


Impact of *Escherichia coli*, *Candida non-albicans*, and *Trichomonas vaginalis* on Semen Chemical and Functional Parameters: an *In-Vitro* Study

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Background: Infertility is a worldwide medical issue in which infection is recognized to play a major role. Pathogens trigger various mechanisms that impact fertility, either directly by affecting the physiological indices of semen or indirectly by disrupting the process of spermatogenesis. In the current work, the effect of *in-vitro* cultivation of *Escherichia coli* (*E. coli*), *Candida non-albicans* (*C. non-albicans*), and *Trichomonas vaginalis* (*T. vaginalis*) (as the most frequently reported sexually transmitted infections) was assessed on the physiological functions of the spermatozoa and the chemical characteristics of the seminal fluid. **Method:** The semen samples were exposed to cultures of *E. coli*, *C. non-albicans*, and *T. vaginalis*. The study analyzed the changes in motility, agglutination, viability, DNA fragmentation index (DFI%), seminal pH, and biochemical parameters at 1/2, 1, 1.5, 2, 2.5, 3.5 and 4 hours.

Results: Incubation of the semen samples with *E. coli* resulted in a progressive increase in agglutination, pH, and nitrite. The seminal glucose and the sperm motility, on the other hand, were reduced. The sperm vitality and seminal protein remained unaffected. *C. non-albicans* induced three forms of agglutination (head-to-head, tail-to-tail, and head-to-tail), lowered pH values and decreased the sperm motility, but did not alter the seminal protein, glucose, nitrite, nor the spermatozoa viability at the different tested time intervals. *T. vaginalis* resulted in increased seminal protein, and reduced glucose, pH, and motility. It also induced minimal agglutination and caused unchanged nitrite and sperm viability. The DFI% was increased in all pathogens with the *C. non-albicans* showing the highest DNA fragmentation index.

Conclusion: Urogenital infection with *E. coli*, *C. non-albicans*, or *T. vaginalis* is assumed to affect the quality of semen through DNA fragmentation, agglutination and altered seminal chemical microenvironment.

Keywords: *E. coli*; *C. non-albicans*; *T. vaginalis*; seminal parameters; DNA fragmentation

Introduction

Infertility is a crucial global medical problem [1]. The male factor has been recognized to account for 20% of the infertility cases [2] with the genitourinary infection representing a significant cause (8–35%) [3]. Various microbial agents including bacteria, fungi, and parasites can cause reproductive disorders. These microbial infections inhibit the acrosomal functionality of spermatozoa, alter the normal sperm morphology and compromise spermatogenesis [4].

Escherichia coli (*E. coli*) is the most common bacteria affecting the urinary tract. *E. coli* is involved in causing non-sexual epididymo-orchitis, thereby representing a major causative agent of infertility [5]. This effect could be related to the toxins and metabolites produced by *E. coli*, or the direct adherence of the bacteria to the spermatozoa [6]. In Egypt, several studies were conducted alarming the epidemiological spread of *E. coli* pathogenicity [7–11].

Candida non-albicans (*C. non-albicans*) is an opportunistic fungal infection that causes urogenital infections [12]. The actual role of yeast has scarcely been investigated due to experimental difficulties [13], yet, the deleterious effects of genital *Candida* infection on semen parameters have recently attracted attention [14]. Urogenital candidiasis has significantly shifted from the *albicans* to the *non-albicans* type in Egyptian patients [15].

Trichomonas vaginalis (*T. vaginalis*) has long been recognized as a mere vaginal parasite. However, it has now been accepted as a cause of mutual urogenital infection in both males and females, representing a significant factor in sexually transmitted diseases (STDs). In men, *T. vaginalis* infection has been connected to infertility through impairment of the functionality and quality of the spermatozoa [16]. Semen was found to provide an adequate medium for the survival of trichomonads [17]. The relationship between *T. vaginalis* and infertility in females has also been reported [18].

The spermatozoon is assumed to be the only cell that accomplishes its physiological functions outside the human body. Seminal plasma is composed of sugars, and proteins in addition to other components such as $\text{HCO}_3^-/\text{CO}_2$, cholesterol, phospholipids, steroids, nitrogenous bases and amino acids. The optimum composition of this seminal microenvironment is crucial for optimizing sperm functions. Pathogens affect the chemical composition of semen and thus can influence the functionality of spermatozoa [19,20]. In addition, the integrity of the paternal DNA in the spermatozoa plays a significant role in initiating and preserving viable gestation, whether via natural conception or *in-vitro* fertilization [21]. Therefore, the WHO has demonstrated the importance of assessing sperm chromatin integrity and considered it an important parameter in semen analysis [22].

In this context, the current study has been designed to investigate the effect of *E. coli*, *C. non-albicans*, and *T. vaginalis* (as the most prevalent community-based transmitted organisms) on the semen functional and chemical parameters and to determine the integrity of the sperm DNA after sperm-pathogen co-incubation.

Materials and Methods

Study Setting

This study was approved by the ethical committee of the Armed Forces College of Medicine, Cairo, Egypt (No. 265 dated 17/6/2023). According to Amsterdam's declarations, all participants have been verbally informed, and a written consent has been signed before any step. All the study procedures followed the ethical standards of the National Research Committee and the 1964 Helsinki Declaration and its ensuing regulations.

The semen ejaculates were obtained from twenty four healthy adult Egyptian males as part of their fertility eval-

uation. The semen samples were gathered by masturbation after sexual abstinence for 4–6 days. The samples were obtained in the clinical setting (Department of Dermatology, Venereology, and Andrology, AFCM Hospital and Laboratories of Almaza, Cairo, Egypt).

Study Design

To execute the *in-vitro* cultivation of the microorganisms with semen, 24 patients (for the acquisition of 24 healthy semen ejaculates) were involved in the study. The semen samples were randomly divided into four groups as follows:

Group 1 (6 samples): Control.

Group 2 (6 samples): Semen incubated with *E. coli*.

Group 3 (6 samples): Semen incubated with *T. vaginalis*.

Group 4 (6 samples): Semen incubated with *C. non-albicans*.

The microorganisms were obtained from the microbiology and immunology department, Faculty of Medicine, Cairo University as pure isolates.

Preparation of Semen Samples

All samples were obtained in sterile plastic cups and incubated at 37 °C for 45 min to liquefy. Semen was centrifuged for 10 min at 1500 rpm at room temperature [23]. The Mycoplasma test revealed negative results. The obtained semen samples were physically assessed for color, viscosity, volume, and pH. Using light microscopy, the count of spermatozoa per mL was calculated in five small squares within the large central square in the improved Neubauer hemocytometer (Germany) [24].

Total and progressive motility was evaluated using the computer-assisted sperm analysis device, SCA® CASA system. The morphology of the spermatozoa was evaluated using an Olympus light microscope (Olympus BX43F, Olympus Corporation, Tokyo, Japan).

The *in-vitro* cultivation process included only the healthy semen ejaculates that were microscopically devoid of trichomonads and microbiologically negative for both bacteria and fungi. The standards for healthy semen samples adopted in the current studies were illustrated in Table 1 as per the WHO criteria [22,25]. None of the participants was under antibiotic therapy at the time of the sample collection or had a history of anti-parasitic/antibacterial medication for the past 3 months. Patients with abnormal semen parameters, contaminated specimens or patients suffering from diabetes mellitus were excluded from the study [26].

In-Vitro Co-Incubation of Semen with Microorganisms

One hundred microliters from each of the *E. coli*, *C. non-albicans*, and *T. vaginalis* cultures (grown for 24 and 48 hours) were mixed with an equal volume of semen samples and incubated at 37 °C in Eppendorf tubes labeled for

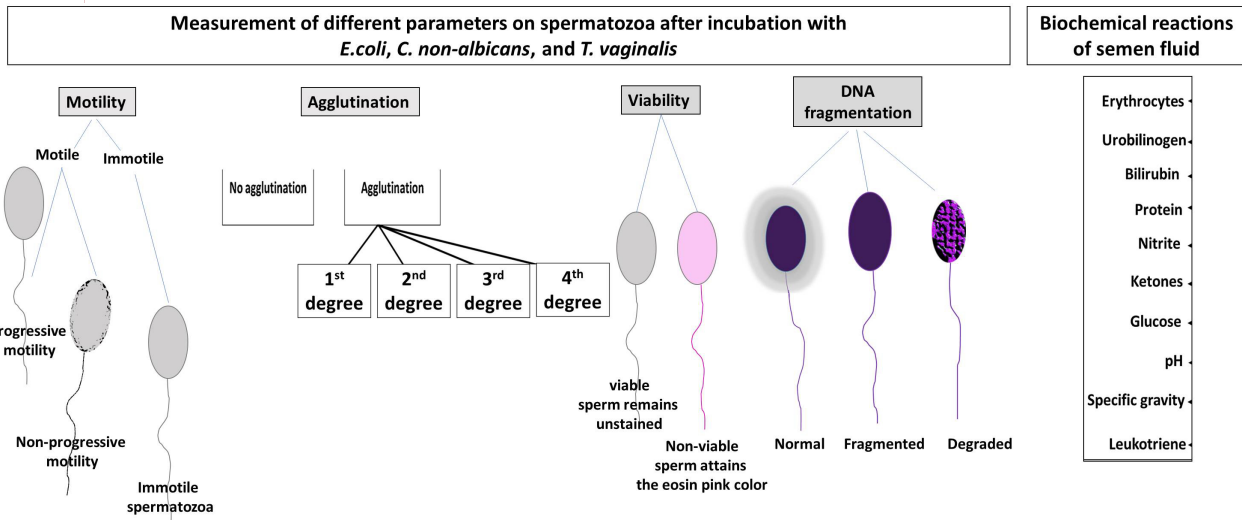


Fig. 1. The paradigm of the different measured parameters was influenced by *Escherichia coli* (*E. coli*), *Candida non-albicans* (*C. non-albicans*), and *Trichomonas vaginalis* (*T. vaginalis*). The figure was generated by the PowerPoint presentation program (version 16, Microsoft, Los Angeles, CA, USA).

Table 1. Standards for healthy semen samples with a reference range of spermatozoa motility, vitality, and count according to the WHO 2021 (the 6th edition).

| Semen parameter in (%) | Reference range |
|--------------------------------|-----------------------|
| Total motility | 42 (40–43) |
| Progressive motility | 30 (29–31) |
| Non-progressive motility | 1 (1–1) |
| Vitality | 54 (50–56) |
| Total sperm count in ejaculate | 39–928 million |
| Ejaculate volume | 1.5–7.6 mL |
| Sperm concentration | 15–259 million per mL |

the different time intervals. The influence of genitourinary pathogenic organisms on different parameters of the semen was assessed as shown in (Fig. 1).

Preparation of *E. coli*

The *E. coli* isolates were grown on sterile plates of blood agar, Chocolate, and MacConkey agar using a sterilized wire loop. All plates were, then, aerobically incubated at 37 °C for 24 hours. Evidence of selective *E. coli* growth was identified by the conventional biochemical reactions. To initiate the experiment, *E. coli* suspension was adjusted approximately at 3×10^6 bacteria/mL in sterile saline (0.85%). Cultures (Brain heart infusion broth, BHI) were finally obtained after 24 and 48 hours.

Preparation of *C. non-albicans*

C. non-albicans isolates were grown on Sabouraud’s dextrose agar (Code: PO0394, Oxoid, Milan, Italy) at 37 °C, overnight, in aerobic conditions, and identified by the API Candida system (version 2.1, bioMerieux SA, Marcy-l’Etoile, France). The *C. non-albicans* suspension was

freshly prepared the day before the experiment by incubation at 37 °C, overnight, in brain heart infusion under aerobic conditions. A final suspension, containing 4×10^6 CFU/mL, was prepared in physiological saline, starting from an initial concentration of 1×10^8 CFU/mL obtained using a 0.5 McFarland standard. The exact concentration of the suspension was verified by a spectrophotometer [13].

Preparation of *T. vaginalis*

The trichomonads were cultured into fresh Diamond’s medium supplemented with antimicrobials (5 Ig/mL of fungizone, 10^3 U of penicillin-G/mL, and 10^2 Ig/mL of streptomycin) until it was free of bacteria. The trichomonads were then cultured on chocolate and mycoplasma agar (Oxoid) for 48 hours in 5% CO₂ at 37 °C. For the experiment, the obtained axenic trichomonads were adjusted at a concentration of approximately 5×10^5 using a Neubauer hemocytometer. Only protozoa cultures with preserved jerky movement $>90\% \pm 10$ were involved and incubated in 5% CO₂ at 37 °C [27].

Assessment of the Influence of Genitourinary Pathogenic Organisms on the Functionality and Physical Parameters of the Semen

Motility

To evaluate the influence of the tested pathogens on sperm motility at diverse time intervals, twenty µL of the semen mixture was obtained on a clean microscopic slide covered with a coverslip and examined underneath an Olympus light microscope (Olympus BX43F, Olympus Corporation, Tokyo, Japan) at magnification 400×. The interpretation was defined as total motility (progressive and non-progressive) or non-motile [28].

Agglutination

The impact of different pathogens on the agglutination of spermatozoa was evaluated using an Olympus light microscope (Olympus BX43F, Olympus Corporation, Tokyo, Japan). Agglutination of motile sperms was identified as head-to-head, tail-to-tail, or head-to-tail agglutination.

Agglutination of mobile spermatozoa to each other was thoroughly examined to exclude non-specific adherence of motile spermatozoa to epithelial debris and adherence of immobile spermatozoa. The interpretation was done as follows; grade 1 (isolated): <10 spermatozoa per agglutinate with many free motile spermatozoa, grade 2 (moderate): 10–50 spermatozoa per agglutinate with many free motile spermatozoa, grade 3 (massive): agglutinates with >50 spermatozoa with only a few free motile spermatozoa, and grade 4 (complete): completely agglutinated spermatozoa with no motile free spermatozoa [25].

Viability

To evaluate the influence of the tested microorganisms on the viability of the spermatozoa, a dye exclusion test using Eosin Y 0.5% aqueous solution was implemented as an indicator dye. For this, equivalent volumes (100 µL) of each semen sample and different microorganisms were thoroughly mixed and incubated at 37 °C. A wet preparation, consisting of 10 µL of the well-mixed reaction mixture and Eosin Y (0.5%), was performed and observed under light microscopy. As a control, the semen sample was mixed and incubated with BHI broth. The results were interpreted as the percentage of live (unstained) spermatozoa and the dead (pink stained).

Biochemical Investigation of the Seminal Fluid

The impact of incubation of urogenital pathogens on the vital biochemical parameters (Protein, Nitrite, Glucose, pH) was qualitatively and semi-quantitatively assessed. The commercially available reagent strip test (Medi-Test Combi 10 ® SGL, item number: MN93067, Macherey-Nagel GmbH & Co., Düren, Germany) was completely immersed in the test tube for 2–3 seconds and immediately read. The diagnostic parameters to each index test were valued using the corresponding cut-off levels (glucose positive at >50 mg/dL, protein at >30 mg/dL whereas nitrite was negative or positive, and pH >7 is alkaline and <7 is acidic [29]. The count of the nitrite positive and nitrite negative samples was estimated using the reagent strip test (Medi-Test Combi 10 ® SGL, item number: MN93067, Macherey-Nagel GmbH & Co., Düren, Germany) after incubation of the semen samples with each of the organisms at the different tested time intervals. Quantitative analysis was done for glucose (RayBio® Glucose Colorimetric Assay Kit, Catalog #: MA-GLU, RayBiotech Inc., GA, USA), protein (colorimetric, Spectrum-Diagnostics Reagents, Catalog #: 310 001, Egyptian Company for

Biotechnology, Cairo, Egypt), and pH (Lit-Control pH Meter, model number: CN175533.9, Devicare, Barcelona, Spain).

Sperm DNA Fragmentation Test

DNA fragmentation was evaluated using a Halosperm® (HT-HS10 batch number 1623, Halotech DNA SL, Madrid, Spain) as per the manufacturer's instructions. The initial acid treatment of the sperm cells denatures the fragmented molecules of the DNA. Thereafter, the lysis solution eliminates the nuclear proteins. In case there is no massive DNA damage nucleoids with large halos emergent from a central core are produced. Spermatozoa with fragmented DNA either do not express a dispersion halo or show a minimal halo from their nucleoids.

$$\text{DNA fragmentation index (DFI\%)} = \frac{\text{small halo+without halo+degraded sperm head}}{\text{total count of spermatozoa}} \times 100\%$$

Interpretation: the reference range is <25% excellent to good DNA integrity, 25–50% fair to poor, and >50% very poor DNA integrity.

Statistics

Statistical package for the Social Sciences (SPSS) version 26 (IBM Corp., Armonk, NY, USA) was used to analyze the data. Data were summarized using mean and standard deviation for quantitative variables and frequencies (number of cases) and relative frequencies (percentages) for categorical variables. Comparisons between groups were done using analysis of variance (ANOVA) with multiple comparisons post hoc Bonferroni test. For comparing categorical data, the Chi-square test was performed [30,31]. *p*-values less than 0.05 were considered statistically significant.

Results

Agglutination

Light microscopy was carried out to assess agglutination in spermatozoa after being incubated with various urogenital pathogens. The results of the control semen sample revealed normal freely mobile spermatozoa with an oval-shaped head and normal neck and tail (Fig. 2A). Incubation of spermatozoa with *E. coli* showed adherence of the bacilli to the tails and heads of the spermatozoa (Fig. 2B) resulting in progressive agglutination at different time intervals. *C. non-albicans* resulted in tail-to-tail agglutination of the spermatozoa during the first hour. Progressive adherence of *C. non-albicans* to the heads of the spermatozoa was also observed starting from the 2nd hour of incubation (Fig. 2C) with triggering of the three forms of agglutination (head-to-head, tail-to-tail, or head-to-tail). *T. vaginalis*, head-to-head agglutination was scarcely seen (Fig. 2D).

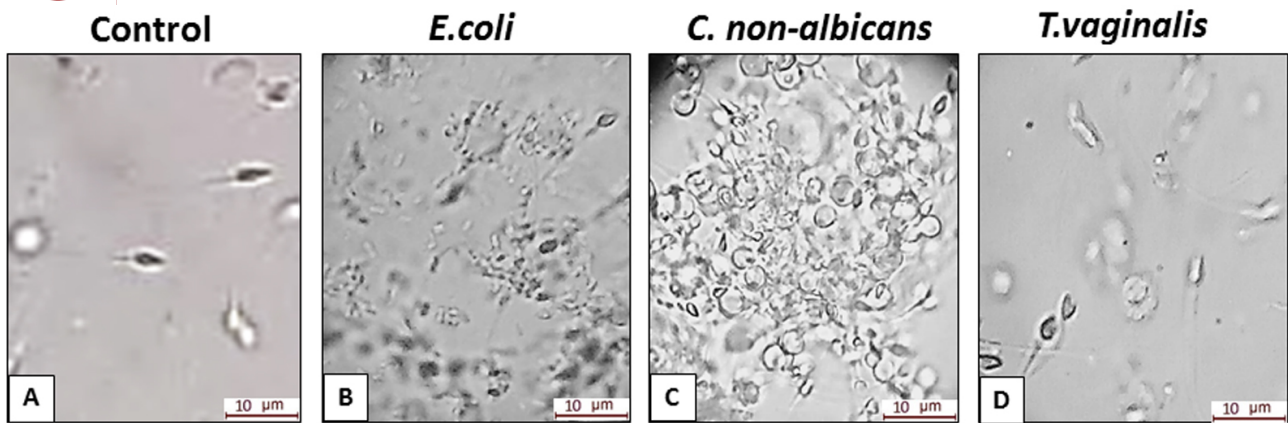


Fig. 2. Photograph showing agglutination and viability of spermatozoa in semen fluid after co-incubation with 24-hour cultures for 30 minutes. (A) Control samples, (B) *E. coli*, (C) *C. non-albicans* and (D) *T. vaginalis*. Note control sample shows no agglutination and viable sperm while the pathogen incubated samples show agglutination of spermatozoa and dead sperm (scale bar: 10 µm) (Images are captured via an Olympus light microscope (Olympus BX43F, Olympus Corporation, Tokyo, Japan)).

E. coli

E. coli-48-hour culture showed significant reductions in seminal glucose (mg/dL) at ½, 3.5, and 4 hours of co-incubation compared with healthy semen (11.5 ± 2.26 versus 14.83 ± 1.47 , 4 ± 1.26 versus 6.83 ± 2.48 , and 2.83 ± 1.47 versus 5.83 ± 2.48 respectively) (p -value < 0.05). Seminal protein (mg/dL) was not altered at different time intervals with the *E. coli* cultures (p -value > 0.05); yet, seminal nitrite was predominantly positive (p -value < 0.05). After 4 hours of co-incubation, *E. coli* cultures resulted in increased pH values (8.8 ± 0.21 and 8.87 ± 0.16 respectively) compared with healthy semen (8.18 ± 0.51) (p -value < 0.05).

Regarding the effect of *E. coli* on agglutination at different time intervals, the first thirty minutes of co-incubation with the *E. coli* cultures showed grade 2 agglutination of spermatozoa (49.17 ± 11.14 and 55.83 ± 9.7 respectively versus 1.83 ± 1.94 in the control samples) (p -value < 0.05). By the end of the fourth hour, agglutination progressed into grade 4 (97 ± 2.61 and 98.5 ± 1.05 respectively versus 15.5 ± 2.07 in the control samples) (p -value < 0.05). Compared with the healthy uninfected semen, incubation with the *E. coli* cultures exhibited a significant reduction in motility ($p < 0.05$) at different time intervals whereas, the percentage of viability was not affected ($p > 0.05$) (Fig. 3).

C. non-albicans

The *C. non-albicans* cultures did not result in significant alterations in the seminal protein and glucose at all the tested time intervals. At 3 and 4 hours of co-incubation, the *C. non-albicans*-24-hour culture showed lower pH values (7.7 ± 0.11 , 7.57 ± 0.1 , and 7.4 ± 0.13 respectively) versus healthy semen (8.18 ± 0.51) with p -value < 0.05 . Also, the *C. non-albicans*-48-hour culture exhibited reduced pH val-

ues at 3.5 hours and 4 hours of co-incubation (7.62 ± 0.08 and 7.45 ± 0.1 , respectively) compared with the healthy semen ($p < 0.05$).

Incubation of the semen samples with *C. non-albicans* revealed a significant increase in agglutination grade to grade 3 (84.17 ± 8.61) and grade 4 (95.5 ± 3.27) agglutination after 4 hours of co-incubation with the 24-hour and 48-hour cultures respectively ($p < 0.05$). The *C. non-albicans* 24-hour and 48-hour cultures resulted in a significant reduction in spermatozoal motility ($p < 0.05$) at the different tested time intervals while the percentage of sperm viability was not significantly altered ($p > 0.05$) (Fig. 4).

T. vaginalis

Incubation of the semen with the 24-hour and 48-hour *T. vaginalis* cultures resulted in significant reductions in seminal glucose (mg/dL) within the 4-hour incubation period (p -value < 0.05). Also, the *T. vaginalis*-48-hour culture showed a significant increase in seminal protein (mg/dL) after 1.5 hours of co-incubation.

The pH values were also significantly reduced with both the 24-hour and 48-hour *T. vaginalis* cultures at 2 to 4 hours of semen co-incubation. Through the experiment, the seminal nitrite was not altered with neither the 24-hour nor the 48-hour *T. vaginalis* cultures.

Sperm agglutination was scarcely seen with both the 24-hour and 48-hour cultures of *T. vaginalis* and was almost always isolated agglutination (grade 1) with a statistically insignificant mean difference relative to the control samples. The 24-hour and 48-hour *T. vaginalis* cultures also resulted in a significant reduction in sperm motility relative to the healthy uninfected semen ($p < 0.05$) at the different tested time intervals. Nevertheless, viability was not affected (Fig. 5).

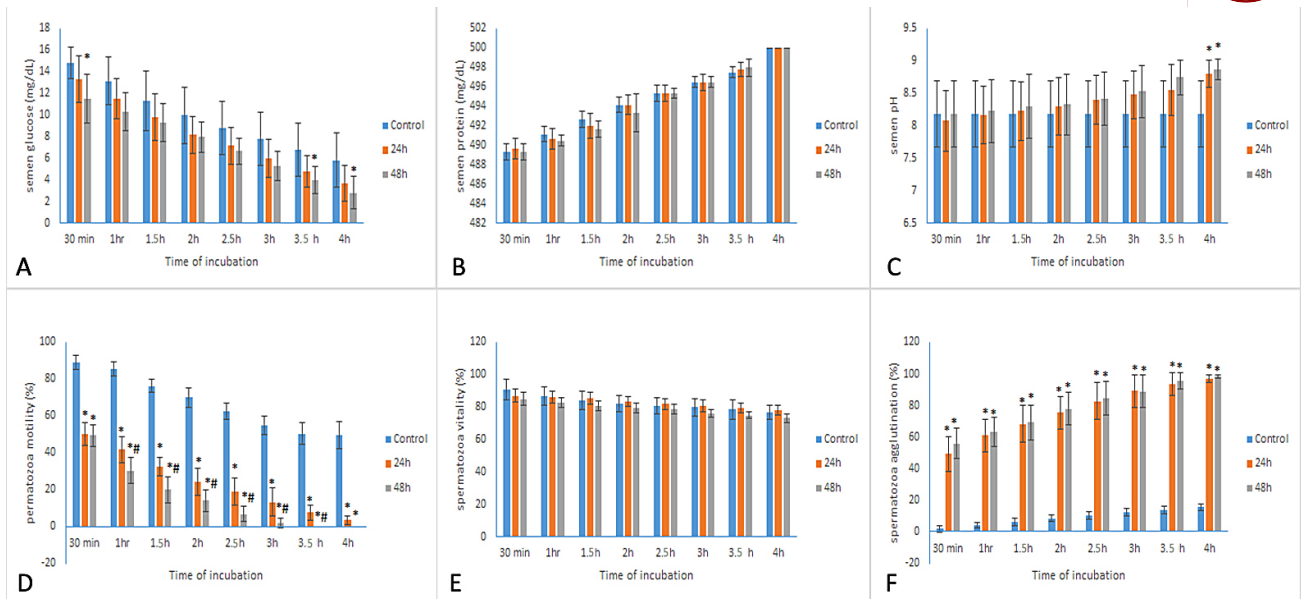


Fig. 3. Alterations in semen parameters following *in-vitro* cultivation with the *E. coli* cultures at different time intervals. (A) Semen glucose. (B) Semen protein. (C) Semen pH. (D) Spermatozoa motility. (E) Spermatozoa vitality. (F) Spermatozoa agglutination. Data are represented as mean \pm SD. (*) Statistically significant compared to the healthy control (n = 6). (#) Statistically significant compared with 24 hours at $p < 0.05$ using analysis of variance (ANOVA) multiple comparisons with post hoc Bonferroni test.

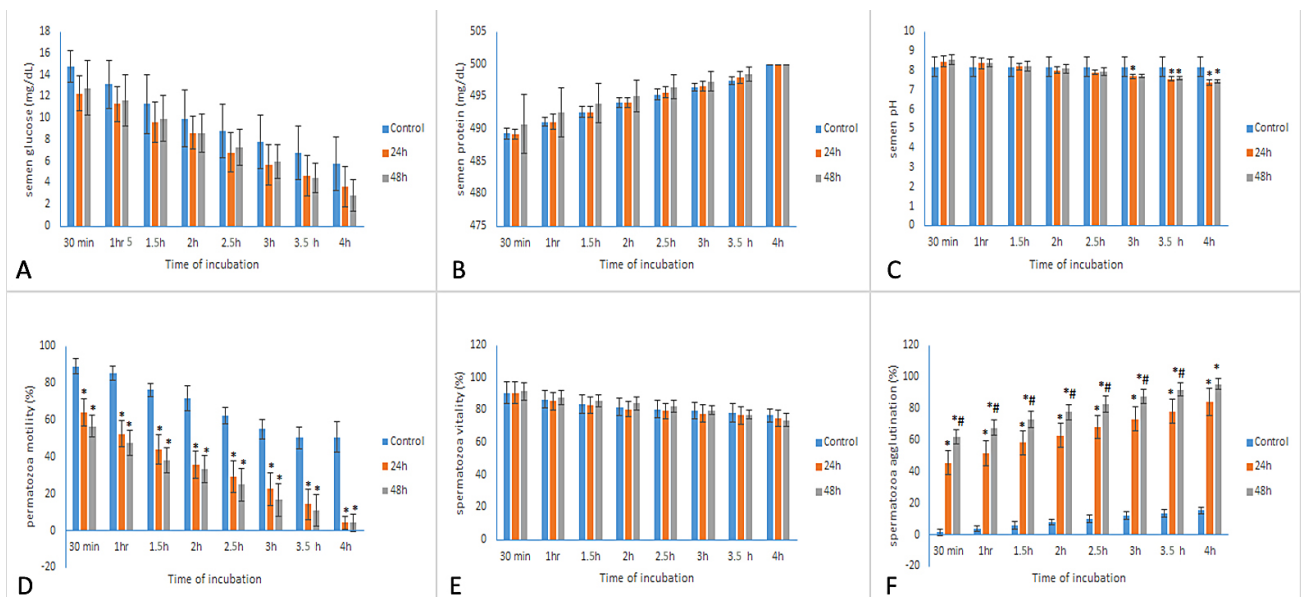


Fig. 4. Alterations in semen parameters following *in-vitro* cultivation with *C. non-albicans* cultures at different time intervals. (A) Semen glucose. (B) Semen protein. (C) Semen pH. (D) Spermatozoa motility. (E) Spermatozoa vitality. (F) Spermatozoa agglutination. Data are represented as mean \pm SD. (*) Statistically significant compared to the healthy control (n = 6). (#) Statistically significant compared with 24 hours at $p < 0.05$ using ANOVA multiple comparisons with post hoc Bonferroni test.

DNA Fragmentation

The spermatozoa incubated with the urogenital pathogens in the current study showed high DFI% indicating poor integrity of the DNA molecule. *C. non-albicans* co-incubated with semen (group 2) showed the highest DFI% (97%) (p -value < 0.05) with thirty percent having degraded DNA whereas a big halo in non-fragmented DNA

was as low as 1%. Nevertheless, there were no significant differences in the average DFI% between sperms incubated with *E. coli* (group 3) (56%) or with trichomonads (57%) (group 4) (p -value > 0.05) (Fig. 6). Values of fragmented DNA (small halo, without halo, and degraded) were also evaluated.

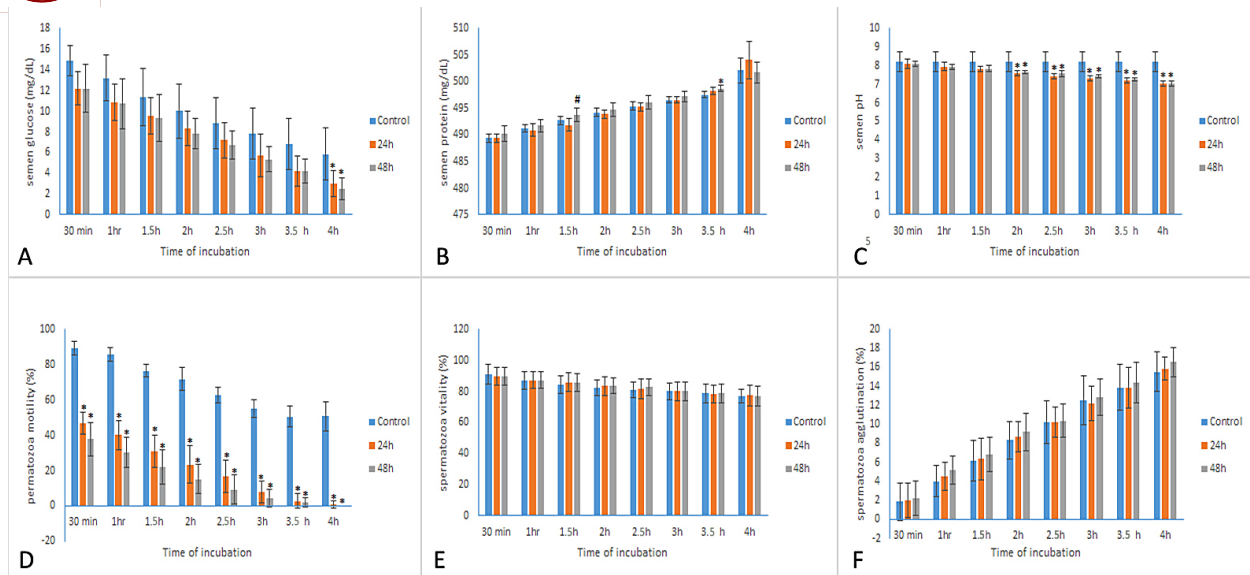


Fig. 5. Alterations in semen parameters following *in-vitro* cultivation with 24-hour and 48-hour-old cultures of *T. vaginalis* at different time intervals. (A) Semen glucose. (B) Semen protein. (C) Semen pH. (D) Spermatozoa motility. (E) Spermatozoa vitality. (F) Spermatozoa agglutination. Data are represented as mean \pm SD (n = 6). (*) Statistically significant compared to the healthy control, (#) Statistically significant compared with 24 hours at $p < 0.05$ using ANOVA multiple comparisons with post hoc Bonferroni test.

Discussion

The current study has been designed to investigate the effect of *in-vitro* incubation of pathogens on the semen functional and chemical parameters. *E. coli* induced progressive agglutination and clustering of spermatozoa in a time-dependent manner. *E. coli* is the most frequent bacteria to induce agglutination of spermatozoa (54.7%) followed by *Proteus sp.*, *Streptococcus agalactiae*, and *Enterococcus fecalis*. This reflects the great capability of *E. coli* to directly interact with the sperms [32,33].

E. coli co-incubation with semen in the current study also resulted in increased pH values. This alkaline pH suppresses the cell division protein (FtsN) in *E. coli* because *E. coli* requires a Na^+/H^+ antiporter to proliferate at alkaline pH [34].

Nitrite increased predominantly in the semen incubated with *E. coli*. This could be explained by the ability of *E. coli* to import Nitrate and reduce it into nitrite. *E. coli* then expels nitrite and reduces it into ammonia. This mechanism protects the cytoplasm of *E. coli* from extreme nitrite toxicity [35]. Increased nitrite in seminal fluid has been previously linked to oligo- and azoospermic [36].

Seminal glucose was also reduced following the co-incubation of semen with *E. coli* in the current study. This is concordant with the previous reports illustrating the strong association between glucose consumption and energy production with the oxidation-reduction reactions, and growth of *E. coli* [37].

In the current work, the 48-hour culture of *E. coli* resulted in a significant reduction in spermatozoa motility when compared with the 24-hour culture and control. This

could be attributed to the soluble factors produced by *E. coli* which can affect sperm functions [38]. The effect of *E. coli* on spermatozoa is dependent on the bacterial load [39,40].

The current work revealed no significant differences in the protein concentration nor the viability of spermatozoa between the *E. coli*-incubated and the control semen samples. The Semenogelin (Sg) protein, a major constituent of the semen coagulum, activates the hyaluronidase enzyme inducing an anti-microbial effect. "Eppin" is another seminal protein component that induces antibacterial activity through alteration of the permeability of the inner and outer membranes of *E. coli*, thereby protecting spermatozoa [41,42].

DNA fragmentation increased up to 56% after incubation with *E. coli* (p -value < 0.05). This could be attributed to the capacity of *E. coli* to directly attach to the sperms and their ability to trigger oxidative stress and apoptotic damage [4]. In a prior study, comparable deterioration in semen quality was recorded after incubation with different concentrations of enterotoxigenic and verotoxigenic *E. coli* [43].

Candida non-albicans cultures in the current study resulted in a significant reduction in the motility of spermatozoa at the different tested time intervals. *Candida sp.* has been previously proven to directly adhere to the sperms and damage their ultrastructure [44]. *Candida* also releases extracellular sperm immobilization factors that hinder sperm motility [45]. Co-incubation with *C. non-albicans* in the current study also induced marked sperm agglutination (tail-to-tail, head-to-head, and head-to-tail). Agglutination is always associated with morphological changes in sper-

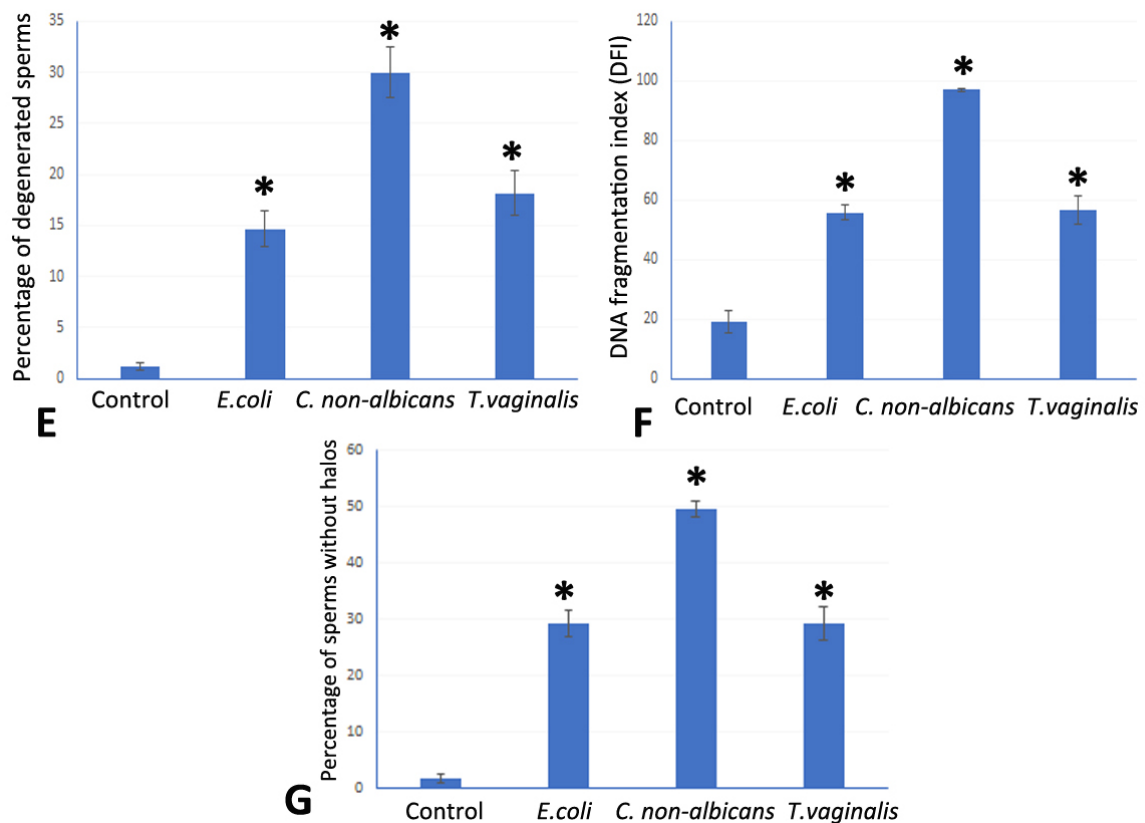
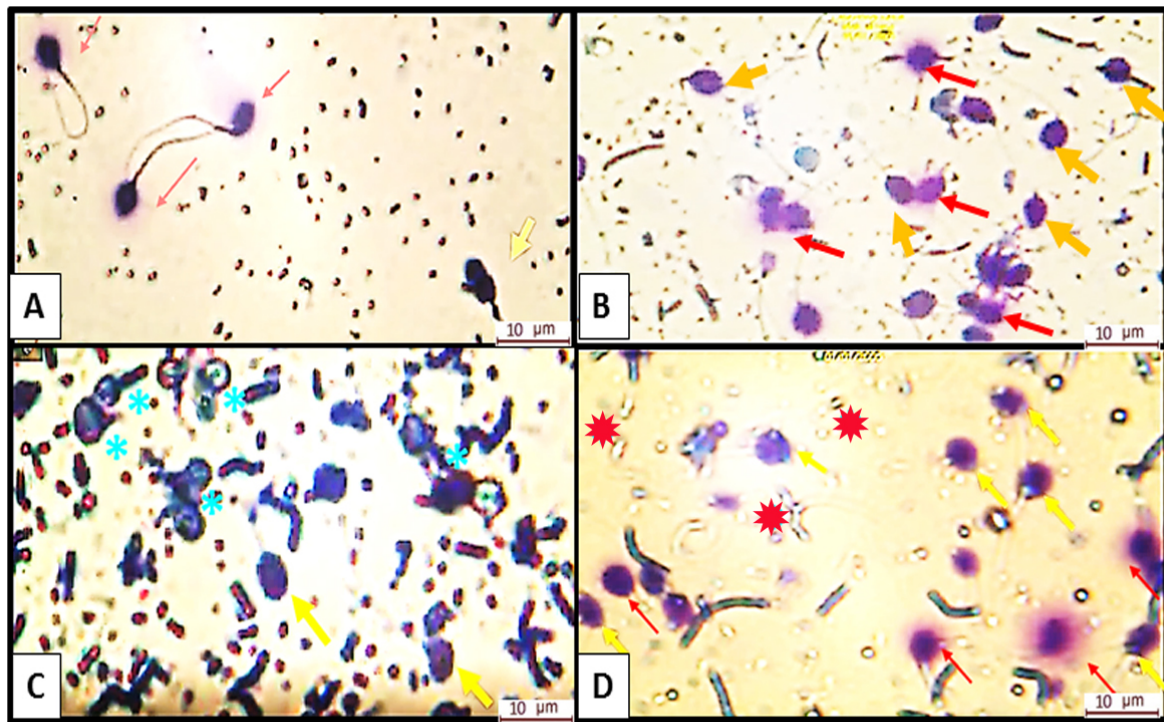


Fig. 6. DNA fragmentation after 4 hours of co-incubation with the 24-hour cultures. (A) Control. (B) *E. coli*. (C) *C. non-albicans*. (D) *T. vaginalis*. Note: Blue asterisks point to *C. non-albicans*, red asterisks point to degraded sperms with pin and double-head, yellow arrows point to sperms without halo, and red arrows point to sperm with a big or medium halo (scale bar: 10 μ m). (E) Percentage of degenerated sperms. (F) DNA fragmentation index (DFI%). (G) Percentage of sperms without halos. Data are represented as mean \pm SD (n = 6). (*) Statistically significant compared to the healthy control at $p < 0.05$ using ANOVA multiple comparisons with post hoc Bonferroni test. Images are captured via an Olympus light microscope (Olympus BX43F, Olympus Corporation, Tokyo, Japan).

matozoa, with essential interfering with sperm motility. This agglutination effect is also dependent on the incubation time and the concentration of the organisms [13,46].

In the current work, the pH of semen has substantially dropped following co-incubation with *C. non-albicans* in a time-dependent manner. The alkaline seminal pH preserves the quality and viability of the spermatozoa [47]. Any modification of the microenvironment of seminal plasma leads to excessive energy consumption and deterioration of sperm motility [48].

In the current study, co-incubation with *C. non-albicans* increased the DFI% up to 97%. DNA fragmentation is closely associated with sperm apoptosis [49] and is regulated by a caspase-regulated DNase complex known as the DNA fragmentation factor [50]. DNA fragmentation predisposes to oxidative stress [51] and latent chromosomal damage [52] because it is closely related to condensation of chromatin material, nuclear fragmentation, and sperm apoptosis [53]. Detection of *Candida* infection has been proposed as an essential factor to avoid poor pregnancy outcomes [54,55].

In the current study, agglutination was hardly seen with the *T. vaginalis* cultures. Despite the ability of *T. vaginalis* to induce tropism of the spermatozoa (for the purpose of phagocytosis), yet, the trichomonads undergo immediate necrosis when they come in contact with the sperms [56]. This comes in agreement with our results which revealed that prokaryotic organisms are more efficient in triggering agglutination rather than the eukaryotic *Trichomonas*.

In the current study, motility was markedly reduced upon incubation with *T. vaginalis*. *T. vaginalis* produces extracellular polymeric substances (EPS) that exert an inhibitory effect on semen motility [57]. Also, the rotation of *T. vaginalis* trophozoites hinders the horizontal movement of the spermatozoa [58]. Co-incubation of semen with *T. vaginalis* cultures showed a reduction in seminal PH. Low pH in semen impairs the cell membrane of the sperm or increases the active content of oxygen, thus in turn distresses the motility of sperm and its capacitation [59].

T. vaginalis co-incubation in the current study resulted in a reduction of the seminal glucose. *T. vaginalis* is recognized to gather considerable amounts of the polysaccharide glycogen that serves as a carbon and an energy stock. Yet, glycogen stores are dependent on the extracellular concentrations of glucose [60]. Glucose is the chief source of ATP generation in the hydrogenosome of the parasite [61]. Interestingly, survival of *T. vaginalis* was found to be enhanced in a glucose-restricted microenvironment where it could benefit from low glucose levels [62]. A recent study demonstrated the presence of Na⁺/K⁺-ATPase in the sperm flagellar membrane. This Na⁺/K⁺-ATPase is important for the uptake of glucose, production of ATP, and induction of sperm motility [63]. This could explain the relation between the reduction of seminal glucose and the impaired sperm motility observed with the *T. vaginalis* cultures in the current study.

Another remarkable observation in the current study was the increased seminal protein after co-incubation with *T. vaginalis*. A previous study has reported that *T. vaginalis* produces protein metabolites that increase the viscosity of seminal fluid and exert a lethal effect on the sperms [64].

The semen samples incubated with *T. vaginalis* in our study recorded a 57% increase in the DFI%. This could be attributed to the excretory/secretory metabolites produced by *T. vaginalis* that induce DNA fragmentation, reduce mitochondrial membrane potential, and initiate sperm apoptosis [65]. Our work revealed that co-incubation with *T. vaginalis*, on the other hand, did not exhibit alteration in spermatozoa viability. The extracellular polymeric substances produced by *T. vaginalis* have been previously reported to affect the viability of sperm in a time-dependent manner [57].

Limitations of the current study are to be considered. We recommend a wider scale of studies on the genome of *C. non-albicans*, *T. vaginalis*, and *E. coli*. Care should also be attained during the interpretation of the *in-vitro* studies into the clinical practice since the situation of genitourinary infections could differ significantly from the *in-vitro* incubation of microorganisms. Future studies should be designed to analyze the semen chemical and functional parameters among different genitourinary infection patients.

Conclusion

Incubation of semen with *E. coli* resulted in a significant reduction in seminal glucose and an increase in seminal pH. The seminal protein remained unaltered throughout the experiment; yet, the seminal nitrite was positive. *E. coli* also induced agglutination through its attachment to the tail of spermatozoa resulting in progressive agglutination that proceeded into grade 4 by the end of the fourth hour. Sperm motility was significantly reduced while viability was not affected. *C. non-albicans* cultures did not show alterations in seminal protein, glucose, nor viability of the spermatozoa. Yet, pH and motility were reduced compared to the control semen samples. Also, *C. non-albicans* resulted in grade 4 agglutination with its three forms (head-to-head, tail-to-tail, or head-to-tail agglutination). *T. vaginalis* reduced seminal glucose, pH, and motility and increased seminal protein. Nitrite and viability appeared not to be affected, and agglutination was scarcely seen. The DFI% was increased in all the tested organisms, yet, the *C. non-albicans* exhibited the highest DNA fragmentation index. The increase in DFI% was statistically non-significant in both *E. coli* and *T. vaginalis*.

Availability of Data and Materials

Data are available upon reasonable request from the corresponding author.

Author Contributions

MA and EAES: designed the research study. RYS, HE, ASG, MFF, BEA and AM: acquisition of data, analysis and interpretation of data. HMA and EAA: conceptualization. All authors contributed to the drafting and critical revision of the manuscript. All authors have read and approved the final manuscript. All authors have participated sufficiently in the work and agreed to be accountable for all aspects of the work.

Ethics Approval and Consent to Participate

This study was approved by the ethical committee of the Armed Forces College of Medicine, Cairo, Egypt (No. 265 dated 17/6/2023). All participants have been informed verbally and a written consent has been signed before any step according to Amsterdam's declarations. All procedures involved in the current study were performed following the ethical standards of the National Research Committee and with the 1964 Helsinki Declaration and its ensuing regulations.

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Conflict of Interest

The authors declare no conflict of interest.

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