

Is carbonyl detoxification an important anti-aging process during sleep?

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Summary Organisms living on the earth may undergo inevitable toxification by biological 'garbage', a variety of bio-metabolites. Such garbage includes a particularly large number of toxic carbonyls, such as α,β -unsaturated carbonyls created by free radicals, glycation, and other post-translational side-reactions during various stresses and diseases. The accumulation of these toxic substances and their crosslinking products leads to the formation of different age pigments, such as lipofuscin, lens cataracts, and crosslinked collagen. The diurnal fluctuation in the concentration of the pineal gland hormone, melatonin, may be responsible for the 'cleaning activities' that reverse the covalently-bound semi-toxified proteins and nucleic acids. This toxification-cleaning cycle may explain the biochemical necessity for sleep of human and animals during aging. © 2000 Harcourt Publishers Ltd

CARBONYL TOXIFICATION

An important characteristic of most stresses, diseases and aging processes is the consistent increase in the concentration of various unsaturated carbonyls, mainly toxic aldehydes, in almost all tissues and body fluids (1–5). Such carbonyls, originating from major biological macromolecules including proteins, lipids, carbohydrates, and nucleic acids, are produced mainly by different post-translational side-reactions, especially oxidative stress and glycation (4–8). The high reactivity of carbonyls, particularly α,β -unsaturated carbonyls, underlies their toxicity by crosslinking or primarily binding to important biomolecules such as structural and functional proteins and nucleic acids. This type of carbonyl-amino crosslinking is now considered to be a key process in the formation of ceroid, lipofuscin (intracellular age pigments), lens cataracts, and crosslinked collagens and elastins (9). It is

also a critical factor in the decline of cell proliferation and various cell membrane functions, in atherosclerosis and amyloid formation, and in many other degenerative alterations during aging (6,7,10).

The amount of biochemically detectable carbonyls in different body constituents, best represented by thiobarbituric acid reactive substances (TBARS), is the most used index of the steady-state levels of the reactive and toxic carbonyls of an organism (11). Although the amount of protein carbonyls is an important index of aging, the high variability of results and the time-consuming nature of analytic techniques hinder accurate determination (11–13). On the other hand, food is another important source of carbonyls. Since most foods and beverages are unprotected during processing and cooking, or only partly protected during storage, they may contain a large amount of reactive carbonyls, which provides an unlimited and unavoidable source of such (unhealthy) substances and their derivatives. In vivo, the concentration of α,β -unsaturated carbonyls, best studied as malondialdehyde (MDA) and 4-hydroxyalkenals (4HA), is under active control of different biological enzymes, such as carbonyl dehydrogenases (14). In mammals, a first line of defense against toxic unsaturated carbonyls from food

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takes place in the digestive system, where hepatic carbonyl dehydrogenases convert carbonyls to other compounds and eventually either to carbon dioxide and water, or to soluble derivatives which are excreted in the urine (6).

REVERSIBILITY OF CARBONYL TOXIFICATION

It is known that the primary reactions between α,β -unsaturated carbonyls and amino compounds are reversible (Fig. 1). When organisms contain relatively high concentrations of toxic carbonyls due to daily activities, functional and structural proteins are likely to react with these substances to form primarily semi-bound protein-carbonyl compounds (6), e.g. amino-propanal, due to 1:1 MDA-protein reactions (Fig. 1). Although heavy stresses lead to the formation of large amounts of toxic carbonyls (as does the ingestion of oxidized food or a high caloric diet), biological defending substances, such as carbonyl dehydrogenases, glutathione and glutathione transferase (6,14,15) all work intensively to reduce the concentration of these toxic components and to keep the carbonyl binding and subsequent crosslinking at reasonably low levels.

POTENTIAL OF CARBONYL TOXIFICATION

Daily activities, particularly intense exercise, staying awake overnight, diseases, and major surgical operations all lead to large increases in the concentration of toxic carbonyls (5–7), thereby driving the reaction equilibrium primarily towards the formation of 1:1 amino-carbonyl compounds, although these are unstable and reversible.

The detected concentration of serum TBARS mainly represents a balance between the formation of carbonyls and their elimination by defense mechanisms (probably also including the fast protection of thiol compounds in addition to enzymatic defense), which may mirror the real reaction situation between carbonyls and amino compounds *in vivo*. The actual concentration of amino compounds is much higher than the concentration of α,β -unsaturated carbonyls (16), e.g. the total amount of free amino acids is as high as 80 mM, although proteins are found to be more readily modified by MDA than free amino acids under physiological conditions (6). Since the reaction potential between α,β -unsaturated carbonyls and proteins, such as enzymes, is fairly low, the possibility of crosslinking should be even lower. Hence, the principal reaction between MDA and functional and structural proteins would be an amino-propanal reaction, rather than a crosslinking reaction, to form amino-imino-propen structures. Although the primary reaction is slow and actively reversed by several defense mechanisms, the accumulation of amino-carbonyls is progressive and affects the normal body functions, probably until the brain cannot tolerate further increase, and signals sleep (by melatonin secretion?).

SLEEP, MELATONIN AND CARBONYL DETOXIFICATION

The observation that melatonin secretion helps to keep down TBARS levels during various oxidative stresses, in addition to its known beneficial effects on sleep as well as endocrine and immune functions (17,18), leads me to raise a very interesting question: how does melatonin interfere with oxidative stress when it is secreted mostly only during sleep? Although melatonin has been shown to prevent damages from oxygen free radicals, it has also been shown that administration of melatonin often causes TBARS levels lower than in control samples (19,20) which may suggest a detoxification function of melatonin. Furthermore, the indole structure of melatonin is similar to the indole used in MDA+4HA measurement in the LPO-586 assay. It is logically unsound to suppose that melatonin has been evolved mainly as a free-radical scavenger, because oxygen free radicals, as well as most hydroperoxides, react so quickly, that they will almost immediately attack any biomaterials in their vicinity (21,22). After the rapid disappearance of oxygen radicals, only various toxic carbonyls and their derivatives remain in tissues and body fluids (3–5). Therefore, one function of melatonin may be to interact with the accumulated amino-propanal-like products produced by various stresses, in order to remove them before they form stable amino-imino-propen crosslinks. When the concentration of melatonin reaches a high level during

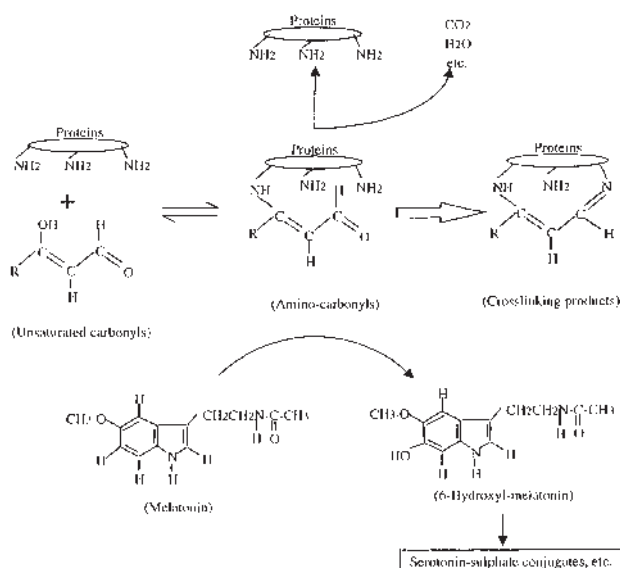


Fig. 1 Mechanisms of carbonyl toxification due to stresses, and proposed melatonin detoxification during sleep.

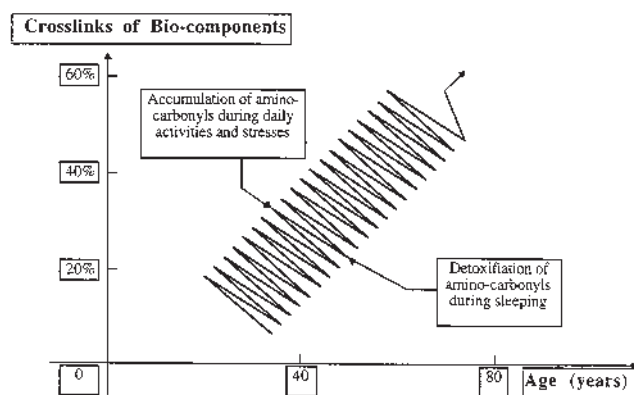


Fig. 2 Carbonyl detoxification during sleep and its hypothesized relation to aging processes.

deep sleep, the carbonyl-amino reaction may be reversed causing a removal of toxic compounds at the expense of the hydroxylation of melatonin (23,24). A sound sleep may thus lead to an efficient clearance of carbonyl complexes, although a 100% clearance may not be attained, especially not in tissues such as collagen where less melatonin is delivered and where crosslinks seem to occur at an average speed of 0.001% of the total amount of structural protein per day in humans. When the brain is free of, or has only a very low level of such amino-carbonyl garbage, we wake up and feel refreshed and alert again, and we are ready to face a new day of activity, during which we will once again accumulate new amounts of noxious substances. This 'contamination-cleaning' cycle repeats every day even as undegradable crosslinked materials gradually accumulate, eventually becoming abundant and 'mature' in the tissues. Such fluorescent, age-related accumulates that are found either intracellularly (lipofuscin) or extracellularly are known as age pigments. When these crosslinked materials become abundant, our bodies do not function well and we enter into the aging period. Figure 2 shows a scheme for garbage accumulation, clearance during sleep and the aging process. Genetic factors are found to regulate the defending, cleaning and repairing systems (e.g. genes for superoxide dismutase, catalase and glutathione transferase, ref. 6,25,26). Differences in the efficiency of these systems would determine the life-span of different species.

This hypothesis, which links sleep and aging, is mainly based on the understanding of oxidative stress, glycation and age pigment formation. It may appear to be an oversimplification of such extremely complicated biological processes as sleep and aging (27,28). However, in the author's opinion, a sound aging theory may eventually be required to provide a comprehensive explanation to the crucial daily need for sleep.

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