

2 Physiological Role of Copper

This chapter begins by discussing the basis of the essentiality of copper. The kinetics of copper in the body and select roles of copper at the cellular and molecular level are described. In addition, factors influencing the bioavailability of copper are presented.

Essentiality

Copper is both essential and toxic to living systems. As an essential metal, copper is required for adequate growth, cardiovascular integrity, lung elasticity, neovascularization, neuroendocrine function, and iron metabolism. An average adult human ingests about 1 mg of copper per day in the diet; about half of which is absorbed (Harris 1997). An expert committee of the World Health Organization recommends 30 micrograms (µg) per kilogram (kg) of body weight per day, which equates to about 2 mg per day for an average adult (WHO 1996). The Food and Nutrition Board (FNB) recommends dietary copper intake of 1.5–3.0 mg per day for adults (NRC 1989). Copper is obligatory for enzymes involved in aerobic metabolism, such as cytochrome c oxidase in the mitochondria, lysyl oxidase in connective tissue, dopamine monooxygenase in brain, and ceruloplasmin. As a cofactor for apo-copper-zinc superoxide dismutase (apoCuZnSOD), copper protects against free-radical damage to proteins, membrane lipids, and nucleic acids in a wide range of cells and organs. Severe copper deficiencies, either gene defects due to mutations or low dietary copper intakes, although relatively rare in humans, have been linked to mental retardation, anemia, hypokinesia, neutropenia, diarrhea, cardiac lymphopathy, bone fragility, impaired immune function, weak connective tissue, impaired central-nervous-system (CNS) functions, peripheral neuropathy, and loss of skin, fur (in animals), or hair color (Hinderly and Goode 1991; Umayi et al. 1998; Ruscione 1998).

Biochemistry and Physiology

Copper taken in through the diet might be absorbed partially in the stomach, where the highly acidic environment frees the bound copper ions from partially digested food particles. However, the largest portion of ingested copper passes into the duodenum and ileum, which are the major sites of absorption. As a result of complexing with amino acids, organic acids, or other chelators, a high fraction of copper is soluble in the intestinal tract. Studies on isolated segments of the duodenum suggest that copper ions enter into the mucosal cells lining the intestine by simple diffusion and exit at the basolateral surface by a different transporter (Bremner 1980). Recent reports indicate that there is a divalent transporter that might transport copper (Rofis and Hedinger 1999).

Basolateral transport is markedly reduced in Menkes disease, which results in systemic copper deficiency. Studies of this disease led to the prediction of a copper-transporting adenosine triphosphatase (ATPase) in the basolateral membrane of mucosal cells. The copper-transporting ATPase presumably discharges the copper into the serosal capillaries where the copper binds to albumin and amino acids for transport to the liver via the portal circulation. From the liver, copper is transported to extrahepatic tissues by albumin, amino acids, and, to a lesser extent, ceruloplasmin (Dunn et al. 1991).

A large fraction of circulating copper is returned to the liver as ceruloplasmin-bound copper. Ceruloplasmin, a sialoglycoprotein, is constantly being secreted from the liver into the blood. When ceruloplasmin returns to the liver, the sialic acid can be removed by the outer endothelial cells followed by ATPase of the desialated protein via the asialoglycoprotein receptor in the liver parenchyma (Irie and Tavassoli 1986). Likewise, removal of copper from ceruloplasmin hastens its uptake by liver parenchymal cells (Holtzman and Gaumnitz 1970).

As discussed in Chapter 4, Wilson disease, a genetic disease characterized by accumulation of copper mainly in the liver and brain, attests to a potential role for ceruloplasmin biosynthesis in liver homeostasis of copper. Copper-containing fragments of ceruloplasmin are found in the bile of normal subjects and are generally absent from the bile of Wilson patients.

Although many Wilson patients do not synthesize sufficient amounts of ceruloplasmin, this decrease in biliary ceruloplasmin is observed even in Wilson patients who have reasonably normal blood concentrations of the protein. That observation suggests that biliary secretion of copper bound to ceruloplasmin accounts for most of the copper that is excreted (Iyengar et al. 1988). In aceruloplasminemia, a genetic defect in ceruloplasmin biosynthesis, ceruloplasmin in the circulation is totally absent (Harris et al. 1995). Yet, the aceruloplasminemic patient does not develop Wilson disease, nor does copper accumulate in the liver in aceruloplasminemia. Thus, factors besides ceruloplasmin are required for biliary copper excretion and for maintaining normal liver copper homeostasis in the liver. A clue to the identity of such factors has come from studies of lipofuscin-like granules that accumulate in the liver of patients who suffer from primary biliary cirrhosis and are unable to excrete copper. These granules apparently arise from primary lysosomes and might contain degradation fragments of metalloprotein-bound copper (Humbert et al. 1982). Studies in Wistar rats suggest that biliary copper excretion occurs by a glutathione-dependent (rapid) phase and a glutathione-independent (slow) phase (Holtzman et al. 1990). No identification has been made of which rat enzyme is associated with ceruloplasmin.

A second copper-transporting ATPase enzyme is required to transport the copper ions into the Golgi compartment for incorporation into apo-ceruloplasmin (Murata et al. 1995). Several types of cells have been shown to have receptors for ceruloplasmin. However, the above observation supports, but does not define a role for ceruloplasmin in copper delivery to tissues. The receptors have been found on many cell types. K562 cells are capable of engaging ceruloplasmin in liver and transporting ceruloplasmin-bound copper to cellular enzymes (Percival and Harris 1991). Unlike transferrin, which delivers iron by a receptor-mediated endocytosis of the whole protein, ceruloplasmin protein is not endocytosed (except in hepatic cells) and delivers its bound copper to components at the cell surface. The role of the protein, therefore, stops at the cell membrane, and transport of copper to the interior requires transport proteins in the membrane. Ascorbate facilitates the release of copper, and chelators of copper prevent its absorption by the cell (Percival and Harris 1989, 1990).

When excess copper is fed to rats, it selectively accumulates in periportal and midzones of the liver lobules (Fuentetaja et al. 1989), suggesting that portal blood flow determines copper disposition (Haywood 1981). In some patients with chronic cirrhosis, copper also accumulates in lysosomes (Humbert et al. 1982), suggesting that those organelles take part in copper storage or, more likely, in copper excretion when copper concentrations are high (Gross et al. 1989).¹ Metallothionein is a small cysteine-rich protein that tightly binds copper. This protein is thought to play roles in both copper storage and internal copper movement (Cousins 1985). At high concentrations, copper can stimulate metallothionein synthesis. Under normal conditions, copper exists at extremely low concentrations as the free ion in the cytosol. Copper-binding ligands, therefore, are key regulators in copper atoms, ATP and the adaptation to toxic effects. The ligands protect against toxicity and can help target the proteins for copper incorporation. Copper-binding ligands include glutathione (GSH), amino acids, metal, and recently identified copper metallochaperones. All of these ligands have been shown to transport copper within the cell from one location to another, and make copper available to intracellular enzymes (Sternlieb 1980; DiSilvestro and Cousins 1983; Holt et al. 1986; Harris 1995).

Numerous biochemical and nutritional studies have focused on the mechanisms of copper absorption and metabolism. The challenge is to explain how extremely small quantities of copper are able to impact cell membranes and into enzymes that require copper for function. Some insight has been obtained from studies in yeast. Yeast models of copper transport with separate low- and high-affinity systems for copper uptake (Cu(I)). The yeast gene *CTR1* (copper transport 1) was the first copper transporter identified to be discovered (Dancis et al. 1994a,b). *CTR1* encodes a transmembrane protein that selectively transports Cu(I). A homologous human gene, *hCTR1*, that encodes a transporter protein was identified in HeLa cells (Zhou and Gitschier 1997). Because the above transporters recognize only Cu(I), a reductase enzyme must reduce Cu²⁺ before transport into the cell. Two reductase genes that can reduce Cu²⁺ for transport were identified in yeast (Hassett and Kosman 1995; Georgatou et al. 1997). A reductase system that uses NAD⁺ as an electron donor to reduce copper has been found in rat liver cells (van den Berg and McArdle 1994).

Cells in a defined culture medium take up copper. High-affinity membrane permeases allow copper ions at micromolar concentrations to move through the phospholipid bilayer and enter the cytoplasm. The rapid uptake is not ATP dependent, suggesting a passive carrier system (Schmitt et al. 1983; Tong and McArdle 1995). In general, chemical form, valence state, and relative concentrations of competing metals influence the quantity of copper that is absorbed. Select amino acids such as histidine and glutamine (Harris and Sasse-Kortsak 1967), and ions, such as sodium, chloride, and bicarbonate (Aida and Garay 1990), stimulate transmembrane copper movement in vitro, whereas zinc and copper chelators, such as phytate, reduce absorption.

Glutathione is a ubiquitous cysteine-containing tripeptide, is present in millimolar quantities in liver, brain, kidney, and other tissues (Denke and Fanburg 1989). Glutathione avidly binds copper. There is a rapid turnover of a GSH-Cu(I) complex in hepatoma cells. That turnover is consistent with GSH involvement in the reduction of Cu(II) to Cu(I), potentially facilitating its binding to metallothionein (Freedman and Peisach 1989; Freedman and Peisach 1989b).

Mammalian and yeast cells have small polypeptides that bind Cu(I) and transfer it to selected recipients. These proteins, called copper chaperones or metallochaperones, transiently bind Cu(I) at a cysteine-rich region in the peptide chain (Valentine and Gralla 1997). Chaperones provide copper ions from one location in the cell to another, sometimes crossing through organelle membranes. The chaperone promotes a rapid exchange of copper with the target proteins (Portnoy et al. 1999). Targets include cytochrome c oxidase in the mitochondria, ATP7B in the trans-Golgi, and apoCuZnSOD in the cell cytosol. Three chaperones first described in yeast are known to have human counterparts. ATOX1 (formerly HAH1) is the human homolog of ATOX1 and is abundantly expressed in all tissues. ATOX1 and ATOX2 specifically deliver copper to the secretory pathway where the targets are membrane-bound copper ATPases that regulate the flow of copper into cell compartments. Examples include Ccc2p, which delivers copper to a multicopper oxidase (FeTsp1) required for iron uptake in yeast, and ATP7B (the Wilson disease gene product), which delivers copper to apo-ceruloplasmin in the trans-Golgi of liver. Proper ATP7B function is essential for the excretion of copper in the bile. COX17 is a chaperone that transports copper to cytochrome oxidase in the mitochondria of yeast cells (Carr-Saunders 1997). The human homolog is hCOX17. LY57, a 27-kilodalton (kDa) protein that delivers copper to the apoCuZnSOD (Culotta et al. 1997, 1999), has a human counterpart designated CCS (copper chaperone for SOD). Human cells deficient in LY57 are unable to incorporate copper into SOD 1 and hence are defective for SOD1 activity. CCS is comparable in size and has a 28% sequence identity with LY57. Both proteins contain a single MHCXXC consensus copper-binding sequence.

Factors Affecting Bioavailability

The amount of copper which is absorbed from the diet can vary considerably depending on other dietary constituents. However, in general approximately half the copper consumed in the diet is typically absorbed by the gastrointestinal (GI) tract. Approximately two-thirds of the copper that is absorbed is rapidly secreted into the bile (Lönnerdal 1998; Turnlund 1998; Wapnir 1998) (Figure 2-1). Thus, approximately 80–90% of dietary copper is typically excreted in the feces. Thus, copper homeostasis is primarily regulated at the GI level, through biliary excretion with the kidney excreting only small amounts of copper. Small amounts of copper are also excreted in hair and sweat. The bioavailability, or the fraction of copper absorbed from the GI tract, has been shown to be influenced by the age of the individual, the amount of copper in the GI tract, and various dietary components. With respect to dietary components, copper in meat has been reported to be more bioavailable than that in vegetables. The bioavailability of copper is also expected to depend on the form of copper present (Baker et al. 1991). Absorption is higher for soluble or ionic forms than for less soluble or insoluble mineral forms or copper deposited in soil.

In adults, absorption varies according to the amount of copper in the diet. Animal studies indicate that absorption rates can be as low as about 10% with very high copper intakes, and as high as around 70% with low copper intakes. The average for typical diets in animals and humans is 30–40% (Lönnerdal 1996, 1998; Turnlund et al. 1998; Wapnir 1998). Turnlund et al. (1988) measured the amount of copper excreted by humans in the feces for 12 days after an oral dose, or intravenous dose, of copper for four subjects and five subjects, respectively. The intravenous dose allows for an estimation of the endogenous excretion into the gut. Retention from oral intake as estimated from copper in the feces was shown to be highest when dietary copper concentrations were lowest: 67% at 0.38 mg per day, 54% at 0.66 mg per day, and 44% at 2.49 mg per day (Turnlund et al. 1988). However, the estimated total percentage that was actually absorbed by the GI tract before endogenous (or biliary) excretion back to the GI tract was 77%, 73%, and 66%, respectively, for the three doses. Thus, changes in endogenous excretion (in the bile), rather than GI absorption, is the primary mechanism of action in regulating total body copper. Specifically, the change in endogenous excretion between the lowest dose and the highest dose varied between 12% and 34% (Turnlund et al. 1988), although those data might not adequately reflect long-term homeostasis because of the short-term nature of the study. Thus, increases in copper excretion in the feces with increasing copper dose is a function of decreased absorption and increased biliary secretion, the latter having a greater effect.

Effect of Age

Studies in rats indicate that the absorption and retention of copper can be particularly high in the neonatal period, and that it decreases by the time of weaning (summarized by Lönnerdal 1996, 1998). Data on absorption in human infants are sparse; however, studies on copper balance (i.e., the amount of copper in the body) show similar decreases in retention (a function of absorption and biliary excretion) of copper with age in full-term infants. Negative copper balance (a loss of copper from the body which exceeds dietary intake) can arise in preterm infants as a consequence of a reduced prenatal storage of copper and low dietary copper intake (Widdowson 1974; Sann et al. 1980; Hillman et al. 1981; Lönnerdal 1996, 1998; Cordano 1998). Both animal and human infant studies indicate that, within the intake range examined, copper absorption and retention increase linearly with intake amount, although the percentage of the dietary copper retained decreases (Lönnerdal et al. 1985; Lönnerdal 1996). For example, in 14-day-old suckling rat pups, the percentage of copper retained from a meal (0.5 mL) containing 0.2 mg of copper per liter was approximately 30%, and it was lower to 20% when the meal contained 2 mg of copper per liter. However, it is important to note that while the percentage of copper retained from the high-copper meal was lower than that from the low-copper meal, the total amount of copper retained from the high-copper meal was 7 times higher than that from the low-copper meal (Lönnerdal et al. 1985).

Due to tissue growth, and an increased expression of some copper proteins during the postnatal period, the percentage of absorbed copper retained by the body is higher in infants than in adults. Little information is available on dietary copper absorption, or copper retention, in toddlers and young children.

Dietary and Other Interactions

Early research with whole animals showed that the rate and amount of copper ions transferred across intestinal epithelia were influenced positively by dietary amino acids, but negatively by ascorbate and competing metal ions (Bremner 1980; Hogan and Rausser 1981; L'Abbe and Fischer 1984; Oestreicher and Cousins 1985; Fields et al. 1986). Chloride ions and bicarbonate appeared to stimulate cellular absorption (Aida and Garay 1990). The transport across the membrane was considered a property of transport proteins, themselves subject to antagonistic effects of competing metals, principally zinc and ferrous iron. Studies in yeast and bacteria have led to the discovery of membrane proteins that transport copper ions across cell membranes.

Copper absorption is reported to be greater in infants who ingest human milk rather than cows milk (Lönnerdal 1996, 1998). That increased bioavailability might be attributed to the greater association of copper with albumin and whey in human milk, than in cow's milk, where much of the copper is bound to casein (Wapnir 1998).

The literature indicates that copper absorption is greater when diets are animal protein rather than plant protein (i.e., vegetarian) (Srikumar et al. 1992; Lönnerdal 1996; Wapnir 1998). Studies in experimental animals found that phytates and dietary fiber generally reduce copper bioavailability; however, effects on bioavailability are less clear in humans (Turnlund 1988; Lönnerdal 1996). Declines in serum copper give some indication that phytates or α-cellulose added to the diet might alter copper utilization or distribution (Turnlund 1988). In general, the effect of phytates and dietary fiber on absorption of copper appears to be less than the effect on absorption of other divalent cations, such as zinc (Wapnir 1998).

Dietary differences have been found in patients with Wilson disease. Observations of two Wilson patients on lacto-vegetarian diets suggest copper bioavailability is reduced by about 25% (Brewer et al. 1993b). The first of two patients on that diet was asymptomatic (i.e., normal liver function and normal slit lamp examination for Kayser-Fleischer rings) for 12 years despite having a typical average daily copper intake and no anticopper therapy. The second patient started on the anticopper therapy for 2 years and then switched to a lacto-vegetarian diet. After switching, her serum transaminase and transpeptidase activity (alanine aminotransferase, aspartate aminotransferase, and liver γ-glutamyltransferase), which were previously elevated, showed improvement over the succeeding year. When last observed about 2 years after starting the diet, the patient remained clinically well. Other Wilson-disease patients who discontinued their anticopper therapy had serious difficulty after 3 to 4 months and serious degeneration in their condition after 1.5 years on average (Brewer and Yuzbasiyan-Gurkan 1992).

Carbohydrates, such as fructose, have been studied for their effect on copper absorption and metabolism. Fructose, or the fructose moiety of sucrose, fed to rats increased fecal and urinary excretion of copper (Lönnerdal 1996). The influence of fructose on copper balance in humans has not been well defined.

Based on studies in rats, ascorbic acid is thought to lower plasma and liver copper levels by reducing copper absorption, and the reduced copper absorption later stimulates copper absorption and depresses biliary excretion (Van den Berg et al. 1994). The decrease in absorption is caused by a reduction of cupric (Cu²⁺) ions to cuprous ions (Cu⁺), which are less well absorbed (Lönnerdal 1996). High levels of ascorbic acid might also decrease ceruloplasmin oxidase activity, although the overall effect of ascorbic acid on absorption and metabolism of copper in humans may be less than in animals (Lönnerdal 1996; Turnlund 1988). Administration of ascorbic acid with zinc at 1 mg per day in patients with Wilson disease showed no interaction of ascorbic acid and zinc, on copper balance, or ⁶⁵Cu absorption (Brewer et al. 1993a). Ascorbic acid had no detectable effect on the efficacy of zinc for copper-balance control in those patients.

Amino acids such as histidine and cysteine reduce copper absorption by forming complexes that are not well absorbed (Lönnerdal 1996). Histidine also enhances the inhibitory effects of zinc on copper absorption in rats (Wapnir and Balkman 1991). On the other hand, complexes of copper with other amino acids and organic acids might result in similar bioavailability to that of soluble copper sulfate (Wapnir 1998).

Based on studies to date, zinc is the primary mineral, and dietary element, which affects copper absorption. The effect of excess zinc on reducing copper absorption is well documented in a number of species (see summaries in Turnlund 1998; Wapnir 1998), and zinc has been used effectively in the treatment of Wilson disease (Hoogenraad et al. 1978; Hoogenraad et al. 1978; Hoogenraad et al. 1987; Brewer and Yuzbasiyan-Gurkan 1992; Brewer et al. 1994; Brewer et al. 1998). In pregnant rats, the consumption of diets containing high levels of zinc can result in fetal copper deficiency (Reinstate et al. 1984). In humans, the consumption of zinc supplements (50 mg per day for 6–8 weeks) can result in reductions in erythrocyte copper-zinc superoxide dismutase activity (Fisher et al. 1984; Yadrick et al. 1989; Johnson et al. 1998), suggesting that the chronic consumption of this level of zinc supplement could result in a condition of marginal copper status. Given the above possibility, the Institute of Medicine recommended that, at least for pregnant women, copper supplements (2 mg) should be provided to women when zinc supplements (25 mg or more) are given (IOM 1990).

In addition to zinc, various other minerals, such as iron, tin, calcium, phosphorus, cadmium, and molybdenum, interact with copper absorption and metabolism, although their effect compared with that of zinc is relatively minor or debatable in humans (Lönnerdal 1996; Turnlund 1988; Wapnir 1998). These minerals are cations that might compete for uptake in the digestive tract, thereby reducing absorption. These minerals might also affect copper utilization and excretion. For example, molybdenum has long been known to result in copper deficiency in ruminants but has little effect in nonruminants. Along with zinc for maintenance therapy, tetraethiomolybdate is now being used in the initial treatment of patients with the neurological or psychiatric form of Wilson disease. Tetraethiomolybdate acts by blocking absorption of copper when given with food and by complexing with serum copper when given separately from food (Brewer et al. 1996).

Similar to other metals, the bioavailability of lead, solubility and suspended particulates in water is likely to be a function of its chemical or surface sorbed form, solubility, and particle size (Davis et al. 1993; Ruby et al. 1996). As demonstrated for lead, soil and bioavailability can vary greatly, depending on the chemical and physical form (Ruby et al. in press). Copper acetate and sulfate are considerably more soluble and thus more bioavailable than cupric oxide (Johnson et al. 1998; Wapnir 1998). Copper sulfides (e.g., chalcocryolite), and other less-soluble minerals. Copper in soil and sediments also adsorbs strongly to soil components, such as clay minerals, hydrous iron, and manganese oxides (Tyler and McBride 1982), resulting in reduced solubility and mobility.

Conclusions

- Copper is an essential nutrient.
- Studies of absorption, transport and metabolism of copper have provided insights into the biochemical mechanisms for coping with copper deficiency and excess.
- The retention of copper from the diet is influenced by age, amount and form of copper in the diet, and genetic background.
- Bioavailability of copper varies with age and diet composition.
- The liver plays a central role in copper homeostasis by varying the excretion of copper into the bile for loss in the stool.
- The newly discovered chaperones for copper have provided insight into how copper ions in cells are guided to their target proteins.

Recommendations

- Studies are needed to elucidate mechanisms of copper absorption, distribution, and excretion in humans.
- Studies are needed on age-related changes in copper absorption and retention.
- Research should be conducted on the genetic basis of the absorption mechanism and on whether variation in absorption efficiency has a genetic basis.
- Research is needed to define extrahepatic processes for uptake and distribution.
- The ability of copper to induce proteins involved in its metabolism and transport should be investigated. Particular emphasis should be given to the investigation of metal response elements on copper transport proteins.
- Research is needed to determine the ontogeny of copper transporters.
- Research is needed to identify how the form of copper (i.e., valence state and complexed forms) influences absorption and distribution.

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