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Effects of Reactive Oxygen Species on Sperm Function, Lipid Peroxidation and DNA Fragmentation in Bovine Spermatozoa

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Objective: To evaluate the effects of the reactive oxygen species (ROS) generated with a xanthine (X) and xanthine oxidase (XO) system on sperm function, the change of sperm characteristics, lipid peroxidation, and DNA fragmentation in bovine spermatozoa.

Materials and Methods: ROS were produced using a combination of 1000 uM X and 50 mU/ml XO. The ROS scavengers: superoxide dismu tase (SOD) (200 U/ml) and catalase (500 U/ml) were also tested. Spermatozoa were incubated for 2 hours in BWW medium with a combination of XXO supplemented with or without ROS scavengers at 37 under 5% CO_2 incubator. Sperm movement characteristics by CASA (computer-aided sperm analysis), HOST (hypoosmotic swelling test), Ca-ionophore induced acrosome reaction, malondialdehyde formation for the analysis of lipid peroxidation, the percentage of DNA fragmentation using the method of TdT-mediated nick end labelling (TUNEL) by flow cytometry were determined after 2 hours incubation.

Results: The action of ROS on bovine spermatozoa resulted in a decreased in capacity for sperm motility, Ca-ionophore induced acrosome reaction and membrane integrity, an increased in malondialdehyde formation and the percentage of sperm with DNA fragmentation. In the effects of antioxidant, catalase completely alleviated the toxic effects induced by the ROS in terms of sperm function and characteristics, however SOD exhibited no capacity to reduce the toxic effects.

Conclusion: The ROS can induce significant damages to sperm functions and characteristics. The useful ROS scavengers can minimized the defects of sperm function and various damages of spermatozoa. **Key Words:** ROS, Sperm function, Lipid peroxidation, DNA fragmentation, Catalase, SOD

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(reactive oxygen species; ROS)7 7 7 (aerobic) 7 7 , ROS7 ROS7 . ROS

superoxide anions (\cdot O₂⁻), peroxyl radical (ROO \cdot) hydroxyl radicals (\cdot OH), hydrogen peroxide (H₂O₂), ROS7 \dagger .

duction . ROS (hyperactivation) phospholipase A2 . ROS super-

,⁷ (cytoskeleton) ⁸ 9 . ROS7

> xanthine (X)-xanthine oxidase (XO) ROS7 , LPO DNA fragmentation

1.

Holstein 0.5 ml straw . 2.

1)

20 mM N-(2-hydroxyethy) piperazine-N'-2-ethane sulphonic acid (HEPES; Sigma, USA), 20 mM NaHCO₃ (Sigma, USA) 0.1% polyvinyl alcohol (PVA; Sigma, USA) 7 Biggers Witten Whittingham (BWW) ¹⁰

2) 37 20 , 5 ml

(500 ×g, 5 min) . 2 ml 2 (300 ×g, 5 min) .

80%

3) (Com puter - Aided Sperm Analysis; CASA) CASA 10 ?1 37 7 makler counting chamber (Sefi medical instrument, Israel) TI-23A CCD (NEC, Japan)7 (Olympus, Japan) CTS-60/200 system (Motion Analysis, USA)

(motility), (straight line velocity; VSL), (curvilinear velocity; VCL), (linearity; LIN), (amplitude of

Table 1. Parameter settings of CTS-60 system

1. Temperature	37
2. Frame rate	60 frames/sec
3. Duration of data capture	30 frames
4. Minimum path length	25 frames
5. Minimum motile speed	10 ? m/sec
6. Maximum burst speed	600 ? m/sec
7. ALH path smoothing factor	7 frames
8. Depth of sample	10 ? m

lateral head displacement; ALH) VCL (hyperactivated sperm) 11 100 ? m/s, LIN<60%, ALH 5 ? m CTS-60/200 system Table 1 4) (Hypo - Osmotic Swe lling Test; HOST) 12 HOST Jeyendran 300 ?1 (20 ×10⁶ spermatozoa/ml) fructose (Sigma, USA) 13.51 gm sodium citrate (Shin-

yo, Japan) 7.35 gm 3 1000 ml 150 mOsm/Kg 5 ml 37 30 . 500 ×g 5 , 10 ?1





Figure 1. Diagrammatic representation of hypo-osmotic swelling patterns. Shaded area denote regions of tail swollen. a, no swelling; $b \sim d$, tail-tip patterns; $e \sim g$, various types of swelling.¹²

5)

(1) Stock solution

Calcium ionophore A23187 (A23187; Sigma, USA) 1.0 mg 382 ?1 dimethyl sulfoxide (DMSO; Sigma, 5 mM/L stock solution USA) eppendorf tube 20 ?1 -20 20 ?1 stock solution 4.98 ml 가 20 ?M/L A23187 4.98 ml DMSO 20 가 ?1 Hoechst 33258 (H33258; Sigma, USA) 1 mg/ml eppendorf tube 20 ?1 -20 stock solution 1 ? g/ml

Fluorescein isothiocyanate conjugated pisum sativum agglutinin (FITC-PSA; Sigma, USA) 1 mg/ ml eppendorf tube 100 ?1 -20 FITC-PSA (1 : 9) 100 ?g/ml

(2)

Cummins ¹³

(20 ×10⁶ spermatozoa/ml) 20 ?M/L A23187 (10 ?M/L) DMSO 30

37 , 5% CO₂ (3) H33258 7 가 1.5 ml phosphate buffered saline (PBS; GibcoBRL, USA) 4 ml 2% polyvinyl pyrrolidone (Sigma, USA) column 500 ×g 5 20 ?1 95% ethanol 5 FITC-PSA (100 ?g/ml) 100 ?1 4 moist chamber 15 propyl gallate mercury mountant epi-illumination module burner (Olympus, Japan) H33258 1000 filter cube U , FITC-PSA

6) (lipid peroxidation) Lipid peroxidation malondialdehyde thiobabituric acid (TBA)

Ca⁺⁺, Mg⁺⁺ Hank's ba-가 20× lanced salt solution 10⁶/ml malondial-. Lipid peroxide 3 dehyde 1 mM ferrous sulphate (Sigma, USA) 5 mM sodium ascorbate (Sigma, USA) 10 ?1 가 1 ml 37 30 . 250 ?1 40% trichloroacetic acid (Sig-가 0 10 lipid ma, USA) 2500 ×g peroxidation 10 250 ?1 1 ml 1% TBA 10 . Malondialdehyde spectrophotometer (Kontron instruments, Switzerland) (optical 532 nm malondialdedensity) $(1.49 \times 10^{5} 1 \cdot mol^{-1} \cdot$ hyde .15 cm⁻¹) 7) (flow cytometry) **DNA** fragmentation DNA fragmentation TUNEL (terminal deoxynucleotidyl transferase (TdT) dUTP nick end APO-DIRECT TM kit (Pharlabelling) (Becton dickinson, mingen, USA) USA) 가 2~4 ×10⁶/ml 0.5 ml PBS 5 ml 1% paraformaldehyde (pH 7.2) 15 5 5 300 ×g PBS ml 2 70% ice-cold ethanol 5 ml -20 30 . 300 **x**g 5 ethanol 1.0 ml wash buffer 300 5 2 ×g FITC-dUTP ΤđΤ enzyme 50 ?1 37 60 5 ml rinse buffer 300 ×g 5

2 . 0.5 ml propidium iodide(PI)/RNas e A 7 30 . 3



Figure 2. Histograms of the TUNEL assay in bovine spermatozoa analyzed by flow cytometry. (a) A typical negative control, in which 0.1% of the spermatozoa are labeled. (b) A positive control treated with DNase I, in which 62.8% of the spermatozoa are labeled.

10¹

FL1-H

0

10⁰

ΤďΤ DNA fragmentation , PBS negative control 0.1 IU DNase I (Sigma, USA) DNA fragmentation positive Figure 2 DNA frag-Figure 2 DNA fragmentation 8) ROS ROS7 xanthine (X) xanthine oxidase (XO) ROS X-XO 250, 500, 1000 uM Х 25, 50 mU/ml XO

(1000 uM X~50 mU/ml XO)



10²

Figure 3. Kinetics of reactive oxygen species generation in the xanthine and xanthine oxidase system used in this study. (A) Luminol-dependent chemiluminescence for time resolution of H_2O_2 generation. (B) Lucigenin-dependent chemiluminescence for time resolution of $\cdot O_2^-$ generation.

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Movement characteristic	Time (min)			
Wovement enaracteristic -	0 30		60	120
Motility (%)	82.4 ±0.8	$49.5 \pm 0.5^*$	$50.8 \pm 1.5^*$	$16.3 \pm 5.0^*$
VSL (? m/s)	66.3 ±7.2	$41.8 \pm 2.5^*$	$19.5 \pm 0.9^*$	$4.8 \pm 4.8^*$
VCL (? m/s)	129.3 ±7.2	$88.8 \pm 3.3^*$	$86.5 \pm 3.1^*$	$23.8 \pm 23.8^*$
LIN (%)	51.5 ±3.1	48.3 ± 2.6	$25.5 \pm 2.3^*$	$5.0 \pm 5.0^*$
ALH (?m/s)	6.0 ±0.2	$4.4 \pm 0.4^*$	$4.8 \pm 0.3^*$	$2.1 \pm 2.1^*$
HA (%)	21.6 ±3.4	$5.1 \pm 1.9^*$	$5.4 \pm 0.8^*$	$0.4 \pm 0.4^{*}$

 Table 2. Influence of reactive oxygen species generation with the xanthine and xanthine oxidase system on the movement characteristics of bovine spermatozoa

Values are mean ±SEM, n=4. *p<0.05 compared with zero time. VSL: straight line velocity; VCL: curvilinear velocity; LIN: linearity; ALH: lateral head displacement; HA: hyperactivation.

Table 3. Effects of antioxidants on the movement characteristics of bovine spermatozoa after 2 h of incubation with the xanthine (X) and xanthine oxidase (XO)

Movement			X-XO +		
characteristic	Control	X-XO	SOD	CAT	SOD + CAT
Motility (%)	52.0 ± 2.7	$16.3 \pm 5.0^*$	17.5 ± 4.6 [*]	$66.8 \pm 4.0^*$	63.3 ± 4.5
VSL (? m/s)	72.5 ±3.9	$4.8 \pm 4.8^*$	$6.5 \pm 3.8^*$	63.3 ±8.1	61.5 ±5.8
VCL (?m/s)	123.0 ± 6.7	$23.8 \pm 23.8^*$	$21.8 \pm 13.4^*$	121.3 ± 6.8	115.3 ± 5.3
LIN (%)	56.8 ±1.1	$5.0 \pm 5.0^*$	$16.8 \pm 10.8^*$	51.3 ± 4.7	52.0 ±3.5
ALH (?m/s)	5.4 ±0.3	2.1 ±2.1	2.7 ±1.8	5.8 ±0.3	5.8 ±0.3
HA (%)	5.6 ±1.3	$0.4 \pm 0.4^{*}$	$0.0 \pm 0.0^*$	$14.4 \pm 2.3^*$	11.3 ± 2.5

Values are mean ±SEM, n=4. *p<0.05 compared with control values. SOD: superoxide dismutase; CAT: catalase; VSL: straight line velocity; VCL: curvilinear velocity; LIN: linearity; ALH: lateral head displacement; HA: hyperactivation.

ROS X-XO superoxide dismutase (SOD) catalase 200, 400, 800 U/ml SOD 500, 1000, 2000 U/ml catalase 가). (200 U/ml SOD 500 U/ml catalase X-XO , X-XO SOD catalase

.

HOST, lipid peroxidation, DNA fragmentation

2

9) ROS

ROS chemiluminescence

.¹⁶ Chemiluminescence signal LB950 luminometer (Berthold, German) 3 integration mode . ROS probe luminol (5-amino-2, 3-dihidro-1, 4phthalazinedione; Sigma, USA) lucigenin (10, 10'-dimethyl-9, 9'-biacridinium dinitrate; Sigma, USA) . Luminol pH ,

> ROS (H_2O_2 , $\cdot O_2^-$, $\cdot OH$) , peroxidase H_2O_2



Figure 4. Effects of superoxide dismutase (SOD) and catalase (CAT) on the acrosome reaction in bovine spermatozoa after 2 h of incubation with the xanthine (X) and xanthine oxidase (XO). Acrosome reaction expressed as the difference between spontaneous and ionophore induced acrosome reaction rates. Values are mean \pm SEM, n=5. * p<0.01 compared with the control.

•	, lucigenin	pН
	, ROS	$\cdot O_2$

probe stock solution DMSO 25 mM X-XO H_2O_2 Luminol X (1000 uM)~XO (50 mU/ml)7 가 400 luminol stock solution 4 ?1 (?1 250 uM) PBS 8 ?1 horseradish peroxidase (12.4 U; Sigma, USA) 가 luminometer 120 signal Lucigenin $\cdot O_2$ X (1000 uM)~ 가 400 ?1 XO (50 mU/ml)가 lucigenin stock solution 4 ?1 (250 uM) 가 luminometer signal 120

3.

SPSS (ver. 7.5) , Mann-Whitney U test . p<0.05



Figure 5. Effects of superoxide dismutase (SOD) and catalase (CAT) on the swelling of bovine spermatozoa under hypo-osmotic condition after 2 h of incubation with the xanthine (X) and xanthine oxidase (XO). Values are mean \pm SEM, n=5. *p<0.05 compared with the control.

1000 uM X	~50 mU	XO/m	1		
ROS			H_2O_2		1
(9020×10^3)	cpm), ·	O_2		10	
(1400 ×1	0^3 cpm)			,	
		(Fi	gure 3).	ROS	
Aitker	9			X-X	O system
	uric aci	d			
			ROS		
X-XO					
30			param	eter	
	(p<0.0)5) (Ta	able 2).		
ROS					
		X-XO)		• O ₂ -
S	SOD H	$_2O_2$		catal	ase
	catalase	SOD)		
Table	3				
			catalas	se	catalase
SOD		X-X	O system	n	
ROS					
	,	SOD			ALH
		Σ	K-XO		



Figure 6. Effects of superoxide dismutase (SOD) and catalase (CAT) on the induction of lipid peroxidation in bovine spermatozoa after 2 h of incubation with the xanthine (X) and xanthine oxidase (XO). Lipid peroxidation expressed as nmol malondialdehyde generated by 1×10^8 spermatozoa. Values are mean ±SEM, n=5. p<0.05 compared with the control.



(p<0.01) (Figure 7).



Figure 7. Effects of superoxidsmutase (SOD) and catalase (CAT) on DNA fragmentation assessed by TU-NEL and flow cytometry in bovine spermatozoa after 2 h of incubation with the xanthine (X) and xanthine oxidase (XO). Values are mean ±SEM, n=5. *p<0.01 compared with the control.



. 60% 7 , 50% 7 docosahexaenoic acid , ROS

 .1.20
 lipid peroxidation
 ROS

 .17
 lipid peroxidation

 .71
 lipid peroxidation

21,22

ROS

lipid peroxidation , 9 lipid peroxidation malondialdehyde DNA 23 ROS lipid peroxidation , 7 , ROS SOD, catalase, glutathione peroxidase

가 SOD catalase SOD $\cdot O_2$ O_2 $H_2O O_2$ H_2O_2 , catalase H_2O_2 ROS X XO . X-XO $\cdot O_2$ H_2O_2 ROS가 . $H_2O_2 \quad 2$ Haber-Weiss cycle 24 · OH ROS catalase

.

ROS

ROS SOD lipid peroxidation 가 catalase SOD $\cdot O_2^ H_2O_2$. Aitken 9 X-XO system ROS가 ROS 가 ROS가 ROS lipid peroxidation $\cdot O_2$ SOD lipid 가 peroxidation H_2O_2 catalas e , lipid peroxidation 25 . Griveau

X-XO system ROS가 ionophore , lipid peroxi-가 dation lipid hydroperoxide H_2O_2 가 , X-XO H_2O_2 glutathione peroxidase glucose-6-phosphate dehydrogenase 26 . Blondin hypoxanthine xanthine oxidase ROS

ROS ROS · SOD 가 , catalase 가

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H_2O_2 ROS가 DNA fragmentation 27 X-XO Lopes ROS DNA ROS DNA fragmentation 가 가 DNA ROS glutathione hypotaurine DNA catalase DNA catalas e ROS DNA fragmentation 가 ROS ROS

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