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MICROBIOLOGY

Free Radicals and Actinobacteria as a Misexplored Goldmine of Antioxidant Compounds

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Abstract: Free radicals are highly reactive unstable molecules, which can be synthesized in different ways, considered harmful and threatening to humans; these chemical species have free traffic throughout the human body, interacting with biological molecules and human body organ tissues. The interaction between free radicals and biological molecules is the main factor for disease development or pre-existing disease symptoms aggravation. Antioxidants are chemical compounds able to donate electric charge to stabilize molecules such as free radicals. Recent studies have proved the benefits of antioxidants intake in health improvement. In this way, the search for natural sources of antioxidants has become an ascending trend. In this field, the microbial sources are considered poorly explored compared to the numerous amount of other compounds obtained from them, especially from Actinobacteria. The searched literature about Actinobacteria highlights an important capacity of producing natural antioxidants; however, there is a lack of in vivo studies of these isolated compounds. In this review, we gathered information that supports our point of view that Actinobacteria is a truly renewable and superficially explored source of natural antioxidants. Furthermore, our purpose is also to point this limitation and stimulate more researches in this area.

Key words: Free radicals, antioxidants, oxidative stress, Actinobacteria antioxidant.

INTRODUCTION

Free radicals are dangerous molecules to cells and the human body. The excess of these molecules causes tissue damage and disease development or worsening. The free diffusion of free radicals throughout the body and the malfunction of antioxidant enzyme systems are concomitant factors that contribute severely to reactivity with cell biomolecules. In a first approach, it was established that free radicals formation was restricted to aerobic metabolism, but actually, transition metals, organelles, and other enzymes can also synthesize free radicals. Some additional factors are involved in free radical syntheses such as xenobiotics, intake of industrial synthetic compounds, and lifestyle.

Antioxidants are molecules that stabilize free radicals donating protons or electrons. They can be a result of a chemically synthetic process or a natural metabolite. Thus, antioxidants are able to mitigate deleterious effects caused by free radicals and protecting the cells and tissues against damages, stopping free radical forming chain reactions, scavenging formed radicals, regenerating damaged structures, or reinforcing cell defendant systems. Lately, the search for natural products has increased; in the same way, the exploration for natural antioxidants has gained attention being directed to plant exploration, mostly.

Actinobacteria is a renewable source of natural bioactive molecules. This group of bacteria is a true reservoir of antimicrobial, antifungal, antitumor, and immunomodulator compounds. Even though, up to this time, there is no mention of any approved Actinobacteria compound with known antioxidant activity for human use. Then, it makes sense to hook up this absence and the emergent exploration of Actinobacteria toward antioxidant approach with some other factors, such as low or non-toxicity of Actinobacteria compounds in counterpart to synthetic antioxidants as described in the following issues.

This review focused on the antioxidant potential of molecules produced by Actinobacteria isolated from different environments and its potential of extinguishment of the most variety of unstable molecules, with many unknown and non-reported compounds as an escape route to this trouble. Data from the literature have demonstrated the quenching of some unpaired molecules such as hydrogen peroxide and cationic radicals by 2,6-dimethoxy terephthalicacidand1-hydroxy-nonrresistomicin as well as cell redox balance maintenance by angeloline A and actinosporines C and D all of them produced by Actinobacteria. Therefore, supported by these findings our purpose is also to show the potential of antioxidant compounds from Actinobacteria for human usage despite some evident limitations in this field.

FREE RADICALS

Origin, action on macromolecules and cell metabolism

Free radicals are widely defined as molecules or atoms with a characteristic imparity of at least one electron in the last atomic orbital. Free radicals are mainly derived from reactions involving molecular oxygen or nitrogen being grouped as Reactive Oxygen or Nitrogen Species (ROS and RNS) (Table I). These chemical species may be negatively or positively charged, have a short half-life and its electric disparity is the main factor to high reactivity with biological molecules in the origin site or outside via secondary mediators cascade (Nikki 2010).

Free radicals are mainly formed through oxidoreduction reactions by electron addition

	Radical	Non-Radical
	Superoxide (O_2^{-})	Hydrogen peroxide (H ₂ O ₂)
Reactive Oxygen Species	Hydroxyl (OH)	Singlet oxygen (O ₂)
(ROS)	Peroxyl (RO ₂ ⁻)	Hypochlorous acid (HClo)
	Alkoxyl (RO ⁻)	Ozone (O ₃)
		Hypobromous acid (Hbro)
Reactive Nitrogen Species (RNS)	Radical	Non-Radical
	Nitric oxide (NO ⁻)	Peroxynitrite (ONOO ⁻)
	Nitrogen dioxide (NO ₂)	Nitrosyl cation (NO⁺)
		Nitrosyl anion (NO ⁻)
		Nitrogen trioxide (NO ₃)
		Nitrogen tetraoxide (NO ₄)
		Nitrous acid (HNO ₂)

Table I. Reactive oxygen species and reactive nitrogen species.

Based on Phaniendra (2015).

or removal and can exist without binders or carriers. The aerobic metabolism (Reaction I) is the main source of ROS being the mitochondria the protagonist of electron transference reactions through oxidative phosphorylation in the electron transport chain. As described by Mourier & Larsson (2011), complex I (NADH ubiquinone oxidoreductase) and complex III (cytochrome bc1) are the main generators of reactive oxygen species.

Some other non-enzymatic reactions such as those involving transition metal ions can also lead to ROS and RNS formation. Fenton reactions (Reactions II - III) describe the interaction between ferrous (Fe²⁺) or ferric iron (Fe³⁺) and hydrogen peroxide (H₂O₂) forming hydroxyl (OH⁻) and hydroperoxyl (HO₂⁻) radicals (Propac et al. 2017). The propagation of the Fenton Reaction is headed by the Haber-Weiss reaction (Reaction IV) which catalyzes the reduction of ferric iron to ferrous iron through mitochondrial superoxide ion (O₂⁻) interaction, leading to a free radical formation looping by Fenton reaction.

ROS overproduction is the main factor for oxidative stress, which leads to diseases development or aggravation by tissue oxidative damage. Regardless of the interaction with transition metals to form RNS, the synthesis of nitric oxide (NO⁻) and peroxynitrite (ONOO⁻) (Reaction V) through L-arginine decomposition by oxide nitric synthase is also reported (Fionda et al. 2016). RNS display a crucial role as antimicrobial agents, immunological modulators, vascular function regulators, and signaling molecules (Adams et al. 2015). However, the excess of RNS generation is noxious to the body in order to nitrosative stress triggering.

Notwithstanding, the mitochondria as the main source of reactive species, other cytoplasmic organelles as peroxisomes (Tripathi & Walker 2016), endoplasmic reticulum (Wang & Kaufman 2016), and lysosomes (Fu et al. 2015) have been reported as free radical producers. Enzymes as cell membrane NADPH-oxidases (NOx) (Kovac et al. 2015), cytochrome P450 oxidase (CyP450) (Brown & Borutaite 2012), superoxide dismutase (SOD) (Nguyen et al. 2020), and nitric oxide synthases (eNOS) (Brown & Borutaite 2012) are also reported as reactive species sources (Figure 1).

Exogenous factors as xenobiotics are described as strong inducers of ROS synthesis (Żukowski et al. 2018). The metabolization of alcohol by cytochrome P450 oxidase yields ROS as a process byproduct (Li et al. 2016). Likewise, polycyclic hydrocarbons derivatives generate ROS by electron transfer to molecular oxygen through UV absorption and quinine metabolization (Yang et al. 2017a). The air pollutant ozone (O_3) is also pointed out as a dangerous ROS to cells (Aulakh et al. 2020).

> **Reaction I:** $O_2 + e^{-}: O_2^{-}$ **Reaction II:** $Fe^{2^+} + H_2O_2$: $Fe^{3^+} + OH^{-}$ **Reaction III:** $Fe^{3^+} + H_2O_2$: $Fe^{2^+} + OOH^{-} + H^{+}$ **Reaction IV:** $Fe^{3^+} + O_2^{-}: Fe^{2^+} + O_2$ **Reaction V:** $NO + O_2^{-}: ONOO^{-}$

Biological molecules are attacked at side chains releasing protons and electrons to stabilize reactive species. Fatty acids are the most prone biological molecules to oxidative damage followed by enzymes, proteins, amino acids, and nucleic acids. Lipid peroxidation is considered a threatening to cell membrane integrity as well as lipid hydroperoxides (LOOH) are dangerous to other macromolecules such as proteins (Zhang et al. 2018a). Among damages are fragmentations or inactivation via protein side-chain or polypeptide backbone attack, moreover post-translational modifications as nitrosylation and glutathionylation are also described (Martínez & Andriantsitohaina 2009. Huang et al. 2016). Many studies have announced the free radicals as mediators of DNA damage and repair response system. They may change

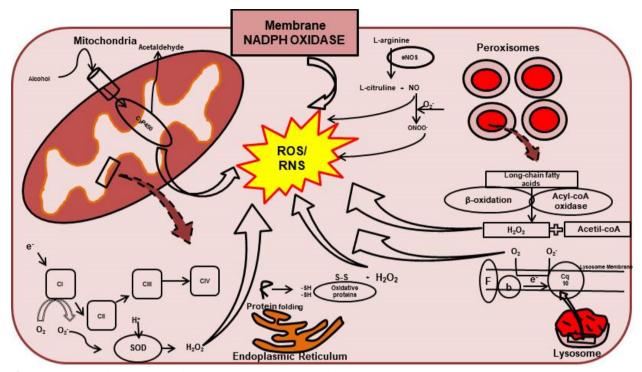


Figure 1. Schematic representation of ROS and RNS production by cell organelles and cell enzymes.

DNA structure by sugar backbone oxidation and nucleoside bases translocation (Srinivas et al. 2019).

The published literature has established free radicals' importance on cellular metabolism processes when under physiological concentrations. ROS and RNS are reported as cellular proliferation regulators (Diebold & Chandel 2016), mediators of necrosis induction (Qin et al. 2015), genic expression modulators (Furukawa et al 2004), NFkβ activators (Topal et al. 2017), autophagy and apoptosis modulators (Redza-Dutordoir & Averill-Bates 2016), MAPkinases activators (Henkler et al. 2010), and mediators of cellular inflammatory responses (Khojah et al. 2016).

FREE RADICALS AND DISEASE PATHOPHYSIOLOGY

Free radicals have a bifold role in a disease context. In the same way that triggers it can

also participate in disease pathophysiology. In this sense, it is still established that antioxidant system imbalance and free radical damage are cofactors for disease development. Currently, approaches for oxidative stress mitigation have demonstrated being an interesting therapeutic target for disease treatment. Based on this, in this section, we gathered recent shreds of evidence about how free radicals interact or prompt diseases in the main human body organs.

Nervous system diseases

The co-participation of free radicals on nervous system pathologies is studied since the end of the last century. The high oxygen intake makes the brain tissue extremely vulnerable to oxidative damage. This condition becomes an open door to the development of neurodegenerative, and other nervous system diseases (Driver et al. 2000). The brain oxygen metabolization is the responsibility of NOx family enzymes resulting in ROS as a process product, which combined with other inherent conditions of nervous system pathologies, contributes to a constant synthesis of free radicals and consequent aggravation of some nervous disease symptoms (Nayernia et al. 2014).

Parkinson's disease is mainly characterized by nigra substance cells' death and dopaminergic neuron loss as a result of ROS action which leads to continuous movement and cognitive losses (Guo et al. 2018a). The ROS also might aggravate chronic cerebral inflammation through inflammasome NLRP3 lineage recruitment by pannexin and connexin hemichannels interaction (Ahmadian et al. 2019). Oxidative stress is proposed as a contributing factor behind fundamental mechanisms of neuronal loss, cell death, and dopamine levels disturbances through oxidation by ROS (Goel & Chaudhary 2020).

Alzheimer's disease has striking characteristics such as cognitive impairment, memory losses, delusions, and mental embarrassment. Other hallmarks are synapse degeneration, hippocampus neuron death, b-amyloid deposition, and cerebral atrophy. The oxidation scenery in Alzheimer's disease is sustained by a combination of metallic ions, amyloidal aggregates, and mitochondrial dysfunction (Cheignon et al. 2018). Oxidative stress is closely linked to neuronal membrane integrity destruction. Even so, brain oxidative imbalance can also favor neurofibrillary tangle formation and increase nucleic oxidation in response to Tau protein conformation change (Wang et al. 2020).

Liver diseases

The liver is one of the body's detoxification centers being also coadjutant on metabolism balance regulation. Due to the susceptibility of hepatic tissue, reactive species attack is harmful and is also involved in physiopathology of hepatic diseases, however, in some of them, the underlying mechanisms are still not clarified (Kim et al. 2018, Liu et al. 2020). In hepatic steatosis, for example, some evidence in the literature points to a central role of mitochondrial dysfunction in liver steatosis, even with some studies highlighting the malfunction of the electron transport chain and oxidative phosphorylation. For instance, they associated the participation of NADPH oxidase with liver diseases, mainly because of the generation of anion superoxide (O_2^{-}) (García-Ruiz & Fernandéz-Checa 2018).

Within the hepatic environment, the free radicals when excessively present may act as signaling molecules driving upper regulation of collagen deposition genes such as TGF-β1 fibrogenic factor in hepatic fibrosis scenario (Tian et al. 2018), stimulating chronic hepatic inflammation through Hbx protein (Ha & Yu 2010), inducing lipid peroxidation that leads to hepatic tissue damage (Thuy le et al. 2017), and lipid deposition on non-alcoholic steatosis once the occurrence of mitochondrial imbalance (Marseglia et al. 2014). On the other side, the metabolization of acetaminophen by cytochrome P450 oxidase yields N-acetyl-pbenzoguinoneimine which disturbs oxidative balance leading to cellular apoptosis of hepatocytes, centrilobular necrosis, and free radicals generation enhancement (Hu et al. 2018).

It is also important to look at the other remarkable factors involved in the physiopathology of free radicals in liver diseases; within this group is alcohol. It was observed that alcohol is determinant to acute hepatic failure and alcoholic hepatic steatosis development once alcohol metabolization byproducts induce lipid peroxidation and free radicals synthesis in the liver contributing to death of hepatocytes, mainly (Wang et al. 2018). Summarily, in spite of the real scenario of a poorly studied physiology of free radicals within liver diseases mentioned at the top of this issue, some pieces of evidence proving the involvement of these species on liver pathologies have raised the proposal of mesenchymal stem cell therapy which is considered a promising strategy in acute liver injury treatment, as well as antioxidant therapy, pointed out as a possible, hopeful and less costly approach that has attracted attention in recent years (Hwang et al. 2019).

Metabolic disorders

Metabolic disorders such as diabetes might have ROS and RNS interaction. Diabetes is a disease characterized by insulin production dysfunction or insulin resistance that led to a raise in blood glucose levels. The glucose autoxidation on constant hyperglycemia causes superoxide radical (O,⁻) and hydrogen peroxide (H₂O₂) generation that mediates b-pancreatic cell autoimmune attack and negative regulation of cellular antioxidant systems aggravating disease symptoms (Newsholme et al. 2016). Some complications of diabetes may have symptoms aggravated by ROS and RNS. In diabetic retinopathy, for example, there is a positive correlation between ROS and retinopathy events such as the increase of vascular permeability, retinal cell apoptosis and death, and retinal vascular thickness reduction by NO removal (Lu et al. 2018). In addition, a recent work observed that REDD1 protein is the regulator of the feedback loop responsible for retinal damage mediated by ROS and visual dysfunction development in retinopathy (Miller et al. 2019).

The published literature evidence that 70% of deaths in diabetes are due to angiopathy. A study using endothelial cell culture observed that cells cultured with high glucose presented ROS accumulation as a necessary factor in

diabetic angiopathy development (Xu et al. 2019a). Similarly, it was found downregulation of synthase nitric oxide enzymes in diabetic aortas might be the mechanism of vessel dysfunction on diabetic angiopathy (Liu et al. 2017). ROS are still reported as important supporters in persistent inflammation on post fractured bone healing by NLRP3 inflammasomes recruitment (An et al. 2019), moreover, ROS are involved in osteoclasts differentiation by RANKL/ RANK/OPG route expression and osteoclasts efferocytosis inhibition favoring osteoporosis in a diabetes context (Hendrijantini et al. 2019). It is of interest to report that other diabetes related-diseases such as obesity also might be influenced by reactive species. Several studies in this field support ROS role on adipose tissue characteristic inflammation as a response to NLRP3 inflammasome lineage recruitment by NADPH oxidase overregulated ROS production and caspase-1 activation route (Engin 2017). A study carried out by Sindhu et al. (2018) reported that ROS could mediate overexpression of TRL10 inflammation marker in obesity of Type-2 diabetes patients.

Pulmonary diseases

There are a few recent pieces of evidence about ROS involvement in pulmonary diseases. In allergic asthma and rhinitis, the determinant role of ROS is on the inflammation process by interleukin gene expression, as well as mucus hypersecretion, airflow restrainment due to smooth muscle contraction, bronchial hyperresponsiveness, and oxidative damage to the accessory structures of epithelial cells like cilium (Shi et al. 2018, Wang et al. 2019). ROS can also damage alveolar-capillary barrier tightjunctions proteins contributing to pulmonary edema and acute lung injury progression and development (Sokolowska et al. 2019). Moreover, extravascular liquid in the lungs has been pointed out as an important condition of acute respiratory distress syndrome (ARDS) where ROS contributes to interstitial liquid accumulation by epithelium (Kellner et al. 2017).

A study investigated the action of bleomycininduced oxidative stress on pulmonary fibrosis by induction of differentiation from pericytes to myofibroblasts (Andersson-Sjöland et al. 2015). In addition, other recent studies (Zhou et al. 2018, Yang et al. 2017b) also demonstrated that ROS mediated: increased inflammatory response, various degrees of edema, and hydroxyproline increase due to characteristic collagen deposition on pulmonary fibrosis. Environmental studies observed the dangerousness of photochemical generated ozone to the pulmonary environment. Ozone causes airway cell damage and disturbs gas exchange in alveoli triggering emphysema development and progression by alveoli destruction (Xu et al. 2019b).

Cardiac diseases

The high consumption of oxygen by cardiomyocytes and upper regulation of NOX family enzymes contribute to cardiac damage and heart dysfunction by ROS intoxication (Panth et al. 2016). The involvement of ROS in cardiac diseases is mainly associated with those triggered by ischemia-reperfusion (Turillazzi et al. 2017). A study using aged rats has proved that occur an imbalance in mitochondrial potential and proton gradient that leads to high production of ROS, after ischemic injury (Rancan et al. 2018). It is also believed that ROS and RNS interact with non-selective mPTP (Mitochondrial Permeability Transition Pore) pore opening that leads to molecules influx and irreversible damage to heart tissue (Cadenas 2018).

A study using isoprenaline hydrochloride as a heart failure inducer proposes mPTP pore opening by ROS as a contributive mechanism of cardiac damage aggravation (Odinokova et al. 2018). In relation to reactive species' action on vascular conditions they may cause endothelial dysfunction by NO removal, vessel smooth muscle hypertrophy, collagen production, and fibronectin deposition stimulation favoring arterial hypertension development and aggravation (Pinheiro & Oliveira-Paula 2019). In addition, experiments with animal models have associated alcohol with endothelial NO removal and hooked up to artery resistance and endothelial inflammatory cytokines recruitment (Simplicio et al. 2016).

As well as in other tissues, in cardiac muscle inflammation, ROS participation on NLRP3 inflammasome recruitment, interleukin secretion, and NF-kB route activation is reported (Guo et al. 2018b, Zhang et al. 2018b). However, there is not paramount evidence on acute heart failure mediated by ROS, there is a slight relationship with acute heart failure diseases as myocardial infarction, for example (Yan et al. 2019). A study using guinea-pig hearts reported an elevation of non-designated ROS in cardiomyocyte tissue (Dev et al. 2018). Moreover, it was demonstrated that oxidative stress via the calmodulin-kinase protein II-dependent route contributes to acute cardiac failure atrial fibrillation (Yoo et al. 2018).

ANTIOXIDANTS: ORIGINS, CHEMISTRY OF ACTION AND IMPORTANCE

As firstly proposed by Haliwell & Gutteridge (1995), an antioxidant is defined as any substance able to suppress, prevent, or delay substrate oxidation under low concentrations. The mechanisms of action include Prevention, stopping radical formation chain reactions and their intermediates, Scavenger, promoting unstable molecules exhaustion cleaning up them, Reparatory, restoring native conformation

ACTINOBACTERIA AS SOURCE OF ANTIOXIDANT COMPOUNDS

of molecular structures through catalytic or non-catalytic ways, or Genetic, stimulating genic expression of enzymes related to cellular antioxidant capacity (Nikki 2010, Kusumawati & Indrayanto 2013). Huang et al. (2005) stated that the chemical reactions behind antioxidant mechanisms are based on two main mechanisms of energy transfer as follows: hydrogen atom transfer (HAT) or oxidoreduction through single electron transfer (ET).

According to the aforementioned, HATbased antioxidants are those with the ability to form non-reactive compounds by single dissociation and ionization of hydrogen with unstable molecules. On the other hand. ETbased antioxidants reduce oxidized compounds interacting with molecules' functional groups (Karadag et al. 2009). Several methodologies were developed to determine the antioxidant potential of compounds using the degree of color and absorbance as a principle of measurement (Figure 2). To determine a substance's antioxidant potential the most diffused methods may request different amounts of oxidant and antioxidant and can be expressed by scavenger potential or equivalent concentration opposite to a molecule with known antioxidant capacity (Huang et al. 2005, Karadag et al. 2009).

The most used methodologies for antioxidant measurement are ABTS⁺⁺ (2,2'-azinobis (3-ethylbenzothiazoline-6-sulfonic acid) cationic radical, DPPH (2,2-diphenyl-1picrylhydrazyl), MO⁶⁺ (Phosphomolydbenum Assay), ORAC (Oxygen Radical Absorbance Capacity), CUPRAC (Cupric Reducing/Antioxidant Capacity), FRAP (Ferric Reducing Antioxidant Capacity), FRAP (Ferric Reducing Antioxidant Capacity), H₂O₂ (Hydrogen Peroxide Assay), NO⁻ (Nitric Oxide Assay), OH⁻ (Hydroxyl Radical Assay), and O⁻₂ (Singlet Oxygen Assay) (Kusumawati & Indrayanto 2013). It is important to highlight the methods CUPRAC and FRAP due to several antioxidants might display a double-edged

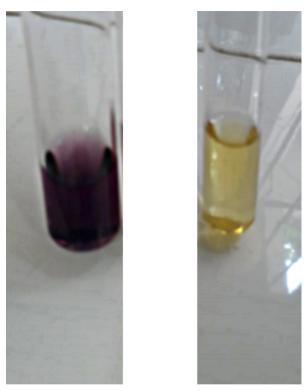


Figure 2. Antioxidant assay by DPPH (Negative control (left): DPPH + methanol; Positive control (right): DPPH + ascorbic acid).

sword role acting as both antioxidants molecules and heavy metals removing agents (Huang et al. 2005).

Antioxidant compounds may have different natures: synthetic (Figure 3), natural (Figure 4), or natural-like. The first is derived from a synthetic chemical process. The second are metabolites or products from microbes, animals, or plants. The third are those mirrored to a natural antioxidant and chemically synthesized due to molecular structure reproducibility with activity maintenance (Carocho et al. 2014). Although synthetic antioxidants are more efficient than natural, there is a lack of information about long-term usage, being also associated with allergic processes (Zaknun et al. 2012) and involvement in carcinogenesis events (Carocho et al. 2014). In contrast, natural antioxidants are preferable by consumers (Karadag et al. 2016),

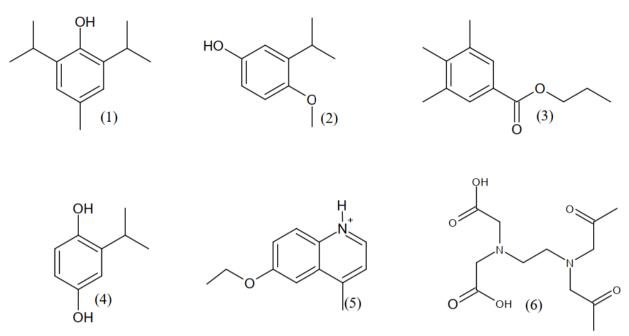


Figure 3. Molecular structures of industrial synthetic antioxidants. (1) Butylated hydroxytoluene (BHT), (2) butylated hydroxyanisole (BHA), (3) propyl gallate (PG), (4) *tert*-butylhydroquinone (TBHQ), (5) ethoxyquin and (6) ethylenediaminetetraacetic acid (EDTA).

safer (Torres-Fuentes et al. 2015), less reactive, more compatible with human metabolism (Akbarirad et al. 2016), and without toxic limit of consumption (Lorenzo et al. 2018).

The main antioxidant application is made through incorporation into product formulations. In dermo-cosmetics, may have anti-aging action stimulating collagen production, skin integrity, elasticity, and UV light blockage (Montenegro 2014). In the food industry, antioxidants are used as conservators and oxidation delayers, especially on oil peroxidation and butter rancidity prevention (Silva & Lidon 2016). In the pharmaceutical field, antioxidant compounds incorporated into food supplements act as nutraceuticals (Nirmala et al. 2018). Recent findings from the literature have shown antioxidants with anti-inflammatory properties (Gomes-Rochette et al. 2016) and potential candidates to help to treat cancer (Atici et al. 2018), metabolic syndrome (Gregório et al. 2016), neurodegenerative diseases (Amato et al. 2019), chronic diseases (Abdali et al. 2015), and cardiovascular diseases (Amit et al. 2015).

ACTINOBACTERIA

Actinobacteria comprise a diverse and heterogeneous group of gram-positive filamentous bacteria with highly conserved genomic content due to the predominance of G+C genomic combination (Nouioui et al. 2018). These bacteria are mainly aerobic and mesophilic, however, some of them are able to grow under extreme conditions of oxygen absence, pH, temperature, and salt concentration. The microscopic morphology of Actinobacteria colonies is similar to fungi, endowed of grouped thin hyphae to form a characteristic mycelium (Barka et al. 2016, Araújo-Melo et al. 2017). Macroscopically, Actinobacteria colonies may have various colors, dried aspect, crinkled appearance, and strong adherence to the surface when cultured on agar plates. The taxonomy of

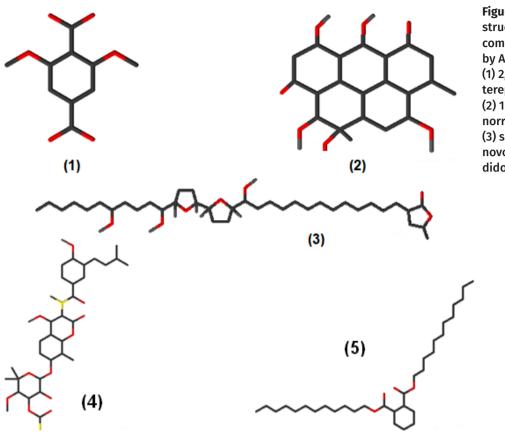


Figure 4. Tridimensional structures of antioxidant compounds produced by Actinobacteria. (1) 2,6-dimethoxyterephthalic acid, (2) 1-hydroxynorresistomycin, (3) squamocin, (4) novobiocin and (5) didodecylphthalate.

this group was based on cell wall content but currently is also focused on 16S and 23S rRNA sequences and different genes (Nouioui et al. 2018, Araújo-Melo et al. 2017).

Actinobacteria became known from the antimicrobial properties of secondary metabolites. The first Actinobacteria antibiotic discovered was streptomycin produced by *Streptomyces griseus* in 1943, followed by the discovery of other bioactive compounds over the years such as chemotherapeuticals, antifungals, immunomodulators, pesticides, and antihelmintics useful in medical routine, agricultural, and veterinarian fields. It is proposed that the metabolic plasticity of Actinobacteria is intrinsically related to this sortive amount of different compounds (Manteca & Yagiu 2018). The adaptation of Actinobacteria to different environments such as glaciers, mangroves, plants, marine environments, deserts, and animals may be also another explanation for a wide range of compound production (Araújo-Melo et al. 2017) (Figure 5).

The research for natural compounds remains under constant ascension being natural antioxidants a branch of this field. Many of the natural compounds produced by actinobacteria have demonstrated antioxidant properties under *in vitro* assays, however, there are few studies using *in vivo* protocols or induction models of oxidative stress. Here, we present a relevant cutting from published literature about how the Actinobacteria must be considered an important fountain of antioxidant molecules, with compounds belonging to the most different chemical classes such as phenols, alkaloids, pyrrolizidines, biphenyls, and other non-reported compounds being its antioxidant activity mostly

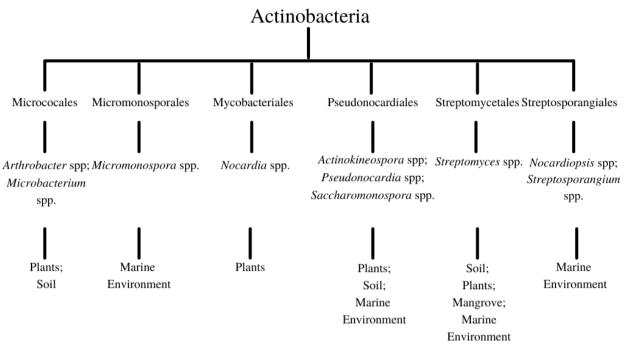


Figure 5. Different isolation environments of Actinobacteria.

related to the structural arrangements of these compounds (Tables II & III). These antioxidant compounds from actinobacteria may be a suitable alternative to the usage of synthetic compounds in the most varied areas whereas they can stabilize strong and weak oxidants.

ACTINOBACTERIA ISOLATED FROM DIFFERENT ENVIRONMENTS

Soil

Actinobacteria are part of the soil microbiome having an important contribution to soil health. These bacteria act on organic matter decomposition, and nutrient cycling, for example (Bhatti et al. 2017). Some similar studies as the one performed by Rani et al. (2018) show up the microenvironment of plant rhizospheres as the pronest opposite to other soil regions for Actinobacteria isolation since the increased amount of nutrients around roots. Other extreme soil environments as those from desert and glaciers have also been reported as an isolation environment of Actinobacteria (Reis-Mansur et al. 2019). Thus, the soil is still considered the first choice for Actinobacteria isolation.

In spite of soil Actinobacteria exploration to find potential antimicrobial agents and other bioactive compounds, a new approach of soil Actinobacteria as a reservoir of antioxidant molecules has emerged and is ascending in the last years, turning the soil also an inexhaustible source of actinobacteria with potential for production of antioxidant molecules. Using this approach Rani et al. (2018) when evaluating Camellia senensis rhizosphere, isolated Streptomyces cellulosae TE217 where its ethyl acetate crude extract post-fermentation showed scavenger potential equal to 91.88%, 78.47%, and 82.08% opposite to ABTS^{**}, DPPH, and radical superoxide methods at 5 mg/mL. Additionally, the phenol compounds found in the crude extract were also able to reduce molybdenum and ferrous ion being equivalent to 76.93 and 231.96 mg of ascorbic acid/100 mg of extract dry weight, respectively.

Chemical Classes	Producer Actinobacteria	Assay Methods	References
Phenols			
4-methyl-benzoic acid	Saccharomonospora oceani VJDS-3	DPPH, ABTS [™]	(Indupalli et al. 2018)
2,6-dimethoxy terephthalic acid	Streptomyces sp. YIM66017	DPPH	(Zhou et al. 2014)
1-hydroxy-1-norresistomycin	Streptomyces variabilis KP149559	DPPH, H ₂ O ₂ , OH, FRAP	(Ramalingam & Rajaram 2016)
Phenol, 2,4-bis(1,1-dimethylethyl)	Arthrobacter sp.	DPPH, FRAP, MO ⁶⁺	(Akshatha et al. 2016)
2,6-dimetoxyphenol	Streptomyces griseorubens NBRC12780	DPPH	(Sengupta et al. 2015)
Phenol, 2,6-bis (1,1-dimethylethyl) -4 - [(4-hydroxy-3,5- dimethylphenyl) methyl	<i>Streptomyces</i> sp. strain KAV2	ABTS ^{+*} , DPPH, O₂ ⁻ , MO ⁶⁺ , FRAP	(Keerthana et al. 2019)
Phenol, 2,2'-methylenebis [6- (1,1-dimethylethyl) -4-methyl	Streptomyces MUM265	ABTS [™] , DPPH, O ₂ ⁻ , Metal chelating	(Tan et al. 2019a)
Alkaloids			
Diazepinomicine	Micromonospora sp. RV115	FRAP, H ₂ O ₂ cell-free system	(Abdelmohsen et al. 2012)
Prodigiosin	Streptomyces sp. WMA- LM31	DPPH, Lipid peroxidation, Protein oxidation	(Sajjad et al. 2018)
N-acetyltyramine	Actinokineospora sp.		(Heidari &
N-acetyltryptamine	UTMC 968	DPPH	Mohammadipanah 2018)

Radhakrishnan et al. (2016) isolated Streptomyces sp. D25 from Thar Desert soil, North of India, able to produce an orangeyellow pigment with scavenger potential of 35.65% and 96.19% when assayed by DPPH and nitric oxide at 500 µg/mL. The work of Kaur & Arora (2017) describes the usage of chloroform and ethyl acetate in the extraction of bioactive molecules from Actinobacteria named OS-6 and TES-25; ethyl acetate proved the best choice. It was detected an IC₅₀ of 49.89 µg/mL and 121.51 µg/mL when applied DPPH method and 2.63 µg/mL and 46.61 µg/mL in ABTS⁻⁺ assay, besides that reducing power of molybdenum ion was equivalent to 83.7 and 74 mg of ascorbic acid.

The search for antioxidant molecules from soil Actinobacteria has yielded the isolation of

pure bioactive compounds such as a prodigiosin produced by Streptomyces sp. WMA-LM31 isolated from Marwart Lakki District Desert. Pakistan, was able to inhibit lipid peroxidation at 25.4% as well as protein autoxidation at 54.82% and scavenging DPPH radical at 62.5% (Sajjad et al. 2018). Heidari & Mohammadipanah (2018) also reported the production of two alkaloids by rare Actinobacteria Actinokineospora sp. UTMC68 isolated from the rhizospheric region on Isfahan Province, Iran, with antioxidant activities and respective IC $_{50}$ of 64.7 and 131.3 μ g/mL when exposed to DPPH radical. The work developed by Reis-Mansur et al. (2019) describes the production of a carotenoid with possible strong antioxidant capacity from Microbacterium sp. LEMMJ01 isolated from Arctic soil. Furthermore, the extracts of Actinobacteria isolated from agricole soil in Egypt proved to be able to upregulate cellular antioxidant system enzymes with reduced activity in aflatoxin-treated animals (El-Nekeety et al. 2017).

Plants

The endophytic microenvironment is indicated as a promising niche for the isolation of nonrelated compounds with pharmaceutical potential. The Actinobacteria have already been isolated from herbaceous, arboreal, ornamental, and medicinal plants being frequent the isolation of rare genus as Nocardia, Pseudonocardia, and Actinomadura (Tanvir et al. 2016). The first Actinobacteria isolated from vegetal tissues were classified as belonging to the Frankia genus (Araújo-Melo et al. 2017). The knowledge about endophytic Actinobacteria has increased, however, this niche is still considered untapped or misexplored. Regarding endophytic Actinobacteria and pharmaceutical potential bioactive compounds, antitumor, antidiabetic, antimicrobial, and antioxidant properties are described. This wide potential can be associated with the stressful conditions inside vegetal tissue which Actinobacteria need to adapt to survive (Sengupta et al. 2015).

The Actinobacteria Streptomyces sp. BO-07 isolated from Boesenbergia rotunda L. radicular tissue was able to produce two compounds belonging to phenylbenzenes with IC_{50} equivalent to 85.84 µg/mL and 88.26 µg/ mL by DPPH method (Taechowisan et al. 2017). In the work developed by Akshatha et al. (2016) strains of Streptomyces globosus (JQ926176) and Arthrobacter sp. (JQ926171) were isolated from four medicinal plant stems, inflorescences, and leaves. The 21 days post-fermentation metabolic liquid extract showed reducing power of ferrous ion of 40.44 µM and 52.44 µM/g of dry weight extract. Equivalently, Fahmy et al. (2016) observed scavenging potential varying from 62 to 85% when evaluated antioxidant potential of ethyl acetate extract from four Actinobacteria isolated from *Asphodelus tenuifolius*.

In a similar investigation, Tanvir et al. (2016) isolated two rare Actinobacteria named Nocardia caishiiiensis and Pseudonocardia carboxydivorans from Ageratum conyzoides and Sonche oleraceus L. They reported an IC₅₀ of 0.552 and 0.670 mg/mL by DPPH. From the evaluation of roots, leaves, and stems of Racomitrium ellipticum, Passari et al. (2017) observed that bioactive molecules present in methanolic extract from Streptomyces olivaceus BPSAC77 and Streptomyces thermocarboxydus BPSAC147 isolated from these parts scavenged 50% of DPPH at 43.2 and 75.4 µg/mL, respectively. Curiously, a carboxyl derivative produced by Streptomyces sp. YIM66017 isolated from Alpinia oxyphylla returned an impressive IC₅₀ of 4.16 µg/mL when applied DPPH method (Zhou et al. 2014). Two acetogenins were isolated from the metabolic liquid of Streptomyces sp. VE2 recovered from Vernonia cinerea, which showed antioxidant potential with the respective IC_{ro} 86.76 and 58.2 µg/mL (Taechowisan et al. 2016).

Mangrove

Brazil holds approximately 7% of global mangrove coverage, which is around 11.942 km². Mangrove is a characteristically hypoxic, highly saline, and malodorous environment due to continuous organic matter decomposition. Mangroves are considered a diversified environment and display a pivotal role in primary productivity, vegetal stabilization, and costal line balance (Ferreira & Lacerda 2016). The exploration of extreme environments as mangrove has been considered an escape route for the isolation of non-described compounds with potential in the pharmaceutical field once depletion of common sources as terrestrial (Sengupta et al. 2015).

Table III. Iminosugars and other antioxidant	compounds produced by Actinobacteria.

Chemical Classes	Producer Actinobacteria	Assay Methods	References
Pirrolizidines			
5- (2,4-dimethyl benzyl) pyrrolidine-one	Streptomyces VITSVK5 spp.	DPPH, MO ⁶⁺	(Sarauv & Kannabiran 2012)
Pyrrolo [1,2-a] pyrazine-1,4-diona, hexahydro	Streptomyces mangrovisoli; Streptomyces sp. MUSC14	ABTS [™] , DPPH, SOD, Metal chelating	(Ser et al. 2015, Kemung et al 2020)
Pyrrolo [1-a] pyrazine-1,4-dione, hexahydro-3- (2-methylpropyl) –	Streptomyces sp. S2A	ABTS ⁺⁺ , DPPH, FRAP, Metal chelating	(Siddharth & Vittal 2018)
Pyrrolo [1,2-a] pyrazine-1,4-dione, hexahydro-3- (2-methylpropyl)	Streptomyces sp. VITMK1	DPPH, FRAP, NO	(Manimaram & Kannabiran 2017)
Other Compounds			
5-amino-2 (6- (2- hydroxyethyl) -3- oxononyl) - cyclohex-2- enone 8 (aminomethyl) -7-hydroxy-1- (1-hydroxy-4 - ((hydroxymethoxy) -2,3-dimethyl (butyl-2-methyl-	Streptomyces coelicoflavus BC01	DPPH, FRAP, MO ⁶⁺	(Rao et al. 2017)
dodecahydrophenanthren-9 (1H) -one 1 ((E) -1-ethylhex-1em-1-yl) 2 - ((E) -2-ethylidenehexyl) cyclohexane- 1,2-dicarboxylate			
Angeloline A	Streptomyces sp. SBT345	Cell-free system by NQO	(Cheng et al. 2016)
(Z) -1 - ((hydroxypenta-2,4-dien-1- ie) oxy) anthracene-9-10-dione	Nocardiopsis alba	DPPH, FRAP, MO ⁶⁺	(Janardhan et al. 2014)
6-hydroxy-4-2 ', 3', 4 "-tetramethyl- p-terphenyl			
4,7-bis (4-methoxyphenyl) -6-hydroxy-5-methoxybenzo [d] thiazole (benzothiazole)	Nocardiopsis gilva YIM90087	ABTS", DPPH, O ₂	(Tian et al. 2013)
Novobiocin			
Cyclo (L-Ile-L-Pro)			
Palmitic acid, betamonoglyceride	Streptomyces globosus	DPPH, FRAP, MO ⁶⁺	(Akshatha et al. 2016)
Actinosporine C and D	Actinokineospora sp. strain EG49	FRAP, H ₂ O ₂ cell-free system	(Grkovic et al. 2014)
3-hydroxy-5-methoxy-3,4- methylenedioxybiphenyl	Standard DO 07	22211	(Taechowisan et al. 2017)
3-hydroxy-5-dimethoxy-3,4- methylenedioxybiphenyl	Streptomyces sp. BO-07	DPPH	
Didodecyl phthalate	Nocardiopsis sp. VITSRTB DPPH, FRAP		(Singh at al 2012)
Hexacosanol acetate			(Singh et al. 2013)
2,5-bis (1,1-dimethyethyl)	Streptomyces sp. MUM212 ABTS", DPPH, O ₂ , Metal chelating, Lipid peroxidation, H ₂ O ₂ cell-free system		(Tan et al. 2017)
2,2'-methylenebis [6- (1,1-dimethylethyl) -4-methyl]			

Rolidecin B	Streptomyces sp. VE2 DPPH	(Taechowisan et al. 2016)	
Squamocin			
Hexacirines	Streptosporangium sp.	DPPH, OH', O_2^-	(Gao et al. 2018)
Bis- (2-ethylhexyl) phthalate	Nocardiopsis sp. SCA21	ABTS ^{+*} , DPPH, Metal chelating	(Siddharth & Rai 2019)
Saccharomonopirone A	Saccharomonospora sp. CNQ-490	ABTS [∵] , DPPH	(Yim et al. 2017)

Table III. Continuation

In this context, several authors have believed that mangrove dynamics as tidal gradient, salinity floating, radiation exposure, and other environmental conditions may be involved in the activation of metabolic pathways for the production of non-reported compounds (Tan et al. 2017).

Surprisingly, an acid-like compound was isolated from the metabolic liquid of Saccharomonospora oceani VJDS-3 and showed an excellent antioxidant activity of 73.88% and 99.74% at very low concentrations of 25 µg/ mL and 20 μ g/mL when applied the methods of DPPH and ABTS[™] (Indupalli et al. 2018). The finding of Rao et al. (2017) corresponds to the molecular structure of three compounds from Streptomyces coelicoflavus BC01. These compounds showed a scavenger percentage of 63%, 61.92%, and 52.39% in the DPPH assay. The work carried out by Tan et al. (2017) reported the scavenger potential of 35.98%, 67.96%, and 79.23% from compounds present in the methanolic extract of Streptomyces sp. MUM292 isolated from mangrove forest soil at Kuala Selangor when assayed through DPPH, ABTS⁺, and radical superoxide. It is still reported the reducing power of these compounds opposite to molybdenum ion as equivalent to ascorbic acid 60.0; 59.53; and 56.66 µM/mL. In addition to reducing power perspective, ferrous ion was also evaluated returning ascorbic acid equivalence of 67.83; 67.0; and 60.0 µM/mL.

Tan et al. (2019a) believed that phenol compounds present in the methanolic extract

of Streptomyces sp. MUM265 were able to inhibit dismutase superoxide in 56%, act as chelating metal agents, with a percentage of 46.02% besides present a scavenger potential varying from 42.33% to 88.5% at the unique concentration of 4 mg/mL. Equally, Tan et al. (2019b) observed that phenol compounds, pyrazine derivatives, and cyclic dipeptides produced by Streptomyces sp. MUM273b showed antioxidant potential with promising applications in the dermo-cosmetics field due to its potential for radiation blocking which causes keratinocytes' death. To evaluate the antioxidant capacity of Streptomyces olivaceus MSU3 ethyl acetate crude extract was assayed by many assay methods the following inhibition percentages of 62%, 87%, 32.5%, 47.99%, and 33.2% were encountered by DPPH, total antioxidant of molybdenum ion, hydroxyl radical, and nitric oxide radical appliance (Sanjivkumar et al. 2016).

Marine environment and related animals

The marine environment is dynamic since receives influences from many outside factors. It is a diverse ecosystem fulfilled by the most different organisms. The importance of the marine environment is closely related to primary production and food withdrawal. Certainly, this environment also holds an unexplored resident microbial diversity either by symbiosis or ground housing (Beygmoradi & Homaei 2017). The tolerance of microbes to salinity and marine stressful conditions may have headed the unprecedented exploration for new drugs whereas depletion of terrestrial sources and lack of novelty is a real threat (Lahoum et al. 2015). Recently, the marine environment has provided a wide variety of unheard-of compounds with promising pharmaceutical applicability, including antioxidant molecules (Beygmoradi & Homaei 2017). In this section, it is proposed to show a small fragment from published literature about antioxidant compounds production by Actinobacteria isolated from marine sediment and related organisms.

One of the isolated antioxidant compounds from marine sources is a new chlorinated quinolone from Streptomyces sp. SBT345 isolated from a Mediterranean marine sponge, which demonstrated antioxidant potential when evaluated under a free-cell antioxidant system (Cheng et al. 2016). Ramalingam & Rajaram (2016) isolated the strain *Streptomyces variabilis* KP149559 from mucus samples of Acropora formosa coral and purified a hydroxylated compound produced by this Actinobacteria with scavenger percentage varying from 75 to 82% at a concentration of 0.5 mg/mL. In the work managed by Manimaran & Kannabiran (2017) it was possible to observe that *Streptomyces* sp. VITMK1 produced a diketopiperazine that scavenged more than 70% of DPPH and nitric oxide radicals at 500 µg/mL. Ravi et al. (2017) working with coastal region sediments of Rameswaram and Dhanushkodi, India, isolated the Gancidine W from chloroformic extract of Streptomyces paradoxus VITALK03 which at 1 mg/mL exhibited antioxidant percentage of 34%, 48%, and 61.5% when evaluated by DPPH, ABTS^{`+}, and radical superoxide.

The crude extracts can be considered the column of antioxidant search. A derivative from pyrrolopyrazines was partially purified from ethyl acetate crude extract of the strain *Streptomyces* sp. S2A isolated from Mannar Gulf sediment samples by Siddharth & Vittal (2018) and pointed it as one of the responsible for the antioxidant activity of 56.5% and 42.4% present in the crude extract. In addition, the extract showed a metal-chelating potential of 59.98% and ferrous ion reducing power with 0.248 of absorbance. Based on the knowledge of internal microbiome existence. Gozari et al. (2018) isolated Actinobacteria strains from the stomach content of sea cucumber (Holothuria *leucospilota*); three strains showed an IC₅₀ varying from 211 to 822 μ g/mL when assayed by stable radical DPPH. Dholakyia et al. (2017) isolated Streptomyces variabilis RD-5 from marine soil of Khambhat Gulf Region, India, acetoethylic extract demonstrated good antioxidant properties with scavenger potential equivalent to 82.86% at DPPH, 89% of chelating activity, and inhibition potential of hydrogen peroxide higher than 85% at 5 mg/mL.

CONCLUSIONS

In this review, we approached the misexplored antioxidant potential of Actinobacteria compounds. Unquestionably, antioxidants compounds have proved being essential to health, since they can restore redox cell balance and avoid cell damages caused by the excess of free radicals. Here, we showed that compounds from Actinobacteria are promising given the surprisingly antioxidant capacity of some of them as well as the discovery of non-reported antioxidant compounds by Actinobacteria. As presented, part of these compounds is strong candidates for synthetic compounds' usage replacement whereas they showed antioxidant capacity without toxicity or damage to cells. Furthermore, some of them may be applicable in the feeding industry such as some antioxidant pigments, which can perform a double role of dyeing and slowing food oxidation. Regardless

that, we would still like to raise an observed limitation in this field. The absence of studies testing Actinobacteria antioxidant compounds using *in vivo* protocols, and in the same way to encourage more works to fill up this gap on the antioxidant exploration of Actinobacteria metabolites in order to assure the safety of these composites to human usage in a brief future.

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This work was carried out through collaboration among all authors. Author THBO and LCBBC managed the literature searches and wrote the manuscript. Authors NBG and LAOS participated in revision of this article and contributed with comments. Authors THBO and LCBBC performed revision of manuscript and English translation. Authors LAOS and LCBBC designed, supervised and managed the study performed. All authors read and approved the final manuscript.

