The Oxidative Stress of Hyperhomocysteinemia Results from Reduced Bioavailability of Sulfur-Containing Reductants

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Abstract: Vegetarian subjects consuming subnormal amounts of methionine (Met) are characterized by subclinical protein malnutrition causing reduction in size of their lean body mass (LBM) best identified by the serial measurement of plasma transhyretin (TTR). As a result, the transsulfuration pathway is depressed at cystathionine-β-synthase (CβS) level triggering the upstream sequestration of homocysteine (Hcy) in biological fluids and promoting its conversion to Met. Maintenance of beneficial Met homeostasis is counterpoised by the drop of cysteine (Cys) and glutathione (GSH) values downstream to CβS causing in turn declining generation of hydrogen sulfide (H$_2$S) from enzymatic sources. The biogenesis of H$_2$S via non-enzymatic reduction is further inhibited in areas where earth’s crust is depleted in elemental sulfur (S$_8$) and sulfate oxyanions. Combination of subclinical malnutrition and S$_8$-deficiency thus maximizes the defective production of Cys, GSH and H$_2$S reductants, explaining persistence of unabated oxidative burden. The clinical entity increases the risk of developing cardiovascular diseases (CVD) and stroke in underprivileged plant-eating populations regardless of Framingham criteria and vitamin-B status. Although unrecognized up to now, the nutritional disorder is one of the commonest worldwide, reaching top prevalence in populated regions of Southeastern Asia. Increased risk of hyperhomocysteinemia and oxidative stress may also affect individuals suffering from intestinal malabsorption or westernized communities having adopted vegan dietary lifestyles.

Keywords: Vegetarianism, malnutrition, sulfur-deficiency, hyperhomocysteinemia, oxidative stress, hydrogen sulfide, cardiovascular diseases, developing countries, Asia.

Homocysteine (Hcy) is a nonproteogenic sulfur-containing amino acid (SAA) generated by the intrahepatic transmethylation (TM) of dietary Met. Hcy may either be recycled to Met following remethylation (RM) pathways or catabolized along the transsulfuration (TS) cascade. Under normal circumstances, the Met-Hcy cycle stands under the regulatory control of three watersoluble B-vitamins: folates (5-methyl-tetrahydrofolates, B$_9$) are regarded as the main factor working as donor of the CH$_3$ group involved in the remethylation process, pyridoxine (pyridoxal-5’-phosphate, PLP, B$_6$) plays the role of co-factor of both cystathionase enzymes belonging to the TS pathway and cobalamins (B$_12$) ensure that of methionine-synthase. The main steps of the Met → Hcy → Met cycle are summarized in Fig. (I).

Hyperhomocysteinemia (HHcy) is an acquired metabolic anomaly first identified by McCully [1], a finding confirmed in the last four decades by a considerable body of experimental and clinical investigations. The current consensus is that dietary deficiency in any of three watersoluble vitamins may operate as causal factor of HHcy. PLP-deficiency may trigger the upstream accumulation of Hcy in biological fluids [2] whereas the shortage of vitamins B$_9$ or B$_{12}$ is held responsible for its downstream sequestration [3,4]. HHcy is regarded as a major causal determinant of CVD working as an independent and graded risk factor unrelated to the classical Framingham criteria such as hypercholesterolemia, dyslipidemia, sedentary lifestyle, diabetes and smoking. Hcy may invade the intracellular space of many tissues and locally generate in a time- and concentration-dependent manner [5] endothelial dysfunction working as early harbinger of blood vessel injuries and atherosclerosis. Although many deleterious mechanisms have been incriminated [6], most investigators contend the view that production of harmful reactive oxygen and nitrogen species (ROS, NOS), notably hydrogen peroxide (H$_2$O$_2$), superoxide anion (O$_2$•-) and peroxinitrite (ONOO•), constitutes a major culprit in the development of HHcy-induced vascular damages [7-10]. Accumulation of ROS is associated with increased risk for cardiovascular diseases [11] and stroke [12], arterial hypertension [6], kidney dysfunction [13], Alzheimer’s disease [14], cognitive deterioration [15], inflammatory bowel disease [16] and bone remodeling [17].

We nevertheless observe that these toxicological effects currently overlook the potential protective roles played by extra- and intracellular reductants such as cysteine (Cys) and glutathione (GSH) in the sequence of events leading from HHcy to tissue damages. Moreover, after the discovery of nitric oxide (NO) and carbon oxide (CO), hydrogen sulfide (H$_2$S) is the third gaseous signaling messenger found in mammalian tissues [18]. H$_2$S is a reducing molecule displaying strong scavenging properties as animal experiments and tissue cultures demonstrate that exogenous supply of the neurotransmitter significantly attenuates [19, 20] or even
Malnutrition, Sulfur-Deficiency and Hyperhomocysteinemia

Fig. (1). Schematic representation of the methionine cycle and homocysteine degradation pathways.

**Compounds:** ATP, adenosyltriphosphate; THF, tetrahydrofolate; SAM, S-adenosylmethionine; SAH, adenosylhomocysteine; Cysta, cystathionine; Cys, cysteine; GSH, glutathione; H\(_2\)S, hydrogen sulfide; Tau, taurine; SO\(_4^{2-}\), sulfate oxyanions.

**Enzymes:** (1) Met-adenosyltransferase; (2) SAM-methyltransferases; (3) adenosyl-homocysteinase; (4) methylene-THF reductase; (5) Met-synthase; (6) cystathionine-\(\beta\)-synthase, C\(\beta\)S; (7) cystathionine-\(\gamma\)-lyase, C\(\gamma\)L; (8) \(\gamma\)-glutamyl-synthase; (9) \(\gamma\)-glutamyl-transpeptidase; (10) oxidase; (11) reductase; (12) cysteine-dioxygenase, CDO.

**Metabolic pathways:** Met molecules supplied by dietary proteins are submitted to TM processes allowing to release Hcy which may in turn either undergo Hcy-Met RM pathways or be irreversibly committed into TS decay. Impairment of C\(\beta\)S activity, as described in protein malnutrition, entails supranormal accumulation of Hcy in body fluids, stimulation of (5) activity and maintenance of Met homeostasis. This last beneficial effect is counteracted by decreased concentration of most components generated downstream to C\(\beta\)S, explaining the depressed C\(\beta\)S- and C\(\gamma\)L-mediated enzymatic production of \(\text{H}_2\text{S}\) along the TS cascade. The restricted dietary intake of elemental S further operates as a limiting factor for its non-enzymatic reduction to \(\text{H}_2\text{S}\) which contributes to downsizing a common body pool (dotted circle). The data show that combined protein- and S-deficiencies work in concert to deplete Cys, GSH and H\(_2\)S from their body reserves, hence impeding these reducing molecules to properly face the oxidative stress imposed by hyperhomocysteinemia.

abolishes [21,22] the oxidative injury imposed by HHcy burden. The endogenous production of the naturally occurring H\(_2\)S reductant depends on Cys bioavailability through the mediation of TS enzymes [23,24]. Production of H\(_2\)S may also occur in human tissues starting from elemental sulfur, a non-enzymatic reaction requiring the presence of Cys, GSH, and glucose [25,26]. It would be worth disentangling the respective roles played by Cys, GSH and H\(_2\)S for the prevention and restoration of HHcy-induced oxidative lesions. This proposal is made in the light of a recently published study showing that the plasma concentration of Cys and GSH is severely depressed in subclinically malnourished HHcy patients [27], hence impeding appropriate biogenesis of H\(_2\)S molecules. The present paper attempts to clarify the biological consequences resulting from the complex interplay existing between the 3 reducing molecules, allowing to
throw deeper insight into the pathophysiologic mechanisms associated with HHcy states.

CLINICAL BACKGROUND

Numerous surveys undertaken throughout the last decades have conclusively shown that the watersoluble vitamin deficiency concept, taken together, provides only partial causal account of the HHcy metabolic anomaly. In contrast, recent studies have highlighted that the components of body composition, mainly the size of lean body mass (LBM), constitutes a critical determinant of HHcy status [28,29]. Because nitrogen (N) and sulfur (S) concentrations maintain tightly correlated ratios in tissues of both healthy subjects [30] and diseased patients [31], it might follow that defective N intake and accretion rate would cause concomitant and proportionate depletion of total body N (TBN) and total body S (TBS) stores [30]. Our clinical investigation undertaken in Central Africa in apparently healthy but nevertheless subclinically malnourished vegetarian subjects has documented that reduced size of LBM could lead to HHcy states [27].

The field study was conducted in the southern part of the Republic of Chad, a sahelian area mainly populated with inhabitants belonging to the Sara ethnic group [27]. In this semi-arid region, the staple food is monotonous, consisting mainly of cassava, sweet potatoes, beans, millets and groundnuts. Participants were invited to fill in a detailed dietary questionnaire whose results were compared with values reported in food composition tables [32-34] allowing to calculate the proportion of energy, fiber, carbohydrate, lipid, protein, and SAAs ingested on a daily basis [27]. Overall health status was determined as well as plasma lipids, watersoluble vitamins and selected CVD risk factors. The dietary inquiry indicates that participants consumed a significantly lower mean SAA intake (10.4 mg.kg⁻¹.d⁻¹)[27] than the Recommended Dietary Allowances (RDAs) (13 mg.kg⁻¹.d⁻¹) [33,34].

The blood lipid profiles of rural subjects were confined within normal ranges ruling out this class of parameters as causal risk factors for CVD disorders. The normal levels measured for pyridoxine, folates, and cobalamins did preclude these vitamins from playing any significant role in the rise of Hcy plasma concentrations [27]. Analysis of plasma SAAs revealed unmodified methioninemia, significantly elevated Hcy values (18.6 μmol/L) contrasting with significantly decreased plasma Cys and GSH values [27]. The significant lowering of classical anthropometric parameters (body weight, BW; body mass index, BMI) together with that of the main plasma and urinary biomarkers of metabolic (visceral) and structural (muscular) compartments point to an estimated 10 % shrinking of LBM [27]. We have attached peculiar importance to the measurement of plasma transsulfhydrin (TTR) since this indicator integrates the evolutionary trends outlined by body protein reserves [35], providing from birth until death an overall and balanced estimate of LBM fluctuations [29]. In the absence of any superimposed inflammatory condition, LBM and TTR profiles indeed reveal striking similarities [29]. Scientists belonging to the Foundation for Blood Research (Scarborough, Maine, 04074, USA) have recently published a large number of TTR results recorded in 68,720 healthy US citizens aged 0-100 yr which constitute a comprehensive reference material to follow the shape of LBM fluctuations in relation with sex and age [29]. TTR concentrations plotted against Hcy values reveal a strongly negative correlation (r = -0.71) already described previously [29,30], confirming that the depletion of TBN and TBS stores plays a predominant role in the development of HHcy states. The body of a reference man weighing 70 kg contains 64 M of N (1,800 g) and 4,400 mM of S (140 g) [36].

Our vegetarian subjects consume diets providing low fat and high fiber content conferring a large spectrum of well-described health benefits notably for the prevention of several chronic disorders such as cancer and diabetes, together with an effective protection against the risk of hypercholesterolemia-induced CVD [37,38]. Plant-based regimens, however, do not supply appropriate amounts of nitrogenous substrates of good biological value which are required to adequately fulfill mammalian tissue needs [30]. More precisely, vegetable items contain suboptimal concentrations of both SAAs [33,34,39] below the customary RDA guidelines. This dietary handicap may be further deteriorated by unsuitable food processing [40] and by the presence in plant products of naturally occurring anti-nutritional factors such as tannins in cereal grains and anti-trypsin or anti-chymotrypsin inhibitors in soybeans and kidney beans [41].

IMPAIRMENT OF THE TRANSSULFURATION PATHWAY

The hypothesis that subclinical protein malnutrition might be involved in the occurrence of HHcy states via inhibition of cystathionine-β-synthase (CBS) activity did first arise in Senegal in 1986 [42] and was later corroborated in Central Africa [43]. The concept was clearly counterintuitive in that it was unexpected that high Hcy plasma values might result from low intake of its precursor Met molecule. Despite the low SAA intake of our vegetarian patients [27], plasma Met concentrations disclosed noticeable stability allowing to maintain the synthesis and functioning of myriads of Met-dependent molecular, structural and metabolic compounds described in detail elsewhere [30,44]. These clinical investigations have received strong support from recent mouse [45] and rat [46] experiments submitted to Met-restricted regimens. At the end of the Met-deprivation period, both animal species did manifest meaningful HHcy states (p<0.001) contrasting with significantly lower BW (p<0.001) reduced by 33 % [45] and 44 % [46] of control, respectively. In comparison with all other indispensable AAs, the uniqueness of Met behavior stands in accordance with balance studies performed on large mammalian species showing that the complete withdrawal of Met from otherwise normal diets causes the greatest rate of body loss, nearly equal to that generated by protein-free regimens [47,48].

This efficient Met homeostatic mechanism is classically ascribed to a PLP-like inhibition of CBS activity exerted through allosteric binding of S-adenosylmethionine (SAM) to the C-terminal regulatory domain of the enzyme [49,50]. The validity of such allosteric mechanism has been recently questioned, at least in Met-depleted animal studies, showing that the loss of CBS activity may develop via a (post)translational defect independently from intrahepatic SAM concentrations [45]. We have postulated the existence
of an independent sensor mechanism set in motion by TBS pool shrinkage and reduced bioavailability of Met – its main building block – working as an inhibitory feedback loop of CβS activity [30]. Such Met-bodystat, likely to be centrally-mediated, and aiming at maintaining unaltered Met disposal in conditions of decreased dietary provision implies the fulfillment of high metabolic priorities of survival value [30,44].

Whereas HHcy may be regarded as the dark side of a beneficial adaptive machinery [43], impairment of the TS pathway also depresses the production of compounds situated downstream to the CβS blockade level, notably Cys and GSH, keeping in mind that Cys may undergo reversible GSH conversion (Fig. 1). The plasma concentration of both Cys and GSH reductants is indeed significantly decreased in our vegetarian subjects by 33 % and 67 % of control, displaying negative correlations (r = –0.67 and –0.37, respectively) with HHcy values [27]. Reduced dietary intake of the preformed Cys molecule [27] and diminished Cys release from protein breakdown in malnourished states [51] may contribute to the lowering effect. The significantly decreased GSH blood levels may similarly be attributed to dietary composition since the tripeptide is mainly found in meat products but is virtually absent from cereals, roots, milk and dairy items [52] and because regimens lacking SAAs may lessen the production of blood GSH and its intrahepatic sequestration [53].

**BIogenesis of Hydrogen Sulfide**

The TS degradation pathway schematically proceeds along two main PLP-dependent enzymatic reactions working in succession (Fig. 1). The first is catalyzed by CβS (EC 4.2.1.22) governing the β-replacement of the hydroxyl group of serine with Hcy to generate Cysta plus H₂O. Cys may however substitute for serine and the β-replacement of its sulfhydryl group with Hcy releases Cysta and H₂S instead of water [54]. The second is regulated by cystathionine-γ-lyase (CγL, EC 4.4.1.1.) hydrolyzing Cysta to release Cys and α-ketobutyrate plus ammonia as side-products [55]. Under the control of CβS or CγL enzymes, Cys may also undergo non-oxidative desulphuration pathways leading to H₂S or sulfane-sulfur production [56]. Alternatively, Cys may be committed into oxidative conversion regulated by cysteine-dioxigenase (CDO, EC 1.13.11.20) which catalyzes the replacement of the SH group of Cys by SO₂⁻ to yield cysteine-sulfinate [56]. This last compound may be further decarboxylated to hypotaurine that is finally oxidized to Tau (67 %) and SO₄²⁻ oxyanions (33 %) [56]. CβS and CγL are both cytosolic enzymes and their relative contribution to the generation of H₂S may vary according to animal strains, tissue specificities and nutritional or physiopathological circumstances [23,24]. CβS and CγL are expressed in most organs such as liver, kidneys, brain, heart, large vessels, ileum and pancreas [57,58] potentially subjected to HHcy-induced ROS injury while keeping the capacity to desulfurate Cys and to locally produce H₂S as cytoprotectant signaling agent. CβS is the principal TS enzyme found in cerebral glial cells and astrocytes [59]. CγL predominates in the vascular system [60] whereas both enzymes are present in the renal proximal tubules [61].

H₂S is the third gaseous substrate found in the biosphere [18] after NO and CO. All three gases are characterized by severe toxicity when inhaled at high concentrations. In particular, H₂S produced by anaerobic fermentation is capable of causing respiratory death by inhibition of mitochondrial cytochrome C oxidase [62]. NO, CO and H₂S are synthesized from arginine, glycine and Cys, respectively, exerting at low concentrations major biological functions in living organisms. Most of our knowledge on these atypical signal messengers [63] are derived from animal experiments and tissue cultures. These transmitter molecules may share some properties in common such as penetration of cellular membranes independently from specific receptors [64]. They are also manifesting dissimilar activities: whereas NO and CO activate guanylyl cyclase to generate biological responses via cGMP-dependent kinases, H₂S induces Ca²⁺-dependent effects through ATP-sensitive K⁺ channels [65]. Some of these potentialities may work in concert while others operate antagonistically. For instance, NO and H₂S express vasorelaxant tone on endogenous smooth muscle [66] but reveal different effects on large artery vessels [67]. The data suggest that these gaseous substances maintain whole body homeostasis through complex interactions and multifaceted cross-talks between signaling pathways.

Elemental S (32.064 as atomic mass) is a primordial constituent of lava flows in areas of volcanic or sedimental origin usually presenting as crown-shaped stable octamolecules – hence its S₈ symbolic denomination – which conglomerate to form brimstone rocks. The vegetable kingdom is unable to assimilate S₈ and requires as prior step its natural or bacterial oxidation to SO₄²⁻ derivatives before launching the synthesis of SAA molecules along narrowly regulated metabolic pathways [30,44]. Distinct anabolic processes are identified in mammalian tissues which lack the enzymatic equipment required to organize sulfate oxyanions but possess the capacity of direct S₈ conversion into H₂S. S₈ is poorly soluble in tap waters [68] may be taken up and transported to mammalian tissues loosely fastened to serum-albumin (SA) [69]. S may also be covalently bound to intracellular S-atoms taking the form of sulfane-sulfur compounds [70] either firmly attached to cytosolic organelles or in untied form to mitochondria [57,58,71,72] to undergo later release in response to specific endogenous requirements [71]. Sulfane-sulfur compounds are somewhat unstable and may decompose in the presence of reducing agents allowing the restitution of S [70,71]. S may either endorse the role of stimulatory factor of several mammalian apoenzyme activities as shown for succinic dehydrogenase [73] and NADH-dehydrogenase [74] or operate as inhibitory agent of other mammalian apoenzymes such as adenylate kinase [75] and liver tyrosine aminotransferase [76]. Elemental S resulting from dietary supply or from sulfane-sulfur decay may be subjected to non-enzymatic reduction in the presence of Cys and GSH [25,26] and/or reducing equivalents obtained from glucose oxidation [25], hence yielding at physiological pH additional provision of H₂S.

The gaseous mediator is a weakly acidic molecule endowed with strong lipophilic affinities. In experimental models, the blockade of the TS cascade at CβS or CγL levels significantly depresses or even abolishes the vitally required production of Cys operating at the crossroad of multiple converging processes (Fig. 1). Addition of Cys to the incubation milieu resumes the generation of H₂S [19] in a Cys concentration-dependent manner [77]. The data indicate that, in
the context of SAA deprivation, the compounds situated downstream both cystathionases keep their functional poten-
tialities but are unable to express their converting Cys → H2S capacities in the absence of precursor substrate. Sum-
mning up, inhibition of CblS activity contributes to promote ef-ficient RM processes and maintenance of Met homeo-
Stasis but entails as side-effects upstream sequestration of Hey
molecules in biological fluids while decreasing the bioavail-
ability of Cys and GSH working as limiting factors for H2S
production. These last adverse effects thus constitute the
Achilles heel of a remarkable adaptive machinery.

**ROLES PLAYED BY HYDROGEN SULFIDE**

The first demonstration that human tissues may reduce S
to H2S was incidentally provided in 1924 when a man given
colloid sulfur for the treatment of polyarthritis did rapidly
exhale the typical rotten egg malodor [78]. H2S may be pro-
duced by the intestinal flora [79] and serves as a metabolic
fuel for colonocytes [80]. Prevention of endogenous poison-
ing by excessive enteral production is insured by the detoxi-
fying activities of mucosal cells [81], hindering any systemic
effect of the gaseous substrate. In tissue culture media, the
concentrations of sulfane-sulfur is situated at any time below
the nanomolar range [71]. The normal H2S concentration
measured in mammalian plasmas usually ranges from 10 to
100 μM with a mean average turning around 40-50 μM
[19,21,82,83]. This H2S plasma level, appearing as the net
product of organs possessing CblS and CyL enzymes and
supplemented by the non-enzymatic conversion of S, flows
transiently into the vasculature and freely penetrates into all
body cells. Supposing that the gaseous reductant is evenly
distributed in total body water (45 L in a 70 kg reference
man) allows an estimate of bioavailable H2S pool turning
around 2 mM which represents, in terms of S participation,
largely less than 1 / 1,000 of TBS. The peculiar adaptive
physiology of vegetaric subjects renders very unlikely that
their TBS pool might be solicited to release S-substrates
prone to undergo conversion to nascent H2S molecules since
they adapt to declining energy and nutrient intakes by
switching overall body economy toward downregulated
steady state activities. The release from TBS of substantial
amounts of S-compounds occurs only during the onset of
hypercatabolic states as documented in trauma patients [31]
and in infectious diseases [84], exacting as preliminary step
cytokine-induced breakdown of tissue proteins, a selective
hallmark of stressful disorders [85]. As a result, the limited
disposal of H2S endogenously produced might be readily
exhausted in fulfilling ROS scavenging tasks at the site of
oxidative lesions. All body organs generating H2S from TS
enzymes are indeed simultaneously producers and consum-
ers of the gaseous substrate whose actual concentration re-
fects the balance between synthetic and catabolic rates ow-
ing to an estimated biological half-life of less than 30 min-
utes in mammalian plasma [86]. The data are consistent with
clinical investigations showing that H2S concentrations
found in cerebral homogenates from Alzheimer’s disease
(AD) patients are very much lower than expected from val-
ues measured in healthy brains [87], suggesting that the
gaseous messenger is locally submitted to enhanced con-
sumption rates reflecting disease severity. The concept is
strongly supported by studies pointing to the negative corre-
lation linking the severity of AD to H2S plasma values [88].

Similar findings are described in pediatric [89] and elderly
[90] hypertensive patients indicating, here gain, that more
severe HHcy-dependent oxidative burden is associated with
more intense H2S uptake rates. These H2S cleansing prop-
eries are mainly exerted by mitochondrial organelles known to
be centrally involved in oxidative disorders [20,91]. Mal-
nourished subjects deprived of Cys and GSH disposal thus
incur the risk of H2S-deficiency rendering them unable to
properly overcome HHcy-imposed oxidative lesions. The
rapid exhaustion of H2S stores have detrimental conse-
quences as shown a contrario by converging animal experi-
timents and tissue cultures disclosing the beneficial effects of
exogenous administration of commonly used sulfide salt
donors (Na2S and NaHS) generating H2S gas once in solu-
tion. Such supply significantly augments H2S plasma con-
centrations allowing to counteract ROS damages.

H2S was primarily recognized as a physiological sub-
strate working as neuromodulator [92] and soon later as
vasorelaxant factor [65]. H2S is now regarded as endowed
with a broader spectrum of biological properties [18], oper-
at ing as a general protective mediator striving against most
degenerative organ injuries, being capable of neutralizing or
abolishing most ROS harmful effects. Table 1 collects find-
ings displaying that H2S may promote the synthesis and ac-
tivity of several anti-oxidative enzymes (catalases, Cu- and
Mn-superoxide dismutases, GSH-peroxidases) and stimulate
the production of anti-inflammatory reactants (interleukin-
10) or conversely downregulate pro-oxidative enzymes (col-
lagenases, elastases), pro-inflammatory cytokines (interleu-
kine-1β, tumor-necrosis factor α) and immune reactions (hy-
perleukocytosis, diapedesis, phagocytosis). It has been calcu-
lated that, at physiological pH (7.4) and body temperature
(37°C), about 18.5% of H2S remains undissociated whereas
81.5% undergoes catabolic disintegration in the form of hy-
drosulfide anion (HS-) or sulfide anion (S 2-) [117]. Since S is
the main element comprised in the diprotonated H2S mole-
cule (34.08 as molecular mass), we postulate that partial or
complete repair of HHcy-induced lesions constitutes the
therapeutic proof that S-deficiency is causally involved in
the development of ROS damages. The concept is sustained
by the observation that all synthetic drugs (diclofenac, indo-
methacin, sildenafil) utilized as surrogate providers of H2S
[64,118] are characterized by a large diversity of molecular
conformations but share in common the presence of S-
atom(s) mimicking, once released, H2S-like pharmacological
properties. It remains to be clarified whether the beneficial
effects of S-fortification to S-deficient subjects are mediated,
among other possible mechanisms, via stimulation [73,74] of
anti-oxidative enzymes or inhibition [75,76] of pro-oxidative
enzymes. The true RDA requirements for elemental S are
presently unknown in human nutrition and it is only very
recently that the essentiality of S has been recognized, caus-
ing HeC elevation in deficient individuals [119]. It is worth
reminding that the gaseous NO substrate may work in con-
cert or antagonistically [66,83] to fine-tuning the helpful
properties exerted by H2S on body tissues. Preliminary stud-
ies suggest for instance that NO operates, in combination
with H2S, as a potential modulator of endothelial remodeling
since NO-synthesize isofoms contribute to the activation of
metalloproteinasves involved in the regulation of the colla-
gen/elastin balance defining vascular elastance [83,120].
Representative models of these socio-economic disparities in ing stagnant on the sidelines of household income growth. Urban areas and underprivileged rural communities remain-middle class population groups mainly located in affluent economic development. Inequalities however persist between owing to appropriate education programs and improved economic development. Inequalities however persist between some countries such as Vietnam [127] and Bangladesh [128] the last decades, significant progresses have been achieved in Latin America appears as the least affected [125,126]. Along Saharan Africa endures medium nutritional distress and indicates that chronic malnutrition remains a worldwide spanning degrees of protein and energy deficiency that is associ- with increased morbidity and mortality rates. The determinants of malnutrition are complex and interrelated, comprising socio-economic and political conditions, insufficient dietary intakes, inadequate caring practices and super-imposed inflammatory burden. Children living in developing countries are paying a heavy toll to chronic malnutrition whereas adult populations are handicapped by feeble physical and working capacities, increased vulnerability to infectious complications and reduced life expectancy. Cross-sectional studies collected in the eighties indicate that chronic malnutrition remains a worldwide scourge with top prevalence recorded in Asia, whereas sub-Saharan Africa endures medium nutritional distress and Latin America appears as the least affected [125,126]. Along the last decades, significant progresses have been achieved in some countries such as Vietnam [127] and Bangladesh [128] owing to appropriate education programs and improved economic development. Inequalities however persist between middle class population groups mainly located in affluent urban areas and underprivileged rural communities remaining stagnant on the sidelines of household income growth. Representative models of these socio-economic disparities in global nutrition and health are illustrated in the two most populated countries in the world, China and India. Large surveys undertaken in 105 counties of China and recently published have concluded that the rural communities haven’t yet reach the stage of overall welfare [129]. In India, similar investigations have documented that extreme poverty still prevails in the northern mountainous states of the subcontinent [130]. Taken together, southern Asian countries fail to overcome malnutrition burden [131]. In some African countries, there exists even upward trends suggesting nutritional deterioration over the years [132] still aggravated by a severe drought.

The assessment of malnutrition in children usually rely on anthropometric criteria such as height-for-age, weight-for-height, mid upper arm circumference and skinfold thickness allowing to draw the degree of stunting and wasting from these estimates. In adult subjects, BW and BMI are currently selected parameters to which some biochemical measurements are frequently added, notably SA, classical marker of protein nutritional status, and creatininuria (u-Cr), held as indicator of sarcopenia. The former biometric approaches are very useful in that they correctly provide a static picture of the declared stages of malnutrition but fail to recognize the dynamic mechanisms occurring during the preceding months and the adaptive alterations running behind. The National Center for Health Statistics / WHO reference Tables must therefore be regarded as theoretical guide-
lines defining minimum risk distribution of underweight [133]. The validity of the latter biological analyses may similarly be questioned since it is known for a while that SA is an insensitive marker of protein-depleted states compared to TTR [134] and a better indicator of population than of individual protein status. u-Cr is likewise a meagerly informative tool as 10% loss of muscle mass is required before it reaches significantly decreased urinary concentrations [135]. The data imply that the magnitude of subclinical malnutrition is largely underscored when classical biometric and laboratory investigations are performed. Moreover, ruling out the protein component involved in HHCy epidemiology and confining solely attention to the B-vitamin triad led to unachieved conclusions. For instance, surveys undertaken in Taiwan [136] and in India [137] did establish HHCy variance turning around 30%, indicating that a sizeable percentage of subjects do not come within the vitamin shortage concept. To our knowledge, there exists only one recent review recommending the use of TTR in vegetarian subjects [138]. The main reason for making the choice of TTR is grounded on the striking similar plasma profile disclosed by this marker with both LBM and Hcy [29]. Under healthy conditions, the 3 parameters indeed show low concentrations at birth, linear increase without sexual difference in preadolescent children, gender dimorphism in teenagers with higher values recorded in adolescent male subjects and thereafter maintenance of distinct plateau levels during adulthood [29,139,140]. Under morbid circumstances, the plasma concentrations of Hcy manifest gradual elevation negatively correlated with LBM downsizing and TTR decline. In vegetarian subjects and subclinically malnourished patients, rising Hcy and diminished TTR plasma concentrations look as mirror image of each other, revealing divergent distortion from normal and allowing early detection of preclinical steps at the very same time both SA and u-Cr markers still remain silent.

Any disease process characterized by quantitative or qualitative dietary protein restriction or intestinal malabsorption may cause LBM shrinking, downregulation of TTR concentrations and subsequent HHCy upsurge. These conditions are documented in frank kwashiorkor [141], subclinical malnutrition [27,43] and anorexia nervosa [142]. In patients submitted to weight-reducing programs, LBM was found the sole independent variable negatively correlated with rising Hcy values [143]. Morbid obesity may be alleviated by medical treatment [143] or surgical gastroplasty [144,145], conditions frequently associated with secondary malabsorptive syndromes and malnutrition [146], rendering an account of acquired HHCy. In westernized countries, subclinical protein-depleted states are illustrated in immigrants originating from developing regions but keeping alive their traditional feeding practices [147] or by communities having adopted, for socio-cultural reasons, strict vegan dietary lifestyles [148].

THE ADDITIONAL BURDEN OF S-DEFICIENCY

After N, K and P, elemental S is recognized as the fourth most important macronutrient required for plant development. The essentiality of S in the vegetable kingdom arose from observations made many decades ago by pedologists and agronomists [149,150] revealing that the withdrawal of sulfate salts from nutrient sources produces rapid growth retardation, depressed chlorophyllous synthesis, yellowing of leaves and reduction in fertility and crop yields. A large number of field studies, mainly initiated for economical reasons, has provided continuing gain in fundamental and applied knowledge and led to the overall consensus that \( \text{SO}_4^{2-} \)-deficiency is a major world-wide problem [151,152]. Field investigations have shown that the concentration of \( \text{SO}_4^{2-} \) oxyanions in soils and drinking waters may reveal considerable variations ranging from less than 2 mg/L to more than 1 g/L, meaning a ratio exceeding 1/500 under extreme circumstances [30]. The main causal factors responsible for unequal distribution of \( \text{SO}_4^{2-} \) oxyanions are geographical distance from eruptive sites and intensity of soil weathering in rainy countries. \( \text{SO}_4^{2-} \)-deficient nutritional deficiencies entail detrimental effects to most African and Latin American crops [151] reaching nevertheless top incidence in Southeastern Asia [151,153]. One of the most depleted area is the Indo-Gangetic plain extending from Pakistan to Bangladesh and covering the North of India and Nepal [154]. Intensive agricultural production, lack of animal manure and use of fertilizers providing N, K and P substrates but devoid of sulfate salts may further aggravate an already complicated situation. As global population increases steadily and the production of staple plants predicted to escalate considerably, \( \text{SO}_4^{2-} \)-deficient disorders are expected to become more pregnant along the coming years [155] with significant harmful impact for mankind. Nevertheless, effective preventive efforts are developed in some countries aiming at fortification of soils mainly by ammonium sulfate or calcium sulfate (gypsum) salts, resulting in meaningful improvements in crop yield, SAAs content and biological value and opening more optimistic perspectives for livestock and human consumption [152,155-158].

Contrasting with the tremendously high amounts of data accumulated over decades by pedologists and agronomists on sulfate requirements and metabolism, the available knowledge on elemental sulfur in human nutrition looks like a black hole. Despite the fact that \( \text{S}_2 \) follows H, C, O, N, Ca and P as the seven most abundant element in mammalian tissues, it appears as a forgotten item. Not the slightest attention is dedicated to \( \text{S}_2 \) in the authoritative “Present Knowledge in Nutrition” series of monographs even though they go over most oligo- and trace-elements in minute detail. The geographical distribution of \( \text{S}_2 \) throughout earth’s crust is not well-known as extreme paucity of measurements in soils and tap waters prevents from reaching a comprehensive overview. Nevertheless, and because \( \text{S}_2 \) is the obligatory precursor substrate for the oxidative production of sulfate salts, a decremental dispersion pattern paralleling those of \( \text{SO}_4^{2-} \) oxyanions is likely to occur with highest values recorded in the vicinity of volcano sources and lowest values found in remote and washed-out areas. Obviously, a great deal of research on elemental \( \text{S} \) remains to be completed by clinical biochemists before rejoining the status of plant agronomy.

Taken together, these data imply that subclinically malnourished subjects living in areas recognized as \( \text{SO}_4^{2-} \)-deficient for the vegetable kingdom also incur increased risks to become \( \text{S}_2 \)-depleted. This clinical entity most probably prevails in all regions, notably Northern India, where protein malnutrition [130] and sulfur-deficiency [154] coexist. Combination of both nutritional deprivations explains why the bulk of local dwellers, including young subjects [159,160], may develop HHcy states and CVD disorders.
characterized by strong refractoriness to vitamin-B supplementation [160] or high incidence of stroke [161] unrelated to the classical Framingham criteria. The current consensus is that “the problem of CVD in South Asia is different in etiology and magnitude from other parts of the world” [162]. These disquieting findings are confirmed in several Asian countries [163] and have prompted local cardiologists to exhort their governments to focus more attention on CVD epidemiology [164].

CONCLUDING REMARKS

The main lesson to learn from the present review is that vegetarian subjects are not protected against the risk of CVD and stroke which should no longer be regarded as solely affecting populations living in westernized societies and whose morbidity and mortality risks are stratified by classical Framingham criteria. Likewise hypercholesterolemia, hyperhomocysteinemia should be incriminated as emblematic risk factor for a panoply of CVD and related disorders. Whereas the causality of cholesterol and lipid fractions largely prevails in affluent societies consuming high amounts of animal-based items, that of homocysteine predominates in population groups whose dietary lifestyle gives more importance to plant products.

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CONFLICT OF INTEREST

None Declared.

REFERENCES


