Oxidative DNA damage and oxidant status in hypertensive patients- A preliminary study

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ABSTRACT

Essential hypertensive patients on monotherapy (Atenolol, a beta-blocker), belonging to the Jat Sikh Punjabi population sub-group, and normotensive healthy, age-, sex- and socio-economic status-matched control individuals belonging to the same population sub-group were assessed for oxidative DNA damage and oxidative stress. Leukocytic oxidative DNA damage (measured using modified enzymatic comet assay/SCGE assay), oxidant status (total oxidative stress, total antioxidant capacity, malondialdehyde levels and oxidative stress index) and lipid profile was determined for each studied participant. The mean oxidative DNA damage and oxidative status of hypertensive patient group was found to be higher than in the control group (p<0.001). Statistically significant increases in total cholesterol, LDL, VLDL and triglycerides were also observed in the hypertensive patient group (p<0.001). Oxidative DNA damage showed a statistically significant positive correlation with blood pressure (SBP and DBP) and total oxidative stress levels and negative correlation with total antioxidant capacity. The findings suggest that oxidative stress could be a cause of observed DNA damage. Studies on a larger cohort are underway to determine confirmation of these findings.

Key Words: Comet assay, Tail DNA percent, antioxidant capacity, malondialdehyde, oxidized purines, oxidized pyrimidines

INTRODUCTION

Essential hypertension (\geq 140/90mmHg) requires lifelong treatment for which combinational therapy is the general prescriptive measure (Sever and Messerli, 2011). However, effects of these levels of treatments may themself prove to be deleterious, adding to the compromised state of health of the hypertensive patients in causing and/or increasing morbidity in terms of increased genetic damage which increases susceptibility to malignancy. This may further be compounded by multidrug therapy. Recommended medications include calcium channel blockers, beta-blockers, ACE inhibitors, angiotensine II receptor blockers and diuretics (IHG, 2013) which may act as antioxidants (Wassmann *et al.*, 2004)and/or as receptor blockers/enzyme inhibitors (JNC7, 2004).

In fact, whether oxidative stress is a cause or a result of hypertension is not clear (Grossman, 2008) though the role of reactive oxygen species (ROS) in vascular function and in the development of hypertension is known (Touyz, 2004; Aggarwal *et al.*, 2013). In the present preliminary study, we report oxidative DNA damage in essential hypertensive patients on monotherapy viz. those on treatment with atenolol (beta-blocker). The significance of the study lies in the fact that monotherapy is an exceptional mode of therapy with few patients being prescribed a single drug. The patients had been prescribed Atenolol (beta-blocker) which acts by blocking beta-1 adrenergic receptors on heart muscle cells, preventing epinephrin and norepinephrin from stimulating the cardiovascular system as they increase heart rate, strength and activity leading to elevated blood pressure (Nezu *et al.*, 1985). This preliminary study further attempted to select a single population sub-group (Jat Sikh) to control for confounders based on variant genetic make-up in order to initiate a higher-order investigation.

MATERIALS AND METHODS:

Essential hypertensive patients on monotherapy (Atenolol) were identified in this hospital-based study to investigate for oxidative DNA damage in their peripheral blood leukocytes (PBL) and compare with genetic damage in normotensive healthy individuals. Oxidative DNA damage was assessed by the enzymatically modified comet assay (Collins *et al.*, 2002) to score for levels of oxidized purines and oxidized pyrimidines. Demographic information and anthropometric measurements for obesity assessment were also taken. In order to find support for etiology of oxidative DNA damage, the oxidative stress in patients and controls was also assessed by estimating serum levels of lipid peroxidation as malondialdehyde, the total antioxidant capacity (TAC) and total oxidative stress (TOS). The lipid profile was also assertained for all participants.

Blood pressure was taken after the study participants had been seated comfortably for ten minutes using a mercury sphygmomanometer as per AMA guidelines (ASH, 1992) and a record of the average of three readings were taken as the systolic blood pressure (SBP) and diastolic blood pressure (DBP) values.

Anthropometric measurements of height, weight, waist circumference (WC) and hip circumference (HC) were taken as per Weiner and Lourie (1981). The abdominal adiposity as a function of WC, waist hip ratio (WHR) and general obesity as a function of body mass index (BMI) were also determined according to recommended cut-offs for Indians (Misra *et al.*, 2009).

Intra-venous blood samples (~10ml from each participant) were used for genetic damage assessment and aliquots for serum separation were analyzed for lipid levels and oxidant status. The levels of the total cholesterol(TC), high density lipoproteins (HDL) and triglycerides(TG) were determined on an automated lipid analyzer (Erba, India) using kits (Angstrom Biotech, India), as per the manufacturer's instructions and the very low density lipoproteins (VLDL=triglycerides*1/5) and low density lipoproteins (LDL=total cholesterol-HDL-VLDL) were then calculated (Friedewald *et al.*, 1972).

Oxidative stress parameters of total antioxidant capacity (TAC), total oxidative stress (TOS) and malondialdehyde (MDA) levels were assessed in blood serum samples by standard methodology (Erel, 2004, 2005; Beuge and Aust, 1978). Oxidative stress index (OSI) was also calculated as ratio of TOS levels to TAC levels.

Genetic damage in PBL was determined using the modified single cell gel electrophoresis (SCGE/comet) assay with foramidopyrimidine DNA-glycolase (FPG) and endonuclease III (Endo III) enzymes (Collins *et al.*, 2002) after checking for cell viability(Coligan *et al.*, 2003). DNA damage was scored using the comet assay Software Program and a record of percent DNA in tail, olive tail moment (OTM), tail moment (TM) and tail length (TL) was made. The values are mean±S.E.M. Test of significance for categorical (χ^2) and continuous (t-test) variables were applied. Correlation and regression analyses were performed and significance levels were set at p≤0.001.Statistical analysis was carried out on SPSS version 16 for windows.

RESULTS

Demographic and clinical information of the study participants (Table 1) were matched in terms of age, gender, height, weight, hip circumference (HC), waist circumference (WC), body mass index (BMI) and waist hip ratio (WHR) but differed in relation to systolic blood pressure (SBP), pulse pressure (PP) and mean arterial pressure (MAP). The participants had no smoking habits and did not use mobile phone.

Serum lipid levels (Table 2) for all individuals were significantly elevated ($p\leq0.01$) in hypertensive patients. Oxidative stress levels of malondialdehyde (MDA), TOS, oxidative stress index (OSI) were significantly ($p\leq0.01$) increased in hypertensive patients while total antioxidant capacity (TAC) was significantly ($p\leq0.01$) reduced in hypertensive patients.

Oxidative damage to purines and pyrimidines was detectable following incubation with enzymes (Table 3) and showed significant elevation ($p \le 0.01$) for the assessed DNA damage parameters in patients from values observed in normotensive controls.

On correlation and regression analyses, DNA damage showed significant association with WHR (r=0.88, p=0.04), SBP and DBP in hypertensive patients while as hypothesized DNA damage was negatively

correlated with TAC and positive with TOS (Table 4) suggesting that increased oxidative stress could be leading cause to DNA damage.

DISCUSSION

The Atenolol- treated hypertensive patients were hyperlipidemic with manifestation of oxidative stress and oxidative DNA damage compared with normotensive individuals. TOS levels were two times raised in hypertensive patients compared to values in healthy individuals, while TAC levels were two times decreased in patient group. Increased oxidative stress manifests as lipid peroxidation, further producing additional ROS that can enhance oxidative damage to proteins and DNA and could lead to severe myocardial cell damage (Chen *et al.*, 1995). The role of ROS in cardiovascular dysfunction along with hyperlipidemia, diabetes, hypertension and chronic heart failure has been known (Taniayama *et al.*, 2003).

Hypertension is often associated with obesity (Wolk et al., 2003) and body mass index (BMI) is an indicator/risk factor for hypertension (Feldstein et al., 2005). With increase in age, BMI values and DNA damage also increases (Krajcovicova et al., 2007). Increase in waist hip ratio (WHR) also significantly increases pulse pressure (Feldstein et al., 2005). There is often an associated increase in various lipid components in hypertensive patients (Maharajan et al., 2008). In accordance with the results of the present study, hypertensive patients were deficient for antioxidant enzymes but with increased lipid peroxidation (MDA) and DNA oxidation byproducts (Redon et al., 2003). Quantitatively increased oxidative DNA damage was represented in hypertensive patients, while treatment decreased urinary 8-OHdG levels and TAS levels in treated hypertensive patients (Yildiz et al., 2008; Subash et al., 2010). Lee et al., (2005) also showed that DNA damage caused by ROS was more common in hypertensive patients than in normotensive while reduction of antioxidant enzymes was associated with increased level of oxidative DNA damage (Dittmar et al., 2008) as many antihypertensive agents are genotoxic and carcinogenic (Brambille et al., 2006). The monotherapy with atenolol in patients of the present study could also be genotoxic. Chromosomal breaks were two times higher in hypertensive patients on betablocker therapy as compared to values in control individuals in a study by (Telez et al., (2010). Thukral et al., (2012) have also reported a significant increase in DNA damage and dyslipidemia in Punjabi hypertensive patients. The increased DNA damage observed in hypertensive patients pose them to be at a risk for developing neoplasia and target organ damage.

CONCLUSION

These preliminary results emphasize the increased oxidative DNA damage and oxidative stress in essential hypertensive patients on atenolol treatment raising the possibility that hypertensive state is

associated with increased free radical damage to DNA. Further studies on a large scale on oxidative DNA damage in hypertensive patients can clarify the role of oxidative stress in the pathogenic mechanism of hypertension.

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Table1.Demographic, anthropometric and physiometric measurements of hypertensive patients and normotensive controls

Parameters	Hypertensive Patients (n=5)		Normotensive Controls (n=3)		
	Range	Mean±S.E.M.	Range	Mean±S.E.M.	
Age	60-78	66.2±3.32	58-70	67.00±5.57	
Height (cm)	152.4-186	166.96±5.53	152.4-186	169.47±9.70	
Weight (kg)	54-90	78.20±6.55	40-90	66.67±14.53	
Hip Circumference (cm)	90-166	123.80±17.13	102-117	108.67±4.41	
Waist Circumference (cm)	94-172	127.40±17.85	102-112	105.33±3.33	
Body Mass Index (kg/m ²)	23.25-33.79	28.04±2.16	17.22-26.01	22.49±2.68	
Waist Hip Ratio	0.55-1.05	0.77±0.11	0.55-0.67	0.62±0.04	
Systolic Blood pressure (mmHg)	140-180	156.00 [*] ±8.12	110-130	120.00±5.77	
Diastolic Blood Pressure (mmHg)	80-120	100.00±7.07	75-85	80.00±2.89	
Pulse Pressure (mmHg)	50-60	56.00 [*] ±2.45	30-45	40.00±5.00	
Mean Arterial Pressure (mmHg)	99.80-139.80	118.48 [*] ±7.34	89.85-99.85	93.20±3.32	

*Statistically significant differences between hypertensive patients and normal control individuals (Students' t-test; $p \le 0.05$)

Variables	Hypertensive Patients (Mean±S.E.M.)	Normotensive Controls (Mean±S.E.M.)	t-value	p-value
Cholesterol(TC)	168.62±10.27	104.97±15.55	3.415*	0.030
High Density lipoproteins (HDL)	47.04±3.68	22.03±5.91	3.825**	0.009
Low density Lipoproteins (LDL)	79.68±10.08	55.25±9.78	1.740	0.159
Triglycerides (TG)	209.48±13.61	38.47±11.94	3.526**	0.012
Very low density lipoproteins (VLDL)	41.89±2.72	27.69±2.38	3.526**	0.012
Malondialdehyde (MDA) (nmol/mL)	10.44±0.61	5.44±1.47	3.721**	0.010
Total Antioxidant capacity (TAC) (mmol Trolox Eqv./L)	0.37±0.04	0.77±0.07	5.621***	0.001
Total oxidative stress (TOS) $(\mu mol H_2O_2 Eq./L)$	0.70±0.06	0.35±0.04	4.113**	0.006
Oxidative stress Index (OSI)	0.20±0.04	0.05±0.01	3.090*	0.021

Table2. Lipid Levels and Oxidative Stress in essential hypertensive patients and normotensive controls

**Statistically highly significant ($p \le 0.01$) when compared to normal control individuals * Statistically significant ($p \le 0.05$) when compared to respective values in normal control individuals

Method of Comet assay	DNA Damage parameters	Hypertensive patients	Normotensive controls	t-value	p-value
Alkaline standard comet	Percent Tail DNA	22.08±1.96	5.25±1.79	5.754**	0.001
	Tail length (TL)	218.40±27.15	56.67±5.49	4.440**	0.004
	Tail Moment (TM)	49.07±8.91	2.90±1.07	3.877**	0.008
	Olive Tail Moment (OTM)	63.64±9.14	7.17±3.14	4.533**	0.004
Comet assay with enzyme Endonucleas e III (Endo III)	Percent Tail DNA	8.96±1.92	1.78±0.50	3.618*	0.018
	Tail length (TL)	51.8±9.73	19.67±0.88	3.290*	0.030
	Tail Moment (TM)	17.45±2.19	1.14±0.53	7.223***	0.001
	Olive Tail Moment (OTM)	20.03±3.24	2.04±0.70	5.420**	0.004
Comet assay with enzyme formamidop yrimidine DNA glycosylase (FPG)	Percent Tail DNA	11.39±2.52	2.15±0.19	3.663*	0.021
	Tail length (TL)	62.40±7.69	19.67±1.20	5.489**	0.005
	Tail Moment (TM)	19.20±2.07	1.36±0.77	8.060***	0.000
	Olive Tail Moment (OTM)	20.95±3.16	2.17±0.83	5.754**	0.003

Table 3. DNA damage parameters in Hypertensive patients and normotensive controls

***Very highly statistically significant ($p \le 0.001$),**Statistically highly significant ($p \le 0.01$), *Statistically significant ($p \le 0.05$) when compared to normal control individuals

Table 4. Pearson's correlation coefficients of blood pressure (SBP and DBP) and Oxidative Stress Parameters with DNA damage related parameters in hypertensive patients

Variables		SBP (mmHg) D		DBP (m	DBP (mmHg)		Total Antioxidant capacity (TAC) (mmol Trolox Eqy./L)		Total oxidative stress (TOS) (µmol H ₂ O ₂ Eq./L)		Oxidative stress index (OSI)	
		r- value	p-value	r- value	p-value	r- value	p-value	r- value	p-value	r- value	p-value	
Alkaline comet	Percent Tail DNA	0.773	0.125	0.656	0.230	-0.751	0.143	0.726	0.165	0.836	0.078	
	Tail length (TL)	0.673	0.213	0.839	0.076	-0.889*	0.044	0.914*	0.03	0.822	0.088	
	Tail moment (TM)	0.866	0.058	0.909*	0.032	-0.987**	0.002	0.991***	0.001	0.993 ^{**} *	0.001	
	Olive tail moment (OTM)	0.884*	0.046	0.885*	0.046	-0.965**	0.008	0.961**	0.009	0.993 ^{**}	0.001	
Comet assay with	Percent Tail DNA	0.950**	0.013	0.846	0.071	-0.797	0.107	0.788	0.113	0.884*	0.046	
enzyme Endonucle	Tail length (TL)	0.820	0.089	0.865	0.058	-0.879*	0.049	0.861	0.061	0.820	0.089	
ase III (Endo III)	Tail moment (TM)	0.872*	0.054	0.728	0.163	-0.733	0.158	0.709	0.180	0.828	0.083	
	Olive tail moment (OTM)	0.970**	0.006	0.902*	0.036	-0.734	0.158	0.751	0.143	0.826	0.085	
Comet assay with	Percent Tail DNA	0.832	0.832	0.667	0.219	-0.637	0.248	0.616	0.268	0.753	0.142	
enzyme formamido	Tail length (TL)	0.911*	0.032	0.910*	0.032	-0.825	0.086	0.816	0.092	0.810	0.096	
pyrimidine DNA	Tail moment (TM)	0.843	0.073	0.662	0.223	-0.532	0.356	0.499	0.392	0.615	0.269	
glycosylase (FPG)	Olive tail moment (OTM)	0.962**	0.009	0.869*	0.056	-0.704	0.185	0.713	0.176	0.800	0.104	

**statistically highly significant correlation (p≤0.01) * statistically significant correlation (p≤0.05)