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The role of S-nitrosogluthione reductase (GSNOR) in human disease and therapy

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ABSTRACT

S-nitrosogluthione reductase (GSNOR), or ADH5, is an enzyme in the alcohol dehydrogenase (ADH) family. It is unique when compared to other ADH enzymes in that primary short-chain alcohols are not its principle substrate. GSNOR metabolizes S-nitrosogluthione (GSNO), S-hydroxymethylglutathione (the spontaneous adduct of formaldehyde and glutathione), and some alcohols. GSNOR modulates reactive nitric oxide (\cdot NO) availability in the cell by catalyzing the breakdown of GSNO, and indirectly regulates S-nitrosothiols (RSNOs) through GSNO-mediated protein S-nitrosation. The dysregulation of GSNOR can significantly alter cellular homeostasis, leading to disease. GSNOR plays an important regulatory role in smooth muscle relaxation, immune function, inflammation, neuronal development and cancer progression, among many other processes. In recent years, the therapeutic inhibition of GSNOR has been investigated to treat asthma, cystic fibrosis and interstitial lung disease (ILD). The direct action of \cdot NO on cellular pathways, as well as the important regulatory role of protein S-nitrosation, is closely tied to GSNOR regulation and defines this enzyme as an important therapeutic target.

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S-nitrosation; ADH5


Introduction

S-nitrosogluthione reductase (GSNOR) is an important regulator of human health and disease. The modulation of protein S-nitrosation by GSNOR contributes to a host of maladies and can be exacerbated by the dysregulation of GSNOR. In recent years, much effort has been dedicated to identifying a safe and efficacious means to alter GSNOR activity. A myopic investigation of GSNOR would reveal little more than its inherent ability to metabolize S-nitrosogluthione (GSNO) (Jensen *et al.*, 1998), S-hydroxymethylglutathione (HMGSH) (Hedberg *et al.*, 2000), and a handful of alcohols (Adinolfi *et al.*, 1984; Jensen *et al.*, 1998). If we look beyond the direct actions of the enzyme itself, it quickly becomes apparent that GSNOR influences several downstream and parallel pathways (Figure 1). One of the most important is GSNOR's regulation of GSNO, and by extension, nitric oxide (\cdot NO) and protein S-nitrosation. \cdot NO is a reactive nitrogen species (RNS) that is critical to the normal function of most cell types (Beckman & Koppenol, 1996; Moncada *et al.*, 1991; Radi *et al.*, 1991; Salvador Moncada, 1994). It is a powerful smooth muscle relaxing agent (Bradley *et al.*, 1998; Buxton *et al.*, 2001; Ricciardolo *et al.*, 2004; Tomita *et al.*, 2002), cardiopulmonary regulator (Liu *et al.*, 2004; Tamargo *et al.*, 2010), neuroeffector (Bredt & Snyder,

1992; Corti *et al.*, 2014) and immune system modulator (MacMicking *et al.*, 1997). \cdot NO is likely carried as GSNO from endothelium, and other sources, and acts as a stable \cdot NO reserve (Broniowska *et al.*, 2013; Smith & Marletta, 2012). GSNO can transfer its \cdot NO moiety to a cysteine thiol, resulting in the posttranslational modification (PTM) S-nitrosation/S-nitrosylation (Stamler *et al.*, 1992). S-nitrosation describes a thiol (e.g. cysteine) converted to a S-nitrosothiol (RSNO) by a one-electron oxidation from the \cdot NO radical (Smith & Marletta, 2012). The term nitrosylation describes addition of an \cdot NO group to a metal centered protein such as guanylyl cyclase (Martínez-Ruiz & Lamas, 2004). Researchers have used both terms to describe \cdot NO addition to a protein thiol. We employ S-nitrosation to refer to protein modifications on cysteine residues. Protein S-nitrosations are also referred to in the literature in a fashion that takes into account protein and non-protein nitrosations (e.g. RSNO). We employ the term RSNO as it appears in the literature.

Alcohol dehydrogenase family overview

The alcohol dehydrogenase (ADH) family of enzymes have been investigated for well over a century

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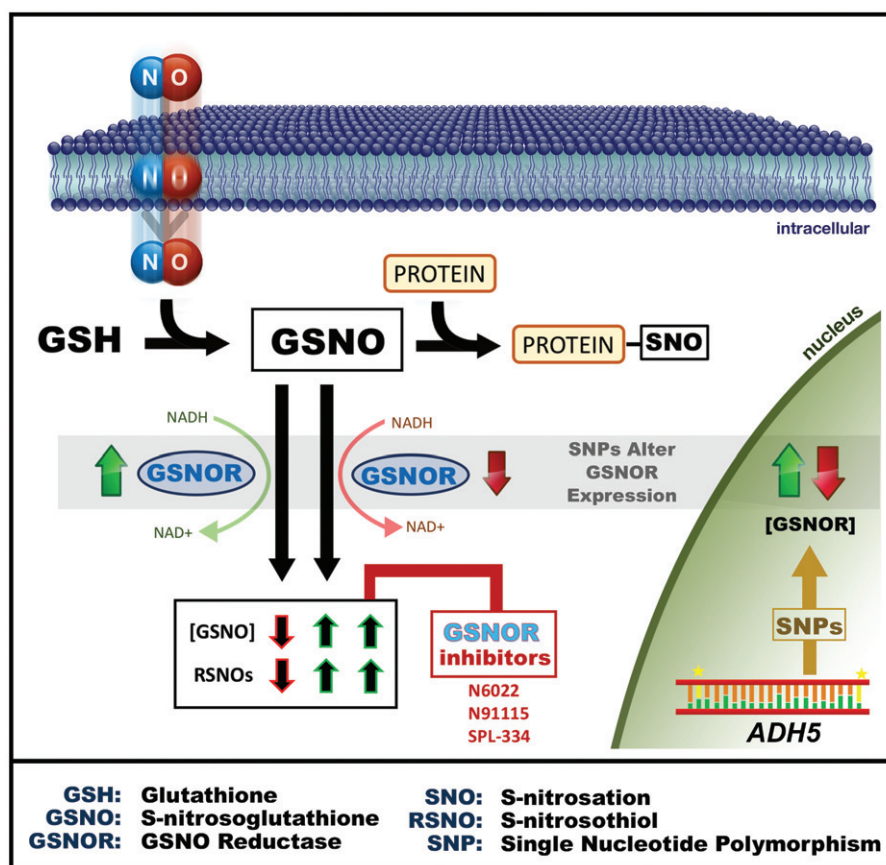


Figure 1. GSNOR in the cell. One of the principle functions of GSNOR is to metabolize GSNO. SNPs in ADH5 can affect expression of GSNOR in the cell, which in turn alters the concentration of GSNO and total levels of RSNOs (term to include nitrosation of cysteine residues, e.g. SNO). Inhibitors of GSNOR increase available GSNO and increase total RSNOs (see colour version of this figure at www.tandfonline.com/ibmg).

(Battelli & Stern, 1910; Daniel, 1909; Lutwak-Mann, 1938). They are evolutionarily conserved from bacteria to man (González-Duarte & Albalat, 2005; Liu *et al.*, 2001) and are categorized into five distinct classes that contain seven known isoforms (Table 1). ADH enzymes perform several important functions in human cells. The most well studied of these is the metabolism of short chain alcohols. Ethanol, being of significant cultural relevance due to its widespread consumption and abuse (Oscar-Berman & Marinkovic, 2003), has led to an extensive investigation of the entire ADH family. Most ADH enzymes have some affinity for ethanol. In hepatocytes, ADH1A (formerly ADH1), ADH1B (formerly ADH2) and ADH1C (formerly ADH3), are responsible for the oxidative catabolism of ethanol to acetaldehyde before further processing in the Krebs cycle, or elimination (Cederbaum, 2013). ADH4, a class II ADH (Svensson *et al.*, 2001) whose sequence is 70% homologous to ADH1, catalyzes the oxidation of retinol (Vitamin A), and bolsters ethanol metabolism in the liver (Ramchandani *et al.*, 2001). Numerous single nucleotide polymorphisms (SNPs) in the genes encoding the ADH family affect the rate of ethanol metabolism. These SNPs have

been linked to some forms of alcoholism and cancer (Edenberg, 2007; Hurley & Edenberg, 2012). Other ADH SNPs have been correlated with schizophrenia, Parkinson's disease, asthma and autism in certain populations (Bowers *et al.*, 2011; Buervenich *et al.*, 2000; Wu *et al.*, 2007; Zuo *et al.*, 2013). GSNOR (ADH5), the focus of this review, is differentiated from other ADH enzymes in that primary short chain alcohols, in particular ethanol, are not its principal substrate. ADH6 has been identified in both fetal and adult livers, but its function remains unclear as this enzyme has yet to be isolated for biochemical analysis (Edenberg, 2007; Östberg *et al.*, 2016). A recent examination of ADH6 has provided evidence that it may act as an S-nitroso-CoA reductase (Anand *et al.*, 2014). Similarly, ADH7's function remains elusive. Available data suggest ADH7 may serve a role in seemingly disparate cellular functions and diseases, such as: first pass gastric metabolism of ethanol (Lee *et al.*, 2006), retinol metabolism (Chase *et al.*, 2009), Parkinson's disease (Buervenich *et al.*, 2000) and even personality traits in some individuals with substance dependence (Luo *et al.*, 2008). Clearly, the ADH family of enzymes performs a diverse and important role in

Table 1. ADH variants. ADHs are most commonly recognized as highly effective metabolizers of ethanol. ADH5 evolved independently from class I and class II ADHs varies from other ADHs in that GSNO and HMGSH are its primary substrates. The identification and analysis of other ADHs continue.

Gene name	Principle substrate	Uniprot Identifier	Subunits	Enzyme class
<i>ADH1A</i>	Ethanol	P07327	α	I
<i>ADH1B</i> (formerly <i>ADH2</i>)	Ethanol	P00325	β	I
<i>ADH1C</i> (formerly <i>ADH3</i>)	Ethanol	P00326	γ	I
<i>ADH1(D-H)</i> ^a	Unkown	–	–	–
<i>ADH4</i>	Ethanol/Retinol	P08319	π	II
<i>ADH5</i>	GSNO/HMGSH	P11766		III
<i>ADH6</i>	Ethanol/S-nitroso-CoA ^b	P28332	μ/σ^c	V
<i>ADH7</i>	Retinol	P40394	σ	IV
<i>ADH(8-14)</i> ^a	Unkown/Retinol ^b	–		–

Source: Modified from Edenberg (2007).

^aNon-human.

^bLimited evidence.

^ccDNA data.

the cellular metabolism of endogenous and exogenous chemicals. Here we focus on the function, significance and therapeutic potential of modulating GSNOR activity.

Nomenclature of alcohol dehydrogenases

The ADH family of enzymes has had several overlapping naming schemes in the past (Holmquist & Vallee, 1991; Staab *et al.*, 2008). This has led to ambiguity in the literature and is due in part to the fact that naming assessments have historically been guided by substrate specificity, phylogenic classification and publication date. GSNOR was not disambiguated from glutathione-dependent formaldehyde dehydrogenase (FDH) until 1989 when it was found that these two proteins were in fact the same enzyme (Koivusalo *et al.*, 1989). A formal attempt to reconcile the nomenclature began in 1999 when it was proposed that ADH proteins use numeric Arabic designators to identify each class of enzyme (Duester *et al.*, 1999). In recent years, the research community has generally adopted the gene naming guidelines put forth by the Human Genome Organization's Gene Nomenclature Committee (Wain *et al.*, 2002). Of all the ADH enzymes, GSNOR naming is particularly convoluted in this respect. While this protein is still sometimes referred to in the literature by its nonstandard name, ADH3 (as in class III ADH), the official gene designator is now *ADH5*, and the protein is GSNOR, ADH5 or alcohol dehydrogenase 5 (class III) χ -polypeptide. It can also be found in the literature under several other monikers: formaldehyde dehydrogenase (FDH or FALDH); alcohol dehydrogenase X (ADHX); alcohol dehydrogenase class-3 (ADH-3); $\chi\chi$ -ADH (homodimeric *chi* ADH); ADH 5; glutathione-dependent formaldehyde dehydrogenase (GSH-FDH); and S-(hydroxymethyl) glutathione dehydrogenase (EC 1.1.1.284). For purposes of clarity,

this review will address the gene as *ADH5*, and the protein as ADH5 or GSNOR.

ADH5: structure/localization

ADH5, the gene that encodes GSNOR, is located on the reverse strand of chromosome 4 (4q23 – chr4:99993567 – 10000985) (Smith, 1986). *ADH5* is tandemly aligned in the same orientation as the other genes that encode for the entire family of ADH enzymes. Phylogenic analysis of the *ADH5* locus revealed that GSNOR evolved independently from class I and II ADH (Adinolfi *et al.*, 1984), and it is highly conserved across most vertebrate species (Foglio & Duester, 1996). GSNOR has a molecular weight of 39,724 Da and is translated to a 374 amino acid enzyme (UniProtKB identifier: P11766) via 9 exons (Hur & Edenberg, 1992). Glu-67 and Arg-368 are highly conserved essential amino acids important to the catalytic mechanism of this enzyme (Sanghani *et al.*, 2006). Splice variants of *ADH5* exist and result in the production of truncated proteins; however, their functional relevance has not been documented (Höög *et al.*, 2001).

GSNOR functions as a homodimer (Figure 2) (Yang *et al.*, 1997) and is localized to the nucleus and cytoplasm (Fernández *et al.*, 2003). Amino acid substitutions in the subunit interacting portions of the coenzyme-binding domain prevent heterodimeric variants from being generated with other ADH enzymes (Julià *et al.*, 1988). Each subunit binds a catalytic and structural Zn²⁺ cofactor (Kaiser *et al.*, 1988; Östberg *et al.*, 2016), for a total of four Zn²⁺ ions per functional enzyme. In addition to Zn²⁺, GSNOR also requires a coenzyme that can vary based upon the substrate. These include: nicotinamide adenine dinucleotide (NAD⁺), its reduced form NADH, NADPH + H⁺ or NAD(P)⁺ (Gupta & Igamberdiev, 2015; Hedberg *et al.*, 2003; Jensen *et al.*, 1998; Sanghani *et al.*, 2000).

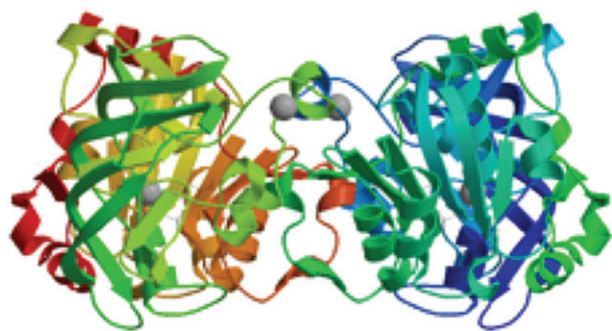


Figure 2. GSNOR quaternary model derived from X-ray diffraction (2.7 Å) and displayed as a functional $\chi\chi$ homodimer with (2) Zn⁺ ions and (1) NADH co-enzyme per subunit. swissmodel.expasy.org SMTL id 1teh.1 (see colour version of this figure at www.tandfonline.com/ibmg).

In general, ADH enzymes are highly expressed in the liver, the upper digestive tract and the kidneys (Zuo *et al.*, 2013). *ADH5* RNA has been recognized in all major human tissue types with protein expression highest in smooth muscle, liver, epididymis, kidney and testis (Giri *et al.*, 1989). GSNOR is an important negative regulator of neuronal differentiation during development (Wu *et al.*, 2014) and is the only known ADH enzyme present in the brain (Beisswenger *et al.*, 1985; Galter *et al.*, 2003). Conversely, GSNOR protein expression is negligible or non-existent in skeletal muscle, lymph nodes, spleen, bone marrow, cerebellum and the lateral ventricle (If & Wb, 2017).

Substrates

S-nitrosoglutathione/formaldehyde

As with most enzymes, GSNOR has a varying degree of affinity for several substrates. The two primary targets of GSNOR are GSNO, and HMGSH, the spontaneous adduct of formaldehyde and glutathione. HMGSH binds at the zinc active site and interacts with the highly conserved residues Arg114/115, Asp55, Glu57 and Thr46 (Engeland *et al.*, 1993; Sanghani *et al.*, 2002). That being said, the rate of substrate conversion (K_{cat}) is about 20-fold higher for GSNO over HMGSH (Green *et al.*, 2012; Hedberg *et al.*, 2003; Salisbury & Bronas, 2015; Sanghani *et al.*, 2000; Staab *et al.*, 2008). Both reactions are dependent on an abundant source glutathione (GSH) in the cell. GSH is the major thiol in mammalian cells and while concentrations can reach as high as 10 mM (Bateman *et al.*, 2008), they are typically 1 mM. Under stress conditions, the concentration can fluctuate dramatically and drive GSNO toward atypical reactions (Figure 3) (Salisbury & Bronas, 2015; Staab *et al.*, 2009). The enzymatic activity of human recombinant GSNOR for GSNO exhibits a K_m of approximately 27 μ M and a

k_{cat} value of between 2400 and 12,000 min^{-1} (Fernández *et al.*, 2003; Hedberg *et al.*, 2003).

Alcohols

GSNOR more readily acts upon alcohols of greater chain length than class I ADH enzymes (Figure 4). This is due in part to a longer span between the binding and active site of the enzyme (Salisbury & Bronas, 2015), as well as amino acid substitutions that affect binding affinity (Julià *et al.*, 1988; Östberg *et al.*, 2016). As a result of these evolutionary divergences, GSNOR is not optimized for metabolizing short-chain alcohols. Consequently, it is not a misnomer to identify GSNOR as an ADH. GSNOR metabolizes both ethanol and medium/long chain alcohols (preferring a double-bond in the beta position). The active site of GSNOR cannot be saturated by ethanol (Beisswenger *et al.*, 1985), and the high activity of class I ADH enzymes toward ethanol minimizes the functional role of ethanol metabolism by GSNOR. Several K_m values for EtOH (all >2 M) (Lee *et al.*, 2003; Sharma *et al.*, 1989) have been reported in the literature, with a k_{cat} of $33 \pm 3 \text{ min}^{-1}$ (Beisswenger *et al.*, 1985; Lee *et al.*, 2003). GSNOR's ability to metabolize EtOH is far surpassed by those of class I ADH enzymes whose K_m values range from 0.05 to 40 mM. As such, medium and long chain alcohols (>4 carbons) (Holmquist & Vallee, 1991; Salisbury & Bronas, 2015; Theorell *et al.*, 1969; Wagner *et al.*, 1984) are more freely oxidized by GSNOR (Staab *et al.*, 2009).

Other substrates

As with most enzymes, the entire cohort of ADH5 substrates is not fully known. Additional classes of molecules such as ω -hydroxy fatty acids (Achkor *et al.*, 2003; Boleda *et al.*, 1993; Moulis *et al.*, 1991) exhibit a limited affinity for the enzyme. The ability of GSNOR to metabolize retinol remains in question. ADH7 (a class IV ADH) is the primary ADH accountable for retinol metabolism (Cederbaum, 2013), but there is evidence to support GSNOR's contribution in the retinoid-signaling pathway. Studies have shown that *ADH5*^{-/-} null mice exhibit reduced retinoic acid production (Molotkov *et al.*, 2002), and the presence of *ADH5* transcript in human fetal lungs correlates with a decrease in the presence of retinol (Coste & Labbe, 2011). Ultimately, the exact nature of relationship between GSNOR and retinol is still under investigation (Boleda *et al.*, 1993; Cañestro *et al.*, 2010; González-Duarte & Albalat, 2005).

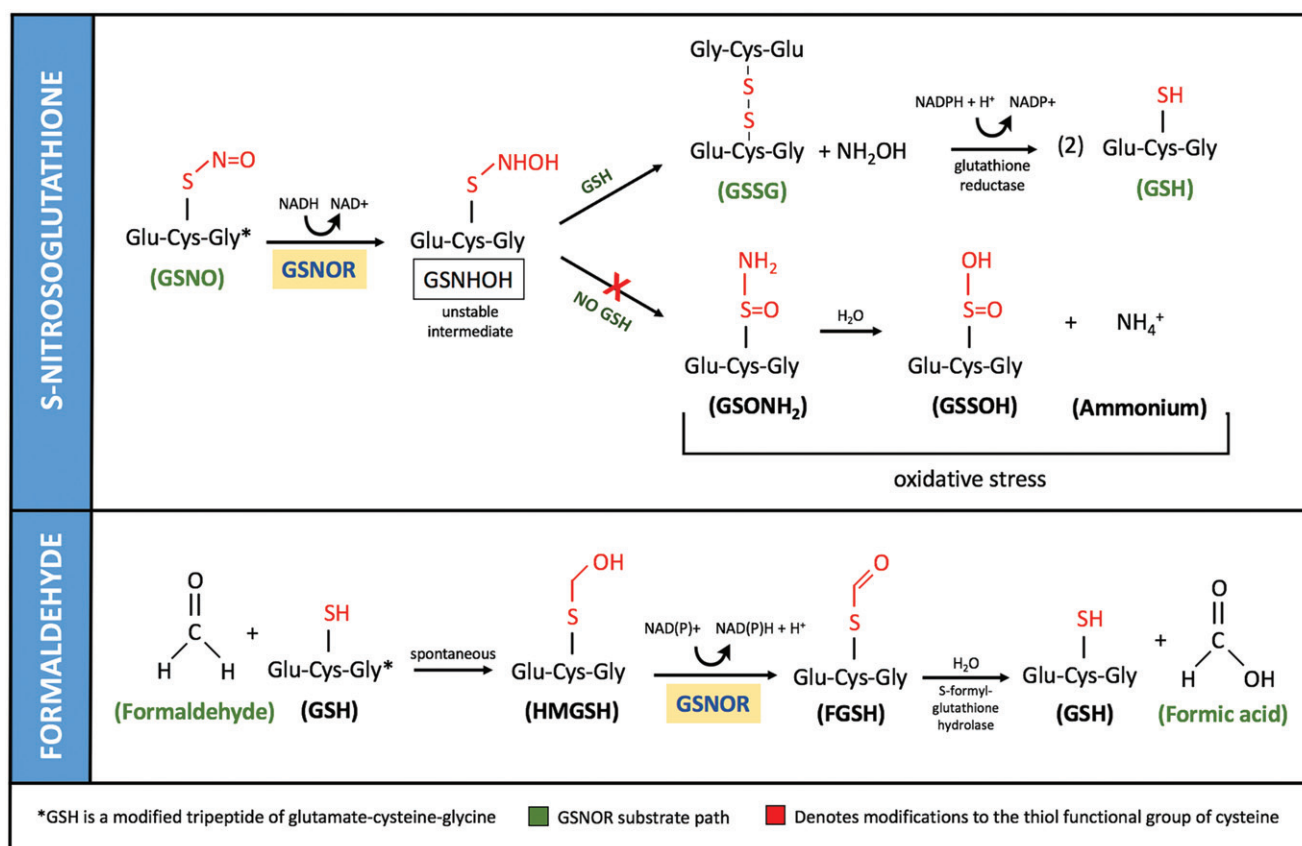


Figure 3. GSNO and HMSGH metabolism by GSNOR – GSNOR metabolizes multiple substrates. S-nitrosoglutathione (GSNO), one of the primary substrates for GSNOR, is first enzymatically degraded to an unstable intermediate, N-hydroxysulfonamide (GSNHOH). In the presence of additional glutathione (GSH), GSNHOH will be converted to glutathione disulfide (GSSG). Under certain conditions, such as high levels of oxidative stress, GSH will not be sufficiently available, and other products, such as glutathione sulfonamide (GSONH₂) and glutathione sulfonic acid (GSSOH) will be formed. GSNOR can also oxidize the spontaneous adduct of formaldehyde and GSH, S-(hydroxymethyl)glutathione (HMSGH), to S-formylglutathione (FGSH) (see colour version of this figure at www.tandfonline.com/ibmg).

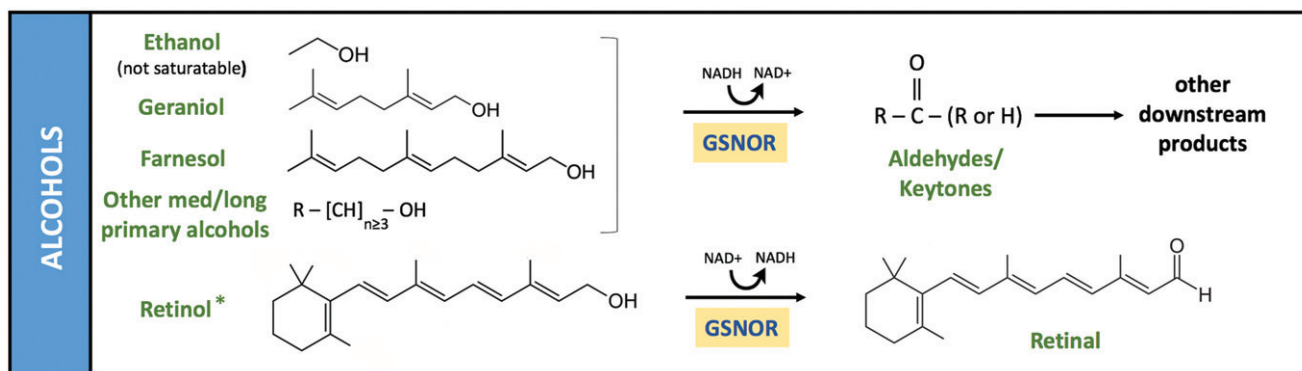


Figure 4. Alcohol metabolism by GSNOR: GSNOR can metabolize medium and long chain primary alcohols to aldehydes and/or ketones before being further processed by other enzymes. It preferentially metabolizes medium and long chain alcohols with a double-bond on the beta carbon, however, its specificity for all alcohols is much lower than for GSNO. *Evidence supporting GSNOR's ability to metabolize retinol to the aldehyde retinal is limited (see colour version of this figure at www.tandfonline.com/ibmg).

GSNOR: health and disease

GSNOR is integral to the modulation of •NO in the cell. •NO is produced enzymatically in many cell types (Schmidt & Walter, 1994). Free •NO is a highly reactive

uncharged radical with a half-life of ~1–5 second *in vivo* (Kelm & Schrader, 1990), and will often establish a stable RSNO equilibrium with GSH in the form of GSNO (Wink & Mitchell, 1998). •NO, and by extension, GSNO,

plays a critical role in smooth muscle relaxation (Bradley *et al.*, 1998; Buxton, 2004; Liu *et al.*, 2016; Ricciardolo *et al.*, 2004; Tomita *et al.*, 2002) cardiopulmonary regulation (Rastaldo *et al.*, 2007; Sears *et al.*, 2004; Tamargo *et al.*, 2010), neuronal signaling (Shahani & Sawa, 2011), as well as dozens of other intra/extracellular functions (Pacher *et al.*, 1995; Salvador Moncada, 1994). The dysregulation of $\cdot\text{NO}$ production and metabolism can lead to drastic changes in protein S-nitrosation (Foster *et al.*, 2003, 2009), an important PTM, and can have numerous other downstream consequences.

Oxidative/nitrosative stress

The dysregulation of GSNO through aberrant GSNOR modulation, when combined with oxidative stress, can further exacerbate disease. During conditions of cellular stress RNS, such as peroxynitrite (ONOO^-), are formed when $\cdot\text{NO}$ reacts with superoxide ($\text{O}_2^{\cdot-}$) (Squadrito & Pryor, 1998). Not only does oxidative stress commandeer available $\cdot\text{NO}$ and GSH (Rahman & MacNee, 2000), but peroxynitrite can cross the cell membrane and react directly with protein thiols (Alvarez & Radi, 2003), which may prevent S-nitrosation. RNS also induce S-glutathionylation of protein thiols (Dalle-Donne *et al.*, 2009), further depleting the GSH pool (Klatt & Lamas, 2000). Decades of research have left little question as to detrimental effects of oxidative/nitrosative stress (Dalle-Donne *et al.*, 2006; Guzik *et al.*, 2002; Münzel *et al.*, 1997), and the mechanistic underpinnings of this process have been thoroughly investigated (Apel & Hirt, 2004; Valko *et al.*, 2007). For the purpose of this review, it should be noted that this process can alter the levels of $\cdot\text{NO}$ and GSH in the cell, which in turn can affect $\cdot\text{NO}$ /GSNO signaling.

GSNO and S-nitrosation

Any investigation into the modulation/activity of GSNOR would not be complete without mention of S-nitrosation. The study of this PTM and its influence on normal cell-signaling and disease has significantly impacted research and medicine for over 25 years (Broniowska & Hogg, 2012; Foster *et al.*, 2009; Stamler *et al.*, 1992).

The detection and quantitation of RSNOs in biological systems are inherently challenging. The biotin switch technique (Jaffrey & Snyder, 2001), in which S-nitrosated cysteines are reduced and biotinylated, provides a simple and elegant method for the qualitative detection of S-nitrosated proteins. An analysis of a wide variety of RSNO measurement techniques, including the biotin switch, has established that artifacts are

common when measuring RSNOs and it is not always possible to identify which thiols have been S-nitrosated (Giustarini *et al.*, 2003). Newer techniques have become available in recent years (Chen *et al.*, 2013; Devarie-Baez *et al.*, 2013), such as tandem mass spectrometry (MS/MS) of S-nitrosated protein thiols (Murray *et al.*, 2012; Ulrich *et al.*, 2013), that are highly quantitative. Beyond the problem of quantitation, it has been proposed that other thiol modifications such as dithiol/disulfide exchange, S-glutathionylation and oxidation, may affect signaling more readily than do RSNOs (Lancaster, 2008), and should be investigated along with S-nitrosation.

As with phosphorylation, S-nitrosation regulates cellular mechanisms and affects protein-protein interactions. The intracellular availability of $\cdot\text{NO}$ and its functional derivatives, like GSNO, affect protein S-nitrosation (Broniowska & Hogg, 2012; Hess *et al.*, 2005; Thomas & Jourdeuil, 2012). GSNOR is a potent negative regulator of GSNO in smooth muscle (Que *et al.*, 2009). The aberrant expression of ADH5, as with many ADH subclasses, is associated with disease (Jelski & Szmitkowski, 2008; Jelski *et al.*, 2009; Laniewska-Dunaj *et al.*, 2013). In fact, the deletion of the *ADH5* gene increases both the levels of GSNO and total protein S-nitrosation *in vivo* (Liu *et al.*, 2001). Protein S-nitrosation is of intense interest to researchers and clinicians as the hypo/hyper-S-nitrosation of a diverse set of proteins, spanning nearly every tissue types, can have a drastic effects in disease (Foster *et al.*, 2009). Some of these include: Type 2 diabetes (Carvalho-Filho *et al.*, 2005), sickle cell anemia (Bonaventura *et al.*, 1999, 2002), ventricular arrhythmia in individuals with the Duchenne muscular dystrophy (Fauconnier *et al.*, 2010), cell death and survival pathways (Iyera *et al.*, 2011), post-infarct cardio-protection (Methner *et al.*, 2014), pregnancy/parturition (Ulrich *et al.*, 2013) and many others. Interestingly, GSNOR itself is a cysteine rich protein that is S-nitrosated by GSNO, which in turn initiates a feedback loop that affects GSNOR expression (Guerra *et al.*, 2016) and activity (Brown-Steinke *et al.*, 2010). Although it is beyond the scope of this review, it should be noted that GSNOR dysregulation in plants can result in significant biotic and abiotic nitrosative events that affect growth, development and survival (Leterrier *et al.*, 2011; Shi *et al.*, 2015; Yun *et al.*, 2016).

GSNOR dysregulation

GSNOR dysregulation has been implicated in numerous disease states (Figure 5). The use of models and *ADH5*^{-/-} knockout animals has uncovered surprising and valuable data related to GSNOR function. RSNO levels, as

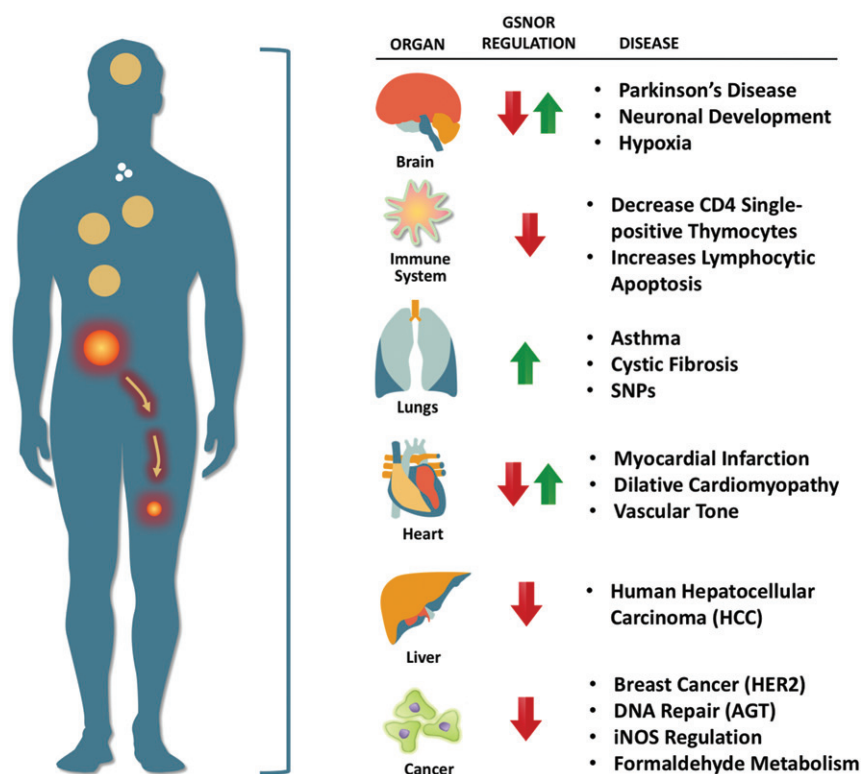


Figure 5. The dysregulation of GSNOR can initiate or exacerbate many disease states. This is due in part to GSNOR's indirect function as a S-nitrosothiol modulator, as well as its ability to mediate canonical NO cell signaling through GSNO metabolism. GSNOR inhibitors are being actively investigated to treat certain disorders in which increased NO availability would be beneficial (see colour version of this figure at www.tandfonline.com/ibmg).

well as canonical NO-mediated pathways, are severely altered when GSNOR activity is modulated.

Cardiovascular health

One of the major organs affected by GSNOR is the heart and surrounding vascularity. It has long been known that •NO and S-nitrosation protect the body from cardiovascular disease. Following myocardial infarction, *ADH5*^{-/-} mice exhibit enhanced cardiac regenerative capabilities as a result of increased cardiac stem cell turnover (Hatzistergos *et al.*, 2015), as well as a reduction in myocardial infarct size and higher coronary vascular density (Lima *et al.*, 2009). Moreover, de-S-nitrosation of cardiac ryanodine receptor 2 (RyR2) in *ADH5*^{-/-} mice results in decreased peripheral vascular tone due to calcium "leak" (Beigi *et al.*, 2012). In skeletal muscle only about 1 in 50 cysteines on the ryanodine receptor are S-nitrosated, indicating that this PTM, even when conservatively distributed, can drastically alter protein function (Sun *et al.*, 2001). Taken together this data suggests that RyR2 S-nitrosation modulates calcium storage in the sarcoplasmic reticulum. There is clearly a complex relationship between the correlative observation of an increase in S-nitrosation and GSNOR dysregulation.

Immune system

GSNOR performs an important protective role in the immune system's development of lymphocytes. *ADH5*^{-/-} KO mice show increased RSNO production that decreases CD4 single-positive thymocyte development, and increases lymphocytic apoptosis (Yang *et al.*, 2010). Damage to immune cells from nitrosative stress in *ADH5*^{-/-} mice results in a significant increase in the animal's susceptibility to pulmonary infection by *K. pneumoniae* as well as multi-fold increases of the bacteria in the spleen and blood, resulting in increased inflammation (Tang *et al.*, 2013a, 2013b). Enhanced nitric oxide synthase (NOS) 2 activity in monocytes and macrophages increases •NO production and elicits a cytostatic or cytotoxic response against bacteria, viruses and other intruders, but also increases inflammation (MacMicking *et al.*, 1997). The bronchoalveolar lavage fluid of asthmatics consists of high macrophage levels as well as significantly increases GSNOR activity (Que *et al.*, 2009). Inhibiting GSNOR in these patients increases total RSNOs and restores inflammatory markers to near baseline levels while limiting ova-induced NFκB activation (Blonder *et al.*, 2014). Ultimately, the balance between GSNOR activation and inhibition is critical in maintaining balance in the immune system.

Brain development and function

GSNOR regulation in the brain affects a broad swath of cellular functions ranging from neural development and maturation to other neurodegenerative diseases more typically associated with adult and geriatric populations. These disease states are often the result of aberrant protein S-nitrosation caused by the dysregulation of GSNOR. For instance, in developing and adult mouse brains the overexpression of GSNOR results in decreased neuronal differentiation in part due to de-S-nitrosation of histone deacetylase 2 (HDAC2) (Wu *et al.*, 2014). Conversely, *ADH5*^{-/-} mice exhibit neuromuscular atrophy as a result of a decrease in muscle mass, while also presenting with neuropathic behavior (Montagna *et al.*, 2014). In *Drosophila*, GSNOR overexpression results in visual pattern memory defects which can be rescued by co-expression of cyclic-GMP dependent protein kinase G (PKG) (Hou *et al.*, 2011). This occurs independently from neuronal development and implies an adjacent regulatory role for GSNOR in the PKG phosphorylation pathway. Neuronal homeostasis is also affected by GSNOR. In a Parkinson's disease model using neuronal (SH-SY5Y) cells, a decrease in GSNOR availability results in activation of nuclear factor Nrf2 ((erythroid-derived 2)-like 2), which regulates the expression of antioxidant proteins (Rizza *et al.*, 2015). Interestingly, GSNOR may also affect the phosphorylated state of platelet-derived growth factor receptor-β (Palmer *et al.*, 2015) in the brainstems of mice during hypoxic exposure. When these data are considered as a whole, it is apparent that deviating GSNOR activity and expression from baseline can have drastic consequences in both the developing and mature brain.

Cancer

The link between GSNOR dysregulation and cancer is not well understood. GSNOR deficiency has been known to affect the rate of genomic mutations in mice by increasing the frequency of A:T to T:A transposition (Leung *et al.*, 2013). This may be the result of a GSNOR-mediated reduction in activity of the DNA repair protein O6-alkylguanine-DNA alkyl transferase which can lead to an increase in the rate of human hepatocellular carcinoma (HCC) (Tang *et al.*, 2012; Wei *et al.*, 2010, 2011). Pharmacologic inhibition of inducible NOS (iNOS) when GSNOR is down-regulated shows strong potential as a therapeutic for those patients with HCC (Tang *et al.*, 2013a). As with HCC, some types of breast cancer are linked to a decrease in GSNOR expression. Specifically, high levels of human epidermal growth factor receptor 2 (HER2)

expression in breast tumors is associated with low GSNOR expression and an increase in apoptosis-related protein S-nitrosation (Cañas *et al.*, 2016). This study also determined that an increase in GSNOR expression in *HER2* tumors correlates with higher patient survival and begs the question as to whether or not NOS inhibition would also serve this population well. These examples are of course complicated by the fact that •NO is a pleiotropic regulator of gene function and the modulation of GSNO by GSNOR can have both cytostatic and cytotoxic effects on tumor survival (Xu *et al.*, 2002). To this point, GSNOR is effective at removing formaldehyde, a known carcinogen, from the cell; however, *ADH5* polymorphisms do not significantly affect an individual's capacity to protect against DNA damage when exposed to formaldehyde (Xie *et al.*, 2010). Furthermore, *ADH5*^{-/-} mice are known to generate DNA damage when formaldehyde forms and adduct with guanine to create N2-hydroxymethyl-dG which can result dysfunction of hepatocytes and nephrons (Pontel *et al.*, 2015).

Asthma and single nucleotide polymorphisms

SNPs can alter the transcriptional output of a gene as well as the structure/function of proteins they encode. Several SNPs in the promoter and 3' UTR of the *ADH5* gene can result in the aberrant expression of GSNOR (Choudhry *et al.*, 2010). Of particular interest is the observation that airway hyperresponsivity in wild-type mice correlates with increased expression of GSNOR and decreased RSNO production, while *ADH5*^{-/-} mice are protected from airway hyperresponsiveness and maintain higher total RSNO levels (Que *et al.*, 2005). In humans, GSNOR upregulation can lead to changes in airway smooth muscle tone in asthmatics (Henderson & Gaston, 2005; Wu *et al.*, 2007). A study involving Mexican children with asthma who possess SNPs in the promoter region of *ADH5* at suspected NF-κB binding sites (rs2602899 and rs2851301), were found to exhibit a decreased relative risk of asthma due to suppressed GSNOR production (Wu *et al.*, 2007). Interestingly, alternative SNPs (rs1154404 and rs28730619) were associated with an increase in childhood asthma risk, although the mechanism behind this correlation has not been determined (Wu *et al.*, 2007). Another study in African American children found that SNPs in *ADH5* and the β2 adrenergic receptor gene are associated with acute response to asthma-specific therapy (Moore *et al.*, 2009).

Looking beyond GSNO-mediated relaxation of airway smooth muscle we may also consider GSNOR's ability to

metabolize formaldehyde, a chemical known to induce bronchoconstriction after long term exposure at low concentrations (Leikauf, 1992). It has been suggested that the presence of formaldehyde in airway smooth muscle may stoichiometrically favor bound NADH/GSNOR, thereby increasing GSNOR metabolism of GSNO, and by extension, promote smooth muscle contraction (Thompson & Grafström, 2007).

Regardless of the mechanism driving GSNOR-mediated consumption of GSNO in airway smooth muscle, it is easy to see why the inhibition of GSNOR has been of particular interest to researchers for its therapeutic potential as a smooth muscle relaxant.

Myoendothelial junctions

GSNOR plays an interesting role at myoendothelial junctions (MEJ) where it co-localizes with the hemichannel Connexin-43 (Cx43). Cx43 hemichannels form gap junctions between cells by linking to hemichannels in opposing membranes to couple endothelial and vascular smooth muscle cells and when Cx43 is S-nitrosated, this pore allows for the free movement of inositol triphosphate from vascular smooth muscle to endothelial cells. Due to the co-localization of GSNOR and Cx43 at the MEJ, basal $\cdot\text{NO}$ availability at this site is blunted, which in turn increases the likelihood that Cx43 will not be S-nitrosated (Straub *et al.*, 2011). This decreases channel permeability until Ca^{2+} levels increase as a result of smooth muscle cell stimulation, which in turn activates eNOS and increases the probability of Cx43 S-nitrosation.

Myometrium

$\cdot\text{NO}$ is an important mediator of relaxation in the myometrium. It has been well established that $\cdot\text{NO}$ relaxes vascular and gastrointestinal smooth muscle by activating soluble guanylyl cyclase (sGC), which in turn converts guanosine triphosphate to cyclic guanosine monophosphate (cGMP), activating PKG, which in turn dephosphorylates the regulatory light chain (MYL9) of myosin *via* the amplified phosphatase activity of MYPT1 (pS695) (Grassie *et al.*, 2011; Nakamura *et al.*, 2007; Puetz *et al.*, 2009; Roux *et al.*, 2012). This is not the dominant $\cdot\text{NO}$ -mediated relaxation pathway in uterine smooth muscle, however, $\cdot\text{NO}$ can relax the myometrium even when sGC has been inhibited (Buxton *et al.*, 2010). The pathway through which $\cdot\text{NO}$ relaxes the myometrium independent of cGMP is unknown, but it is likely that the S-nitrosation of contractile proteins plays a role. It has been determined that the state of labor (full

term versus preterm) can vastly alter the S-nitrosated protein landscape in uterine smooth muscle after exposure to GSNO (Ulrich *et al.*, 2012). It is also well known that S-nitrosation can vary significantly based upon the cytoplasmic availability GSNOR (Broniowska & Hogg, 2012; Hess *et al.*, 2005; Thomas & Jourd'heuil, 2012). Regardless of the pathway through which $\cdot\text{NO}$ acts to relax uterine smooth muscle, it does beg the question as to whether or not inhibiting GSNOR, and thereby increasing intracellular availability of GSNO, may serve as an effective tocolytic strategy by promoting uterine quiescence through $\cdot\text{NO}$ -mediated relaxation pathways. This notion is supported by data showing an increased expression of GSNOR in patients delivering spontaneously preterm.

Therapeutic inhibition of GSNOR

GSNOR is an attractive therapeutic target. GSNOR inhibition increases GSNO availability in the cell and in turn facilitates $\cdot\text{NO}$ -mediated signaling pathways. Dozens of small molecules have been identified that can inhibit GSNOR to varying degrees (Green *et al.*, 2012; Jiang *et al.*, 2016; Sanghani *et al.*, 2009; Sun *et al.*, 2011a, 2011b, 2012). Two of these, N6022 (3-(5-(4-(1H-imidazol-1-yl) phenyl)-1-(4-carbamoyl-2-methylphenyl)-1H-pyrrol-2-yl) propionic acid) and N91115 from Nivalis Pharmaceuticals show promise as potentially safe and effective GSNOR inhibitors that have undergone clinical trial for both the treatment of mild asthma (clinicaltrials.gov – NCT01316315), and cystic fibrosis in individuals who are heterozygous for the cystic fibrosis transmembrane conductance regulator (CFTR) gating mutation CFTR Δ F508 + (clinicaltrials.gov – N6022: NCT01746784; N91115: NCT02724527). Endogenous GSNO levels are low in the airways of cystic fibrosis patients (Grasemann *et al.*, 1999) and GSNOR inhibition is an appealing alternative to the direct administration of GSNO (Snyder *et al.*, 2002; Zaman *et al.*, 2001, 2013). N6022 is well tolerated with minimal side effects, even at high concentrations, in both animals (Blonder *et al.*, 2014; Colagiovanni *et al.*, 2012) and humans (clinicaltrials.gov – NCT01147406, NCT01746784). Another GSNOR inhibitor, SPL-334 (4-[[2-[(2-cyanobenzyl) thio]-4-oxothieno[3,2-d]pyrimidin-3(4H)-yl]methyl]benzoic acid) from SAJE Pharmaceuticals (Baltimore, MD), is being tested as a therapeutic to treat allergic asthma and interstitial lung disease (ILD). Using an allergic asthma mouse model, intranasally administered SPL-334 decreased CD4⁺ Th2 cytokines, eosinophils, and mitigated the lung inflammatory response (Ferrini *et al.*, 2013). Likewise, in a mouse model of ILD, SPL-334 functions as both a prophylactic agent and a therapeutic to

attenuate profibrotic cytokines and collagen accumulation in the lungs (Luzina *et al.*, 2015). Unlike N6022 and N91115, SPL-334 is not in human clinical trials.

FDA-approved drugs are also being tested as potential GSNOR inhibitors. Nebivolol, a β_1 -adrenergic receptor antagonist used for the management of hypertension, has been shown to increase total RSNO levels in animal and cell models (Jiang *et al.*, 2016). Our own investigation of GSNOR fails to confirm Nebivolol as an inhibitor of GSNOR in an enzyme activity assay. Since there are no FDA-approved GSNOR inhibitors, the repurposing of existing therapeutic agents that inhibit GSNOR and/or modulate GSNO and RSNOs is of interest.

When considering GSNOR inhibitors as therapeutic agents, it should be taken into consideration that enzymes other than GSNOR modulate •NO availability in the cell. •NO is critical to the normal function of most cell types, and as is often the case, there are multiple concurrent and complementary mechanisms to regulate •NO and RSNOs (Benhar *et al.*, 2009; Liu *et al.*, 2001). Two of the most well-known are thioredoxin-1 (Sengupta & Holmgren, 2012a, 2012b) and carbonyl reductase (Bateman *et al.*, 2008). NOS, the predominate source of •NO in the body, can also be dysregulated in certain disease states, as can its substrate, L-arginine (Ckless *et al.*, 2007). For instance, after stimulation of the cavernous nerve in *ADH5*^{-/-} mice, eNOS phosphorylation did not increase as predicted (Musicki *et al.*, 2016). Modulating GSNOR activity may insufficiently control, or even aggravate some conditions if these alternate •NO-regulators are the source of the disorder. Unfortunately, direct application of endogenous •NO-donors, such as GSNO, Cys-NO or SNO-albumin, as well as some exogenous donors, are of limited clinical value because they either degrade rapidly, cause intolerable side effects, or lead to a toxic systemic build up nitrates (Al-Sa'doni, 2005).

The therapeutic inhibition of GSNOR to treat •NO-mediated disorders should be weighed carefully against potential contraindications. For example, the inhibition of GSNOR may increase a patient's susceptibility to bacterial or viral infection. The inhibition of GSNOR will also increase total RSNO levels and this can have adverse effects in the body, especially if the drug is administered systemically and not targeted to a specific tissue type through means such as liposomal delivery. GSNOR regulation varies widely in different cancer types (Cañas *et al.*, 2016; Tang *et al.*, 2013a). Inhibiting GSNOR may lead to a further increase in GSNO at the tumor site which can favor angiogenesis (Prudente *et al.*, 2017). Conversely, with disorders such as asthma and hypertension, GSNOR inhibition results in the desired relaxation of the smooth muscle.

Conclusions

•NO, and by extension •NO-donors, have been investigated intensely for over a century as therapeutics (Schmidt & Walter, 1994). •NO modulation not only affects traditional pathways connected to this highly reactive molecule, but it also drastically alters S-nitrosation levels in the cell. GSNOR is unique among the ADH family of enzymes in that it targets GSNO and varies the body's response to endogenously generated •NO carried as GSNO. *ADH5*^{-/-} animal and cell models have provided a unique window into the importance of GSNOR in nearly every tissue type. The up/down regulation of GSNOR in humans has also provided invaluable data to the medical and research communities concerning its role in disease states. There are currently no FDA-approved modulators of GSNOR; however, several drugs are being investigated, and some are in clinical trial. Indeed, our understanding of the dysregulation of GSNOR and its effect on protein S-nitrosation and other glutathione/•NO-mediated events is in its infancy. Further investigations into the role of GSNOR in health and disease are needed to reveal the most effective therapeutic options.

Disclosure statement

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