

Molecular BioSystems

Accepted Manuscript



This is an *Accepted Manuscript*, which has been through the Royal Society of Chemistry peer review process and has been accepted for publication.

Accepted Manuscripts are published online shortly after acceptance, before technical editing, formatting and proof reading. Using this free service, authors can make their results available to the community, in citable form, before we publish the edited article. We will replace this *Accepted Manuscript* with the edited and formatted *Advance Article* as soon as it is available.

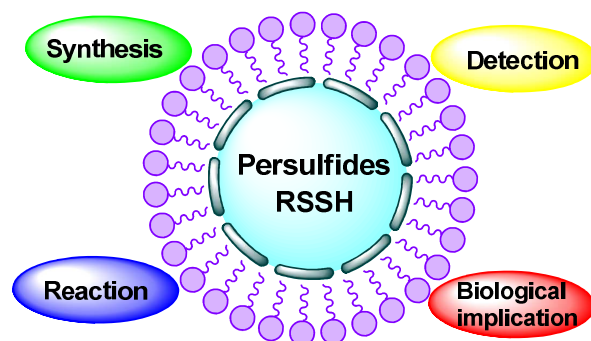
You can find more information about *Accepted Manuscripts* in the [Information for Authors](#).

Please note that technical editing may introduce minor changes to the text and/or graphics, which may alter content. The journal's standard [Terms & Conditions](#) and the [Ethical guidelines](#) still apply. In no event shall the Royal Society of Chemistry be held responsible for any errors or omissions in this *Accepted Manuscript* or any consequences arising from the use of any information it contains.



www.rsc.org/molecularbiosystems

This review article summarizes known knowledge about both small molecule persulfides and protein persulfides, including their preparation/detection methods, reactions, and biological implications.



ARTICLE

Persulfides: Current Knowledge and Challenges in Chemistry and Chemical Biology

Cite this: DOI: 10.1039/x0xx00000x

Chung-Min Park,^a Laksiri Weerasinghe,^a Jacob J. Day,^a Jon M. Fukuto,^b and Ming Xian^{a,*}Received 00th January 2012,
Accepted 00th January 2012

DOI: 10.1039/x0xx00000x

www.rsc.org/

Recent studies conducted in hydrogen sulfide (H₂S) signaling have revealed potential importance of persulfides (RSSH) in redox biology. The inherent instability of RSSH makes these species difficult to study and sometimes controversial results are reported. In this review article we summarize known knowledge about both small molecule persulfides and protein persulfides. Their fundamental physical and chemical properties such as preparation/formation and reactivity are discussed. The biological implications of persulfides and their detection methods are also discussed.

Introduction

Cysteine is a redox sensitive amino acid residue in proteins. The -SH group of cysteine is electron rich and its *d*-orbitals enable multiple oxidation states (from -2 to +6) leading to an array of redox modifications that play important roles in regulating protein structures and functions.^{1,2} A number of endogenous molecules and exogenous chemicals are known to react with cysteines. Among these, the reactions with reactive oxygen species (ROS) and reactive nitrogen species (RNS) have received considerable attention in redox biology.^{3,4} ROS mainly include superoxide (O₂^{•-}), hydrogen peroxide (H₂O₂), and hydroxyl radical (•OH). These species are generated from incomplete reduction of oxygen during normal respiration in aerobic organisms and by enzymes such as NADPH oxidase. RNS include molecules such as nitric oxide (NO) and peroxynitrite (ONOO⁻). Both ROS and RNS can react with cysteine residues in the active and allosteric sites of proteins. The resultant post-translational modifications are known to be important redox signaling mechanisms. It should be noted that not all cysteines in a protein are equally sensitive to ROS/RNS. Normally the low pK_a thiols, often known as “reactive cysteines”, are more susceptible. The low pK_a thiols are caused by local environment (such as neighboring positively charged amino acids, hydrogen bonding, and location at the N-terminal end of an α -helix) and ionized to form thiolates under physiological pH. Because of their strong nucleophilic character, thiolates are extremely vulnerable to oxidation. A number of possible thiol derivatives can be produced endogenously and the ones being most well studied include S-nitrosation (-SNO), S-glutathionylation (-SSG), and S-sulfenylation (-SOH).⁴ These oxidized species can be reduced to form free thiols (-SH) by the antioxidant defense network or converted to further oxidized forms depending on the cellular redox-state. The oxidative posttranslational modification depends on many factors including the reactivity of the

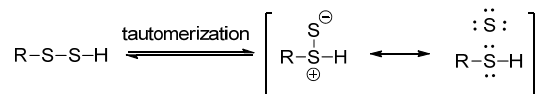
individual Cys residue, its surroundings, and the composition of the local redox-environment.

Recently a new cysteine-based posttranslational modification, i.e. S-sulphydration, has emerged as a hot research area and been linked to redox signaling. S-sulphydration was initially suggested to be the result of H₂S-mediated cysteine modification. As H₂S is considered an endogenously generated gasotransmitter,⁵⁻⁷ S-sulphydration might be a possible way by which H₂S alters the functions of a wide range of cellular proteins.^{8,9} However, recent studies reveal that other reactive sulfur species (RSS) such as sulfane sulfurs or polysulfides can also lead to S-sulphydration and these species may be even more effective than H₂S in S-sulphydration.¹⁰⁻¹⁴ In addition, the detection of S-sulphydration is still challenging. Some commonly used methods are problematic (*vide infra*). As such, more work is needed to elucidate how endogenous S-sulphydration occurs and what their functions/roles are in biology. The products of S-sulphydrations are cysteine persulfides (R-S-SH), also referred to as perthiols or hydrodisulfides. Unlike their thiol analogs (RSH), the fundamental chemistry and chemical biology of persulfides are still poorly understood. Two reasons contribute to this lack of knowledge: 1) the biological significance of RSSH has only been recently discovered; and 2) persulfides are unstable species, especially in aqueous buffers, which make them difficult to study. Persulfides readily decompose in a pH-dependent manner and possible products include the corresponding thiol, disulfides, polysulfide, and elemental sulfur (S₈). H₂S can also be a decomposition product. In this review article, we wish to summarize known information about persulfides. The preparation/formation, reactivity, and biological implication of both small molecule persulfides and protein persulfides will be reviewed.

General physical/chemical properties of persulfides

Persulfides have significantly different chemical properties compared to their structurally related thiol analogs (RSH). For example, the difference in S-H bond energies for persulfides versus thiols is significant.¹⁴⁻¹⁷ The disulfide bond in persulfides weakens the RSS-H bond by ~22 kcal/mol relative to RS-H. This energy difference can be attributed to the increased stability of the perthiyl radical (RSS[•]) compared to thiyl radical (RS[•]). The adjacent sulfur atom can stabilize the odd electron of RSS[•] via resonance effect while this is unavailable for RS[•]. The perthiyl radical has an absorbance at 374 nm ($\epsilon \sim 1700 \text{ M}^{-1}\text{cm}^{-1}$) which can be used to determine the presence of perthiyl in reactions.¹⁸

The pKa of persulfides is approximately 6.2, which makes them stronger acids compared to thiols (pKa ~7.6).¹⁹ At physiological pH, thiols predominantly exist the protonated forms but persulfides (RSSH) exist as the deprotonated anions (RSS⁻). Therefore, the conversion of cysteine residues in proteins to persulfides (i.e. S-sulfhydration) should increase the nucleophilicity of the reactive sulfur atom and this may be associated with higher reactivity as reducing agents. It is believed that tautomerization of persulfides exists, leading to the formation of thiosulfoxides (RS(=S)H) (Scheme 1).¹⁴ Tautomerization would generate a reactive ‘singlet sulfur’ that would be very electrophilic. However, thiosulfoxides, for the most part, are thought to be thermodynamically uphill compared to the persulfide tautomer. Persulfides belong to the sulfane sulfur family (note: sulfane sulfur is defined as a sulfur bonded sulfur atom formally with six electrons²⁰ – e.g. the terminal sulfur atom of a persulfide is a sulfane sulfur). The characteristic reaction with CN⁻ to form SCN⁻ and RSH has been used as evidence for the existence of a persulfide.

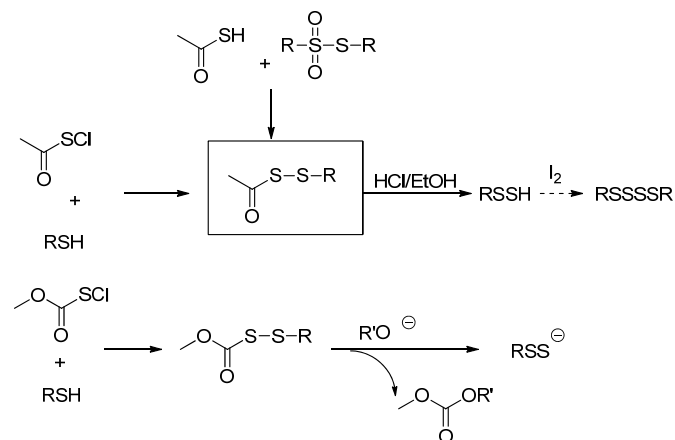


Scheme 1 Tautomerization between persulfides and thiosulfoxides

Small molecule persulfides: synthesis and reactions

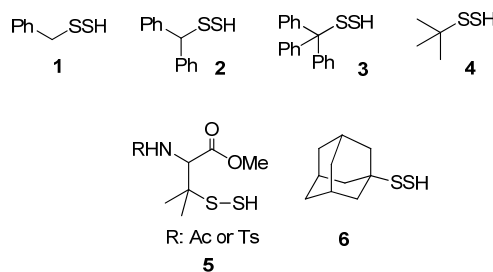
In 1954, Böhme and Zinner reported the first preparation of aryl- and alkyl-persulfides (R-SSH).²¹ The key intermediates were acyl disulfides, which were prepared from sulfonyl chloride (RC(=O)SCl) and mercaptans (Scheme 2). Acid (usually HCl) mediated hydrolysis of acyl disulfides provided the persulfide products, which were quantified as the stable tetrasulfide upon iodine oxidation. So far, acid-mediated hydrolysis of acyl disulfides has become a general method for the preparation of small molecule persulfides, while different methods have been developed to prepare the acyl disulfide intermediates. For example, alkyl persulfide has also been synthesized from both dialkyl thiosulfones with thioacid.²² Persulfides can also be prepared from alkoxide-induced displacement of the RSS⁻ anion from methoxycarbonyl disulfides (RSSO₂CH₃),¹⁹ which in turn can be prepared from

the condensation of thiol with methoxycarbonyl disulfonyl chloride (Scheme 2).



Scheme 2 Persulfide generation reactions

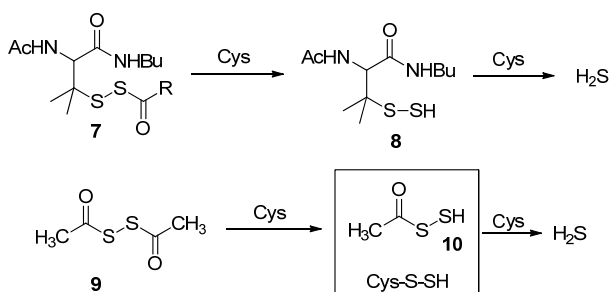
So far a limited number of small molecule persulfides have been prepared and some of them are only partially characterized. The structures shown in Scheme 3 are most often seen in literature, which include benzyl/diphenylmethyl/trityl persulfides (**1-3**)²³⁻³⁶, *t*-butyl persulfide **4**^{36,37}, penicillamine-derived persulfides **5**^{38,39}, and adamantyl persulfide **6**.⁴⁰ In general more sterically hindered persulfides are more stable. For example, adamantyl persulfide can be purified by distillation and stored in a sealed ampule under ambient conditions for several months. Aryl persulfides (Ar-SSH) were also reported⁴¹ but these compounds seem to be less stable compared to the ones shown in Scheme 3 and not well characterized.



Scheme 3 Representative small molecule persulfides

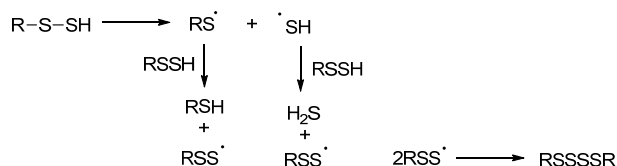
Persulfides are in general transient species. Both acidic and basic conditions were found to promote the decomposition.³⁹ The major decomposition products are polysulfides (RSS_nSR). H₂S was reported to be a by-product of decomposition in some studies^{35,39,42} while other studies claimed H₂S was not the product.⁴³ Solvents or reaction conditions may determine the decomposition products. Unstable persulfides can serve as potential H₂S donors. For example, penicillamine-derived acyl disulfides **7** can release H₂S in the presence of cysteine or glutathione (Scheme 4).⁴⁴ In this process penicillamine persulfide **8** is the key intermediate. Similarly dithioperoxyanhydride **9** was proved to be another type of thiol-

activated H₂S donor with acyl persulfide **10** and cysteine persulfide proposed to be the active intermediates.⁴⁵



Scheme 4 Persulfide intermediates as H₂S donors

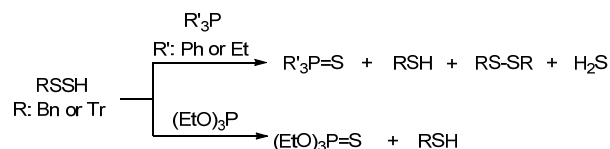
Persulfides are excellent radical precursors allowing some of their reactions to proceed via free radical mechanisms. For example, the thermal decomposition in organic solvents is initiated by homolytic dissociation of the S-S bond into two thiyl radicals (RS[•] and HS[•]), which further react with persulfides to form the final products: H₂S, thiol, and polysulfide (Scheme 5).³⁵ Persulfides can also undergo hydrogen transfer (from their RSSH forms) or electron transfer (from their anionic forms RSS⁻, normally under physiological pH) to generate perthiyl radicals (RSS[•]) that undergo subsequent reactions. An excellent review by Everett and Wardman summarized some important radical scavenging reactions of persulfides as antioxidants.¹⁸



Scheme 5 Formation of perthiyl radicals

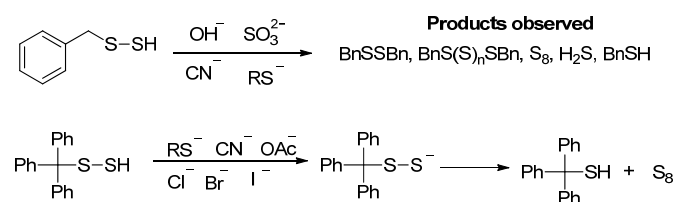
Reactions with phosphines/phosphites: Persulfides can be considered as the thio-analogs of hydroperoxides. Their reactions with tertiary phosphines (PPh₃ or PEt₃) were studied and compared with that of hydroperoxides.²⁴ Since empty *d*-orbitals of the sulfur atom may participate in bond formation by accepting electrons from phosphorus, sulfur should be more susceptible to attack by phosphine than oxygen. Such reactions led to the corresponding phosphine sulfide and thiol as the major products (Scheme 6),²⁴ along with small amounts of disulfides and H₂S. Both sulfur atoms in persulfide could be attacked by the phosphine. Steric effects in the persulfide and thiophilicity of the phosphines are the factors governing which sulfur atom is attacked. In another study, phosphites (such as P(OEt)₃) were found to react with persulfides in a similar way, producing the corresponding thiol and phosphorothionate as the major products.³³ In an analogous study the reactions between persulfides and tertiary arsines (Ph₃As or Et₃As) were tested and the corresponding thiols (RSH) and arsine sulfides were the major products.³² It should be noted that persulfides belong to the sulfane sulfur family (also include polysulfides and

elemental sulfur). The reaction of phosphines to form phosphine sulfide is generic for sulfane sulfurs. This reaction has been used to detect sulfane sulfurs in biological samples.^{46,47}



Scheme 6 Reactions of persulfides with phosphines and phosphites

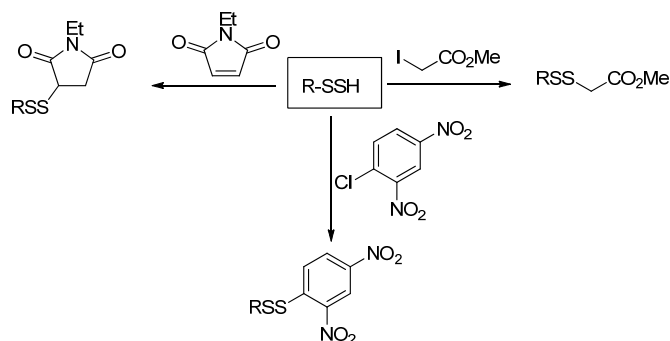
Reactions with nucleophiles: The reactions between RSSH and nucleophiles have been well studied.^{25,48,49} Theoretically four pathways are possible when RSSH is treated with a nucleophile: 1) deprotonation of the sulfhydryl group; 2) nucleophile attacking the outer sulfur atom; 3) nucleophile attacking the inner sulfur atom; and 4) deprotonation of the alpha-carbon. The reactions of benzyl persulfide (BnSSH) with various nucleophiles such as hydroxide, sulfite, cyanide, and thiolate, were first studied.⁴⁹ Products were H₂S, benzyl disulfide, benzyl polysulfide, and elemental sulfur (Scheme 7). Benzylthiol (BnSH) was also found for cyanide and thiolate. It was postulated that less nucleophilic reagents (hydroxide and sulfite) preferably attack the inner sulfur atom while the stronger nucleophilic reagents (cyanide and thiolate) may attack both sulfur atoms. Steric hindrance on the persulfide substrates should also be considered. In the case of diphenylmethyl persulfide (Ph₂CHSSH), attack on the outer sulfur was usually more predominant. Interestingly a recent work by Pluth et al found that a more steric hindered persulfide-Ph₃CSSH did not undergo nucleophilic reactions with nucleophiles (such as thiolate, CN⁻, I⁻, Br⁻, AcO⁻).³⁰ In organic solvents, these nucleophiles promoted deprotonation of the persulfide to generate Ph₃CSS⁻, which then disproportionated into Ph₃CSS⁻ and S⁰. The corresponding thiol (Ph₃CSSH) and S₈ were found to be the final products.



Scheme 7 Reactions of persulfides with various nucleophiles

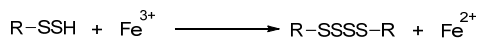
Amines can act as bases and nucleophiles in the presence of persulfides. The reactions between persulfides (mainly PhCH₂SSH and Ph₂CHSSH) and amines (aniline, pyridine, butylamine, morpholine, etc.) were studied by Tsurugi et al.²⁶ Products were found to be H₂S, disulfide, polysulfide, thiol, and S₈. It was concluded that weaker bases (pK_a < 5.7) behave as nucleophiles while stronger bases (pK_a > 8.2) behave as bases. For sterically hindered persulfide like Ph₃CSSH, amines mainly acted as bases. Deprotonation and subsequently disproportion led to the thiol and S₈ as the major products.³⁰

Reactions with electrophiles: Although persulfides (RSSH) are believed to be more nucleophilic than thiols (RSH) their reactivity toward electrophiles have not been well studied. A large number of thiol (-SH) blocking reagents such as methyl methane sulfonate (MMTS), iodoacetamide (IAM), N-ethylmaleimide (NEM), etc. are known, so all of these compounds should also react with RSSH. However, very few such reaction examples can be found in literature. So far only NEM³⁰, 2,4-dinitrochlorobenzene (DNCB)^{40,50}, and iodoacetate²⁷ have been reported to derivatize RSSH (Scheme 8). Surprisingly in many cases the yields of the products formation were not reported, perhaps due to low isolated yields. It was found that the reaction between electrophiles (such as NEM) and RSSH was also rather slow.³⁰ The addition of bases (such as Et₃N) could drastically increase the reaction rate. However bases could promote the decomposition of RSSH, presumably leading to decreased product yields.



Scheme 8 Reactions of persulfides with selected electrophiles

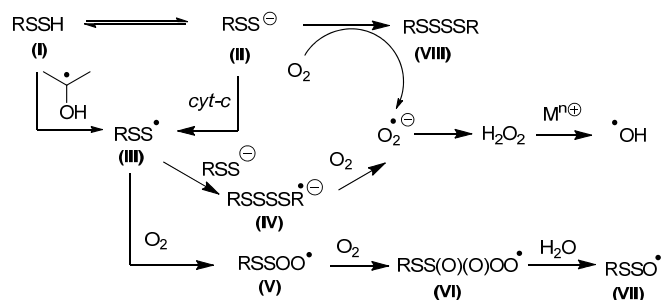
Reactions with iron salts: The redox reactions of persulfides with iron salts were studied and compared with hydroperoxides (ROOH).⁵¹ In the absence of oxygen, the reaction of persulfides led to the disappearance of the color of ferric ion. Tetrasulfides were obtained in good yields (Scheme 9). The redox behavior of persulfides is clearly different from that of hydroperoxides. The former acts as a reductant, which easily reduces ferric ion, while the latter is an oxidant.



Scheme 9 Persulfide reaction with iron (III)

Persulfides as pro-oxidants and antioxidants: The biological relevance of persulfides has been discussed in terms of persulfides being both pro-oxidative and anti-oxidative species (summarized in Scheme 10). The predominant anion form (II), at physiological pH, is expected to be oxidized by the molecular oxygen to form reactive oxygen species ($\text{O}_2^{\bullet-}$, H_2O_2 and HO^\bullet).^{27,52} These reactive oxygen radicals were found to induce DNA damage in the Gates's Leinamycin model.^{53,54} Furthermore Fe(III)-cytochrome *c* mediated-oxidation of II formed the more thermodynamically stable perthiyl radical (III).⁵⁵ This radical III can lead to the generation of superoxide radical anion via the formation of a putative tetrasulfide radical

anion (IV). However, no experimental evidence has been reported supporting the existence of such a species.⁵⁶ Molecular oxygen can react with radical III to generate perthiyl peroxy radical species V, VI and VII. The oxidative properties attributed with those species could cause biological damage.⁵⁷



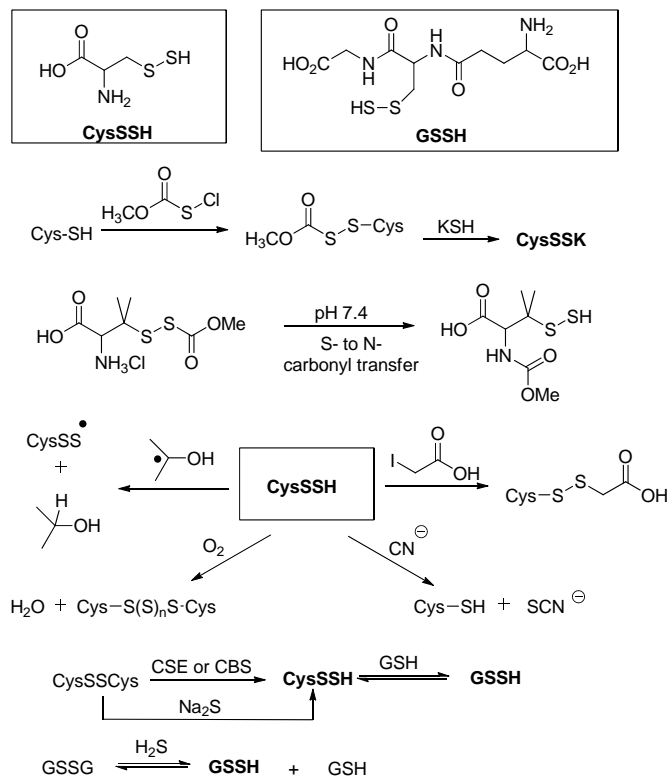
Scheme 10 Persulfides as pro-oxidants and antioxidants

The radical scavenging role of persulfides was also reported and based on the ability of RSSH to donate a hydrogen atom.¹⁸ The more stable perthiyl radical is responsible for H-atom donation and reductive electron transfer to an oxidant. Therefore, persulfides are stronger antioxidants than thiols. At lower pH RSSH (I) could donate hydrogen to a carbon radical, which is responsible for its anti-oxidant property.¹⁹ However the diminished rate of hydrogen transfers to $(\text{CH}_3)_2\text{COH}$ radical with increasing pH is evidence that under physiological pH, hydrogen donation by RSSH may not be an efficient process. The formation of more stable radical (III) from the anion (II) via electron donation could be dominant at physiological pH. Therefore RSSH's antioxidative properties could still be the result of electron donation to oxidizing species. Indeed, Fukuto et al found that GSSH has much stronger reducing ability for ferric-cytochrome *c* than H_2S and GSH.⁵⁸

Cysteine and glutathione persulfides (CysSSH, GSSH)

The existence of cysteine persulfide (also known as thiocysteine) has been recognized for a long time. CysSSH is suggested to be the product of cystine (CysSSCys) upon reacting Na_2S or with pyridoxal and Cu^{2+} .^{59,60} It is also an intermediate in the cystathionase catalyzed degradation of the substrate cystine.^{61,62} Synthetically Rao and Gorin reported that Na_2S can react with cystine under strong basic conditions to form CysSSH.⁶³ Smith et al reported a method to prepare CysSSH by treating methoxycarbonylcysteine disulfide with potassium hydrosulfide (KSH) (Scheme 11).⁶⁴ A unique S- to N- carbonyl transfer was also used by Galardon et al to prepare cysteine persulfide analogs (penicillamine persulfides).⁴³ CysSSH appears to be quite unstable. Polysulfides, cysteine, and S_8 are found to be the decomposition products. It should be noted that H_2S was not found to be the decomposition product but the presence of thiols can lead to H_2S formation.^{43,44} Theoretically the reactivity of CysSSH should be similar as other small molecule persulfides. For example, CysSSH can be

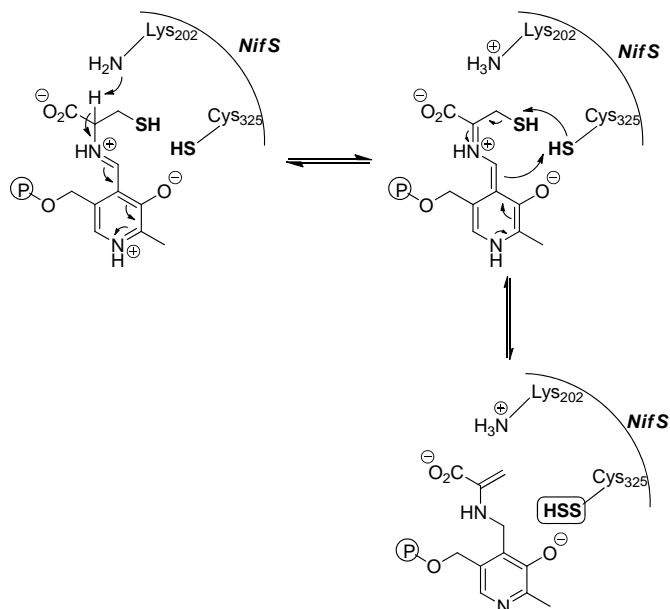
trapped by iodoacetate to form the corresponding disulfide derivative.⁶¹ Endogenous electrophiles like 8-nitro-cGMP can also trap CysSSH or GSSH.¹² Some reactions mentioned here are summarized in Scheme 11.



Although CysSSH has been known for a long time, its significance in thiol-related redox biology has been recognized only recently. Ida et al. described a mass spectrometric method to quantitate persulfide (RSSH) in cells.¹² With this detection method, they demonstrated the high levels (up to > 100 μM) of cysteine- and glutathione-persulfides in cells, tissues, and plasma. They showed that cystathionine- β -synthase (CBS) and cystathionine γ -lyase (CSE) can convert cystine to CysSSH, and subsequently lead to the formation of GSSH and polysulfides. GSSH can also be produced by glutathione reductase (GSR)-mediated glutathione polysulfide reduction. Glutathione (GSH) is a potent antioxidant in cells. However its antioxidant activity is typically mediated by specific enzymes such as GSH-dependent peroxidase. Without this assistance, GSH is relatively inert with low nucleophilicity and reacts poorly with electrophilic oxidants like H_2O_2 . However GSSH has much stronger nucleophilic/antioxidant activity.^{12,58} The strong H_2O_2 -scavenging activity of GSSH was confirmed by Ida et al.¹² Thus the presence of CysSSH and GSSH may provide a primary and potent antioxidant defense in cells. In addition, the high in vivo concentrations of persulfides may indicate that these species are the real players in cellular signaling and regulation.

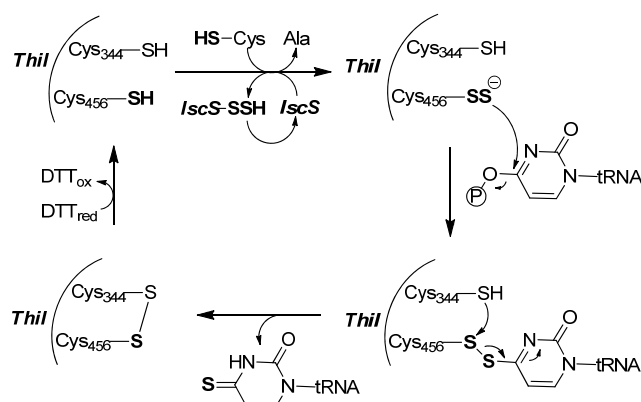
Persulfides in biosynthesis of sulfur-containing biomolecules

Sulfur is an important element of living systems. Strategies to incorporate sulfur atom into biomolecules involve biosyntheses of iron sulfur clusters, cofactors such as thiamin, molybdopterin, biotin and lipoic acid, and the thio-modification of tRNA via enzymatically generated persulfide species at enzyme active sites. In these processes the persulfide adducts in proteins are generated by cysteine desulfurases (such as NifS, IscS, SufS) and sulfurtransferases (like rhodanese and mercaptopyruvate sulfurtransferase).^{65,66} Cysteine desulfurases are pyridoxal 5'-phosphate (PLP)-dependent enzymes that catalyze the conversion of Cys to Ala and sulfane sulfur via the formation of a protein-bound persulfide intermediate on a conserved cysteine residue. NifS was the first cysteine desulfurase found to generate an enzyme-based persulfide as a crucial intermediate. NifS provides a sulfur atom for the iron-sulfur clusters in nitrogenase.⁶⁷ The catalytic reaction of NifS is initiated by Schiff base formation between cysteine and pyridoxal 5'-phosphate (PLP) (Scheme 12). Lys202 and Cys325 residues work together and transfer the sulfur atom from Schiff base adduct to Cys325 to form a persulfide. The terminal sulfur can then be transferred to other acceptor proteins and subsequently produce sulfur-containing molecules.



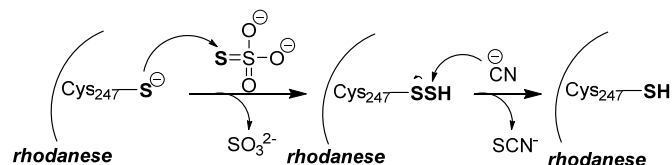
IscS is biologically similar as NifS.^{68,69} IscS is needed for the synthesis of 4-thiouridine ($s^4\text{U}$) in tRNA as well as the incorporation of sulfur into the thiazole ring of thiamine.^{70,71} The synthesis of $s^4\text{U}$ in tRNA is initiated by the transfer of cysteine sulfur to IscS, forming an IscS persulfide (like NifS-persulfide formation). The IscS persulfide then transfers the sulfhydryl sulfur to Cys465 of Thil (a rhodanese-like sulfurtransferase), forming a Thil persulfide.⁷⁰ In the presence of ATP-Mg, the activated uridine residue reacts with Thil-persulfide (Cys456-SSH) to form a disulfide intermediate and expel AMP. Another active site cysteine (Cys344) then attacks

the Thi1-tRNA disulfide to produce s^4U in tRNA and form an enzymic disulfide bond, which can be reduced for another round of catalysis.⁷² (Scheme 13).



Scheme 13 Enzymatic formation of Thi1 persulfide and the sulphur transfer into tRNA

Persulfide residues are also found in the active sites of rhodanese and similar enzymes (Scheme 14). Thiosulfate (SSO_3^{2-}) is the common substrate. The formation of persulfide was confirmed by X-ray crystallography.⁷³ The reaction between persulfide and cyanide (CN^-) to form thiocyanate (SCN^-) accounts for detoxification of rhodanases.⁷⁴⁻⁷⁶



Scheme 14 Generation of rhodanese persulfide and the reaction with cyanide

Protein persulfide formation and H_2S -related signaling

Persulfide formation on specific proteins (also known as S-sulfhydration or S-persulfidation) is a recently defined oxidative posttranslational modification that leads to either activation or inhibition of protein activity by regulating protein structures and ultimately their functions. This concept was formulated along with the characterization of H_2S as a signaling molecule and S-sulfhydration is believed to be involved in H_2S -based signal transduction. In this regard, the mechanism behind S-sulfhydration is still poorly understood. Some hypotheses have been proposed. It is well accepted that H_2S cannot directly react with protein cysteine residues (-SH) to form SSH in the reducing intracellular environment. The oxidative forms of cysteine, for example, sulfenic acid (-SOH), disulfide (-SS-), nitrosothiol (-SNO), could react with H_2S to form persulfides. Hydrogen polysulfides (H_2S_n), formed by two-electron oxidation of H_2S are possible cellular sulfhydration reagents. Other cellular sulfane sulfur species accumulated with H_2S biosynthesis or sulfur catabolism may also play a role in S-

sulfhydration.⁷⁷ Nevertheless a large number of S-sulfhydrated proteins have been identified so far and sulfhydration-mediated functional changes have been studied. For example sulfhydration induced by transfection of HEK293 cells with CSE was studied by Snyder et al.⁷⁸ Three S-sulfhydrated proteins were characterized: glyceraldehyde-3-phosphate dehydrogenase (GAPDH), β -tubulin, and actin. Sulfhydration altered actin-dependent cytoskeletal rearrangements in HEK293 cells and enhanced actin polymerization. In another study the same group found that S-sulfhydration of the p53 subunit of nuclear factor κB (NF- κB) facilitated its translocation to the nucleus and increased interaction with ribosomal protein S3 (rps3), resulting in enhanced transcription of anti-apoptotic gene.⁷⁹ Wang et al reported that S-sulfhydration of Kelch-like ECH-associated protein-1 (Keap1) at Cys151 stimulated its dissociation from nuclear factor erythroid 2-related factor 2 (Nrf2).⁸⁰ This process enables Nrf2 to translocate to the nucleus and regulates the expression of cytoprotective genes. Tonks et al suggested that S-sulfhydration of protein tyrosine phosphatase 1B (PTP1B) inhibited its activity, thus regulating the endoplasmic reticulum (ER) stress response.⁸¹ Yang et al discovered that S-sulfhydration at Cys611 and Cys614 located in the DNA binding domain (DBD) of androgen receptor (AR) inhibited its transactivation.⁸² It was proposed that S-sulfhydration of AR might shatter zinc-sulfur clusters and cause a structural modification of AR, resulting in inhibition of the formation of abnormal AR dimerization and its DNA binding. Table 1 summarizes selected S-sulfhydrated protein examples. It is expected that more S-sulfhydrated proteins will be discovered.

Table 1 Examples of sulfhydrated proteins: outcomes and detection methods

Proteins	Outcomes of sulfhydration	Detection methods
GAPDH ^{78,83}	activation of GAPDH	^a MBS, ^b TS, ^c LC-MS
Actin ⁷⁸	activation of actin polymerization	MBS
Kir6.1 subunit of K_{ATP} ⁸⁴	reducing ATP binding, activation	MBS
Hsp70 ⁸³	N/A	TS
Protein disulfide isomerase ¹²	N/A	TS
Aldo-keto reductase ¹²	N/A	TS
Phosphoglycerate kinase ¹²	N/A	TS
NF- κB ^{79,85}	activation of RPS3 binding, increasing transcription of genes	MBS, LC-MS
Keap1 ^{80,86}	activation of Nrf2	MBS

Parkin ⁸⁷	activation of parkin ligase	MBS, LC-MS
Androgen receptor ⁸²	inhibition of its dimerization	MBS
MEK1 ⁸⁸	activation of PARP-1	MBS
eNOS ⁸⁹	Activation of its dimerization	MBS
Ca ²⁺ TRP channels ⁹⁰	activation of Ca ²⁺ influx	MBS
IK _{Ca} ⁸⁴	activation of IK _{Ca}	MBS
P66Shc ⁹¹	Inhibition of PKCβII binding	MBS
Cu/Zn-SOD ⁹²	inhibition of its oxidation-induced aggregation	Cyanolysis, ^d Maldi-MS
Papain ⁵⁸	Inhibition of activity	Cyanolysis
Platelet proteins ⁹³	activation of its activity	MBS

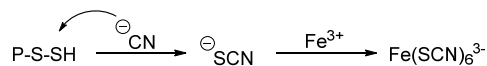
^aMBS: modified biotin switch assay, ^bTS: tag-switch assay, ^cLC-MS/MS: liquid-chromatography mass spectrometry, ^dMaldi MS: matrix-assisted laser desorption/ionization mass spectrometry

Methods for the detection of protein persulfides

Persulfide residues in proteins seem to have better stability than those in small molecules, presumably due to steric hindrance of second order decomposition pathways. Nevertheless the detection of protein persulfides is still challenging, especially in intact cells. One should consider similar reactivity of other sulfur species when applying certain methods in analysis, such as being a nucleophile similar to thiols (-SH) and being an electrophile similar to S-nitrosothiols (-SNO) and sulfenic acids (-SOH). This section summarizes currently available methods for identifying protein persulfides.

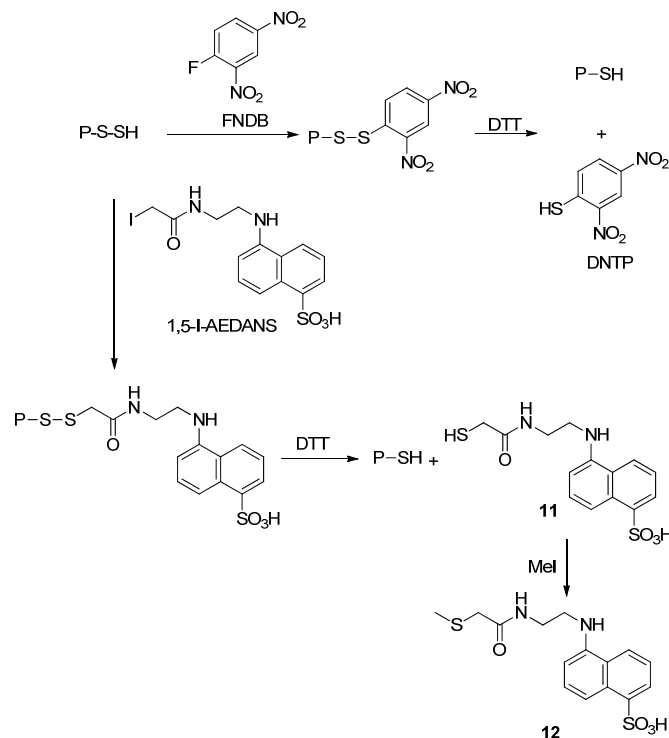
Cyanolysis: Cyanolysis was the first method that was used in the identification of protein sulfides. For example, the presence of persulfides in the active sites of xanthine oxidase^{94,95} and aldehyde oxidase⁹⁶ was confirmed by cyanolysis. In this method, thiocyanate (SCN⁻) is formed by nucleophilic attack of cyanide anion (CN⁻) on the terminal sulfur (Scheme 15). The thiocyanate formed can be determined spectrophotometrically following reaction with ferric nitrate reagent.⁹⁷ However, the nucleophilic attack of CN⁻ may also occur on the inner sulfur, releasing the terminal sulfur as sulfide (HS⁻). This competing reaction may cause underestimation of the persulfides in proteins. Another problem is that cyanolysis cannot distinguish persulfides and polysulfides as polysulfides can also react with CN⁻ to form SCN⁻. As such, this method may also cause

overestimation of persulfides. Moreover this method is usually used for individual/purified proteins, and cannot be used for identifying persulfide-containing proteins in complex systems like cell lysates.



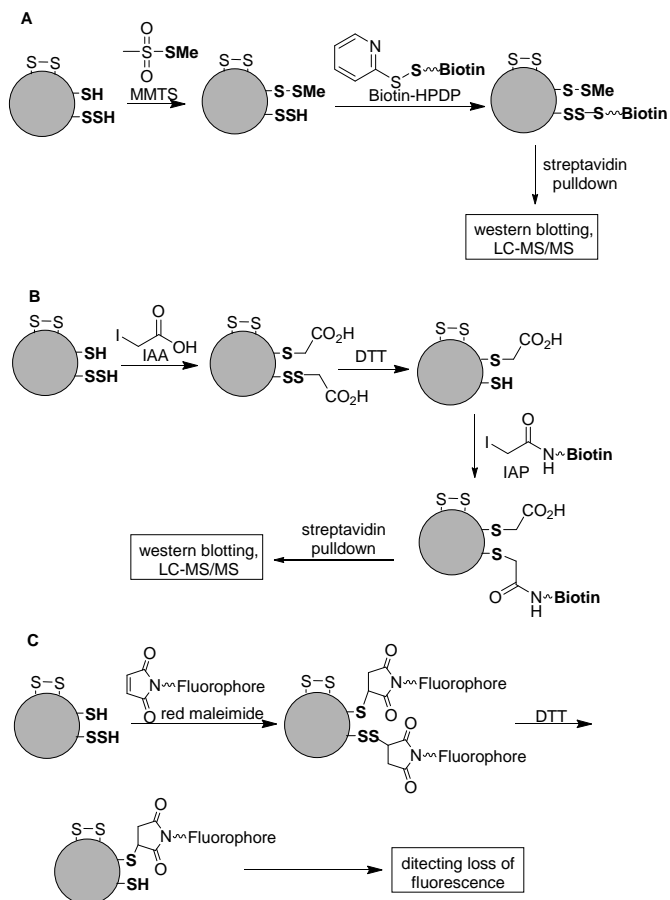
Scheme 15 Detection of protein persulfides by cyanolysis

2,4-Dinitrothiophenol (DNTP) based methods: The presence of persulfides in proteins can also be detected by persulfide-mediated nucleophilic aromatic substitution. 1-Fluoro-2,4-dinitrobenzene (FDNB) was a probe based on such a reaction.⁹⁸ As shown in Scheme 16, this method involves derivatization of protein SSH with FDNB to form a dinitrobenzene disulfide, which is then treated with dithiothreitol (DTT) to release DNTP. The concentrations of DNTP can be measured by spectroscopic methods and reflect the original persulfide concentrations. This technique can distinguish persulfides from thiols (-SH), thioethers, and polysulfides. However it is again limited to individual and purified proteins. Other thiol-alkylation reagents can also be used in similar derivatization-cleavage sequences to identify protein persulfides. For example, *N*-(iodoacetaminoethyl)-1-naphthylamine-5-sulfonic acid (1,5-I-AEDANS) was used to identify NifS-bound persulfide.⁹⁹ This reagent blocked NifS-SSH to form a disulfide product, which was then cleaved by DTT to give NifS-SH and a small molecule thiol **11**. Methylation of **11** provided a stable product **12** for quantification.



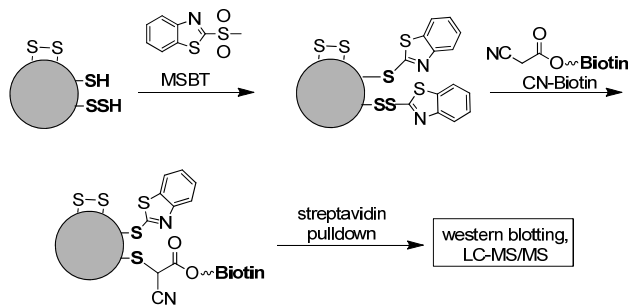
Scheme 16 Trapping protein persulfides by aromatic substitution with FDNB and alkylation with 1,5-I-AEDANS

Modified biotin switch assay: In order to identify and isolate S-sulfhydrated proteins in cell lysates a modified biotin-switch assay was reported by Snyder et al in 2009.⁷⁸ The goal of this method is to label protein persulfide residues with detectable molecules like biotin. As shown in Scheme 17-A, the first step is to block protein free thiols (-SH) with methyl methane sulfonate (MMTS). It was claimed that persulfides did not react with MMTS but could be tagged by a thiol specific biotinylating agent, biotin-HPDP (N-[(6-biotinamido) hexyl]-3'-(2'-pyridyldithio) propionamide) which seems improbable. This method allows capture and identification of S-sulfhydrated proteins by immunoblotting or LC/MS analysis. Using this method, proteins in mouse liver were analyzed and 25-50% of hepatic proteins were found to be sulfhydrated. The three most prominent proteins were glyceraldehyde-3-phosphate dehydrogenase (GAPDH), β -tubulin, and actin.⁷⁸ Despite these interesting findings, the specificity of the method is questioned as it is hard to justify the selectivity of MMTS for thiols (-SH) vs persulfides (-SSH). It is normally accepted that SSH should also be reactive for thiol-blocking reagents. Carroll et al proved that protein persulfides could react very effectively with commonly used thiol-blocking reagents including IAM, NEM, MMTS, 5,5'-dithiobis-(2-nitrobenzoic acid) (DTNB), *N*-acetylcysteine pyridyldisulfide (NACP).¹⁰⁰ These results suggest that protein persulfides behave quite differently from small molecule persulfides in reactions with electrophiles (see Scheme 8 and related discussions). Perhaps protein persulfides are more stable and their self-decomposition is less likely to happen. In another study Tonks et al used a different biotin switch strategy for analyzing S-sulfhydrated proteins.⁸¹ They first blocked both thiols and persulfides with iodoacetic acid (IAA). Then the disulfide adducts formed on persulfides were reduced by dithiothreitol (DTT) to give free thiol groups, and subsequently labeled with iodoacetamide-linked biotin (IAP) (Scheme 17-B). Using this method protein tyrosine phosphatase 1B (PTP1B) was found to be reversibly inactivated by H₂S, resulting in regulating the endoplasmic reticulum (ER) stress response. More recently, Snyder et al reported a similar method employing fluorescent maleimide to detect and label persulfides in proteins.⁷⁹ As such, the proteins of interest were immunoprecipitated and treated with a fluorescent derivative of maleimide, which selectively labeled both free thiols and persulfides. One portion of the reaction mix was then treated with DTT to release the maleimide moiety, resulting in a decrease in fluorescence. The samples were run on a polyacrylamide gel and scanned to detect the fluorescent proteins. The decrease in fluorescence could be quantified using an image analysis software.



Scheme 17 Modified biotin switch assays for S-sulfhydration detection

Tag-switch assay: Another method suitable for proteomic studies of protein S-sulfhydration is the tag-switch assay.^{12,83} A two-step sequence can selectively label persulfides. In the first step, free thiols (-SH) and persulfide (-SSH) are blocked by methylsulfonyl benzothiazole (MSBT) to give the corresponding thioether (-S-BT) and disulfide (-SS-BT) adducts. Benzothiazole-containing disulfides are highly reactive towards certain carbon nucleophiles. Therefore a tag-switch step using cyanoacetate-biotin conjugate can convert the benzothiazole disulfides to biotin tags. Benzothiazole thioesters are not affected in this step. Finally the biotinylated proteins can be captured by streptavidin agarose beads for pull-down and detected by Western blotting or LC/MS. The selectivity of this method was confirmed by Gpx3 and BSA, two established protein-SSH models.⁸³ The method was also applied in identification of endogenous S-sulfhydrated proteins in A549 cells.¹² The proteomic analysis was able to identify a series of sulfhydrated proteins including protein disulfide isomerase, heat shock proteins, aldo-keto reductase, GAPDH, enolase, and phosphoglycerate kinase.



Scheme 18 Tag-switch assay

Summary

The discovery that various persulfide species (i.e. CysSSH, GSSH, protein-SSH, etc) are formed in living organisms and exert specific roles in redox signaling is rather attractive. This review presents some general descriptions of persulfide formation and reactivity from small molecules to proteins. Despite recent advances in understanding persulfide chemistry and chemical biology, a number of important questions remain unanswered, such as what the intracellular targets are and to what extent such reactions can impact signaling. The mechanisms that confer sulfhydration specificity (i.e. what proteins are sulfhydrated and why) also needs to be addressed as well as mechanisms of the reversal of sulfhydration back to the thiol species. Further understanding of biological roles of persulfides will require a better understanding of their fundamental chemistry (which is currently underdeveloped). Chemical tools that allow easy and reliable access to persulfides (both in small molecules and proteins) should be critical for this field. Novel and ‘easy to use’ detection methods for protein S-sulfhydration are also necessary. There is little doubt that the general area of investigation of persulfide chemistry and chemical biology will be a topic of significant research interest for many years to come.

Acknowledgements

This work is supported by an American Chemical Society-Teva USA Scholar Grant and NIH (R01GM088226 and R01HL116571).

Notes and references

^a Department of Chemistry, Washington State University, Pullman, WA 99164, United States. Fax: +1-509-335-8867; Tel: +1-509-335-6073; E-mail: mxian@wsu.edu

^b Department of Chemistry, Sonoma State University, Rohnert Park, CA 94928, United States.

1. K. G. Reddie and K. S. Carroll, *Curr. Opin. Chem. Biol.*, 2008, **12**(6), 746-754.
2. R. Wani, A. Nagata and B. W. Murray, *Front Pharmacol*, 2014, **5**(224), 1-8.
3. H. S. Chung, S. B. Wang, V. Venkatraman, C. I. Murray and J. E. Van Eyk, *Circ. Res.*, 2013, **112**(2), 382-392.

4. C. E. Paulsen and K. S. Carroll, *Chem. Rev.*, 2013, **113**(7), 4633-4679.
5. C. Szabo, *Nature reviews. Drug discovery*, 2007, **6**(11), 917-935.
6. L. Li, P. Rose and P. K. Moore, *Annu. Rev. Pharmacol. Toxicol.*, 2011, **51**, 169-187.
7. O. Kabil and R. Banerjee, *J. Biol. Chem.*, 2010, **285**(29), 21903-21907.
8. B. D. Paul and S. H. Snyder, *Nat. Rev. Mol. Cell Biol.*, 2012, **13**(8), 499-507.
9. R. Wang, *Physiol. Rev.*, 2012, **92**(2), 791-896.
10. H. Kimura, *Antioxid. Redox Signal.*, 2015, **22**(5), 347-349.
11. H. Kimura, *Molecules*, 2014, **19**(10), 16146-16157.
12. T. Ida, T. Sawa, H. Ihara, Y. Tsuchiya, Y. Watanabe, Y. Kumagai, M. Suematsu, H. Motohashi, S. Fujii, T. Matsunaga, M. Yamamoto, K. Ono, N. O. Devarie-Baez, M. Xian, J. M. Fukuto and T. Akaike, *Proc. Natl. Acad. Sci. U. S. A.*, 2014, **111**(21), 7606-7611.
13. J. I. Toohey, *Anal. Biochem.*, 2011, **413**(1), 1-7.
14. K. Ono, T. Akaike, T. Sawa, Y. Kumagai, D. A. Wink, D. J. Tantillo, A. J. Hobbs, P. Nagy, M. Xian, J. Lin and J. M. Fukuto, *Free Radical Biol. Med.*, 2014, **77**, 82-94.
15. P. Wardman, L. P. Candeias, S. A. Everett and M. Tracy, *Int. J. Radiat. Biol.*, 1994, **65**(1), 35-41.
16. S. A. Everett, C. Schoeneich, J. H. Stewart and K. D. Asmus, *J. Phys. Chem.*, 1992, **96**(1), 306-314.
17. S. W. Benson, *Chem. Rev.*, 1978, **78**(1), 23-35.
18. S. A. Everett and P. Wardman, *Methods Enzymol.*, 1995, **251**, 55-69.
19. S. A. Everett, L. K. Folkes, P. Wardman and K. D. Asmus, *Free Radical Res.*, 1994, **20**(6), 387-400.
20. J. I. Toohey, *Biochem. J.*, 1989, **264**(3), 625-632.
21. H. Böhme and G. Zinner, *Justus Liebigs Annalen der Chemie*, 1954, **585**(1), 142-149.
22. J. Tsurugi, Y. Abe and S. Kawamura, *Bull. Chem. Soc. Jpn.*, 1970, **43**(6), 1890-1892.
23. T. Nakabayashi, J. Tsurugi and T. Yabuta, *J. Org. Chem.*, 1964, **29**(5), 1236-1238.
24. J. Tsurugi, T. Nakabayashi and T. Ishihara, *J. Org. Chem.*, 1965, **30**(8), 2707-2710.
25. S. Kawamura, T. Kitao, T. Nakabayashi, T. Horii and J. Tsurugi, *J. Org. Chem.*, 1968, **33**(3), 1179-1181.
26. J. Tsurugi, Y. Abe, T. Nakabayashi, S. Kawamura, T. Kitao and M. Niwa, *J. Org. Chem.*, 1970, **35**(10), 3263-3266.
27. T. Chatterji, K. Keerthi and K. S. Gates, *Bioorg. Med. Chem. Lett.*, 2005, **15**(17), 3921-3924.
28. S. Kawamura, T. Horii and J. Tsurugi, *Bull. Chem. Soc. Jpn.*, 1971, **44**(10), 2878-2879.
29. R. Franzi, M. Geoffroy and G. Bernardinelli, *Mol. Phys.*, 1984, **52**(4), 947-954.
30. T. S. Bailey, L. N. Zakharov and M. D. Pluth, *J. Am. Chem. Soc.*, 2014, **136**(30), 10573-10576.
31. T. Nakabayashi, T. Horii, S. Kawamura and J. Tsurugi, *Annu. Rep. Radiat. Cent. Osaka Prefect.*, 1968, **9**, 64-67.
32. J. Tsurugi, T. Horii, T. Nakabayashi and S. Kawamura, *J. Org. Chem.*, 1968, **33**(11), 4133-4135.
33. T. Nakabayashi, J. Tsurugi, S. Kawamura, T. Kitao, M. Ui and M. Nose, *J. Org. Chem.*, 1966, **31**(12), 4174-4178.
34. T. Nakabayashi, S. Kawamura, T. Kitao and J. Tsurugi, *J. Org. Chem.*, 1966, **31**(3), 861-864.
35. T. Nakabayashi and J. Tsurugi, *J. Org. Chem.*, 1963, **28**(3), 813-816.

36. G. Derbesy, D. N. Harpp, B. Rather and G. Carroll, *Sulfur Lett.*, 1992, **14**, 199-204.
37. D. F. Aycock and G. R. Jurch, *J. Org. Chem.*, 1979, **44**(4), 569-572.
38. N. E. Heimer and L. Field, *J. Org. Chem.*, 1984, **49**(8), 1446-1449.
39. N. E. Heimer, L. Field and R. A. Neal, *J. Org. Chem.*, 1981, **46**(7), 1374-1377.
40. N. E. Heimer, L. Field and J. A. Waites, *J. Org. Chem.*, 1985, **50**(21), 4164-4166.
41. J. Tsurugi, S. Kawamura and T. Horii, *J. Org. Chem.*, 1971, **36**(24), 3677-3680.
42. J. Tsurugi and T. Nakabayashi, *J. Org. Chem.*, 1959, **24**(6), 807-810.
43. I. Artaud and E. Galardon, *ChemBioChem*, 2014, **15**(16), 2361-2364.
44. Y. Zhao, S. Bhushan, C. Yang, H. Otsuka, J. D. Stein, A. Pacheco, B. Peng, N. O. Devarie-Baez, H. C. Aguilar, D. J. Lefer and M. Xian, *ACS Chem. Biol.*, 2013, **8**(6), 1283-1290.
45. T. Roger, F. Raynaud, F. Bouillaud, C. Ransy, S. Simonet, C. Crespo, M.-P. Bourguignon, N. Villeneuve, J.-P. Vilaine, I. Artaud and E. Galardon, *ChemBioChem*, 2013, **14**(17), 2268-2271.
46. U. Hannestad, S. Margheri and B. Sörbo, *Anal. Biochem.*, 1989, **178**(2), 394-398.
47. C. Liu, F. Zhang, G. Munske, H. Zhang and M. Xian, *Free Radical Biol. Med.*, 2014, **76**, 200-207.
48. S. Kawamura, Y. Otsuji, T. Nakabayashi, T. Kitao and J. Tsurugi, *J. Org. Chem.*, 1965, **30**(8), 2711-2714.
49. S. Kawamura, T. Nakabayashi, T. Kitao and J. Tsurugi, *J. Org. Chem.*, 1966, **31**(6), 1985-1987.
50. T. Nakabayashi, S. Kawamura, T. Horii and M. Hamada, *Chem. Lett.*, 1976, **5**(8), 869-874.
51. S. Kawamura, Y. Abe and J. Tsurugi, *J. Org. Chem.*, 1969, **34**(11), 3633-3635.
52. T. Chatterji and K. S. Gates, *Bioorg. Med. Chem. Lett.*, 2003, **13**(7), 1349-1352.
53. K. Mitra, W. Kim, J. S. Daniels and K. S. Gates, *J. Am. Chem. Soc.*, 1997, **119**(48), 11691-11692.
54. K. S. Gates, *Chem. Res. Toxicol.*, 2000, **13**(10), 953-956.
55. W. A. Prütz, *Int. J. Radiat. Biol.*, 1992, **61**(5), 593-602.
56. I. Kende, T. L. Pickering and A. V. Tobolsky, *J. Am. Chem. Soc.*, 1965, **87**(24), 5582-5586.
57. M. D. Sevilla, D. Becker and M. Yan, *Int. J. Radiat. Biol.*, 1990, **57**(1), 65-81.
58. N. E. Franconeri, S. J. Carrington and J. M. Fukuto, *Arch. Biochem. Biophys.*, 2011, **516**(2), 146-153.
59. D. Cavallini, C. De Marco and B. Mondovi, *Arch. Biochem. Biophys.*, 1960, **87**(2), 281-288.
60. M. Villarejo and J. Westley, *J. Biol. Chem.*, 1963, **238**(12), 4016-4020.
61. M. Flavin, *J. Biol. Chem.*, 1962, **237**, 768-777.
62. D. Cavallini, C. De Marco, B. Mondovi and B. G. Mori, *Enzymologia*, 1960, **22**, 161-173.
63. G. S. Rao and G. Gorin, *J. Org. Chem.*, 1959, **24**(6), 749-753.
64. D. J. Smith and V. Venkatraghavan, *Synth. Commun.*, 1985, **15**(10), 945-950.
65. E. G. Mueller, *Nat. Chem. Biol.*, 2006, **2**(4), 185-194.
66. H. Mihara and N. Esaki, *Appl. Microbiol. Biotechnol.*, 2002, **60**, 12-23.
67. L. Zheng, R. H. White, V. L. Cash, R. F. Jack and D. R. Dean, *Proc. Natl. Acad. Sci. U. S. A.*, 1993, **90**(7), 2754-2758.
68. L. Zheng, V. L. Cash, D. H. Flint and D. R. Dean, *J. Biol. Chem.*, 1998, **273**(21), 13264-13272.
69. Y. Takahashi and M. Nakamura, *J. Biochem.*, 1999, **126**(5), 917-926.
70. P. M. Palenchar, C. J. Buck, H. Cheng, T. J. Larson and E. G. Mueller, *J. Biol. Chem.*, 2000, **275**(12), 8283-8286.
71. R. Kambampati and C. T. Lauhon, *J. Biol. Chem.*, 2000, **275**(15), 10727-10730.
72. K. Nilsson, H. K. Lundgren, T. G. Hagervall and G. R. Bjork, *J. Bacteriol.*, 2002, **184**(24), 6830-6835.
73. J. H. Ploegman, G. Drent, K. H. Kalk, W. G. Hol, R. L. Heinrikson, P. Keim, L. Weng and J. Russell, *Nature*, 1978, **273**(5658), 124-129.
74. J. Westley, *Adv. Enzymol. Relat. Areas Mol. Biol.*, 1973, **39**(327-368).
75. C. J. Knowles, *Bacteriol. Rev.*, 1976, **40**(3), 652-680.
76. D. L. Nandi, P. M. Horowitz and J. Westley, *Int. J. Biochem. Cell. B.*, 2000, **32**(4), 465-473.
77. M. Iciek and L. Wlodek, *Pol J Pharmacol*, 2001, **53**(3), 215-225.
78. A. K. Mustafa, M. M. Gadalla, N. Sen, S. Kim, W. Mu, S. K. Gazi, R. K. Barrow, G. Yang, R. Wang and S. H. Snyder, *Sci. Signal.*, 2009, **2**(96), ra72.
79. N. Sen, B. D. Paul, M. M. Gadalla, A. K. Mustafa, T. Sen, R. Xu, S. Kim and S. H. Snyder, *Mol. Cell*, 2012, **45**(1), 13-24.
80. G. Yang, K. Zhao, Y. Ju, S. Mani, Q. Cao, S. Puukila, N. Khaper, L. Wu and R. Wang, *Antioxid. Redox Signal.*, 2013, **18**(15), 1906-1919.
81. N. Krishnan, C. Fu, D. J. Pappin and N. K. Tonks, *Sci. Signal.*, 2011, **4**(203), ra86.
82. K. Zhao, S. Li, L. Wu, C. Lai and G. Yang, *J. Biol. Chem.*, 2014, **289**(30), 20824-20835.
83. D. Zhang, I. Macinkovic, N. O. Devarie-Baez, J. Pan, C. M. Park, K. S. Carroll, M. R. Filipovic and M. Xian, *Angew. Chem. Int. Ed.*, 2014, **53**(2), 575-581.
84. A. K. Mustafa, G. Sikka, S. K. Gazi, J. Stepan, S. M. Jung, A. K. Bhunia, V. M. Barodka, F. K. Gazi, R. K. Barrow, R. Wang, L. M. Amzel, D. E. Berkowitz and S. H. Snyder, *Circ. Res.*, 2011, **109**(11), 1259-1268.
85. J. Du, Y. Huang, H. Yan, Q. Zhang, M. Zhao, M. Zhu, J. Liu, S. X. Chen, D. Bu, C. Tang and H. Jin, *J. Biol. Chem.*, 2014, **289**(14), 9741-9753.
86. C. Guo, F. Liang, W. Shah Masood and X. Yan, *Eur. J. Pharmacol.*, 2014, **725**, 70-78.
87. M. S. Vandiver, B. D. Paul, R. Xu, S. Karuppagounder, F. Rao, A. M. Snowman, H. S. Ko, Y. I. Lee, V. L. Dawson, T. M. Dawson, N. Sen and S. H. Snyder, *Nat. Commun.*, 2013, **4**, 1626.
88. K. Zhao, Y. Ju, S. Li, Z. Altaany, R. Wang and G. Yang, *EMBO Rep.*, 2014, **15**(7), 792-800.
89. Z. Altaany, Y. Ju, G. Yang and R. Wang, *Sci. Signal.*, 2014, **7**(342), ra87.
90. Y. Liu, R. Yang, X. Liu, Y. Zhou, C. Qu, T. Kikuri, S. Wang, E. Zandi, J. Du, I. S. Ambudkar and S. Shi, *Cell Stem Cell*, 2014, **15**(1), 66-78.
91. Z.-Z. Xie, M.-M. Shi, L. Xie, Z.-Y. Wu, G. Li, F. Hua and J.-S. Bian, *Antioxid. Redox Signal.*, 2014, **21**(18), 2531-2542.
92. M. D. de Beus, J. Chung and W. Colón, *Protein Sci.*, 2004, **13**(5), 1347-1355.

93. E. Grambow, F. Mueller-Graf, E. Delyagina, M. Frank, A. Kuhla and B. Vollmar, *Platelets*, 2014, **25**(3), 166-174.
94. V. Massey and D. Edmondson, *J. Biol. Chem.*, 1970, **245**(24), 6595-6598.
95. D. Edmondson, V. Massey, G. Palmer, L. M. Beacham, 3rd and G. B. Elion, *J. Biol. Chem.*, 1972, **247**(5), 1597-1604.
96. U. Branzoli and V. Massey, *J. Biol. Chem.*, 1974, **249**(14), 4346-4349.
97. B. Sörbo, *Biochim. Biophys. Acta*, 1957, **24**, 324-329.
98. T. Sawahata and R. A. Neal, *Anal. Biochem.*, 1982, **126**(2), 360-364.
99. L. Zheng, R. H. White, V. L. Cash and D. R. Dean, *Biochemistry*, 1994, **33**(15), 4714-4720.
100. J. Pan and K. S. Carroll, *ACS Chem. Biol.*, 2013, **8**(6), 1110-1116.