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Protein S-nitrosylation in health and disease: a current perspective

Matthew W. Foster¹, Douglas T. Hess¹, and Jonathan S. Stamler^{1,2}

¹Department of Medicine, Duke University Medical Center, Durham NC 27710

²Department of Biochemistry, Duke University Medical Center, Durham NC 27710

Summary

Protein *S*-nitrosylation conveys a large part of the ubiquitous influence of nitric oxide on cellular signal transduction, and accumulating evidence indicates important roles for *S*-nitrosylation both in normal physiology and in a broad spectrum of human diseases. Here we review recent findings that implicate *S*-nitrosylation in cardiovascular, pulmonary, musculoskeletal and neurological (dys)function, as well as in cancer. The emerging picture shows that, in many cases, pathophysiology correlates with hypo- or hyper-*S*-nitrosylation of specific protein targets, rather than a general cellular insult due to loss of or enhanced nitric oxide synthase activity. In addition, it is increasingly evident that dysregulated *S*-nitrosylation can result not only from alterations in the expression, compartmentalization and/or activity of nitric oxide synthases but can also reflect a contribution from denitrosylases, including prominently the *S*-nitrosoglutathione (GSNO)-metabolizing enzyme, GSNO reductase. Finally, because exogenous mediators of protein *S*-nitrosylation or denitrosylation can substantially affect the development or progression of disease, potential therapeutic agents that modulate *S*-nitrosylation could well have broad clinical utility.

Keywords

nitric oxide; S-nitrosylation; S-nitrosoglutathione; cysteine; GSNOR; heart failure; asthma; nitrosative stress

Introduction

In mammalian cells, the L-Arg-dependent nitric oxide (NO) synthases – neuronal NOS (nNOS, NOS1), inducible NOS (iNOS, NOS2) and endothelial NOS (eNOS, NOS3) – are the major sources of endogenous NO, and the stimulus-coupled activation or induction of NO synthases has been shown to mediate or modulate a broad range of cellular signaling pathways. The physiological influence of NO is exerted predominantly through the posttranslational modification and functional regulation of proteins. It was first established that nitrosylation of heme iron within soluble guanylate cyclase activates the enzyme to generate cyclic GMP, and thereby subserves NO-based vasoactivity. However, hemes do not generally elicit cellular signaling involving posttranslational modification of proteins, and

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To whom correspondence should be addressed: Jonathan S. Stamler, M.D., Box 2612, Department of Medicine, Duke University Medical Center, Durham, NC 27710, Phone: 919-684-6933, FAX: 919-684-6998, staml001@mc.duke.edu.

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thus an explanation for most NO-based bioactivity was not apparent. Subsequently, a large body of experimental evidence has demonstrated that *S*-nitrosylation of Cys residues within a broad functional spectrum of proteins conveys a large part of the ubiquitous influence of NO on cellular function [1].

Expression of iNOS is induced in many mammalian cell types by a variety of stressors or injury, and the cytotoxic action of NO generated in particular by phagocytic cells raised the possibility that NO generated at relatively high and sustained levels by iNOS could compromise cellular function through a generalized nitrosative stress. However, the emerging recognition that NO is involved in a multiplicity of cellular signal transduction pathways, through protein S-nitrosylation, pointed to the possibility that dysregulated Snitrosylation could contribute to pathophysiologies characteristic of a wide range of disease states [2]. The relatively recent development of improved methods for the analysis of protein S-nitrosylation [3,4] has allowed the identification of numerous S-nitrosylated proteins (SNO-proteins) whose levels of S-nitrosylation can be altered in disease, and the emerging picture shows that hypo- or hyper-S-nitrosylation of these specific protein targets (which result in alterations in protein function) are directly implicated in the etiology and symptomatology of an increasing number of human diseases, including prominently disorders of the cardiovascular, musculoskeletal and nervous systems (Table 1). In a number of cases, the specific Cys residues that are the loci of (patho)physiological regulation by Snitrosylation have been identified.

The molecular mechanisms underlying the (dys)regulation of S-nitrosylation, and possible approaches to therapeutically altering SNO-protein levels, have also begun to receive increasing attention. Although NOS expression and activity are obvious governors of Snitrosylation, the co-localization of NOS enzymes with target proteins, including direct interactions, appears in many cases to be an important determinant of S-nitrosylation under physiological conditions, and accordingly, aberrant NOS localization appears to be involved in at least several diseases (Fig. 1). Such deficits can have a genetic basis. For example, in a variant of long OT syndrome, a mutation in an nNOS-scaffolding protein results in disinhibition of nNOS and aberrant S-nitrosylation of a cardiac ion channel [5]. In addition to localization, it has been established that the transfer of NO groups between proteins and glutathione governs a cellular equilibrium between small-molecular-weight and protein Snitrosothiols (SNOs) (Fig. 2). In mouse models, genetic ablation of S-nitrosoglutathione reductase (GSNOR), the enzyme that is principally responsible for GSNO metabolism, results in enhanced levels of SNO-proteins and significantly attenuates experimental asthma and heart-failure [6,7], while increasing the severity of endotoxic shock [8]. Finally, the therapeutic potential of agents that affect S-nitrosylation is being explored with promising results (Table 2). These agents have the potential to restore deficient SNO-proteins to physiological levels or to otherwise influence cellular signaling pathways that are mediated or modulated by S-nitrosylation. For example, SNO-repleting agents have been shown to be highly efficacious in the setting of inflammation, including experimental models of lung injury, stroke and multiple sclerosis in which S-nitrosylation appears to play a major role in expression of the innate immune response [9–12]. This review focuses on the numerous proteins and signaling pathways that are regulated by S-nitrosylation, in the context of diseases in which aberrant S-nitrosylation has recently been implicated.

Modulation of SNO production: NO synthases and organic nitrates

In tissues and extracellular fluids, SNO levels are likely to reflect NOS activity and, accordingly, can be modulated through altered expression or activity of enzymes that control the availability of endogenous NOS substrates (e.g. l-Arg) or endogenous NOS inhibitors (e.g. asymmetric dimethylarginine, ADMA). Additionally, a number of enzymes in the l-

Arg/NO pathway are also targets of regulatory *S*-nitrosylation [1], including arginase, which catabolizes l-Arg. Arginase 1 (Arg1; cytosolic arginase) is activated through enhanced multimerization resulting from *S*-nitrosylation at cysteine 303 (C303) [13]. Arg1 *S*-nitrosylation and activation coincide with increased iNOS expression in the aortic endothelium of aging rats [13] and may contribute to age-related endothelial dysfunction and vascular stiffness [14]. Levels of the bronchodilator GSNO are low in asthmatic airways [2], and a causal link between SNO depletion and arginase levels has been suggested [15]: Arg1 and Arg2 are upregulated in human asthma and in experimental asthma models [16,17], and single nucleotide polymorphisms (SNPs) in genes encoding Arg1 and Arg2 are associated with increased risk of childhood asthma [18]. Finally, pharmacological inhibition of arginase leads to increase inflammation [15], arginase inhibition correlates with protection from allergen-induced airway hyperresponsivity (AHR) in several asthma models [16,19]. These data suggest that altering NO homeostasis and SNO levels might ameliorate the asthmatic phenotype.

NO or low-mass SNOs can also be produced pharmacologically from organic nitrovasodilators, chief among which is nitroglycerin (glyceryl trinitrate; GTN). The vasodilatory properties of GTN arise from its metabolism by mitochondrial aldehyde dehydrogenase (mtALDH) and the concerted production of NO bioactivity by mitochondria [20]. However, prolonged use of GTN results in tachyphylaxis to the drug itself (mechanism-based tolerance) as well as diminished responsiveness to other nitrovasodilators (cross-tolerance). S-nitrosylation is increasingly implicated in these effects [21], with soluble guanylate cyclase (sGC) serving as a major locus of effect (Box 1). Both exposure to low-mass SNOs and eNOS activation result in S-nitrosylation of sGC, and the modification coincides with a reduction in NO-stimulated sGC activity [22]. Two cysteines residues in the sGC alpha/beta heterodimer are the targets of S-nitrosylation, C243 of the alpha-chain (α C243) and C122 of the beta-chain (β C122). Of these, β C122 is in proximity to the hemebinding site, although mutation of either residue partially prevents SNO-induced sGC densitization [22]. These effects of NO/SNO are essentially recapitulated in cellular and animal models of nitrate tolerance, and, in addition, partial sGC denitrosylation and restoration of sGC activity are observed after treatment with N-acetylcysteine (NAC) [23]. Thus, denitrosylation of SNO-sGC might underlie some of the clinical benefit of NAC therapy for nitrate tolerance.

Therapy with the NO donor isosorbide dinitrate (in combination with hydralazine) markedly reduces mortality in African-Americans with heart failure [24]. Morbidity and mortality in heart failure patients results from either arrhythmia or ventricular dysfunction that is due at least in part to downregulation of β -adrenergic receptors (β -AR) and dysregulation of calcium homeostasis. It is therefore of interest that NO bioactivity has been strongly associated with both arrhythmia and pump failure [25–27], and recent studies support the possibility that *S*-nitrosylation of cardiac proteins ameliorates heart failure by reversing calcium leakage [28] and normalizing β -AR expression [27].

GSNO reductase and its regulation by NO/SNO – implications for asthma

A major mechanism for protein *S*-nitrosylation *in vivo* is the reversible transnitrosylation of protein thiols by GSNO, the predominant low-mass SNO (Figure 2) [2]. GSNO is the preferred physiological substrate for glutathione-dependent formaldehyde dehydrogenase (class III alcohol dehydrogenase, ADH III, which in methylotropic bacteria may also metabolize formaldehyde), whereasalcohols are apparently not physiological substrates of ADH III. Thus the enzyme has been renamed GSNO reductase (GSNOR). GSNOR metabolizes GSNO to glutathione disulfide and a glutathione N-hydroxysulfenamide

intermediate, which is ultimately reduced to ammonia (Figure 2). Analyses in GSNORknockout mice have helped elucidate the physiological roles of both GSNO and protein *S*nitrosylation. GSNOR-knockout mice have markedly increased levels of SNO-proteins, which demonstrates a role for GSNO/GSNOR in SNO-protein homeostasis. These mice exhibit increased mortality in endotoxic shock, and SNO-protein levels and mortality are attenuated by iNOS inhibitors. In contrast GSNOR-knockout mice are protected from experimental myocardial infarction, due at least in part to *S*-nitrosylation-mediated stabilization of hypoxia-inducible factor HIF-1 α and increased angiogenesis ([7]; see below).

There is also accumulating evidence from both human studies and experimental models that GSNOR plays a significant role in the etiology and symptomatology of asthma. It is well established that airway SNOs are depleted in asthma [2], and it has recently been reported that this decrease is associated with increased GSNOR levels and activity in airway lining fluid (obtained by bronchoalveolar lavage) of adult asthmatics [29]. In addition, SNPs in GSNOR are associated with a decreased (1 or 2 copies of minor allele of SNP rs1154404) or increased (homozygotes of minor allele of SNP rs28730619) risk of childhood asthma; haplotypes with the major allele of rs1154404 and minor allele of rs28730619 have an increased risk for childhood asthma [30]. Further, one SNP (rs1154400) is also associated with decreased responsiveness to {Greek beta}-agonist therapy in African-American children with asthma [153]. The effects of these SNPs on GSNOR protein expression, localization and/or activity have not yet been characterized. However, GSNOR activity in the lungs of 36 human asthmatics correlated directly with airway hyperreactivity [29]. In an experimental asthma model, GSNOR-knockout mice exhibit lower basal airway tone and less AHR to methacholine challenge following sensitization with ovalbumin (OVA) [6]. This protection is associated with an increase in iNOS-derived SNO-proteins in the lungs of GSNOR-knockout mice following OVA challenge [6], and is reversed by NO synthase inhibitors. Lungs from GSNOR-knockout mice also have increased levels of β -adrenergic receptors (β -ARs) and their airways are resistant to β -agonist-induced desensitization in *vitro*, suggesting that GSNO promotes airway relaxation through the regulation of β -AR signaling (Box 2) [6].

NO also appears to modulate GSNOR expression. Inhibition of constitutive NOS activity increases GSNOR mRNA in mouse lungs [31], whereas overexpression of vascular endothelial growth factor (VEGF) results in large, NO-dependent increases in GSNOR transcript levels that appear to be mediated through induction of eNOS and iNOS [31]. The increase in GSNOR expression (although not verified at the protein level) suggests a mechanism for VEGF-induced AHR. GSNOR transcription is stimulated by specificity factor 1 (Sp1) and repressed by Sp3/Sp4 [32]. By analogy to the GSNO-dependent regulation of cystic fibrosis trans-membrane conductance regulator (CFTR) transcription by Sp3/Sp1 [33], low levels of NO/SNO (arising from constitutive NOS activity) might repress GSNOR transcription through augmented Sp3 binding, whereas high levels of NO/SNO (e.g. induced by VEGF) might activate GSNOR transcription by simultaneously inhibiting Sp3 binding and potentiating Sp1 binding [33]. Collectively, these studies demonstrate a role for GSNO in SNO-protein homeostasis, suggest that (dys)regulation of GSNOR expression may be important in the etiology of airway disease and strongly support a role for protein *S*-nitrosylation in NOS-dependent (patho)physiology.

Ras S-nitrosylation – adaptive immunity and tumor maintenance

Many GTPases within the Ras superfamily contain redox-sensitive Cys residues that are susceptible to *S*-nitrosylation [34]. In the cases of H-, K- and N-Ras, NO promotes the conversion by guanine nucleotide exchange of inactive, GDP-bound Ras to its active, GTP-

bound form. Ras activation is coincident with *S*-nitrosylation of C118, which resides within the nucleotide-binding domain; mutation of C118 abolishes NO-induced Ras activation [34]. Although *S*-nitrosylation of Ras is closely coupled to NOS activity, it has been proposed that higher oxides of NO (e.g. NO_2^{\bullet}) affect guanine nucleotide exchange by C118 thiyl radicalcatalyzed oxidation and, finally, release of the guanine cofactor [35,36]. In this context, *S*nitrosylation might prevent or promote further nucleotide turnover (depending on whether Cys radical formation is inhibited or enhanced) and/or serve as a marker of NO-based activation.

In T-lymphocytes, eNOS undergoes rapid and sustained activation by both phosphoinositide 3-kinase/protein kinase B (PI3K–Akt)- and Ca²⁺-dependent pathways upon binding of the T-cell receptor (TCR) to antigen-presenting cells (APCs) (Figure 4) [37]. In T-cells, eNOS is localized to the Golgi complex, and its activation (at the immunological synapse between the T-cell and APC) drives N-Ras-dependent ERK activation [37]. Importantly, only Golgi-localized N-Ras, and not plasma membrane-associated K-Ras, is activated and *S*-nitrosylated by eNOS, demonstrating a requirement for compartmentalization of eNOS with its target [38,39]. Mutation of C118 abrogates N-Ras *S*-nitrosylation and activation as well as TCR-dependent apoptosis [38], suggesting a role for *S*-nitrosylation in the negative selection of highly reactive T-cells in the thymus.

Ras *S*-nitrosylation is also implicated in the initiation of tumorigenesis and maintenance of established tumors. In tumor cells, oncogenic (mutant) K-Ras activates eNOS by PI3K–Akt-dependent phosphorylation of eNOS at S1117, which results in *S*-nitrosylation of wild-type N-and H-Ras [40]. S1117-phosphorylated eNOS is elevated in tumors isolated from patients with pancreatic cancer, and overexpression of either S1117A eNOS, or C118S N- or H-Ras inhibits pancreatic tumor growth [40]. Although C118 of Ras appears to be a principal locus of action of tumorigenic nitric oxide, NO might play a multifaceted role in cancer [41], and *S*-nitrosylation of numerous proteins could influence the cancer phenotype (Box 3; [42]).

Aberrant hemoglobin S-nitrosylation and the human respiratory cycle – sickle cell anemia, banked blood and pulmonary arterial hypertension

Accumulating evidence supports a role for red blood cell (RBC) SNOs, which originate from *S*-nitroso-(β Cys93)hemoglobin (SNO-Hb), in mediating oxygen tension (PO₂)dependent vasodilatory activity [43] within the human respiratory cycle. In the resultant 3gas (NO, O₂, CO₂) model of the respiratory cycle, RBCs liberate NO-based bioactivity to enable efficient O₂ delivery (which is primarily a function of blood flow). This activity of RBCs appears to require the transfer of SNO from Hb to RBC low-mass and/or membrane protein thiols (e.g. erythrocyte band 3) and finally to circulating or vascular thiols [44–46]. *Ex vivo* arterial vasodilation by human RBCs, triggered by hypoxia (e.g. 1% O₂), does not require NOS activity or intact endothelium, but depends critically upon RBC-SNO levels [47]. Moreover, this vasodilatory activity is actuated in a graded manner across a PO₂ range of 60-7 mm Hg, representative of physiological O₂ gradients in the microcirculation [48].

The loss of SNO-Hb, or defective transfer of SNO from Hb to acceptor thiols, might underlie numerous pathological states [2], and accumulating data suggest that the restoration of SNO levels ('SNO repletion') can ameliorate these conditions. Sickle cell Hb (HbS; Glu6Val Hb) is impaired in both its ability to form SNO-HbS and to transfer the SNO to the RBC membrane, resulting in impaired hypoxic vasodilation, and the magnitudes of these deficits correlate with clinical severity in patients with sickle cell anemia [49]. Hypoxic vasoactivity (and membrane SNO content) of sickle RBCs can be at least partially restored by treatment with NO [49]. In addition, storage of blood (under conditions employed in blood banking) leads to a time-dependent loss of SNO-Hb and impairment of hypoxic

vasodilation by RBCs, both of which are restored upon SNO repletion [50,51]. These data provide a rationale for the well-established inability of stored blood to fully correct anemiaassociated deficits in tissue oxygen delivery, and they suggest that SNO repletion might be effective in reducing morbidity and mortality associated with transfusion. Finally, RBCs exposed to prolonged hypoxia are defective in SNO-Hb formation, and in the lung, exhibit enhanced hypoxic pulmonary vasoconstrictive activity and are deficient in increasing blood oxygenation; these deficiencies are reversed by SNO repletion [52]. Consistent with these findings, hypoxemic patients with pulmonary aterial hypertension (PAH) have markedly reduced SNO-Hb levels, and hypoxic vasodilation by their RBCs is impaired both *ex vivo* and *in vivo* [52]. Treatment of PAH patients with the *S*-nitrosylating gas, ethyl nitrite (ENO), normalizes SNO-Hb levels and RBC vasodilation, and improves oxygenation and lowers pulmonary arterial pressures [52]. In addition, ENO has shown clinical benefits in the syndrome of persistent hypertension of the newborn [53].

Control of hypoxic signaling – tumor radiotherapy, protection from myocardial ischemia and development of pulmonary arterial hypertension

HIF-1, a master transcriptional regulator, is activated at low oxygen tension through the stabilization of its alpha subunit (HIF-1 α). Under normoxia, hydroxylation of HIF-1 α by an O₂-dependent prolyl hydroxylase promotes its binding with and ubiquitylation by the E3 ligase complex containing pVHL and subsequent proteasomal degradation. Accumulating evidence suggests that, under normoxia, NO/SNO stablizes HIF-1 α (thus mimicking hypoxia) [1].

S-nitrosylation-dependent HIF-1 α stabilization is implicated in the resistance of solid tumors to ionizing radiation [54]. Irradiation of tumors results in increased HIF-1 activity and VEGF levels, coincident with macrophage- and iNOS-dependent stabilization of HIF-1 α through its oxygen-dependent degradation (ODD) domain. S-nitrosylation appears to prevent HIF-1 α degradation by inhibiting the binding of the Von Hippel-Lindau disease tumor suppressor pVHL to the hydroxylated ODD, and the effects of exogenous SNO or iNOS (on both the ODD–pVHL interaction and HIF-1 α stability) require S-nitrosylation of C533 in the ODD domain [54]. In addition, NOS inhibitors act synergistically with radiotherapy to reduce the rate of tumor growth in mouse models of breast cancer and melanoma, and tumor growth is substantially slower in radiation-treated iNOS-knockout versus wild-type mice [54]. These data are consistent with a role for tumor vascularization, mediated by NO, in tumor survival [41]. Although NOS inhibitors may have therapeutic potential in cancers in which paracrine (macrophage-derived) or autocrine (eNOS-derived; see discussion of Ras S-nitrosylation) NO plays a role, it should be noted that NO may also sensitize tumors to chemo-, radio- or immuno-therapies [55].

HIF-1 α stabilization, resulting from increased endogenous GSNO, also appears to underlie amelioration of myocardial infarction. GSNOR-knockout mice have reduced myocardial infarct size and preserved ventricular systolic and diastolic function and maintain tissue oxygenation following left coronary artery ligation [7]. These effects are associated with increases in myocardial capillary density, increased levels and activity of HIF-1 α and elevated HIF-1 α *S*-nitrosylation in GSNOR-knockout mice [7]. These data identify a role for endogenous SNO in angiogenesis and myocardial protection and, more generally, suggest novel therapeutic approaches to modulate angiogenesis and preserve cardiac function. It should be noted that these data do not rule out the possibility that additional, pro-angiogenic proteins and/or pathways might be (dys)regulated by *S*-nitrosylation. For example, *S*nitrosylation of dynamin stimulates its GTPase activity [56,57] and potentiates endothelial cell survival signaling [57], and *S*-nitrosylation of JNK3 phosphatase MKP7, which promotes endothelial proliferation [58], might also promote (S)NO-dependent angiogenesis.

HIF-1 α stabilization by S-nitrosothiols is also implicated in the development of PAH. Mice treated with NAC or S-nitroso-N-acetylcysteine (SNOAC) for several weeks develop PAH, mimicking the effects of chronic hypoxia [46]. eNOS-knockout mice are protected from the increases in relative right ventricular weight and systolic pressure induced by NAC (but not SNOAC), implicating SNOAC in the effects of NAC [46]. The bioactivity of NAC appears to originate from SNO-Hb, as SNOAC is formed from NAC in blood coincident with declines in SNO-Hb, and SNO-Hb levels are lower and SNOAC formation from blood is impaired in eNOS-knockout mice [46]. Declines in SNO-Hb and increases in SNOAC are facilitated by increasing hypoxia, a consequence of SNO-Hb instability upon deoxygenation. NAC/SNOAC treatment stabilizes whole-lung HIF1 DNA-binding activity; C162 of pVHL, a residue known to be essential for binding of the adaptor protein elongin C, is implicated in the effects of SNOAC [46]. These data suggest that PAH may be caused either by deficiency in SNO-Hb and/or by excess in low-mass S-nitrosothiols: deficiency of SNO-Hb might impair hypoxia-mediated release of NO bioactivity to counter pulmonary vasoconstriction, whereas excess SNO simulates sustained hypoxia, which ultimately increases pulmonary pressure (secondary to vascular remodeling). PAH might also be a potential side effect of chronic NAC therapy.

(Dys)regulated S-nitrosylation in the heart through differential NOS expression and localization – contractility, ischemia and long QT

The importance of NOS subcellular compartmentalization is well-exemplified in the heart, where eNOS and nNOS are differentially localized and exhibit opposite effects on myocardial contractility [25,59,60]. eNOS, associated primarily with sarcolemmal caveolae, attenuates β -AR-dependent myocardial contractility [61]; inhibition of the L-type Ca²⁺ channel (LTCC), at least in part by S-nitrosylation [26], appears to underlie this effect. By contrast, nNOS is primarily localized to the sarcoplasmic reticulum (SR) and activates the cardiac ryanodine receptor/Ca²⁺ channel (RyR2) by S-nitrosylation [62], resulting in release of Ca^{2+} to the cytosol and enhanced catecholamine-stimulated contractility. Accordingly, nNOS- but not eNOS-knockout mice exhibit hypo-S-nitrosylation of RyR2 and diastolic Ca^{2+} leakage with arrhythmia characteristic of sudden cardiac death (SCD) [28]. Sex-related differences in NOS expression/localization also appear to modulate Ca²⁺ handling. Hearts [63] and myocytes [26,63] from female mice show less isoproterenol (ISO)-induced SR Ca²⁺ loading compared with males, evidently owing to the translocation of nNOS to the sarcolemma and S-nitrosylation of the LTCC [26]. The redistribution of nNOS by ISO, as well as the upregulation of NOS isoforms (due at least in part, to 17β-estradiol-dependent gene expression [64,65]), appears to underlie the protection of female hearts from ischemia/ reperfusion (I/R) injury [26]. However, the redistribution of nNOS to the plasma membrane is also observed in the hearts of humans with idiopathic dilated cardiomyopathy [66,67] and of rodents with experimental myocardial infarction [66,68]. Thus, aberrant RyR2 and LTCC S-nitrosylation might also explain some of the characteristic dysfunction of the failing heart.

Dysregulated *S*-nitrosylation that results from altered NOS-target associations has also been linked to long QT syndrome (LQTS), an inherited disease that is characterized by a prolonged QT interval, which can result in fainting and sudden cardiac death from arrhythmia. A mutation in α1-syntrophin (A390V SNTA1), identified in a patient with LQTS, results in release of the plasma membrane calcium-ATPase 4b (a negative regulator of nNOS activity) from a complex containing SNTA1, nNOS and the cardiac sodium channel SCN5A [5]. Heterologous expression of A390V SNTA1 results in NOS-activity-dependent increases in sodium channel gating and in both peak and late sodium currents (*I*na). These effects coincide with hyper-*S*-nitrosylation of SCN5A, which is suggested as a mechanism for regulation of nNOS activity) and both altered QT interval [69,70] and

risk of SCD [70], although it is unknown whether aberrant NOS activity or ion channel *S*nitrosylation underlies these phenotypes. Endogenously derived NO has been shown to influence QT interval through *S*-nitrosylation of the slowly activating delayed-rectifier K⁺ channel (I_{Ks}) [71,72]. Gender differences in QT duration and susceptibility to ventricular arrhythmia have been linked to I_{Ks} currents [73]. Thus, sex-related differences in cardiac (dys)function may reflect differences in NOS activity/localization and *S*-nitrosylation of cardiac ion channels/transporters.

S-nitrosylation in disorders of skeletal muscle

The skeletal muscle ryanodine receptor (RyR) isoform (RyR1) is activated by Snitrosylation of C3635, which reduces the inhibitory effect of Ca^{2+} -CaM on the channel [74]. A number of mutations in RyR1 (e.g. Y522S) are associated with malignant hyperthermia and related diseases that are characterized by involuntary muscle contractures, tissue lysis (rhabdomyolysis) and sudden death in response to elevated environmental temperatures [75,76]. Heterozygous RyR1^{Y522S/wt} mutant mice, which have a malignant hyperthemia phenotype, have elevated ratios of oxidized-to-reduced glutathione, and RyR1 is hyper-S-nitrosylated and -S-glutathionylated in these mice, in association with temperature-dependent increases in cytosolic Ca²⁺ levels [75]. Prolonged Ca²⁺ leakage ultimately results in inadequate force generation and mitochondrial damage. NOS inhibitors and ascorbate treatment, which reverse NO- but not GSH-based modifications of the channels, ameliorate the classic temperature-sensitivity of the channel. Furthermore, NAC treatment in vitro and prolonged administration in vivo reverse NO-based modifications of the channel and ameliorate the phenotype [75]. These data suggest a cycle in which the mutation in RyR1 causes Ca²⁺ leakage, leading to increased NOS activity and further Snitrosylation and activation of the channel (i.e. a SNO-induced feed-forward mechanism). Thus, S-nitrosylation is primarily responsible for the increased temperature sensitivity of the RyR characteristic of malignant hyperthermia; oxidative stress appears to be a secondary consequence of increased NOS activity.

Hyper-S-nitrosylation of RyR1, and consequent channel leakage, is also observed in mice (and humans) subjected to repeated exercise [77] as well as in dystrophin-deficient (mdx)mice [78] that serve as a model of Duchenne muscular dystrophy (DMD), a genetic disorder characterized by progressive neuromuscular degeneration and eventual paralysis and death. Exercise-induced RyR1 S-nitrosylation appears to result from an increase in eNOS expression [77], whereas *mdx*-associated hypernitrosylation is mediated by an RyRaffiliated iNOS that is upregulated by inflammatory cytokines. In addition to altering RyR1-CaM interactions, hyper-S-nitrosylation of RyR1 correlates with the loss of binding between RyR1 and the peptidyl-prolyl cis-trans isomerase calstabin1 (FKBP12/FKBP1A; a protein that stabilizes the closed channel conformation), which contributes at least in part to channel leakage and decreased exercise capacity [77]. A novel small molecule (S107), which stabilizes the interaction of FKBP12 and RyR1, largely rescues these effects [77]. Human DMD results from mutations in the gene encoding dystrophin, which forms a complex with nNOS and syntrophin, and it has long been known that nNOS is mostly absent from skeletal muscle in DMD patients [79]. However, in mdx mice, iNOS is upregulated (and coimmunoprecipitates with RyR1), coincident with RyR1 hyper-S-nitrosylation [78]. These effects also lead to destabilization of the FKBP12-RyR1 complex, and treatment of mdx mice with S107 for two weeks restores RyR1-bound FKBP12, decreases Ca²⁺ leakage, ameliorates muscle damage and improves muscle function [78]. It will be of interest to determine whether S107 can alter other pathologies associated with RyR1 hyper-Snitrosylation, including malignant hyperthermia.

Although S107 apparently acts to reverse the deleterious effects of iNOS induction in the mdx mouse, overexpression of eNOS also appears to be ameliorative. Expression of constitutively active eNOS restores muscle differentiation and mimics the effects of histone deactylase (HDAC) inhibitors, which appear to induce myogenesis through transcriptional activation of follistatin [80]. HDAC2 is upregulated in *mdx* skeletal muscle, and Snitrosylation of HDAC2 by eNOS inhibits its deacetylase activity [81]. This contrasts with the report that S-nitrosylation of HDAC2 (at C262 and C274) by nNOS in neurons facilitates its release from chromatin without altering deacetylase activity [82]. Whether these mechanistic differences reflect NO source, cell type or additional parameters (and whether nNOS is a physiological regulator of HDAC2 in skeletal muscle) has not been determined, and these factors may impact the approach to therapy; in particular, whereas all isoforms of NOS might hyper-S-nitrosylate RyR, which is likely to have deleterious consequences, the underlying cause of hypernitrosylation, and the extent to which the nitrosative stress is localized to the sarcoplasmic reticulum, or is more generalized, will vary in different muscle disorders. It remains possible that appropriately targeted NO therapies, perhaps in conjunction with treatments that ameliorate RyR hypernitrosylation, will have beneficial effects in one or more disorders.

iNOS-induced S-nitrosylation in diabetes

Accumulating evidence links insulin resistance in type 2 diabetes to NO production and protein S-nitrosylation [83]. iNOS expression is increased in mouse models of diabetes, and iNOS-knockout mice fed a high-fat diet develop obesity but have improved glucose tolerance and normal insulin sensitivity (in vivo) and insulin-stimulated glucose uptake (ex vivo) compared with diet-induced obese wild-type mice [84]. iNOS-dependent Snitrosylation of protein kinase B (PKB/Akt), the insulin receptor β -subunit (IR β) and insulin receptor substrate 1 (IRS-1) are all markedly increased in diabetic mouse models and might underlie the effects of iNOS on insulin signaling [85,86]. NO/SNO inhibits insulinstimulated Akt activity, and mutation of C224 within Akt abolishes both Akt S-nitrosylation and its inhibition by low-mass SNOs [85]. iNOS activity or exogenous GSNO inhibit the tyrosine kinase activity of IR β [86]; a specific target of regulatory S-nitrosylation has not been identified, although the enzyme possesses at least one potential regulatory cysteine (C860) [87]. Finally, S-nitrosylation of IRS-1 might promote its ubiquitylation-dependent degradation [86,88]. S-nitrosylation of each of these substrates is also increased in skeletal muscle of wild-type mice treated with lipopolysaccharide (LPS) to stimulate iNOS induction but not in iNOS-knockout mice, and LPS-stimulated insulin resistance is also abrogated in the iNOS knockout [89]. These data support a link between protein S-nitrosylation and abnormal toll-like receptor (TLR) signaling in type 2 diabetes [90,91], and suggest that blocking iNOS induction might provide a mechanism for restoring insulin sensitivity. It has been proposed that the beneficial effects of thiazoledinediones, peroxisome proliferatoractivated receptor-y agonists that are used clinically for the treatment of diabetes mellitus, might be attributable to their inhibition of cytokine-induced iNOS expression [84]. Indeed, in diet-induced obese rats, the thiazoledinedione rosiglitazone largely abrogates Snitrosylation of IRB, IRS-1 and Akt, concomitant with a reduction in iNOS expression, amelioration of insulin resistance and restoration of insulin signaling [86]. Similar effects could underlie the improvement in insulin signaling by acute physical exercise, which reduces iNOS expression as well as IRS-1 and Akt S-nitrosylation [92].

Nitrosative stress in neurodegenerative diseases

Dysregulated protein *S*-nitrosylation (which can result from over-activation of NMDA receptors) appears to be prevalent in neurodegenerative disorders characterized, in particular, by the accumulation of misfolded proteins [2,93]. Hyper-*S*-nitrosylation of

parkin, protein disulfide isomerase (PDI), peroxiredoxin 2 (Pdx2), X-linked inhibitor of apoptosis (XIAP) and dynamin-related protein 1 (Drp1) is observed in brains from patients with neurological disorders (including Alzheimer's, sporadic Parkinson's and diffuse Lewy body diseases; Table 1), and S-nitrosylation of these proteins is implicated in stress-induced neuronal death [94–99]. S-nitrosylation initially appears to increase the E3 ligase activity of parkin and promote its auto-ubiquitylation [95], and ultimately results in its inhibition [94,95]. Zn-binding Cys residues in the RING domain of parkin have been identified as targets of S-nitrosylation [95]. NO inhibits both the ubiquitin-dependent degradation of parkin substrates (including synphilin-1) and the protection by parkin from cell death induced by protease inhibitors in α -synuclein and synphilin-1-expressing cells [94,95]. PDI is S-nitrosylated at one or both of the Cys thiols in each of its dithiol (Cys-Gly-His-Cys) active sites, resulting in inhibition of disulfide isomerase activity and accumulation of misfolded synphilin-1 in the endoplasmic reticulum [96]. Cumulative S-nitrosylation of parkin and PDI might underlie the accumulation of misfolded and ubiquitylated proteins that ultimately leads to cell death. S-nitrosylation of Cys residues within the baculoviral IAPrepeat (BIR) motifs of XIAP is implicated in inhibition of the antiapoptotic function of the enzyme (which requires a direct interaction with caspases), but in contrast to parkin, lowmass SNOs do not compromise the E3 ligase activity of XIAP [98]. Pdx2, a 2-Cys peroxiredoxin, is inactivated by S-nitrosylation of both its catalytic and resolving cysteine residues (C51 and C172, respectively), sensitizing dopaminergic neurons to H₂O₂dependent cell death [97]. Increased nitrosative stress, and Pdx2 S-nitrosylation, might contribute to the loss of dopaminergic neurons in Parkinson's disease [97]. It is notable that compounds that inhibit mitochondrial complex I (rotenone) or induce a Parkinson-like phenotype (MPTP/MPP⁺) also induce S-nitrosylation of each of these four proteins in vitro or *in vivo* [95–98]. Although a specific mechanism has not been elucidated, these data support the idea that, in neurons, nitrosative and oxidative stress might be linked. Finally, it has recently been shown that S-nitrosylation of Drp1 is enhanced in the brains of Alzheimer's (but not of Parkinson's) patients [99]. Similar to the effects of S-nitrosylation on dynamin (Box 1), S-nitrosylation of C644 within Drp1 promotes its multimerization and thereby mitochondrial fission, which causes neuronal damage [99]. Furthermore, exposure of nNOS-expressing cells to β-amyloid protein (a crucial effector of neurotoxicity in Alzheimer's disease) results in Drp1 S-nitrosylation, suggesting a role for S-nitrosylation in β-amyloid-associated neurotoxicity.

SNO-GAPDH and cell death – therapeutic implications in neurodegeneration and cancer

Nuclear accumulation of glyceraldehyde-3-phosphate dehydrogenase (GAPDH) is a hallmark of stress-induced cell death [100]. Stimulation of nNOS in neurons or of iNOS in macrophages (by NMDA and cytokines, respectively) triggers GAPDH *S*-nitrosylation at its active site, C145, which promotes the association of GAPDH with the E3 ubiquitin ligase Siah1, resulting in nuclear translocation of the SNO-GAPDH–Siah1 complex [101]. Within the nucleus, SNO-GAPDH stabilizes Siah1, which facilitates the ubiquitylation and degradation of nuclear proteins [101], and independently SNO-GAPDH is acetylated by and consequently activates the acetyltransferase p300/CREB binding protein, leading to the acetylation/activation of downstream targets, including the tumor supressor p53 [102]; both of these mechanisms promote cell death.

Accumulating evidence suggests that this pathway might be therapeutically relevant. Deprenyl and CGP-3466, drugs that protect neurons from stress-induced apoptosis [103] and slow the progression of Parkinson's disease [104], bind to GAPDH [105] and inhibit both GAPDH *S*-nitrosylation and the GAPDH–Siah1 interaction in MPTP-treated mice [103]. Deprenyl also induces the expression of thioredoxin [106], a denitrosylase for GAPDH

[107]. Conversely, BCNU (carmustine), a chemotherapeutic agent often used to treat brain and other cancers, increases SNO-GAPDH levels [108], providing a mechanism for sensitization of tumor cells to GAPDH-dependent death. *S*-nitrosylation of GAPDH and its nuclear accumulation, as well as NOS-dependent apoptosis, are also observed in thyroid carcinoma cells treated with TNF-related apoptosis-inducing ligand (TRAIL) [109], a promising cancer therapeutic [110]; iNOS is most likely the source of NO [109,111]. It has been suggested that this mechanism contributes to the tumor-selective effects of TRAIL [110].

Regulation of prostaglandin synthesis – preconditioning in ischemiareperfusion injury

Prostaglandin synthesis by cyclooxygenase (COX) is coupled to iNOS activation in cells and tissues, and *S*-nitrosylation is implicated in this effect. iNOS binds directly to and activates COX-2 by *S*-nitrosylation of a single cysteine (C526) proximal to the substrate (arachidonic acid; AA) binding site [112], and COX-2 *S*-nitrosylation and prostaglandin E₂ formation are attenuated by blocking iNOS–COX-2 interaction [112]. Similarly, binding and *S*-nitrosylation of COX-2 by nNOS appears to mediate NMDA receptor-induced neurotoxicity [113]. Upstream of COX-2, the formation of AA by cytosolic phospholipase $A_{2\alpha}$ (cPLA₂ α) is similarly regulated. In particular, iNOS activates cPLA₂ α by *S*nitrosylation of C152 [114], located within the N-terminal regulatory domain [115]. However, rather than interacting directly with iNOS, cPLA₂ α forms a ternary complex with COX-2 and iNOS, and induction of COX-2 is crucially required for cPLA₂ α *S*-nitrosylation and activation by iNOS [114].

These findings suggest that blocking the *S*-nitrosylation of COX-2 and/or cPLA₂ α might prove ameliorative in settings of inflammation, cancer and neurotoxicity. By contrast, the activation of COX-2 by *S*-nitrosylation is implicated in the cardioprotective effects of 3hydroxy-3-methylglutaryl-CoA reductase inhibitors (statins) [116,117], and in the protective effects of ischemia- or drug-induced preconditioning [118,119] in ischemia-reperfusion (I/ R) injury [120,121]. Although the molecular targets of prostanoids have not been elucidated fully [118], the protective effect of the statin atorvastatin (ATV) in reducing myocardial infarct size (when administered before ischemia or upon reperfusion) appears to involve both the up-regulation and PI3K/Akt-dependent activation of eNOS [117], the NF- κ Bdependent induction of iNOS and COX-2, and finally COX-2 *S*-nitrosylation and activation [116,117]. Consistent with these findings, the ATV-dependent reduction in infarct size after I/R is lost in both eNOS-knockout and iNOS-knockout mice (in which COX-2 upregulation and/or *S*-nitrosylation are abrogated) [117], and, in wild-type mice, ATV-dependent increases in myocardial 6-keto-PGF_{1 α} are blocked by an iNOS inhibitor without concomitant changes in COX-2 levels [116].

Anti-inflammatory activities of endogenous and exogenous S-nitrosothiols

Protein *S*-nitrosylation by iNOS-derived NO occurs downstream of TLR activation and can be proinflammatory; targets of TLR-dependent *S*-nitrosylation include surfactant protein D, which is increasingly implicated in lung inflammation [122,123]. However, numerous additional studies point to a role for *S*-nitrosylation in the feedback inhibition of TLRmediated signaling. NOS and/or low-mass SNOs inhibit the expression both of iNOS [124– 126] and of multiple cytokines (interleukin-1 β (IL-1 β) [127], IL-12 p40 subunit [128], IL-6, IL-8 and MIP-2 [129]), and induce the expression of the anti-inflammatory mediators IRAK-M [130] and SOCS-1 [131]. Possible mechanisms include: 1) inhibition of poly(ADP-ribose) polymerase isoform 1 (PARP-1) binding to the iNOS promoter via PARP-1 *S*-nitrosylation [(the site(s) of *S*-nitrosylation are unknown) [125]; 2) attenuation of

NF- κ B p50-p65 activity, including its binding to iNOS promoter DNA, by *S*-nitrosylation of a conserved Cys residue (C38 of p65) at the NF- κ B–DNA interface [120,124]; 3) *S*nitrosylation (by eNOS) at C216 of the adaptor protein MyD88 (which couples TLR to NF- κ B activation), resulting in the inhibition of its LPS-induced membrane association and binding to the sorting adaptor TIRAP, an interaction crucial for the recruitment of MyD88 to TLRs [129]; 4) *S*-nitrosylation at C179 of the inhibitory κ B kinase (which phosphorylates and promotes the degradation of the NF- κ B inhibitory protein, I κ B), resulting in its inhibition and the attenuation of NF- κ B activation through I κ B [121]. In addition to providing molecular bases for the anti-inflammatory actions of low-mass SNOs, these findings also suggest multiple potential targets for SNO-based therapy.

Anti-inflammatory and ameliorative roles for GSNO have been demonstrated in numerous diseases and disease models. For example, in a model of multiple sclerosis (experimental autoimmune encephalomyelitis; EAE), oral GSNO greatly improves clinical scores, attenuates infiltration of immune cells into the CNS, protects against severe demyelination and causes downregulation of proinflammatory cytokines and iNOS [12]. The protective role of GSNO in EAE might reflect disruption of leukocyte trafficking, inasmuch as GSNO pretreatment inhibits monocyte adhesion to TNFα-activated endothelial cells in vitro. This effect appears to be mediated through downregulation of endothelial cell adhesion molecules and S-nitrosylation of the p65 subunit of NF-KB [12]. Similarly, GSNO protects against experimental autoimmune uveitis, a model of human uveitis, a leading cause of blindness, with concomitant inhibition of proinflammatory cytokine expression [132]. Finally, GSNO is implicated as the barrier-inducing factor produced by enteric glia to regulate intestinal barrier function and inflammation [133]. Under conditions of glial cell ablation, GSNO protects against intestinal permeability, intestinal inflammation, TNF- α expression and mortality, and GSNO also inhibits human intestinal mucosal barrier permeability and restores colonic permeability in colon biopsies from patients with Crohn's disease [133]. Whether GSNO deficiency, due to a disruption in the enteric glial network, plays a role in Crohn's disease pathology is as yet undetermined.

The gaseous S-nitrosylating agent ethyl nitrite (ENO, which can be delivered by inhalation) might be more efficacious in some diseases than either GSNO, which requires processing (to S-nitrosocysteine or S-nitroso-cysteinylglycine) for cellular uptake, or inhaled NO (iNO), which has lower target specificity and a propensity to produce toxic byproducts. Antiinflammatory effects of ENO are observed in models of bronchopulmonary dysplasia (BPD) and acute lung injury/acute respiratory distress syndrome (ALI/ARDS). Whereas experimental BPD is induced in newborn rats exposed to hyperoxia, simultaneous exposure to hyperoxia and ENO protects rat pups from BPD, reduces cytokine expression and inhibits bronchiolar epithelial NF- κ B activation [9]. Consistent with its higher specificity for thiols, the effects of ENO are superior to those of iNO, which has not been proven to reduce the incidence of BPD [134] and shows a very narrow window of efficacy. ENO also offers protection against lung inflammation and injury in mice exposed to aerosolized LPS – a model of ALI/ARDS [10]. Interestingly, LPS alone induces protein denitrosylation in the lung, although the effect on inflammation is unknown [10]. ENO increases protein SNOs in airway lining fluid and in the lung, increases NF-KB p65 S-nitrosylation and inhibits LPSinduced NF- κ B DNA-binding [10]. These findings point to the potential for ENO as a preventative measure in ALI/ARDS, in contrast to iNO, which has failed to show a reduction in mortality or other outcome benefit [135,136], and which might even be associated with increased morbidity [136]. Thus, although NO production by one or more forms of NOS has been implicated in disease etiology (in particular as a proinflammatory agent), the onset of disease and/or the severity of symptoms can be attenuated by Snitrosylating agents.

The broad purview of protein S-nitrosylation in normal and disturbed cell function presents, in principle, novel therapeutic opportunities in a wide range of human diseases. These opportunities remain largely untapped. In addition, the improvement and dissemination of new methodologies for the analysis of protein S-nitrosylation – which have enabled many of the recent discoveries detailed here and which have been reviewed recently [137,138] – will allow the role of aberrant S-nitrosylation to be elucidated in additional, related and dissimilar disease states. The available data already indicate that, (i) in the case of skeletal muscle, a collection of disorders share in common dysregulated S-nitrosylation of RyR1 [75,77,78]; (ii) in the case of RBCs, seemingly disparate disorders share in common dysregulated S-nitrosylation of hemoglobin, which can influence disease symptomatology and which may be viewed as a new type of hemoglobinopathy; and (iii) in the case of Parkinson's disease, dysregulated S-nitrosylation of multiple disparate proteins might contribute to a common phenotype. However, it should be noted that, in many cases, it remains to be determined whether alterations in S-nitrosylation are causal in the development of disease. In addition, a number of disease-modifying drugs, including statins, deprenyl and BCNU, can exert ameliorative effects at least in part through modulation of Snitrosylation.

Further progress may be expected from the rapid and continuing accumulation of knowledge in several salient areas. It is increasingly apparent that protein *S*-nitrosylation is spatially and temporally regulated not only by the direct interaction or compartmentation of SNO targets with NO synthases, but also by the enzymes that remove SNO from glutathione (e.g. GSNOR) and proteins (e.g. thioredoxin [107]). These discoveries suggest novel mechanisms for the dysregulation of protein *S*-nitrosylation in disease and add to an expanding list of potential therapeutic targets. Finally, the related elucidation of a genetic basis in some cases of dysregulated *S*-nitrosylation, although still in its infancy, points to the possibility that molecular medicine will offer new avenues for treatment through the targeted repair (or pharmacological rescue) of the genetic defects that seem likely to underlie a significant number of pathophysiologies in which dysregulated *S*-nitrosylation plays a role.

Box 1. Protein S-nitrosylation can be regulated independently of changes in NOS expression, activity or colocalization with substrates

- Anion Exchange Protein 1 (AE1/band 3). Decreased transnitrosylative transfer of NO from SNO-Hb to the erythrocyte membrane protein band 3 in sickle cell anemia, resulting in decreased NO-based vasoactivity of red blood cells [49].
- Guanylyl cyclase. Nitroglycerin-mediated NO production by mitochondrial aldehyde dehydrogenase (mtALDH) and *S*-nitrosylation of both mtALDH and guanylate cyclase in tolerance to nitrovasodilators [23].
- Microbial proteins. Transition metal-dependent protein *S*-nitrosylation in yeast, implicated in nitrosative stress that subserves innate immunity [139].
- Hemoglobin. Superoxide dismutase (SOD)-catalyzed S-nitrosylation of hemoglobin within RBCs, proposed as a mechanism to regulate hypoxic vasodilation and SNO homeostasis [140]; the O₂-catalyzed (allosterically-coupled) auto-S-nitrosylation of Hb wherein NO transfers from oxidized heme to thiol, proposed as an integral aspect of the human respiratory cycle [48,141].
- NMDA receptor. Copper-induced S-nitrosylation of neuronal NMDA receptors, suggested as a mechanism for copper-dependent protection against NMDA receptor-mediated excitotoxicity [142]

 Mitochondrial proteins and GAPDH. Expression of mutant SOD results in decreased S-nitrosylation of mitochondrial proteins and GAPDH in an experimental model of amyotrophic lateral sclerosis [152].

Box 2. Regulation of G protein-coupled receptor signaling—pulmonary and cardiovascular consequences

Several lines of evidence point to a role for S-nitrosylation in the regulation of GPCR desensitization (loss of G-protein signaling) and downregulation that characterize heart failure. Diminished responsiveness (tachyphylaxis) to β-agonists, as assessed by vasodilation or cardiac contractility, is potentiated in vivo by NOS inhibitors and can be reversed by administration of low-mass SNOs [27,143]. GSNOR-knockout mice are protected from tachyphylaxis to β -agonists [6], and hearts and lungs from these mice display increased basal β -AR expression [27]. In addition, GSNO inhibits isoproterenolinduced downregulation of cardiac β -ARs in a mouse model of 'heart failure' [27]. In vitro, exogenous SNOs or eNOS overexpression inhibit desensitization and internalization of the β_2 -AR, in concert with inhibition of β -AR phosphorylation and β arrestin translocation to the plasma membrane [27]. These effects are consistent with inhibition of the G protein-coupled receptor kinase, GRK2, by S-nitrosylation of a crucial Cys (C340) in the catalytic domain, mutation of which abolishes GPCR-mediated Snitrosylation of GRK2 as well as the eNOS-dependent regulation of β_2 -AR internalization [27]. These data establish a possible mechanism for the tonic inhibition of cardiac and vascular β -AR desensitization by endogenous NO, and identify GRK2 as a regulatory locus. Inasmuch as both asthma and heart failure are characterized by attenuated β -AR signaling that results from increased receptor desensitization and downregulation, these disease states might reflect dysregulated GRK2 S-nitrosylation, one potential underlying mechanism being diminished NOS activity (which could result in part from inhibition of eNOS by GRK2 itself [144]). In addition, S-nitrosylation of dynamin and β-arrestin2 (βarr2) regulates GPCR internalization [56,145]. S-nitrosylation of C607 within dynamin promotes its mutimerization and activation, and S-nitrosylation of C410 within βarr2 promotes its interaction with clathrin/AP-2. Notably, SNO-βarr2 levels, and the affiliations of Barr2 with clathrin/AP-2, are enhanced in GSNORknockout mice [145].

Box 3. Targets of S-nitrosylation in cancer

S-nitrosylation of a number of proteins can be linked directly or indirectly to tumor proliferation (see [42] for review):

- Pro-angiogenic targets include HIF-1α, dynamin, Ras, COX-2 (see text) and PTEN [146].
- Anti-apoptotic targets include NF-κB p65 (see text), caspases [107], p53 [147] and Bcl-2 [148].
- Additional proliferative targets include methionine adenosyl transferase [149].
- DNA repair enzymes suppressed by S-nitrosylation include 8-oxoguanine glycosylase (Ogg1) and O⁶-alkylguanine-DNA alkyltransferase (AGT) [150,151].

S-nitrosylation has also been implicated in the effects of several anti-cancer therapeutics:

- BCNU (see text) and auranofin [107] are likely to inhibit cellular denitrosylation and induce general nitrosative stress.
- TRAIL appears to induce iNOS and activate SNO-GAPDH-dependent carcinoma cell death (see text).

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Fig. 1. NOS-dependent mechanisms of S-nitrosylation

Three principal mechanisms for regulation of NO synthase (NOS)-dependent protein *S*nitrosylation and the (patho)physiology associated with the induction or disruption of these mechanisms are shown. (a) Binding of a transcription factor (TF) to NOS promoter induces expression of the gene encoding NOS. (b) (Left) influx into the cytosol of extracellular or internal store-derived Ca²⁺ promotes Ca²⁺-calmodulin (CaM) binding and activation of eNOS and nNOS. Alternatively (right), phosphoinositide-3 kinase (PI3K) activates protein kinase B (Akt), which phosphorylates and activates eNOS. (c) Subcellular compartmentation (co-localization) of NOS and its substrates, which may involve a direct interaction (as in the illustrated case of a membrane-intercalated ion channel) is an important

determinant of the target specificity of S-nitrosylation, and dysregulated co-localization can result in hyper- or hypo-S-nitrosylation.



Fig. 2. Regulation of SNO homeostasis by S-nitrosoglutathione reductase (GSNOR)

(a) In cells, the low-mass *S*-nitrosothiol, *S*-nitrosoglutathione (GSNO), is in equilibrium with a subset of protein *S*-nitrosothiols. GSNO is metabolized by the enzyme GSNOR, and cells and tissues lacking GSNOR exhibit increased levels of SNO-proteins. (b) Analyses in GSNOR-knockout mice (GSNOR^{-/-}) have revealed numerous roles for protein *S*-nitrosylation. In particular, knockout animals exhibit low systemic vascular resistance and increased cardiac output under basal conditions, and are protected from mocardial infarction and allergen-induced airway hyperreactivity. (c) Mutations in *GSNOR*, as well as increased airway GSNOR expression and activity, are associated with human asthma.

Table 1

S-nitrosylated peptides and proteins that have been implicated in mammalian (patho)physiology.

	Adaptive immunity	Angiogenesis	Alzheimer's disease	Asthma	Cystic fibrosis	Diabetes (Type 1)	Diabetes (Type 2)	Duchenne muscular dystrophy	Efficacy of transfused blood	Endothelial dysfunction	Fatigue	Heart failure	Long QT syndrome	Lung inflammation	Malignant hyperthermia	Nitroglycerine tolerance	Parkinson's disease	Preeclampsia	Pulmonary arterial hypertension	Sickle cell disease	Stroke	Sudden cardiac death	Tumor maintenance	Tumor radiosensitivity
Arginase								-																
Cardiac sodium channel SNC5a																					с. — ў.			
Dynamin-related protein 1																								
Glutathione																								
Glyceraldehyde 3-phosphate dehydrogenase													2 2 2 2											
Hemoglobin																								
Hif-1α																								
Insulin receptor β																								
Insulin receptor substrate 1																								
Matrix metalloproteinase 9																								
Parkin							8																	
Peroxiredoxin 2																								\square
Protein disulfide isomerase																								
Protein kinase b																								
Ras																								\square
Ryanodine receptor 1																								\square
Ryanodine receptor 2																	-							\square
Serum albumin																								
Soluble guanylate cyclase				<u>, , , , , , , , , , , , , , , , , , , </u>			9 0			3 - 2 -		< - 24	8 - 0		1	1					5 C			
Surfactant protein D															1								\square	
X-linked inhibitor of apoptosis																							Π	\square

Table 2

Amelioration of disease/tissue injury in humans and in animal models by treatments that affect S-nitrosylation.

	Treatment	Disease/Injury (Model)	Refs.
	Ethyl nitrite:: gaseous <i>S</i> -nitrosylating agent	Pulmonary arterial hypertension (PAH) ¹ Persistent PAH of newborns ¹ Bronchopulmonary dysplasia Acute lung injury/acute respiratory distress syndrome	[52] [53] [9] [10]
Increased SNO	S-nitrosoglutathione: S-nitrosylating agent	Heart failure Experimental autoimmune encephalomyelitis Experimental autoimmune uveoretinitis Crohn's disease	[27] [12] [132] [133]
	Nitric oxide ²	Loss of vasodilation in stored red blood cells Sickle cell disease	[50,51] [49]
	eNOS adenovirus	Duchenne muscular dystrophy	[81]
	Atorvastatin ³	Myocardial infarction/preconditioning	[116,117]
No.	N-acetyl cysteine ⁴	Malignant hyperthermia, environmental heat stroke	[75]
reased S	Rosiglitazone ⁵ Exercise ⁵ iNOS antisense oligo	Type 2 diabetes	[86,92]
Dec	Deprenyl ⁶	Neurodegeneration (experimental Parkinson's disease)	[103]
	S107 ⁷	Duchenne muscular dystrophy Exercise-induced fatigue	[77] [78]

¹Effects in humans.

 2 Ex vivo repletion of *S*-nitroso-hemoglobin in red blood cells.

 3 Induces *S*-nitrosylation (e.g. of COX-2) by eNOS and iNOS.

⁴A glutathione (GSH) precursor that reverses nitrosative stress.

⁵ Lowers iNOS-mediated S-nitrosylation.

 $^{6}{\rm Binds}$ to GAPDH and blocks iNOS-induced S-nitrosylation.

⁷Potentiates the interaction between SNO-RyR1 and FKBP12 to ameliorate calcium leak.