# **Effects of dietary protein source on plasma lipids, HMG CoA reductase activity, and hepatic glutathione levels in gerbils**

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*It has recently been postulated that soy protein-induced hypocholesterolemia is induced by increased production of reduced glutathione (GSH). The objective of the current study was to determine the influences of dietary proteins and amino acids on plasma lipid concentrations, hepatic GSH, and HMG CoA reductase activity. Twenty-eight adult male, inbred gerbils were fed diets similar in all respects except that dietary protein source was from either soy protein isolate (SOY), casein (CAS), L-amino acid patterned after SOY (SAA), or L-amino acid patterned after CAS (CAA). Results indicated that gerbils fed SOY had lower plasma total cholesterol concentrations than gerbils fed CAS or CAA. Low density lipoproteincholesterol was also lowest in SOY-fed animals compared with all other groups. High density lipoprotein (HDL)-cholesterol concentrations were highest in CAA-fed animals. Plasma triglycerides were unaffected by dietary treatment. Hepatic GSH concentrations were lowest in SOY-fed gerbils, but were not different between SAA- and CAA-fed groups. HMG CoA reductase activity was highest in gerbils fed SO Y compared with CAS; however, when amino acids were fed the opposite occurred-reductase activity was higher in gerbils fed CAA compared with animals fed SAA. Correlation analysis showed no association between GSH and HMG CoA reductase activity. Positive correlations were present between GSH and total and HDL-cholesterol concentrations. These results indicate that lipid metabolism is influenced by dietary protein source, but these changes are not likely to be mediated by a GHS-mediated modification of cholesterol biosynthesis.* 

**Keywords:** soy protein; casein; cholesterol; GHS; HMG CoA reductase

## **Introduction**

The hypocholesterolemic effect of soy protein compared with animal protein has been demonstrated in humans and a variety of animal models.<sup>1-3</sup> In rabbits, feeding L-amino acids patterned after casein produces hypercholesterolemia similar to that produced by feeding intact casein. Feeding L-amino acids patterned after

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soy protein isolate produces hypocholesterolemia; however, values were not as low as those produced by feeding the intact protein. 4,5 This indicates that the cholesterolemic changes induced by dietary proteins are **partially** due to their amino acid composition.

One of the main differences in amino acid composition between casein and soy protein is the proportionality of sulfur-containing amino acids. The ratio of cysteine to methionine in casein is 0.34:2.9, while in soy protein isolate, it is 1.3:1.3. It has been postulated that the higher cysteine content of soy protein may contribute to the resultant hypocholesterolemia by enhancing glutathione production.<sup>6</sup> There are indications that reduced glutathione (GSH) is a cofactor in cholesterol metabolism. 7.8 In addition, we have reported that adding cysteine (0.33%) to soy protein isolate lowers serum cholesterol levels, while adding methionine (0.33%) in-

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creases levels in rats. 9 The current investigation was designed to examine the effects of dietary proteins and amino acids on plasma lipid concentrations, hepatic GSH concentration, and HMG CoA reductase activity; and to examine if there are associations between hepatic GSH content and activity of HMG CoA reductase.

## **Methods and materials**

## *Experimental design*

Gerbils were chosen as the animal model for this study due to reported lipemic responses to dietary fat and cholesterol.<sup>10,11</sup> Male, inbred gerbils *(Meriones unguiculates,* strain designation Mon/Tum; Tumblebrook Farms, Brent Lake, NY USA) were fed diets similar in all respects with the exception of dietary protein source coming from soy protein isolate (SOY), casein (CAS), L-amino acid patterned after SOY (SAA), or L-amino acids patterned after CAS (CAA).

## *General procedures*

Gerbils weighing approximately 60 g each were individually housed in an environmentally controlled room  $(23^{\circ} \text{ C})$  with an alternating 12-hr light/12-hr dark cycle. Upon arrival, all animals were fed a pulverized commercial pelleted diet (Purina Rat Chow, Ralston Purina, St. Louis, MO USA) for 1 week. Animals were then randomly assigned to dietary treatment groups (eight animals/group). Composition of basal diet is provided in *Table 1.* Amino acid composition of soy protein

**Table 1 Basal diet** composition

Component	% by weight
Protein <sup>a</sup>	18.00
Dextrose	53.90
Lard	16.00
Safflower oil	1.00
Cellulose	5.00
Vitamin mix <sup>b</sup>	1.00
Mineral mix <sup>c</sup>	5.00
Cholesterol	0.10

aSoy protein isolate, PP610 (91.5% protein based on manufacturer's analysis, calculated as gNx6.25 (30); Protein Technologies Int., St. Louis, MO USA); Casein, vitamin-free test (91% protein based on manufacturer's analysis, calculated as gNx6.38 (30); Teklad, Madison, WI USA); L-amino acid mixture patterned after soy protein isolate PP610 based on manufacturer's analysis *(Table 2);* L-amino acid mixture patterned after vitamin-free test casein based on manufacturer's analysis *(Table 2).* 

bTeklad vitamin mix. Vitamin mix supplied the following vitamins in mg/kg of diet: vitamin B-12 (0.1% trituration in mannitol), 30; calcium pantothenate, 66; dihydrogen citrate, 3500; folic acid, 2; inositol, 110; menadione (vitamin K), 50; niacin, 99; pyridoxine HCI, 22; riboflavin, 22; thiamin HCI, 22; dry retinyl palmitate (500,000 IU/g), 40; dry ergocalciferol (500,000 IU/g), 4.4; dry tocopherol acetate (500 IU/g), 2423.

cMineral mix, Hegsted IV (Teklad). Mineral mix supplied the following minerals in g/kg of diet: calcium carbonate, 15.0; potassium phosphate, diabasic, 16.1 ; calcium phosphate, diabasic, 3.7; magnesium sulfate, 5.1; sodium chloride, 8.4; ferric citrate (USP 16.7% Fe), 1.4; potassium iodide, 0.034; manganese sulfate, 0.19; zinc chloride, 0.012; cupric sulfate, 0.015; nonnutritive fiber, 0.061.

isolate and casein is provided in *Table 2.* Feed and water were provided ad libitum, and feed intake and weight gain were monitored throughout the project.

After a 28-day experimental period, animals were sacrificed via decapitation starting 2 hours after initiation of the dark cycle and completed 4 hours after initiation to allow for peak HMG CoA reductase activity.<sup>12</sup> Blood was collected into heparinized tubes and plasma was separated. High density lipoprotein (HDL) and low density lipoprotein  $+$  very low density lipoprotein (LDL + VLDL) (LDL) lipoprotein fractions were separated, and plasma and fractions were frozen at  $-20^{\circ}$  C for subsequent analysis. Portions of livers were prepared for GSH analysis, while microsomes were isolated from the remainder of the livers and stored in liquid nitrogen until microsomal HMG CoA reductase activity was assayed.

#### *Blood lipid analysis*

HDL and LDL lipoprotein fractions were separated by hepafin-bound agarose columns (Liposep, Isolabs, Inc., Akron, OH USA). Total, HDL, and LDL cholesterol were measured enzymatically by a modification of the method described by Allain et al.<sup>13</sup> using a commercially available reagent (Sigma) Diagnostics, St. Louis, MO USA). Total triglyceride concentrations were measured enzymatically, based on the method described by McGowen et al.<sup>14</sup>, also using a commercially available reagent (Sigma Diagnostics).

## *Hepatic GSH concentration*

Livers were removed from animals and minced. From this, 0.2 g samples were taken and homogenized in 800  $\mu$ L of icecold  $1.15\%$  KCl. Aliquots of the homogenate were deproteinized with 10% sulfosalicylic acid, and reduced GSH was analyzed according to methods described by Asaoka and Takahashi. 15 Briefly, conjugated GSH and 0-dinitrobenzene, upon being acted on by glutathione S-aryltransferase, release nitrite ions that diazo-couple with sulfanilamide and N-(1-

Table 2 Composition of amino acid mixtures<sup>a</sup>

	<b>SAA</b>	CAA∘
∟-Alanine	4.3	2.8
L-Arginine HCI	7.6	3.4
L-Aspartate	11.6	6.3
L-Cysteine <sup>d</sup>	1.3	0.3
∟-Methionine	1.3	2.9
<b>Total Sulfur Amino Acids</b>	2.6	3.2
L-Glutamate	19.1	20.5
Glycine	4.2	1.6
L-Histidine HCI H <sub>2</sub> O	2.6	2.5
L-Isoleucine	4.9	4.7
L-Leucine	8.2	8.2
L-Lysine HCI	6.3	7.2
L-Phenylalanine	5.2	9.5
L-Serine	5.2	5.0
L-Threonine	3.8	3.8
∟-Tryptophan	1.3	1.6
L-Tyrosine	3.8	4.7
L-Valine	5.0	6.0

ag/loo g protein.

**bBased on amino acid composition of soy protein isolate, PP610** (Protein Technologies, Int., St. Louis, MO USA).

cBased on amino acid composition of vitamin-free test casein (Teklad, Madison, WI USA).

naphthyl) ethylene diamine dihydrochloride. The resulting diazo product was then measured at 540 nm.

## *HMG CoA reductase activity*

Microsomes were prepared from the remaining liver as described by Shapiro et al.<sup>16</sup> HMG CoA reductase activity was assessed using  $^{14}$ C HMG CoA as a substrate and  $^{3}$ H mevalonate (New England Nuclear, Boston, MA USA) as an internal standard.<sup>17</sup> The reaction was carried out at  $37^{\circ}$  C for 20 minutes, stopped by the addition of 10 N HC1, and allowed to lactonize for 30 min at  $37^{\circ}$  C. <sup>14</sup>C-mevalonate produced was separated by thin-layer chromatography (Silica Gel G, Alltech, Deerfield, IL USA), counted, and quantified. Microsomal protein was quantified using the bicinchoninic acid (BCA) copper reagent (Pierce, Rockford, IL USA) method for use in the presence of sulfhydryl reagents.<sup>18</sup>

## *Statistical analysis*

Data were analyzed using the Statistical Analysis System (SAS, Inc. Cary, NC USA) by a one-way analysis of variance. Differences between treatments were determined using contrast analysis, and an alpha level of 0.05 was used to indicate statistical significance. Pearson's correlation coefficients between hepatic GSH and HMG CoA reductase activity, plasma lipids, feed intake, and weight gain were also computed. 19

# **Results**

## *Feed intake and weight gain*

Mean weight gain and feed intake are summarized in *Table 3.* Weight gain was unaffected by dietary treatment; however, feed intake was lower in CAA-fed animals compared with those fed CAS ( $P < 0.05$ ).

# *Plasma lipid concentrations*

Mean plasma lipid concentrations are summarized in *Table 4.* Plasma total cholesterol concentrations were

Table 3 Weight gain and feed intake (g/28 days)<sup>a</sup>

Treatment	Weight gain	Feed intake	
SOY	$9.9 \pm 2.2$	$159 \pm 16^{\circ}$	8
CAS	$8.6 \pm 2.3$	$174 \pm 43^{\circ}$	8
<b>SAA</b>	$8.7 \pm 1.3$	$153 + 14$ <sup>a</sup>	
CAA	$8.5 \pm 2.1$	$141 \pm 10^{6}$	

Values represent mean  $\pm$  SD. Means in a column with a different superscript are significantly different at  $P < 0.05$ .



lower in gerbils fed SOY compared with those fed CAS or CAA  $(P < 0.05)$ , or SAA  $(P = 0.06)$ . LDL-cholesterol concentrations were lower in the SOY-fed group compared with all other groups ( $P < 0.05$ ). When the L-amino acid mixtures simulating SOY and CAS were fed, no differences in plasma lipid concentrations were observed, with the exception of CAA-fed gerbils having highest HDL-cholesterol values ( $P < 0.05$ ). Total triglycerides were unaffected.

# *Hepatic GSH and HMG CoA reductase activity*

Mean hepatic GSH concentrations and HMG CoA reductase activities are summarized in *Table 5.* Concentrations of hepatic GSH were lowest in animals fed SOY compared with all other groups ( $P < 0.05$ ). When Lamino acid mixtures were fed, there were no differences in hepatic GSH concentrations.

With regard to HMG CoA reductase, animals fed SOY exhibited highest activities compared with those fed CAS and SAA  $(P < 0.01)$ . When L-amino acid mixtures were fed, reductase activity was higher in the CAA group compared to the SAA group ( $P < 0.05$ ).

## *Correlations*

Significant correlations were found between GSH and plasma total  $(R = 0.44, P = 0.03)$  and HDL-cholesterol concentrations  $(R = 0.42, P = 0.03)$ . There were no statistically significant correlations between GSH and HMG CoA reductase activity, feed intake, or weight gain.

## **Discussion**

Data from the present study indicate that variations in dietary protein source influence lipid metabolism in

Table 5 Hepatic GSH concentrations and HMG CoA reductase activities<sup>1</sup>

GSH <sup>2</sup>		HMG CoA <sup>3</sup>	
SOY	$5.7 \pm 0.8^{\circ}$	$147.8 \pm 17.2$ <sup>a</sup>	
CAS	$7.4 \pm 1.5^{\circ}$	$81.2 + 26.1$	
<b>SAA</b>	$7.7 \pm 1.9^{\circ}$	$83.8 \pm 25.3$ <sup>b</sup>	
CAA	$7.2 \pm 0.8^{\circ}$	$118.3 \pm 32.3^{\circ}$	

 $v$ alues represent mean  $\pm$  SD. Means in a column with a different superscript are significantly different at  $P < 0.05$ .  $2$ SOY n = 6; CAS n = 7; SAA n = 7; CAA n = 7.

 $3$ SOY n = 4; CAS n = 6; SAA n = 6; CAA n = 7.



Values represent mean \_+ SD. Means **in a** column with a different superscript are significantly different at **P < 0.05.**  \*SOY versus SAA ( $P = 0.06$ ).

gerbils. Consumption of soy protein isolate resulted in significant decreases in plasma total and LDL-cholesterol concentrations in comparison with values for other treatment groups. When L-amino acids patterned after SOY or CAS were fed, neither plasma total nor LDL-cholesterol concentrations were demonstrated to be significantly different, although total cholesterol values were numerically lower in the *SAA*fed animals in comparison with those fed CAA. HDLcholesterol, however, was found to be highest in CAA-fed animals, which may represent a beneficial partitioning of lipoproteins even though total cholesterol was numerically highest in this group. Both total and LDL-cholesterol values were higher in gerbils fed the L-amino acids patterned after soy than when they were fed soy protein isolate. This indicates that the hypocholesterolemic effect of soy protein in gerbils may be mediated by a nonprotein constituent, possibly a phytochemical.

The mechanism for soy protein-induced hypocholesterolemia is not known; however, it has been postulated that GSH concentrations could be increased by feeding adequate amounts of soy protein due to the higher content of cysteine.<sup>6</sup> Hepatic GSH level is closely dependent on diet, especially the amount of ingested cysteine, because this amino acid is rate limiting in GSH biosynthesis.<sup>20</sup> In addition to HMG CoA reductase, cholesterol 7- $\alpha$ -hydroxylase,<sup>21</sup> the ratelimiting enzyme in bile acid synthesis, $22$  has been reported to be influenced by GSH.

GSH is also believed to be a cofactor in conversion of thyroxine (tetraiodothyronine,  $T<sub>4</sub>$ ) to triiodothyronine  $(T_3,$  the active form of thyroid hormones).<sup>23</sup> In vitro data indicate that the release of  $T_3$  and  $T_4$  is dependent on thyroid GSH breaking disulfide bonds of thyroglobulin.<sup>24</sup> Forsythe<sup>10</sup> reported increases in  $T<sub>4</sub>$  concentrations in gerbils fed isolated soy protein compared with casein, and we have noted an increase in plasma  $T<sub>4</sub>$  (without concurrent increases in plasma thyroid-stimulating hormone) in humans consuming isolated soy protein compared to a basal mixed protein diet.<sup>25</sup> Furthermore, hepatic GSH concentrations in hyperthyroid rats have been reported to be lower and plasma GSH higher than in euthyroid rats.<sup>26</sup> This may indicate that hepatic GSH is being depleted by increases in extrahepatic transport, possibly to the thyroid, facilitating thyroid hormone release. The actions of thyroid hormones on cholesterol metabolism are to increase biosynthesis, bile acid synthesis, and LDL receptor activity.<sup>24</sup> These same responses in cholesterol homeostasis are seen upon consumption of soy protein.<sup>27</sup>

Our findings do not support the hypothesis that feeding soy protein increases hepatic GSH concentrations, thereby increasing activity of cholesterol metabolic enzymes. In fact, hepatic GSH concentrations were lowest while HMG CoA reductase activities were highest, and correlation analysis failed to show a significant association between these two parameters. Perhaps concentrations of GSH in the thyroid and periphery are more reflective of the possible role of

GSH in the hypocholesterolemic effect of soy protein than hepatic concentrations.

Others have reported increased growth rates and hepatic total glutathione in rats fed diets containing  $16\%$  protein from casein versus soy protein isolate.<sup>28</sup> However, when level of protein intake increased to 24%, growth rates and GSH concentrations were similar. In the current investigation, older animals were used, producing minimal growth rates that were similar between treatment groups. It is not clear why gerbils fed only the SOY diet and not the SAA diet had depressed hepatic GSH concentrations if sulfuramino acid intake alone is the determining factor for hepatic GSH concentrations. The reason for this divergence in response may be a result of differential absorption of amino acids between the intact proteins and their amino acid counterparts or, possibly, the action of an unidentified phytochemical present in soy protein isolate that either inhibits protein absorption or affects synthesis of GSH directly.

Increased HMG CoA reductase activity in soy protein-fed rats has been reported by Nagata et a1.,29 but when amino acids replaced intact proteins, higher reductase activities were observed in rats fed amino acids patterned after casein. In our study, when the amino acid mixtures simulating SOY (SAA) or CAS (CAA) were fed reductase activity was also higher in CAA-fed gerbils compared with those fed SAA. Nagata et al. 29 propose that this variation in response between the intact and simulated proteins is a function of diurnal variation. However, in our experiments animals were sacrificed randomly, at specific time points corresponding to peak reductase activity. It is possible that slight differences in time points and/or stress involved in sacrificing could have caused the observed differences between gerbils fed the intact versus simulated proteins. This does not explain why both the Nagata group and our group observed higher HMG CoA reductase activities with intact soy protein in comparison with casein and lower HMG CoA reductase activities with SAA in comparison with CAA. These observations again may be more reflective of differences in absorption of the amino acids or the action of a nonprotein constituent of soy protein isolate resulting in variations in HMG CoA reductase activity.

In conclusion, data from the present study show that dietary protein source influenced lipid metabolism in gerbils. Our data also indicate that hepatic GSH concentration and HMG CoA reductase are influenced by dietary protein source. It is unlikely that changes in plasma lipids associated with feeding soy protein are a result of changes in hepatic GSH concentrations causing an effect on activity of HMG CoA reductase. However, it is clear that feeding intact soy protein isolate produces lower hepatic GSH concentrations and hypocholesterolemia-responses that are not present when L-amino acids patterned after soy protein are fed. Further studies looking into GSH concentrations in the plasma, thyroid, and possibly skeletal muscle along with thyroid hormone activation and release warrant further investigation.

#### *Research Communications*

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