

Sulfenic acid in human serum albumin

Review Article

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Summary. Sulfenic acid (RSOH) is a central intermediate in both the reversible and irreversible redox modulation by reactive species of an increasing number of proteins involved in signal transduction and enzymatic pathways. In this paper we focus on human serum albumin (HSA), the most abundant plasma protein, proposed to serve antioxidant functions in the vascular compartment. Sulfenic acid in HSA has been previously detected using different methods after oxidation of its single free thiol Cys34 through one- or two-electron mechanisms. Since recent evidence suggests that sulfenic acid in HSA is stabilized within the protein environment, this derivative represents an appropriate model to examine protein sulfenic acid biochemistry, structure and reactivity. Sulfenic acid in HSA could be involved in mixed disulfide formation, supporting a role of HSA-Cys34 as an important redox regulator in extracellular compartments.

Keywords: Thiol – Human serum albumin – Sulfenic acid – Peroxynitrite – Hydrogen peroxide – Free radicals

Abbreviations: HSA, human serum albumin; HSA-SH, the thiol of HSA; BSA, bovine serum albumin; SH/HSA, amount of thiol per albumin molecule; RSOH, sulfenic acid; NBD-Cl, 7-chloro-4-nitrobenzo-2-oxa-1,3-diazol; DTNB, 5,5'-dithio-bis(2-nitrobenzoic acid); dimedone, 5,5-dimethyl-1,3-cyclohexanedione; GSH, glutathione; GSSG, glutathione disulfide.

Introduction

The overproduction of reactive oxygen species including nitrogen oxides under conditions of oxidative stress may damage various biomolecules leading to cell and tissue toxicity. In addition, there is increasing evidence that these reactive mediators play a central role in signal transduction under physiological and physiopathological conditions, leading to new proposals on the definition of ox-

idative stress in terms of disruption of redox signaling and control (Hansen et al., 2006a).

Thiols are preferential targets of reactive species. With one- and two-electron reduction potentials $E^{\circ'}$ (RS^{\bullet}/RSH) of 0.92 V and $E^{\circ'}$ ($RSSR/2RSH$) of -0.24 V for cysteine (Surdhar and Armstrong, 1986; Keire et al., 1992), thiols can be oxidized by a wide spectrum of radical and non-radical species, usually at rates several orders of magnitude faster than other amino acids. At the high concentrations that thiols can achieve in different cell compartments, thiols can effectively compete with other targets. Last, several thiol oxidation products can be efficiently repaired.

In the last few years, the reactions of reactive species with thiols, far from constituting simple scavenging pathways, are being recognized to participate in modulatory and signaling mechanisms, coupling changes in the redox state of the milieu to biochemical and cellular processes (Schafer and Buettner, 2001). Among the different posttranslational modifications generated by thiol oxidation, sulfenic acid is receiving increased interest since it has been identified in a growing list of proteins, where it serves catalytic or regulatory functions. In this paper we will focus on the formation of sulfenic acid in human serum albumin and revise its properties and possible biological function.

Thiol oxidation states and sulfenic acid formation

Oxidation of low molecular weight or protein thiols typically results in the formation of disulfide bonds (RSSR).

sine phosphatase 1B after reaction with an amide nitrogen of a neighboring residue (van Montfort et al., 2003; Salmeen et al., 2003; Salmeen and Barford, 2005).

Alternatively, thiol oxidation by one-electron pathways results in a sulfur-centered radical (thiyl radical, RS^\bullet) which is highly reactive and can combine with another thiyl radical to form disulfides, or react with a thiolate to yield a reductive disulfide radical anion ($RSSR^{\bullet-}$) that forms superoxide upon reaction with molecular oxygen. Thiyl radicals have also been shown to react with oxygen to form the peroxy radical ($RSOO^\bullet$), an unstable intermediate that rapidly derives in sulfinyl radical (RSO^\bullet) which after hydrogen abstraction from reductants (AH) can finally yield sulfenic acid through a free radical pathway (Quijano et al., 1997; Bonini and Augusto, 2001; Giles and Jacob, 2002).

In addition to the oxyacids listed above, thiol exposure to reactive species derived from nitric oxide can also yield nitroso ($RSNO$) and presumably nitrothiols ($RSNO_2$), which can also be sulfenic acid precursors (Stamler et al., 1992; Wink et al., 1994; van der Vliet et al., 1998).

Undoubtedly, several of the products of thiol oxidation by reactive species are highly reactive themselves (*i.e.* RS^\bullet , $RSOH$). This has led to the definition of an additional group of redox active molecules named reactive sulfur species (RSS), complementary to the list of reactive oxygen and nitrogen species (Giles and Jacob, 2002).

The formation of disulfides, sulfenic acid and nitrothiol can be reverted to thiol non-enzymatically or enzymatically, with the aid of the glutaredoxin and thioredoxin systems, or to glutathione disulfide by GSNO reductase (Liu et al., 2001; Sahoo et al., 2006). In contrast, the formation of higher oxidation states such as sulfinic (RSO_2H) and sulfonic acids (RSO_3H) has been considered to be irreversible, although recent evidence indicates that reduction of the sulfinic form of several peroxiredoxins can occur enzymatically *in vivo* by cysteine sulfinyl reductases in a rather complex ATP-consuming mechanism (Biteau et al., 2003; Budanov et al., 2004).

Sulfenic acid stabilization and reactivity

Sulfenic acids exhibit both nucleophilic and electrophilic reactivity in biological milieus. Consequently, sulfenic acid can react with another thiol providing a mechanism for disulfide formation or condense with a second sulfenic acid to form a thiosulfinate ($RS(O)SR$) (Kice and Cleveland, 1973; Torchinsky, 1981; Claiborne et al., 1993). Since there are few electrophilic functional groups in proteins,

the fact that nucleophiles can react with the sulfur atom of sulfenic acid is remarkable.

The biological formation of low molecular weight sulfenic acid is difficult to detect due to its high reactivity and consequent instability, therefore it has long been thought to be a transient intermediate in the oxidation of thiols. More recently, however, the formation of a relatively stable sulfenic acid has been identified in several proteins and enzymes and is receiving increased interest since it has been recognized to play a crucial role in catalytic, redox regulatory and signaling functions (Claiborne et al., 1999, 2001; Finkel, 2000). Indeed, sulfenic acid is involved in the catalytic cycle of several enzymes. Peroxiredoxins, involved in peroxide detoxification, contain a conserved cysteine that in the catalytic cycle is oxidized to a sulfenic intermediate, which then condenses with another cysteine to form a thioredoxin-reducible disulfide. In addition to this catalytic cycle, sulfenic acid can also be overoxidized to sulfinic acid in an inactivation process that can be eventually reverted by cysteine sulfinyl reductases.

Sulfenic acid stabilization depends on steric hindrance and the local environment of the thiol involved. The most important factor to account for sulfenic acid stabilization is the absence of other proximal thiol groups, as well as intramolecular hydrogen bonding interactions with suitable hydrogen bonding partners, ionization of sulfenic acid to stabilize the corresponding sulfenate, together with an apolar microenvironment with limited solvent access (Claiborne et al., 1999).

Human serum albumin

Human serum albumin (HSA) is the most abundant protein in the vascular compartment (0.6 mM) where it represents ~60% of the total proteins in plasma. It is interesting to note that an important amount of albumin is localized extravascularly (Peters, 1996). Actually, the albumin pool present in extravascular/extracellular compartments represents more than double of the intravascular pool. This globular protein of 66 kDa consists of a non-glycosylated single chain polypeptide containing 585 amino acids cross-linked by 17 disulfide bridges with one tryptophan, six methionines, 18 tyrosines and only one free cysteine, Cys34. The crystallographic structure of HSA with its reduced cysteine has already been resolved (Carter et al., 1989; Carter and He, 1990). Heart-shaped, it contains 67% α -helix and no β -sheet with three homologous domains that contain common structural motifs (Fig. 2).

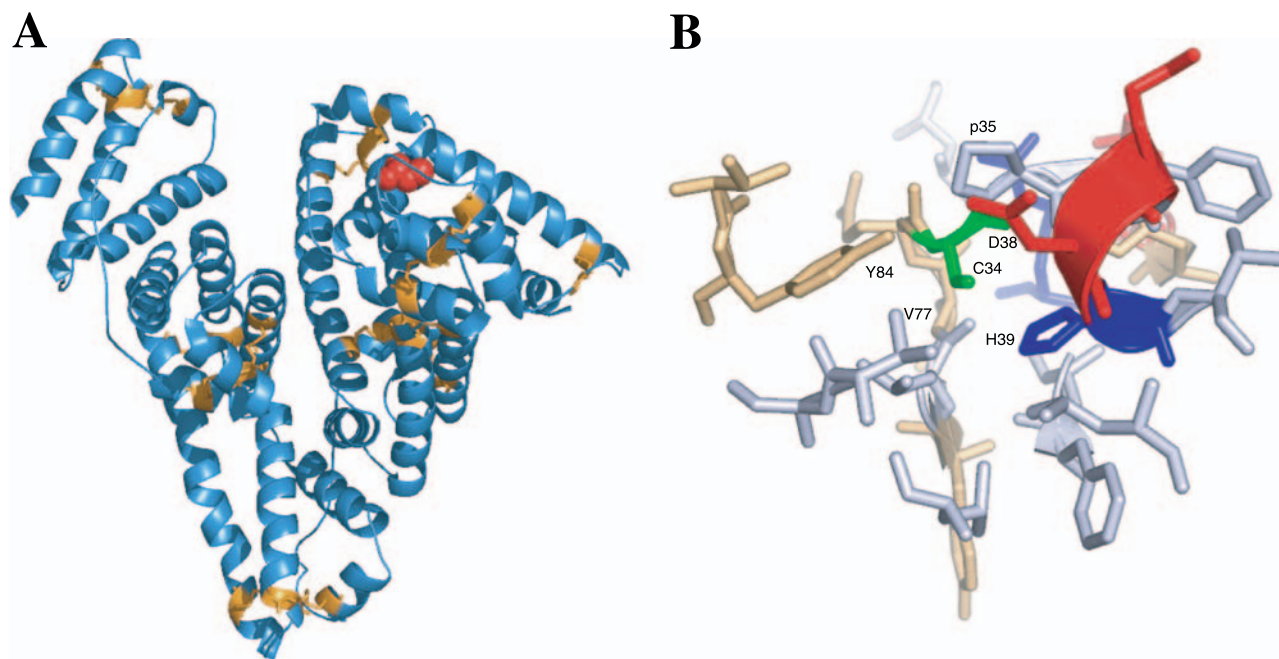


Fig. 2. Three-dimensional structure of HSA and location of Cys34. **A** Domain structure of human serum albumin. The locations of the disulfide bonds are shown in orange and Cys34 is shown in red. The amino terminus is at the right and the carboxyl terminus at the left. **B** Structural features of groups within 8 Å of the Cys34 site showing Asp38, Pro35, His39, Val77 and Tyr84. Polar residues are shown in light orange and non polar residues are shown in light blue. Cys34 is shown in green. Polar hydrophilic residues with positive and negative charge are colored in red and blue, respectively. HSA structural data was downloaded from the Protein Data Bank, accession code 1BMO (Sugio et al., 1999) and the figure was generated using PyMOL Molecular Graphics System (DeLano, 2002)

In addition to its physiological functions in plasma that include the maintenance of colloid osmotic pressure and the binding and transport of several ligands such as fatty acids, hormones and drugs, albumin is also proposed to serve antioxidant functions in the vascular compartment. This protective role is mainly attributed to its single thiol, Cys34, which accounts for ~80% of reduced thiols in human plasma and was determined to be the preferential plasma scavenger of the diverse family of oxygen and nitrogen reactive species (Halliwell, 1988; Halliwell and Gutteridge, 1990; Radi et al., 1991a; Alvarez et al., 1999).

Albumin is heterogeneous with respect to the thiol content. In plasma samples, the main HSA fraction called mercaptalbumin contains the reduced free thiol (~70% of total HSA). An important fraction of non-mercaptalbumin contains mixed disulfides between albumin thiol (HSA-SH) and low molecular weight thiols. In fact, about one-fourth of circulating albumin is present as mixed disulfides with cysteine, cysteinylglycine, homocysteine, γ -glutamylcysteine or glutathione. It has been proposed that in the remaining plasma non-mercaptalbumin fraction, the thiol is oxidized to higher oxidation states including sulfinic and sulfonic acids non-reducible by thiol

reagents (Janatova et al., 1968; Noel and Hunter, 1972; Era et al., 1988).

This balance between reduced and oxidized forms of HSA can shift in pathological states (Kawai et al., 2001; Tomida et al., 2003), with intense exercise (Inayama et al., 1996; Imai et al., 2002) and during ageing (Leto et al., 1970; Era et al., 1995). For example, an age-dependent decrease in the amount of thiol per human serum albumin molecule (SH/HSA) was observed, consistent with oxidative age-related damage.

Reactivity of HSA-SH with reactive species

Previous work from our group showed that the thiol in albumin was the preferential plasma target of the reactive species generated by xanthine oxidase, which include superoxide, hydrogen peroxide and hydroxyl radical (Radi et al., 1991b). We also demonstrated that among all amino acids, the thiol was the preferential target of peroxynitrite and its derived radicals (Radi et al., 1991a; Alvarez et al., 1999).

The thiol in HSA reacts with hydrogen peroxide and peroxynitrite with rate constants of 2.26 and $3.8 \times 10^3 \text{ M}^{-1} \text{ s}^{-1}$ at 37°C and pH 7.4 (Alvarez et al., 1999;

Carballal et al., 2003). This reactivity is comparable to that of low molecular weight thiols such as cysteine, glutathione and homocysteine (Radi et al., 1991a; Koppenol et al., 1992; Winterbourn and Metodiewa, 1999; Trujillo and Radi, 2002). Unlike other proteins that have thiols several orders of magnitude more reactive, enhanced by the protein microenvironment and their low pK_{ASH} values, such as peroxiredoxins or tyrosine phosphatases (Takakura et al., 1999; Bryk et al., 2000; Trujillo et al., 2004), the thiol in HSA is not particularly reactive exhibiting an apparent pK_{ASH} of 8.3–8.6, similar to those of cysteine (8.36) and glutathione (8.75) (Wilson et al., 1980; Torchinsky, 1981). Although the thiol in HSA is oxidized at relative slow rates, it still constitutes an important scavenger due to its high concentration in plasma (0.4–0.5 mM).

The oxidation of HSA-SH with hydrogen peroxide and peroxyxynitrite did not lead to the formation of disulfide bridged HSA dimers (Radi et al., 1991b; Carballal et al., 2003). Steric restrictions and the location of the single thiol of albumin in a crevice preclude dimer formation. This suggested that HSA-SH was oxidized to higher oxidation states such as sulfenic acid or further oxidized to sulfinic or sulfonic acids.

Sulfenic acid detection in HSA

Since sulfenic acids do not possess distinguishing absorbance or fluorescence features and are unstable, different tools and chemical methods have been developed for trapping and identifying such functional groups.

An important characteristic of sulfenic acid is the reactivity towards nucleophilic reagents such as benzylamine or dimedone (5,5-dimethyl-1,3-cyclohexanedione), that react specifically with sulfenic acid to form a stable thioether (Allison, 1976). These reagents can only be identified by mass spectrometry or using the radioactively labeled compounds. Recently, modified dimedones have been designed that react with sulfenic acids yielding fluorescent signals (Poole et al., 2005). While this reaction remains the standard for detection of sulfenic acids, reactions of dimedone with other groups such as aldehydes may be possible, therefore mapping the alkylation to a specific cysteine residue is strongly recommended.

Sulfenic acid can also be determined using thionitrobenzoate, another nucleophilic reagent that forms mixed disulfides and results in a loss in the absorbance at 412 nm (Ellis and Poole, 1997b; Boschi-Muller et al., 2000; Peshenko and Shichi, 2001).

The electrophilic reagent 7-chloro-4-nitrobenzo-2-oxa-1,3-diazol (NBD-Cl) has also been employed to detect sul-

fenic acid in proteins (Ellis and Poole, 1997a; Denu and Tanner, 1998; Carballal et al., 2003), but its ability to react with other protein nucleophile groups is a matter of concern.

Last, sulfenic acid can be detected after its reaction with arsenite, since arsenite reduces sulfenic acid back to thiol but does not reduce disulfides. Recently, proteomic approaches have been devised that allow to detect sulfenic acid based on its specific reduction with arsenite, followed by thiol detection with biotin-maleimide (Saurin et al., 2004).

The formation of sulfenic acid in albumin was first suggested by arsenite reduction of hydrogen peroxide-oxidized BSA-SH (Radi et al., 1991a, b). Later, its formation was also proposed after exposure of HSA to nitric oxide (DeMaster et al., 1995) as well as platinum complexes (Kratochwil et al., 1999). More recently mass spectral analyses using dimedone confirmed that sulfenic acid was formed by hydrogen peroxide-oxidation of HSA Cys34 and provided the definitive evidence for the formation of this derivative in human albumin (Carballal et al., 2003).

The reaction of sulfenic acid with thiols such as glutathione to form disulfides is a useful tool for characterizing the formation of sulfenic acid in HSA. Indeed, in the experiment described in Table 1 we show that, when peroxyxynitrite-treated HSA was incubated with reduced glutathione, the concentration of the latter decreased in the ultrafiltrates due to its reaction with sulfenic acid, in line with the initial formation of HSA-SOH by peroxyxynitrite which oxidizes thiols by two electrons. In the presence of carbon dioxide peroxyxynitrite decomposes to produce nitrogen dioxide and carbonate radical anion ($\text{CO}_3^{\cdot-}$) in

Table 1. Oxidation of GSH by peroxyxynitrite and peroxyxynitrite/ CO_2 -treated HSA

Time (min)	GSH oxidized (nmol) ^b	
	– CO_2	+ CO_2
3 ^a	156 ± 7 ^b	85 ± 8
60	152 ± 20	70 ± 15
120	157 ± 28	56 ± 11

^aThe amount of HSA thiols decreased upon exposure to peroxyxynitrite in the absence and in the presence of CO_2 from 0.563 mM to 0.187 mM and 0.357 mM, respectively

^bThe results are expressed as mean ± S.D. ($n = 3$)

HSA (0.56 mM) was incubated with ONOO^- (1 mM) in the presence and absence of CO_2 (25 mM NaHCO_3) at 37 °C, pH 7.4. At different time points, aliquots (0.5 mL) were removed and incubated with glutathione (0.48 mM) for 30 min. Then, the samples were ultrafiltered and glutathione oxidation was determined as reported previously (Carballal et al., 2003). As control, the addition of decomposed peroxyxynitrite to HSA led to no thiol oxidation. Catalase was added to the aliquots to prevent oxidation of glutathione by residual hydrogen peroxide

35% yield. These species can oxidize HSA-SH to thiyl radical (RS \cdot) which has been detected previously through EPR during the oxidation of BSA and low molecular weight thiols (Gatti et al., 1994; Bonini and Augusto, 2001) and can initiate an oxygen-dependent radical chain reaction. Remarkably, peroxynitrite-treated HSA in the presence of carbon dioxide also led to a decrease in the concentration of reduced glutathione, in agreement with the formation of sulfenic acid under these conditions, although to a lower extent. In Fig. 1 we show that the peroxy radical derived from the reaction of thiyl radical with oxygen (RSOO \cdot) can undergo homolytic cleavage to a sulfinyl radical (RSO \cdot) which can then oxidize another residue and finally yield sulfenic acid. Thus, our results support the hypothesis that sulfenic acid in HSA can also be formed through a free radical pathway.

Stability of sulfenic acid in HSA

Although sulfenic acids are generally unstable and highly reactive, HSA-SOH generated from exposure of albumin to peroxynitrite was remarkably stable (Table 1), similar as reported for hydrogen peroxide-treated HSA where more than 85% of sulfenic acid formed remained after 120 min (Carballal et al., 2003). In the presence of carbon dioxide, sulfenic acid was less stable, probably due to the free radical mechanism involved. Since stabilization depends on the local environment of the thiol in the native protein, the three-dimensional structure of HSA provided by X-ray diffraction confirms that Cys34 is located in a partially solvent-protected cleft in a hydrophobic crevice of 9.5–10 Å with no other adjacent thiol, surrounded by side chains of Pro35 and Val77 which could help to limit solvent accessibility and contribute to the apolar microenvironment. In addition, the imidazole ring from His39, the carboxylate group of Asp38 and the hydroxyl group of Tyr84 could also stabilize sulfenic acid with suitable electrostatic and hydrogen bonding interactions (Fig. 2) (He and Carter, 1992; Sugio et al., 1999; Stewart et al., 2005). It is possible that sulfenic acid will be present as the sulfenate anion at neutral pH, since its pK_a is expected to be lower than that of the thiol, based on the pK_as of the oxyacids of cysteine (2.1 and 1.3 for cysteine sulfinic and sulfonic acids, respectively) and those reported for stable low molecular weight sulfenic acids (4.8–6.3) (Claiborne et al., 1993).

Sulfenic acid in the context of the plasma environment

In contrast to the intracellular situation, where a reducing environment is maintained by a high concentration of

glutathione (5–10 mM) in the reduced state (GSH/GSSG \sim 100), the extracellular plasma environment is more oxidizing, with total ratios of reduced/oxidized thiols close to 1–2.

The most abundant reduced thiol in plasma is that of HSA (400–500 μ M), while the concentration of reduced low molecular weight thiol compounds only totals 10–20 μ M. Indeed, low molecular weight thiol compounds such as cysteine (265 μ M), cysteinylglycine (32 μ M), homocysteine (12 μ M) and glutathione (7.5 μ M) are present at total concentrations of \sim 300 μ M and mostly oxidized, with ratios of reduced *versus* total disulfides close to 0.01–0.1. The exception is glutathione, where the ratio reduced/total disulfide is close to 2 (5.1 μ M GSH versus 1.5 μ M low molecular weight glutathione disulfides and 1.6 μ M glutathione protein disulfides) (Mansoor et al., 1992; Andersson et al., 1993). In turn, a very important fraction of the low molecular weight thiol compounds (65%) are forming mixed disulfides with proteins, mainly with HSA, so that the reduced/disulfide ratio is close to 3 for HSA.

Thus, it is interesting to note that, analogously to the intracellular situation, the different plasma thiol/disulfide pools are not in equilibrium, and therefore could function as control nodes for different redox sensitive processes (Hansen et al., 2006b), although the kinetic barriers accounting for such differences are not yet elucidated. It is most likely that exchanges with intracellular compartments probably play a critical role in maintaining the thiol pools, since it has been observed that cultured cells and perfused tissues are able to regulate the redox state of extracellular compartments (Moriarty-Craige and Jones, 2004; Hansen et al., 2006a). In this line of thought, it is interesting to note that oxidized conformationally denatured albumin, which occurs *in vitro* and under oxidative stress conditions, is taken up by endothelial cells at increased rates (Schnitzer et al., 1992; Bito et al., 2005).

The numbers given above highlight the fact that the albumin thiol will be an important scavenger of reactive species, while low molecular weight thiols will not be able to compete with albumin for oxidants. This scavenging by Cys34 could represent a catalytic activity if the oxidation of the thiol in HSA is reverted by suitable yet undetermined reductants. In addition, it has been recently reported that the HSA disulfide between Cys392 and Cys438 sustains lipid peroxidase activity *in vitro* in the presence of thioredoxin (Cha and Kim, 2006), but its physiological significance in plasma needs to be confirmed. Another possible target for peroxides will be plas-

matic glutathione peroxidase, but since glutathione is scarce, the system may be easily overcome (Brigelius-Flohe, 1999).

While albumin circulates mostly in the reduced state, it can certainly form mixed disulfides. Assuming that albumin is released into the circulation from the liver with Cys34 mostly in its reduced form (Peters, 1996), the fact that 25% of HSA circulates as mixed disulfides with low molecular weight thiols, mainly cysteine, can be accounted for by three possible non-enzymatic mechanisms: thiol-disulfide exchange, radical reactions or reactions of sulfenic acid with a thiol (Di Simplicio et al., 2005). Indeed, one possible role for the stable sulfenic acid in albumin could be as intermediate in mixed disulfide formation (Carballal et al., 2003).

Sulfenic acid in HSA as well as mixed disulfides, could represent reversible modifications able to regenerate reduced HSA thiol with the appropriate reductant. This can be considered a protective mechanism to prevent overoxidation of cysteine to sulfinic and sulfonic acid. The detection of sulfenic acid *in vivo* remains a challenge. Also, whether an enzymatic system is operative in plasma for reducing sulfenic acid or mixed disulfides, and what the ultimate reductants are, needs to be established. Hopefully, the characterization and quantification of the oxidative modifications in albumin Cys34 will have utility as biomarkers of oxidative reactions in the vascular compartment and will yield useful information concerning the role and/or significance of the stable sulfenic acid in HSA.

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