Rhodanese: One of Natures "Sulphur" Biotransformation Machinery

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Abstract: Rhodanase (EC2.8.1.1) is a sulphur transferase enzyme. It is widely distributed across all levels of living organism; microbes, fungi, plants and animals. Sulphur is a basic requirement for existence and sustenance of life, as notable in the myriad of chemical substances in which it is incorporated such as; proteins containing cysteine and/or methionine amino acid, proteins containing iron-sulphur center prosthetic groups, glutathione, sulphated carbohydrates, hydrogen sulphide, and thiosulphate. Rhodanese plays an important role in the metabolism of sulphur in cells of organism. It helps to maintain homeostatic balance of sulphur pool in cell. The functional roles of sulphur also include cyanide detoxification, hydrogen sulphide detoxification, neuro-modulation, repair of iron sulphur centres, and maintenance of respiratory rates. Further study of this enzyme is desired for more insight into its role cellular metabolism and possible medical application.

1. Introduction

In order to understand the ubiquitous nature of rhodanese, it would be essential to briefly provide an insight into the chemistry of sulphur in relation to life. The iron-sulphur world theory states that sulphur was crucial for the origin of life itself (Wächtershäuser, 1988, 1992). The oxidation state of sulphur ranges from -2 (sulphide) to +6 (sulphate) and undergoes a constant cycle of oxidation and reduction reactions. Sulphur is introduced into the biomass by assimilatory sulphate reduction to the state of sulphide (an endergonic reaction) that is restricted to prokaryotes, fungi and plants, which is then incorporated into a stable organic molecule: L-cysteine (Stockdreher, 2014). This amino acid is the central precursor for other sulphur containing biomolecules in the cell, i.e. proteins, cofactors, carbohydrates, sulpholipids, vitamins and antibiotics (Brüser et al., 2000; Dahl et al., 2002; Kessler, 2006).

Lang (1933) reported that certain tissues of mammals contained an enzyme which is able to catalyze *in vitro* the reaction between thiosulphate and cyanide leading to the production of thiocyanate and sulphite. The responsible enzyme was designed 'rhodanese', from the German name for thiocyanate ('rhodanid'), with the ending 'ese' indicating that this compound was formed by enzymatic reaction (Cipollone *et al.*, 2007b). According to the official nomenclature rules, the proper name of the enzyme is thiosulphate:cyanidesulphur transferase (TST, EC 2.8.1.1), based on the reaction catalyzed *in vitro*; however, the trivial name of rhodanese is commonly used.

In vitro, rhodanese catalyzes the transfer of sulphur atom from thiosulphate to a nucleophilic acceptor like cyanide and reduced lipoate (Villarejo and Westley, 1963; Villarejo and Westley, 1963a) via a ping pong reaction with the formation of an enzyme- sulphur complex intermediate (Green and Westley 1961; Westely and Nakamoto, 1962).

The thiosulphate sulphur transferase (TST) gene encoding human rhodanese, is located on chromosome 22 (Aita *et al.*, 1997) and two single nucleotide polymorphisms (SNP), c.306A>C and c.853C>G leading to amino acid substitutions, E102D and P285A, have been reported

(Billaut-Laden *et al.*, 2006). The allelic frequencies for the SNPs in the French Caucasian population are 1 (E102D) and 5 (P285A) percent, respectively (Libiad *et al.*, 2015).

This review tries to highlight the distribution, structure, mechanism, roles in nature and medical relevance.

Distribution of rhodanese in nature

Rhodanese has been studied from numerous of sources, which include bacteria, yeast, plants, and animals. The expression of rhodanese has been detected in many tissues of animals like the liver, proventriculus, oesophagus, gizzard, cecum, brain, large intestine, duodenum, crop, spleen, trachea, pancreas, heart, kidney, lung (Dudeck et al., 1980; Westley 1981; Drawbaugh and Marrs 1987; Aminlari and Gilanpour 1991; Aminlari and Shahbazi 1994; Aminlari et al., 1994, 2000, 2002; Al-qarawi et al., 2001; Agboola et al., 2006; Baghshani and Aminlari 2009). The enzyme which was originally detected in the mitochondrion has now been found to be located in the cytosol and other organelles (Nagahara and Nishino 1996; Agboola and Okonji, 2004). Although some cyanide may be formed in vivo, this is not sufficient to explain the abundance and the ubiquity of rhodanese (Finazzi Agro et al., 1971).

The existence of rhodanese has been controvertial in many plants due to its low activity (Kakes and Hakvoort, 1992; Lieberei and Selmar, 1990). Hatzfeld and Saito (2000) were the first to isolate and characterize in plants two cDNAs encoding rhodanese isoforms in *Arabidopsis thaliana*, AtRDH1 and AtRDH2. Rhodanese purified from tapioca leaves showed properties similar to that of bovine rhodanese (Boey *et al.*, 1976; Hatzfeld and Saito, 2000).

Rhodanese activity has been demonstrated in chloroplasts from several plants and its activity correlate with the labile sulphide concentration in the plant (Tomati, 1972). Rhodanese has been purified from cabbage leaves and shown to be able to reactivate ferredoxin from apoferredoxin (Tomati *et al.*, 1974a; 1974b). Ehigie *et al.* (2013) investigated and detected the expression of rhodanese in crude plant extracts of nine randomly selected plant tubers which includes sweet potato (*Ipomoea batatas*), yellow yam, Irish potato (*Solanum tuberosum*), bitter yam (*Diascorea*

bulbifera), cocoyam, sweet yam (*Diascorea esculentum*), water yam (*Diascorea alata*) and cassava (*Manihot esculentum*).

Structure of rhodanese

Rhodanese from the mitochondrial of bovine (Bos taurus) was the first and best characterized and conventionally referred to as Rhobov (Sorbo, 1953; Westley et al., 1983; Nandi, et al., 2000). Rhobov primary structure composition is 293 amino acids residues long. The protein consists of two equally sized globular domains, the inactive N-terminal and the catalytic C-terminal rhodanese domain, showing identical α/β topology and each domain is about 120 amino acids residues long with a low sequence homology. A connecting loop separates the two domain of the enzyme (Ploegman et al., 1978a). The two domains have a highly homologous tertiary structure, but show little sequence similarity (Hatzfeld and Saito, 2000). The active site comprises of six amino acid residues that hosts at its first position the Cys residue involved in the catalytic process which is situated at the C- terminal domain (Cipollone et al., 2007b). This conserved cysteine binds the sulfane moiety of thiosulfate at the active site (Ploegman et al., 1978a). The counterpart of the Cys residue in the N-terminal domain is an Asp residue with no catalytic property (Ploegman et al., 1978b) (Figure 2.3). The uniqueness of rhodanese is resident in two patterns of amino acid residues sequences called 'rhodanese signatures', that can be recognized at the Nterminal end ([F/Y]-X3-H-[L/I/V]-P-G-A-X2-[L/I/V/F]) and at the C-terminal region of the protein ([A/V]-X2-[F/Y]-[D/E/A/P]-G-[G/S/A]-[W/F]-X-E-[F/Y/W]). The 'rhodanese signatures' have a remarkable degree of conservation amongst rhodaneses and therefore they are used for the recognition of these proteins encoded by different genomes (Cipollone et al., 2007b).

Mechanism of action of rhodanese

Rhodanese catalytic activity process occurs through a double displacement (ping-pong) mechanism involving the stable formation of a persulphide-containing enzyme intermediate (ES) (Horowitz and Criscimagna, 1983a; Pagani et al., 2000; Cipollone et al., 2004) as illustrated inFigures1, 2 and 3. It follows a non-sequential ping-pong mechanism in the generation of thiocyanate from thiosulphate and cyanide (Jack et al., 2015). The reaction proceed in two half reactions (Figures 1, 2 and 3). In the first half reaction, the sulphane sulphur is transferred from the substrate to the active site cysteine to form a persulphide enzyme intermediate, while in the second half reaction, a thiophilic acceptor attacks the enzyme-bound persulphide intermediate forming product and regenerating the free form of the enzyme (Schlesinger and Westley, 1974; Westley et al., 1983). The transferring sulphur is bound to an invariant catalytic Cys residue and is transferred formally as S^0 by nucleophilic reaction with cyanide, to yield thiocyanate thereby regenerating the active Cys residue for a new round of catalysis. The chemical species intervening in sulphur delivery is sulphane sulphur, a sulphur atom covalently bound to the sulfur atom of Cys, which has an apparent oxidation state of 0 or -1 (Horowitz and Criscimagna, 1983b; Pagani et al., 2000; Cipollone et al., 2004).



Figure 1: Representative reactions catalyzed by rhodanese. Thiosulphate is the sulphane sulphur donor and an enzymebound persulphide intermediate (E-SSH) is formed whereas

cyanide is the sulphur acceptor (Libiad et al., 2014).



Figure 2: Representative reactions catalyzed by rhodanese. GSSH is the sulphur donor and an enzyme-bound persulphide intermediate (E-SSH) is formed whereas sulphite is the sulphur acceptors (Libiad *et al.*, 2014).



Figure 3: Representative reactions catalyzed by rhodanese. Thiosulphate is the sulphane sulphur donor and an enzymebound persulphide intermediate (E-SSH) is formed whereas glutathione (GSH) is the sulphur acceptor (Libiad *et al.*, 2014)

2. Functional role of rhodanese

Cyanide detoxification

Historically, the biological importance of rhodanese was ascribed to cyanide detoxification via transfer of the sulphane sulphur from thiosulphate to cyanide to generate thiocyanate and sulphite (Figure 1). It has been understood for over eight decades that most of the sub-lethal amount of cyanide absorbed by either inhalation or ingestion is metabolized and detoxified by cyanide reaction with reactive sulphur. In eukaryotes, rhodaneses and mercaptoethanol sulphur transferases (MSTs) are postulated to be involved in detoxification of cvanide. The high concentration of tandemdomain TSTs in tissues and organs exposed to cyanide supports the hypothesis of cyanide detoxification (Sylvester and Sander, 1990). The level of rhodanese expression in different tissues of animals and plants has been found to correlate with the level of cyanide exposure (Wood, 1975; Lewis et al., 1992). Although the pattern of distribution of tandem-domain TSTs in various tissues is species specific, rhodanese activity are usually high in liver cells that are in close proximity to the hepatic blood supply, in epithelial cells covering the bronchioles (entry route for gaseous cyanide) and in proximal tubule cells of the kidney (facilitating cyanide detoxification and elimination as thiocyanate in urine) (Aminlari and Gilanpour, 1991). Rhodanese is expressed at its peak in the epithelium of rumen, omasum, and reticulum in cattle (Aminlari and Gilanpour, 1991). In these stomach compartments, cyanide is liberated after ingestion of plant cyanogenic glucosides; rhodanese activity is significantly higher in the stomach than in the liver, supporting the role of cyanide detoxification (Aminlari and Gilanpour, 1991).

However, since rhodanese activity is majorly restricted to the mitochondrial matrix where thiosulphate permeates with low efficiency (Westley et al., 1983), it has been suggested that rhodanese recruits other unknown sulphur source(s) for detoxification of cyanide (Nagahara, et al., 2003). Alternatively, based on compartmental distribution of sulphur transferases (STs), it has been suggested that rhodaneses detoxify cyanide together with mercaptoethanol sulphur transferases (MSTs) (Nagahara et al., 2003). MSTs are localised both in cytoplasm and in mitochondria (Nagahara et al., 2003). When cyanide diffuses into the cytoplasm, MSTs in the cytosol may catalyze a primary sulphuration reaction of cyanide, while the cyanide that eventually escapes this detoxification step may be catalysed by mitochondrial rhodanese, thereby preventing the lethal inhibition of cellular respiration (Nagahara et al., 2003). The therapy for acute cyanide poisoning is the intravenous administration of sodium nitrite and sodium thiosulfate Sodium combination therapy. nitrite oxidizes oxyhemoglobin to methemoglobin whose affinity for cyanide is higher compared to cytochrome c oxidase; on the other hand sodium thiosulfate is the substrate of rhodanese Moreover, (Cipollone et al., 2007b). the rapid decomposition of mercaptopyruvate intravenously administered makes it not potent for cyanide detoxification (Nagahara et al., 2003). Thus, rhodanese and MSTs specific roles in physiological detoxification of cyanide remain controversial (Cipollone et al., 2007b).

Interestingly, detoxification of cyanide which was first attributed to Rhobov in eukaryotes (Sorbo 1953; Westley et al., 1983) has also been confirmed in prokaryotes (Cipollone et al., 2007a, Cipollone et al., 2006). Cyanide detoxification was greatly increased in Bacillus stearothermophilus mutants with about 5-6 folds rhodanese activity than the wild type (Atkinson, 1975). Moreover, RhdA from the P. aeruginosa has been reported to participate in detoxification of cyanide (Cipollone et al., 2004, Cipollone et al., 2007a, Cipollone et al., 2006). Although RhdA is characterised with low affinity for both thiosulfate and cvanide in vitro, it has been demontrated to provide protection against cyanide toxicity when overexpressed in E. coli as heterologous host (Cipollone et al., 2006). The viability of P. aeruginosa growth under cyanogenic conditions is promoted by RhdA (Cipollone et al., 2007a). Hence, in vitro biochemical studies which suggest a minor role of RhdA in cyanide detoxification may underestimate the actual role of the enzyme in vivo (Cipollone et al., 2007b).

Plants are exposed to cyanide from numerous exogenous and endogenous sources (Most and Papenbrock, 2015). The major source of cyanide in the environment comes from human activities, for example, soil polluted by various industrial wastes containing up to 11,000 mg cyanide kg-DW soil (Henny et al., 1994). The natural exogenous sources include bacteria, fungi, algae, and plants that are cyanogenic in significant amounts (Most and Papenbrock, 2015). The endogenous source of cyanide in plants arise from the conversion of 1-amino-cyclopropane-1-carboxylic acid to ethylene which liberate cvanide in equimolar amounts as ethylene and is increased during ripening and senescence of the fruit (Yip and Yang, 1988). Cyanide is a potent inhibitor of cytochrome c oxidase in the respiratory chain. Plants can readily uptake cyanides available in their root zone (Doucleff and Terry, 2002). Cyanide stimulation of NADPH oxidase and inhibition of antioxidant enzymes for example catalase induces the production of reactive oxygen species (ROS) and also triggers the production of hydrogen peroxide (H_2O_2) in embryonic axes of sunflower (Helianthus annuus L.) (Oracz et al., 2009). In higher plants, two metabolic pathways are involved in the assimilation and detoxification of excess cyanide (Most and Papenbrock, 2015). Firstly, the sulphur transferase (Str) pathway as observed in bacteria and mammalians. In mammals, Strs play an important role in the catalysis of cyanide to less toxic thiocyanate that is mainly excreted in the urine (Ressler and Tatake, 2001). In Pseudomonas aeruginosa, mitochondrial rhodanese has been demonstrated to be involved in the protection of aerobic respiration from cyanide poisoning through the transfer of sulphane sulphur from thiosulphate to cyanide to produce less toxic thiocyanate (Cipollone et al., 2008). However, in plants, the contribution of Str to cyanide detoxification may be incidental or negligible (Meyer et al., 2003). More recently, it was observed that the β -cyano-L-alanine (β -CAS) pathway is the main mechanism for maintaining cyanide homeostasis in higher plants (Machingura and Ebbs, 2014). Previously, it was suggested that β -cyano-L-alanine synthase (CAS) plays a more crucial role in cyanide detoxification than Str activity in A. thaliana (Meyer et al., 2003). In β-cyano-L-alanine synthase (CAS) pathway, the cyanide is first substituted for the sulfhydryl group of cysteine to form β -CAS with the

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liberation of hydrogen sulphide (Hatzfeld *et al.*, 2000). Subsequently, the β -CAS produced is hydrolyzed by the gene product of NIT4, which is a dual enzyme complex with nitrilase and nitrile hydratase activity, generating asparagine, aspartate and ammonia, respectively (Piotrowsk *et al*, 2001; Machingura and Ebbs, 2014).

Sulphur and selenium metabolism

Rhodanese-like proteins may have a role in sulphur and selenium metabolism. The expression of the P21 rhodaneselike protein from A. ferrooxidans (a chemolithotrophic bacterium) seems to be induced during growth on various oxidizable sulphur compounds to yield thiosulphate that is used as electron donor (Ramirez et al., 2002). The disruption of the gene encoding the rhodanese-like enzyme CysA from Saccharopolyspora erythraea results in cysteine auxotrophy (Donadio, et al., 1990). However in the aforementioned cases above, there is no significant effect on the cellular TST activity. It is noteworthy to know that the pathway for cysteine biosynthesis in S. erythraea differs from that established, for example, in E. coli, just as the former includes thiosulphate as an intermediate (Cipollone et al., 2007b). Hence, P21 and CysA may perform the specific function to synthesizing thiosulphate (Ramırez et al., 2002). Ogasawara and co-workers (2001) reported that free enzyme form of bovine rhodanese can bind selenium at ratio 1:1 in vitro, resulting in the generation of the stable perselenide form (E-Se) of rhodanese. The reaction of SeO_3^{2-} and glutathione at physiologically meaningful amounts produced perselenide form (E-Se) of rhodanese (Cipollone et al., 2007b). Therefore, perselenide form of rhodanese has been proposed to be necessary to produce the reactive form of selenium for the biosynthesis of selenophosphate (SePO₃³⁻) which is the active selenium-donor compound required by prokaryotes and eukaryotes for the synthesis of SeCystRNA, the precursor of selenocysteine in selenoenzymes (Cipollone et al., 2007b).

Synthesis or repair of iron-sulphur proteins / cellular enzymatic antioxidant system

Rhodanese possibly contribute to the native architecture of reconstituted iron-sulphur protein(s) by mobilizing sulfur for the generation or repair of iron-sulphur clusters (Cipollone et al., 2007b). Tomati et al. (1974b) suggested the reactivation of ferredoxin from apoferredoxin by rhodanese. The ironsulphur clusters electron carriers of ferrodoxins, succinate dehydrogensase, and mitochondrial NADH dehydrogenase can be repaired via incubation rhodanese, a sulphur donor, a sulphur acceptor (dihydrolipoate), and an iron source (Bonomi et al., 1977; Pagani and Galante, 1983; Pagani et al., 1984). The participation of rhodanese in the formation of iron-sulfur clusters has been challenged due to the strong evidence that Nif/Isc-related proteins are involved in the mobilization of sulphur from cysteine for the synthesis of iron-sulphur clusters (Urbina et al., 2001). However, Cereda et al. (2009) in their preliminary phenotypic characterization of the gene of rhodanese from Azotobacter vinelandii (rhdA) mutant suggested that rhodanese from Azotobacter vinelandii (RhdA) may protect over Fe-S enzymes, which are labile targets for oxidative damage. Conversely, rhodanese is a constituent of the cellular enzymatic antioxidant system. The possible role of RhdA as a redox switch which helps A.vinelandii in maintaining the cellular redox homeostasis was studied through the use of an *in vitro* model system that showed reversible chemical modifications in the highly reactive RhdA Cys^{230} thiol (Cereda *et al.*, 2009).

Aerobic energy metabolism

Probably, the main physiological role of rhodanese is to supply of sulphur for the production of iron sulphur centre proteins of the respiratory chain in mammalian tissues, while cyanide detoxification may be a secondary benefit (Oke, 1973; White et al., 1981). An expression study in E. coli demonstrated that rhodanese is subject to catabolite repression and was summarised to play a role in aerobic energy metabolism (Alexander and Volini, 1987). Rhodanese forms stable complexes via disulfide bonds with membrane-bound enzymes, and catalyzes the production of iron-sulfur centers in mitochondria (Ogata and Volini, 1990). Rhodanese activity is inactivated by phosphorylation and may convert it into a sulfurase that remove the sulfur from iron-sulfur centers (Ogata et al., 1989). Rhodanese regulate the respiration rate, by controlling of the status of the iron-sulfur centers of enzymes of the electon transport chain (Ogata and Volini, 1990; Ogata et al., 1989). The rhodanese activity is regulated by а protein kinase/phosphatase and would be the terminal step of hormonal or neurotransmitter signaling pathways acting on oxygen utilization or oxidative phosphorylation pathways (Hatzfeld and Saito, 2000).

H₂S detoxification

It has been reported that rhodanese in the large intestine is the main enzyme involved in Hydrogen sulphide (H₂S) detoxification which is generated normally in the intestine (Picton *et al.*, 2002). H₂S is a potent poison normally present in the colonic lumen which may play a role in ulcerative colitis (UC) (Picton *et al.*, 2002).

Neuromodulator

Moreover, rhodanese has been detected in mouse brain (Wrobel, *et al.*, 2006), and discovered to be localized in areas that synthesize cyanide, suggesting a neuromodulatory role (Borowitz, *et al.*, 1997). Therefore, a regulatory role of rhodanese in tuning (via conversion of cyanide to thiocyanate) the neuromodulator properties of cyanide has been postulated (Cipollone and Visca, 2007a).

Mitochondrial sulphide oxidation pathway

Recently, the role of rhodanese in the mitochondrial sulphide oxidation pathway has been recognized, where it catalyzes the transfer of sulphane sulphur from glutathione persulfide (GSSH) to sulphite to form thiosulphate (Figure 2.6) (Hildebrandt and Grieshaber, 2008; Libiad *et al.*, 2014). It has also been proposed that the role of rhodanese or another thiol sulfurtransferase in the mitochondrial sulphide oxidation pathway is to utilize thiosulphate, forming GSSH instead (Figure 2.7) (Jackson *et al.*, 2012; Melideo, *et al.*, 2014). Kinetic data combined with simulations at physiological substrate concentrations have established that the first enzyme in the mitochondrial sulphide oxidation pathway, sulphide quinone oxidoreductase, preponderantly catalyzes the biosynthesis of GSSH, which is a substrate for rhodanese (Libiad *et al.*, 2014).

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3. Medical significance of rhodanese

- 1) Shahbazkia et al. (2009) related the high activity of rhodanese in testis and ovary of Felis cattus to its role in steroidogenesis through formation of iron-sulphur proteins (Shahbazkia et al., 2009). During steroidogenesis process iron-sulphur proteins are involved in the oxidative cleavage of the cholesterol side-chain occurring in ovary and testis by cholesterol side-chain cleavage (SCC) enzyme complex (Baron, 1975). The components of cholesterol side-chain cleavage (SCC) enzyme complex are the iron-sulphur proteins adrenodoxin, cytochrome P- 450, and NADPH:ISP reductase (Trzeciak, et al., 1986).
- 2) Serum rhodanese activity may be a possible candidate for liver function test. In sheep and dog rhodanese activity was elevated after induction of hepatic necrosis (Aminlari *et al.*, 1994a). Shahbazkia *et al.* (2009) suggested that serum rhodanese determination can be used as a hepatic function test in animals. It may also be applicable to humans.
- 3) Rhodanese protects against continuous oxidative stress, including that induced by radiation (Nakajima, 2015). Persistent oxidative and inflammatory reactions induce changes in hepatic metabolism and may lead to cancer and other pathophysiological conditions (Szabo and Bala, 2013; Yoshimoto *et al.*, 2013). Internal or External stresses induce continuous oxidative stresses around mitochondria that may lead to induction of rhodanese activity and the induced rhodanese, collaborating with MST, promotes sulphane sulphur, GSH, or thioredoxin regulations, resulting in activation of anti-oxidative stresses (Nakajima, 2015).
- 4) Rhodanese has also been reported to be a cancer biomarker candidate (Birkenkamp-Demtroder *et al.*, 2002; 2005). Interestingly, the activity of rhodanese has been reported to decrease in hepatic tumors, for example in Ehrlich ascites tumor-bearing mice compared to control mice (Iciek *et al.*, 2007). Probably, if its activity is restored, it may repress tumor growth. Rhodanese inducers include many nutrients and it is feasible that its induction could be readily applied clinically (Nakajima, 2015).
- 5) Rhodanese induction may protect normal tissues from acute high-dose irradiation in the case of radiocancer therapy like antioxidants, which have been investigated for their protective action (Demiryilmaz *et al.*, 2012; Kaya *et al.*, 2014)
- 6) It has been demonstrated that the decrease in rhodanese expression depicts an increase of oxidative stresses and predicts mortality in hemodialysis patients; therefore, its expression might be a prognostic indicator (Krueger *et al.*, 2010).

4. Conclusion

The relevance of rhodanese in the homeostatic balance of sulphur pool in cells of living organisms cannot be accidental, looking at the myriad of roles this enzyme is playing to the cells so as to sustain life. However, more study on this enzyme is needed to unravel in detail how rhodanese clearly perform its function inside the cellular milieu.

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