New Vistas on Oxidative Damage and Aging

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Abstract: Age-associated rises in oxidative damage are assumed to be a central phenomenon of aging. Their attenuation is an aim for both healthy aging and life extension. This review intends to critically discuss the potential of anti-oxidant actions, but even more to direct the attention to the modes of radical avoidance and to regulatory networks involved. Mitochondria seem to play a decisive role in radical formation and cellular decline. Avoidance and repair of disruptions in the electron transport chain reduce electron leakage and, thus, oxidative damage. Several low molecular weight compounds, such as melatonin, its metabolite N^1 -acetyl-5-methoxykynuramine, resveratrol, α -lipoic acid, and various mitochondrially targeted nitrones are capable of supporting mitochondrial electron flux. Some of them have been successfully used for extending the lifespan of experimental animals. Importantly, chemopreventive effects of these substances against cancer development should not be confused with a slowing of the aging process. We also focus on connections between these compounds and mitochondrial biogenesis, including the roles of sirtuins and signaling via peroxisome proliferator-activated receptor- γ coactivator-1 α , the participation of the circadian oscillator system in radical avoidance, as well as the potentially beneficial or detrimental effects of NO, as either a regulator or a source of mitochondrial dysfunction. Especially in the central nervous system, anti-excitatory actions by melatonin, kynurenic acid and theanine are discussed, which seem to prevent calcium overload that results in mitochondrial dysfunction. New findings on direct binding of melatonin to the amphipathic ramp of Complex I may indicate an additional regulatory role in the avoidance of electron leakage.

Keywords: Aging, anti-oxidants, electron leakage, melatonin, mitochondria, PGC-1α, sirtuins.

INTRODUCTION

Aging is a complex phenomenon, in which slow, lingering processes of steady declines in metabolism and physiological functions are superimposed by single events of deterioration, which are caused by diseases and followed by secondary impairments. In the worst case, infarction, stroke, renal failure or cancer, and sometimes also the medicinal treatment, represent profound individual catastrophies, which impair the function of organs and cells and, thereby, contribute to the acceleration of the more continual processes of aging, even if the primary treatment is per se successful. However, less severe diseases may also affect the health state in a way to make aging more rapidly progressing. Much of theoretical work has been published on the mechanisms of aging. Although the increase of age-related morbidity has some explanatory plausibility for the consequences of the step-wise events of impairment, it does not seem to fully comprise the causes and mechanisms of the more steady functional declines. This aspect is multi-faceted. It includes the maintenance of immunological functions for safeguarding the individual's resistance to germs and viruses [1,2], the preservation of healthy endothelial surfaces and also efficiently working mitochondria. In part, the organismal decline seems to be associated with reduced capacities for

cell division because of telomer shortening, but even more with progressing losses of adult stem cells, which leads to a reduced capacity for tissue repair. However, this cannot be equally valid for every organ, because of enormous differences in the replacement of cells and the survival of partially dysfunctional cells. Despite this obvious diversity, the key to the slow processes of impairment has to be sought in the metabolism of an organism. This view is supported by the existence of aging-suppressor genes, such as those for sirtuins [3-8] and klothos [9-14], which act as cellular energy sensors and regulators of cellular and organellar functions.

A common theme to many of the age-related dysfunctions is the damage by reactive oxygen and nitrogen species (ROS and RNS). Different sources of these potentially damaging intermediates are of relevance. Oxidants induced especially by UV light are important to the skin. Macrophages and other oxidant-producing leukocytes can contribute to the damage of epithelial surfaces, and atherosclerotic incrustations lead to further damage by nitric oxide (•NO) derived free radicals, especially radical pairs formed from peroxynitrous acid (ONOOH \rightarrow ·NO₂ and ·OH, i.e., nitrogen dioxide and hydroxyl radical) or from the peroxynitrite-CO₂ adduct (ONOOCO₂⁻ \rightarrow ·NO₂ and CO₃·⁻, i.e., nitrogen dioxide and carbonate radical) [15,16]. In this scenario, superoxide anions $(O_2 \cdot \overline{})$ required for peroxynitrite formation are generated at high rates in cells compromised by insufficient oxygen supply, whereas 'NO is produced in the partially unsuccessful attempts of causing vasodilation, and CO₂ is present in supranormal concentrations because of the

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reduced blood flow [16]. If not formed in the oxidative burst of an activated leukocyte, the major sources of O_2 . are usually mitochondria. Electrons are leaking, mostly at Complexes I and III, when secondary bottlenecks transiently impede electron flux, e.g., at Complex IV under conditions of insufficient oxygen supply [17,18]. Also in other scenarios, electron leakage from mitochondria is a main cause of oxidant formation. $O_2^{\bullet^-}$ is converted to more destructive radicals, either via peroxynitrite or dismutation to O_2 and H₂O₂, this latter compound being another source of the devastating \cdot OH. Moreover, CO₂ is abundantly available in the mitochondrial matrix, where it is produced. Therefore, elevated mitochondrial generation of O_2 . by electron leakage from the electron transport chain is likely to result in high rates of ONOOCO₂⁻ formation and the radicals formed by its decay.

Mitochondrial dysfunction obviously occurs during normal aging, but seems to be cell type- or tissue-specific. It can even vary within a cell, as observed in cardiomyocytes, in which subsarcolemmal mitochondria did not show profound signs of dysfunction in the course of aging, whereas interfibrillary mitochondria exhibited severe decreases in Complex III and IV activities and increased electron leakage, mainly from the Qo site of Complex III. However, antioxidative protection especially of cardiolipin, which is required for the function of Complexes III and IV, largely restored interfibrillary mitochondrial activities [19-25].

The involvement of mitochondria in the aging processes has already been assumed in the earlier free radical theory of aging [26,27]. In its more recent versions, it has developed towards a model centered on mitochondrial damage and dysfunction [28-30]. The paradigm has meanwhile changed from the premises that these organelles are both major source and target of destructing radicals to the new perspectives of metabolism modification and avoidance of electron leakage. Therefore, the new questions to be answered are those of how mitochondrial function can be supported and maintained in the long run, how electron dissipation can be avoided, and which molecules may be suitable for this purpose.

OXIDATIVE DAMAGE DURING AGING AND ITS RELATIONSHIPS TO CHRONOBIOLOGY

Enhanced oxidative damage of membrane lipids and biological macromolecules during aging, including the consequences for cell death, are amply documented [31-48]. In the absence of experimentally induced radical formation, e.g., for testing the rises in vulnerability, it is frequently difficult to distinguish between damage as a cause, a cophenomenon and a consequence of aging processes. Presumably, all of these possibilities have to be equally considered. Diseases may not only weaken the body's resistance, but also lead to enhanced radical generation. The causes may be either cell damage and death or oxidants formed in the organism's defense responses. Rises in oxidant formation, as occurring in the course of aging, may increase the dysfunction of structures relevant to free-radical generation, in particular, mitochondria, so that a vicious cycle between cause and consequence is initiated.

It is not the aim of this review to repeat discussions on the numerous findings concerning progressive oxidative damage during aging, nor to refer to the countless attempts of pharmacological interventions. However, some aspects shall come into focus which are, in the broadest sense, bridging chronobiology and oxidant generation.

Damage to mitochondria, a major source of oxidants, during aging is a long-known phenomenon [19-28,44,49]. This damage has been found to progress more rapidly in senescence-accelerated mice [50]. However, caution is due with regard to interpretations based on the original freeradical theory of aging, including the classic assumptions of a purely mitochondrial vicious cycle. It has been criticized with good reason that many of such conclusions are derived from pharmacological experiments, which are far from real life during normal aging [51]. Moreover, investigations in mitochondrial DNA (mtDNA) mutator mice revealed that, despite amplified mitochondrial mutations in the course of aging, no increased ROS production was demonstrable [52]. This should mean that the original assumption of mitochondrial mutations as a source of enhanced ROS generation is not supported. Moreover, it seems important to dismiss the idea that mtDNA were particularly vulnerable because of a lacking chromatin structure. In fact, mtDNA is densely loaded with proteins, in particular, the mitochondrial transcription factor A (TFAM), a high mobility group (HMG) related molecule, which also fulfills functions in nucleoid structure, damage sensing and mitochondrial replication [53-56]. Several other proteins, including anti-oxidant enzymes, are bound to the mtDNA as well [56]. Nevertheless, the lack of a clearly demonstrable relationship between mitochondrial mutations and radical generation does not immediately imply that a vicious cycle between mitochondrial dysfunction and free radical production does not exist. It only does not include the damage to mtDNA as a necessary component. It rather seems that enhanced radical formation causes damage to the electron transport chain (ETC). This leads, in turn, to enhanced electron leakage and, thus, oxidants affecting both the ETC and, upon dissipation into other compartments, the rest of the cell. Therefore, the vicious cycle should be understood as a self-enhancing, in the cybernetic sense "positive", feedback loop between electron leakage and the causes of further leakage from the damaged ETC.

The relationship between radical formation to chronobiology has several aspects, (i) age-associated declines in the amplitude of circadian oscillations, (ii) dysphasing, as occurring by progressive shortening of the spontaneous circadian period and leading to age-related advances of the clock, and (iii) the steady decline in the nocturnal rhythmic secretion of melatonin.

The importance of circadian rhythms in aging may not be a matter of general awaress. Nevertheless, the evidence for its relevance is convincing. An important, but frequently overlooked finding was obtained in aged Syrian hamsters, which received transplants of suprachiasmatic nuclei (SCN) [57], the hypothalamic structures which represent the circadian pacemaker. Although such a transplant cannot form a structure fully equivalent to the recipient's own SCN, it did not only restore the circadian time pattern of locomotor activity, which had previously become irregular and partially disrupted in the aged animals, but it also extended the lifespan of the recipients [57].

Relationships between circadian rhythms, oxidants and anti-oxidative protection are obvious under various aspects. Oxidant formation is rhythmic, already as a result of rhythms in metabolism, in light-exposed species also as a consequence of radiation. Organisms have adapted to this periodic challenge by generating circadian oscillations of anti-oxidant enzymes and concentrations of low molecular weight antioxidants (LMWAs), findings that were obtained in many different species from phylogenetically distant taxa, including unicellular algae, invertebrate and vertebrate animals [58]. In Drosophila melanogaster, a rosy mutant (ry^{506}) unable to synthesize the endogenous LMWA urate showed strongly elevated levels, enhanced circadian amplitudes and alterations in the temporal pattern of protein carbonyl, an oxidative protein modification [59]. The relevance of circadian rhythms for attenuating oxidative damage became even more evident in animals carrying mutations in clock genes of the circadian core oscillator. In the arrythmic per^{0} mutants of D. melanogaster, protein carbonyl did not only show the absence of a circadian pattern, but was, addi-tionally, markedly elevated [59]. Moreover, per^0 flies exhibited an elevated susceptibility to exogenously induced oxidative stress [60]. Interestingly, increased oxidative damage to proteins was also observed in the short period mutant per^s, when the animals were maintained under a light/dark cycle of 12:12 hours, i.e., a Zeitgeber period substantially deviating from the spontaneous period of the *per^s* animals [59]. A corresponding observation was made in another short period mutant of a different organism, the tau mutant of the Syrian hamster. In this case, both protein carbonyl and lipid peroxidation were strongly elevated in the Harderian gland [61], an organ that is particularly vulnerable to oxidative damage [62]. Collectively, these findings indicate that not only the mere existence of a circadian rhythm is decisive for avoiding damage by oxidants, but also an appropriate phasing that allows detoxification of free radicals in those circadian phases when enhanced anti-oxidative protection is required, e.g., in conjunction with elevated metabolism. Moreover, findings in *tau* hamsters revealed a relationship to lifespan. Again, the aspect of appropriate rhythmic coordination comes into focus: Homozygous tau hamsters showed only a moderate statistical tendency of reduced lifetime (mean survival 15.8 vs. 17.5 months in wild-type], but heterozygous *tau* animals had a substantially shortened life span (10.9 months) [57]. This may be taken as a sign of internal malcoordination, due to differently acting oscillators.

Enhanced oxidative damage because of dysphased rhythms may not only occur in clock mutants, but also result from circadian phase shifts, as occurring in shift work or after transmeridian flights. In fact, repeated phase shifts were already shown decades ago to reduce the lifespan of flies [63]. Similar results were obtained in cardiomyopathic hamsters [64]. Shift work as a source of health problems is beyond any doubt [65-70] and may also affect longevity, at least in a subpopulation of individuals. However, this statement should not be misunderstood in terms of an oversimplification, since the impairment by shift work

cannot be reduced to oxidative stress. Many other factors, such as sleep difficulties, associated problems of working conditions, and the role of circadian clock genes in preventing cancer contribute to statistically demonstrable health problems. The relationship between circadian dysregulation and cancer has received substantial support during the last years, After the original discovery that mice carrying a mutation in the per2 gene are cancer-prone [71,72], numerous studies have now shown that several core oscillator proteins, including their variants, such as PER1, PER2, PER3, CRY1, CRY2, CLK, BMAL1 and CK1E are downregulated in various forms of cancer, and/or act as cancer suppressor genes and sensors of DNA damage [73-82]. Although it is far from being clear to what extent oxidative stress may be involved in this phenomenology, and although down-regulation of clock genes may be largely due to hypermethylation in the respective promoters [73,83,84], the consequence of the circadian dysregulation is unavoidably an insufficient management of redox control, resulting in increased oxidative damage and higher probability of mutations.

Another aspect of circadian rhythmicity in relation to redox balance concerns melatonin. When thoroughly analyzed, this turns out to be a highly multi-faceted host of actions. In its role as a hormone of the pineal gland, melatonin acts as a regulator of numerous circadian functions. In its multiple roles as hormone, tissue factor, paracoid and autocoid [85], it also acts as an agent protecting against oxidative damage [86-95]. Various modes of action have been identified at different levels, from anti-excitatory/antiexcitotoxic and anti-inflammatory effects to regulation of anti- and pro-oxidative enzymes and direct radical detoxification, as summarized elsewhere [91,96]. Moreover, melatonin has profound actions on mitochondria and, thereby, strongly influences electron flux and leakage [17,18,58,97-105]. Importantly, such findings were not only obtained under experimental conditions designed to challenge mitochondrial functions, but also in otherwise untreated animals. Beneficial effects on mitochondrial electron flux, respiratory efficacy and avoidance of damage to the ETC were especially observed in aged animals [99-101,106-112].

In the relationship between melatonin and aging, a crucial observation is the age-dependent decline in the nocturnal peak of melatonin secretion by the pineal gland. Although this is not equally pronounced in every individual, it is clearly demonstrable on a statistical basis and can be described as a dramatic change occurring in the majority of subjects [113-123]. The waning nocturnal melatonin peak has been interpreted as a consequence of general circadian deterioration, largely caused by progressing dysfunction of the pacemaker [124]. This may not be the only possible reason, since the onset of the decrease is already observed in middle-aged subjects [118,119,121], in which deteriorations of pacemaker function are not yet obvious. Nevertheless, the changes in melatonin may strongly contribute to a weakening of the intra-organismal circadian organization and to losses in the efficacy of protective functions, at the levels of anti-oxidative defense, the immune system and mitochondria as well.

RADICAL DETOXIFICATION, RADICAL AVOI-DANCE AND RESISTANCE TO OXIDATIVE STRESS

The classic view of anti-oxidative protection is based on the assumption that radicals already formed have to be detoxified as efficiently as possible. This concept was supported by numerous pharmacological investigations, in which oxidants or oxidant-generating compounds were administered and their deleterious effects antagonized by usually unphysiologically high quantities of radical scavengers. Although such approaches are not without value, they may tell little about the processes underlying the aging process. They may be, in part, helpful in the combat against chronic inflammation associated with aging, but the aging process *per se*, in terms of a steady, lingering decline may remain unaccessible. In fact, the idea of prolonging life by overexpression of anti-oxidant enzymes has been only of limited success, and some of the initial reports on efficacy were later corrected or withdrawn. Moreover, knockout mice for major anti-oxidant enzymes, including those located in mitochondria, frequently failed to substantially decrease longevity, although oxidative damage to DNA and the incidence of cancer were enhanced [125-128]. Radical detoxification by scavengers and anti-oxidant enzymes is certainly a necessity, and a deficiency of anti-oxidants, e.g., by malnutrition, will cause the development of diseases, but, in relation to aging, this does not seem to touch the core of the problem. A possible access to unveiling the relevant processes may be sought in the comparison between longlived and short-lived species of approximately same size. In the past, differences in the efficacy of the radical-detoxifying enzymes have been in the focus and several reports have, in fact, pointed into this direction. Although this may contribute to longevity, it does not necessarily identify the crucial differences. In a recent study, muroid species of similar size, but differing considerably in lifespan were compared, the white-footed mouse (Peromyscus leucopus) and deer mouse (Peromyscus maniculatus) with a maximal lifespan of up to 8 years and the house mouse (Mus musculus) having a maximal lifespan of 3.5 years [129]. Although Peromyscus was shown to possess a more efficient anti-oxidant defense than Mus, two additional features became evident, namely, a superior DNA repair system, and a lower rate of mitochondrial oxidant formation. A more pronounced oxidative stress resistance was also reported for two other small, longlived mammals, the naked mole rat (*Heterocephalus glaber*) and the little brown bat (Myotis lucifugus) [130].

The triad of anti-oxidative defense, radical avoidance and oxidative stress resistance may be a key to especially healthy aging, but the relative contribution to lifetime remains to be clarified. In the cases of anti-oxidative protection and stress resistance, it is of utmost importance to distinguish between chemopreventive effects of anti-oxidants and changes in the aging process itself. This is of particular relevance in the use of inbred laboratory mouse strains, in which mortality from cancer prevails and, therefore, apparent changes in average lifespan are often caused by a reduction or delay of carcinogenesis [131]. However, in conjunction with the chemopreventive effects, a more youthful appearence of aging mice has been observed when compounds such as melatonin [29,132] or resveratrol [133] were administered, even in those investigations in which no significant effects on lifespan were demonstrated. Therefore, natural compounds or drugs that reduce oxidative damage can be of value for healthy aging, even in the absence of life extension.

However, these considerations clearly indicate the premier importance of selecting a suitable object for aging studies. In organisms or strains in which cancer is not a major cause of death and, therefore, chemoprevention not decisive, effects on lifespan may be entirely different. This is of particular relevance in the case of the highly pleiotropic agent melatonin, which is not only a pineal hormone but also present in various tissues, often at much higher quantities than in the pineal gland [85,93,96,132,134]. The selection of senescence-accelerated mice, SAMP8, which can be also compared with SAMR1, a normally aging strain carrying the same genetic background [135], can lead to results substantially different from those obtained in other strains. In SAMP8 mice, melatonin caused a clearly demonstrable prolongation of life [111]. A main question raised by the studies on SAMP8 is that of the reasons for the more rapid aging. These may not be identical with the normal causes of aging, although such an idea is highly attractive. As an example, one should remember progerias caused by mutations in lamin genes [136,137]. No one would assume that breakdown of the nuclear lamina could be a major cause of normal aging, despite the enormous acceleration of aging in these laminopathies. In SAMP8, a higher sensitivity towards oxidative damage was demonstrated [138]. Therefore, anti-oxidant treatment should be effective, including the possibility of reducing oxidative damage by radical avoidance, but the question remains whether it counteracts the normal aging process or "only" a cause of accelerated aging. Nevertheless, as long as this question has not been decided, studies on senescence-prone mice remain of high gerontological interest.

SAMP8 mice may also provide substantial information to a more comprehensive understanding of aging since they exhibit more rapidly developing changes in mitochondrial metabolism, which are clearly reduced by melatonin. This aspect will not be discussed here in any detail, because of a parallel publication by another group on this issue. As outlined in other reviews dealing with mitochondrial function and melatonin [17,18,58], numerous results collectively show that changes observed at advanced age of the senescence-accelerated animals are reverted or attenuated by this indoleamine, such as, reductions in the respiratory control index (RCI); in state 3 respiration; in dinitrophenoluncoupled respiration, which reflects to some extent the respiratory capacity; in the ADP/oxygen ratio; in the ATP level; in the activities of Complexes I and IV. However, it is necessary to distinguish in studies on SAMP8 mice between processes occurring during aging in otherwise unchallenged animals and those intending to identify alterations in vulnerability, inducibility of apoptosis, and effects of inflammation. Though all these latter changes may be relevant in a specific context, they may not properly reflect the aging process per se.

Although caution is due concerning the very meaning of experimental interventions in senescence-accelerated mice, these animals provide valuable hints on radical avoidance, since they show age-related mitochondrial malfunctions in electron flux and energy efficiency, which are associated



Fig. (1). The vulnerable sites of the mitochondrial electron transport chain (ETC) and the main sites of electron leakage. Arrows in pink color: electron flux through the ETC; red arrows: superoxide formation by electron donation to O_2 ; flashes and text in violet color: effects leading to interruption of the electron transport and enhanced electron dissipation. The amphipathic ramp of Complex I is particularly vulnerable to modifications because of its exposure to reactive intermediates arriving from the aqueous matrix. Abbreviations: Com = Complex; CoQ = coenzyme Q; Cyt C = cytochrome c.

with enhanced electron dissipation and, thus, free radical formation [17,18]. In this context, it seems worth-while to look at the changes which naturally occur during aging in the absence of pharmacological treatments. As already outlined in the Introduction, rises in mitochondrial free-radical formation are observed during aging, which are related to impairments of electron flux and lead to electron dissipation. The causes for transient, local blockades of the ETC (Fig. 1) may be mainly sought in (i) more frequent interactions of elevated NO with irons in the ETC complexes; (ii) destructive actions of peroxynitrite-derived radicals, again a consequence of elevated NO; (iii) oxidative or nitrosative modifications at the major sites of electron leakage, i.e., (a) the especially vulnerable amphipathic ramp of Complex I and (b) Complex III, in which the interruption of the intramonomer electron transfer between the two $b_{\rm L}$ hemes at the Qo site seems to be decisive; (iv) cardiolipin peroxidation leading to dysfunction of Complexes III and IV; and (v) relative O₂ deficiency that may result from atherosclerosis and which causes a secondary bottleneck of electron flux at Complex IV [17,18]. All these changes can occur concomitantly. A realistic picture of electron dissipation is only possible by considering its dynamics. Electron flux is by no means a steady and uniform homeostatic process occurring at more or less constant rates in a given metabolic situation. Recent investigations have shown that electron leakage happens in a flash-like fashion [139]. Therefore, bursts of superoxide anions, a source of H2O2- and peroxynitritederived radicals, can be expected to be much stronger at reduced electron flux capacity, whereas basal rates should be

less affected. The lesson from these considerations should be that maintenance of electron flux capacity is decisive for reducing electron dissipation and, thus, radical formation. With some likelihood, the management of radical avoidance by safeguarding electron flux may be much more efficient than the detoxification of radicals already formed [17,18,58,91].

MECHANISMS OF RADICAL AVOIDANCE AND POSSIBLE INTERVENTIONS

Numerous causes of enhanced radical generation can be identified, also beyond toxicological and inflammationrelated effects. Already within the frame of physiological, lifestyle-dependent, genetically or age-related pathophysiological processes, elevated radical formation is possible. The chronobiological influence on oxidative damage has already been discussed. Correspondingly, the avoidance of unnecessary phase shifts and a chronohygiene in favor of sustained circadian amplitudes should be of advantage. The frequently observed age-related declines in SCN function and in nocturnal melatonin secretion may be seen as a possibility for interventions that could lead to reduced radical formation. Circadian amplitudes may be enhanced by repeated exposures to strong Zeitgeber signals, such as bright light in the morning, and/or melatonin, alternately other melatoninergic agonists, at fixed times in the evening. Additionally, these treatments favor appropriate circadian phasing relative to external time and, in the case of melatoninergic agonists, promote sleep onset in humans, which may be also of advantage for aged individuals [140]. As far as a substitution therapy is intended in cases of strongly reduced nocturnal melatonin, the natural hormone may be given daily, provided that no contraindications exist. If only enhancements of the circadian amplitude are required, melatonin or one of its synthetic analogs may be used for a few days, a procedure which should be repeated at intervals as individually required. In the relation to antioxidative protection, the use of melatonin instead of its analogs may be recommendable because of its additional beneficial effects, but this has to be distinguished from the chronobiotic actions concerning radical avoidance.

Reduced damage by free radicals is especially desirable in the central nervous system and may contribute to the



Fig. (2). Radical avoidance: multiple sites of intervention by several protective compounds. A simplified overview of a complex network. \downarrow : Stimulation or chemical reaction; \perp : inhibition. Colors of text and arrows, black: processes leading to enhanced formation of free radicals; brown: actions of resveratrol; green: actions of melatonin and its metabolite AMK; blue: main action of kynurenic acid; olive: main action of L-theanine; pink: responses to metabolic sensing; red: pathways influenced by multiple factors. Abbreviations: AMK = N^1 -acetyl-5-methoxykynuramine; AMPK = adenosine 5'-monophosphate-activated protein kinase; Com = complex (of respiratory chain); ETC = electron transport chain; FenR = Fenton reaction; GSH = reduced glutathione; PGC-1 α = peroxisome proliferator-activated receptor- γ coactivator-1 α ; PPAR γ = peroxisome proliferator-activated receptor- γ ; SIRT1 = sirtuin-1; SODs = superoxide dismutases.

prevention or delay of neurodegeneration. In this context, the avoidance of neuronal overexcitation, in the worst case excitotoxicity, is of particular importance (Fig. 2). Longlasting neuronal excitation is usually associated with elevated calcium influx, leads to extended periods of high NO levels and, thus, peroxynitrite-induced damage. It may cause mitochondrial calcium overload, with the possible consequence of breakdown of the mitochondrial membrane potential and apoptosis. All these effects described are antagonized by melatonin [91,93,102-107]. Another compound highly efficient in suppressing neuronal overexcitation is kynurenic acid [141,142], which acts in multiple ways, by binding to the glycine_B site of the NMDA receptor [143,144] and by decreasing extracellular glutamate levels by blocking presynaptic α 7 nicotinic receptors (α 7nAChRs) at glutamatergic nerve endings [145,146]. An obstacle for the direct application of kynurenic acid is its poor penetration of the blood-brain barrier. However, this can be overcome by using indole-3-pyruvic acid, which is sufficiently taken up by the aromatic amino acid transporter and readily forms kynurenic acid via free-radical reactions [147-149]. In fact, administration of indole-3-pyruvic acid leads to rises in brain kynurenic acid [150]. The precursor has been clinically tested and was found to be suitable in reducing signs of chronic stress and mild anxiety, in accordance with the assumed anti-excitatory action of the product, kynurenic acid [151,152]. From the antagonism to overexcitation one might also deduce decreases in radical formation and, thus, a mitigation of oxidative stress.

While the above-mentioned mechanisms are widely related to intercellular signaling, direct effects on mitochondria may be expected to be even more effective in reducing radical formation. Again, melatonin is a compound of premier interest because of its actions as a mitochondrial metabolism modifier [58]. The indoleamine, previously mainly regarded as a radical scavenger and regulator of redox-active enzymes, exerts several additional actions at the mitochondrial level that exceed the earlier findings (Fig. 2). Due to its amphilicity melatonin easily crosses membranes, but is also water-soluble and, therefore, available in any compartment and in the lipid environment as well. Moreover, it has been shown to accumulate in mitochondria [153,154]. Again, melatonin acts in multiple ways, which collectively contribute to radical avoidance. First, melatonin interferes with NO metabolism by down-regulating the inducible nitric oxide synthase (iNOS), especially a mitochondrially targeted iNOS subform [154-159], by inhibiting neuronal NOS (nNOS) [107,160-164], scavenging of NO and peroxynitrite-derived radicals [89,107,165,166] (Fig. 3). Moreover, melatonin's metabolite, N^1 -acetyl-5methoxykynuramine (AMK) acts as a potent scavenger of NO [167-170], of ROS formed from peroxynitrite [167], and as a highly efficient inhibitor of nNOS [171]. Therefore, any NO-related partial ETC blockade is antagonized by melatonin and its product. Second, melatonin supports the function of the ETC by preventing lipid peroxidation in mitochondrial membranes, in particular that of cardiolipin, which is crucial for the appropriate function of Complexes III and IV [17,18,112,172,173] (Fig. 2). Lipids, in particular, cardiolipin, are additionally protected by melatonin because of its influences on glutathione metabolism and the upregulation of mitochondrial glutathione peroxidase (GPx) [105-109,111,174-176]. This has been also demonstrated in SAMP8 mice, in which melatonin antagonized the agerelated decreases in GPx [99,109,111]. Although these studies did not specifically relate the effects observed to the isoform GPx4, this enzyme should have been involved. Moreover, cytochrome c release was shown to be strongly affected by GPx4 levels, and cardiolipin was protected in



Fig. (3). Some aspects of NO metabolism, which are related to mitochondrial effects and counteractions by melatonin (Mel) and its metabolite N^1 -acetyl-5-methoxykynuramine (AMK). Red arrows and symbols: down-regulation or inhibition; violet symbols: scavenging.

transgenic mice overexpressing this isoenzyme, whereas, in heterozygous knockouts, cardiolipin peroxidation was exacerbated [177]. Third, melatonin modulates mitochondrial metabolism by directly interacting with the ETC [17,18].

The direct effects of melatonin at the ETC have remained unclear for quite some time, and more specific data were obtained not before recently. One of the ideas for explaining the reduction in electron leakage caused by melatonin had been that of an electron shuttle, in which the indole acts as an electron donor to the ETC and the resulting melatonyl radical as an electron acceptor [58,91,93]. A corresponding action was assumed for AMK [58,91]. Electron donation to the ETC was, in fact, observed at cytochrome c, but the oxidation process studied led to AFMK, at least under the experimental conditions used [178]. Therefore, the idea of an electron shuttle bridging electron acceptor sites and those of electron dissipation would still require experimental support. However, this concept had not been entirely speculative, because it was designed on the basis of experience with nitrones [58]. These compounds are known as spin traps, but, compared to melatonin, they are relatively poor radical scavengers. Nevertheless, they are effective in supporting the electron flux through the ETC, thereby reducing oxidative stress [179-181], findings which may be interpreted in terms of decreased electron leakage and improved radical avoidance. Since the compound used in earlier studies, α phenyl-tert-butylnitrone (PBN), had some unfavorable properties with regard to mitochondrial entrance and decomposition, derivatives were developed which were specifically designed for mitochondrial targeting, e.g., by making the compounds more amphiphilic [181-185]. Several of these newly developed nitrones were much more potent scavengers than PBN, but, more importantly, they protected from H_2O_2 , peroxynitrite and doxorubicin toxicity with high efficiency [182-185]. In a gerontological model organism, the rotifer Philodina acuticornis odiosa, some of these compounds exerted remarkable effects of life extension. The amphiphilic nitrone LPBNAH was capable of more than doubling mean and maximal lifespan [182]. This effect was obviously not caused by slowing metabolic processes, as known from life-extending, ETC-related mutants of Caenorhabditis elegans [17,186,187], since the longer-lived rotifers grew to much larger sizes than the normal-lived animals [182]. Unfortunately, corresponding experiments in mice were not successful, presumably because of decomposition of these nitrones in the murine metabolism differing from that of rotifers, and formation of harmful products. However, the nitrone effects in Philodina may be taken as a proof of principle for the efficacy of mitochondrial metabolism modifiers.

The distinction between mitochondrial metabolism modification and free-radical scavenging seems to be of particular importance, although several of the protective compounds combine the two properties, foremost melatonin and its metabolite AMK. Experiments with L-theanine, a substance present in green tea leaves, but being almost devoid of radical scavenging properties, was found to efficiently prolong life in outbred mice not prematurely dying from cancer, in Fischer 344 rats and in *Philodina*. L-Theanine also supported healthy aging of mice, restored brain ATP levels to normal in 24 months old rats, prevented neurodegeneration in the hippocampal areas CA1 and CA3

and restored the hippocampal density of serotoninergic innervation [B Poeggeler, pers commun]. These data by MA Pappolla and B Poeggeler, which are not yet published in detail, but were briefly cited elsewhere [132], could be related to the modification of mitochodrial metabolism in the central nervous system. The efficacy of the compound is obviously of indirect nature, due to the modulation of glutamatergic neurotransmission, and presumably includes the prevention of overexcitation [188] (Fig. 2).

The mitochondrial effects of melatonin may be partially independent of redox reactions. In addition to the previously known low-affinity binding site related to the inhibition of the mitochondrial permeability transition pore [189], a different, high-affinity mitochondrial binding site exists. Earlier studies in pigeon brain had shown that 39% of 2iodomelatonin was bound to mitochondria [190]. More recently, the high-affinity site ($K_d = 150 \text{ pM}$) was found to be localized at the amphipathic ramp of Complex I and seems to be associated with melatonin-dependent rises in the activity of the iron-sulfur cluster N2 [17,18,96,132,134,191]. These data by MA Pappolla, B Poeggeler and B Pucci have not yet been published in detail, but some characteristics have been reported [191]. Displacement experiments with specific ligands indicated binding near the N2 cluster. These results strongly favor the idea that melatonin may directly control electron flux at Complex I and, thereby, reduce electron dissipation at this site, an action of radical avoidance.

CONNECTIONS BETWEEN REDOX, SENSING MECHANISMS AND AGING-SUPPRESSOR GENES?

The conclusion that anti-oxidative protection is much more than radical scavenging and regulation of anti- and prooxidant enzymes, but rather has to consider the attenuation of free-radical generation, necessarily leads to the question of how mitochondrial function is controlled and dysfunction avoided. This complex of problems remains to be elucidated in many details. Nevertheless, various indications for future studies exist. The processes of aging have been shown to be influenced by mechanisms of both redox and nutrient sensing [192]. Since nutrients are also related to the energy balance and, thus, mitochondrial function, interconnections between redox- and nutrient-regulated pathways have to be expected. Their terminal actions seem to control electron flux and dissipation, but also mitochondrial biogenesis.

Calorie restriction has been frequently used as a means for prolonging life [4,192-195]. Nutrient sensing had been related to the signaling pathway of growth hormone (GH) and insulin-like growth factor-1 (IGF-1), whose disruption robustly favored longevity [193]. However, an additional relationship between caloric restriction and the aging suppressor SIRT1 became apparent [4,193,194]. Sirtuins are known to promote longevity in numerous organisms, from yeast to insects and vertebrates. The seven mammalian subforms, SIRT1 to SIRT7, are multiply involved in mitochondrial function. At least SIRT3, SIRT4 and SIRT5 are mitochondrially localized [196]. Like another aging suppressor, klotho, SIRT3 modulates the FoxO signaling pathway by interacting with the mitochondrial FoxO3a homolog, daf-16 [197]. Due to its function as an NAD⁺- dependent lysine deacetylase [198], SIRT3 prevents mitochondrial lysine hyperacetylation [196]. In fibroblasts, it was shown to physically interact with the Complex I subunit, the 39-kDa protein NDUFA9, to enhance Complex I activity and ATP levels [199]. The functional relationship of sirtuins to mitochondria goes beyond their presence within these organelles, comprises additional regulatory effects and also mitochondrial biogenesis [4] (Fig. 2). The subform SIRT1, which is not mitochondrially localized, also modulates NO formation, the insulin/IGF-1 pathway, activates FoxO subforms and, thereby, anti-oxidant enzyme expression [4,200]. A connection to free radical metabolism is, therefore, evident. The requirement of SIRT1 for mitochondrial activity becomes obvious by its dependence on the AMP level, an indicator of ATP deficiency. Normally, SIRT1 as well as AMP-activated protein kinase (AMPK) simultaneously respond to elevated AMP, but these two important regulators also act concordantly in situations of stress, starvation or calorie restriction [201] (Fig. 2). SIRT1 seems to have additional effects on mitochondrial electron transport capacity, including the amount of mitochondrial volume per cell [4,200]. Increased mitochondrial biogenesis was found to be associated with rises in AMPK, in NO - which, at moderate concentrations not leading to detrimental peroxynitrite levels, should not be generally harmful - and, more in particular, with the up-regulation of the peroxisome proliferator-activated receptor- γ coactivator-1 α (PGC-1 α) [202,203] (Fig. 2). A further connection exists with regard to circadian rhythmicity, which we had discussed in its relevance to radical avoidance. SIRT1 was shown to modulate chromatin remodeling via the clock gene protein CLK and seems to directly influence at least peripheral oscillators by interacting with the CLK/BMAL1 complex [204,205]. These actions should lead, for fundamental reasons, to numerous secondary effects, also beyond chromatin structure.

It seems that the relationships between mitochondrial regulators and the anti-oxidative defense are more complex than previously thought. AMPK was shown to be upregulated by peroxynitrite, so that the activation of this enzyme was concluded to represent an early warning signal of oxidative stress [206]. Resveratrol, which is, among other properties, a radical scavenger, anti-oxidant and anti-inflammatory agent [207-209], has also been identified as a dietary modulator and ligand of SIRT1 [4,210] (Fig. 2). Resveratrol, being the most potent natural SIRT1 activator [8], loses its chemopreventive efficacy in SIRT1 knockouts, so that the most potent pertinent actions are obviously mediated via this sirtuin isoform [211]. Stimulation of mitochondrial biogenesis via SIRT1 signaling involves PGC-1a [193,195, 212]. In endothelial cells, resveratrol increased mitochondrial mass and mtDNA content via SIRT1 and up-regulation of eNOS. Knockdown experiments showed that the blockade of either SIRT1 expression or NO formation prevented these effects [212]. Although the involvement of NO may be celltype specific, it demonstrates again that moderate levels of this gaseous mediator are not detrimental per se. Another anti-oxidant, a-lipoic acid, also acts via PGC-1a [203]. However, it remains to be clarified whether the protective effects of this compound are mainly exerted through this signaling pathway or through lipoylation of mitochondrial proteins, which is also required for the normal functioning of the ETC, especially Complex I [213]. With regard to mitochondrial actions of sirtuins and their common endpoints with mela-tonin, we had suggested to investigate their connection to the indoleamine [191]. Thereafter, first indications were obtained for such a relationship: in SAMP8 mice, melatonin was reported to up-regulate SIRT1 [214]. Recently, more detailed studies have supported the influence of melatonin on SIRT1. In a model using sleep-deprived rats [215], mela-tonin was shown to favor the hippocampal expression of SIRT1. In another investigation, neuronal cultures from young and aged rats were compared with regard to effects of melatonin [216]. In neurons from aged rats, melatonin up-regulated SIRT1 expression and also enhanced the deacety-lation of various SIRT1 substrates, such as PGC-1 α , FoxO1, NF κ B, and p53, effects which were largely reverted by the SIRT1 inhibitor sirtinol [216]. The melatonin-induced deacetylation of PGC-1a indicates that mitochondrial biogenesis might be stimulated by the indoleamine in vivo.

CONCLUSION

Multiple connections exist between oxidative damage and aging. However, it seems important to avoid simplifications. Although the supply with anti-oxidants may favor a healthy aging, one should not expect to arrive at substantial extensions of lifespan by only detoxifying free-radicals and other oxidants formed, even not when this comprises upregulations of anti-oxidant enzymes. With regard to the crucial roles of mitochondria in the generation of free radicals and other oxidants, as well as in the aging-related metabolic decline, the support of mitochondrial function turns out to be of highest relevance. A well-operating ETC does not only provide sufficient amounts of ATP, but, at the same time, avoids electron dissipation and thereby radical formation. A compromised ETC may be regarded as a source of further damage and cellular decline, but this should not be interpreted in terms of the original version of the free radical theory of aging. If a vicious cycle exists, it is not that one between the radical producing ETC and the mtDNA, but rather another one within the ETC itself. Damage to the ETC causes oxidant formation, which leads to further impairment of electron flux. Radical avoidance as an aim of antioxidative protection intends to safeguard electron flux. Several low molecular weight molecules, such as resveratrol, α -lipoic acid, melatonin and its metabolite AMK may be suitable for helping sustain the electron transport through the ETC and for reducing radical formation. Although these compounds have been regarded in the beginning mainly as direct anti-oxidants and, later, also as regulators of anti- and pro-oxidant enzymes, their role in the control of ETC components and mitochondrial biogenesis may be decisive. It remains fascinating to see evidence accumulating, which reveals how some of these chemically entirely different compounds merge at same pathways to promote mitochondrial growth. The earlier findings indicating an important role of the circadian oscillator system in avoiding oxidative damage now receives further support by the demonstration that PGC-1a, a central regulator of mitochondrial biogenesis, is, at the same time, involved in the circadian control. Melatonin, which does not only regulate anti- and pro-oxidant enzymes, combines properties of a major circadian regulator and a mitochondrial metabolism modifier. It may now be seen in this larger context of a regulatory network centered at the support of mitochondrial function.

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