

# Carbamylation –induced inactivation of glyceraldehydes 3 –phosphate dehydrogenase and thioltransferase in bovine lens

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## Abstract

• **AIM:** To investigate whether potassium cyanate can inactivate glyceraldehydes 3-phosphate dehydrogenase (GAPDH) and thioltransferase (TTase) in bovine lens.

• **METHODS:** Fresh intact bovine lenses were incubated with 100mmol/L potassium cyanate (KCNO) for 7 and 12 days respectively. Then all lens were incubated in 50mmol/L DMEM solution. The proteins in the water-soluble fractions from the normal control and the cyanate-modified lens were extracted. The activity of GAPDH and TTase in the water-soluble fraction after incubation at 37°C was measured by spectrophotometer.

• **RESULTS:** GAPDH activity was significantly lower in the cyanate-modified lens proteins than that of the normal control ( $P < 0.01$ ), and considerably diminished in protein incubated with 100mmol/L potassium cyanate for 12 days. There were statistically significant differences in the activity of TTase between the normal control lenses and the carbamylated lenses incubated for 7 days ( $P < 0.05$ ) and 12 days ( $P < 0.01$ ). However, there was no statistical difference between the samples incubated with 100mmol/L KCNO for 7 and 12 days ( $P = 0.19296$ ).

• **CONCLUSION:** This study provides evidence to show carbamylation is able to inactivate GAPDH and TTase in bovine lenses. This may have implications for the susceptibility of lenticular GAPDH and TTase to carbamylation, and also for

the research on pathogenesis of cataract.

• **KEYWORDS:** carbamylation; cataract; potassium cyanate; glyceraldehydes 3-phosphate dehydrogenase; thioltransferase

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## INTRODUCTION

Post-translational modifications, known to occur during aging, such as glycation, carbamylation, oxidation, deamidation, phosphorylation and truncation, can cause conformational changes and thus cataract [1]. Carbamylation, an important post-translational modification of proteins occurring *in vivo*, may contribute to cataract formation in populations with high levels of blood urea. Urea in blood and other body fluids, including aqueous humour is always in equilibrium with isocyanic acid, the reactive species of carbamylation. Cyanate reacts with proteins at physiological pH by reversible binding to thiol groups, or by covalent binding to amino groups [2]. Binding of cyanate to lysyl residues, by eliminating positive charges, would alter protein-protein or protein-water interactions, and destabilize the protein. Such changes of protein surface charge produced by carbamylation lead to opacification of the lens [3]. Our earlier studies showed that the primary targeted lens proteins of carbamylation were  $\alpha$ A-,  $\beta$ B2- and  $\gamma$ S-crystallin. It indicates that these lens proteins are prone to be exposed to isocyanate (data were not published). The importance of carbamylation of lens crystallins, its effect on protein conformation and aggregation, and the potential to induce cataract have been discussed extensively [4-8], but little is known about the inactivation of enzymes induced by carbamylation.

Glyceraldehyde 3-phosphate dehydrogenase (GAPDH), a

key glycolytic enzyme, oxidizes and subsequently phosphorylates glyceraldehydes 3-phosphate in the glycolytic pathway. Its activity decreased in clear human lenses with aging and in cataract lenses<sup>[9,10]</sup>. *In vitro* incubations of GAPDH with sugar, cyanate and prednisolone-21-hemisuccinate, will lead to significant loss of enzyme activity<sup>[11]</sup>.

Thioltransferase (TTase), which is a member of the oxidoreductase family, regulates thiol/disulfide homeostasis<sup>[12]</sup>. Its active site sequence (Cys-Pro-Tyr-Cys) is responsible for this activity<sup>[13]</sup>. TTase has been postulated to repair oxidant-sensitive sulfhydryls by facilitating thiol-disulfide interchange between glutathione (GSH) and protein thiol mixed disulfides<sup>[12,14,15]</sup>. It is likely that this mechanism keeps the lens crystallins and membrane proteins in the reduced form by preventing their cross-linking or inactivation, and initial event in cataractogenesis.

In this communication, this is the first evidence to show carbamylation is able to inactivate GAPDH and TTase in bovine lenses. It may have implications for the susceptibility of lenticular GAPDH and TTase to carbamylation. The relationship between carbamylation and cataract pathogenesis is explored.

### MATERIALS AND METHODS

**Materials** Six 2-year-old bovine lenses were obtained from a local abattoir (Oxford, UK). Potassium cyanate and all other chemicals and enzymes were obtained from Sigma-Aldrich (Poole, Dorset, UK), and were of the highest purity available.

### Methods

**Incubation of bovine lens with potassium cyanate** The intact bovine lenses were dissected, and then four clear bovine lenses were incubated with 100mmol/L potassium cyanate (KCNO) for 7 and 12 days respectively in a 37°C shaking waterbath<sup>[4,16]</sup>. All lens incubations were carried out in 50mmol/L DMEM with 15mmol/L Hepes, D-199 and added antibiotics (penicillin, streptomycin and amphotericin B, Sigma). At the end of the incubation, each intact bovine lens was decapsulated and gently homogenized in a hand-held homogenizer over ice with 7 times the volume of double purified water (containing 13mmol/L EDTA) of its wet weight, to extract the water-soluble fraction of the lens proteins. And then it was centrifuged at 22 440g for 40 minutes at 4°C. The supernatant containing the water-soluble

proteins was dialyzed against distilled water at 4°C for 24 hours and freeze-dried, then stored at -20°C before enzyme determination and used for all the experiments described.

**GAPDH activity assay** The activity of glyceraldehyde 3-phosphate dehydrogenase (GAPDH) was determined spectrophotometrically by measuring the decrease in absorbance at 340nm against water within 1 minute at 30°C. The reaction mixture (3.0mL) consisted of 2.258mL of 0.1mol/L triethanolamine buffer with pH adjusted to 7.6, using 1mol/L NaOH, 200µL of 50g/L solution of glycerate-3-p, 100µL of 20g/L ATP, 100µL of 10g/L EDTA solution, 50µL of 0.1mol/L magnesium sulphate, 50µL of 10g/L NADH, 12µL of 3-phosphoglyceric phosphokinase suspension containing 7 units and finally 230µL of the incubation sample (containing 3mg lens protein). All of these reagents were pipetted into a 3mL quartz cuvette in a Hitachi U-2001 spectrophotometer with 1cm light path, and then the absorbance values were recorded. The enzyme activity was calculated from the linear decrease in absorbance between time 0 and 1 minute. By definition, one international unit GAPDH reduces 1.0µmol of 3-phosphoglycerate to D-glyceraldehyde 3-phosphate in a coupled system with 3-phosphoglyceric phosphokinase at pH 7.6 in 1 minute at 30°C.

**TTase activity assay** TTase was assayed following the method of Mieyal *et al*<sup>[17]</sup>. Reaction mixtures without the cell homogenate were used as blanks in enzyme assay. The reaction was carried out in the presence of NADPH, GSH and glutathione reductase (GR) with the synthetic disulfide, hydroxyl ethyl disulfide (HEDS), as substrate. The decreased A340nm for NADPH was monitored for enzyme activity. The enzyme activity was measured in an order by which one sample from each group (controls, incubated for 7 days, and incubated for 12 days) was measured in succession, to ensure that all the samples were treated similarly. The change in absorbance was corrected for any change in parallel assays without supernatant.

**Statistical Analysis** All experiments were repeated at least three separate times and differences were assessed by student's paired *t*-test where relevant. All comparisons were made with control solution unless stated otherwise.

### RESULTS

**Activity of GAPDH from Normal Control Lenses and Carbamylated Lenses** The activity of GAPDH in clear

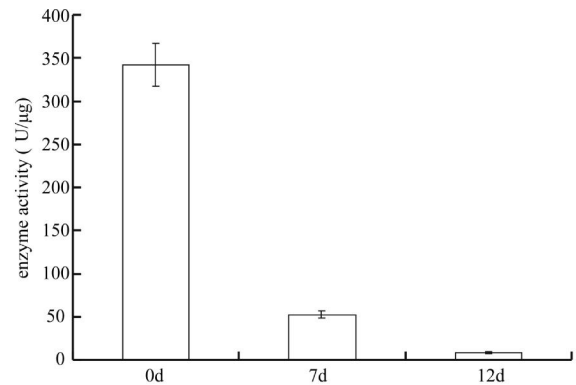
lenses was tested and compared with samples containing clear lenses incubated with 100mmol/L KCNO for 7 and 12 days respectively. Potassium cyanate at concentration of 100mmol/L inactivated GAPDH in a time-dependent manner (Figure 1). GAPDH activity in control group was very high,  $341.6 \pm 24.7 \text{ U}/\mu\text{g}$  lens protein in samples containing clear lens supernatant incubated without KCNO. All the carbamylated groups offered a statistically significant decrease in activity. The significant decrease in activity ( $52.4 \pm 4.4 \text{ U}/\mu\text{g}$  lens protein) was achieved in the group incubated with potassium cyanate for 7 days ( $P < 0.01$ ), whereas the greatest decrease ( $8.5 \pm 1.5 \text{ U}/\mu\text{g}$  lens protein) was achieved in the group incubated for 12 days ( $P < 0.01$ ) compared with the normal control group. All of the results were highly statistically significant.

**Activity of TTase from Normal Control Lenses and Carbamylated Lenses** Potassium cyanate (100mmol/L) inactivated TTase after an incubation period of 7 days and 12 days. The activity of TTase was greater in the normal control lenses than that in the carbamylated lenses. However, there was no statistical difference between the samples incubated with 100mmol/L KCNO for 7 and 12 days ( $P = 0.19296$ ). After an incubation period of 7 days, the activity of TTase decreased to  $32.1 \pm 2.1 \text{ U}/\mu\text{g}$  lens protein compared with  $38.7 \pm 2.7$  in the normal control lenses ( $P < 0.05$ ). The activity of TTase lowered to  $30.1 \pm 0.4 \text{ U}/\mu\text{g}$  lens protein after an incubation of 12 days ( $P < 0.01$ ) compared with the normal control group (Figure 2).

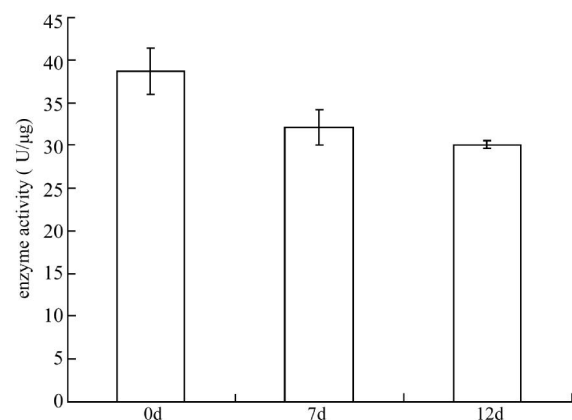
## DISCUSSION

Our data showed that the activity of GAPDH in carbamylated supernatant was much lower than that in clear lens supernatant. The activity in lenses obtained by incubation with 100mmol/L potassium cyanate for 12 days was lower than that of the lenses supernatant incubated with same concentration of potassium cyanate for 7 days, which was found to be time-dependent.

The role of GAPDH in many biochemical processes has been studied extensively. GAPDH activity decreases in experimentally induced cataracts and in human diabetic cataract [3]. This is paralleled by increased glycation of the lens proteins of diabetics [18] or modification of glucocorticoids in the case of patients who had undergone glucocorticoid therapy [19]. Earlier studies reported that GAPDH activity



**Figure 1** GAPDH activity was measured in bovine lenses in the absence of, or presence of 100mmol/L potassium cyanate for 7 days or 12 days



**Figure 2** TTase activity was measured in bovine lenses in the absence of, or presence of 100mmol/L potassium cyanate for 7 days or 12 days

was significantly decreased in pooled lenses from cataract groups compared to clear lenses of similar average age [20]. Carbamylation of essential lysine residues can inhibit the normal function of proteins. GAPDH contains a highly conserved and reactive lysine near its active site [21]. Reaction of this residue with a small molecule could lead directly to its inactivation.

TTase activity tested in this study was only slightly reduced when clear lenses were exposed to 100mmol/L potassium cyanate for 7 days, while its activity was significantly affected by potassium cyanate after incubation for 12 days.

The physiological function of TTase has been speculated to be a repair enzyme for oxidation damaged proteins/enzymes through its dethiolation capability. The lens depends on a balanced redox state and SH/S-S- ratio for maintaining its transparency. TTase can specifically dethiolate protein-S-S-glutathione and restore protein free SH groups for proper enzymatic or protein functions and maintain the function

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of the lens effectively [22]. This is the first evidence to show carbamylation is able to inactivate TTase in bovine lenses.

A unifying mechanism by which environmental cataracts may develop could be accelerated by post-translational modification of lenticular proteins such as carbamylation. It is possible that inactivation of these enzymes was due to cumulative post-translational modification which induces conformational changes of proteins and affects the active sites, resulting in the inactivation of enzymes.

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