Epididymal Toxicity Associated with Vincristine Treatment

T. SONAWANE*, S. AZAZ¹, K. HEMANT¹ AND T. LIJI¹

Amity Institute of Biotechnology, Amity University, Mumbai-410 206, ¹School of Biotechnology and Bioinformatics, D. Y. Patil Deemed to be University, Mumbai-400 706, India

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Vincristine, a major player in front line combination chemotherapy of cancer reduces testosterone levels contributing to reproductive toxicity. Much is known about testicular toxicity of vincristine as compared to its effect on epididymis; hence, the present study aimed to evaluate the epididymal toxicity associated with vincristine treatment, which also contributes to the overall reproductive toxicity associated with vincristine. Vincristine was intraperitoneally injected to adult male Wistar rats of proven fertility with a dose of 40 mg/kg/day dissolved in 0.5 ml of physiological saline for 30 days. The epididymal weight was found to be unaltered after treatment whereas sperm count was reduced significantly. Significant changes were noted in ion concentrations of cauda and caput of epididymis with changes in protein profile of the tissue, sperm and luminal protein from cauda and caput region, which plays a significant role in sperm maturation and sperm transport. Infertility associated with vincristine could be attributed to its effects on various epididymal proteins involved in sperm protection and various stages of sperm development such as cytoplasmic extrusion and membrane stabilization, which had contributed to the abnormal sperm count and impaired function.

Key words: Vincristine, epididymis, toxicity, ion concentration, sperm count

Vincristine (VCR) also known as leurocristine is a *Vinca* alkaloid derived from the *Catharanthus roseus* (Madagascar periwinkle) formerly *Vinca rosea*. VCR has been widely used as an anticancer drug because of its ability to inhibit cell division through prevention of polymerization of tubulin of the microtubules, which constitute the spindle fibre and the disassociation of the existing microtubules^[1]. Although being a potent anticancer drug its use is limited due to a wide spectrum of side effects such as neurotoxicity, alopecia, colicky abdominal pain, myocardial infarction, optic atrophy and diplopia^[2,3]. The male reproductive system consisted of actively dividing cells has also shown to be affected by VCR, thus compromising fertility^[4].

Epididymis being the site for sperm maturation is very important for the development of mature and motile sperm with fertilizing ability. The array of proteins and ions in the epididymis plays a vital role in maintaining the epididymal function^[5]; even a slight modification would effects the metabolic pathways and ultimately result in compromised epididymal function. Several reports have suggested the impairment of epididymal functions and spermatogenesis on VCR administration^[6] thereby affecting the levels of sperm maturation and transit proteins.

It has also been shown that apart from conventional male reproductive toxicity assessment by measuring the weight of reproductive organs such as testis, epididymis and prostate, analysis of sperm motility and viability, macroscopic and histological examination and changes in hormone levels^[7,8], the protein profile can also be used to evaluate the male reproductive toxicity^[9-12]. The present study was aimed to evaluate the alterations in the sperm count, ion concentration and pattern of protein profile in cauda and caput after treatment with VCR thus leading to compromised epididymal functions.

MATERIALS AND METHODS

Male Wistar rats of proven fertility weighing 150-200 g

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(Bharat Serum and Vaccines Ltd, Mumbai) were maintained under standard laboratory conditions and were handled according to the institutional legislation regulated by the Committee for the Purpose of Control and Supervision of Experiments on Animals. They were allowed free access of food and water *ad libitum* (Amrut Feed Ltd).

Drug and dose:

VCR (Cytocristin, Cipla Ltd, Goa, India) was purchased from a local pharmacy and rats (n=5) were injected intraperitoneally with 40 μ g/kg/d of VCR in 0.5 ml of physiological saline while the control rats (n=5) received 0.5 ml of saline (ip) for 30 d. At the end of the treatment period, animals were sacrificed under diethyl ether anaesthesia. The epididymis were quickly removed, blotted free of blood, weighed, snap frozen in liquid nitrogen and stored at -80° until used for further studies.

Sperm count:

The epididymal fluid was obtained by mincing of tissues in phosphate-buffered saline (PBS) followed by incubation of 30 min and centrifugation at 2500 rpm for 2 min. Sperm count was assayed by the method of Belsey *et al.*^[13]. The total number of sperms was calculated using the following formula, total number of cell counted×20 (dilution factor)/0.4 mm² (total volume of the squares in which the cells were counted)

Estimation of Na⁺, K⁺ and Mg²⁺:

Luminal ions from caput and cauda of epididymis were extracted by the method described by Aranha *et al.*^[14]. The extract was used for estimating Na⁺, K⁺ and Mg²⁺ using an atomic absorption spectrophotometer (Chemito Instruments Pvt. Ltd, Worli, Mumbai, India).

Protein extraction and quantification:

The epididymal tissue samples were sequentially thawed, and dissected to separate individual segments, the cauda and caput. The epididymal segments were minced and incubated in 2 ml of 0.1 M PBS (pH 7.4) containing 0.1 M phenylmethylsulfonyl fluoride (PMSF) for 30 min at 4° to facilitate the release of sperms. The suspension was then pelleted by centrifugation at 2500 rpm for 2 min. The washing procedure was repeated at least 12 times until the tissue became sperm free.

The sperms were separated from the luminal proteins in sperm suspension by centrifuging at 4000 rpm for 10 min. The supernatant containing luminal proteins were concentrated, dialyzed and stored at -20° in aliquots. Sperm protein extracts were prepared by lysing sperm pellet with a 1.5 ml solubilisation buffer (Tris, 1 % sodium dodecyl sulphate (SDS), 10 % glycerol, 10 mM PMSF) at 4° at a final concentration of 0.75–4×10⁵ sperm/µl. The suspension was then vigorously vortexed for 3 min.

Tissue extracts were prepared by homogenizing the tissues with 1.5 ml solubilisation buffer (Tris, 1 % SDS, 10 % glycerol, 10 mM PMSF) at 4°. The cellular debris was pellet at 11 000 rpm at 4° for 15 min. Samples were concentrated, dialyzed and stored at -20° before further treatments. The protein concentration of samples was estimated by Lowry's method^[15] using bovine serum albumin as the standard.

Electrophoresis:

About 30, 20 and 15 µg of tissue, sperm and luminal protein samples were denatured in equal volume of gel loading buffer (0.5 M Tris-Cl+10 % SDS+glycerol+βmercaptoethanol+0.025 % bromophenol blue), in boiling water bath at 95° for 5 min. Proteins were electrophoresed on a gradient SDS-polyacrylamide gel electrophoresis (PAGE) of 8-20 % resolving gel and 5 % stacking gel with a constant current of 35 mA until dye front entered the resolving gel, which was the increased to 55 mA thereafter till the dye front reached the end of the gel. Standard medium range molecular weight markers (Bangalore Genei, Mumbai) were also run along with tissue sample to determine the approximate molecular weight of sample proteins. Coomassie staining was carried out in order to observe the band pattern. Densitometry analysis of each sample and the standard markers were carried out on densitometer (Bio imaging, Syngene).

Statistical analysis:

Quantitative results are expressed as mean \pm SEM. The data were analysed by using independent student-t test and the differences in the control and treated groups were considered significant at p<0.05.

RESULTS AND DISCUSSION

The epididymal weights of Wistar rat was found to be unaltered after treatment with VCR, at a dose of $40 \,\mu$ g/kg for a period of 30 d, which is in contradiction with the previous reports, which has shown a decrease in the epididymis weight^[16]. This variation is probably because of the difference in the drug dose administered

to the rats. Sperm count is an important indicator of male fertility^[17] hence assessment of this parameter provides evidence on the effects of VCR on male fertility. A significant decrease (p<0.05) in the sperm count was observed in cauda (27.30 %) and caput (36.04 %) of treated rats compared with that of the control rats (fig. 1). The reduced sperm count observed in the current study could be due to the toxic effect of VCR on Leydig cells and testosterone levels, which was reported to cause sloughing of germinal epithelium and spermatogenesis^[18].

The concentration of sodium, potassium and magnesium in the cauda and caput regions of epididymis of control and VCR-treated rats are shown in Table 1. The metabolism, flagellar beating and acrosome reaction of spermatozoa are regulated by ion flux across the plasma membrane in mammals. Apical cells of epididymis play a major role in electrolyte transport for maturation events^[19]. Sodium ions are the principal cations that induce motility of spermatozoa and potassium has additive effects depending on concentration, with higher concentration being inhibitory^[14]. The decrease in sodium levels in caput and increase in magnesium



Fig. 1: Sperm count of cauda and caput segments of epididymis of control and treated rats

(**•**) Control, (**•**) vincristine-treated; the values are expressed as mean±SEM (p<0.05), '*' indicates statistically significant change. The unit for total sperm count is millions/ml

TABLE 1: CONCENTRATION OF CATIONS IN CONTROL AND VCR-TREATED RAT EPIDIDYMAL CAUDA AND CAPUT

Туре	lon concentration (ppm)		
	Sodium	Potassium	Magnesium
Control	0.794±0.05	0.82±0.005	0.104±0.02
Treated	0.753±0.05	0.78±0.04	0.144 [*] ±0.01
Control	1.221±0.08	0.72±0.03	0.120±0.02
Treated	0.808 [*] ±0.04	0.68±0.04	0.136±0.02
	Type Control Treated Control Treated	Ion co Sodium Control 0.794±0.05 Treated 0.753±0.05 Control 1.221±0.08 Treated 0.808'±0.04	Ion co-rentration Sodium Potassium Control 0.794±0.05 0.82±0.005 Treated 0.753±0.05 0.78±0.04 Control 1.221±0.08 0.72±0.03 Treated 0.808*±0.04 0.68±0.04

The values are expressed as mean \pm SEM (p<0.05), '*' indicates statistically significant change

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concentration of cauda seen in the present study could lead to an inhibitory effect on sperm motility of both caput and cauda region on VCR treatment, as optimal concentrations of these cations are required for sperm motility, which is specific for each species.

The total protein concentrations of tissue, sperm and luminal tissue from 3 treated and 3 control rat cauda and caput were determined using Folin Lowry method. No significant changes were observed in treated samples when compared with control (Table 2). Though there were no significant changes observed in total protein concentration, alteration in protein profile of these samples were noted when tissue, sperm and luminal extracts of epididymis was separated on gradient (8-20 %) SDS PAGE and band patterns were analysed using densitometry (Bio Imaging System, Syngene).

In all the samples, separation of the proteins revealed the presence of approximately 10-16 polypeptides ranging in size 12-280 kDa with major bands lying between 14-120 kDa (fig. 2). In cauda tissue protein, the densitometry graph (fig. 2A) showed that the polypeptide of molecular weight 66 kDa of treated samples did not show any change in the expression levels with respect to control samples. On the other hand peptides from the treated sample with molecular weight of 160, 95, 80, 41, 38, 27, 24, 19, 16 and 13 kDa showed increase whereas the one with the molecular weight of 31 kDa showed slight decrease in expression levels when compared to control samples. In caput tissue, the densitometry graph (fig. 2B) revealed that no polypeptides of treated samples showed any change in the expression levels with respect to control samples except that hose with molecular weight of 28, 20 and 14 kDa showed slight increase in the treated samples as compared to control.

In cauda sperm proteins, the densitometry graph (fig. 2C) showed that the polypeptide of molecular weight 17 kDa of treated samples did not show any change in the expression levels with respect to control samples. On the other hand, rest of all peptides from the treated sample showed increase when compared with control samples. In cauda sperm proteins, the densitometry graph (fig. 2D) showed that the polypeptide of molecular weight 17 kDa of treated samples did not show any change in the expression levels with respect to control samples. On the other hand, rest all peptides from the treated sample showed increase in expression levels when compared with control samples.

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Fig. 2: Gradient-SDS PAGE (8-20 %) gel electrophoresis showing protein profiling and densitometry graph A: control and vincristine-treated rat epididymal cauda tissue protein; B: control and vincristine-treated rat epididymal caput tissue protein; C: control and vincristine-treated rat epididymal cauda sperm protein; D: control and vincristine-treated rat epididymal caput sperm protein; E: control and vincristine-treated rat epididymal cauda sperm protein; D: control and vincristine-treated rat epididymal caput sperm protein; E: control and vincristine-treated rat epididymal cauda luminal protein; F: control and vincristine-treated rat epididymal caput luminal protein. Lane 1: corresponds to medium range marker. Lanes 2, 3, 4 display tissue protein profiles of epididymal cauda of control rats, lanes 5, 6, 7 display tissue protein profiles epididymal cauda of treated rats. 30 µg of proteins of control and treated were loaded in each well

TABLE 2: PROTEIN LEVELS OF CAUDA AND CAPUT OF CONTROL AND TREATED RATS

Tissue	Туре	Average protein concentration (mg/g of tissue)		
		Tissue protein	Sperm protein	12.16±2.78
Cauda	Control	10.82±0.93	3.55±0.04	8.64±0.61
	Treated	12.51±0.72	4.35±0.05	11.93±1.85
Caput	Control	18.17±0.84	6.96±0.11	12.64±0.87
	Treated	15.55±1.87	5.99±0.08	12.16±2.78

The values are expressed as mean±SEM (p<0.05)

In cauda luminal proteins, the densitometry graph (fig. 2E) showed that the polypeptides of molecular weights 50 and 45 kDa of treated samples did not show any change in the expression levels with respect to control samples. On the other hand, peptides with molecular weights 93, 70, 27, 24 and 19 kDa from the treated sample showed increase whereas the polypeptides with molecular weights 81, 55 and

43 kDa showed slight decrease respectively in expression levels when compared with control samples. In caput luminal proteins, the densitometry graph (fig. 2F) showed increase expression of peptides with molecular weights 65, 53 and 14 kDa whereas the polypeptides with molecular weights 61. 5, 43, 41, 35, 29, 21 and 18 kDa showed slight decrease respectively in expression levels when compared with control samples.

The 24 kDa protein level in tissue, sperm and lumen of cauda was increased in response to VCR treatment. On the basis of molecular weight comparison it likely to be glutathione peroxidase type 5 (24 kDa; GPX5), present in principal cells, sperm and lumen of caput and cauda, which plays a vital role in protecting the membranes of spermatozoa from the damaging effect of lipid peroxidation (by inactivating hydrogen peroxides and organic hydroperoxides) and/or preventing premature acrosome reaction^[20]. The increase in the antioxidant

enzyme, GPX5 protein suggested that the epididymis is probably exposed to oxidative stress on VCR treatment.

Two proteins of molecular weight 19 kDa and 14 kDa from tissue, sperm and lumen of caput, which were found to be depleted on VCR treatment, could most likely be glycosylated isoform of cystatin-related epididymal spermatogenic (CRES) and non-glycosylated isoform of CRES. CRES defines a new subgroup in the family 2 cystatins of cysteine protease inhibitors by virtue of its reproductive specific expression^[21]. It inhibits the serine protease pro-hormone convertase 2 (PC2), a subtilisin/ Kex2p-like endoprotease involved in prohormone processing required for spermatogenesis^[16]. CRES increase might halt spermatogenesis due to increased inhibition of PC2^[22]. This could be one of the probable reasons for depletion of sperm in epididymis on treatment with VCR.

In case of treated caput and cauda tissue, the level of a protein of molecular weight 27.5 kDa was increased, which is most likely to be galectin-3 (27.5 kDa), an IgE binding protein endogenous soluble lectin within the family called galectins that bind beta-galactosides. It is found to be localized primarily in the nucleus of the duct epithelial cells and has shown to be a necessary factor in the splicing of nuclear pre-mRNA^[23]. A 30.41 kDa caudal tissue protein that showed decrease on VCR treatment was thought to be HOXBES2 protein (30 kDa). Its decrease could lead to abnormalities in sperm physiology. In previous studies, a novel 30 kDa sperm homeoprotein, termed as HOXBES2 (HOXB2 homeodomain containing epididymis-specific sperm protein) has potential GPI modification site, which is required for proper acquisition of luminal proteins by spermatozoa during maturation^[24]. The decrease in the protein could lead to hamper in acquisition of protein by spermatozoa leading to impaired sperm maturation.

The luminal proteins from caput region, showed a decrease in 61.5 kDa protein which is probably gamma glutamyl-transpeptidase (GGT), that plays a vital role in sperm thiol oxidation required for sperm maturation and motility^[23]. This reduction in GGT could be because of depletion seen in 40 kDa luminal protein from both caput and cauda region, which is most likely to be basic-fibroblast growth factor found to be involved in regulation of GGT activity and GGT protein level in epididymis via MAP kinase pathway^[25]. A 26.8 kDa caudal luminal protein was found to be increased on treatment with VCR. This protein might be triosephosphate isomerase-1, which plays a role in sperm activity by acting on its metabolism^[26].

The analysis of sperm proteins revealed an increase in caudal sperm proteins of molecular weight 60 and 27 kDa in treated rats as compared to control, which are likely to be fertilin β and phosphotyrosine phosphatase (PTP), respectively. Fertilin β is a member of a growing family of sequence-related proteins called the metalloproteinase-like, disintegrin-like, cysteine-rich (MDC) family, and acts in regulation of spermatogenesis, sperm-oocyte binding and sperm-oolemma membrane fusion^[27] whereas PTP negatively controls the level of any particular protein phosphotyrosine that regulates sperm capacitation and motility^[28,29]. However, a 84.8 kDa caudal sperm protein showed a significant decrease on treatment, which could probably be Gelsolin (85.4 kDa) that plays a key role in regulating vacuolar V-ATPase recycling, which in turn helps in maintaining optimum pH required for sperm maturation and storage^[30]. A caput sperm protein of 37 kDa, showed a slight decrease on treatment with VCR, which is probably lactate dehydrogenase. It shows primary metabolism activity and probably helps in the maturation of sperms. Apart from these, changes were also observed in sperm proteins of molecular weight 90 and 74 kDa from caput and cauda, respectively, which could be β -D-galactosidase, an exo-glycosidase in intracellular digestion of specific terminal glycosyl residues from glycoproteins and glycolipids^[31].

Epididymis being the site of sperm development and maturation, any change in the ion concentration and the proteins that might play a role in various processes of sperm maturation and transit through the epididymal duct, would eventually lead to impaired sperm count as observed in the present study and further possibly to compromised sperm function. The rate of sperm movement and the protein acquisition by the sperms during its transit through epididymal lumen governs the sperm maturation event. Thus, alteration in proteins involved in sperm transport could lead to incomplete development of sperms. Additionally, variations in concentrations of sodium, potassium and magnesium ions could also be detrimental to sperm motility. These alterations in proteins involved in sperm protection and various stages of sperm development such as cytoplasmic extrusion and membrane stabilization might further contribute to the abnormal sperm count and impaired function, thus leading to infertility.

Conflicts of interest:

The authors declare that there is no conflict of interest.

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