AMAZING ROLE OF SELENIUM IN BIOLOGICAL SYSTEM

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INTRODUCTION

Most essential micronutrients appear to play opposing roles in biological processes depending on their concentrations. While adequate dietary micronutrient intake is beneficial, deficient or excessive intake often leads to biological malfunctions resulting in the development of a wide diversity of diseases. Selenium, an essential micronutrient, recycles through the food chain and its concentration at each stage is basically determined by the amount remaining in the soil. The geographic distribution of selenium varies widely from selenium-deficient to selenium-rich regions. As with most essential micronutrients, selenium exhibits various biological functions according to its intake concentrations. At adequate concentrations, selenium exists as selenocysteine in the catalytic site of antioxidant proteins, including glutathione peroxidase and thioredoxin reductase, and is involved in the regulation of cellular redox status. A deficiency in selenium has been linked to many clinical...
selenium. Blind staggers is an generation of oxygen radicals that can be its cytotoxicity and anti, their target DNA sites. 

Selenium concentrations that prevent deficiency symptoms and are sufficient for exerting beneficial effects are close to those that lead to toxicity. Additionally, the safety margin of selenium dosage depends on various factors including age, gender, the chemical form present in the diet, transport capacity through cellular cytoplasmic membranes, efficiency of bioconversion from the inorganic to organic forms as well as exposure duration, frequency, and route. For these reasons, it is difficult to determine the exact safety levels of dietary selenium. Collectively, selenium seems to have bimodal roles in various biological processes. Several questions have been raised about how selenium simultaneously exerts beneficial and harmful effects under different circumstances. This article briefly discusses the diverse dose-dependent biological events associated with selenium and summarizes the biochemical roles of selenium in the regulation of cellular redox status focusing on the dual characteristics of selenium as an antioxidant and a pro-oxidant.

SELENIUM STATUS AND ITS BIOLOGICAL ACTIONS

Adequate selenium intake, approximately 100 μg daily in adult humans20 which does not exceed 1 mg, greatly reduces the rates of death from viruses such as coxsackie, hepatitis, and HIV. Moreover, adequate selenium intake reduces the incidence of cancer and is important for male reproduction12,14 These preventive and protective effects supposedly represent the antioxidant function of selenium-containing selenoproteins. Most of the selenoproteins, such as glutathione peroxidase (GPx), thioredoxin reductase (TR), and methionine-R-sulfoxide reductase 1 (MsrB1), serve as antioxidants.4,5,12 Furthermore, selenium can be used for protection against allergen-induced asthmatic inflammation1,6 and cancer therapy27 This chemopreventive effect is thought to result from the combined actions of the antioxidant functions of selenoproteins and the thiol modification of targeted proteins essential for allergen-mediated signaling pathways and cancer processing. An additional possible mechanism for the anti-carcinogenic effects of selenium is thought to be its cytotoxicity and anti-proliferative effects in malignant cells, due to oxidative stress resulting from thiol oxidation of cellular reductant molecules and oxygen radical production. Chronic exposure to high doses of dietary selenium in the range of milligrams per kilogram causes alkali diseases and blind staggers in livestock that graze in pasture rich soil8,9 Since plants absorb selenium more easily when grown in alkali soil, alkali disease frequently occurs in livestock that feed on selenium-tolerant plants with high selenium levels ranging from 5 to 50 milligram per kilogram of mass. Animals with this disease present abnormal symptoms, including hoof malformation, lameness, anemia, and stiffness. Blind staggers is an acute selenium poisoning disease appearing in cows and sheep, and is characterized by altered behavior, impaired vision, weight loss, ataxia, and respiratory failure. Symptoms of human selenium poisoning include loss of hair and nails, tooth decay, dermatitis, and gastroenteritis.1,8,10 Selenium toxicity develops in mammals with a daily intake of one milligram of selenium per kilogram of body weight, and in cultured cells exposed to micromolar concentrations of selenium1,13 Selenium toxicity is believed to be a result of low cellular redox status due to the oxidation of protein thiol groups and glutathione, and the generation of oxygen radicals. The toxic mechanism underlying oxidative damage by selenium is discussed in detail in the following section.

SELENIUM-INDUCED THIOL OXIDATION AND ROS GENERATION

Selenium is capable of negatively affecting cellular redox status by directly oxidizing thiols and indirectly generating reactive oxygen species (ROS), leading to a decreased reduction status in cells and cellular damage. Selenium reacts with essential thiol groups of proteins and cysteine residues of reduced glutathione (GSH) to form an intramolecular disulfide bond, a selenitrusulfide bond (S-Se-S), and a selenenylsulfide bond (S-Se)14. This can inactivate signaling molecules by oxidizing redox-sensitive cysteine residues present within the enzymatic active site of the catalytic domain. Selenium-induced oxidation of cysteine residues in transcription factors such as NF-κB and AP-1 leads to reduced binding affinity of these factors to their target DNA sites6,15,16 Furthermore, other redox-sensitive enzymes shown to be targeted for thiol oxidation by selenium include caspase-
3. Cdk2, protein kinase C, JNK, Na+-K+-dependent ATPase, glucocorticoid receptors, prostaglandin D synthase, human squalene monooxygenase, and mitochondrial proteins. Among these proteins, NF-κB, AP-1, caspase-3, Cdk2, protein kinase C and JNK are known to act as redox-dependent signal molecules.\[^{5,7,16}\] Therefore, deregulation of target molecules by selenium-induced thiol modification may be involved in disrupting various signal transduction pathways that control cell survival and apoptosis. ROS, which are chemically reactive molecules, including superoxide anions, hydrogen peroxides, hydroxyl radicals, and nitric oxide derivatives, are generated in all aerobic organisms through several pathways as summarized in Fig. 2. The superoxide anion (O\(^2-\)) is produced intracellularly by transferring electrons leaked from the electron transport chain in mitochondria and from NADPH cytochrome P450 reductase in the endoplasmic reticulum to oxygen, as well as by the action of several enzymes including NADPH oxidase, lipoxygenase, cyclooxygenase, flavoenzymes (e.g., xanthine oxidase), and uncoupled nitric oxide synthase.\[^{7}\] In particular, the superoxide anion is also formed endogenously by the reaction of selenium compounds such as selenite, selenium dioxide, diselenides, and selenocysteine, and with thiols such as reduced GSH or L-cysteine.\[^{3,11,48}\] The generation of superoxide anions by selenium has been confirmed by treating cells with exogenous GSH or selenite; this has also been observed by adding isolated mitochondria to selenium-containing compounds, including selenite, selenocysteine, selenocystamine, and selenodioxide, to GSH, or to both.\[^{1,4,10}\] The superoxide anion is rapidly converted to hydrogen peroxide (H\(_2\)O\(_2\)) via superoxide dismutate (SOD) followed by the conversion of hydrogen peroxide to a highly reactive hydroxyl radical (HO\(^•\)) in the presence of Fe\(^2+\) through the Fenton reaction. Subsequently, the hydroxyl radical reacts with NO\(^•\) to yield the more reactive species NO\(_2^•\) and HO\(^•\) through an ONOO- intermediate. Collectively, selenium induces a redox shift toward more oxidizing environments as follows: (i) selenium directly reacts with essential thiols, resulting in thiol oxidation, (ii) serial reductive reactions of selenium with GSH produce superoxide anions after which the superoxide anions generate additional reactive molecules (H\(_2\)O\(_2\), HO\(^•\), HO, ONOO-, NO\(_2^•\), and HO\(^•\)) via subsequent reactions, such as dismutation and Fenton reaction, and (iii) subsequent reaction of selenium with GSH results in GSH depletion from the export of cellular oxidized GSSG via a transporter. A more oxidizing environment produced by selenium can damage most biomolecules, DNA, proteins, and lipids, thus inducing cellular cytotoxicity.

**SELENIUM-INDUCED APOPTOSIS**

Numerous studies have suggested that selenium might be a preventive and effective anticancer agent for several human cancer cells including those of the prostate, colon, mammary, bladder, lung, liver, ovarian, oral, and bone or bone marrow.\[^{5,1-5}\] Selenium induces both ROS generation and oxidation, and cross-linking of protein cysteine residues, resulting in impaired protein function and apoptotic cell death. This element also inhibits cancer cell growth and induces cancer cell apoptosis in vitro at molar concentrations known to be toxic. Programmed cell death is marked by cellular morphological changes, including membrane blebbing, nuclear breakdown, chromatin condensation, and formation of apoptotic bodies that are readily eliminated by phagocytosis. This process also initiates activation of caspases, a cystinyl aspartate-specific protease that is stimulated in the process of mitochondrial-dependent or -independent apoptosis, and internucleosomal DNA fragmentation.\[^{5}\] In most cases, apoptosis induced by selenium is associated with typical features commonly observed in cells undergoing this process. Selenium-induced apoptosis in vascular endothelial cells, leukemia HL-60 cells, prostate DU-145 cancer cells, and murine monocyctic RAW264.7 cells activates caspase enzymes.\[^{3,5,7}\] DNA fragmentation during the selenium-induced apoptotic process has been reported in various human cancer cell lines including HT29 and SW480 (colonic carcinoma), HepG2 (hepatic carcinoma), A172 and T98G (glioma), and HL-60 (leukemia), as well as the murine monocyctic RAW264.7 cell line.\[^{3,6}\] Cells undergoing apoptosis due to selenium occasionally show characteristics of the mitochondrial-mediated apoptotic pathway, such as oxidative damage of mitochondrial protein thiol groups, a decrease in mitochondrial membrane potential, and mitochondrial release of cytochrome c, that can stimulate caspases.\[^{49,50}\] Cyclosporine A, an immunosuppressive mitochondrial membrane permeability transition inhibitor, blocks
mitochondrial swelling, and dithiothreitol restored the aggregation between intra- and inter-proteins by cross-linking mitochondrial protein thiol groups and mitochondrial swelling, implying that selenium-induced oxidative stress is involved in mitochondrial dysfunction. In summary, selenium-induced apoptosis is accompanied by ROS generation, oxidation of thiol groups in mitochondrial proteins, changes in mitochondrial membrane potential, cytochrome c release, caspase activation, and DNA fragmentation.

**BIOLOGICAL FUNCTIONS OF SELENIUM-CONTAINING ANTIOXIDANT PROTEINS**

Normal cellular oxygen metabolism in aerobic organisms leads to the generation of ROS. The disruption of intracellular redox equilibrium results in a state of oxidative stress that can easily damage biologically significant macromolecules. In order to scavenge harmful ROS, aerobic organisms have a wide variety of antioxidant enzymes to protect cells from oxidative stress. Among these, selenoproteins are primarily implicated in maintaining redox homeostasis and reversing apoptotic cell death induced by oxidative stressors, indicating that selenoproteins may act as a safeguard against oxidant-induced toxicity in cells.

Based on bioinformatic data, it has recently been reported that there are 25 selenoproteins in humans and 24 in rodents. Selenium is incorporated into these proteins as the 21st amino acid selenocysteine encoded by a UGA codon normally recognized as a stop codon within the mRNA open reading frame. Selenoproteins with known functions include the antioxidants glutathione peroxidase (GPx), thioredoxin reductase (TR), and methionine-R-sulfoxide reductase 1 (MrSB1). These antioxidant selenoproteins serve as central components for the regulation of cellular redox status by maintaining cysteine residues of redox-sensitive proteins in the reduced state and promoting normal cellular functions. GPx catalyzes the reduction of hydrogen peroxide to water by glutathione, TR, with a selenocysteine residue in the conserved C-terminal sequence glycine-cysteine-selenocysteine-glycine, is known to reduce thioredoxin by using NADPH and as a result maintains the reduced state of intracellular proteins. MrSB1, in which the active site contains a selenocysteine, contributes to the reduction of methionine sulfoxides in many proteins. Together, these enzymes help rescue oxidatively-damaged proteins and perform housekeeping redox functions that allow cells to maintain a favorable intracellular redox status.

**CONCLUSIONS**

Although it is difficult to determine the difference between beneficial and toxic selenium levels, the current recommended daily intake of selenium to maintain normal cellular functions in adult humans is approximately 100 μg. Selenium can induce both beneficial and harmful cellular responses, and serves as both an oxidant and antioxidant. A severely selenium-deficient diet leads to oxidative stress due to decreased levels of antioxidant selenoproteins such as GPx, TR, and MsRB1. On the other hand, excessive dietary intake of selenium induces a redox shift towards a more oxidizing cellular environment by directly oxidizing and cross-linking protein thiol groups and indirectly generating oxygen radicals, resulting in apoptotic cell death. Selenium cytotoxicity is now believed to be caused by oxidative stress. The dual functions of selenium essential for its beneficial antioxidant and toxic pro-oxidant properties make this element useful for decreasing the incidences of selenium-deficiency disorders as well as therapies for treating and preventing cancer.

**REFERENCES**