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Effect of dietary ascorbate on covalent binding of benzene to bone marrow and hepatic tissue *in vivo**

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Benzene produces aplastic anemia in humans and is a human leukemogen [1, 2]. The biochemical mechanics whereby benzene produces these effects are presently unknown. However, several reports have demonstrated a relationship between covalent binding in the bone marrow, the target site, and benzene-induced blood dyscrasia in laboratory animals [3, 4]. Toxicity appears to be the result of the metabolic activation of benzene [5, 6]. Ascorbate has been shown to be effective in preventing the *in vitro* covalent binding of the metabolites of benzene and phenol, the major metabolite of benzene [7–9]. We have demonstrated previously that, when phenol and hydrogen peroxide are incubated with bone marrow preparations isolated from guinea pigs with low ascorbate tissue concentration, there is a 4-fold increase in *in vitro* covalent binding of phenol equivalents to bone marrow tissue when compared to guinea pigs on a high ascorbate intake [9]. Ascorbate cannot be synthesized in the guinea pig or human and must be procured through dietary intake, and the concentrations of ascorbate in tissue can vary widely from individual to individual due to their respective dietary intake [10]. In the present communication, we have investigated the effect of ascorbate on covalent binding of [¹⁴C]benzene metabolites *in vivo*. To evaluate the effect of ascorbate on covalent binding *in vivo*, guinea pigs were placed on different dietary intakes of ascorbate followed by i.p. administration of [¹⁴C]benzene.

Covalent binding was inversely related to dietary ascorbate intake and to the concentration of ascorbate in both liver and bone marrow. Covalent binding was altered by 2- and 1.4-fold in the liver and bone marrow respectively. The concentration of ascorbate in bone marrow and hepatic tissue ranged from 0.12 to 2.63 $\mu\text{moles/g}$ tissue depending upon the dietary intake.

Three groups of four male Hartley guinea pigs (200–250 g) were placed on the following dietary intakes of ascorbate: 2.0, 0.35 and 0.05 mg ascorbate/ml drinking water for 4 weeks prior to the i.p. administration of benzene. All guinea pigs received ascorbate-deficient guinea pig chow *ad lib.* (ICN Nutritional Biochemicals, Cleveland, OH). The growth rate for each group of guinea pigs was not significantly different. The guinea pigs on the lowest intake showed no signs of a scorbatic condition. After 4 weeks on the diets, each guinea pig was injected with 660 mg/kg [¹⁴C]benzene intraperitoneally twice, 12 and 6 hr before termination. The specific activity of [¹⁴C]benzene was either 52.7 or 15.2 $\mu\text{Ci}/\mu\text{ol}$. [¹⁴C]Benzene was injected in 0.5 ml corn oil. Six hours after the second injection the guinea pigs were terminated by decapitation. Each liver was removed, and a 20% homogenate was made with 100 mM sodium phosphate buffer, pH 7.4. Femurs were removed from each guinea pig, and the bone marrow was scraped out of the femoral cavity. Each marrow sample was weighed and a 10% homogenate was prepared. Hepatic homogenate (250 μl) and bone marrow homogenate (1.2 ml) were extracted, and the covalently bound benzene equivalents in these tissues were determined by the method

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of Tunek *et al.* [11]. Bone marrow and hepatic ascorbate concentrations were measured in these guinea pigs by the method of Zannoni *et al.* [12].

The concentrations of ascorbate in bone marrow and liver of the benzene-treated guinea pigs are shown in Table 1. There was approximately a 10-fold difference in the ascorbate tissue concentration in both liver and bone marrow when the low dietary intake of ascorbate was compared to the high. As shown in Fig. 1, covalent binding of benzene metabolites to hepatic tissue was decreased as the hepatic ascorbate tissue concentration was increased. Covalent binding to hepatic tissue was decreased in the order of 2-fold when the hepatic ascorbate tissue concentration was increased approximately 10-fold. Covalent binding of benzene equivalents to bone marrow tissue also varied inversely with ascorbate tissue concentration (Fig. 2). Bone marrow ascorbate tissue concentration could also be modulated over 10-fold, while covalent binding could be decreased 1.4-fold.

In the present communication we have demonstrated that ascorbate is capable of altering the covalent binding of benzene metabolites to hepatic or bone marrow tissue *in vivo*. Covalent binding was inversely related to ascorbate intake and to the concentration of ascorbate in these tissues. The difference in covalent binding between groups may be important with regard to the myelotoxicity of benzene since several investigators have shown that a decrease in covalent binding to bone marrow tissue results in the decreased toxicity of benzene [3, 4].

The differences in covalent binding are small when compared to the differences in the concentration of ascorbate in these tissues. We have demonstrated previously that ascorbate via its antioxidant activity is capable of inhibiting the *in vitro* covalent binding of the metabolites of phenol by over 90% and, when benzene is employed as the substrate, covalent binding is inhibited by 35% and this changes only slightly when the concentration of ascorbate is increased from 1 to 5 mM [9]. Similarly, in the present study the *in vivo* covalent binding of benzene equivalents after i.p. administration of benzene was decreased 30%, while the bone marrow ascorbate concentration was altered 10-fold. These data may suggest that the reactive species responsible for the observed covalent binding is only slightly refractory or non-refractory to the antioxidant effect of ascorbic acid. This is in keeping with the concept that the major species responsible for the observed *in vivo* covalent binding may not be the oxidized forms (hydroquinone, catechol or phenol) since we have demonstrated previously that ascorbic acid is very effective in reducing these metabolites and thus preventing their interaction with tissue macromolecules [8, 9]. Another possible explanation to consider is that the cellular distribution of ascorbate is such that it is not at a substantial concentration at the site of reactive metabolite formation. Nevertheless, the dietary intake of ascorbate does alter the covalent binding of benzene metabolites to both bone marrow and hepatic tissue *in vivo*, and this event in the bone marrow is associated with the myelotoxicity of benzene [3, 4].

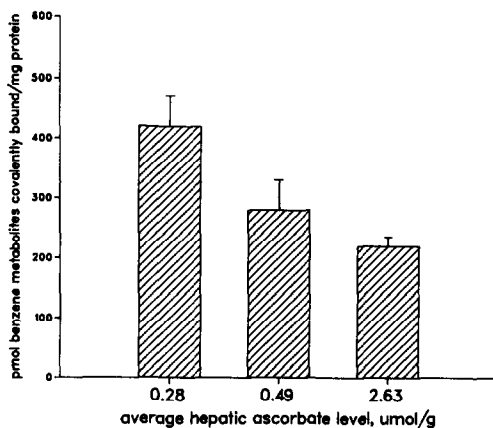


Fig. 1. Covalent binding of the metabolites of benzene in the liver after i.p. administration of [14 C]benzene. [14 C]Benzene (660 mg/kg) was administered 12 and 6 hr before termination to three groups of guinea pigs on different dietary intakes of ascorbate. Ascorbate and binding values were determined for each guinea pig, and the values are expressed as the mean \pm standard deviation. $P < 0.01$ with respect to 0.28 μ mole/g ascorbate to 2.63.

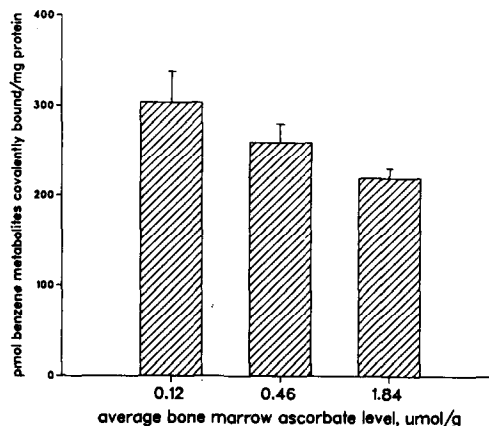


Fig. 2. Covalent binding of the metabolites of benzene in the bone marrow after i.p. administration of [14 C]benzene. [14 C]Benzene (660 mg/kg) was administered 12 and 6 hr before termination to three groups of guinea pigs on different dietary intakes of ascorbate. Ascorbate and binding values were determined for each guinea pig and the values are expressed as the mean \pm standard deviation. $P < 0.01$ with respect to 0.12 μ mole/g ascorbate to 1.84.

Table 1. Ascorbate concentration in liver and bone marrow of guinea pigs on the three dietary regimens of ascorbate

| Dietary regimens | Ascorbate tissue concentration (μ moles/g) | |
|-------------------------------------|---|-----------------|
| | Liver | Bone marrow |
| 2.0 mg Ascorbate/ml drinking water | 2.63 \pm 0.49 | 1.84 \pm 0.24 |
| 0.35 mg Ascorbate/ml drinking water | 0.43 \pm 0.11 | 0.46 \pm 0.17 |
| 0.05 mg Ascorbate/ml drinking water | 0.28 \pm 0.06 | 0.12 \pm 0.02 |

Values are expressed as the mean \pm standard deviation of four guinea pigs/dietary intake.

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Endocytosis of proteins by kidney tubule cells: inhibition by the aminoglycoside gentamicin

Nephrotoxicity is a major limitation to the long term use of aminoglycoside antibiotics such as gentamicin. This nephrotoxicity is associated with damage to both kidney glomerulus and tubule and is dependent on such factors as dose, duration of exposure, age and sex [1]. Early renal dysfunction has been related to the accumulation of aminoglycosides in the lysosomes of proximal tubule cells [2, 3]. Gentamicin also accumulates in lysosomes of other cell types such as fibroblasts [4] as has been shown for other weak bases [5].

Proximal tubule cells in the kidney are responsible for the reabsorption and degradation of protein remaining after ultrafiltration in the glomerulus [6]. It has been suggested [7] that an initial event in normal tubular protein reabsorption is an interaction between the protein and an anionic binding site at the proximal tubule brush border. Additionally, competition for such sites by polycationic aminoglycosides appears to take place [8]. Thus the effect of gentamicin on the renal handling of proteins is of interest.

We have recently used a kidney cell line—Madin Darby Canine Kidney (MDCK) cells—to study the selectivity in adsorptive endocytosis of different size proteins by tubular cells [9]. The present study utilises this cell line to investigate the effects of gentamicin on the uptake and degradation of rat plasma proteins.

Materials and methods

MDCK cells were maintained in monolayer culture in Eagles minimum essential medium (MEM) (Flow Laboratories) supplemented with 10% heat inactivated foetal calf serum and antibiotics. Rat serum (Miles Laboratories, Bucks, U.K.) was inactivated by heating at 56° for 30 min and then separated into three molecular size fractions by gel filtration on a Sepharose 6B column (Pharmacia, Sweden). Gel electrophoresis on 7.5% non-denaturing gel in Tris-glycine pH 8.9 showed the largest molecular size fraction to contain predominantly macroglobulins. The intermediate size fraction comprised gammaglobulins with

some albumin and the smallest molecular size fraction contained albumin and the smaller proteins. Aliquots of concentrated fractions were iodinated using the iodogen method [10] and the reaction products dialysed against phosphate buffered saline (PBS) after which less than 2% of the radioactivity was soluble in 5% trichloroacetic acid (TCA). Competition studies (data not show) indicated that the iodinated protein was recognised in a similar manner to the unlabelled protein as would be expected with the mild iodination procedure employed. All reagents were of the best available commercial grade.

MDCK cells were seeded at low density and allowed to grow to confluence over a period of 5 days in the presence of gentamicin (Flow Laboratories) in the concentration range 250–1000 µg/ml of culture medium. For the measurement of endocytosis, [³H]-sucrose (Amersham International) or ¹²⁵I-labelled proteins were then presented to the resultant confluent cell monolayers in fresh serum supplemented culture medium containing gentamicin such that gentamicin was present throughout the experimental period at the same concentrations as used in the preincubations. At the commencement of uptake the confluent monolayer consisted of viable cells and viability was maintained throughout the experiments as judged by morphology and retention of cell protein in the monolayer.

In order to measure uptake and degradation at chosen times after addition of the substrate, the media were harvested and the cells washed four times with ice-cold PBS to remove extracellular radioactivity. The cells were then lysed by the addition of 0.1% Triton X-100 in PBS. To measure uptake and degradation, media and cell fractions were precipitated with 5% TCA and TCA soluble and insoluble fractions counted separately by a gamma counter (Ultragamma 1280, LKB) or, for [³H]-sucrose uptake, aliquots counted directly in a liquid scintillation counter (Packard Tricarb 460C). An aliquot of each cell sample was assayed for protein by the method of Lowry *et al.* [11] using bovine serum albumin as standard. Uptake was