

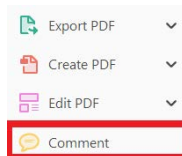
USING e-ANNOTATION TOOLS FOR ELECTRONIC PROOF CORRECTION

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This will open up a ribbon panel at the top of the document. Using a tool will place a comment in the right-hand panel. The tools you will use for annotating your proof are shown below:



1. Replace (Ins) Tool – for replacing text.

Strikes a line through text and opens up a text box where replacement text can be entered.

How to use it:

- Highlight a word or sentence.
- Click on .
- Type the replacement text into the blue box that appears.

...n of nutritional conditions, and landmark events are monitored in populations of relatively homogeneous single n of *Saccharomyces*, and is initiated after carbon source [1]. S are referred to as mei n of meiosis-specific g *revisiae* depends on th inducer of meiosis) [3 I functions as a repre repression, the genes *pression*) and *RGR1* at ase II mediator subur osome density [4]. *SIM* irectly or indirectly re

jstaddon Reply X
05/05/2017 15:32 Post

2. Strikethrough (Del) Tool – for deleting text.

Strikes a red line through text that is to be deleted.

How to use it:

- Highlight a word or sentence.
- Click on .
- The text will be struck out in red.

... experimental data if available. For ORFs to be had to meet all of the following criteria:

1. Small size (35–250 amino acids).
2. Absence of similarity to known proteins.
3. Absence of functional data which could n the real overlapping gene.
4. Greater than 25% overlap at the N-termin terminus with another coding feature; ove both ends; or ORF containing a tRNA.

3. Commenting Tool – for highlighting a section to be changed to bold or italic or for general comments.

Use these 2 tools to highlight the text where a comment is then made.

How to use it:

- Click on .
- Click and drag over the text you need to highlight for the comment you will add.
- Click on .
- Click close to the text you just highlighted.
- Type any instructions regarding the text to be altered into the box that appears.

...nformal invariance: or A: Math. Gen., Vol. 12, N

...lified theory for a matrix. 'ol. 8, 1984, pp. 305–323 :d manuscript, 1984. ching fractions for $D0 \rightarrow K+K$ relation in $D0$ decays' Phys

jstaddon Reply X
This needs to be bold
05/05/2017 15:40 Post

4. Insert Tool – for inserting missing text at specific points in the text.

Marks an insertion point in the text and opens up a text box where comments can be entered.


How to use it:

- Click on .
- Click at the point in the proof where the comment should be inserted.
- Type the comment into the box that appears.


... Meiosis has a central role in the sexual reproduction of nearly all eukaryotes. *Saccharom* analysis of meiosis, esp by a simple change of n conveniently monitored cells. Sporulation of *Sae* cell, the a/α cell, and is of a fermentable carbon sporulation and are refe [2b]. Transcription of me meiosis, in *S. cerevisiae* activator, *IME1* (inducer of the gene *RME1* funct Rme1p to exert repress of *GAL1* gene expression) and *HGR1* are required [1, 2, 3, 4]. These ge

jstaddon Reply X
Yeast.
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5. Attach File Tool – for inserting large amounts of text or replacement figures.

 Inserts an icon linking to the attached file in the appropriate place in the text.


How to use it:

- Click on  .
- Click on the proof to where you'd like the attached file to be linked.
- Select the file to be attached from your computer or network.
- Select the colour and type of icon that will appear in the proof. Click OK.


The attachment appears in the right-hand panel.

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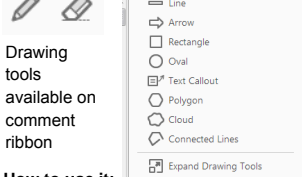
6. Add stamp Tool – for approving a proof if no corrections are required.

 Inserts a selected stamp onto an appropriate place in the proof.

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- Click on  .
- Select the stamp you want to use. (The **Approved** stamp is usually available directly in the menu that appears. Others are shown under *Dynamic, Sign Here, Standard Business*).
- Fill in any details and then click on the proof where you'd like the stamp to appear. (Where a proof is to be approved as it is, this would normally be on the first page).

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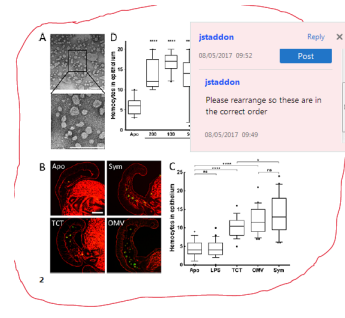


7. Drawing Markups Tools – for drawing shapes, lines, and freeform annotations on proofs and commenting on these marks.

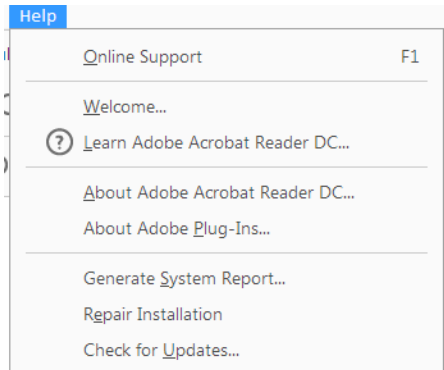
Allows shapes, lines, and freeform annotations to be drawn on proofs and for comments to be made on these marks.

How to use it:

- Click on one of the shapes in the **Drawing Markups** section.
- Click on the proof at the relevant point and draw the selected shape with the cursor.
- To add a comment to the drawn shape, right-click on shape and select **Open Pop-up Note**.
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15	AUTHOR: Please provide the page range for reference Hill (2012).	
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Viewpoint

Recommendations on terminology and experimental best practice associated with plant nitric oxide research

Summary

Nitric oxide (NO) emerged as a key signal molecule in plants. During the last two decades impressive progress has been made in plant NO research. This small, redox-active molecule is now known to play an important role in plant immunity, stress responses, environmental interactions, plant growth and development. To more accurately and robustly establish the full spectrum of NO bioactivity in plants, it will be essential to apply methodological best practice. In addition, there are some instances of conflicting nomenclature within the field, which would benefit from standardization. In this context, we attempt to provide some helpful guidance for best practice associated with NO research and also suggestions for the cognate terminology.

Recommendations on terminology

S-Nitrosylation or S-nitrosation?

The reversible, covalent modification of cysteine thiols by nitric oxide (NO) is termed S-nitrosylation. According to the most extensive comprehensive dataset to date, the Arabidopsis proteome contains 1195 endogenously S-nitrosylated peptides belonging to 926 proteins (Hu *et al.*, 2015), which implies the important biological relevance of this post-translational modification (PTM). There has been recent debate in the field over whether S-nitrosylation is an appropriate term, as this name implies an enzymatic function. Nitrosylation involves direct addition of NO to a reactant and is derived from chemistry terminology that describes the coordination of NO to a metal centre leading to formation of a metal nitrosyl complex (Ford *et al.*, 2005). These metal nitrosyls can also be formed by other chemical reactions. For instance, a transition metal can react with acidified nitrite via a multistep reaction also leading to the formation of a metal nitrosyl complex (Ford, 2010). Hence, the more chemically orientated term, S-nitrosation, has been

proposed as an alternative expression to that of S-nitrosylation (Heinrich *et al.*, 2013), where the addition of a nitrosonium ion (NO⁺) to a nucleophilic group takes place. In the context of proteins, transfer of an NO⁺ molecule is a predominant mechanism for oxidation of protein cysteine (Cys) thiols, although the formation and subsequent role of this molecule in S-nitrosothiol (SNO) formation also depends on the cellular conditions and the chemical environment surrounding the target Cys embedded within the given protein, respectively. In this case, we suggest S-nitrosation is therefore a more applicable expression for this chemical process.

In mammals, a handful of proteins have been identified, for example glyceraldehyde-3-phosphate dehydrogenase (GAPDH), that interact with specific protein targets and transfer their NO moiety, resulting in SNO formation at a specific Cys residue on the target protein (Kornberg *et al.*, 2010). The proteins driving this PTM can therefore be considered as nitrosylases (Seth *et al.*, 2018). In this case, the term *trans*-nitrosylation might be more appropriate.

Further, in both mammals and plants, thioredoxin (Trx) enzymes have been shown to directly and selectively remove a SNO from target proteins functioning as de-nitrosylases (Wu *et al.*, 2011; Kneeshaw *et al.*, 2014). Clearly, this process is mediated by enzyme activity, thus de-nitrosylation rather than de-nitrosation appears a more appropriate term in this context.

Nitrosative stress or nitro-oxidative stress?

The term nitrosative stress refers to a secondary stress condition characterized by a parallel, unregulated increase in the generation of both NO and reactive oxygen species (ROS) (Valderrama *et al.*, 2007). Further, these redox active molecules can react with each other to form additional molecules, such as peroxynitrite (ONOO⁻), formed by the interaction of NO and superoxide (O₂⁻). Collectively, these molecules can trigger irreversible damage to different biomolecules, such as proteins, lipids and nucleic acids.

In a similar manner to that of protein carbonylation, which is considered a major hallmark of oxidative stress (Fedorova *et al.*, 2014), an increase in protein tyrosine nitration has been proposed as a plausible marker for nitrosative stress (Corpas *et al.*, 2007). Tyrosine nitration involves an oxidative and a nitrative step and is directly driven by radicals derived from peroxynitrite. Additionally, different antioxidant enzymes, such as catalase, ascorbate peroxidase, monodehydroascorbate reductase and superoxide dismutases, were found to be negatively affected by nitration, further supporting a close relationship between NO and ROS, especially under stress conditions.

Thus, nitro-oxidative stress may be considered a suitable expression to describe cellular events resulting from detrimental

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1 accumulation of and interaction between ROS and reactive
2 nitrogen species (RNS).

3 Nonsymbiotic hemoglobins or phytoglobins?

4 Leghemoglobins (Lb) and symbiotic hemoglobins (sym-Hb) are
5 present in either legume species or actinorhizal/nonlegume sym-
6 biotic nodules, respectively. The presence of hemoglobin-like
7 proteins in organs not associated with symbiotic interactions led to
8 the term, nonsymbiotic hemoglobins (Hill, 2012; Rubio *et al.*,
9 2019). These nonsymbiotic hemoglobins are thought to function
10 as key scavengers of NO under various environmental and stress
11 conditions (Hill, 2012). Therefore, these proteins may play an
12 important role in NO homeostasis within various organs and are
13 also involved in the hemoglobin–NO cycle, which increases energy
14 efficiency under hypoxia, by oxidizing NAD(P)H to enhance
15 proton pumping and concomitant ATP production. At the 2014
16 XVIII Conference on Oxygen-Binding and Sensing Proteins,
17 several prominent research groups focusing on heme proteins
18 reached a consensus to rename these proteins as ‘Phytoglobins’
19 (Hill *et al.*, 2016). In this context, *phyto* means plant (including
20 algae and land plants) and *globin* refers to a heme-containing
21 protein fold similar to the myoglobin structure of the sperm whale,
22 where heme-Fe is invariably coordinated at the proximal site by His
23 F8. Hence, in agreement with Hill *et al.* (2016), we recommend the
24 term ‘Phytoglobin’ when referring to nonsymbiotic hemoglobins
25 in plants. This terminology applies to hexacoordinated, nonsym-
26 biotic hemoglobin 1/class 1 (Phytogb1), pentacoordinated non-
27 symbiotic hemoglobins 2/class 2 (Phytogb2) and penta/
28 hexacoordinated, nonsymbiotic hemoglobin 3/truncated
29 hemoglobins (Phytogb3) (Hill, 2012).

30 It has been common practice to describe the reaction between
31 oxyhemoglobin and NO as a ‘dioxygenase’ reaction or ‘dioxyge-
32 nase’ activity, including Phytogb1 and Phytogb2, ascribing enzyme
33 function to hemoglobin. There is no evidence of any hemoglobin
34 acting as an enzyme, except for flavohemoglobin (Gardner
35 *et al.*, 1998) which is a bifunctional protein with true enzyme
36 activity.

37 Nomenclature of nitric oxide synthase-like activity in plants

38 Nitric oxide synthase (NOS) is the main enzymatic source for NO
39 in metazoans. This enzyme catalyses the production of both NO
40 and L-citrulline from L-arginine using two co-substrates (NADPH,
41 oxygen) and several cofactors including two flavins (FMN, FAD),
42 calmodulin and a pterin (tetrahydro-L-biopterin). In contrast to
43 several algal species, land plants do not possess a typical NOS
44 (Jeandroz *et al.*, 2016). However, several lines of evidence suggest
45 that activity resembling that of an NOS is present in land plants
46 (Reviewed in Corpas *et al.*, 2009) and the identification of the
47 protein(s) catalysing this activity is a major goal (Del Castello *et al.*,
48 2019).

49 In aggregate, we therefore suggest employing the terms NOS-
50 like activity or NOS-like enzyme when referring to this enzymatic
51 process and to the corresponding unidentified enzyme(s). We
52 advocate that these terms can be utilized to describe L-Arg-

dependent activities, as the NOS measured in land plants requires
L-arginine, NADPH, calcium and calmodulin, also essential
prerequisites of mammalian NOS enzymes. It has been suggested
that this plant activity could be a result of cooperation between
separate proteins, which, when combined, biochemically resemble
the NADPH:oxygen oxidoreductases of animal NOSs (Corpas &
Barroso, 2017).

53 Suggested best practice for NO detection methods

54 Due to the rapid chemical reactions exhibited by the NO free
55 radical with a wide range of biological targets, the detection and
56 quantification of this molecule in plant samples is routinely
difficult. Further, the current methods deployed differ in terms of
both their selectivity and specificity (Vishwakarma *et al.*, 2019).
Unfortunately, to date, there is no entirely satisfactory method for
the quantification of NO. Each of the current methods has specific
limitations. However, by employing best practice, reliable results
can be obtained, enabling successful interpretation of NO function
(Gupta & Igamberdiev, 2013). Thus, the available methods of NO
measurement performed carefully can typically provide accurate
and robust results *in vitro*, using NO or chemical compounds
including NO donors. However, there are significant limitations
when these approaches are applied to complex biological matrices
without proper validation and control assays.

Therefore, the given technique, whether direct or indirect,
should be selected with caution, given that all the current methods
have both advantages and disadvantages. Those relevant to plant
samples include colorimetric assays (based on Griess reactions and
oxyhemoglobin), fluorimetric assays using different diaminofluo-
rescein (DAF) dyes, photo-acoustic laser detection, electron
paramagnetic resonance (EPR) spectroscopy with an NO-specific
spin trap and ozone chemiluminescence detection utilizing an NO-
specific electrode (Mur *et al.*, 2011). Due to its NO-specificity, spin
trap EPR is considered one of the most specific methods for
demonstrating the unequivocal presence of NO. In plants, this
method has been useful in detecting the presence of NO in both
plant extracts and purified organelles (Maskall *et al.*, 1977; Caro &
Puntarulo, 1999; Corpas *et al.*, 2004; Jasid *et al.*, 2006).

Chemiluminescence is best suited for measurement of emitted
NO but in order to measure oxidized forms of this molecule, one
has to employ indirect chemiluminescence, where samples should
be injected with solutions such as vanadium(III) chloride (VCl₃) to
reduce oxidized forms of NO. Recently, genetically encoded NO
probes have been described (Eroglu *et al.*, 2016; Calvo-Begueria
et al., 2018). Such proteins are an optimal tool for NO detection/
quantification *in vivo* and have the potential to revolutionize the
field of plant NO research. Calvo-Begueria and colleagues
monitored NO production via formation of a nitrosyl–leghe-
moglobin complex (Lb2 + NO), which can be detected by EPR
spectroscopy. Further, Eroglu *et al.* (2016) fused a bacteria-derived
NO-binding domain adjacent to different fluorescent protein
variants, enabling both direct observation and quantification of
NO. These genetically encoded NO probes provide a specific real-
time, read-out of cellular NO dynamics and, hence, potentially
open a new era for NO bioimaging.

1 In addition to these methods, the application of microelectrodes
2 is also an effective approach, for example, in tracking NO in the
3 extracellular media of cell suspensions. These electrodes consist of a
4 platinum/iridium (Pt/Ir) wire sealed in a glass or plastic capillary in
5 which a thin film of nickel phthalocyanine is electrodeposited. An
6 outermost layer constituted from Nafion and *o*-phenylenediamine
7 increased the selectivity of the electrodes against possible interfering
8 molecules (Griveau *et al.*, 2016). This method is considered as one
9 of the most specific methods in animals and has also been
10 successfully used to measure NO production in plant cell
11 suspensions (Besson-Bard *et al.*, 2008). With appropriate controls:
12 NO deficient mutants such as *nia1 nia2*, *atnoa1* or NO scavenging
13 lines, including those overexpressing *Pgb1* or *S*-nitrosoglutathione
14 reductase (*GSNOR*) (Yun *et al.*, 2011), one can accurately
15 determine the endogenous NO level. Also, the application of
16 NO donors or scavengers as controls, can function as key controls in
17 the determination of NO.

18 DAF-based dyes

19
20
21 Fluorescence-based methods for the detection of NO and other
22 RNS are commonly utilized (Mur *et al.*, 2011). The technique
23 relies on the presence of a nonfluorescent probe which can be
24 located to the source of NO and subsequently becomes fluorescent
25 on reaction with this molecule or a related RNS. Therefore, a major
26 advantage to this approach is that it can provide spatial information
27 regarding the accumulation of the specific RNS under study.

28 There are however, numerous potential problems with this
29 approach and often such issues are not considered. The technique
30 relies on the measurement of fluorescent light, which does not
31 readily lend itself to quantitation and is thus usually reported as
32 pixel intensity, not molarity. Therefore, measured light relies on
33 efficient penetration of the excitation light and efficient release of
34 emitted light, both which can be problematic within deeper
35 samples. This approach can also be affected by autofluorescence of
36 the sample, relatively common in plant material and is also prone to
37 photobleaching. It also noteworthy that RNS accumulation is
38 typically not static, but can be repositioned, as reported for
39 peroxynitrite (ONOO⁻) moving through membranes (Denicola
40 *et al.*, 1998). Created fluorescent RNS-adducts can also move, so
41 spatial data is not always reliable.

42 However, one of the biggest concerns regarding DAF-based dyes
43 is specificity and selectivity. Fluorescent probes rely on redox
44 chemistry and it would not be unusual for such probes to be
45 oxidized by a range of endogenous redox active molecules. For
46 example, 2,7-dichlorodihydrofluorescein (DCFH) oxidation
47 yields 2,7-dichlorofluorescein (DCF), a reaction which can be
48 driven by the presence of RNS but also ROS. Indeed, DCFH is
49 commonly utilized to detect ROS. Issues and limitations of the
50 fluorescent-based approach to RNS measurements have been the
51 subject of several reviews (Kalyanaraman *et al.*, 2012; Li & Wan,
52 2015).

53 However, despite these potential limitations, the application of
54 these reporter dyes can often be effective. The most commonly used
55 probes are based on DAF: that is DAF-2DA and DAF-FM DA,
56 both readily commercially available. On entering the cell,

intracellular esterases cleave these dyes to DAF-2D or DAF-DM
and subsequently, react with RNS leading to the formation of a
nitrosated form, DAF-2 triazole (DAT-2T), which is fluorescent.
Importantly, however, DAF does not react directly with NO. It
reacts with oxidized forms such as NO⁺ or N₂O₃. Although DAF is
relatively specific to RNS, DAF-2DA can also react with ROS
(Balcerczyk *et al.*, 2005). Hence, it is essential to check the
specificity of fluorescence employing an NO scavenger, to confirm
the detected fluorescence is resulting from NO accumulation, as
ROS and NO are often produced in parallel. An ideal NO
scavenger to utilize in this case is 2-(4-carboxyphenyl)-4,4,5,5-
tetramethylimidazoline-1-oxyl-3-oxide potassium salt (cPTIO, see
later). It should be also considered that DAF-2 itself is weakly
fluorescent, so in some instances the observed increased fluores-
cence can result from its accumulation inside cells and not by its
reaction with RNS. This potential scenario can be checked by using
4-aminofluorescein DA (which is converted to 4-AF), as a negative
control, which cannot react with NO (Beligni *et al.*, 2002).

A variation of this approach can also be employed to measure
exogenous or released RNS. DAF-2DA is not fluorescent, but if the
de-esterified version is used it is unable to penetrate cells but can
react with RNS and become fluorescent and so extracellular RNS
can be estimated. Such measurements can then be corroborated by
more difficult approaches, including EPR.

As well as DAF-based probes, there are other fluorescent dyes
available for RNS measurements, such as the copper(II) fluorescein
(CuFL) complex (Lim *et al.*, 2006), the diamino-rhodamine-4M
probes (DAR-4M, Kojima *et al.*, 2001), or the Pyrene-Based
Fluorescent Nitric Oxide Cheletropic Traps (FNOCTs, D ppe
et al., 2010). CuFL complex has the advantage of reacting directly
with NO itself rather than a derivative RNS and is an interesting
alternative to that of DAF. For the measurement of peroxynitrite,
dihydrorhodamine 123 (DHR) can be employed as it yields the
fluorescent compound rhodamine 123 (RH) on oxidation. There
are numerous other fluorescent probes, such as those based on
aromatic boronates (Kalyanaraman *et al.*, 2012). The application
of one of these dyes for RNS detection as the sole method is not
recommended and other techniques should be employed in parallel
to ensure robustness of RNS data (Gupta and Igamberdiev 2011).

NO scavenger controls

cPTIO is a widely used NO scavenger to confirm any observed
DAF fluorescence is attributed to NO. cPTIO oxidizes the NO
molecule leading to formation of the NO₂ radical (NO + cPTIO
→ NO₂ + cPTI). The produced NO₂ radical can react with NO to
form N₂O₃ (NO₂ + NO → N₂O₃) with which DAF-2 reacts
leading to formation of fluorescent DAF-2T (Table 1). This
implies, to some extent, cPTIO has the capability to increase
fluorescence rather than masking it, if used at a low concentration.
At a high concentration, cPTIO reacts rapidly with NO and
oxidizes NO to NO₂ + PTI. Hence, a concentration of > 200 μM
of cPTIO is recommended (Vitecek *et al.*, 2008). Nevertheless,
cPTIO is also known to quench DAF-2T fluorescence due to its
intense blue colour (Arita *et al.*, 2006). Thus, the optimization of
cPTIO concentrations for any given experimental setting is highly

advised (Goldstein *et al.*, 2003; D'Alessandro *et al.*, 2013). It is noteworthy that the reaction product, cPTI, has been reported to possess biological activity without NO scavenging, both in animal and plant models (Cao & Reith, 2002; Planchet *et al.*, 2006), pointing to cautious interpretation of data obtained using DAF-based dyes and PTIO compounds. cPTIO can also be used as a spin trap in EPR to detect NO; however, this approach has also been shown to have considerable limitations (D'Alessandro *et al.*, 2013). Recently, it has been demonstrated that commercially available hemoglobins can be used as a control instead of cPTIO for attributing the fluorescence based MnIP-Cu probes specificity to NO (Singh & Bhatla, 2019).

Collectively, therefore careful consideration should be given before embarking on experiments employing NO scavengers.

Application of NO donors

Treatment of plants with gaseous NO requires special equipment and special care to prevent gas leakage, so the application of NO releasing chemicals (NO donors) provides a more easily executable way of NO treatment. Therefore, supplying plants with different NO donors is a common practice to mimic NO production and potentially rescue NO deficient phenotypes. Different NO donors have different kinetics, mechanisms and environmental conditions for optimal NO release, thus some care should be taken during the choice of NO donor.

In plant research, the most commonly deployed NO donors are sodium nitroprusside (SNP), *S*-nitrosopenicillamine (SNAP), *S*-nitrosoglutathione (GSNO) and diethylamine NONOate (DETA/NO). One should note that these donors differ in the form of NO release. For instance, SNP releases nitrosonium cation (NO⁺) whereas SNAP and GSNO typically release NO in the form of a radical ([•]NO), but under certain environmental conditions these NO donors can also release the nitrosonium cation (NO⁺). Accumulating evidence suggest that the form of NO emitted by various donors plays a key role in switching on appropriate metabolic modifications (Arasimowicz-Jelonek *et al.*, 2011).

Another relevant difference between donors is the kinetics of NO release. In aqueous solution, DETA/NO and SNAP produce transient NO bursts (seconds to minutes), while the NO-releasing effect of SNP is more extended (Floryszak-Wieczorek *et al.*, 2006; Planchet & Kaiser, 2006; Mur *et al.*, 2013). In a similar fashion, GSNO also delivers NO over a longer time period, typically several hours (Floryszak-Wieczorek *et al.*, 2006; Mur *et al.*, 2013). SNP is among the most widely studied NO donors, which is justified by its capability of producing persistent NO (Mur *et al.*, 2013) and by its cost-efficiency. However, the application of SNP has several drawbacks. Firstly, the release of NO requires light and the illumination may influence plant samples in an unwanted way. Secondly, NO release from SNP is associated with the production of toxic gases like hydrogen cyanide (HCN) (Table 1) (Bethke *et al.*, 2006). This compound can inhibit molybdenum-based enzymes and also can inhibit cytochrome c oxidase leading to inhibition of respiration. However, Shishido & de Oliveira (2001) reported that SNP releases CN following ultraviolet (UV) radiation, while illumination with longer wavelengths results in

selective NO release from SNP, which supports the utility of this NO donor under typical light conditions. Reflecting these differences between NO donors, these compounds may have different biological effects: SNP induced the accumulation of *Ferritin* transcripts, while SNAP inhibited the expression of this gene. Similarly, SNP induced cell death and inhibited antioxidant gene expression but other NO donors showed opposing effects (Murgia *et al.*, 2004).

Furthermore, the NO releasing capacity of various donors depends on cellular redox and antioxidant status (Floryszak-Wieczorek *et al.*, 2006) and also on the actual concentration of the NO donor applied (Ederli *et al.*, 2009). Development of precise NO releasing compounds is an important task for future NO research. Until this has been accomplished, it is recommended to test different concentrations of NO donors to determine their NO releasing effects *in vitro* and *in vivo* under the same experimental conditions. Appropriate control treatments (e.g. potassium cyanide, reduced glutathione) are also needed in order to support the NO releasing capacity of the different NO donors. The application of an NO scavenger (e.g. cPTIO) together with an NO donor can provide useful information regarding the NO releasing character of the donor.

Another experimental option is subjecting the biological system under study to NO gas (Palma *et al.*, 2018). Currently, the NO scientific community is searching for 'elicitors' which promote endogenous NO release, enabling more physiological responses. The development of either genetically encoded or chemically based, organelle specific NO reporters, would be also an important future advance.

Enzyme inhibitors of NO metabolism

In the context of the pharmacological approach, several types of compounds have been employed to study the involvement of specific plant enzyme(s) in NO production or signalling pathways. This extended practice might result in hard to interpret data, due to known or unknown unspecific effect of these compounds to other plant proteins or enzymes and partly also due to their application in relatively high concentrations, often required to achieve any observable effects.

A good example of this is tungstate, which can inhibit nitrate reductase (NR) activity through molybdenum displacement and has been used to confirm involvement of NR in observed NO production (Chamizo-Ampudia *et al.*, 2017). However, tungstate is known to interfere with other molybdenum-containing enzymes and also plant developmental processes (Xiong *et al.*, 2012). As tungstate is known to affect both plant NR activity and gene expression (Deng *et al.*, 1989), experiments using tungstate to test NR-dependent NO production should also involve determination of NR activity.

A high number of plant studies have employed chemical substances developed as effective inhibitors of well-characterized animal NOS isoforms. This practice, based on diverse L-arginine derivatives such as (N^G-monomethyl-L-arginine, L-NMMA and N^ω-nitro-L-arginine, L-NAME), has been a subject of long-term criticism (Planchet & Kaiser, 2006). This is mainly for two reasons:

Table 1. Overview of commonly used nitric oxide (NO) donors, reactants, detection reagents and their reactions.

Compound	Chemical name	Reaction(s)	Comment
NO donors			
Nitrite	NO ₂ ⁻	NO ₂ ⁻ + e ⁻ + 2H ⁺ → NO + H ₂ O	Rate of NO release is highly pH-dependent
SNP	Sodium nitroprusside	Na ₂ [Fe(CN) ₅ NO] ²⁻ → NO [·] + CN ⁻ + Na ₂ [Fe(CN) ₄] ⁻ Na ₂ [Fe(CN) ₄] ⁻ Na ₂ [Fe(CN) ₅ NO] ²⁻ → NO ⁺ + CN ⁻ + Na ₂ [Fe(CN) ₄] ⁻	Light-dependent reaction, SNP can also release nitrosonium and free iron
GSNO	S-Nitrosoglutathione	2 GSNO → 2 NO [·] + GSSG (decomposition)GSNO + RSH → RSNO + GSH (transnitrosation)	Reaction catalysed by light, heat and metal ions (Cu ²⁺ , Hg ²⁺), under certain conditions also nitrosonium (NO ⁺) can be formed
SNAP (and other S-nitrosothiols)	S-Nitroso-N-acetyl-DL-penicillamine	2 RSNO → 2 NO [·] + RSSR	Reaction catalysed by light, heat and metal ions (Cu ²⁺ , Hg ²⁺), under certain conditions also nitrosonium (NO ⁺) can be formed
DEA NONOate (and other NO-amine adducts)	Diethylamine NONOate	R ₂ N-NO-NO + H ⁺ → 2 NO [·] + R ₂ -NH ₂	Rate of NO release from NONOate is highly pH-dependent
Peroxyntirite donors			
SIN-1	3-Morpholinosydnonimine	SIN-1 → NO [·] + O ₂ ⁻ + SIN-1C → ONOO ⁻ + SIN-1C	Spontaneous decomposition in presence of oxygen
NO reactions in biological milieu			
Oxygen	O ₂	NO + O ₂ → NO ₂	End-products: NO ₂ ⁻ , NO ₃ ⁻ (in presence of hemoglobins)
Superoxide anionradical	O ₂ ^{-·}	NO + O ₂ ^{-·} → ONOO ⁻	End-products: NO ₃ ⁻
Thiols	R-SH	NO + O ₂ → NO ₂ + NO → N ₂ O ₃ N ₂ O ₃ + RSH → RSNO + NO ₂ ⁻ + H ⁺	End-products: S-nitrosothiols, nitrite, disulphides or mixed sulphides
Oxyphytoglobins	HbFe ²⁺ O ₂	HbFe ²⁺ O ₂ NO	End-products: NO ₃ ⁻ , metaphytoglobin,
NO detection			
Reaction partner	Chemical name	Reaction(s)	Comments
O ₃	Ozone	NO + O ₃ → NO ₂ * NO ₂ * → NO ₂ + light	Reaction exploited in specialized instruments such as a chemiluminescence detector for analysis of NO, nitrites and S-nitrosothiols
cPTIO	(2-(4-Carboxyphenyl)-4,4,5,5-tetramethylimidazole-1-oxyl-3-oxide potassium salt)	NO [·] + cPTIO → NO ₂ ⁻ + cPTI	Carboxy derivative (cPTIO) is preferably used as NO scavenging controls due to higher pH stability
DAF-2/DAF-2 DA	4,5-Diaminofluorescein diacetate	NO + O ₂ → NO ₂ NO ₂ ⁻ + DAF-2 → DAF-2T	Reaction of NO with difluorescein-based probes is O ₂ -dependent
DAF-FM/DAF-FM DA	4-Amino-5-methylamino-2',7'-difluorofluorescein diacetate	NO + O ₂ → NO ₂ NO ₂ ⁻ + DAF-FM → DAF FM- 2T	Reaction of NO with DAF-FM probes is O ₂ and pH dependent

the application of high concentrations of these compounds (orders of magnitude higher compared to animal NOS studies) and possible inhibitory effects on other plant enzymes, such as arginase or arginine decarboxylase (Reisser *et al.*, 2002), iron-containing enzyme (Peterson *et al.*, 1992) and NR (Rasul *et al.*, 2012). Therefore, as standard good experimental practice, the use of L-arginine derivatives should also include their inactive D-enantiomers as a control. However, as their true molecular targets in land plants still remain enigmatic, corresponding caution is advised in the interpretation of results derived from the application of these NOS inhibitors.

Conclusions

It is apparent that significant methodological improvements are required in plant NO research to support more robust data

acquisition. The plant research community should also be open to the adaption of methods and approaches from animal studies, but these should be applied with care. In the meantime, the existing procedures and methods should be deployed in a careful and thoughtful fashion to mitigate their disadvantages, following whenever possible the recommendations as summarized here:


- NO/RNS detection and/or quantification should include at least two different methods based on different principle/reaction mechanisms.
- Application of a pharmacological approach, such as chemical NO/RNS donors, scavengers or inhibitors, should include available negative and/or positive controls in a range of concentrations.
- Methods transferred directly from the animal NO field, should be subjected to careful testing and validation of their applicability on specific plant species.


• Data interpretation should take account of known methodological limitations, including possible unspecific reactions and interference by ROS and other plant reactive compounds. The detection limits of the employed methods and their nonquantitative nature in certain experimental settings should also be considered.


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
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
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