Vitamin A Deficiency Increases Protein Catabolism and Induces Urea Cycle Enzymes in Rats

Guillermo Esteban-Pretel, M. Pilar Marín, Francisco Cabezuelo, Verónica Moreno, Jaime Renau-Piqueras, Joaquín Timoneda, and Teresa Barber

Introduction

Vitamin A (all-trans retinol) is an essential nutrient involved in many physiological functions, including vision, immunity, cell differentiation, embryological development and growth, and it also acts as a physiological antioxidant (1). Retinol is the precursor in the body for 2 especially important metabolites, retinaldehyde and retinoic acid (RA), particularly all-trans and 9-cis RA. RA exerts its broad range of biologic effects mainly by controlling gene expression. RA binds to and activates heterodimeric nuclear receptors, which comprise 2 subfamilies, RA receptors and retinoid X receptors. Moreover, retinoid X receptors form heterodimers with other members of the nuclear receptor family, including PPAR, and vitamin D and thyroid hormone receptors (2).

Abstract

Chronic vitamin A deficiency induces a substantial delay in the rates of weight and height gain in both humans and experimental animals. This effect has been associated with an impaired nutrient metabolism and loss of body protein. Therefore, we analyzed the effect of vitamin A deficiency on endogenous proteolysis and nitrogen metabolism and its reversibility with all-trans retinoic acid (RA). Male weanling rats, housed in pairs, were pair-fed a vitamin A-deficient (VAD) or control diet until they were 60 d old. A group of deficient rats were further treated with daily intraperitoneal injections of all-trans RA for 10 d. Final body and tissue (i.e., liver and heart) weights were significantly lower and tissue:body weight ratios were similar in VAD rats and in controls. Conversely, the epididymal white fat:body weight ratio and the plasma concentrations of alanine aminotransferase and adiponectin were significantly higher in VAD rats, which also had hepatic macrovesicular lipid accumulations. Plasma and gastrocnemius muscle 3-methylhistidine, urine nitrogen, and plasma and urine urea concentrations were all significantly higher in the VAD group. The expression of the genes encoding urea cycle enzymes and their activities increased in VAD livers. These changes were partially reverted by all-trans RA. We propose that fuel partitioning in vitamin A deficiency may shift from fatty acids to protein catabolism as an energy source. Our results emphasize the importance of vitamin A on the energy balance control system and they provide an explanation for the role of vitamin A in protein turnover, development, and growth. J. Nutr. 140: 792–798, 2010.

Vitamin A deficiency is still a nutritional problem in the world, with important implications for global health policies. This deficiency has a wide range of undesirable effects, among which are visual problems, increased susceptibility to severe infection, and disturbances in growth (3). Since its discovery, several studies have linked vitamin A deficiency with delayed rates of weight and height gain in both humans and experimental animals (3–7). However, the underlying mechanism involved in the relationship between vitamin A status and growth has not been convincingly stated. Former studies (8) have reported that vitamin A deficiency causes a loss of appetite with a subsequent reduction in food intake and that gastrointestinal disturbances associated with this deficiency could lead to a decreased absorption of nutrients (8,9). These features might influence growth rate after chronically withdrawing vitamin A from the diet. However, thorough in vivo studies of body composition and nutrient absorption in rats using the paired feeding method have concluded that these factors could not justify the weight loss and growth failure observed with vitamin A deficiency. The effect of vitamin A deficiency on body weight appears to be mostly due to alterations in nutrient utilization and metabolism after absorption; moreover, difference in weight was significantly due to protein loss in deficient rats (8).
Recently, in vivo studies have established a link between retinoid status and the expression of those genes encoding several enzymes of intermediary metabolism. The results suggest that vitamin A deficiency may induce a general catabolic state that could relate with resistance to growth hormone (GH) effects (10). However, despite the interest in understanding the metabolic effects of vitamin A deficiency, particularly those concerning delays in tissue growth rate and development, the effect of this deficiency on protein metabolism has not yet been convincingly stated.

Proteolysis along with protein synthesis is a major process that contributes to body protein turnover and tissue development, and a slight decrease in synthesis or a slight increase in degradation rates, if sustained, can result in a marked loss of mass in the organism as a whole. Experimental and clinical studies have pointed out that the most important feature in the muscle protein loss in several catabolic states, including sepsis, cancer, burn injury, metabolic acidosis, diabetes, or hyperthyroidism, is enhanced muscle protein degradation, especially of myofibrillar proteins (11). Under such conditions, muscle protein catabolism increases to provide the organism with amino acids for energy production, gluconeogenesis, or new protein synthesis (11), and in vivo studies have shown that increased amino acid utilization responds mostly to a decreased availability of nonprotein substrates as an energy supply (12). In this context, it is worth mentioning that retinoid status modulates fatty acid utilization in different tissues. In fact, chronic vitamin A deficiency decreases the expression of the genes involved in mitochondrial fatty acid oxidation in liver, whereas treatment with RA increases the lipid oxidation capacity in liver, skeletal muscle, and white adipose tissue (13–16). The adaptive response in vitamin A deficient (VAD) rats to a lower capacity of lipid oxidation may result in an increased utilization of proteins as an alternative energy source, leading to progressive protein loss and greater nitrogen excretion as urea, a nontoxic metabolite synthesized by the liver from amino acid catabolism.

In this work, we studied the effect of chronic vitamin A deficiency on protein catabolism and its reversibility by all-trans RA treatment. We evaluated growth, myofibrillar protein catabolism by determining 3-methylhistidine concentration in plasma and muscle, total nitrogen urinary excretion, urea production, and hepatic urea cycle enzyme activities and their mRNA. Moreover, because the liver is involved in most aspects of vitamin A homeostasis, expresses the RA-regulated transcription factors at high levels, plays a key role in lipid balance, and is a singular tissue in that it has a complete metabolic capability of detoxifying ammonia to urea, we also evaluated the histological ultrastructure of livers from control and VAD rats.

Materials and Methods

Animals and diets. All procedures using rats were carried out in accordance with the NIH Guide for the Care and Use of Laboratory Animals. The protocol was approved by the Ethics Committee for Animal Research and Welfare at the University of Valencia. VAD rats were randomly divided into 2 groups. The first was fed a complete purified diet (control 21-d lactation shortened the interval of vitamin A depletion in pups (19)). Feeding dams the vitamin A-free diet during the light-dark cycle. On the day after delivery, dams with their litters were taken and processed as previously described (21). Heart, kidney, lung, and epididymal-adipose tissue were then excised and weighed. For the ultrastructural studies, whole anesthetized rats were perfused with 0.5% glutaraldehyde and 4% formaldehyde in 0.1 mol/L cacodylate buffer (pH 7.4) (20). For urine collection, rats were placed in individual metabolic cages at the end of the experimental period for 24 h, as described previously (22).

Retinoids determination. Plasma retinol and liver retinoids were extracted, dissolved in methanol:ethanol (1:1, v:v) (20), and measured using an isotropic HPLC method (23).

Ultrastuctural studies. Small fragments of rat liver were postfixed in 1.5% glutaraldehyde and 1% formaldehyde in 0.1 mol/L cacodylate buffer (pH 7.4) for 60 min at 4°C, dehydrated in graded concentrations of ethanol, and embedded in Epon 812 (6 blocks per sample) (24). Ultrathin sections were examined in a Philips CM 100 transmission electron microscope.

Liver protein. Liver total protein was measured by the method of Lowry et al. (25).

Plasma biochemistry. Plasma glucose, triglycerides, cholesterol, and alanine aminotransferase activity were measured in 18- to 24-h food-deprived rats using enzymatic methods following the manufacturer’s instructions (Spinreact). Adiponectin was measured with a competitive ELISA from Linco Research according to the manufacturer’s manual.

Plasma and urine nitrogenous compounds. Creatinine, urea, and total N were measured in plasma and in 24-h urine samples, as described in Barber et al. (22).

Plasma and gastrocnemius muscle 3-methylhistidine. Methylhistidine from the plasma and muscle extracts was determined by the isotropic HPLC method of Nagasawa et al. (26). Muscle methylhistidine was extracted as described (27).

Urea cycle enzyme assays. Liver urea cycle enzyme activities, carbamoyl-phosphate synthetase I, ornithine carbamoyltransferase, argininosuccinate synthetase, argininosuccinate lyase, and arginase I were measured as described (28). One unit of enzyme activity is defined as the amount of enzyme that catalyzes the formation of 1 μmol of product/h at 37°C.

Real-time PCR. RNA extraction and real-time PCR were carried out as described previously (20). One microgram of total RNA was treated with DNase I (Roche) and reverse transcribed to cDNA using Ready-To-Go You Prime First-Strand beads (Amersham Pharmacia Biotech). Primers for PCR were designed by the Primer3 program and synthesized by Sigma. These were: carbamoyl-phosphate synthetase I, sense: 5′-TATGTTACCTACAATGGCCAGGAG-3′, antisense: 5′-GATACTGGAGCACGACACAGAATC-3′; ornithine carbamoyltransferase, sense: 5′-GACAGCACACCAATC-3;
RESULTS

Retinoid concentrations in plasma and liver. The rats fed a vitamin A-free diet from birth (through the dam’s milk) until reaching the age of 60 d had plasma retinol concentrations that were 5\% of the concentration in controls. In parallel, liver retinyl palmitate, the main retinoid found in this tissue, was 10\% of the control value (Table 1). The VAD rats treated with all-trans RA had almost negligible plasma retinol and liver retinyl palmitate concentrations (Table 1).

The results obtained with the vehicle-treated group did not significantly differ from the VAD group and, for simplicity, they are not shown.

Body and tissue weights. VAD rats grew more slowly than those fed the control diet and their body weight was lower at the end of the study (Fig. 1A). We detected growth retardation in VAD rats by the age of 56 d (wk 8) despite equivalent food consumption. Differences in body weight were significant by wk 9, at which time VAD rats had reached a weight plateau.

At the end of the experiments (d 60), VAD rats were smaller, but their liver weight was also significantly lower than control rats (Table 2); accordingly, the liver weight as a percentage of the total body weight (liver:body weight ratio) was similar in both groups (2.8\% of body weight). The liver protein content was also similar in both the control and VAD rats (Table 2). The absolute and relative heart weights of both groups followed the same pattern as that in the liver (heart weight: 0.89 ± 0.04 g in control rats, 0.75 ± 0.02 g in VAD rats, \( P < 0.05 \); heart:body weight ratio: 0.35\% in both the control and VAD groups); lung and kidney weights did not significantly differ between groups and, for simplicity, they are not shown. Interestingly, we found a greater accumulation of white fat, as the epididymal fat pad weight/body weight ratio was significantly higher in VAD rats than in control rats (Fig. 1B) despite the similar food intake of both groups.

Liver ultrastructural studies. Because the liver plays a key role in lipid homeostasis, we examined representative sections of the livers from both the control and VAD rats (Fig. 1C). Electron microscopic studies showed numerous lipid vacuoles in the hepatocytes of VAD rats, which were very scarce in control rats. In a previous study (29), we also noticed vacuole formation in VAD livers. Such vacuolization is often considered an index of steatosis, which has also been described by other authors in humans and rats with decreased hepatic vitamin A content (15,30).

Plasma biochemistry. The lower body and tissue weights and the greater fat accumulation in VAD rats (Table 2; Fig. 1B,C), despite similar food intake, suggested a shift in total energy
TABLE 2  Body and liver weights, liver protein concentration, and plasma biochemistry of rats fed control or VAD diets for 60 d1

<table>
<thead>
<tr>
<th></th>
<th>Control</th>
<th>VAD</th>
</tr>
</thead>
<tbody>
<tr>
<td>Final body weight, g</td>
<td>273.1 ± 6.8 (10)</td>
<td>212.6 ± 4.1 (10)*</td>
</tr>
<tr>
<td>Liver weight, g</td>
<td>8.2 ± 0.3 (10)</td>
<td>6.5 ± 0.2 (10)*</td>
</tr>
<tr>
<td>Liver proteins, mg/g</td>
<td>215 ± 12 (8)</td>
<td>242 ± 9 (8)</td>
</tr>
<tr>
<td>Glucose, mmol/L</td>
<td>7.1 ± 0.3 (10)</td>
<td>5.9 ± 0.2 (12)*</td>
</tr>
<tr>
<td>Triglycerides, mmol/L</td>
<td>0.70 ± 0.05 (5)</td>
<td>0.42 ± 0.03 (6)*</td>
</tr>
<tr>
<td>Total cholesterol, mmol/L</td>
<td>1.73 ± 0.15 (5)</td>
<td>1.75 ± 0.18 (5)</td>
</tr>
<tr>
<td>Alanine aminotransferase, U/L</td>
<td>12.7 ± 1.8 (8)</td>
<td>40.3 ± 9.9 (7)*</td>
</tr>
</tbody>
</table>

1 Values are means ± SEM (n). Means in a row without a common letter differ, P < 0.05.
2 One unit of alanine aminotransferase activity is defined as the amount of enzyme that catalyzes the transformation of 1 μmol of substrate/min at room temperature.

balance. Therefore, we analyzed the relevant plasma indicators of energy metabolism. The biochemical studies showed lower plasma concentrations of glucose and triglycerides in VAD diet-fed rats compared with control diet-fed rats (Table 2). Plasma total cholesterol concentrations did not differ between the groups. However, the alanine aminotransferase level, a marker of liver injury, was significantly higher in VAD rats than in controls (Table 2).

Plasma adiponectin concentration. Fat tissue regulates nutrient homeostasis by various mechanisms, including adipokine release. In fact, altered adipokine signaling has been implicated in the pathogenesis of fatty liver and insulin resistance (31). Therefore, we measured the changes taking place in the adiponectin of VAD rats. The plasma adiponectin concentration in VAD rats (Table 2; Fig. 2C) did not differ from the control group compared with the control diet-fed rats (Table 2). Plasma total cholesterol concentrations did not differ between the groups. However, the alanine aminotransferase level, a marker of liver injury, was significantly higher in VAD rats than in controls (Table 2).

Concentrations of major N compounds in plasma and urine. Urinary N excretion was significantly higher in the VAD group compared with the control group (Table 3). The difference in urinary N excretion can account for an increase in urea excretion, the major N compound found in the urine of rats. The concentration of creatinine in plasma and the 24-h urine creatinine excretion were similar in the 2 groups. Consequently, the renal clearance of creatinine was not affected by vitamin A deficiency. However, the plasma urea concentration was higher in VAD rats (Table 3; Fig. 2C). Treatment with RA did not change the plasma urea concentrations of VAD rats (Fig. 2C).

3-Methylhistidine concentrations in plasma and gastrocnemius muscle. The dietary deficiency of vitamin A for 30 (Fig. 2B) or 60 d (Fig. 2D) brought about a significant increase in the plasma 3-methylhistidine concentration. The plasma urea concentration did not differ between the groups at 30 d (Fig. 2A) but was higher in the VAD group than in controls at 60 d (Fig. 2C). As in plasma, the concentration of 3-methylhistidine in gastrocnemius muscle at 60 d was higher in VAD rats (6.2 ± 0.8 nmol/g) than in control rats (4.6 ± 0.3 nmol/g) (P < 0.005). RA did not modify the plasma urea concentrations of VAD rats (Fig. 2C) but normalized the plasma methylhistidine concentrations (Fig. 2D).

Liver urea cycle enzyme activities. The urea cycle enzyme activities in the VAD group were significantly higher than in the control group, except for ornithine carbamoyltransferase, which did not differ between the groups (Table 4). Results were similar when enzyme activities were expressed relatively to body weight. Consistent with the plasma urea data, RA treatment of VAD rats did not affect the hepatic enzyme activities (Table 4).

mRNA levels of the liver urea cycle enzymes. In agreement with the differences in the enzyme activities, the amount of mRNA for the urea cycle enzymes was 70–130% greater in VAD rats than in control rats (Table 4). As with the enzyme activities, RA treatment did not normalize mRNA levels in the VAD rats (Table 4).

Discussion

Most research into the pathogenesis of vitamin A deficiency has centered on abnormal fetal development and delayed growth in both humans and experimental animals (3–5,7,32). The effect of vitamin A deficiency on body size and weight during development appears to be associated with an alteration in nutrient utilization after absorption; moreover, N retention is decreased despite strictly controlling dietary intake (8). Therefore, the main aim of our study was to analyze the effect of chronic vitamin A deficiency on body weight, protein catabolism, ammonia detoxification, and N excretion during development. Our data clearly show that, despite the similar food intake in the VAD and control groups, vitamin A deficiency resulted in lower body weight, which was significant by wk 9 and had reached a weight plateau by this time. VAD rats also had lower liver and heart weights, but not lower white adipose tissue weight, than control rats. Moreover, protein catabolism was greater with vitamin A deficiency and was detectable at 30 d, clearly before body weight declined and when food intake was still normal. In parallel to the greater protein catabolism, N excretion was also higher, but no changes were noted in the renal function, which agrees with a greater removal of urea, the end-product of amino acid catabolism. These data indicate that the metabolic response to vitamin A deficiency implies a catabolic phase that is preceded by the increased production of 3-methylhistidine, suggesting that protein catabolism may be an important factor in the growth retardation induced by vitamin A deficiency.

Other authors have proposed that vitamin A deficiency may induce a catabolic state associated with GH resistance and the decreased expression of insulin-like growth factor 1 (IGF-1) (10,33). Although the effect of RA on pituitary GH synthesis in vivo has been controversial (34), several studies have also established a close linkage between the secretion of GH, the principal hormone stimulus for IGF-1 and vitamin A status in
mean 60 d treated further with RA for 10 d (VAD + RA) (control and VAD diets for 30 (796 Esteban-Pretel et al. reported that IGF-I blocks muscle proteolysis, an effect that glucose uptake by different tissues (38). In fact, it has been such as inhibition of muscular proteolysis and stimulation of deposition of fat, and these effects depend on IGF-1 secretion, which is delivered mostly by the liver (38). As mentioned above, the expression of hepatic IGF-1 has been reported to be depressed with vitamin A deficiency, and this may explain at least in part the impaired growth and fat accumulation in the VAD group. Moreover, IGF-1 has additional specific actions, which is in agreement with their long half-lives (40). Although not to control levels, RA diminished significantly (P < 0.02) and tended to diminish (P = 0.1) the mRNA for argininosuccinate synthetase and carbamoyl-phosphate synthetase, respectively. Some authors have found an upregulation of the mitochondrial urea cycle enzymes induced by mild oxidative stress (42) and we have shown in a recent report that RA treatment increases the oxidative stress generated by VAD (17,20). Whether this oxidative status is responsible for the elevated urea cycle enzyme mRNA remains to be clarified.

Additionally, fat accumulation and a histologically apparent fatty liver with hepatocytes revealing large lipid droplets were evident in the VAD group. Consistent with these data, several studies have shown that retinoid signaling plays an important role in adipocytes and energy balance. At the molecular level, a
could be an important factor in growth stimulation by IGF-I (39). Thus, the endocrine status induced by vitamin A deficiency (10,33) and the role ascribed to IGF-1 agree with the significantly greater muscle proteolysis that we found in retinol-depleted rats and suggest a possible mechanism for protein loss and delayed growth in retinol deficiency.

Our work shows that vitamin A deficiency induces a coordinated change of the urea cycle enzymes despite being located in different cellular compartments, which reflects quite closely that of the change in the urinary N output. In agreement with this finding, recent studies using Affymetrix oligonucleotide microarrays have observed that retinol deficiency induces the expression of the urea cycle enzymes in the depleted liver (10). Concerted changes in N excretion and in all urea cycle enzymes have been described under different physiological and pathological conditions (22,40,41). Specifically, the increased amino acid availability induced by high-protein diets leads to an adaptive increase in amino acid utilization, N and urea excretion, and urea cycle enzyme activities (40,41). However, we found higher N excretion and induction of the urea cycle enzyme in rats that had the same N intake as controls, thus supporting the idea of a greater availability of amino acids in the deficient group as a result of the enhanced catabolism of the endogenous proteins.

After RA treatment, protein catabolism decreased to the levels in control rats. Conversely, the urea cycle enzyme activities remained elevated, which is in agreement with their long half-lives (40). Although not to control levels, RA diminished significantly (P < 0.02) and tended to diminish (P = 0.1) the mRNA for argininosuccinate synthetase and carbamoyl-phosphate synthetase, respectively. Some authors have found an upregulation of the mitochondrial urea cycle enzymes induced by mild oxidative stress (42) and we have shown in a recent report that RA treatment increases the oxidative stress generated by VAD (17,20). Whether this oxidative status is responsible for the elevated urea cycle enzyme mRNA remains to be clarified.

Additionally, fat accumulation and a histologically apparent fatty liver with hepatocytes revealing large lipid droplets were evident in the VAD group. Consistent with these data, several studies have shown that retinoid signaling plays an important role in adipocytes and energy balance. At the molecular level, a

Table 4 Activities and mRNA for the urea cycle enzymes in the liver of rats fed control or VAD diets for 60 d and fed a VAD diet for 60 d treated further with RA for 10 d (VAD+RA)1,2

<table>
<thead>
<tr>
<th>Activity</th>
<th>Control</th>
<th>VAD</th>
<th>VAD + RA</th>
</tr>
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<tbody>
<tr>
<td>Carbohydrate-phosphate synthetase I</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Activity, U/g of liver</td>
<td>363.7 ± 39.9a (7)</td>
<td>647.5 ± 152.6b (8)</td>
<td>617.3 ± 20.9a (4)</td>
</tr>
<tr>
<td>mRNA</td>
<td>1.1 ± 0.2a (5)</td>
<td>1.8 ± 0.3b (4)</td>
<td>1.4 ± 0.2a (4)</td>
</tr>
<tr>
<td>Ornithine carbamoyltransferase</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Activity, U/g of liver (×10^-5)</td>
<td>28.2 ± 9.2 (7)</td>
<td>31.2 ± 10 (8)</td>
<td>29.3 ± 8.4 (4)</td>
</tr>
<tr>
<td>mRNA</td>
<td>4.9 ± 0.9b (8)</td>
<td>7.3 ± 0.8a (4)</td>
<td>6.0 ± 1.3a (4)</td>
</tr>
<tr>
<td>Argininosuccinate synthetase</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Activity, U/g of liver</td>
<td>101.8 ± 24.1b (7)</td>
<td>145.0 ± 31.2a (8)</td>
<td>111.3 ± 22.0b (4)</td>
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<tr>
<td>mRNA (×10^-5)</td>
<td>10.8 ± 1.3a (5)</td>
<td>20.8 ± 1.5a (4)</td>
<td>18.4 ± 0.5a (4)</td>
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<tr>
<td>Argininosuccinate lyase</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Activity, U/g of liver</td>
<td>129.6 ± 24.2a (7)</td>
<td>199.1 ± 45.3a (8)</td>
<td>201.3 ± 22.2a (4)</td>
</tr>
<tr>
<td>mRNA (×10^-5)</td>
<td>8.9 ± 0.9a (5)</td>
<td>21.4 ± 2.5a (4)</td>
<td>21.5 ± 12.3b (4)</td>
</tr>
<tr>
<td>Arginine I</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Activity, U/g of liver</td>
<td>54.3 ± 24.3b (7)</td>
<td>91.3 ± 18.9a (8)</td>
<td>90.6 ± 5.3a (4)</td>
</tr>
<tr>
<td>mRNA (×10^-5)</td>
<td>9.7 ± 3.4b (6)</td>
<td>19.1 ± 1.6a (4)</td>
<td>16.5 ± 7.6b (4)</td>
</tr>
</tbody>
</table>

1 Values are means ± SEM (n). Labeled means in a row without a common letter differ, P < 0.05.
2 Amounts of mRNA were normalized to β2-microglobulin.
cross-talk is well established between retinoid signaling and PPAR, which are the nuclear receptors that play a central role in the regulation of energy and lipid metabolism (43).

RA triggers an upregulation, and vitamin A deficiency a downregulation, of the PPARγ gene in liver and white adipose tissue, which may be a key event in the regulation of fatty acid oxidative metabolism by retinoid status in different tissues (15,44). In this sense, the functional loss of RA in transgenic mice expressing the RA receptor α-dominant negative form in hepatocytes developed both massive hepatic steatohepatitis, as a result of impaired mitochondrial fatty acid oxidation, and liver tumor (13). RA treatment also induced a dose-dependent increase in the skeletal muscle mRNA expression levels of PPARδ in vivo, which was associated with not only an increased lipid oxidative capacity in this tissue but also decreased muscle lipid content (16). Moreover, retinaldehyde and/or RA antagonized PPAR-γ activity in vivo producing a loss of body adipose tissue with changes including a reduction of adipocyte size, triacylglycerol content, and lipogenic capacity (44,45). Transgenic mice that did not express retinaldehyde dehydrogenase, which consequently have elevated retinaldehyde concentrations, had decreases in subcutaneous and visceral adipose tissue, liver lipid deposition, and plasma adiponectin (45). In agreement with these data, our results show that loss of retinoid signaling in vitamin A deficiency induces the opposite effect, i.e. greater relative epididymal-adipose tissue and liver lipid deposition and higher concentrations of alamine aminotransferase and adiponectin in plasma. Plasma elevations of alamine aminotransferase have also been described in several models of hepatic steatosis, including those induced by an altered retinoid status (13,46).

The increased expression and release of the PPARy target gene adipocytokine adiponectin (also known as apM1, adipQ, Gbp28, and Acrp30) and enhanced triglyceride accumulation are specific and distinct indicators of adipogenesis (45). Therefore, the fact that vitamin A deficiency induces an increase in fat depots with enlarged hepatic lipid droplets and a rise in adiponectin plasma concentrations suggests that vitamin A deficiency induces adipogenesis.

In conclusion, our work shows that chronic dietary vitamin A deficiency, without changing the intake of dietary protein, increases protein catabolism and raises the gene transcription rate and the corresponding activities of hepatic urea cycle enzymes. The elevation of both the muscle protein catabolic response and urinary nitrogen excretion provides insight into the origin of protein losses following vitamin A deficiency. This elevated protein catabolism was accompanied by higher fat mass and lower body and tissue weight in VAD rats than in control rats. Thus, in the absence of vitamin A, fuel partitioning may shift from fatty acids to protein catabolism as an alternative energy source, leading to progressive protein loss. RA partially reversed the changes observed in VAD rats. The integration of all these results could contribute to an understanding of the complex interaction between vitamin A status and the transcriptional regulation of metabolism.

The accelerated protein breakdown, increased N elimination, and abnormal deposition of triacylglycerols in tissues may contribute to disease development in chronic vitamin A deficiency. Moreover, we have previously reported that vitamin A deficiency produces hepatic oxidative stress (17), which may play an important role in the initiation and/or progression of hepatic lesion (13), and also that it results in profound alterations in the basement membrane of different tissues (20,24). All these could be contributing factors to tissue malfunction in vitamin A deficiency.

Acknowledgments
We thank A. Catharine Ross for her valuable advice on the induction and handling of VAD rats. G.E-P., M.P.M., F.C., and V.M. conducted the research; G.E-P. analyzed the data; J.R-P., J.T., and T.B. designed the research; and J.T. and T.B. wrote the paper and had primary responsibility for the final content. All the authors read and approved the final manuscript.

Literature Cited

Increased proteolysis in vitamin A deficiency


