

# Biotin: From Nutrition to Therapeutics

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## Abstract

Although frank symptomatic biotin deficiency is rare, some evidence suggests that marginal biotin deficiency occurs spontaneously in a substantial proportion of women during normal human pregnancy and might confer an increased risk of birth defects. Herein I review 1) advances in assessing biotin status, including the relation between acylcarnitine excretion and biotin status; 2) recent studies of biotin status in pregnancy; 3) advances in understanding the role of biotin in gene expression and the potential roles of biotinylated proteins that are neither histones nor carboxylases; and 4) novel large-dose biotin supplementation as therapy for multiple sclerosis. The review concludes with a summary of recent studies that have reported potentially dangerous erroneous results in individuals consuming large amounts of biotin for measurements of various plasma hormones for common clinical assays that use streptavidin-biotin technology. *J Nutr* 2017;147:1487–92.

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## Introduction

Hamid Said, an eminent investigator in biotin nutrition and physiology, asked the following in the *American Journal of Clinical Nutrition* more than a decade ago: “Biotin bioavailability and adequate intake: why bother?” (1). This article reviews studies that have provided evidence that the bother is worthwhile and presents observations concerning cautions for individuals consuming large amounts of biotin. The reader is referred to several excellent reviews for a more thorough discussion of the traditional role of biotin as an essential cofactor for the 5 biotin-dependent carboxylases (2–7).

## Historical Perspective

Frank symptomatic biotin deficiency is a rare occurrence. The only well-documented cases have occurred in association with total or near-total intravenous feeding without biotin supplementation during the chronic consumption of undenatured egg white and with inborn errors of metabolism that lead to biotin wasting (2, 6). A single case that does not fit any of these established associations is that of an infant fed a rice-based formula that was presumably very low in biotin (6). However, some studies have suggested that the absence of

overt biotin deficiency does not imply optimal biotin nutritional status (8–14).

## Brief Review

Biotin-dependent carboxylases are initially synthesized as enzymatically inactive apocarboxylase proteins that become active holocarboxylases after the covalent attachment of biotin. The attachment of biotin is catalyzed by holocarboxylase synthetase (HLCS); an amide bond is formed between the carboxyl group of the valeric acid side chain of biotin and the  $\epsilon$ -amino group of a specific Lys residue in each of the 5 apocarboxylases. These regions contain sequences of amino acids (e.g., Met-Lys-Met at the attachment site) that are highly conserved for the individual carboxylases, and both the N and C terminus in holocarboxylase synthetase are important for the recognition of the apocarboxylases (15). In mammals, biotin serves as an essential cofactor for each of the 5 biotin-dependent carboxylases: acetyl-CoA carboxylase (ACC) 1, ACC2, methylcrotonyl-CoA carboxylase (MCC), pyruvate carboxylase, and propionyl-CoA carboxylase (PCC) (Figure 1). These roles have been extensively reviewed previously (2–7). Additional roles are being actively investigated as described in the following sections.

## Does Biotin Deficiency Occur in Pregnancy?

Despite the rarity of symptomatic biotin deficiency, pregnancy is a clinical condition of particular concern with respect to biotin status. A post hoc analysis of data from a large multivitamin supplementation study (17) provided interesting, although indirect, evidence that the marginal degree of biotin deficiency

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Abbreviations used: ACC, acetyl-CoA carboxylase; AI, adequate intake; BTBGD, biotin-thiamin-responsive basal ganglia disease; HLCS, holocarboxylase synthetase; HSP, heat-shock protein; MCC, methylcrotonyl-CoA carboxylase; PCC, propionyl-CoA carboxylase; 3HIA, 3-hydroxyisovaleric acid; 3HIAc, 3-hydroxyisovaleryl carnitine.



lead to a disruption of the esterified CoA:free CoA ratios and, ultimately, to potentially lethal mitochondrial toxicity (20, 21). To prevent this, 3HIA-CoA is detoxified by carnitine trans-esterification to 3-hydroxyisovalerylcarnitine (3HIAc). The reaction is catalyzed by carnitine acetyltransferase, which is one of a family of enzymes with varying chain-length specificity and organelle and organ distribution; these enzymes act in concert with the transfer of the acylcarnitine out of the mitochondria by carnitine-acylcarnitine translocase to defend the CoA ratios (21). Accumulating 3HIAc is transferred across the inner mitochondrial membrane by the translocase, leading to increased plasma and urinary 3HIAc and 3HIA (12, 13) and potentially to secondary carnitine deficiency (22). 3HIA likely arises preferentially from the hydrolysis of 3HIAc in the cytosol.

## Indicators of Biotin Status

Urinary 3HIA and 3HIAc are sensitive indicators of marginal biotin deficiency in healthy adults in whom biotin deficiency is induced experimentally (12). Interestingly, the urinary excretion of 3HIAc was actually lower among pregnant women than among nonpregnant control women in Perry et al. (19), indicating that urinary 3HIAc is not a reliable indicator of marginal biotin deficiency in pregnancy. Our group has observed the same unreliability of urinary 3HIA as an indicator of biotin deficiency in pregnancy (A Bogusiewicz, G Boysen, DM Mock, University of Arkansas for Medical Sciences, unpublished results, 2017) and, in a hepatocyte culture, has confirmed the metabolic pathogenesis presented previously by inducing separate and combined biotin and carnitine deficiency (23); the results were consistent with Perry et al. (19) in that urinary 3HIAc did not increase in parallel with urinary 3HIA because of the functional deficiency of hepatic carnitine. This observation also raises the possibility that this functional carnitine deficiency might impair this important cellular detoxification mechanism in pregnancy.

## Biotin Catabolism

Perry et al. (19) also offered intriguing observations concerning biotin catabolism. Lactating women excreted substantially more of the inactive catabolite bisnorbiotin, which is created by the oxidation of the valeric acid side chain, than did control women. This observation suggests that lactation accelerates biotin turnover and loss. Indeed, the accelerated biotransformation of biotin to bisnorbiotin has been reported in early pregnancy (24); however, in that study, bisnorbiotin excretion returned to normal by late pregnancy.

## Utility of Indicators of Biotin Status

In the context of these seemingly conflicting observations, Eng et al. (25) reported that the abundance of holo-PCC and holo-MCC in peripheral blood lymphocytes determined by gel densitometry and fluorescent-labeled streptavidin seem to be the best indicators of marginal biotin deficiency. Theoretically, the activity of PCC and MCC (2 of the 5 biotin-dependent carboxylases) in peripheral blood lymphocytes should be as sensitive and specific as the abundance of holo-PCC and holo-MCC. The assay quantitates the catalysis by PCC or MCC of  $^{14}\text{C}$ -bicarbonate incorporation into acid-precipitable material (i.e., methylmalonyl-CoA and methylglutaconyl-CoA, respectively) (26). Unfortunately, this assay is technically demanding, and the blood samples require special handling and storage,

making gel densitometry abundance measurement more practical for field studies. Likewise, the urinary excretion of 3HIA and other organic acids characteristic of multiple carboxylase deficiencies as well as the related acylcarnitines seem to be less robust as indicators of biotin status in field studies (25), perhaps because of the variation in the dietary intake of precursors (e.g., Leu) in the hours before the collection of the urine sample (14, 27).

## Role of Biotin in Gene Regulation

Hymes and Wolf (28) suggested that the posttranslational biotinylation of histones might play a role as a covalent modifier in the epigenetic code that regulates DNA transcription. However, subsequent work showed that <0.001% of human histones (primarily H3 and H4) are biotinylated, suggesting that the abundance is too low to elicit biological effects in vivo (29, 30). However, HLCS is located prominently in the nucleus (31), and the knockout of *HLCS* in *Drosophila* as well as in human and other mammalian cells in culture produces distinct phenotypes, including the derepression of long terminal repeats and chromosomal instability; aspects of this *HLCS* knockout phenotype have been attributed to the effects on gene expression rather than reduced activities of the biotin-dependent carboxylases (2).

## HLCS Exerts Some of Its Roles in Gene Regulation through the Formation of a Multiprotein Gene-Repression Complex in Human Chromatin

Zempleni and colleagues (32, 33) have proposed that the biological effects of biotin on gene expression are caused by a multiprotein complex, including proteins involved in DNA methylation, histone methylation, and histone deacetylation. They proposed that the docking of HLCS in chromatin causes an occasional biotinylation of histones ("marks") near the various HLCS-binding sites. Their studies provide evidence that HLCS enters the nuclear compartment and is recruited to chromatin through physical interactions with DNA methyltransferase 1 and methyl CpG-binding protein 2. Chromatin-bound HLCS has been shown to recruit the eukaryotic histone H3 methyltransferase euchromatic histone lysine *N*-methyltransferase 1, which creates abundant Lys9-methylated histone H3 gene repression marks. In addition, HLCS interacts with nuclear receptor corepressor, a protein known to facilitate the binding of histone deacetylases (HDACs) in chromatin. HDACs remove histone acetylation marks and thus play a critical role in gene repression. Overall, emerging data suggest histone biotinylation marks are a side effect of HLCS being in close physical proximity to histones and play no direct role in gene repression, despite contributing toward chromatin condensation (32).

## Biotinylated Proteins Other Than Carboxylases and Histones: Evidence That the Number of Proteins Containing Covalently Bound Biotin Is Larger Than Previously Appreciated

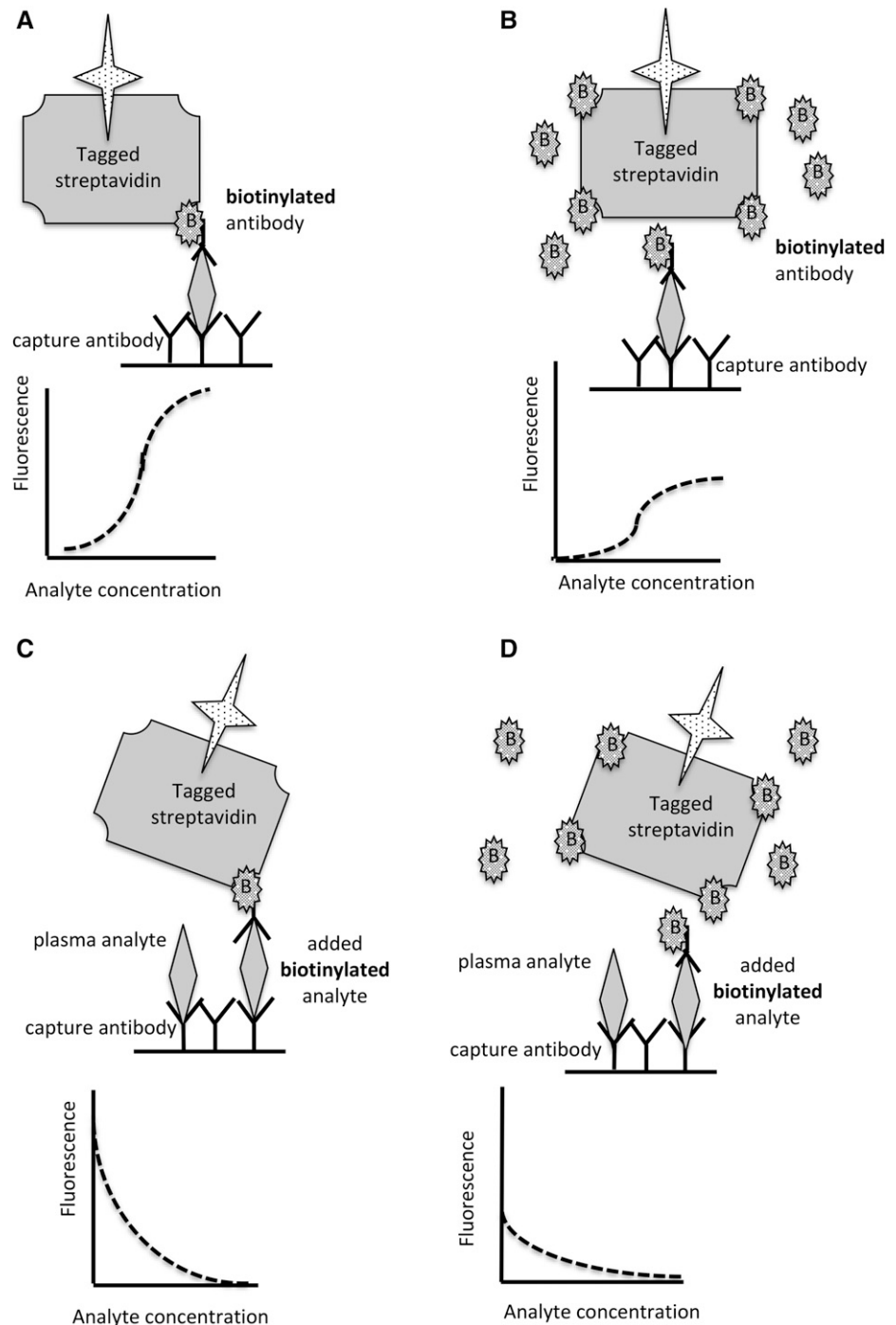
A recent study (34) used MS to identify 108 novel biotinylation sites in the human embryonic kidney cells; members of the heat-shock

protein (HSP) superfamily were overrepresented among the novel biotinylated proteins. Approximately 50% of the biotinylated proteins displayed increased Met oxidation; for biotinylated HSPs, Met sulfoxidation approached 100%. Protein structure analysis suggested that Met sulfoxides localized in close physical proximity to the biotinylated Lys residues on the protein surface and that the likelihood of Met sulfoxidation is greater when one of the adjacent Lys residues is biotinylated. HSP60 knockdown and biotin-depletion experiments supported a synergistic role of biotinylation of Lys residues in the Met defense against reactive oxygen species and further observed that high concentrations of Met sulfoxidation coincided with cell-cycle arrest in biotin-depleted cells.

## High-Dose Biotin Therapy in Profound Neurologic Diseases

Biotin-thiamin-responsive basal ganglia disease (BTBGD) is a genetic disorder that affects the nervous system, including a group of structures in the brain termed the basal ganglia that help control movement. BTBGD is caused by a defect in the solute carrier family 19 member 3 gene that encodes for thiamin transporter 2 (35, 36) and is characterized by subacute episodes of encephalopathy often triggered by febrile illness and characterized by confusion, epilepsy, external ophthalmoplegia, dysphagia, and generalized stiffness; this disease eventually results in coma and death (37). Before elucidating the genetic

**FIGURE 2** Sandwich assay with the use of fluorescent-tagged streptavidin detection (A, B). (A) Normal conditions: bound antibody captures the analyte, which then binds a biotinylated antibody raised against a second epitope on the analyte. The biotinylated secondary antibody binds to fluorescent-tagged streptavidin. The greater the concentration of plasma analyte, the greater the fluorescent signal. (B) Excess plasma biotin blocks streptavidin binding to the biotinylated antibody, leading to an underestimation of the analyte. (C, D) Competition assay with the use of fluorescent-tagged streptavidin detection. (C) Normal conditions: plasma analyte competes with the added biotinylated analyte. The greater the concentration of the analyte, the less streptavidin is bound and the smaller the signal. (D) Excess biotin in plasma blocks streptavidin binding to the biotinylated analyte, leading to an overestimation of the plasma analyte.



pathogenesis, high doses of biotin alone ( $5\text{--}10\text{ mg} \cdot \text{kg}^{-1} \cdot \text{d}^{-1}$ , which is  $\sim 10,000$ -fold greater than the AIs for children and adults) had been used in the successful treatment of this disease; the addition of thiamin improved the clinical prognosis for some patients (38). To our knowledge, the therapeutic mechanism for high-dose biotin in BTBGD remains unknown, but interesting clues are emerging from observations in multiple sclerosis.

## High-Dose Biotin May Slow the Advancement of Progressive Multiple Sclerosis

In 2011, Sedel et al. (39) reported on 5 patients who were suffering from optic neuropathies and leukoencephalopathy. Encouraged by the success of high-dose biotin in BTBGD, biotin therapy was instituted; all responded clinically. Thereafter, 1 of the 5 patients was diagnosed as actually having secondary progressive multiple sclerosis (40). High doses of biotin ( $100\text{--}300\text{ mg/d}$ ) were then tested in 23 additional patients with primary or secondary progressive multiple sclerosis with the use of an open-label design; improved clinical outcomes were reported in nearly all participants (40). In 4 patients with prominent visual impairment related to optic nerve injury, visual acuity improved considerably, and visually evoked potentials improved in 2 patients. Proton magnetic resonance spectroscopy in 1 patient showed an improved choline:creatine ratio. One patient with left-sided blindness for both eyes steadily improved from 2 to 16 mo of biotin supplementation. Of the 18 patients with prominent spinal cord involvement, 16 improved; a blinded review of a videotaped clinical examination was possible for 9 patients and confirmed improvement in all of them. Preliminary results from multicenter double-blind placebo-controlled trials in Europe and the United States are encouraging (41).

Thus, high-dose biotin supplementation may represent a therapeutic option in progressive multiple sclerosis; the mechanism(s) remains to be elucidated (42). The gradual worsening of neurologic disability in patients with progressive multiple sclerosis is caused by progressive axonal loss or damage. The triggers for axonal loss in multiple sclerosis likely include both inflammatory demyelination of the myelin sheath and primary neurodegeneration caused by a state of virtual hypoxia within the neuron. High-dose biotin might increase myelin production by increasing the generation of long-chain FAs (e.g., via effects on ACC1), increasing energy production via the tricarboxylic acid cycle in neuronal cells, or both (42). In the context that the tricarboxylic acid cycle is heavily dependent on acyl-CoA intermediates, reports that chickens fed a diet deficient in pantothenic acid (a required precursor of CoA) developed spinal nerve damage associated with the degeneration of the myelin sheath are intriguing (43). Early results from high-dose biotin treatment in animal models of multiple sclerosis favor increased myelin production.

## Biotin Supplementation Can Interfere with Clinical Laboratory Assays That Use Streptavidin Technology

In both adults and children, pharmacologic doses of biotin are being used with increasing frequency for the legitimate indications discussed previously as well as for off-label conditions (44–47). This trend is likely to continue; the chances of interference with clinical diagnostic tests that depend on streptavidin-biotin technology will likely increase as well. Immunoassays that use

streptavidin-biotin technology are now commonly used for hormone measurements and are common in high-throughput analytical platforms. Reports of erroneous analytical results are appearing with increasing frequency; most are encountered in individuals who received biotin supplementation. Pretreating the plasma sample with streptavidin microbeads normalizes assay results (47). However, a few cases were caused by anti-streptavidin antibody (45). For those receiving biotin therapy, the interference may induce either falsely increased or falsely decreased results, depending on the assay design. As depicted in Figure 2A, B, excess plasma biotin in competition assays characteristically lead to an overestimation of the analyte; as depicted in Figure 2C, D, excess plasma biotin in double-epitope “sandwich” detection and capture immunoassays cause underestimation. Unfortunately, the artifacts acting on 2 hormone assays in the same sample can simulate a seemingly coherent hormonal profile; e.g., falsely increased free thyroxine and falsely decreased thyroid-stimulating hormone in the same plasma sample suggests hyperthyroidism (45). Nutritionists, physicians, nurses, and other healthcare providers should include biotin in the nutritional history in patient groups that might have instituted biotin therapy, such as those with multiple sclerosis.

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