of sperm for withstanding cryopreservation and the resistance of the gamete plasma membrane to hypotonic shock are maintained, if not increased, during the spawning period (Malejeac *et al.* 1990). Thus, as already proposed by Billard (1976), the sperm duct which is the organ of sperm storage, most likely plays an important role in conservation of sperm fertility. However, when males are considered individually, unexplained apparently erratic varia-

tions are observed in sperm quality and in the properties of the gamete membrane (Billard 1976, 1983; Maise *et al.* 1988; Malejac *et al.* 1990).

In the lobular lumen and in the sperm duct, spermatozoa are suspended in seminal fluid which inhibits sperm motility and provides the components needed for sperm metabolism. The chemical composition of seminal fluid, which is different from that of blood plasma, is probably due to the existence of a blood-testis barrier (Steyn and van Vuren 1986). The seminal fluid of teleosts is a product of the testes (Clemens and Grant 1964) and of the sperm duct (Billard *et al.* 1971), the epithelium of which is secretory. The ionic composition of the seminal plasma has been established for several teleosts (Billard and Jalabert 1974; Plouidy and Billard 1982; Morisawa 1985; Piironen 1985; Steyn and van Vuren 1986). The concentration of various organic components such as total proteins, amino acids, total lipids, glucose, fructose, enzymes, has

been determined in some fish species (Plouidy and Billard 1982; Billard and Menezo 1984; Piironen 1985; Steyn and van Vuren 1986). In particular, lipids were detected in salmon seminal fluid (Piironen 1985) and phospholipids and cholesterol were identified in seminal plasma of cyprinid species (Belova 1982; Plouidy and Billard 1982). The form in which these molecules are present is not known.

Although seminal fluid has a protective effect on the gametes (Billard 1983), little is known about its proteinaceous components and their physiological functions. Since sperm possess limited biosynthetic capabilities, it has been documented in mammals that interactions of the spermatozoa with the seminal fluid would play a major role in the surface membrane changes which accompany epididymal sperm maturation. Events similar to such a maturation do not seem to occur in the vas deferens of trout. Indeed, in normal males, sperm acquire the ability to fertilize in the testes before being transferred to the sperm duct (Billard 1976). However, it is unknown whether intermale differences in the fertilizing ability of sperm (using either fresh or cryopreserved milt) are related or not to the quantity and/or quality of the organic components of the seminal fluid.

In order to improve our knowledge of trout seminal plasma, we analysed the proteins of blood serum, seminal fluid and gametes and using immunological techniques, we tentatively determine which antigens are common to the 2 fluids and gametes, and we identified and characterized lipoproteins present in the seminal fluid.

Materials and methods

Animals and sample preparation

One to 3 year-old males of rainbow trout were used. They were kept at 12–15°C, in recycled freshwater, under natural photoperiod.

Sperm was collected by hand stripping, taking care to avoid contamination by blood and feces, then stored on ice. Sperm samples were centrifuged (500 × g for 20 min at 4°C, then 1000 × g for 10 min; these conditions affect neither sperm motility

nor morphology) and the seminal fluid was collected. Spermatozoa were resuspended in seminal fluid mineral medium (SFMM, Billard and Jalabert 1974) pH 9, then washed once or twice by centrifugation. Seminal fluid was centrifuged either at 1000 × g for 20 min or at 100,000 × g for 40 min before being stored at –30°C until needed. No differences was apparent between the 2 procedures.

Blood samples were collected from the caudal vein and serum was prepared in the classic way.

Polyacrylamide gel electrophoresis

SDS-polyacrylamide slab gel electrophoresis was carried out according to Laemmli (1970) on 6–16% gradient gels. Samples of seminal fluid were diluted 1:2 with electrophoresis samples buffer containing 2% SDS and 5% mercaptoethanol; 70 µl was loaded onto the gels. Blood serum was diluted 1:10 with sample buffer and 10 µl of this was loaded. Lipoprotein samples were diluted 3:4 with concentrated sample buffer (×4) and 100 µl of this was loaded. Before loading, all samples were boiled for 5 min. Pharmacia calibration kit proteins were used as molecular weight markers. Proteins were stained with 0.2% R250 Coomassie Blue. Gels were scanned in a Vernon PHI5 densitometer. In some cases, the relative amount of protein bands was estimated by integration of the area under the whole electrophoregram and that under the considered bands.

Polyacrylamide gel electrophoresis under non-denaturing conditions was carried out as above, excepted that SDS and mercaptoethanol were absent in all the solutions and that gels were pre-electrophoresed for 4 hours.

Preparation of antisera and immunological procedures

Polyspecific antisera were raised in rabbits for serum proteins, for seminal fluid proteins and for saline-soluble spermatozoa proteins. Trout serum was diluted with 0.140 M NaCl to obtain 5 mg protein/ml. Seminal fluid was concentrated to the same protein concentration by dialysis against

0.140 M NaCl under vacuum. Washed spermatozoa were homogenised in 0.140 M NaCl in 0.02 M Tris HCl buffer pH 7.4, then centrifuged at $1500 \times g$ for 30 min and the supernatant was collected and adjusted to 5 mg protein/ml. For every antigen, a mixture of 0.9 ml of antigen solution and 0.9 ml Freund's complete adjuvant (2 first injections) then incomplete adjuvant was injected intradermally into the abdomen of 2 rabbits, seven times biweekly. Antisera were tested by the Ouchterlony method and by rocket immunoelectrophoresis performed following routine procedures (Johnstone and Thorpe 1982).

Presence of immunologically-related antigens in blood serum and in seminal fluid on the one hand and in seminal fluid and in spermatozoa on the other was checked using crossed immunoelectrophoresis (Johnstone and Thorpe 1982) and immunoblotting procedure. Following slab gel electrophoresis, carried out with denaturing conditions either present or absent, proteins were transferred to nitrocellulose sheets (Gelman, $0.2 \mu\text{m}$) by electrophoresis in the presence of 0.03% SDS and 20% methanol. Efficiency of transfer was checked by staining the transferred gel with Coomassie Blue and by staining the sheets with 0.2% Ponceau red in 3% trichloroacetic acid before immunological detection. Antigens on the sheets were detected by indirect specific labelling performed at 37°C . Unspecific binding was prevented by using swine serum diluted 1:10 (TBSS; 1 h incubation) in 0.02 M Tris HCl buffer pH 8, 1% NaCl (TBS). The specific immunoserum was diluted 1:100 in TBSS (2 h incubation). The second ligand was antisera to rabbit IgG raised in swine and conjugated with horseradish peroxidase (Dako) diluted 1:500 in TBSS (2 h incubation). After each incubation with the antisera, sheets were washed twice for 10 min with 0.05% Tween 20 in TBS. Peroxidase conjugates were detected by reaction with 4-chloro-1 α -naphthol and hydrogen peroxide.

Anion-exchange HPLC of proteins in blood serum and in seminal fluid

The method used to fractionate proteins of the 2 biological fluids on a Mono Q column ($5 \times 50 \text{ mm}$

i.d., Pharmacia) was modified from Cheng *et al.* (1986). Briefly, we used, at room temperature, a chromatographic system from Waters equipped with an automated gradient controller and 2 pumps. Sample (pools of blood serum or seminal fluid collected at the same time from the same males) volume was 2 ml and protein concentration was 10 mg/ml (seminal fluid was concentrated by vacuum dialysis). Protein elution was performed using a linear gradient 0 to 0.480 M NaCl in 0.02 M Tris HCl buffer pH 7.4 with 3 steps at 0.78, 0.120 and 0.246 M NaCl. The effluent was monitored by UV absorbance at 280 nm. Seven milliliter fractions were collected, dialysed against 0.02 M Tris HCl buffer pH 7.4 then analysed by SDS gel electrophoresis. Each experiment was repeated 3 times.

Preparation of plasma membrane proteins

Direct extraction with sodium desoxycholate: Spermatozoa were washed once more with SFMM then pellets were resuspended in 12 volumes 0.2% Na desoxycholate in 0.25 M sucrose, 0.001 M phenylmethylsulfonide fluoride (PMSF), 0.005 M Hepes buffer pH 8.0 at 4°C for 45 min (Jones *et al.* 1983). Unlike boar sperm (Klint *et al.* 1985), the increase of desoxycholate concentration up to 0.6% or the addition of 0.001 M dithiothreitol did not change the electrophoretic pattern of membrane proteins. After centrifugation ($1000 \times g$, 20 min), supernatants were removed then centrifuged at $25,000 \times g$ for 1 h. Proteins were collected by centrifugation after precipitation by mixing overnight with 9 volumes of ethanol at 4°C . Proteins were washed with ethanol, dried, then solubilised in electrophoresis sample buffer at a concentration of 1.5 mg/ml.

Plasma membrane isolation using nitrogen cavitation: Membranes vesicles were prepared as follows: after washing, 4.8×10^{11} spermatozoa were resuspended in 30 ml 0.02 M Tris-HCl pH 7.8, 0.003 M MgCl_2 , 0.001 M PMSF, 20 μg Soybean trypsin inhibitor/ml, 0.1 μg pepstatin/ml (Tris- MgCl_2), containing 0.25 M sucrose. Nitrogen cavitation was performed under 900 PSI for 20 min. Nuclei and flagella were removed from the homogenate by centrifugation onto 0.65 volumes of 1 M

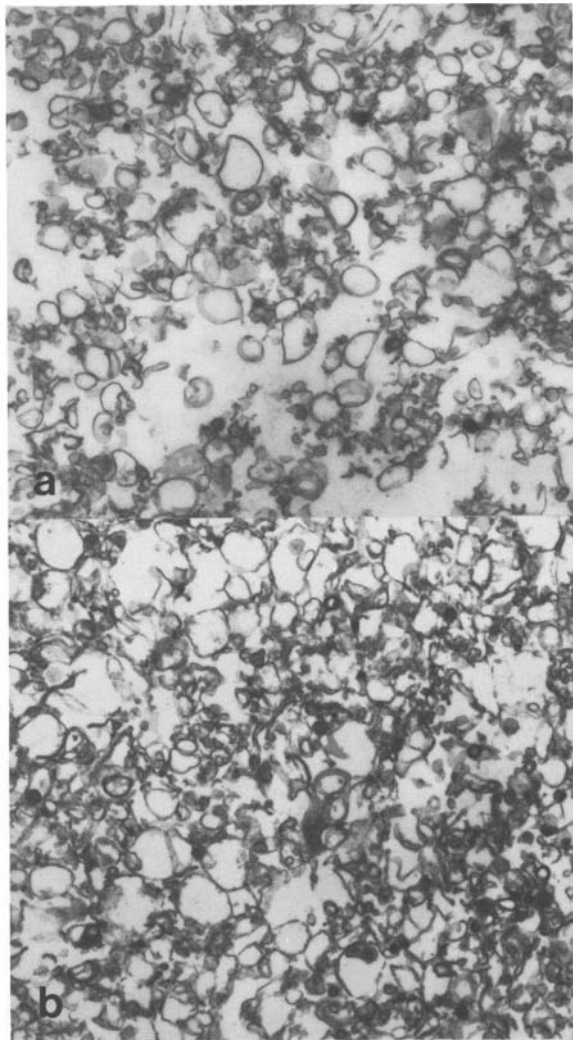


Fig. 1. Electron micrographs ($\times 8,200$) of membrane pellets obtained after centrifugation on continuous sucrose gradients of the homogenate resulting from disruption of trout spermatozoa by nitrogen cavitation. a: band 1 material (0.25 M sucrose); b: band 3 material (0.62 M sucrose).

sucrose at $500 \times g$ for 10 min, then at $1500 \times g$ for 10 min. The suspension was then fractionated by centrifugation ($80,000 \times g$, 2 h at 4°C) on a continuous 0.25–1.8 M sucrose gradient. Fractionated material was collected, diluted at least 1:3 in Tris- MgCl_2 and pelleted at $100,000 \times g$ for 30 min. As confirmed by electron microscopy (Fig. 1), by determination of lactic dehydrogenase and succinic dehydrogenase activities, of cholesterol, phospholipid and triglyceride, membrane vesicles were

recovered without detectable contamination by other spermatozoa organelles in three bands (0.25, 0.37 and 0.62 M sucrose).

Preparation of flagellar proteins

After treatment with sodium desoxycholate, the pellets of demembrated spermatozoa were re-suspended in the desoxycholate solution then sonicated until nearly all the heads and flagella were disconnected. Nuclei were discarded by pelleting ($1,000 \times g$, 4 min). The flagella suspension was adjusted to 0.9 M sucrose then centrifuged ($100,000 \times g$, 3 h) on a discontinuous 1.15–2.2 M sucrose gradient. Completely demembrated flagella (control with electron microscopy) were collected at the interface 1.70–2.00 M sucrose, diluted at least 1:3 in SFMM, 0.001 M PMSF and pelleted at $25,000 \times g$ for 25 min.

Electron microscopy

Aliquots of membrane and flagella suspensions were fixed in 4% glutaraldehyde in phosphate buffered saline (PBS) pH 7.5, washed in 0.2 M cacodylate buffer pH 7.5 and post-fixed in 1% osmium tetroxide in 0.1 M cacodylate buffer plus 1% potassium ferricyanide. Material was embedded in 2% agar, dehydrated in acetone and embedded in Epon-araldite. Sections were double-stained and observed with a 100 CX Jeol electron microscope.

Isolation and analysis of seminal fluid lipoproteins

Total lipoproteins were isolated from seminal fluid by flotation according to Fremont and Marion (1982) modified for seminal fluid lipoproteins. The density of seminal fluid was raised from 1.006 g/ml (1.0064 ± 0.0012 , $n=7$) to 1.27 g/ml by addition of solid KBr. Preliminary experiments had shown that a very low proportion of the expected amount of lipoproteins was recovered when the density was adjusted to 1.21 g/ml, while most of the lipoproteins were obtained without increased contamina-

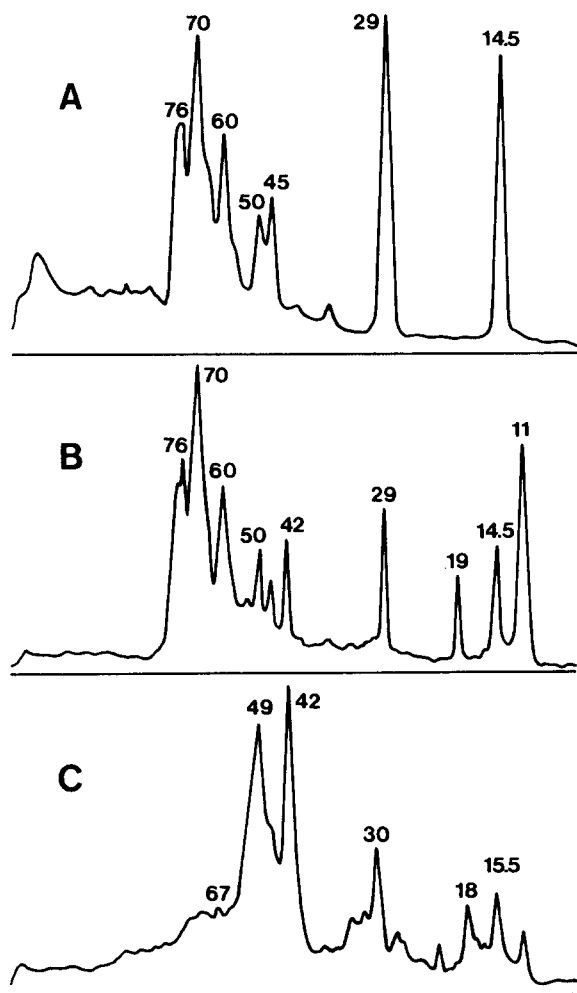


Fig. 2. Electrophoregrams (SDS gels) of the proteins present in the antigen solutions used to immunize the rabbits. A: blood serum; pool from males at the beginning of the spermiation period. B: Seminal fluid; pool from same males. C: saline-soluble sperm proteins; pool from same males.

tion by other proteins after adjustment to 1.27 g/ml. Polycarbonate tubes were filled with 22.5 ml sample (containing 0.03 M Na azide and 0.001 M PMSF), overlaid with 6.5 ml saline solution at 1.27 g/ml. These were centrifuged at $143,000 \times g$ for 44 h at 10°C . Lipoproteins were collected from the top of the tubes. An aliquot was also collected from the bottom. All the samples were exhaustively dialysed against 0.185 M NaCl, 0.03 M Na azide and 0.001 M EDTA.

Total lipoproteins, LDL and HDL were isolated

from 2 pools of blood serum obtained from mature males, according to Fremont and Marion (1982).

Polyacrylamide-gel electrophoresis of lipoproteins was performed in discontinuous three-step gradient gels (2.8, 3.4 and 6.2%, pH 8.9; Leger *et al.* 1979) after prestaining the lipoproteins with Sudan black B.

Other methods

Protein was determined by the method of Lowry *et al.* (1951) using bovine serum albumin as the standard. Protein in lipoprotein samples was measured according to Fremont and Marion (1982), in the presence of 5% Na desoxycholate. Enzymatic methods were used to quantify total cholesterol, triglyceride and phospholipid (Boehringer).

Results

Electrophoretic analysis of blood serum and seminal fluid proteins

Most of the proteins identified in blood serum by 1D SDS gel electrophoresis (Fig. 2) were present in roughly the same relative amounts whatever the sampling time during the spermiation period, while the relative amount of some of the others was slightly variable. Major protein bands were: 14.5, 29, 45, 50, 60, 68–70, 76 and 78 kD. Some of the serum proteins co-migrated with apolipoproteins present in serum total lipoproteins (Fig. 3) and in serum HDL and LDL: 14.5 kD (apo. AII), 29 kD (apo. AI), 44 kD, 54 kD, 76 kD and 250 kD (apo. B).

The mean protein concentration in seminal fluid, determined for 45 sperms collected throughout the spermiation period, was $1.737 \text{ mg/ml} \pm 0.792$.

No protein was significantly detectable above 85 kD in seminal fluid (Figs. 2 and 3). Proteins from 45 to about 85 kD and proteins 14.5 and 29 kD co-migrated with proteins present in serum and their amounts roughly paralleled those of the serum proteins. Various seminal proteins were usually present in variable relative amounts depending on the sampling time during the spermiation period. It was the case for four proteins (11, 19, 37 and 42 kD) un-

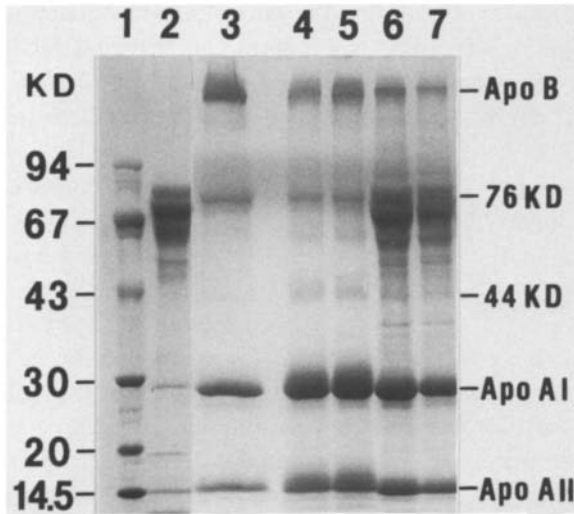


Fig. 3. Electrophoretic patterns of blood serum, seminal fluid and serum apolipoproteins, SDS polyacrylamide gel. 1: standard molecular weights. 2: pool of seminal fluid. 3, 4, 5: serum total lipoproteins; pools of serum from males. 6, 7: blood serum; pools from males.

related to serum. Protein 42 KD which co-migrated with the major membrane protein was in higher amounts at the beginning and especially at the end of the spermiation period (additional proteins co-migrating with membrane proteins were also observed at this time), rather than in the middle of the period. The relative amounts of proteins 29 and 14.5 KD as percentage of total seminal proteins (estimated by integrative densitometry of the bands after analysis on SDS gels) were followed for at least 3 weeks for 10 males considered individually. It appeared that they varied slightly during this period but depended on the males.

Immunochemical identity between seminal proteins, serum proteins and sperm proteins

The antiserum specific for serum proteins was tested for antigenicity of seminal fluid, firstly using crossed immunoelectrophoresis. Six to 13 rockets were observed (Fig. 4). Secondly, we used the immunoblotting technique (Fig. 5). Seminal proteins 11, 14.5, 16.5, 29, 34, 40, 44–45 and 50 to 80 KD were regularly recognized by the antiserum for se-

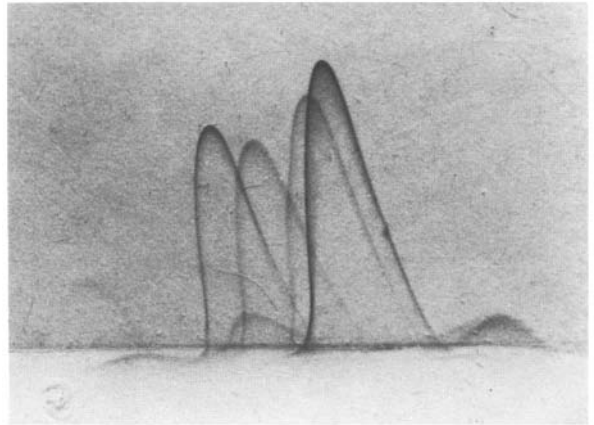


Fig. 4. Crossed immunoelectrophoresis of seminal proteins (pool of seminal fluid; males at beginning of spermiation; identical to pool 2 in Fig. 5). The second dimension contained antiserum specific for serum proteins.

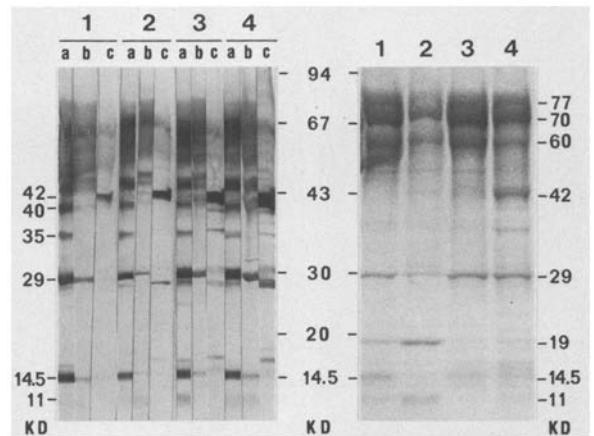


Fig. 5. Immunoblots (left) obtained after electrophoresis of 4 pools of seminal fluid on SDS gel and transfer to nitrocellulose sheet (right, after staining with Ponceau red). a: immunodetection with antiserum specific for serum proteins. b: antiserum for seminal proteins. c: antiserum for NaCl-soluble sperm proteins. Pool 1: males in mid-spermiation period. Pools 2 and 3: males at beginning of spermiation. Pool 4: males at the end of spermiation period; seminal fluid collected from testes.

rum proteins. When immunodetection was carried out after transfer of seminal fluid proteins from non-denaturing gels, identification of the protein bands by the molecular weight was not possible, but about as many bands were recognized by antiserum for serum proteins as after transfer from SDS gels.

Depending on the pools of seminal fluid which were checked by immunoblotting, variable num-

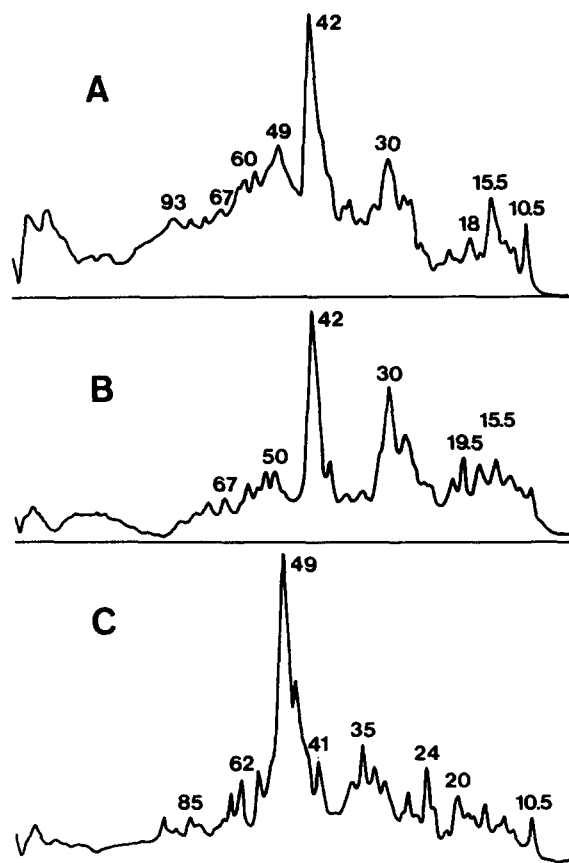


Fig. 6. Electrophoregrams (SDS gels) of, A: proteins extracted from spermatozoa by Na desoxycholate; B: proteins extracted from membrane vesicles obtained in band 2 after centrifugation of spermatozoa disrupted by nitrogen cavitation onto sucrose density gradients; C: proteins present in flagella after demembration of spermatozoa by Na desoxycholate.

bers of protein bands were recognized by the antiserum for NaCl-soluble sperm proteins. At least protein bands 42 and 49 to 70 (Fig. 5) were detected in fluids collected in the middle of the spermiation period. At the beginning and the end of this period, additional protein bands were recognized: 15.5, 17, 18, 27, 28, 30, 35, 39, 45, 48 and around 130 KD. Immunological crossreactivity of seminal fluid proteins (individual samples) with the antiserum for NaCl-soluble sperm proteins was also checked by rocket immunoelectrophoresis. Six to a dozen rockets were observed depending on the samples. The height of the tallest rocket was roughly proportional to the total protein concentration in the sample and was unrelated to the relative amount of protein 42

KD (0 to 13%). The 19 KD seminal fluid protein was never recognized by any of the 3 antisera.

HPLC fractionation of serum and seminal fluid proteins

A total of respectively 11 peaks and 9 peaks were detected after fractionation of blood serum and seminal fluid proteins. The seminal protein peaks co-eluted with serum peaks but their heights were different. Electrophoretic analyses indicated that most of the major proteins common to the fluids and which have the same molecular weight were eluted by the same NaCl molarities.

Proteins of sperm plasma membrane and flagella

Most of the proteins extracted from spermatozoa by sodium desoxycholate had molecular weights ranging between 10.5 and 110 KD (Fig. 6). The predominating protein was 42 KD large.

Proteins present in the three bands obtained after membrane isolation using nitrogen cavitation and fractionation on sucrose gradient, were similar to those obtained using desoxycholate. However 48–110 KD proteins were present in lower amounts and there was few proteins between 32 and 40 KD (Fig. 6). This suggests that in desoxycholate extracts all these proteins corresponded partly to detergent-soluble non-membranous (cytoplasmic?) proteins.

Among the flagellar proteins, one (49 KD) predominated. It most likely corresponded to the α and β chains of tubulin. The other numerous protein bands were scattered between 10.5 and 120 KD. Major protein bands were 20, 24, 35, 47, 56 and 62–70 KD.

Most of the saline-soluble proteins of spermatozoa (Fig. 2) co-migrated with membrane proteins and with some of the major flagellar proteins (Figs. 2 and 6). After transfer of saline-soluble proteins to nitrocellulose sheets, protein bands 14.5, 29, 45 and 63–80 KD were commonly recognized by both antisera for seminal proteins and for serum proteins. Protein bands 30 and 42 KD were detected only by antiserum for seminal proteins.

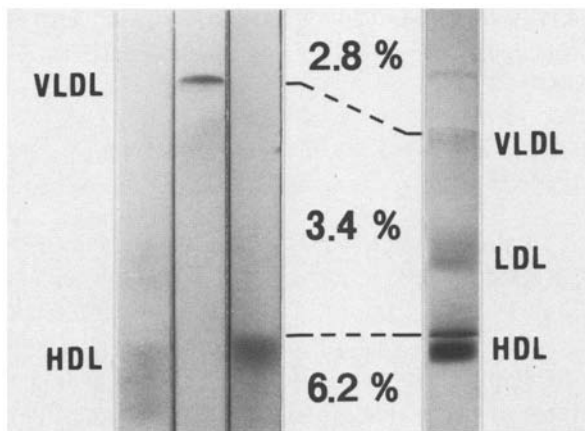


Fig. 7. Electrophoresis of trout lipoproteins in polyacrylamide step gradient gels (2.8, 3.4 and 6.2% acrylamide). 1: pool of seminal fluid; males in mid-spermiation period. 2: serum VLDL; same animals. 3: lipoproteins isolated from a pool of seminal fluid; males in mid-spermiation period. 4: serum total lipoproteins. Gels 1–3 and gel 4 were not run together.

Lipoproteins of seminal fluid

When lipoproteins isolated from pools of seminal fluid were analysed by electrophoresis in discontinuous three-step gradient gels, lipoproteins were observed only in the 6.2% gel where they co-migrated with serum HDL (Fig. 7).

When they were analysed on SDS gels, 2 protein bands were observed at 29 and 14.5 KD, the latter one being predominant. They co-migrated with AI-like and AII-like apoproteins present in serum HDL and LDL (Fig. 8). In addition, minor bands 12–14, 54 and 76 KD were either present or absent depending on the pool of fluid. No detectable amounts of 29 and 14.5 KD proteins remained in the bottom of the tubes after centrifugation of seminal fluid to isolate lipoproteins.

We determined protein (P), total cholesterol (C), triglyceride (T) and phospholipid (PL) in lipoproteins isolated from 9 pools (7 to 17 males) of seminal fluid collected weekly during the spermiation period. The mean lipoprotein concentration was $32.9 \pm 9.5 \mu\text{g/ml}$ seminal fluid (mean \pm SEM). The mean distribution of protein and lipid components as percentage of total lipoprotein weight was P: $50 \pm 6.3\%$, C: $20 \pm 4.2\%$, T: $13 \pm 2\%$ and PL: $17 \pm 3.5\%$. The ratio C/P (w/w) was equal to

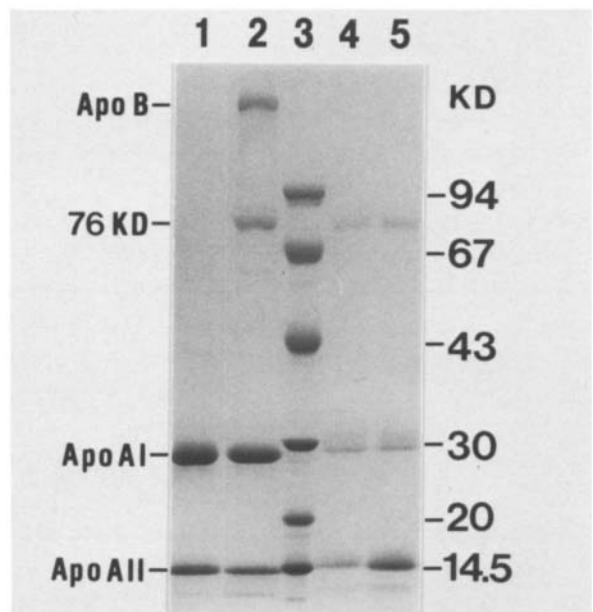


Fig. 8. Electrophoretic patterns (SDS-polyacrylamide gels) of trout apolipoproteins. 1: serum HDL. 2: serum LDL. 4 and 5: 2 pools of seminal fluid. 3: standard molecular weights.

0.41 ± 0.13 and the ratio T/P (w/w) to 0.28 ± 0.07 .

Discussion

In the rainbow trout, the protein concentration in seminal fluid is about 20 times lower than in blood serum (mean value in mature males of rainbow trout: 33 mg/ml according to Borchard 1978 and 36 to 41 mg/ml ($n=3$ pools) in our experiment). In mammals, protein levels in rete testis fluid are about 100 times less than in plasma (Setchell and Wallace 1972).

Our immunological data obtained in denaturing and non-denaturing conditions indicate that a lot of seminal proteins are antigenically-related to serum proteins. These are 11, 14.5, 16.5, 29, 34, 40, 45 and 50 to 80 KD proteins. After HPLC, each protein of both origin was eluted using the same NaCl molarity. These data suggest the presence in the 2 biological fluids of homologous proteins. Matsubara *et al.* (1985) have concluded that, in chum salmon, a number of serum proteins are present in

the coelomic fluid, this suggesting that coelomic fluid could be derived at least in part from blood. In mammals, proteins present in rete testis fluid are also similar to serum proteins. However, even if some of them are derived from the passage of proteins from blood and lymph, it has been demonstrated that several serum-like proteins are produced by Sertoli cells (Mather *et al.* 1983). In the case of the trout seminal fluid, no data are available regarding the possibility of testis cells and/or cells lining the sperm duct producing some of the serum-like proteins. Immunological studies will permit the presence of such proteins in intralobular testis cells as well as in Sertoli cell culture medium to be identified.

Seminal proteins, present in variable relative amounts or even absent, depending on the samples, were recognized by the immunoserum for spermatozoa proteins. Among these recognized proteins, the following were not recognized by the immunoserum for serum proteins: 15.5, 17, 18, 27, 28, 30, 35, 39, 42 and 49 KD. In addition, these proteins co-migrated with constantly present major membranous or flagellar sperm proteins. They are present, sometimes in conspicuous amounts, in seminal fluids collected, in particular, at the end of the spermiation period. These data suggest that they are proteins of spermatozoan origin which were released into the fluid by dying gametes. This assumption is corroborated by the observation of Maisee *et al.* (1988) and of Malejac *et al.* (1990) that the trout ejaculates which have the best fertility after cryopreservation contain spermatozoa with a high functional integrity of the plasma membrane and lack 42 KD protein in the seminal fluid.

Other seminal proteins recognized by the immunoserum for saline-soluble sperm proteins co-migrated with sperm proteins, which themselves were recognized by both immunosera for serum proteins and for seminal proteins. These are 45 and 63–80 KD proteins. This could be due to the simultaneous presence in these bands, identified in 1D gels, of 2 proteins: one serum-like, the other one of spermatozoan origin. As an alternative explanation, these proteins may be serum-like seminal proteins which were bound to the sperm cells used to immunise the rabbits. This hypothesis is favoured

by the fact that the 14.5 and 29 KD saline-soluble sperm proteins were also recognized by the immunoserum for serum proteins. Thus, we propose that some of the major serum-like seminal proteins are present on the spermatozoa. Spermatozoa that were used were washed twice with SFMM after removal of the seminal fluid. Because such a washing treatment is perhaps not sufficient to remove proteins absorbed onto the gamete surface, we do not know whether these proteins were incorporated in the plasma membrane or only adsorbed. However, it may be noted that 48 to 61 KD seminal proteins were not detected on the spermatozoa. This suggests that not all serum-like seminal proteins interact with the gamete membrane. Such interactions have been documented in mammals (Jones *et al.* 1983; Dacheux *et al.* 1989) as participating in the changes which occur in the protein composition of the sperm membrane during epididymal transit. In the trout, it is questionable whether, during sperm storage in the sperm duct, interactions between seminal proteins and the gamete membrane could be important to maintain the fertility of gametes.

Finally, only one major seminal protein was not characterized for its type or origin. It is the 19 KD protein. It co-migrates with a protein present in trout coelomic fluid (data not shown), it differs from the coelomic fluid-specific protein identified by Matsubara *et al.* (1985) and it is questionable whether it is a gonad-specific protein.

Our data show that small amounts of HDL-like lipoproteins are present in trout seminal fluid. Lipids were detected in seminal fluid in cyprinid species (Belova 1982; Plouidy and Billard 1982), in the catfish *Clarias gariepinus* (Steyn and van Vuren 1986) and in *Salmo salar* (Piironen 1985), but not in *Onchorhynchus keta* (Morisawa *et al.* 1979). The presence of lipoproteins in seminal fluid has been suggested in mammals (Clavert *et al.* 1980), but only demonstrated in cockerels (Blesbois and Hermier, unpublished data). In trout as in cockerels, only HDL (or VHDL) have been identified. While in mature males of trout there is 10.13 mg HDL/ml serum (Fremont and Marion 1982), there is a mean of 33 μ g HDL-like lipoproteins/ml seminal fluid, *i.e.*, 160 times less than in serum. However, because we have determined phospholipid levels by an en-

zymatic method which underestimates values compared to phospholipid phosphorus determination, the actual HDL concentration may be around 40 $\mu\text{g/ml}$. It is lower than that found for total lipids (around 200 $\mu\text{g/ml}$ as a mean) in seminal plasma of *Salmo salar* (Piironen 1985), but it is only twice lower than in cockerels (66 $\mu\text{g/ml}$). These seminal HDL-like lipoproteins have a higher buoyant density (1.27 g/ml) than the serum HDL (1.21 g/ml; Fremont and Marion 1982). This disagrees with the fact that there is a trend for the T/P and C/P ratios to be slightly higher in seminal HDL than in serum HDL (respectively, 0.28 vs. 0.21 and 0.41 vs. 0.30).

In biological fluids, lipoproteins act as lipid carriers and HDL are the main vehicles for biomembrane lipid components (phospholipid and cholesteryl ester). Thus, we propose as a hypothesis, that the HDL-like lipoproteins present in seminal fluid interact with the sperm plasma membrane to maintain its lipid composition during storage in the sperm duct. This proposal is partly sustained by the observation that apo AI (29 KD) and apo AII (14.5 KD) were observed to be bound to spermatozoa.

At the present time, the possible origin of the various seminal HDL components is unknown. Although apo AI and AII were recovered in the human pieces of sperm ducts which were extensively washed then incubated for 24 h, the 2 extremities being closed, we were unable to label these proteins with ^{14}C -amino acids added to the medium surrounding the pieces of sperm ducts (unpublished data).

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