

Reactive oxygen species, total antioxidant concentration of seminal plasma and their effect on sperm parameters and outcome of IVF/ICSI patients

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Abstract

Objective The purpose of this study was to determine and compare the concentration of reactive oxygen species (ROS) and total antioxidant (TAS) in seminal plasma of IVF (in vitro fertilization) and ICSI patients, to establish their effect on sperm quality (count, vitality, HOS, morphology, maturity, DNA strand breaks) and assess the fertilization potential of spermatozoa and IVF/ICSI outcome.

Method IVF/ICSI patients ($n = 48$) 26 IVF and 22 ICSI were included in this study. A spermogram was generated from each patient one-hour post ejaculation and smears were made from each semen sample to evaluate the morphology, sperm maturity (Chromomycin CMA₃) and DNA strand breaks (Terminal deoxyribonucleotidyl transferase-mediated dUTP nick-end labelling, TUNEL-assay).

Results In both groups a negative correlation was found between ROS concentration in seminal plasma and sperm vitality ($r = -0.111$; $P = 0.453$); membrane integrity and morphology (-0.141 ; $P = 0.340$) and fertilization rate ($r = -0.0290$; $P = 0.045$). However, TAS in seminal plasma correlated positive with fertilization rate ($r = 0.081$; $P = 0.584$). In addition, an inverse correlation was found between sperm DNA strand breaks (TUNEL-test) and spermatozoa global and progressive motility, vitality, and membrane integrity. Furthermore, the mean percentage of

normal condensed spermatozoa (CMA₃) was significantly higher ($P = 0.0001$) in patients undergoing IVF compared to ICSI. Spermatozoa of male ICSI patients were more susceptible to acid denaturation (acridine orange staining) compared to spermatozoa of male IVF patients ($P = 0.041$). However, ROS concentration was higher in IVF patients compared to ICSI patients (94.73 ± 102.84 vs. 54.78 ± 39.83 $\mu\text{mol/l}$), whilst TAS levels (1.43 ± 0.28 vs. 1.53 ± 0.22) and fertilization rate (67.26 vs. 67.26) were similar in both groups.

Conclusion ROS concentration and other sperm parameters were higher in IVF compared to ICSI patients. TAS concentration was comparable between the two groups. However, the fertilization rate was similar in IVF and ICSI patients. Therefore, ROS concentration in seminal plasma affects the quality of spermatozoa but does not affect the fertilization rate in IVF/ICSI cycles.

Keywords ROS · TAS · IVF · ICSI · Spermatozoa

Introduction

Reactive oxygen species (ROS) are free radicals that have a significant role in many of the sperm physiological processes such as capacitation, hyperactivation, and sperm-oocyte fusion [5, 6, 17]. In the presence of polyunsaturated fatty acids such as sperm plasma membrane, ROS triggers a chain of chemical reactions called lipid peroxidation [41, 81]. Furthermore, it has been shown that ROS can damage DNA by causing deletions, mutations, and other lethal genetic effects [55, 75]. Moustafa et al. [55] demonstrated that infertile patients with high ROS levels in their seminal plasma had a higher percentage of apoptosis than normal healthy donors.

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There are two main sources of ROS in semen: leukocytes and immature spermatozoa [25]. Gomez et al. [28] demonstrated that levels of ROS produced by spermatozoa were negatively correlated with the quality of sperm in the original semen. Poor sperm quality is linked to increased ROS generation as a consequence of the presence of excess residual cytoplasm. Cytoplasmic droplets after spermiation are thought to be immature and functionally defective [8] and are capable of producing increased amounts of ROS [65]. A strong positive correlation exists between immature spermatozoa and ROS production, which in turn negatively affects the sperm quality [26, 59]. However, the cellular antioxidant mechanism present in almost all tissues and their secretions are likely to quench these ROS and protect against oxidative damage [38]. In addition, many studies have demonstrated that infertile men have an impaired seminal plasma non-enzymatic antioxidant capacity than fertile men, suggesting an association between decreased total antioxidant capacity and male infertility [47, 64]. Pasqualotto et al. [57] reported that levels of antioxidants in seminal plasma from infertile men were significantly lower than levels in fertile controls.

Excessive ROS production that exceeds critical levels can overwhelm all the antioxidant defence systems of spermatozoa and seminal plasma causing oxidative stress [16, 65]. Oxidative stress is a condition in which the elevated levels of ROS damage cells, tissues and organs [53, 62, 63] and is thought to be a mediator of sperm dysfunction and male infertility [2].

The semen quality score and ROS levels in semen samples appear to be strongly associated with male factor infertility, since both these parameters are more sensitive than individual sperm parameters in identifying male factor infertility [56]. A recent meta-analysis by Agarwal et al. [1] found that ROS levels correlated significantly with in vitro fertilization (IVF) rate (estimated overall correlation -0.374 , 95% CI, -0.520 to -0.205).

Therefore, the purpose of this study was to determine and compare ROS and TAS concentration in seminal plasma of IVF and ICSI cycles, their effect on sperm quality and consequently the fertilization potential of spermatozoa in both IVF and ICSI cycles.

Materials and methods

Materials

Forty eight male partners of females undergoing controlled ovarian hyperstimulation for IVF ($n = 26$) or ICSI ($n = 22$) were included in this study. Semen was collected by masturbation after 3 days of sexual abstinence.

Methods

Basic sperm parameters evaluation

In all patients, a standard semen analysis was performed according to WHO guidelines [79]. Morphology was assessed according to strict criteria described by Kruger et al. [44]. Semen samples were loaded onto a 45 over 90% discontinuous PureSperm[®] density gradient (Nidacon International AB, Sweden) and centrifugated at $500\times g$ for 20 min at room temperature. The supernatant (seminal plasma) was immediately separated, and examined before storage to rule out the presence of spermatozoa in the supernatant. The seminal plasma was aliquoted into storage ampoules and stored at -80°C until the assay was performed (within 3 months). Repeated assays on fresh and frozen seminal plasma samples from the same patient showed that samples stored at -80°C did not significantly influence ROS, TAS concentrations in the seminal plasma.

The pellet containing the normal spermatozoa was washed, layered with 1 ml of culture media and incubated at 37°C . The motile sperm was used for injection (ICSI) or insemination (IVF) of oocytes.

Smears were prepared pre and post semen processing for morphological examination and chromatin condensation analysis, visible after staining (Chromomycin CMA₃ DNA single strand breaks (acridine orange) staining and by terminal deoxynucleotidyl transferase (TDT) mediated dUTP nick end labelling (Terminal deoxyribonucleotidyl transferase-mediated dUTP nick-end labelling, TUNEL-test).

Eosin test (viability assessment)

The eosin test was performed according to the method described by Eliasson and Treich [20]. Briefly, one drop of semen was mixed on a slide with 1 drop of the 0.5% aqueous yellowish eosin solution and one drop of Nigrosin (10% in distilled water) and covered with a cover slip. After 1–2 min, the dead spermatozoa stained red, whilst live sperm remained unstained. Nigrosin was used as a counter-stain to facilitate visualization of the unstained live cells. One hundred sperm from each semen sample were evaluated per slide.

Assessment of membrane integrity of spermatozoa (hypoosmotic swelling test; HOS-test)

Hypoosmotic swelling test was carried out according to the method described by Jeyendran et al. [37]. One hundred microliters of sperm suspension added to 1 ml of hypoosmotic solution (equal parts of 150 mOsmol fructose and 150 mOsmol sodium citrate solutions), followed by 60 min incubation at 37°C . After incubation, a minimum of 200

spermatozoa were examined per slide under the light microscope and the percentage of spermatozoa that showed typical tail abnormalities (curly tail) indicative of swelling were calculated.

Assessment of sperm morphology

Morphology was evaluated according to strict criteria described by Kruger et al. [44], and taking into consideration the specific effects, e.g., size alteration of a specific staining method. Briefly, a total of 5 μ l of semen were pipetted onto a slide to make thin smears. The slide was air-dried for 3 min, then fixed for 15 s in methyl alcohol. It was stained with Papanicolaou staining. Bright field illumination with a magnification of 100 \times under an oil immersion objective was used for the evaluation. Two hundred spermatozoa were evaluated in each sample. At least ten high power fields from different areas of the slide were examined. The following criteria for normal spermatozoa were used: (1) smooth, oval head, 5–6 μ m in length and 2.5–3.5 μ m in diameter; (2) well defined acrosome, 40–70% of the sperm head; (3) mid-piece without defect, slender, axially attached with <1 μ m; (4) tailpiece uniform, free from kinks, uncoiled, width thinner than mid-piece, length-45 μ m; and (5) cytoplasmic droplets (remnants), which compromise less than half the head area, are acceptable but must be retained in the mid-piece region only. Multiple defects in individual cells were classified as amorphous unless there were only multiple tail defects. The semen samples were either classified as normal (\geq 14% morphologically normal spermatozoa) or abnormal (<14% morphologically normal spermatozoa).

Assessment of chromatin condensation (chromomycin CMA₃) of spermatozoa

CMA₃ was performed as previously described by Bianchi et al. [12]. Briefly, a semen aliquot was washed in Dulbecco's Ca–Mg free phosphate buffer saline (PBS) and centrifuged at 300 \times g for 10 min. The spermatozoa were washed, fixed in methanol/glacial acetic acid (3:1) at 4°C for 5 min and then spread on clean slides. Each slide was treated for 20 min with 100 μ l of CMA₃ solution at 25°C in the dark (Sigma St Louis, MO, USA).

Slides were then washed and rinsed in PBS buffer and mounted with buffered glycerol. Fluorochrome was examined using a Zeiss photomicroscope III using a combination of excite dichroic barrier filters of BP 436/10:FT 580:LP 470. A total of 200 spermatozoa were evaluated on each slide. Evaluation of CMA₃ staining is done by distinguishing spermatozoa that stain bright yellow (CMA₃ positive) from those staining a dull yellow (CMA₃ negative) [31].

Assessment of chromatin structure integrity of spermatozoa, single strand breaks (acridine orange-staining)

Ten microliters of each semen sample was spread onto a pre-cleaned slide that was allowed to dry at room temperature. Slides were then fixed overnight in freshly prepared Carnoy's solution (three parts methanol and one part glacial acetic acid) and allowed to air dry for a few minutes before being stained with AO (CI 46005; sigma Chemical Company, St Louis, MO, USA).

Slides were stained in AO solution for 5 min at ambient temperature in the dark in an aluminium foil covered 500 ml glass container, gently rinsed with deionised water, and mounted with phenylenediamine mounting medium. The AO slides were kept in the dark and analysed by means of a fluorescence microscope (Leitz, Oberkochen, Germany) equipped with a 490-nm excitation filter and 530-nm barrier. At least 200 spermatozoa were assessed per slide. Spermatozoa with normal DNA content revealed a distinct green fluorescent colour, whereas sperm heads displaying a fluorescent spectrum varying from yellow–green–red were considered as being denatured (Abnormal DNA; 74].

Assessment of sperm DNA strand breaks (TUNEL-assay)

DNA fragmentation was assessed using the TUNEL assay. A detection kit (Apoptosis Detection System Fluorescence; Promega, Mannheim, Germany) was used. The procedure was done according to the manufacturer's instructions.

Briefly Ejaculated sperm samples were separated from seminal plasma by slow speed centrifugation (250 \times g; 10 min). The supernatant was discarded, and the remaining pellet was washed in phosphate-buffered saline (PBS), pH 7.4. A droplet of this sperm suspension was smeared onto pre-treated slides (Superfrost; Menzel, Braunschwig, Germany), air dried and fixed by immersion in freshly prepared 4% methanol-free formaldehyde in phosphate-buffered pH 7.4 saline at 4°C for 25 min. The labelling reaction was carried out for 1 h in the dark at 37°C. The slides were then washed in fresh PBS for 25 min at 4°C. The slides were then washed further in fresh PBS for 5 min at room temperature, treated with 2% triton X-100 in PBS for 5 min, and rinsed several time in PBS for another 5 min at room temperature.

After the final rinse, excess water was drained off using a drop of anti fade solution and the sample were covered with a cover slip. Two hundred spermatozoa from various fields of view were immediately analysed with an epifluorescence microscope (Zeiss, Oberkochen, Germany) at 1,000 \times magnification. The percentage of green-fluorescing sperm (TUNEL-positive) was determined. Negative controls without TdT enzyme were prepared for each batch of analyzed slides.

Measurement of ROS concentrations in seminal plasma

The concentration of ROS was measured by a Colorimetric assay for the quantitative determination of peroxides in EDTA–plasma, serum and other biological fluids (Oxy Stat; Cat. No. BI-5007 Biomedica Medicine product GmbH & Co KG, Wien Austria).

Assay characteristics The Biomedica Oxy Stat assay measures the total concentration of peroxides, which is formed in the propagation-phase of the low-density lipoprotein oxidation process using peroxidase/TMB, 12 × 8 well microtiter plate format. Ninety-six tests per micro titer plate. Reference values EDTA plasma <400 μmol/l, serum <350 μmol/l. Measuring range 7–6,000 μmol/l; detection limited 7 μmol/l. Sample volume 10 μl/test. Assay time 30 min.

Measurement of TAS concentrations in seminal plasma

Semen antioxidant capacity was determined by a method described by [52], and Rice-Evance and Miller [58], developed for the evaluation of antioxidant capacity in blood plasma. This is a Colorimetric assay using ABTS[®] (2, 2'-Azino-di-[3ethylbenzthiazoline sulphonate], and a commercially available kit, TAS (Randox Laboratory, Ltd Krefeld, Germany). Range 1.30–1.77 mmol/l plasma [52].

IVF/ICSI procedure

The female partners underwent controlled ovarian hyperstimulation, as described earlier by Hammadeh et al. [30] by pituitary desensitization using gonadotrophin-releasing hormone analogue (Gn-RHa) in the mid luteal phase. Thereafter, stimulation with follicle stimulating hormone (FSH Gonal-F, Serono, Germany) was performed. When at least three follicles were >18 mm in diameter and the serum oestradiol concentration > 800 pg/mL, ovulation was induced by administration of 10,000 IU human chorionic gonadotrophin (hCG). Transvaginal follicle aspiration was carried out 36 h post hCG. Retrieved oocytes were cultured in Vitro Life culture media (Sweden) at 37 h, in 5% CO₂. The female patients underwent either IVF (*n* = 26) or ICSI (*n* = 22) treatment. Patients that achieved a pregnancy were aged 32.5 ± 4.0 years and patients that were unsuccessful were aged 34.0 ± 3.0 years.

Four hundred and eight oocytes were retrieved (278 from IVF and 200 from ICSI patients). Fertilization was recorded after 18–24 h post pronuclei assessment. Fifteen ongoing pregnancies were achieved (31.3%). The study protocol was approved by the institutional review board, University of Saarlandes, Germany, and written consent was obtained from all patients participating in this study.

Statistical analysis

Statistical analysis was performed using SPSS 11 for Windows Software Package (SPSS Inc., Chicago, IL, USA).

The relationship between ROS, TAS and DNA integrity, motility and morphology and their effect on fertilization rate after IVF and ICSI treatment was analysed. The Mann–Whitney (*U* test) was used for non-paired data. The results were presented by the mean ± SD values. Correlations were analysed by means of chi-square test or exact Fisher-test. Variants analysis was performed with the Kruskal–Wallis test and Levene-test. Differences were considered statistically significant at *P* < 0.05.

Results

The mean concentration of spermatozoa (73.9 ± 34.9 mill/ml), motility (42.9 ± 18.6%) and vitality (48.9 ± 23.8%) of IVF patients were significantly higher than in ICSI patients (45.9 ± 41.4 mill/ml; *P* = 0.013), motility 19.6 ± 11.5%; *P* = 0.0001) and (30.3 ± 25.1%; *P* = 0.012) (Table 1). In addition, the mean percentage of normal condensed spermatozoa (CMA₃) was significantly higher (*P* = 0.0001) in patients undergoing IVF (79.88 ± 11.62%) in comparison to ICSI patients (60.6 ± 21.6%). However, spermatozoa of male partners of patients undergoing ICSI therapy were more susceptible to acid denaturation (acridine orange staining) in comparison to spermatozoa of male partners undergoing IVF treatment (98.9 ± 2.7 vs. 94.4 ± 7.5; *P* = 0.041) (Table 1).

ROS was higher in IVF compared to ICSI patients (94.73 ± 102.84 vs. 54.78 ± 39.83 μmol/l, *P* = 0.093). Whereas, TAS concentration was similar in both groups (1.43.0 ± 0.28 vs. 1.53 ± 0.22 mmol, *P* = 0.153). However, no significant difference between both groups could be found either for ROS or TAS concentration (Table 1). An inverse correlation was found between ROS concentration in seminal plasma and sperm vitality (*r* = −0.111; *P* = 0.453), membrane integrity (*r* = −0.042; *P* = 0.778) and morphology (*r* = −0.141; *P* = 0.340) as well as between TAS concentration and global (*r* = −0.313; *P* = 0.030), progressive motility (*r* = −0.409; *P* = 0.004) and morphology after semen processing (*r* = −0.321; *P* = 0.026) (Table 2).

However, in both groups a positive correlation was found between chromatin condensation (CMA₃) of spermatozoa and sperm concentration (*r* = 0.437; *P* = 0.002), global and progressive motility (*r* = 0.299; *P* = 0.039), vitality (*r* = 0.336; *P* = 0.019), and morphology after semen processing (*r* = 0.319; *P* = 0.027). Nevertheless, an inverse correlation was shown between spermatozoa DNA strand breaks (TUNEL) and global motility (*r* = −0.078;

Table 1 Ejaculate parameters of patients undergoing either in-vitro-fertilization (IVF) or intracytoplasmic sperm injection (ICSI)

| Parameters | IVF + ICSI (<i>n</i> = 48) M ± SD median | IVF (<i>n</i> = 26) M ± SD median | ICSI (<i>n</i> = 22) M ± SD median | <i>P</i> -value between IVF and ICSI groups |
|---|--|---------------------------------------|--|--|
| Volumen (ml) | 3.51 ± 1.51 3.5 | 3.63 ± 1.43 3.5 | 3.36 ± 1.63 3.0 | 0.542 |
| pH | 8.57 ± 0.34 8.5 | 8.55 ± 0.34 8.5 | 8.58 ± 0.33 8.6 | 0.747 |
| Concentration (mill/ml) | 60.84 ± 40.25 57.5 | 73.85 ± 34.92 75.0 | 45.47 ± 41.44 30.0 | 0.013 |
| Global motility (%) | 32.19 ± 19.51 30 | 42.88 ± 18.56 40.0 | 19.55 ± 11.54 15.0 | 0.0001 |
| Progress motility (%) | 3.33 ± 6.95 | 5.38 ± 8.71 | 9.1 ± 2.51 | 0.018 |
| Vitality (eosin-test) (%) | 40.33 ± 25.91 37.5 | 48.85 ± 23.85 50.0 | 30.27 ± 25.09 20.0 | 0.012 |
| Membrane integrity (HOS-test) (%) | 61.25 ± 17.12 60.0 | 61.73 ± 18.65 65.0 | 60.68 ± 15.53 60.0 | 0.835 |
| Sperm density after semen preparation (%) | 24.59 ± 24.95 16.5 | 29.68 ± 25.06 19.0 | 18.59 ± 24.00 12.5 | 0.126 |
| Progressive motility after semen processing (%) | 60.31 ± 30.08 75.0 | 78.08 ± 13.57 80.0 | 27.5 | 0.0001 |
| DNA chromatin condensation (CMA ₃) (stained) (%) | 71.02 ± 19.39 75.00 | 79.88 ± 11.62 84.00 | 60.55 ± 21.64 66.5 | 0.0001 |
| DNA single stand breaks (acridine orange) (green) (%) | 96.38 ± 5.70 98.50 | 98.04 ± 2.74 99.50 | 94.41 ± 7.51 98.0 | 0.041 |
| DNA strand breaks (TUNEL-test) (%) | 5.02 ± 4.38 4.00 | 4.38 ± 3.82 3.50 | 5.77 ± 4.95 5.0 | 0.279 |
| Morphologically normal spermatozoa (%) | 8.65 ± 6.03 7.5 | 9.58 ± 5.67 10.5 | 7.55 ± 6.38 6.5 | 0.249 |
| ROS (μmol/l) | 76.42 ± 82.09 51.94 | 94.73 ± 102.84 62.15 | 54.78 ± 39.83 44.85 | 0.093 |
| TAS (mmol/l) | 1.48 ± 0.26 1.45 | 1.43 ± 0.28 1.45 | 1.53 ± 0.22 1.45 | 0.153 |

$P = 0.600$) progressive motility ($r = -0.167$; $P = 0.258$), vitality ($r = -0.108$; $P = 0.467$) membrane integrity ($r = -0.237$; $P = 0.105$) and motility after semen processing ($r = -0.111$; $P = 0.454$) (Table 3).

In the IVF group, membrane integrity (HOS-test) correlated negatively with acridine orange test (DNA-strand break). CMA₃ Chromatin integrity correlated positively with all investigated sperm parameters. Whereas, DNA strand breaks (TUNEL) correlated negatively with motility before and after sperm processing, vitality (Eosin test) and membrane integrity. In the ICSI group, DNA strand breaks (TUNEL) showed a negative correlation with volume, motility, membrane integrity, density after semen processing and morphology (Table 3).

Embryo transfer was performed in 48 patients who were included in this study, 15 patients achieved a pregnancy and 33 patients were unsuccessful (31.25% pregnancy rate) (Fig. 1).

However, there was no significant difference between the mean number of retrieved, fertilized oocytes, fertilization rate between IVF and ICSI groups (10.7 ± 7.2 , 5.5 ± 6.6 and 67.3 vs. 9.1 ± 4.1 , 5.4 ± 3.6 and 67.9% , respectively). In the IVF group, eleven pregnancies were achieved (42.3%) with a 47.8% implantation rate per embryo transferred. In the ICSI group four pregnancies were achieved (18.2%) with a 40.7% implantation rate per embryo transferred. TAS concentration, morphologically normal spermatozoa and DNA integrity (non-fragmented DNA) as assessed by (Tunel) correlated positively with fertilization rate in both groups (Table 4; Figs. 2, 3, 4).

Discussion

Many studies have reported a connection between oxidative stress and DNA damage [15, 18, 35, 77]. The generation of

Table 2 Correlation between ROS, TAS, and others semen parameters, of patients undergoing (IVF) or intracytoplasmic sperm injection (ICSI) treatment

| | Both groups (<i>n</i> = 48) | | IVF (<i>n</i> = 26) | | ICSI (<i>n</i> = 22) | |
|---|------------------------------|--------|----------------------|--------|-----------------------|--------|
| | ROS | TAS | ROS | TAS | ROS | TAS |
| Volume | −0.002 | 0.070 | −0.164 | 0.007 | 0.343 | 0.204 |
| | 0.991 | 0.637 | 0.423 | 0.974 | 0.118 | 0.363 |
| pH | 0.232 | −0.282 | 0.516 | −0.449 | −0.510 | −0.073 |
| | 0.112 | 0.052 | 0.007 | 0.021 | 0.015 | 0.746 |
| Concentration | 0.188 | 0.108 | 0.361 | 0.165 | −0.482 | 0.251 |
| | 0.201 | 0.463 | 0.07 | 0.42 | 0.023 | 0.260 |
| Global motility | 0.083 | −0.313 | −0.108 | −0.499 | 0.028 | 0.370 |
| | 0.575 | 0.030 | 0.601 | 0.009 | 0.901 | 0.090 |
| Progress motility | 0.173 | −0.409 | 0.12 | −0.492 | −0.096 | 0.091 |
| | 0.241 | 0.004 | 0.558 | 0.011 | 0.670 | 0.686 |
| Vitality (eosin) | −0.111 | −0.074 | −0.288 | 0.045 | −0.107 | 0.334 |
| | 0.453 | 0.616 | 0.153 | 0.826 | 0.635 | 0.129 |
| Membrane integrity HOS-test | −0.042 | 0.209 | −0.039 | 0.102 | −0.108 | 0.430 |
| | 0.778 | 0.154 | 0.852 | 0.62 | 0.632 | 0.046 |
| Density after treatment | −0.078 | −0.069 | −0.132 | −0.373 | −0.217 | 0.514 |
| | 0.597 | 0.64 | 0.522 | 0.06 | 0.333 | 0.014 |
| Motility after treatment | 0.169 | −0.15 | 0.125 | −0.157 | −0.125 | 0.067 |
| | 0.251 | 0.310 | 0.542 | 0.444 | 0.579 | 0.767 |
| Morphology (normal) | −0.141 | −0.321 | 0.297 | −0.354 | 0.017 | −0.230 |
| | 0.340 | 0.026 | 0.140 | 0.076 | 0.939 | 0.303 |
| DNA single strand breaks (AO pos.) | 0.150 | −0.066 | 0.310 | −0.091 | −0.098 | 0.050 |
| | 0.308 | 0.654 | 0.123 | 0.660 | 0.664 | 0.824 |
| DNA strand breaks Tunel (green) | −0.010 | −0.117 | −0.005 | −0.216 | 0.132 | −0.098 |
| | 0.948 | 0.426 | 0.981 | 0.290 | 0.557 | 0.665 |
| Chromatin condensation CMA ₃ | 0.228 | 0.010 | −0.082 | 0.114 | −0.299 | −0.356 |
| | 0.119 | 0.944 | 0.691 | 0.580 | 0.177 | 0.104 |
| TAS | −0.039 | 0.0 | −0.003 | 0.0 | 0.076 | 0.0 |
| | 0.791 | | 0.987 | | 0.735 | |
| ROS | 0.0 | −0.039 | 0.0 | −0.003 | 0.0 | 0.076 |
| | | 0.791 | | 0.987 | | 0.735 |

high level of oxygen radicals can mediate the occurrence of high frequencies of single and double-strand DNA breaks commonly observed in the spermatozoa of infertile men [3, 7, 42]. A significant positive correlation between ROS and DNA fragmentation ($r = 0.40$; $P = 0.02$) was reported [10]. Saleh et al. [62] demonstrated a positive correlation between DFI levels and the concentration of ROS in the seminal plasma.

However, seminal plasma provides sperm cells with crucial protection against oxidative attack by different forms of ROS [9]. Lewis et al. [48] reported reduced ascorbate levels in seminal plasma of asthenozoospermic individuals and increased ROS activity. Eskenazi et al. [21] reported that high antioxidant intake was associated with better semen quality, especially motility. In addition, several reports

have suggested that increased generation of ROS and/or variation in the levels of antioxidants defence are involved in the occurrence of oligospermia, sperm motility defects and/or abnormal sperm morphology [14, 40, 72].

In the present study, sperm concentration (73.85 ± 34.92 vs. 45.47 ± 41.44 mill/ml, $P < 0.013$), motility (42.88 ± 18.56 vs. $19.55 \pm 11.54\%$; $P < 0.0012$), vitality (48.85 ± 23.85 vs. $30.27 \pm 25.09\%$, $P = 0.012$), DNA integrity (98.04 ± 2.74 vs. $94.41 \pm 7.51\%$, $P < 0.041$), chromatin condensation (CMA₃) (79.88 ± 11.62 vs. $60.55 \pm 21.64\%$, $P < 0.0001$) were significantly better in IVF patients compared to ICSI patients (Table 1). However, ROS concentration was higher in IVF than in ICSI groups (94.73 ± 102.84 vs. 54.78 ± 39.83 $\mu\text{mol/l}$, $P = 0.093$). Though these difference was not statistically significant. Whereas, TAS

Table 3 Correlation between standard ejaculate parameters and chromatin concentration (Chromomycin CMA₃), single strand breaks (acridine orange) and strand breaks (TUNEL) of patients undergoing IVF/ICSI

| | Both groups | | | IVF (<i>n</i> = 26) | | | ICSI (<i>n</i> = 22) | | |
|------------------------------------|-------------------------|--------------------|--------|-------------------------|--------------------|--------|--------------------------|--------------------|--------|
| | Acridine orange (green) | CMA3 Non-lightning | Tunel | Acridine orange (green) | CMA3 Non-lightning | Tunel | Acridine orange positive | CMA3 Non-lightning | Tunel |
| Volume | 0.085 | −0.028 | 0.197 | 0.359 | −0.290 | 0.629 | −0.050 | 0.026 | −0.119 |
| | 0.568 | 0.849 | 0.179 | 0.071 | 0.151 | 0.001 | 0.826 | 0.909 | 0.597 |
| pH | 0.073 | −0.064 | 0.193 | −0.023 | 0.00 | −0.076 | 0.162 | −0.081 | 0.444 |
| | 0.623 | 0.664 | 0.189 | 0.910 | 0.998 | 0.713 | 0.472 | 0.719 | 0.038 |
| Concentration | 0.260 | 0.437 | 0.020 | 0.205 | 0.392 | 0.094 | 0.167 | 0.294 | 0.075 |
| | 0.074 | 0.002 | 0.891 | 0.314 | 0.048 | 0.646 | 0.458 | 0.185 | 0.740 |
| Global motility | 0.192 | 0.457 | −0.078 | −0.096 | 0.375 | 0.094 | 0.065 | 0.139 | −0.077 |
| | 0.191 | 0.001 | 0.600 | 0.640 | 0.059 | 0.649 | 0.772 | 0.536 | 0.733 |
| Progress motility | 0.110 | 0.299 | −0.167 | 0.024 | 0.159 | −0.143 | −0.008 | 0.403 | −0.175 |
| | 0.456 | 0.039 | 0.258 | 0.906 | 0.439 | 0.487 | 0.972 | 0.063 | 0.437 |
| Vitality (eosin) | 0.172 | 0.336 | −0.108 | −0.204 | 0.175 | −0.091 | 0.184 | 0.214 | 0.022 |
| | 0.241 | 0.019 | 0.467 | 0.318 | 0.393 | 0.657 | 0.412 | 0.338 | 0.921 |
| Membrane integrity (HOS) | −0.235 | 0.272 | −0.237 | −0.45 | 0.321 | −0.164 | −0.223 | 0.321 | −0.326 |
| | 0.108 | 0.061 | 0.105 | 0.021 | 0.11 | 0.424 | 0.318 | 0.146 | 0.138 |
| Density after processing | 0.170 | 0.144 | 0.120 | 0.113 | −0.063 | 0.474 | 0.122 | 0.108 | −0.135 |
| | 0.248 | 0.329 | 0.418 | 0.583 | 0.758 | 0.015 | 0.587 | 0.633 | 0.548 |
| Motility after processing | 0.257 | 0.569 | −0.111 | 0.168 | 0.337 | −0.101 | 0.049 | 0.380 | 0.027 |
| | 0.078 | 0.000 | 0.454 | 0.411 | 0.092 | 0.624 | 0.830 | 0.081 | 0.905 |
| Morphologically normal spermatozoa | 0.204 | 0.319 | 0.127 | 0.260 | 0.0140 | 0.382 | 0.140 | 0.435 | −0.025 |
| | 0.164 | 0.027 | 0.391 | 0.199 | 0.944 | 0.054 | 0.534 | 0.043 | 0.914 |

Table 4 Correlation between reactive oxygen species (ROS) total anti oxidants (TAS), chromatin condensation Chromomycin CMA₃, single strand breaks (Acridine orange), strand breaks (TUNEL), morphology and IVF/ICSI outcome

| Parameters | Both groups | | IVF group | | ICSI group | |
|--|--------------------|-----------|--------------------|-----------|--------------------|-----------|
| | Fertilization rate | Pregnancy | Fertilisation rate | Pregnancy | Fertilization rate | Pregnancy |
| ROS (μmol/l) | −0.290 | 0.051 | −0.267 | 0.006 | −0.241 | −0.092 |
| | 0.045 | 0.730 | 0.187 | 0.976 | 0.280 | 0.683 |
| TAS (mmol/l) | 0.081 | −0.064 | −0.031 | 0.005 | 0.184 | −0.040 |
| | 0.584 | 0.664 | 0.881 | 0.980 | 0.412 | 0.859 |
| DNA single strand breaks (AO) | −0.115 | 0.067 | 0.064 | 0.074 | −0.144 | 0.075 |
| | 0.435 | 0.652 | 0.755 | 0.718 | 0.522 | 0.742 |
| Chromatin condensation (CMA ₃) | −0.060 | 0.070 | 0.126 | 0.200 | −0.011 | −0.308 |
| | 0.683 | 0.639 | 0.540 | 0.327 | 0.961 | 0.164 |
| DNA stand breaks (Tunel) (%) | 0.108 | −0.045 | 0.231 | −0.046 | −0.106 | 0.047 |
| | 0.466 | 0.763 | 0.256 | 0.822 | 0.639 | 0.837 |
| Morphologically normal spermatozoa | 0.138 | 0.040 | 0.203 | 0.191 | 0.152 | −0.268 |
| | 0.349 | 0.787 | 0.319 | 0.350 | 0.500 | 0.228 |

concentration was similar in both groups (1.43 ± 0.28 vs. 153 ± 0.22 mmol/l, $P = 0.153$) (Table 1).

In addition, ROS in seminal plasma of IVF patients was negatively correlated with semen volume, global motility

vitality, membrane integrity, density after semen processing, chromatin condensation, DNA double stand breaks and TAS concentration. However, in ICSI patients, a negative correlation was observed between ROS level and sperm

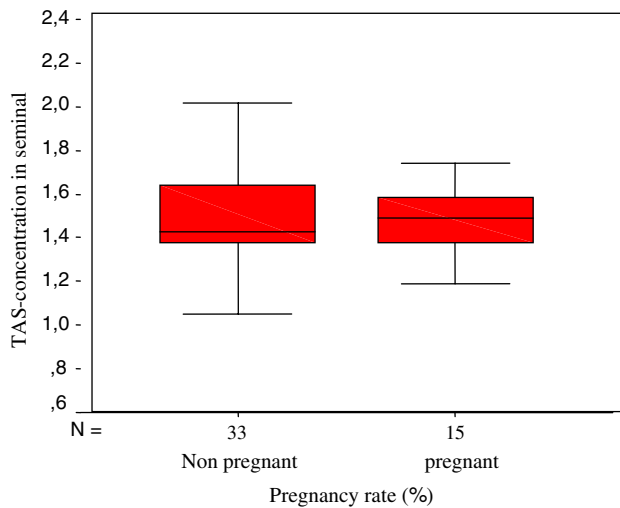


Fig. 1 Comparison between patients who became pregnant and those who did not regarding TAS concentration of seminal plasma (25, 75% percentile). No significant difference was shown between the groups

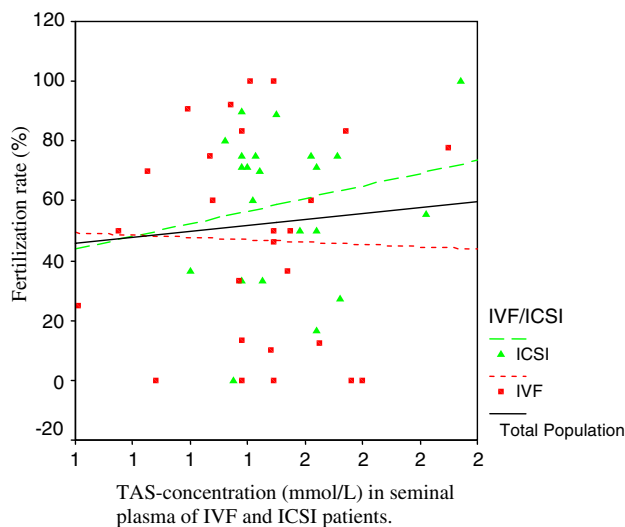


Fig. 2 Correlation between TAS concentration in ejaculate and fertilization rate of IVF/ICSI patients

concentration, progressive motility, vitality, membrane integrity, sperm density and motility after semen processing, chromatin condensation and DNA single stand breaks (Table 2). Kessopoulou et al. [39], and Henkel and Schill [32] previously reported a correlation between sperm concentration and increased production of ROS in infertile patients. Zorn et al. [82] found that ROS concentration in seminal fluid was negatively correlated with progressive motility, normal morphology and positively correlated with abnormal sperm head morphology. Iwasaki and Gagnon [36] and Mazzilli et al. [50] also reported a positive correlation between superoxide anion levels and sperm morphology abnormalities. According to the investigation of Aziz et al. [8] ROS positively correlates with the sperm deformity

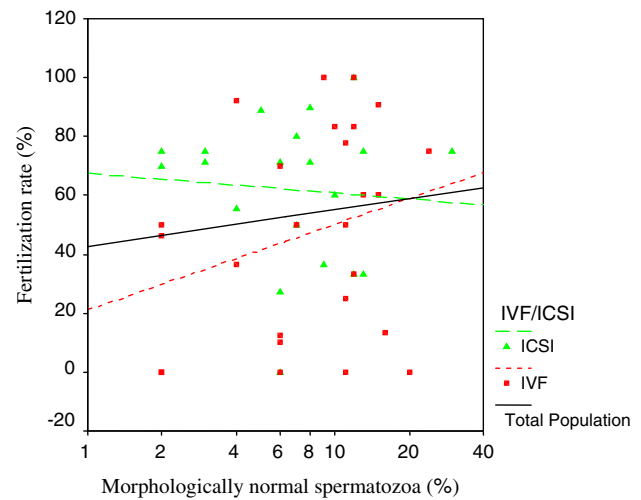


Fig. 3 Correlation between normal morphology of spermatozoa and fertilization rate of IVF/ICSI patients. Mean number of morphologically normal spermatozoa correlated positively with the fertilization rate

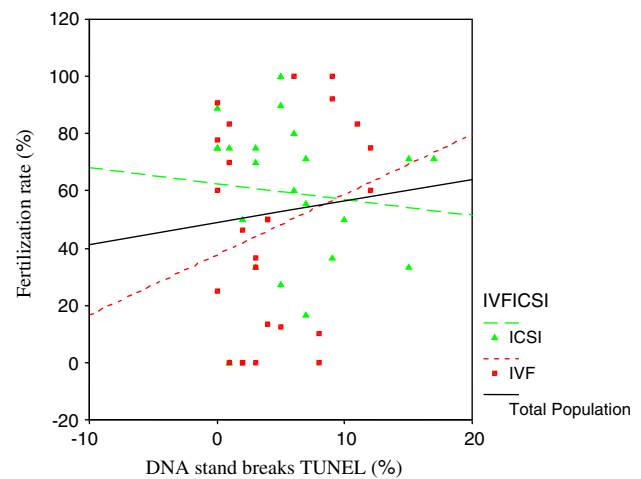


Fig. 4 Correlation between DNA fragmentation of spermatozoa and fertilization rate of IVF/ICSI patients

index, calculated by dividing the total number of deformities observed by the number of sperm evaluated [8].

Besides, many studies indicated that DNA damage is associated with abnormalities in conventional semen parameters [11, 13, 51, 61, 76]. Irvine et al. [35] have demonstrated significant negative correlations between semen quality and sperm DNA integrity. They have found an inverse correlation between DNA and sperm concentration (-0.54), motility (-0.38) and morphology (-0.38). Saleh et al. [62] have also shown that DNA damage exhibits a significant inverse correlation with motility (-0.47).

In the present study, DNA integrity was analysed by chromomycin CMA₃, acridine orange stain and tunel assay, in order to gain complete insight into fertilization and pregnancy failure due to DNA damage. In both groups, a

significantly positive correlation was found between the condensed nucleus chromatin and the sperm motility ($r = 0.457$; $P = 0.001$), morphology ($r = 0.319$; $P = 0.027$) and vitality ($r = 0.336$; $P = 0.019$). DNA strand breaks (Tunel) correlates negatively with the motility ($r = -0.078$; $P = 0.006$) and vitality ($r = -0.108$; $P = 0.467$), but not with morphology ($r = 0.127$; $P = 0.391$) and density ($r = 0.020$; $P = 0.891$) (Table 3). These findings in the present study, confirm several previous studies [27, 67, 80] who found an association between DNA damage and sperm motility. The correlation between DNA strand breaks and sperm concentration and total sperm count is consistent with the previous findings [19, 24, 45, 68].

In addition, negative correlations were found between single strand breaks and global motility, vitality and membrane integrity, progressive motility, vitality, membrane integrity and motility after semen processing in IVF group. In spermatozoa of ICSI patients, similar correlations were observed as well as a negative correlation between DNA strand breaks and morphology (Table 3).

In contrast, Chen et al. [13] studied 59 men undergoing IVF treatment, and Donnelly et al. [19] studied 17 fertile men and 40 infertile men undergoing IVF or ICSI. Both studies reported no statistically significant correlation between sperm DNA damage and sperm morphology scored according to strict criteria. Morris et al. [54] reported a positive association between DNA damage and the percentage of sperm with abnormal morphology and a negative association between DNA damage and sperm concentration.

Said et al. [60] suggested that the sperm deformity index (SDI) may be a useful tool to detect the prevalence of sperm DNA damage and to identify potential infertile men. Infertile patients with a semen sample containing a high proportion of morphological abnormalities, specifically cytoplasmic droplets, may be more susceptible to develop ROS-mediated sperm DNA damage [60]. Therefore, excessive ROS production is one of the best predictor of low fertilization in conventional IVF [43, 70, 71].

On the otherhand, spermatozoa with DNA fragmentation can fertilise oocytes, but the pregnancy process may be disturbed. This explains the positive association of the DNA strand breaks (Tunel) with fertilization rate ($r = 0.108$; $P = 0.466$) and the negative correlation with the number of pregnancies ($r = -0.045$; $P = 0.763$) (Table 4). In accordance with these results, Henkel and Schill [33] did not find an association of the percentage of Tunel positive spermatozoa in the ejaculate with fertilization ($r = -0.087$; $P = 0.269$), but with pregnancy ($r = 0.228$; $P = 0.006$). A further inverse correlation was shown between single strand breaks (acridine orange staining) and fertilization rate after ICSI ($r = -0.144$; $P = 0.522$) (Table 4). The mean percentage of morphologically normal spermatozoa correlated positively

with fertilization in IVF ($r = 0.203$; $P = 0.319$) and ICSI ($r = 0.152$; $P = 0.500$) (Table 4; Fig. 3). In accordance with these results, it has been shown by several working groups that DNA damage in ejaculated spermatozoa as determined by TUNEL assay or other techniques such as the comet assay is negatively correlated with fertilization and pregnancy in IVF [34, 54, 73] and ICSI [49] program. It is well documented that there is a negative correlation between defective sperm chromatin structure (DNA-break) and fertility, in vivo [69] and in IVF cycles [22, 23, 29, 45, 46, 78]. Therefore, correct chromatin packaging around the protamine core seems to be necessary for optimal expression of the male gamete fertility potential. However, this condition does not seem mandatory for successful fertilization as demonstrated by ICSI, where normal fertilization and pregnancy rates can be achieved with cells that have not completed spermiogenesis, such as epididymal and testicular spermatozoa [29, 66]. In the present study, ROS are negatively associated with fertilization rate in conventional IVF and ICSI groups ($r = -0.267$; $P = 0.187$ and $r = -0.241$; $P = 0.280$, respectively). Whereas, a positive correlation was shown between TAS concentration and fertilization rate in both groups ($r = 0.081$; $P = 0.584$) (Table 4; Fig. 2). These results could indicate the influence of ROS on the plasma membrane and as a consequence the membrane fusion in IVF or sperm membrane swelling and chromatin decondensation after intracytoplasmic sperm injection. Besides, high levels of semen ROS are associated with low pregnancy rates in vivo [4]. Besides, the results of the present investigation also revealed that the mean number of retrieved oocytes (10.69 ± 7.24 vs. 9.09 ± 4.14 $P = 0.346$), number of fertilized oocytes (5.46 ± 6.59 vs. 5.36 ± 3.62 $P = 0.951$), fertilization rate (67.26 vs. 67.87%) and number of transferred embryos (1.73 ± 1.00 vs. 2.18 ± 0.79 , $P = 0.097$) were similar in both IVF and ICSI groups. However, 15 patients achieved a pregnancy whilst 33 patients were unsuccessful (31.25% pregnancy rate) (Fig. 1). In IVF group ($n = 26$) 11 pregnancies were achieved (42.3%) with a 47.8% implantation rate per embryo transferred. In ICSI patients ($n = 22$) four pregnancies were achieved (18.2%) with a 40.7% implantation rate per embryo transferred (Fig. 5) as previously thoroughly described [31].

In conclusion ROS and TAS concentration in seminal plasma did not differ significantly between the patients undergoing IVF or ICSI therapy. A negative correlation was shown between ROS concentration in seminal plasma and sperm vitality membrane integrity sperm density Chromatin condensation, DNA single stand breaks in both IVF and ICSI groups. TAS level in seminal plasma was positively correlated to the fertilization rate of patients undergoing IVF/ICSI therapy. However, the potential role of non-enzymatic antioxidants in treatment of male infertility is still questionable.

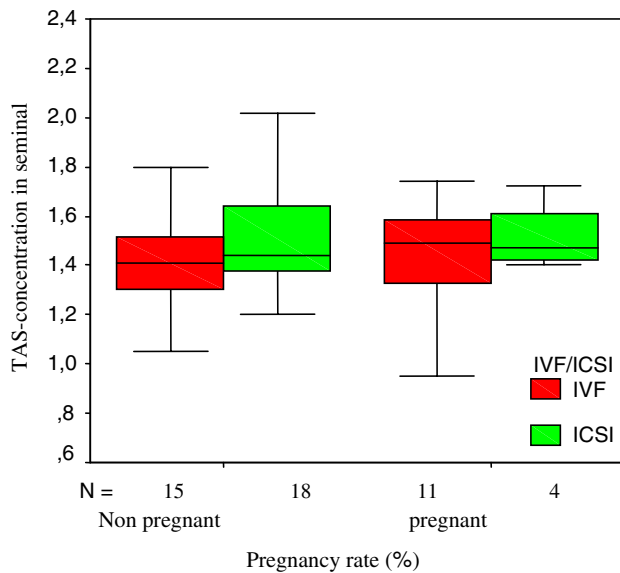


Fig. 5 Relationship between TAS concentration in seminal fluid and pregnancy rate of patients undergoing IVF/ICSI treatment. Higher TAS concentration in seminal plasma indicated increase in pregnancy rate (value represents median and interquartile (25, 75% percentile). No significant difference was shown between the groups

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