

Augmented cytotoxic, mutagenic and genotoxic response triggered by carvedilol and celecoxib combinations

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It is understood that drugs regardless of their order of administration can exhibit drug interactions. Established on the fact that treatment of hypertension may last for decades and prolong usage of multiple drug regimen may induce substantial pathophysiological changes. Hence, This study was designed to evaluate the possible synergistic toxic effects of anti-hypertensive (carvedilol), and anti-inflammatory drug (celecoxib) alone and in combinations. Well-established MTT assay, Single Cell Gel Electrophoresis (SCGE) and Ames assay were employed to evaluate the toxicity at cellular level. Results from MTT assay on Vero cell line revealed that drug combinations have more pronounced anti-proliferative activity with combine IC₅₀ value of 13.7:47.8 µg/mL. Likewise, exposure of peripheral blood mononuclear cells with drug combinations revealed significant (P<0.05) DNA damage (Class 3) in a dose dependent manner at concentrations ≥ 0.78: 2.34 µg/mL. However, carvedilol and celecoxib were non mutagenic against either mutant strain (TA 100 and TA 98) and combinations have also shown mild to moderate mutagenic potential. Nevertheless, upon addition of metabolic activation enzyme, concentration <12.5:37.5 µg/plate exhibited significant (P<0.05) mutagenicity against both tester strains. In conclusion, this study provides additional genotoxicity and mutagenicity data that could be used in considering options for formulating regimens with reduced mutagenic potential.

Keywords: Carvedilol. Celecoxib. Mutagenicity. Genotoxicity. Cytotoxicity.

INTRODUCTION

Hypertension affects seventy six million people in the United States and 1 billion worldwide each year and projection depicts that by 2030 prevalence of hypertension will increase by 7.2% from estimates (Mills *et al.*, 2016). Essential hypertension, or hypertension of obscure cause, represents beyond 90% of instances of hypertension. It tends to cluster in families and represents a collection of genetically based diseases or syndromes with several resultant inherited biochemical abnormalities (Renna *et al.*, 2013). The resulting phenotypes can be modulated by various environmental factors, thereby altering the severity of blood pressure

elevation and the timing of hypertension onset. Numerous pathophysiologic factors have been implicated in the genesis of essential hypertension, obesity and diabetes among the most common ones. Hypertension has been considered as a silent killer amongst all other ailments due to its diminished signs and symptoms. Consequently, if hypertension remains undiagnosed (Blood Pressure [BP] failing to be checked previously), known to the patient, however, left untreated, and known hypertensive patients with low adherence to both antihypertensive drugs and lifestyle modification can invariably lead to chronic kidney disease (CKD) (Mancia *et al.*, 2009). Beta-Blockers are antihypertensive medication and, together with diuretics, have been the foundation of studies signifying their advantages on cardiovascular morbidity and mortality. Carvedilol is a vasodilating non-cardio specific third-generation β-blocker, unlike conventional beta blockers carvedilol has negative hemodynamic and

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metabolic effects (Fergus, Connell, Ferdinand, 2015). Due to these pleotropic activities carvedilol is becoming a preferred therapeutic option among medical practitioners. Invariably, uncontrolled hypertension often comes with co-morbidities. Musculoskeletal pain, associated with the inflammatory diseases has the high prevalence in hypertensive patients (Panoulas *et al.*, 2008). Studies have shown that vascular inflammation is one of the leading cause for the development of hypertension and arteriosclerosis (Gerli *et al.*, 2005; Savoia, Schiffrin, 2006). Henceforth, inflammation and hypertension go hand in hand. In this scenario, celecoxib, an over the counter non-steroidal anti-inflammatory drug (NSAID) with mark anti-inflammatory and analgesic properties, has become physician's drug of choice. It is approved for the treatment of rheumatoid arthritis, osteoarthritis, and acute pain (van Walsem *et al.*, 2015). The anti-inflammatory and analgesic properties of celecoxib result from inhibition of prostaglandin (PG) synthesis by selective inhibition of PG G/H synthase-2 usually stated as cyclooxygenase 2 (COX-2) (Tarr, Reuter, 2015). The use of carvedilol and celecoxib combination has been very common in clinical settings and regardless of their order of administration such drugs can exhibit drug interaction. Moreover, it is impossible to predict the outcome of these interactions of the drugs, based on the individual effect, in their combinations. Established on the fact that duration of treatment of hypertension may last for decades and prolong usage of multiple drug regimens may induce significant pathophysiological changes at cellular level. The overriding goal of this effort is to provide additional genotoxicity and mutagenicity data that could be used in considering options for formulating prophylaxis/treatment regimens with reduced mutagenic potential and thus reduced long-term risk for cancer.

MATERIAL AND METHODS

Shodhana Laboratories Private Limited, Hyderabad India, kindly donated analytical grade samples of carvedilol and celecoxib. Dulbecco's modified eagle medium (DMEM) (Cat no. 12120014, Caisson Lab, USA); Fetal bovine serum (Cat no. S181H-500, Bio-west, USA); MTT [3-(4,5-dimethylthiazolyl-2)-2,5-diphenyltetrazolium bromide] (Cat. no. 29893-1, Sigma-Aldrich, USA); dimethylsulfoxide (DMSO) (Cat. no. 0000179027, Paneac QuimicaSAU, Spain); metabolic activation system (S9) (Cat. no. 505189D, EBPI environmental Bio-detection products Inc., Canada); *Salmonella typhi* TA100 and TA98 tester stains (Environmental Bio-detection Products Incorporation (EBPI), Canada); Histopaque /

Lymphocyte Separating Medium (Cat no: S11230L0560, Biowest, France, Roswell Park Memorial Institute Medium (RPMI-1640) (Cat no; 0214003, Caisson, USA); Low melting point agarose (LMPA) (Cat. no. 10011055, Fine chemicals HT Company, UK); Normal melting point agarose (NMPA) (Cat no. 42083100, Fine chemicals HT Company, UK); Triton-X100 (Cat no. 242639, Panreac, Spain); sodium hydroxide (Cat. no. 58641, Batch no. 77620, Scharlau, Germany); sodium chloride (Cat. NOCHB3266, Riedel-de-Haen, Germany); hydrochloric acid (Cat. no. AB61, Batch no. 02110133, Labscan, Ireland); Nutrient broth (Cat. No. AM5077, LOT no. 1101, UK); purified Ager (Cat. no. 306200012, Duplus, UK); hydrochloric acid (HCl) (Batch no. 0240133, Cat. no. AB61, Lab scan, Ireland); ethidium bromide (LOT no. 1004DU23819; Cat. no. 1239-45-8, Bio basic Inc., Canada); histidine (Cat. no. 101937, MP biomedical, France); biotin (Cat. no, 20284101, Daejung Chem Ltd, Korea); sodium azide (Lot no. V11052400, Cat. No. 41900064-1, Bioworld, USA); phosphate buffer (Lot no. 2431B77, Cat. no. 41620016-1, Bioworld, USA); potassium dihydrogen phosphate (Lot. no. PO02570500, Batch no. 11650701), Methanol (Lot no. 2839070L, Analar (BDH) UK); Trisbase (Cat. no. 240202362, Lot. no. 26089, Bioworld, USA) and EDTA (Lot. no. 62720, RDH Lab, USA) were provided by the department of Pharmacology and Toxicology, University of Veterinary and Animal Sciences, Lahore, Pakistan.

Experimental design

Toxic concentrations tested in this experiment were carefully selected keeping all the following parameters under consideration; 1) maximum repeatable dose (MRD) which provides a preview of the toxicology of new and old drug combinations; 2) Toxicokinetic (TK) range which gives a formal data of drug action at peak plasma concentrations of unknown drugs, and 3) Dose incrimination (two fold increase) process to evaluate acute, sub-chronic and chronic drug toxicity. Finally relevance to effective dose of Carvedilol ($ED_{50}=0.04 \mu\text{g/mL}$) (Cheng *et al.*, 2007) and celecoxib ($ED_{50}=1.14 \mu\text{g/mL}$) (Ouellet, Riendeau, Percival, 2001) and clinical indications were also taken under consideration (Buckley, Dorato, 2009). Hence, selective range of cumulative dilutions of carvedilol (200 $\mu\text{g/mL}$, 150 $\mu\text{g/mL}$, 100 $\mu\text{g/mL}$, 50 $\mu\text{g/mL}$, 25 $\mu\text{g/mL}$, 12.5 $\mu\text{g/mL}$, 6.25 $\mu\text{g/mL}$, 3.12 $\mu\text{g/mL}$, 1.56 $\mu\text{g/mL}$, 0.78 $\mu\text{g/mL}$) and celecoxib (600 $\mu\text{g/mL}$, 450 $\mu\text{g/mL}$, 300 $\mu\text{g/mL}$, 150 $\mu\text{g/mL}$, 75 $\mu\text{g/mL}$, 37.5 $\mu\text{g/mL}$, 18.75 $\mu\text{g/mL}$, 9.34 $\mu\text{g/mL}$, 4.68 $\mu\text{g/mL}$, 2.34 $\mu\text{g/mL}$) were evaluated, alone and in combinations of increasing proportion of both drugs.

Cytotoxicity assay (MTT assay)

Treatment of cell line

DMEM supplemented with 20% fetal bovine serum was used to cultivate adherent Vero cell lines. Adherent cells were treated with each test dilutions individually and incubated for 48 h in 5% CO₂ 95% air at 37°C. After 48 h, viability of Vero cells was measured by adding 20 µL MTT dye solution (50 mg/10 mL) in each well and cells were further incubated for 3 h (Ohno, Abe, 1991). Once this incubation time was completed, 100 µL of 10% DMSO were added to the 96 wells. The numbers of the viable cells in each well was proportional to the intensity of the absorbance of light, which was then read in an ELISA plate reader at 570 nm (Mosmann, 1983). Moreover, cell survival percentage values were used to evaluate the IC₅₀ values of carvedilol, celecoxib and their combinations using prism graphpad 5.

$$\text{Cell Survival Percentage (CSP)} = \frac{\left(\text{Mean optical density of test chemical} \right) - \left(\text{Mean optical density of negative control} \right)}{\text{Mean optical density of positive control}} \times 100$$

Genotoxicity (comet assay)

Lymphocytes separating media was used for the extraction of peripheral mononuclear cells (PMN). After incubating the PMNs with drug concentrations, group of single cells were transferred on frosted cavity slides. These pits were filled with Normal melting ager

(NMA), 0.6% agar layer on top of a 0.6% agar base layer, a third top layer comprising of 0.6% low liquefying point agar. Slides were placed in lysis solution (2.5 M NaCl, 100 mM EDTA, 10 mM Tris, 1%, 10% DMSO and 1% Triton) for 18-24 h and later were submersed in electrophoresis buffer (0.3 M NaOH, 1 mM EDTA) for 40 min, this allowed the DNA strand to break and unwind. Later, the power supply was adjusted at 24 volts and 300 mA. Slides were electrophoresed for 30 min. Slides were taken out after 30 minutes and placed on a dry surface. These slides were then neutralized and stained with ethidium bromide (10 mg/50 mL). DNA damage was determined by measuring the DNA comet length from the core. Quantitation was carried out on a fluorescent image analysis system that compared the overall length of intact nuclei versus damaged nuclei with a comet tail. The comets were observed under fluorescent microscope (BX41, product no. 5M16529 TF, Olympus, Japan) and pictures were taken with digital microscope camera (DP70, product no. 6B05172, Olympus, Japan) and analyzed through Image J software for percentage fragmentation. Finally comets are characterize in four different classes (Singh *et al.*, 1988).

Class 0 = Undamaged Nuclei

Class 1 = Tail length ≤ Undamaged Nuclei head diameter

Class 2 = Tail length > Undamaged Nuclei head diameter but < 2 X Nuclei head diameter

Class 3 = Tail length > 2 X Undamaged Nuclei head diameter.

TABLE I - Cell viability of Vero cell line was measured in term of optical density using MTT assay after treatment with various doses of Carvedilol, Celecoxib and their combinations with respective controls for 48 hours.

Carvedilol		Celecoxib		Carvedilol: Celecoxib		Controls	
Conc. ^a	OD ^b ±S.D	Conc.	OD±S.D	Conc.	OD±S.D	Conc.	OD±S.D
0.78	0.89±0.06	2.34	0.86±0.01	0.78:2.34	0.79±0.02	Positive	0.06±0.08
1.56	0.74±0.05	4.68	0.83±0.02	1.56:4.68	0.77±0.08	Control (Blank wells)	
3.12	0.69±0.07	9.37	0.77±0.02	3.12:9.37	0.74±0.11		
6.25	0.64±0.01	18.75	0.70±0.03	6.25:18.75	0.65±0.01*		
12.5	0.59±0.01	37.5	0.67±0.02	12.5:37.5	0.56±0.07*	Negative	0.97±0.07
25	0.49±0.001*	75	0.56±0.002	25:75	0.49±0.09*	Control (DMEM)	
50	0.43±0.04*	150	0.45±0.03*	50:150	0.41±0.05**		
100	0.35±0.03*	300	0.39±0.01**	100:300	0.38±0.03***		
150	0.33±0.02**	450	0.35±0.02***	150:450	0.34±0.03***		
200	0.29±0.06**	600	0.32±0.02***	200:600	0.29±0.01***		

^aConcentration = µg/ml, ^bOD= Mean optical density of the cells against the control exposed to that concentration in terms of mean± standard deviation. Statistical significance was analyzed using the two-way ANOVA followed by Post Hoc Duncan, *P < 0.05, **P < 0.01, ***P < 0.001 vs positive control, * P<0.05, **P < 0.01, ***P < 0.001 vs positive control.

Damage index = No. Of cells in Class 1 + (2 × No. of the cells in Class 2) + (3 × No. of cells in Class 3) (2)

Mutagenicity (AMES assay)

Pre-incubation assay was utilized for mutagenicity evaluation of Carvedilol and Celecoxib. Mutagenic potential of carvedilol, celecoxib and combinations were evaluated on two *Salmonella typhi* tester strains TA100 and TA98. Both tester strains TA 98 and TA 100 were inoculated in a fresh sterilized nutrient broth for 15-18 h prior to the assay performance. The Bacterial number was balanced at 0.5 mcfarland (mcf) by adjusting the optical density using spectrophotometer. Each drug dilution was exposed with 0.1 mL of bacterial suspension (2 × 10⁸ bacteria), incubated for 20 minutes and then spread over GM ager plates. Plates were then incubated for 48 h at 37 °C. Likewise, the procedure was replicated with metabolic activation system (S9 mix) by adding 0.5 mL of the S9 fraction in each drug dilution. Finally, histidine independent revertant colonies were counted manually and results were calculated in term of mutagenic index by the following formula (Mortelmans, Zeiger, 2000)

Mutagenic Index

$$\frac{\left(\text{Number of revertant colonies per plate with test chemical dose} \right)}{\left(\text{Number of natural revertant colonies of negative control plate} \right)}$$

Negative Control × 2 = Possible Mutagen, Negative Control × 3 = Significant Mutagen, Negative Control × 4 and above = Highly Significant Mutagen

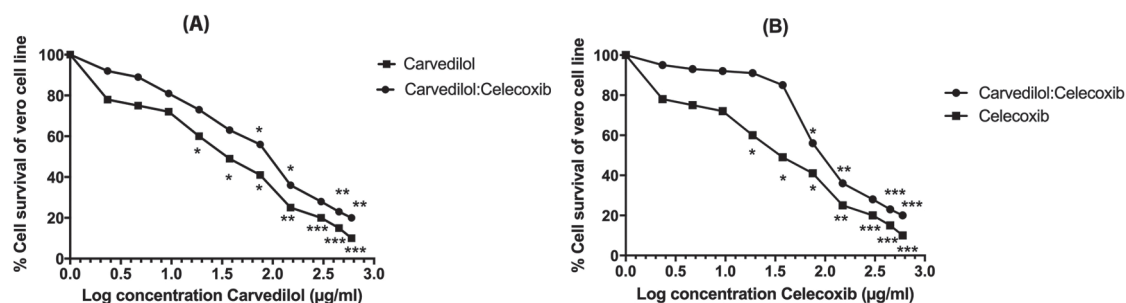


FIGURE 1 - Concentration dependent response in cell survival percentage (CSP) of Vero cells (2 × 10⁴ cells/0.1 mL/well) assessed by MTT assay after 48 hours of incubation with test chemicals, (A) comparison of CSP after carvedilol and combine concentration of carvedilol and celecoxib (B) comparison of CSP after celecoxib and combine concentration of carvedilol and celecoxib. The graph shows that anti-proliferative effect of combine concentration carvedilol:celecoxib was significant at lower level as compare to both carvedilol and celecoxib alone. Each value represent of mean three replicates (n=3). The asterisks indicate significant difference at three levels (*p<0.05, **p<0.01, ***p<0.001) between cells cultured after 48 hours of treatments.

Statistical analysis

The results were analyzed by Statistical Package Of Social Sciences (SPSS for windows version 16, SPSS Inc., Chicago IL USA). The results were explained in terms of means ± SD. The results were considered significant if the (p<0.05), (p<0.01) and (p<0.001). Analysis of variance (TWO WAYANOVA) along with Post Hoc Duncan test was applied to the data to find out the relation between the DNA damage index, Cell survival percentage and the relationship between the means of mutagenic index and dose response among the treatments (Aghbali *et al.*, 2014).

RESULTS

MTT assay

After incubating the cells for 48hrs with test samples, results were presented in form of cell survival percentage (CSP) (Figure 1). All three treatments significantly (P<0.05) inhibited the proliferation of Vero cell in a concentration-dependent manner. Recorded IC₅₀ for carvedilol and celecoxib were 20.4±0.2 µg/mL and 84.7±0.3 µg/mL, respectively, whereas their combinations have shown increase pattern of cytotoxicity with a combine IC₅₀ of 13.7:47.8 µg/mL. The effects of combinations were more cytotoxic as compared to individual drugs (Figure 1).

Comet assay

The genotoxic potentials of carvedilol, and celecoxib were evaluated on peripheral mononuclear cells and extent of the DNA damage was expressed in terms of percentage fragmentation of DNA (Table II). Carvedilol has exhibited

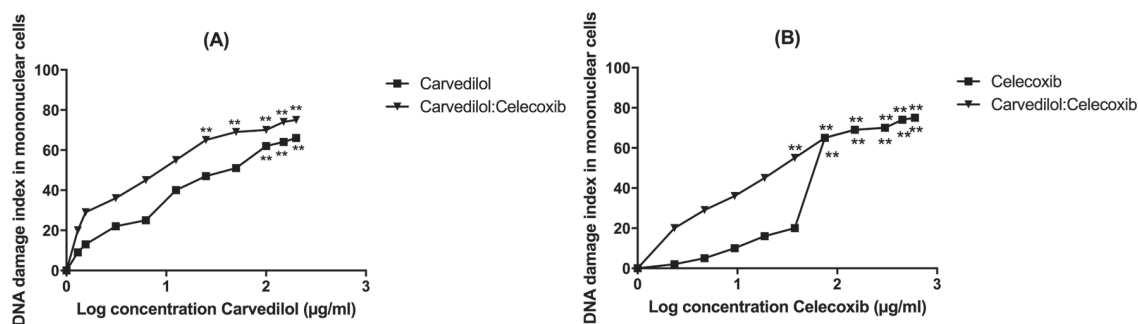


FIGURE 2 - DNA damage index of carvedilol, celecoxib and their combination (carvedilol:celecoxib) in peripheral blood mononuclear cells (PBMCs) measured by Single Cell Gel Electrophoresis, (A) comparison of DNA damage index after carvedilol and combination (B) comparison of DNA damage index after celecoxib and combination. The graphs shows that the DNA damage induced after treatment with carvedilol and combine concentrations were significant. Nevertheless DNA damages induced by >75 µg/mL of celecoxib were also genotoxic when used alone and combinations. Each Value represent of mean three replicates (n=3). The asterisk indicate significant difference at **p<0.01 between cells.

maximum DNA damage at 200 µg/mL leading to Class 3 tail formation (Figure 3). Moreover, at lowest tested concentration of Carvedilol (0.78 µg/mL), head to tail ratio was approximately 2:1 with 10% DNA fragmentation and class 1 damage (Figure 3). Whereas for celecoxib damage was only observed at concentrations ≥ 75 µg/mL with class 3 damage and 100% fragmentation (Figure 3, Table II). Furthermore, maximum tail length formation was observed with carvedilol and celecoxib combinations, where an average of class 3 damage with as average of 30% fragmentation was recorded at concentration range of 0.78:2.34 µg/mL to 200:600 µg/mL (Table II), suggesting that combination of carvedilol and celecoxib has exhibited synergetic genotoxic potential (Figure 3).

AMES assay

Mutagenic potential of Carvedilol, Celecoxib and combinations were evaluated on two *Salmonella typhi* tester strains TA100 and TA98. After incubating the mutant strains for 48hrs with the test concentrations, results were presented in terms of number of revertant colonies/plate and mutagenic index as shown in Table III. As expected, carvedilol and celecoxib failed to produce histidine independent revertant colonies in both TA98 and TA 100. However, mild to moderate mutagenicity was observed when combination of Carvedilol and Celecoxib were incubated with mutant strain and metabolic activation system was added to mimic the *in vivo* condition. Upon addition of metabolic activation system histidine independent revertant colonies were observed at concentrations ≥ 3.12 : 9.38 µg/plate, with mutagenic index of ≥ 2.0 (Table III). On the other hand, the combinations of both drugs were not mutagenic at

therapeutic concentrations of 0.04 µg/mL and 1.14 µg/mL of carvedilol and celecoxib, respectively (Table III).

DISCUSSION

The superior therapeutic efficacy of carvedilol in lowering blood pressure and of celecoxib on inflammatory pains motivated physician to prescribe this combination for multiple years and in some cases for entire life. The possible cytotoxic, genotoxic and mutagenic effect of combination of carvedilol and celecoxib were evaluated in this study using well-established MTT, Ames and SCGE *in vitro* assay. Carvedilol has shown promising results in several anti-proliferative studies. Proposed mechanisms includes (a) Ca²⁺ release from the endoplasmic reticulum which leads to apoptosis (Cheng *et al.*, 2007) and (b) direct mitochondrial damage causing cells arrest at G0/G1 phase (Erguven *et al.*, 2010). In this experiment, carvedilol has shown substantial cytotoxicity at concentrations above 12.5 µg/mL, and IC₅₀ 20.4±0.22 µg/mL (Figure 1). The following concentrations of carvedilol; 25 µg/mL, 50 µg/mL, 100 g/mL, 150 µg/mL and 200 µg/mL have statistically significant difference in CSP as compared to control (Figure 1), whereas celecoxib results suggested that concentrations >75 µg/mL are potential cytotoxic for given cell line with IC₅₀ 84.7±0.34 µg/mL (Table I). Literature suggests that its COX-2 growth-inhibitory and apoptosis-stimulatory potential is primarily responsible for its noticeable anti-proliferative activity (Zuo *et al.*, 2015). Furthermore, test concentrations of combinations produced synergistic cytotoxic potential and cell viability of Vero cell line plunged abruptly upon exposure. Seven combined concentrations (6.25:18.75 µg/mL, 12.5:37.5 µg/mL, 25:75 µg/mL, 50:150 µg/mL, 100:300 µg/mL,

TABLE II - Percentage fragmentations of DNAs in the peripheral blood mononuclear cells (PBMCs) measured by single cell gel electrophoresis (comet assay) after 3 h incubation with various concentrations of Carvedilol, Celecoxib and their combination compared to respective controls.

Treatment	Concentration (µg/ml)	Mean DNA Head Length (µm)	Mean Tail length (µm)	%Fragmentation
Carvedilol	200	1.97 ±0.65	8.46±2.54	100**
	150	1.96±0.80	8.09±2.41	92**
	100	2.15±0.16	7.36±2.10	92**
	50	2.16±0.98	6.21±2.41	84**
	25	2.20±0.55	5.00±1.95	76
	12.6	2.41±0.31	4.68±2.56	68
	6.25	3.51±0.40	3.418±1.48	60
	3.125	3.57±1.48	2.59±0.57	52
	1.56	3.76±1.91	2.09±1.59	48
	0.78	3.99±1.06	2.33±1.18	10
Celecoxib	600	2.59 ±0.392	10.81±0.701	100**
	450	2.66±0.410	10.07±0.805	100**
	300	3.06±0.1903	7.03±2.134	100**
	150	4.34±01.468	6.21±2.411	100**
	75	2.20±0.559	4.36±0.662	92**
	37.5	4.15±1.601	0.482±1.279	36
	18.75	4.16±0.573	0.489±1.125	36
	9.375	4.50±1.125	0±0.00	10
	4.69	5.22±0.941	0±0.00	9
	2.34	5.76±1.063	0±0.00	5
Carvedilol:Celecoxib	200:600	1.68 ±0.81	9.46±1.49	100**
	150:450	1.77±1.49	8.28±0.94	100**
	100:300	2.01±0.53	7.70±0.66	100**
	50:150	2.59±0.67	6.06±1.52	100**
	25:75	3.21±0.68	4.92±1.58	92**
	12.5:37.5	3.33±1.12	4.51±0.95	88**
	6.25:18.75	3.42±0.63	4.07±1.12	76
	3.125:9.375	4.50±1.12	3.78±1.31	64
	1.56:4.69	5.22±0.94	3.27±1.12	56
	0.78:2.34	5.76±1.06	3.11±0.56	36
Positive Control	20 % DMSO	0.78±0.011	13.3±0.333	78
Negative Control	RPMI Medium	5.78±0.166	0.37±0.002	8

150:450 µg/mL, 200:600 µg/mL) showed statistically significant difference in CSP as compare to control cells (Table I) exhibiting an increase cytotoxicity of combination. Previously, both drugs have been analyzed in combination with known anticancer drugs. Carvedilol has shown greater cytotoxicity with imatinib against brain tumor (Erguven *et al.*, 2010) and with doxorubicin against

breast cancer (Jonsson *et al.*, 1999). Data suggested that similar attribute has been associated with Celecoxib as well, combination of Cisplatin and Celecoxib has enhanced therapeutic response against chemotherapy resistant gastric cancer (Xu, Shen, Lv *et al.*, 2016), with Imatinib, Celecoxib has shown superior efficacy against Chronic myelogenous leukemia (CML) by suppressing

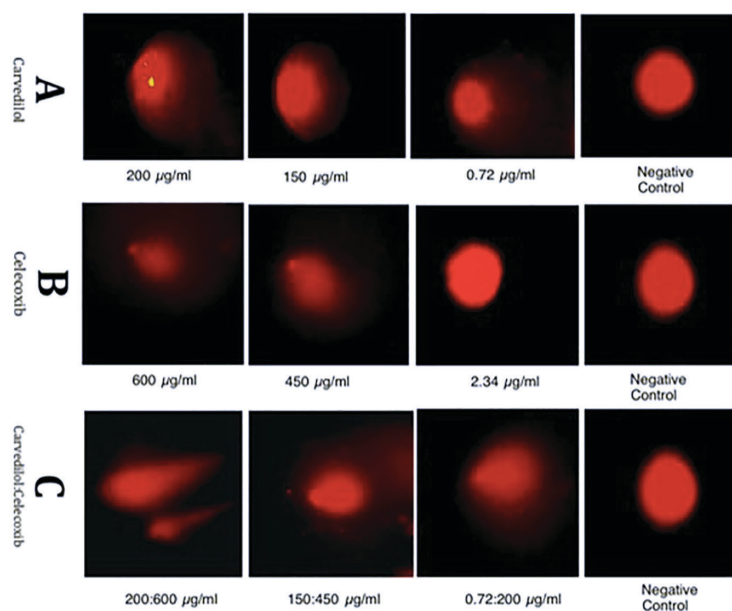


FIGURE 3 - Representative comets with increasing concentrations of three treatment groups. [A] shows the comets protrude by carvedilol 200 µg/mL, 150 µg/mL, 0.72 µg/mL and negative control (Class 0 Damage) [B] Shows the comets protrude by celecoxib 600 µg/mL, 450 µg/mL and 2.34 µg/mL and negative control [C] Shows the comets protrude by combine concentrations of carvedilol and celecoxib 200:600 µg/mL, 150:450 µg/mL, 0.72:2.34 µg/mL and negative control (Class 0 Damage) respectively.

TABLE 3. Shows the mutagenic potential in terms of mean number of revertant colonies/plate, standard deviation (SD) and mutagenic index (M.I) recorded against *salmonella typhi* TA-98 and TA-100, in the presence (+S9) and absence (-S9) of metabolic activation system.

Treatment µg/Plate	Mean Number of revertant (M ± SD)/plate and MI in parenthesis				
	TA 98		TA 100		
	-S9	+S9	-S9	+S9	
Carvedilol	0.78	54±2 (0.83)	79±3 (0.89)	89±3 (0.90)	103±5 (0.88)
	1.56	63±4 (0.96)	106±6 (1.21)	99±4 (1.10)	110±6 (0.94)
	3.12	69±7 (1.06)	124±3 (1.41)	104±3 (1.06)	131±4 (1.11)
	6.26	73±3 (1.12)	134±7 (1.53)	130±5 (1.33)	177±8 (1.50)
	12.5	88±5 (1.36)	149±4 (1.70)	154±6 (1.57)	622±3 (1.88)
Celecoxib	2.34	52±3 (0.81)	81±3 (0.92)	81±2 (0.82)	98±5 (0.83)
	4.69	60±2 (0.91)	88±1 (1)	93±3 (0.94)	114±4 (0.96)
	9.37	63±6 (0.96)	95±7 (1.07)	101±7 (1.03)	141±8 (1.19)
	18.75	79±1 (1.21)	144±9 (1.64)	156±6 (1.59)	198±5 (1.67)
	37.5	93±8 (1.43)	156±5 (1.77)	169±8 (1.72)	216±11(1.83)
Carvedilol:Celecoxib	0.78:2.34	88±2 (1.35)	146±3 (1.66)	155±5 (1.58)	205±7 (1.74)
	1.56:4.68	100±4 (1.55)	157±4 (1.79)	175±4 (1.78)	219±4 (1.86)
	3.12:9.73	115±7 (1.77)	172±8 (1.96)*	189±8 (1.92)*	244±8 (2.07)*
	6.25:18.75	145±4 (2.24)**	218±4 (2.48)**	213±5 (2.17)*	333±6 (2.82)*
	12.5:37.5	166±7 (2.55)**	248±6 (2.82)**	268±6 (2.73)**	353±9 (2.99)***
	Positive Control	540±65 ^b	1220±70 ^c	756±53 ^b	1540±88 ^c
Negative Control	65±4 ^a	88±5 ^a	98±4 ^a	118±6 ^a	

M±SD= mean and standard deviation; MI= mutagenic index; ^aNegative control: Dimethylsulfoxide (50µl/plate);^bPositive control sodium Azide with out S9 (5µg/plate); ^c2-amino Anthracycline (800µl/plate) with S9. Control-×2*, Control-×3**, Control-×4 and above ***. *=Possible Mutagen; **=Significant Mutagen; ***Highly Significant Mutagen

the autophagy and increasing cytotoxicity. Celecoxib with radiotherapy has shown promising results by G2-M cell phase arrest and enhancement of cell apoptosis (Zhang *et al.*, 2014). Hence, molecular target therapy can be used to target advanced malignancies of certain types and carvedilol in combination with celecoxib can be an option in the treatment or prevention of neoplasm particularly in hypertensive patients.

The genotoxicity evaluation of carvedilol and celecoxib were assessed using single cell gel electrophoresis (Comet assay) on peripheral blood mononuclear cells. There is a paucity of information on the toxicity of carvedilol in literature. However, few shreds of evidence have been found in literature to support our data (Table II). Aruna and Krishnamurthy (1986) Suggested induction of a significant increase in the frequency of micronuclei observed in erythrocyte at higher dose levels of propranolol. Many years later Sedigh-Ardekani, Saadat and Saadat (2013) shared the same concern about the non-selective beta blockers. Our data indicates that significant tail formation was observed in all the test concentrations range of 0.78 µg/mL-200 µg/mL of Carvedilol, when compared with negative control (Figure 3). On the contrary, DNA damage was only observed at concentrations ≥ 75 µg/mL for celecoxib. But at concentrations < 75 µg/mL the tail formation decreased momentarily, showing class 0, insignificant damage (Figure 3). Moreover, the combination of carvedilol and celecoxib were subjected to SCGE for evaluation of their genotoxicity potentials. Out of all the tested concentrations four have shown significant ($p < 0.05$) DNA damage (Table II). The reason of these concentrations for being conspicuous is because reported genotoxic concentrations fall under threshold value of combined IC_{50} (13.4:47.9 µg/mL). Henceforth, these concentration may bring about DNA damage yet not kill them. The consolidated impact of DNA damage, DNA repair defects, and an inability to stop or slow down the cell cycle (Lord, Ashworth, 2012). Our results also suggest a synergetic genotoxicity when our test drugs were used in combinations (Figure 3). Previous studies carried out on effects of drugs in combinations also support our data (Carter *et al.*, 2007, Guimaraes *et al.*, 2003).

DNA damage and chromosomal aberration can be a major cause for the development and progression of cancer (Hagmar *et al.*, 2004). To validate this claim we tested carvedilol and celecoxib against *Salmonella typhi* TA98 and TA 100 for the evaluation of their mutagenicity potential alone and in combinations (Table III). Individual test concentrations of carvedilol and celecoxib did not show any signs of mutation against either strains of

Salmonella typhi. Nevertheless, upon addition of metabolic activation system significant rise in mutagenic index was observed. The concentrations, which were mutagenic for TA-100, were almost equally mutagenic for TA-98. The three out of five concentrations used in this project have exhibited mild to moderate mutagenic potential (Table III). Carvedilol and Celecoxib combinations have shown significant ($P < 0.05$) mutagenic potential, against negative control at following concentrations 3.12.9.73 µg/mL, 6.25:18.75 µg/mL and 12.5:37.5 µg/mL. Several reports regarding the mutagenic synergism by combination of drugs are in support of our data. (Mansky, 2003, Torres *et al.*, 2007). For instance, in recent work of Fatima *et al.* (2013) emphasized that all anti-TB drugs, specifically in combination, possess substantial cytotoxic and mutagenic potential making TB patient more susceptible to cytotoxic and mutagenic effects of anti-TB drugs, which could produce further health complications. These collective data suggest genomic instability is one of the most pervasive characteristics of tumor cells and is probably the combined effect of DNA damage, tumor-specific DNA repair defects, and a failure to stop or stall the cell cycle before the damaged DNA is passed on to daughter cells, hence our drug combinations may cause host cell DNA damage and mutations, and impose a cancer risk.

CONCLUSION

These collective data suggest Genomic instability is one of the most pervasive characteristics of tumor cells and is probably the combined effect of DNA damage, tumor-specific DNA repair defects, and a failure to stop or stall the cell cycle before the damaged DNA is passed on to daughter cells, hence our drug combinations may cause host cell DNA damage and mutations, and impose a cancer risk. It was concluded from the results that combination of Carvedilol and Celecoxib can be mutagenic, genotoxic and cytotoxic. Professional medical advice should be recommended before taking these medications in combination. These drugs should be taken at low concentrations for a shortest period of time.

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CONFLICT OF INTEREST

We report no conflict of interest for the present paper

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