

REVIEW

Oxidative depolymerization of polysaccharides by reactive oxygen/nitrogen species

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Reactive oxygen species (ROS) and reactive nitrogen species (RNS) are constantly produced and are tightly regulated to maintain a redox balance (or homeostasis) together with antioxidants (e.g. superoxide dismutase and glutathione) under normal physiological circumstances. These ROS/RNS have been shown to be critical for various biological events including signal transduction, aging, apoptosis, and development. Despite the known beneficial effects, an overproduction of ROS/RNS in the cases of receptor-mediated stimulation and disease-induced oxidative stress can inflict severe tissue damage. In particular, these ROS/RNS are capable of degrading macromolecules including proteins, lipids and nucleic acids as well as polysaccharides, and presumably lead to their dysfunction. The purpose of this review is to highlight (1) chemical mechanisms related to cell-free and cell-based depolymerization of polysaccharides initiated by individual oxidative species; (2) the effect of ROS/RNS-mediated depolymerization on the successive cleavage of the glycosidic linkage of polysaccharides by glycoside hydrolases; and (3) the potential biological outcome of ROS/RNS-mediated depolymerization of polysaccharides.

Keywords: degradation / nitric oxide / polysaccharide / superoxide

Introduction

Reactive oxygen species (ROS) is defined as chemically active products generated by the partial reduction in oxygen, whereas reactive nitrogen species (RNS) refers to reactive products derived from reactions with nitric oxide (NO). Because an extensive overlap and cross-talk exist in the production, function, and

decomposition of ROS and RNS, these species are often referred to interchangeably (Winterbourn 2008), and in this minireview collectively termed ROS/RNS. Both reactive species can be generated from either exogenous (γ - or UV-irradiation and ionization, etc.) or endogenous (mitochondrial respiration, ROS/RNS-producing enzymes, etc.) sources (Kohen and Nyska 2002).

In living organisms, the outcome of the ROS/RNS action may be either beneficial or deleterious. On one hand, ROS/RNS function as intra- and extracellular messengers playing an important role in various signal transduction pathways and in transcriptional regulation (Fialkow et al. 2007; Yao et al. 2007). On the other hand, prolonged elevated levels of ROS/RNS can cause intense oxidative stress, a deleterious process that can be an important mediator of damage to various cellular structures, including lipids and membranes, proteins, and DNA (Valiko et al. 2007).

Polysaccharides are biopolymers possessing repeating units (mono- or oligosaccharide) joined together by glycosidic bonds with an enormous diversity of structure. Abundant polysaccharides are biosynthesized by animals, fungi, algae, microorganisms, and plants. These polysaccharides, in addition to being storage polymers or structure forming macromolecules, are increasingly recognized as key substances with versatile bioactivities and broad applications (Heinze et al. 2006). Studies have established that the functional roles of a given polysaccharide from both prokaryotic and eukaryotic cells are linked to its degree of polymerization (molecular size; Paoletti et al. 1992; Kasper et al. 1996; Wessels et al. 1998). In the past several decades, accumulating evidence has shown that ROS/RNS attack can result in the fragmentation of polysaccharides, and thus presumably alter the functions of these molecules. This paper will focus on the oxidative cleavage of polysaccharides by ROS/RNS and its potential outcome.

Chemical depolymerization of polysaccharides by ROS/RNS

ROS/RNS can be classified as either free radicals, which contain one unpaired electron or nonradicals. The former generally includes superoxide ion radical (O_2^-), hydroxyl radical ($OH\cdot$), peroxy ($ROO\cdot$), alkoxy radicals ($RO\cdot$), and NO radical ($NO\cdot$). The latter contains hydrogen peroxide (H_2O_2), organic peroxide ($ROOR'$), ozone (O_3), hypochlorous acid (HClO), singlet oxygen (1O_2), aldehydes (HCOR), and peroxynitrite (ONOOH), etc. (Kohen and Nyska 2002). Several ROS/RNS can individually cause the backbone scission of polysaccharides, generating smaller fragments.

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Ozone

Ozone can directly oxidize organic polymers possessing olefinic or double bonds. In addition, ozone breakdown to dioxygen gives rise to oxygen-free radicals, which are highly reactive and capable of damaging many organic molecules (Rao and Davis 2001). Furthermore, ozone was reported to react with acetal functions in a specific manner to give esters in which one of the alkoxy residues in the acetal was retained in the acyloxy group (Deslongchamps and Moreau 1971; Wang et al. 1998). The polymerization of polysaccharides by ozone in aqueous solution involves three mechanisms (Wang et al. 1999): (1) selective ozonolytic oxidation of β -D-aldosidic linkages, (2) nonselective oxidative degradation by radical species, and (3) nonselective acid hydrolysis. Among them, the first is the predominant reaction in which aldonic acid esters are formed and are spontaneously saponified to yield smaller fragments (Scheme 1). The ozonolytic oxidation of aldoses proceeds under a strong stereoelectronic control and prefers the aglycone conformation, in which each oxygen has one of its lone-pair orbitals antiperiplanar to the alkylidene C–H bond. Therefore, glycosidic linkages with different conformations can have different reaction rates with ozone, allowing for selectivity in cleaving β -D-linkages of polysaccharides (Wang et al. 1999).

Hypochlorite

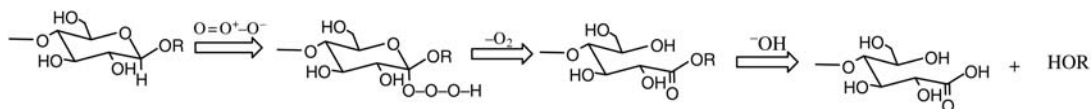
N-acetyl, NH_2 , $\text{N-SO}_3\text{H}$ residues on glycosaminoglycans (GAGs) can react with hypochlorite to generate polymer-derived *N*-chloro derivatives (chloramines, dichloramines, chlorosulfonamides) (Rees et al. 2003, 2004; Rees and Davies

2006; Scheme 2). In the presence of transition-metal ions and/or superoxide, the decomposition of these derivatives gives rise to nitrogen-centered radicals. These radicals undergo rapid intramolecular abstraction reactions to give carbon-centered radicals at C-2 on the amino sugar rings (via a 1,2-hydrogen atom shift; Rees and Davies 2006) and/or at C-4 on the neighboring glycosidic residues (via 1,5-hydrogen atom shifts; Rees et al. 2004). These products finally cause the cleavage of glycosidic bonds through β -scission (Rees et al. 2004; Rees and Davies 2006).

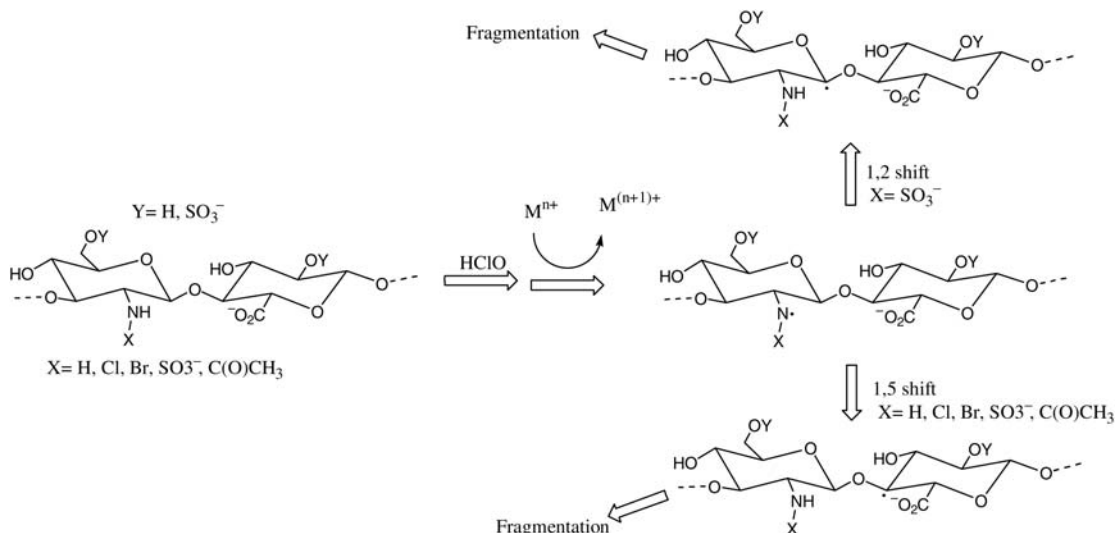
OH radical and peroxyxynitrite

The hydroxyl radical is the most reactive ROS. This radical can abstract hydrogen atoms at all ring C–H bonds of aldoses, uronic acids, and other sites on carbohydrates except C-2 of *N*-acetyl hexosamine (Gilbert et al. 1981; Hawkins and Davies 1996). The abstraction of hydrogen atom will generate carbon-center radicals. The radicals at carbons which form glycosidic bonds will undergo a β -scission reaction resulting in the breakdown of polysaccharide chains (Gilbert et al. 1981; Hawkins and Davies 1996; Rees et al. 2008). Notably, sulfated polysaccharides are shown more resistant to hydroxyl radical attack (Moseley et al. 1995).

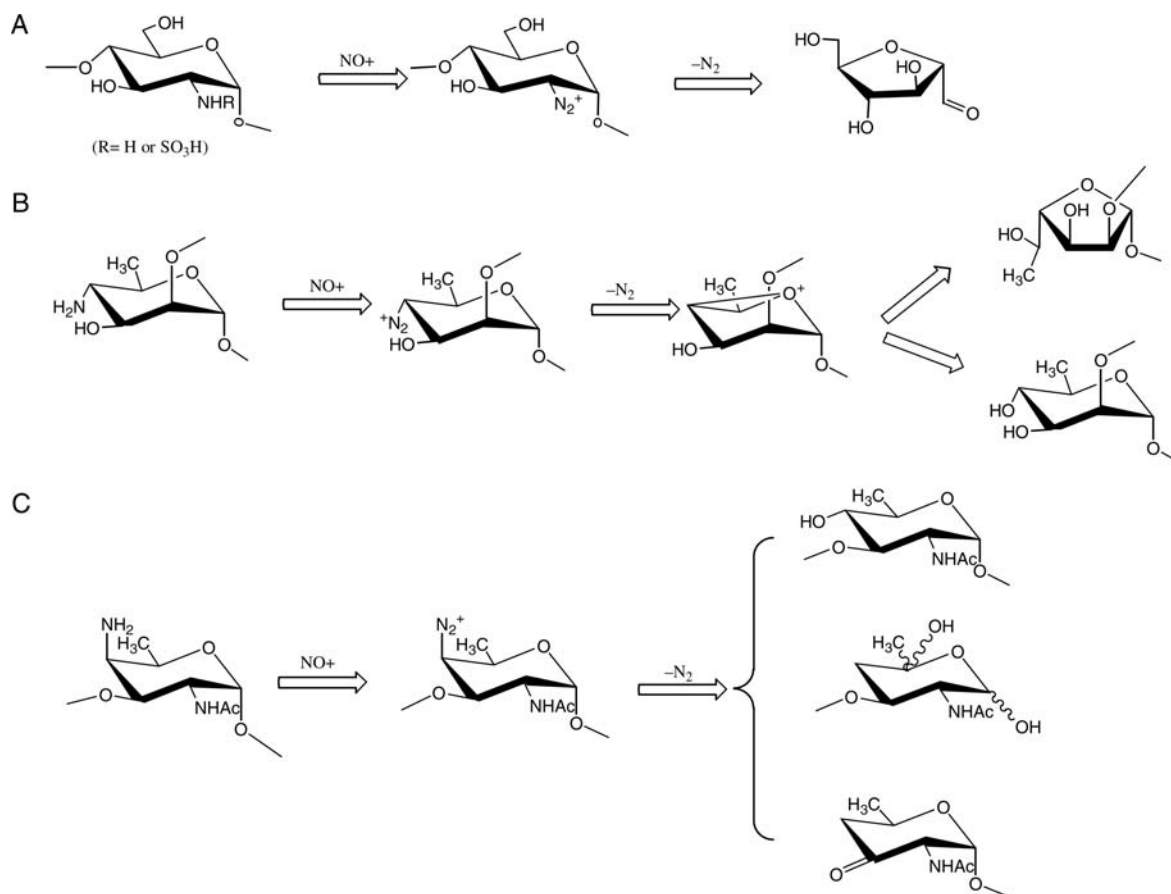
Peroxyxynitrite has been shown to degrade hyaluronic acid (HA) and chondroitin sulfate via a hydroxyl radical-like mechanism (Li et al. 1997; Al-Assaf et al. 2003; Kennett and Davies 2007). The susceptibility of the various GAGs to degradation by peroxyxynitrite appears to depend on their degree of sulfation, for example, hyaluronan (lacking sulfate) was the most susceptible and heparin (highly sulfated) was



Scheme 1. The ozonolytic cleavage of β -D-linked glycoside (adapted from Wang et al. 1998, 1999).



Scheme 2. The cleavage of glycosidic bonds in GAGs through β -scission induced by hyperchlorite (adapted from Rees et al. 2004; Rees and Davies 2006).



Scheme 3. The deamination of 2-amino or 4-amino hexamine residues (adapted from Lindberg et al. 1980; Kenne et al. 1982; Vilar et al. 1997).

the least susceptible to peroxynitrite-mediated degradation (Kennett and Davies 2007).

Singlet oxygen

Singlet oxygen was considered to be involved in the initiation step of the photodegradation of polymers (Jan et al. 1974). In a model system, singlet oxygen was produced by xanthine oxidase, and as a result, there was a dramatic decline in the solution viscosity (30–50%) of plant polysaccharides, such as methylcellulose, starch, pectin, and guar gum (Kon and Schwimmer 1977). The decreased viscosity suggested that the degradation of the polysaccharide chains had taken place. Further experiments of ROS quenchers and scavengers proved that this process was dependent on hydroxyl radicals and singlet oxygen (Kon and Schwimmer 1977). In another study, singlet oxygen generated from irradiation in the presence of methylene blue or riboflavin (a singlet oxygen sensitizer) was shown to generate a slight depolymerization of HA, but did induce substantial changes in the HA tertiary structure (Andley and Chakrabarti 1983).

NO and nitrous acid

NO and HNO₂ share one common intermediate, the nitrosonium cation (NO⁺) to depolymerize GAGs (Shively and Conrad 1976; Vilar et al. 1997; Hassan et al. 1998). The nitrosonium cation

nitrosates free amino groups at ~pH 4 or *N*-sulfo groups at ~pH 1.5, respectively, but not *N*-acetyl groups. The nitrosation of amino groups or *N*-sulfo groups causes loss of nitrogen gas with a ring contraction of 2-amino-2-deoxy sugars to 2,5-anhydro sugars coupled to elimination of the aglycone (Scheme 3a). In addition, 4-amino-4-deoxy sugars are also amenable to the deamination reaction through a reaction sequence similar to 2-amino-2-deoxy sugars (Lindberg et al. 1980; Kenne et al. 1982). However, whether NO-mediated deamination of 4-amino sugars results in the cleavage of the glycosidic bonds depends on the favored conformation of the 4-amino group and the glycosidic linkage of 4-amino-4-deoxy sugars in the carbohydrate polymers. For example, the deamination of 4-amino groups of 2-linked 4-amino-4, 6-deoxymannopyranosyl residues yields 2-linked rhamnopyranosyl and 2-linked 6-deoxyallofuranosyl residues, but the glycosidic linkages stay intact (Scheme 3b; Kenne et al. 1982). In contrast, the deamination of 4-amino groups of 3-linked 2-acetamido-4-amino-2,4,6-trideoxygalactopyranosyl (AATp) residues yields cleavage at both C-3 and C-1 of AATp (Scheme 3c; Lindberg et al. 1980).

Polysaccharide depolymerization by cell-derived ROS/RNS

Although numerous individual ROS/RNS have been shown to depolymerize polysaccharides in vitro, there are at least three

factors to take into account when considering the damage to biological substrates including carbohydrates *in vivo* (Kohen and Nyska 2002; Winterbourn 2008): (1) whether substrates are located at adjacent ROS/RNS production sites, since the high reactivity of these species usually has a relatively short life span; (2) whether the concentration of ROS/RNS is high enough; and (3) whether the levels of antioxidant are efficient to decompose ROS/RNS at the attack sites. The following section will describe the depolymerization of various polysaccharides by ROS/RNS generated from different cell types.

Neutrophils

During defense against infections, neutrophils survey, ingest, and destroy invading microorganisms in mammals, and secrete potent antimicrobial products that includes ROS/RNS, proteinases, and antimicrobial peptides (Fialkow et al. 2007). These ROS/RNS are produced intracellularly and can be released extracellularly (Freitas et al. 2009).

The GAGs, such as hyaluronan, chondroitin 4-sulfate, and dermatan sulfate, were shown to be degraded with loss of both uronic acid and hexosamine residues when cultured with PMN stimulated by phorbol myristyl acetate (PMA). The degradation pattern of GAG was very similar to the pattern of degradation observed following hydroxyl radical attack (Moseley et al. 1997). The ability to degrade hyaluronan appears to be specific to neutrophils stimulated with PMA, not with fMLP, concanavalin-A, or digitonin (Saari et al. 1993; Rees et al. 2008). Degradation of hyaluronan of joint synovial fluid has been linked to rheumatoid arthritis. In the synovial fluid of rheumatoid arthritis, the fragmented hyaluronan was considered to be due to the attack by ROS such as the hydroxyl radical (McCord 1974; Grootveld et al. 1991), and activated neutrophils were considered as the major cell types to produce ROS degrading hyaluronan (Greenwald and Moak 1986). Furthermore, ROS-mediated hyaluronan degradation generated products with a polydisperse size and possibly with different conformational characteristics due to radical-featured repolymerization (McNeil et al. 1985). Interestingly, rat mast cell granules-associated heparin could even be degraded by normal neutrophils. Moreover, neutrophils from patients with chronic granulomatous disease [deficiency of nicotinamide adenine dinucleotide phosphate (NADPH) oxidase] had a dramatic decrease in their ability to degrade heparin (Metcalf et al. 1990). This provided direct evidence demonstrating that ROS in neutrophils are the major reactive species responsible for heparin degradation.

Antigen presenting cells

Polysaccharide A (PSA, see structure in Figure 1), the most immunodominant and highly conserved zwitterionic polysaccharide from the anaerobic bacterium *Bacteroides fragilis* plays a key role in both the pathogenic and the beneficial biological effects of this microbe. On the one hand, PSA facilitates the induction of abscesses when *B. fragilis* enters a normally sterile site (Tzianabos et al. 2000); on the other hand, PSA specifically directs maturation of the mammalian immune system and protects animals from inflammatory bowel disease through the induction of interleukin-10-producing CD4⁺ T cells (Mazmanian et al. 2005, 2008). Both

of these important outcomes are directly attributable to the ability of PSA to activate CD4⁺ T cells. T cell activation requires presentation of PSA by antigen presenting cells (APCs) through the major histocompatibility complex II (MHCII) pathway (Cobb et al. 2004). In marked contrast to the processing of protein antigens in the acidic endolysosomal compartment by proteases, processing of PSA depended on the presence of inducible NO synthase (iNOS), but not NADPH oxidase in the APCs (Cobb et al. 2004). PSA induces iNOS mRNA expression and NO radical production in dendritic cells (DCs) through toll-like receptor 2 activation (Wang et al. 2006; Duan et al. 2008). The degradation of PSA was greatly suppressed in iNOS-deficient DCs when compared with wild type (WT) DCs. In contrast, *N*-acetylated PSA (complete *N*-acetylation of all free amino groups in PSA) was degraded equally between WT and iNOS-deficient DCs (Duan et al. 2008). In addition, native PSA, but not *N*-acetylated PSA was significantly degraded with NO in MPO [myeloperoxidase, which restricts the production of NO (Kumar et al. 2005)] deficient macrophages when compared with WT macrophages. These observations suggested that iNOS-derived RNS is the predominant species to degrade the native PSA molecule with deamination probably being the only responsible mechanism. As opposed to the native PSA, a partially deaminated product PSA-NO (~16 kDa), which has an intact zwitterionic motif bypasses this cleavage step and can directly bind to MHCII for presentation to the $\alpha\beta$ T cell receptor (Figure 1). Similarly, capsular polysaccharide SP1 form type I *Streptococcus pneumoniae* was also degraded in endosome/lysosome of APCs in an NO-dependent manner (Velez et al. 2009). In contrast to PSA degradation by RNS, dextran was depolymerized in the endolysosome, and this degradation attributed to hydrogen peroxide- and superoxide-derived ROS (Duan et al. 2008).

Endothelial cells

After incubation of heparin with human umbilical vein endothelial cells (HUVECs), the degradative products of heparin were recovered in the medium (Vilar et al. 1997). This degradation depended on NO, which is synthesized constitutively by endothelial NO synthase. In contrast, hyaluronan stayed intact when cultured with HUVECs, although other reactive species, such as hydroxyl radical and peroxynitrite, can chemically depolymerize hyaluronan (Li et al. 1997). Deamination is one possible mechanism for NO-mediated degradation of heparin, but not hyaluronan because in the latter molecule all free amino groups are *N*-acetylated.

Cell-associated proteoglycans containing heparan sulfate (HS) are involved in the development, tissue repair, and tumorigenesis (Bernfield et al. 1999). In particular, HS side chains are capable of binding and/or activating and/or transporting many growth factors, cytokines, enzymes, viral proteins, and polyamines (Bernfield et al. 1999). HS proteoglycans on the surface are continuously endocytosed, degraded, and newly resynthesized through a recycling process in vascular endothelial cells (Mani et al. 2000), normal fibroblasts (Fransson et al. 1995), and carcinoma cells (Mani et al. 2007). During recycling, in addition to heparanase-catalyzed degradation, the HS side chains are

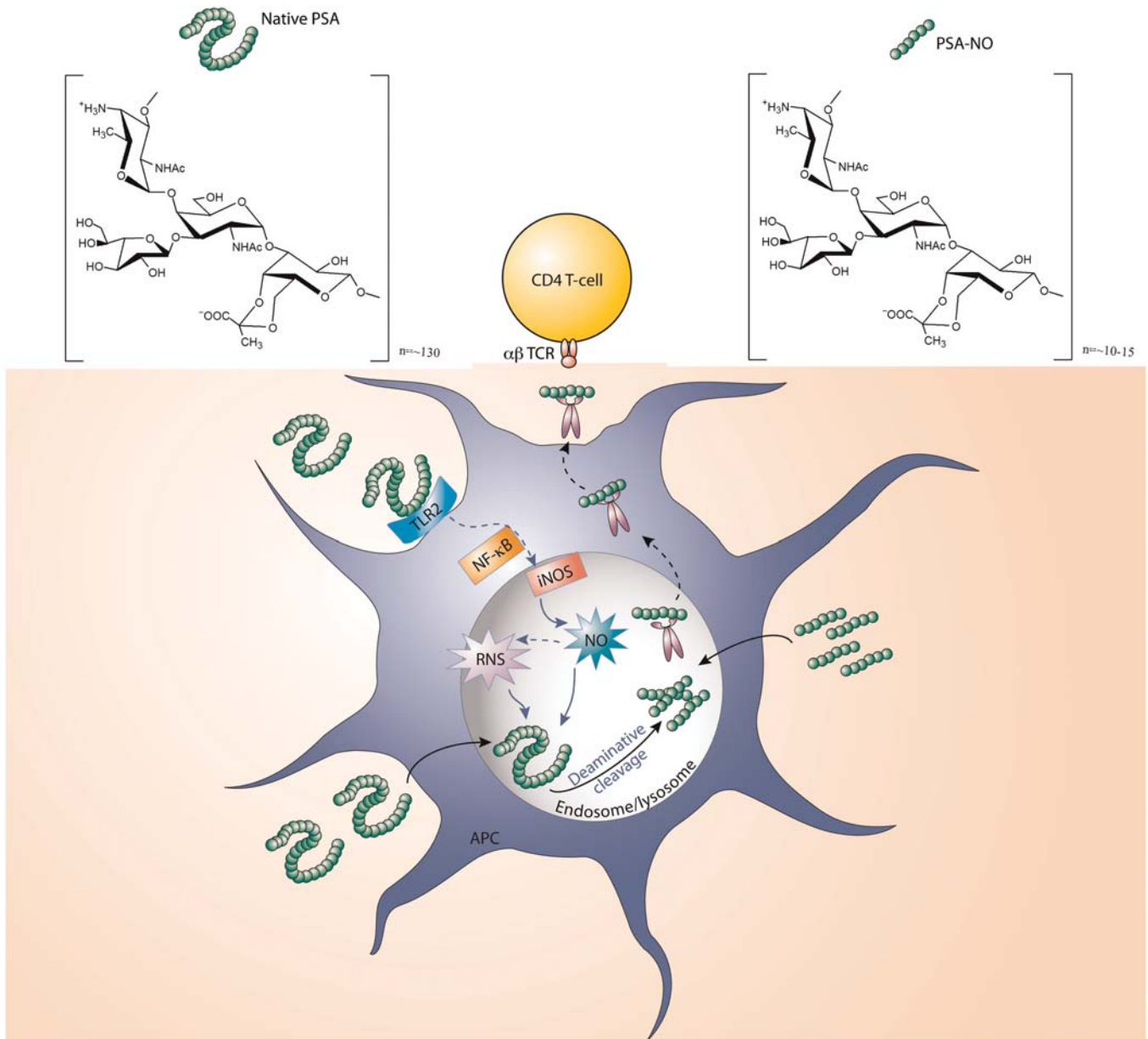


Fig. 1. A proposed model of APC processing and presentation of *B. fragilis* PSA and its pre-processed products. The native PSA molecule (~130 repeating units) produced by *B. fragilis* comes into contact with a DC and initially activates the cell through toll-like receptor 2-mediated signaling resulting in increased NF-κB-mediated transcription of iNOS. iNOS upregulation results in enhanced production of NO by the cell. NO is the chemical responsible for the deaminative cleavage of PSA to its product PSA-NO (~10 repeating units). Native PSA requires deaminative cleavage prior to binding to MHCII and subsequent presentation to the αβ T cell receptor of CD4⁺ T cells. If experimentally, PSA is pre-processed by reacting it with NO in vitro, the product PSA-NO bypasses this cleavage step and can directly bind to MHCII for presentation to the αβ-T cell receptor.

degraded via an NO-dependent deaminative cleavage at N-unsubstituted glucosamine residues. Deaminative cleavage will generate a reducing terminal sugar anhydromannose, which could be visualized by a specific fluorophore-labeled monoclonal antibody (Cheng et al. 2002). In this process, NO/HNO is originated from preformed S-nitroso groups in the core protein of HS proteoglycans, and the degradation is found to start in early endosomes and complete in late endosomes (Mani et al. 2007).

Plant cells

Cellulose, hemicelluloses (mainly xyloglucan in the plant), and pectin are the major carbohydrates making up the primary (growing) plant cell wall. Cell wall loosening is an important developmental process in key stages (seed germination, elongation growth, and fruit ripening) of the plant life cycle (Muller et al. 2009). This process requires structural changes in the cell wall. Enzymatic hydrolysis of cell wall polysaccharides by glycosidases, such as β-1,3-glucanase,

β -1,4-mannanase, xyloglucan endotransglucosylase/hydrolases, etc., is essential for wall modification (Chen et al. 2002; Leubner-Metzger 2002; da Silva et al. 2004). As an additional plant cell wall-loosening agent, ROS-like hydroxyl radical (OH) presumably can target cell wall components including polysaccharides, leading to the breakage of cell wall polysaccharides (Muller et al. 2009). In fact, apoplastic hydroxyl radical and superoxide radicals were detected in cress caps and radicles in vivo during seed germination. By ways of electron paramagnetic resonance spectroscopy and ^3H fingerprinting, it was found that hydroxyl radical attacks tissue-specific, not random cell wall polysaccharides (Schopfer 2001). In addition to ROS, plant tissues also synthesize NO via the nonenzymatic reduction in apoplastic nitrite (Bethke et al. 2004), whereas it remains unknown whether NO-derived RNS such as peroxyxynitrite is involved in the depolymerization of cell wall polysaccharides in vivo.

Polysaccharide depolymerization by ROS/RNS in concert with enzymes

The catabolism of polysaccharide is mainly achieved by glycoside hydrolases in vivo. Genetic deficiency in enzymes essential for cleaving carbohydrate chains, for example, mucopolysaccharidoses, will cause serious diseases characteristic of polysaccharide accumulation in mammalian tissues (Dorfman and Matalon 1976). However, as an alternative and supplementary mechanism, especially in inflammatory conditions, ROS/RNS might play a role in the depolymerization of polysaccharides. In fact, there is evidence suggesting that the oxidatively cleaved polysaccharides by ROS/RNS could serve as the substrates for the enzymatic cleavage that follows, thus facilitating depolymerization of the polysaccharide (Greenwald and Moy 1980; Metcalfe et al. 1990). For instance, prior treatment with ROS/RNS rendered hyaluronan more susceptible to degradation by lysosomal carbohydrases such as β -glucuronidase or β -*N*-acetylglucosaminidase (Greenwald and Moy 1980). One explanation is the partial degradation of polysaccharides by ROS/RNS would overcome steric factors which prevent the interaction between enzymes and highly polymerized substrates (Greenwald and Moy 1980). Notably, the substantial alternation in the structure of unfragmented polysaccharide by ROS/RNS was shown to slow down the cleavage of glycosidic bonds by enzymes with high specificity. Compared with original hyaluronan having a similar molecular size, the polymer modified by ROS/RNS originating from ultrasonic treatment was more resistant to enzymatic cleavage by specific hyaluronidase (Chabreck et al. 1991; Kohen and Nyska 2002).

Brown rot basidiomycetes, the highly destructive wood decay fungi, such as *Gloeophyllum trabeum* and *Postia placenta*, can produce significant quantities of extracellular ROS to degrade cellulose (Hammel et al. 2002; Baldrian and Valaskova 2008). In the case of *P. placenta*, endoglucanases were only active on cellulose after oxidative pretreatment (Ratto et al. 1997). Thus it was proposed that the initial depolymerization probably facilitates the penetration and entry of oxidative and hydrolytic enzymes which further degrade celluloses (Hammel et al. 2002). Actually, one recent study

(Vaaje-Kolstad et al. 2010) had discovered that a chitin-binding protein CBP21 oxidatively catalyzed the cleavage of glycosidic bonds in crystalline chitin in the presence of an external electron donor. The resulting oxidized chain ends effectively promoted further degradation of chitin by chitinases.

Potential outcome of ROS/RNS-mediated scission of polysaccharide chains

The classical immunological paradigms of antigen presentation has MHCII molecules presenting extracellular antigens, whereas MHCI molecules presenting antigens located in the intracellular compartment (Cresswell 1994; Pamer and Cresswell 1998). The molecules that these pathways are known to present are proteins or protein conjugates. In the MHCI pathway, intracellular protein antigens are processed by the cytosolic proteasome in APCs and subsequently presented to CD8+ T cells. Similarly, extracellular protein antigens are degraded by endosomal and lysosomal proteases in APCs before being presented by MHCII molecules to CD4+ T cells (Abbas and Lichtman 2005). Glycolipids have been shown to be presented by CD1 molecules directly or after being degraded by an enzymatic process (Schaible and Kaufmann 2000; Prigozy et al. 2001). Interestingly, polysaccharides are depolymerized via a ROS/RNS-dependent manner in APCs (Duan et al. 2008). In particular, zwitterionic polysaccharides, i.e. PSA from commensal bacteria *B. fragilis*, are processed to a significantly smaller molecular size (\sim 10–15 kDa) within endosome/lysosome through a deaminative mechanism. Thus, MHCII pathway-mediated carbohydrate antigen processing in APCs could be achieved by ROS/RNS-mediated oxidative reactions.

In mammals, hyaluronan, which is an important component of the extracellular matrix involved in the structure of connective tissues, can modulate a variety of cellular and tissue functions (Raines et al. 2000). Hyaluronan is a linear polymer composed of repeating units of the disaccharide [$\text{-D-glucuronic acid-}\beta$ -1,3-*N*-acetyl-D-glucosamine- β -1,4-]. Differently sized hyaluronans trigger different signal transduction pathways (Jiang et al. 2005; Jiang et al. 2007). For example, tetrasaccharides are antiapoptotic and inducers of heat shock proteins (Jiang et al. 2007). Hyaluronan oligomers with 8–16 repeating units, stimulated angiogenesis in vivo, and endothelial proliferation in vitro (Stern et al. 2006); smaller hyaluronans (<500 kDa), but not the native hyaluronans (>1000 kDa) induce proinflammatory responses in macrophages (Jiang et al. 2005). Larger hyaluronan (1000–5000 kDa) suppress angiogenesis, immune responses, and phagocytosis (Jiang et al. 2007). In rheumatoid arthritis, neutrophil-derived ROS attack has been shown to be responsible for the fragmentation of hyaluronan in joint synovial fluid (McCord 1974; Grootveld et al. 1991). The differently sized hyaluronans might be important in regulating the disease process.

In plants, intact chain length of cell wall polysaccharides is a requisite for structural integrity of the plant cell wall (Herron et al. 2000). The depolymerization of these polysaccharides leads to loosening of the cell wall and plays a major role in fruit softening during ripening (Prasanna et al. 2007)

in which ROS/RNS-like hydroxyl radical were shown to be involved (Schopfer 2001; Muller et al. 2009).

Conclusions and future directions

Oxidative degradation of polysaccharides might result in non-specific and/or specific scission of carbohydrate chains. This process could be an alternative and/or supplementary way for glycoside hydrolases-mediated hydrolysis to depolymerize polysaccharides. Indeed, ROS/RNS can degrade polysaccharides and lead to the subsequent enzymatic cleavage of glycosidic bonds. Cellular levels of ROS/RNS can effectively breakdown polysaccharides, and the data suggest that oxidative cleavage of carbohydrate chains is biologically relevant. It appears that ROS/RNS can also modify the monosaccharide composition of polysaccharides and potentially generate polymer fragments possibly having properties substantially different from those of the original macromolecule. However, in a biological context, the consequence of this change remains unclear.

In encapsulated pathogenic bacteria, polysaccharides on their surfaces contribute to host virulence by many mechanisms including: antiphagocytic and antibacteriolytic activity; immune evasion; immune modulation; and biofilm formation (Comstock and Kasper 2006). These capsular polysaccharides always have very large molecular size, and their immunogenicity has been found to be directly related to the chain length (Kabat and Bezer 1958; Howard et al. 1971; Kasper et al. 1982; Kalka-Moll et al. 2000). For instance, dextran with a molecular mass <80,000–90,000 Da does not elicit an antibody response in humans (Kabat and Bezer 1958). Given ROS/RNS are released massively during the crosstalk between host and invading organism, the fragmented products of capsular polysaccharides presumably would be generated intracellularly and/or extracellularly. It remains unknown whether these decomposed carbohydrates could elicit more progressive or suppressive humoral immune responses when compared with those elicited by native polysaccharides. In APCs, it has been found that the predominant products (~10 kDa) of the degradation of polysaccharides by RNS/ROS within endosome/lysosome are produced over a period of time. However, either prolonging treatment of ZPS or dextran with exogenous ROS/RNS or increasing the ROS/RNS concentration results in formation of smaller degraded products in vitro. These data suggested that the ROS/RNS-induced depolymerization of polysaccharide is finely controlled in APCs (Duan et al. 2008). More recently, it has been shown that the oxidation of endocytosed PSA release protons which in turn inhibits the breakdown of PSA. That is, the oxidative processing of carbohydrate antigens is operated in a self-regulatory feedback mechanism (Lewis and Cobb 2010). At the present, it awaits to be understood whether this feedback mechanism is universal to the processing of other carbohydrates in APCs.

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Abbreviation

AATp, 3-linked 2-acetamido-4-amino-2,4,6-trideoxygalactopyranosyl; APCs, antigen presenting cells; DC, dendritic cell; GAG, glycosaminoglycan; HA, hyaluronic acid; HS, heparan sulfate; HUVECs, heparin with human umbilical vein endothelial cells; iNOS, inducible NO synthase; MHCII, major histocompatibility complex II; MPO, myeloperoxidase; NADPH, nicotinamide adenine dinucleotide phosphate; NO, nitric oxide; PMA, phorbol myristyl acetate; PSA, polysaccharide A; RNS, reactive nitrogen species; ROS, reactive oxygen species WT, wild type.

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