




# Biosurfactants produced by *Serratia* species: Classification, biosynthesis, production and application

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## Abstract

Biosurfactants are surface-active molecules that are synthesised non-ribosomally by a wide range of microorganisms including bacteria, yeast and filamentous fungi. The bacterial genus *Serratia* is gaining international interest, as biosurfactants produced by this genus have emerged as a promising source of antimicrobial, antifouling and antitumour compounds that possess emulsification and surface activity. Various species of *Serratia* have been identified as biosurfactant producers, including *Serratia marcescens*, *Serratia rubidaea* and *Serratia surfactantifaciens*. Members of the *Serratia* genus have been reported to principally produce two classes of biosurfactants, namely lipopeptides and glycolipids. Lipopeptides produced by *Serratia* species include serrawettins and stephensiolides, while identified glycolipids include rubiwettins and rhamnolipids. This review will primarily focus on the classification of biosurfactants produced by *Serratia* species and the genes and mechanisms involved in the biosynthesis of these biosurfactant compounds. Thereafter, an indication of the primary growth conditions and nutrient composition required for the optimum production of biosurfactants by this genus will be outlined. An overview of the latest advances and potential applications of the biosurfactants produced by *Serratia* in the medical, pharmaceutical, agricultural and petroleum industries is also provided.

**Keywords** Biosurfactants · *Serratia* · Application · Production · Biosynthesis

## Introduction

Biosurfactants play important physiological roles in cellular metabolism, motion and the defence mechanisms of microorganisms (Banat et al. 2014). Accordingly, various genera, such as *Acinetobacter*, *Alcanivorax*, *Arthrobacter*, *Bacillus*, *Candida*, *Corynebacterium*, *Flavobacterium*, *Lactobacillus*, *Mycobacterium*, *Nocardia*, *Pseudomonas*, *Rhodococcus*, *Rhodotorula*, *Serratia*, *Streptomyces* and *Thiobacillus*, amongst others, secrete various classes of biosurfactants as secondary metabolites (Rahman and Gakpe 2008; Zhang et al. 2012; Santos et al. 2016). The production of biosurfactants by microorganisms is often triggered by the presence of hydro-

phobic substrates and aids in the survival of these microorganisms in nutrient poor or highly contaminated environments (Bodour et al. 2003). This is achieved by increasing the bioavailability of nutrients, thus promoting the uptake and metabolism of less soluble substrates (Fiechter 1992). Furthermore, it has been hypothesised that anionic biosurfactants are able to protect microbial cells by forming complexes with positively charged toxic heavy metals present in the environment (Ron and Rosenberg 2001). Therefore, biosurfactant-producing microorganisms are routinely isolated from highly contaminated sites, such as metal- or hydrocarbon-contaminated soil and water environments, and wastewater treatment plants (Bodour et al. 2003; Ndlovu et al. 2016).

Of the numerous biosurfactant-producing genera isolated from different environments, biosurfactants produced by members of the *Serratia* genus are gaining increased scientific interest as they have been shown to display emulsification, surface, antifouling, antitumor and antimicrobial activity (Escobar-Díaz et al. 2005; Dusane et al. 2011; Nalini and Parthasarathi 2014; Su et al. 2016). They are also able to modify the hydrophobicity of the cell surface, which plays an important role in the adhesion of these bacteria to various

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surfaces and contributes to enhancing the surface spreading of bacteria in nutrient-poor environments (Bar-Ness et al. 1988; Wei et al. 2004; Matsuyama et al. 2011; Su et al. 2016). Various biosurfactant-producing *S. marcescens*, *S. rubidaea* and *S. surfactantfaciens* strains as well as *Serratia liquefaciens* (recently reclassified as a *S. marcescens* strain) have subsequently been isolated from hydrocarbon-contaminated soil and rhizosphere soil, surface water, marine environments and wastewater treatment plants (Matsuyama et al. 1985, 1990; Lindum et al. 1998; Anyanwu et al. 2010; Dusane et al. 2011; Nalini and Parthasarathi 2013, 2014; Ndlovu et al. 2016; Su et al. 2016; Almansoori et al. 2017).

*Serratia* species predominantly produce various glycolipids and lipopeptides (Desai and Banat 1997; Banat et al. 2010; Santos et al. 2016), which have been reported to display antibacterial, antifungal and antiprotozoal activities (Kadouri and Shanks 2013; Su et al. 2016; Ganley et al. 2018). In addition, glycolipids produced by *Serratia* species display biofilm disrupting and antiadhesive activity against bacterial and fungal strains (Dusane et al. 2011). The glycolipids and lipopeptides produced by this genus are thus of interest due to their potential biomedical and therapeutic applications (Cameotra and Makkar 2004). Moreover, the robustness and environmentally friendly nature of biosurfactant compounds in general allows for several potential applications in a number of different industrial fields, such as the petroleum, agricultural, food, cosmetic and pharmaceutical industries (Aparna et al. 2012).

## The genus *Serratia*

*Serratia* species are Gram-negative, facultative anaerobic bacteria that belong to the Enterobacteriaceae family (Su et al. 2016). The genus is comprised of 18 species, including the type strain *Serratia marcescens* which is used as a biological marker because of its easily distinguishable red colonies (Khanna et al. 2013; Su et al. 2016). Although *S. marcescens* was first assumed to be non-pathogenic, it was later found to be an opportunistic pathogen associated with nosocomial infections, such as urinary tract, respiratory tract, surgical wound and blood stream infections (Khanna et al. 2013). Other species within the *Serratia* genus such as *Serratia plymuthica*, *S. rubidaea* and *S. nematodiphila* are also capable of producing the non-diffusible red pigment, prodigiosin, during secondary metabolism (Su et al. 2016). Prodigiosin has been reported to display antimalarial, antibacterial, antifungal, antiprotozoal, antitumour and immunosuppressant activities (Stankovic et al. 2014). In addition to prodigiosin, various members of the genus *Serratia* are known to produce other secondary metabolites, such as biosurfactants, oocycin A, carbapenem, althiomycin,

bacteriocins and serratin (Foulds 1972; Srobel et al. 1999; Matsuyama et al. 2011; Gerc et al. 2012; Wilf and Salmond 2012; Luna et al. 2013). Although *Serratia* species produce a number of bioactive secondary metabolites, this review will focus on the glycolipid and lipopeptide biosurfactant compounds produced by members of this genus.

The primary lipopeptides produced by *Serratia* species include serrawettins (W1, W2 and W3) and stephensiolides (A to K) (Matsuyama et al. 1985; Dwivedi et al. 2008; Su et al. 2016; Ganley et al. 2018). *Serratia* species have also been reported to produce the glycolipids, rubiwettins (R1 and RG1) and rhamnolipids (Matsuyama et al. 1990; Nalini and Parthasarathi 2014). A few additional glycolipids, including a sucrose lipid, an arabinolipid and a glycolipid composed of a glucose attached to a palmitic acid, have also been detected (Pruthi and Cameotra 2000; Bidlan et al. 2007; Dusane et al. 2011). In addition, a study conducted by Ndlovu et al. (2016) isolated *S. marcescens* ST29 from a wastewater treatment plant sample and found the strain to contain genes encoding for the biosynthesis of both surfactin and iturin. However, the chemical characterisation of the biosurfactant compounds produced by this strain was not investigated.

## Lipopeptides produced by *Serratia* species

Lipopeptides represent a class of low molecular weight compounds composed of a hydrophilic peptide attached to a hydrophobic lipid or fatty acid (Banat et al. 2014). A wide range of lipopeptide structures have been identified which display variation in the length and conformation of the lipid moiety resulting in different homologues, while analogues exist due to variation in amino acid composition within the peptide moiety (Banat et al. 2014). As indicated, lipopeptides produced by *Serratia* species include serrawettin W1 (initially referred to as serratomolide A) with identified homologues serratomolide B to G (Wasserman et al. 1961; Matsuyama et al. 1985; Dwivedi et al. 2008; Zhu et al. 2018). In addition, studies have identified serrawettin W2 (and its analogues) (Matsuyama et al. 1992; Su et al. 2016) and serrawettin W3 has been partially characterised (Matsuyama et al. 1986). In addition to serrawettins, Ganley et al. (2018) reported the discovery of antimicrobial lipodepsipeptides known as stephensiolides A to K.

## Serrawettin family

Serrawettins are solely produced by members of the *Serratia* genus; serrawettin W1 is produced by strains of *S. marcescens*, serrawettin W2 is produced by *S. marcescens* and *S. surfactantfaciens* strains, and the partially characterised serrawettin W3 is also produced by a *S. marcescens* strain (Matsuyama et al. 1985, 1992; Matsuyama and Nakagawa

1996; Su et al. 2016). The first serrawettin was isolated in 1985 from a *S. marcescens* strain and was found to be identical to serratamolide A that was previously identified by Wasserman et al. (1961) (Matsuyama et al. 1985). The general structure of serrawettin W1 (also known as serratamolide A) includes a symmetric dilactone structure composed of two L-serine amino acids linked to two  $\beta$ -hydroxy fatty acids (comprised of 3-hydroxydecanoic acids) (molecular weight is 514.66 Da) (Fig. 1a) (Eckelmann et al. 2018). The diversity in the structure of serrawettin W1 (serratamolide A) exists due to the variation in the length of the fatty acid chain ( $C_8$  to  $C_{14}$ ) and the presence or absence of double bonds, resulting in homologues of serrawettin W1 (serratamolide A), namely serratamolide B to G (Dwivedi et al. 2008; Zhu et al. 2018). The molecular weight of the homologues range within 486.61 to 665.40 Da (Dwivedi et al. 2008; Zhu et al. 2018). In contrast to serrawettin W1, the general structure of serrawettin W2 includes five amino acids (D-leucine/isoleucine-L-serine-L-threonine-D-phenylalanine-L-isoleucine/leucine), bonded to a  $\beta$ -hydroxy fatty acid moiety (molecular weight of 731.93 Da) (Fig. 1b) (Matsuyama et al. 1992; Motley et al. 2016).

The diversity in the structure of serrawettin W2 results from variation at the first, second or fifth amino acid positions or the length of the fatty acid chain ( $C_8$  or  $C_{10}$ ), resulting in the detection of analogues with a molecular weight ranging from 703.3 to 759.3 Da (Matsuyama et al. 1986; Motley et al. 2016; Su et al. 2016). While the exact chemical composition of serrawettin W3 has yet to be elucidated, the cyclodepsipeptide was found to be composed of a fatty acid (one dodecanoic acid) and five amino acids, including threonine, serine, valine, leucine and isoleucine (Matsuyama et al. 1986). Serrawettins are described as non-ionic biosurfactants as they have no amino acid residues with ionic hydrophilicity (Matsuyama et al. 2011).

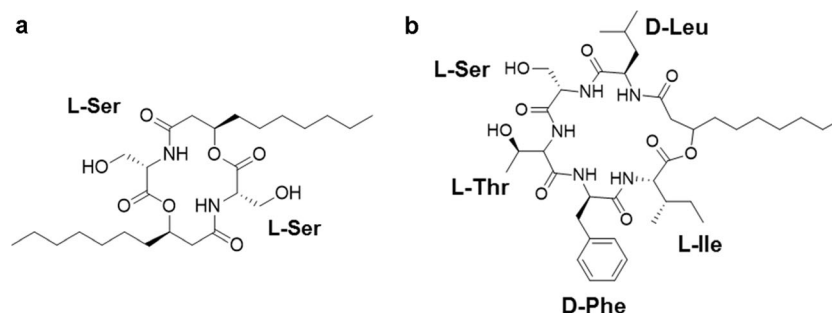
### Stephensiolide family

Stephensiolide lipodepsipeptides (A to K) are produced by a *Serratia* strain that was isolated from the midgut and salivary glands of an *Anopheles stephensi* mosquito (Ganley et al.

2018). Although stephensiolide mimics the overall structure of serrawettin W2 (as both structures are cyclic pentapeptides), the peptide sequence of the stephensiolides differs. The general chemical structure of stephensiolide includes five amino acids (threonine-serine-serine-valine/isoleucine-isoleucine/valine) attached to a long alkyl chain. The diversity in the structure of stephensiolides exists due to variation in the length of the acyl chain, variation in amino acid residues (presences of either isoleucine or valine at the fourth and fifth amino acid residues) or the presence or absence of a double bond in the fatty acyl chain, resulting in congeners of stephensiolide A to K (Ganley et al. 2018). This results in the molecular masses of the stephensiolides ranging from 599 to 695 Da depending on the congener (A to K). Although these lipopeptides have only recently been discovered, preliminary antimicrobial testing of the stephensiolide mixture (A to K) revealed activity against *Bacillus subtilis* and the malaria parasite, *Plasmodium falciparum* (Ganley et al. 2018).

### Glycolipids produced by *Serratia* species

Glycolipids are a class of low molecular weight biosurfactants, which are comprised of a hydrophilic carbohydrate attached to a hydrophobic aliphatic or hydroxyl-fatty acid (Shekhar et al. 2015). A number of structurally diverse glycolipid structures have been identified and are produced by a wide range of bacterial and fungal genera. A microorganism can produce homologues of the same glycolipid due to variation in the length and conformation of the fatty acid moiety, while the carbohydrate moiety is comprised of mono-, di-, tri- or tetra-saccharides (Banat et al. 2014). Previous studies have isolated glycolipids produced by *Serratia* species, including rubiwettin R1, rubiwettin RG1 and rhamnolipids (Matsuyama et al. 1990; Nalini and Parthasarathi 2014). In addition, a number of studies have detected glycolipids such as a sucrose lipid (Pruthi and Cameotra 2000), a glycolipid composed of glucose and palmitic acid (Dusane et al. 2011) and an arabinolipid (Bidlan et al. 2007).



**Fig. 1** The chemical structure of serrawettins **a** W1 [cyclo(D-3-hydroxydecanoyl-L-seryl)<sub>2</sub>] (adapted from Matsuyama et al. 1985) and **b** W2 [D-3-hydroxydecanoyl-D-leucyl-L-seryl-L-threonyl-D-phenylalanyl-L-isoleucyl lactone] (adapted from Su et al. 2016)

## Rubiwettin family

*Serratia rubidaea* ATCC 27593 is currently the only strain reported to produce rubiwettins and was found to produce both rubiwettin R1 and rubiwettin RG1 (Matsuyama et al. 1990). The general structure (undetermined carbohydrate moiety) and molecular weight of rubiwettin R1 have yet to be fully elucidated. However, a mixture of linked 3-hydroxy fatty acids comprised of major components, including 3-(3'-hydroxytetradecanoyloxy) decanoate and 3-(3'-hydroxyhexadecanoyloxy) decanoate, and minor molecular isomers have been identified and a proposed structure of the fatty acid moiety (Fig. 2a) was provided by Matsuyama et al. (1990). The general structure of rubiwettin RG1 (Fig. 2b) was also proposed and consists of  $\beta$ -D-glucopyranosyl 3-(3'-hydroxytetradecanoyloxy) decanoate minor fatty acid isomers (molecular weight of 576.77 Da) (Matsuyama et al. 1990). Therefore, RG1 was found to have a rhamnolipid-like glycolipid structure; however, rhamnose is substituted with a glucose moiety (Matsuyama et al. 1990).

## Rhamnolipid family

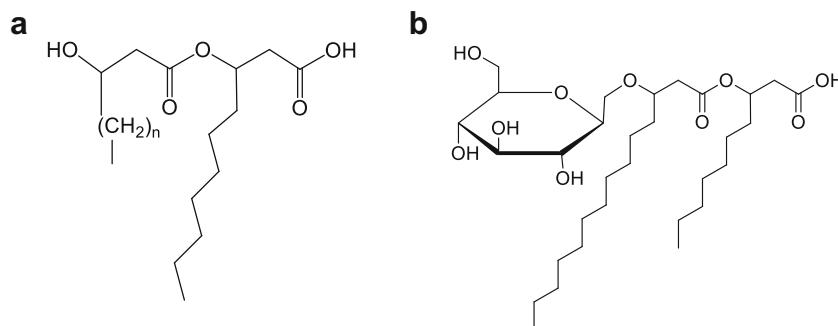
Rhamnolipids consist of one (mono-rhamnolipid) or two (di-rhamnolipid) rhamnose sugars bonded to lipid moieties by an O-glycosidic linkage. Although rhamnolipids are primarily produced by *Pseudomonas* species, studies by Nalini and Parthasarathi (2013, 2014) isolated a *S. rubidaea* SNAU02 strain from hydrocarbon-contaminated soil that was found to produce this class of glycolipid. Chemical characterisation of the compounds produced indicated that *S. rubidaea* SNAU02 was able to produce eight rhamnolipid congeners with varying  $\beta$ -hydroxy fatty acid chains ranging from C<sub>8</sub> to C<sub>16</sub> (Nalini and Parthasarathi 2013, 2014). Amongst the detected mono- and di-rhamnolipids, the di-rhamnolipid rha-rha-C<sub>10</sub>-C<sub>8</sub> ( $\beta$ - $\alpha$  rhamnosyl (1  $\rightarrow$  2) rhamnosyl- $\beta$ -hydroxydecanoyl- $\beta$ -hydroxyoctadecanoic acid) (Fig. 3a) was found to be the most abundant (Nalini and Parthasarathi 2014). The chemical structure of another major component, a mono-

rhamnolipid produced by *S. rubidaea* SNAU02, was also determined, which was rha-C<sub>10</sub>-C<sub>10</sub> (rhamnosyl- $\beta$ -hydroxydecanoyl- $\beta$ -hydroxydecanoic acid) (Fig. 3b) (Nalini and Parthasarathi 2014).

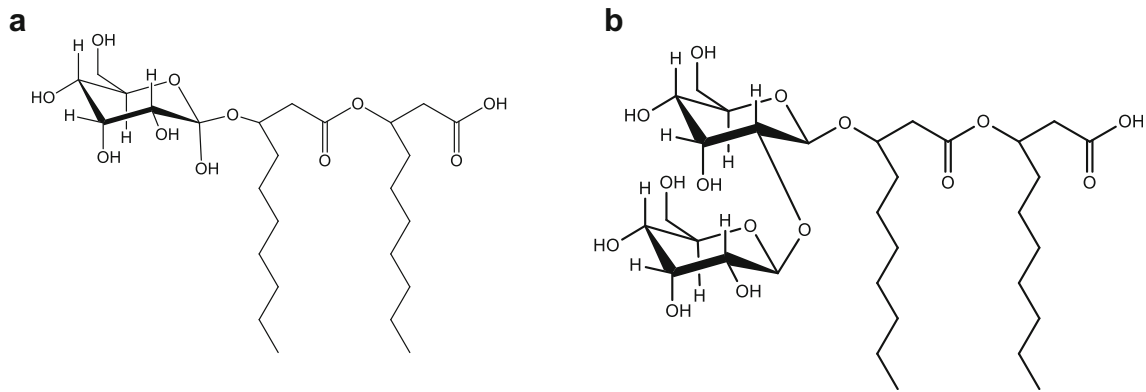
## Biosynthesis of biosurfactants produced by *Serratia* species

The biosynthesis of lipopeptides involves multistep processes mediated by various non-ribosomal peptide synthetase (NRPS) enzymes which catalyse the condensation and selection of amino acid residues to yield various metabolites (including lipopeptides). The mechanisms involved in the biosynthesis of serrawettin W1, serrawettin W2 and the stephensioides have been identified, while the mechanisms involved in the biosynthesis of serrawettin W3 and the glycolipids produced by *Serratia* species are not fully elucidated. Investigation into the biosynthesis of serrawettin W1, serrawettin W2 and stephensioides revealed open reading frames (ORF), namely *swrW*, *swrA* and *sphA* (Fig. 4), respectively, that displayed high homology with the NRPS family (Marahiel et al. 1997; Li et al. 2005; Ganley et al. 2018). The ORF is comprised of a unimodular or multimodular region, where each module consists of specific condensation (C), adenylation (A), thiolation (T) and thioesterase (TE) domains in functional order (Fig. 4) (Marahiel et al. 1997; Li et al. 2005).

The systematic functioning of NRPSs involved in the biosynthesis of secondary metabolites has been determined and indicates that the precursor molecules (e.g. amino acids) are linked to the phosphopantetheinyl moiety of each thiolation domain in the multimodular enzyme (Matsuyama et al. 2011). A previous study by Sunaga et al. (2004) revealed a novel gene, *pswP*, in *S. marcescens* that encodes for 40-phosphopantetheinyl transferase (PPTase). PPTase is the activator of peptidyl carrier protein (PCP) (also known as the thiolation domain in the NRPS family) and was shown to be essential for the biosynthesis of serrawettin W1 (Sunaga et al. 2004). The PPTase is presumed to be involved in the



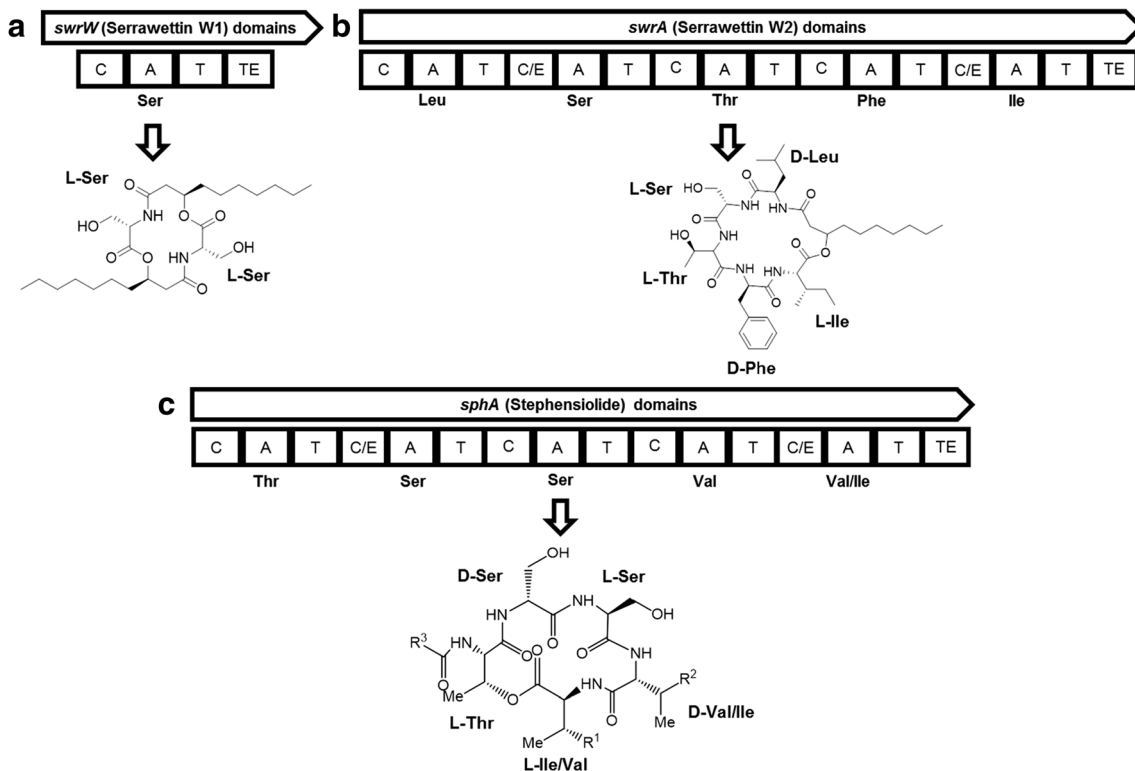
**Fig. 2** The chemical structure of rubiwettin **a** R1 [3-(3'-hydroxytetradecanoyloxy) decanoate and 3-(3'-hydroxyhexadecanoyloxy) decanoate] and **b** RG1 [ $\beta$ -D-glucopyranosyl 3-(3'-hydroxytetradecanoyloxy) decanoate minor fatty acid isomers] (adapted from Matsuyama et al. 1990)



**Fig. 3** The proposed chemical structures for the **a** mono-rhamnolipid (Rha-C<sub>10</sub>-C<sub>10</sub>) and **b** di-rhamnolipid (Rha-Rha -C<sub>8</sub>-C<sub>10</sub>) produced by *S. rubidaea* SNAU02 (adapted from Nalini and Parthasarathi 2014)

incorporation reaction of L-serine as a molecular component of serrawettin W1 (Sunaga et al. 2004). Therefore, the thiolation domain must be activated by acquiring the phosphopantetheinyl moiety by the action of PPTase for the accurate functioning of serrawettin W1 synthetase (Li et al. 2005). After the activation by PPTase, the adenylation domain is able to adenylate L-serine to an activated form. This activated L-serine will then bind as a thioester to the thiolation domain (that has already been phosphopantetheinylated by PPTase) (Li et al. 2005). The amino group of the L-serine

bound to the thiolation domain will react and create an amide linkage with the 3-D-hydroxydecanoyl moiety by detaching from the acyl carrier protein (ACP). This reaction results in the formation of a serratamic acid linked to the thiolation domain (Li et al. 2005). The serratamic acid is transferred to the thioesterase domain and the biosynthesis of a second serratamic acid bound to the free thiolation domain will follow. The two neighbouring serratamic acids will form an intramolecular linkage which results in the release of a symmetric and circular product, serrawettin W1, from the *swrW* gene



**Fig. 4** Comparison of the proposed biosynthetic pathways for the biosynthesis of **a** serrawettin W1 encoded by *swrW*, **b** serrawettin W2 encoded by *swrA* and **c** stephensiolide encoded by *sphA* with their

respective domain regions [condensation (C), adenylation (A), thiolation (T) and thioesterase (TE)] and general structures produced by each gene cluster (adapted from Ganley et al. 2018)

region (Li et al. 2005). The serrawettin W2 biosynthetic pathway has a similar mechanism; however, it is encoded by the *swrA* gene (Su et al. 2016).

The biosynthesis of serrawettin W2 was investigated in *S. surfactantifaciens* sp. YD25<sup>T</sup> and a hybrid polyketide synthases (PKS)-NRPS gene cluster putatively involved in the biosynthesis of serrawettin W2 was identified (Su et al. 2016). The biosynthetic pathway was determined based on the presence of PKS and NRPS encoding genes. The precursor molecule, a C<sub>10</sub> fatty acid (FA), is synthesised by the PKS SwrEFG gene cluster and additional undetermined proteins and is released as a fatty acyl-CoA (Su et al. 2016). The NRPS gene cluster encodes the core W2-peptide chain (5-amino acid peptide moiety) and contains five modules (Fig. 4b) (Su et al. 2016).

Similar to serrawettin W1 and serrawettin W2, the biosynthesis of stephensiolides was investigated in a *Serratia* strain using bioinformatic analysis (sequencing of the *Serratia* sp. genome) and a NRPS gene cluster encoded by the *sphA* gene putatively involved in the biosynthesis of stephensiolides was identified (Fig. 4c) (Ganley et al. 2018). Further bioinformatics analysis indicated that seven sequenced *S. marcescens* strains contained a homologue with identical predicted domain regions as the *sphA* gene (Ganley et al. 2018). Stephensiolides contain D- and L-amino acids and these lipopeptides are cyclised through a macrolactone ring. Both of these characteristic properties provide an indication that the peptides are synthesised by NRPSs. Although both serrawettin W2 and stephensiolide are cyclic pentapeptides and are similarly biosynthesised, they are cyclised in a different manner. Serrawettin W2 is cyclised through a 3-hydroxyl group of the fatty acid, while stephensiolides are cyclised through a hydroxyl group of the threonine (Ganley et al. 2018). Therefore, serrawettin W1, serrawettin W2 and stephensiolides are synthesised by various NRPS enzymes. The biosynthesis of these biosurfactants by *Serratia* species is influenced by the producer strain, growth conditions and nutrient composition.

## Physicochemical characterisation of biosurfactants produced by *Serratia* species

In recent years, there has been an increased scientific interest in the isolation of microorganisms that produce biosurfactants with unique physicochemical properties due to their potential application in diverse industries and in bioremediation processes (Fracchia et al. 2012). The amphiphilic nature of biosurfactants allows for their accumulation at the interface between immiscible fluids or between a fluid and a solid, thereby reducing surface (liquid-air) tension and interfacial (liquid-liquid) tension (Varjani and Upasani 2017). The accumulation of these compounds at surfaces or interfaces also

decreases the repulsive forces (cohesive forces that hold water molecules together) between two immiscible phases, such as water and oil (Peele et al. 2016). This results in the dispersion of one liquid into another leading to the emulsification of the two immiscible liquids (Soberón-Chávez and Maier 2011). The physicochemical properties of biosurfactants thus include their ability to reduce surface tension, form hydrocarbon emulsions (emulsification) and thus enhance the water solubility of hydrophobic compounds (Desai and Banat 1997).

Surface tension is considered to be the measure of free energy per unit area associated with a surface or interface and is measured using a tensiometer (DuNouy ring method). This is a common screening method to detect the presence of biosurfactant compounds produced by a microorganism (Satpute et al. 2010). Typically, a biosurfactant is considered effective if it can reduce the surface tension between water and air from 72 to 35 mN/m and the interfacial tension between water and *n*-hexadecane from 40 to 1 mN/m (Soberón-Chávez and Maier 2011). Furthermore, a bacterial strain is considered to be a good biosurfactant producer if it is able to reduce the surface tension of a growth medium by  $\geq 20$  mN/m compared to distilled water (Walter et al. 2010). Previous studies have identified lipopeptides and glycolipids capable of reducing the surface tension of a growth medium. For example, Dusane et al. (2011) isolated a *S. marcescens* strain that produced a glycolipid that was able to reduce the surface tension of the growth medium from 52.0 to 27 mN/m. Similarly, previously characterised serrawettin W1, serrawettin W2 and serrawettin W3 produced by *S. marcescens* ATCC 13380, NS 25 and NS 45 strains, respectively, were capable of reducing the surface tension of water to 32.2, 33.9 and 28.8 mN/m, respectively (Matsuyama et al. 2011). In addition, Matsuyama et al. (2011) indicated that rubiwettin R1 and rubiwettin RG1 produced by *S. rubidaea* ATCC 27593 reduced the surface tension of water to 25.5 and 25.8 mN/m, respectively.

Biosurfactants are also known to increase the solubility and bioavailability of hydrophobic organic compounds (Mnif and Ghribi 2015). Emulsification activity is thus an indirect method often used to screen for biosurfactant-producing microorganisms and the emulsification index can be calculated by measuring the emulsion height divided by the total height of the solution (equal volume of hydrocarbon to cell-free broth culture). Although kerosene is the most commonly used to test for emulsification, previous studies have tested the ability of biosurfactants produced by *Serratia* species to emulsify various hydrocarbons, such as diesel and crude oil (Pruthi and Cameotra 2000; Wei et al. 2004; Ibrahim et al. 2013). For example, Wei et al. (2004) identified a pigmented *S. marcescens* SS-1 strain that was able to emulsify both kerosene (72%) and diesel (40%).

The amphiphilic nature and structure of these compounds thus confer a diverse range of useful properties, such as emulsification activity, wetting, foaming, dispersion traits, surface activity and the reduction in viscosity of heavy liquids, allowing for their application in many industrial and commercial processes (Satpute et al. 2010; Aparna et al. 2012).

## Production of biosurfactants by *Serratia* species

Biosurfactants are promising alternatives to synthetic surfactants and have been incorporated into commercialised products, such as Bio Surfactants ACS-Sophor® (sophorolipids) produced by Allied Carbon Solutions Co., Ltd., NatSurFact (rhamnolipids) produced by Logos Technologies, LLC and Yashinomi Vegetable Wash (sophorolipids) produced by Saraya Co. Ltd., amongst others (Geetha et al. 2018). However, the increased global biosurfactant market has resulted in the need for cost-effective, industrial-scale production and purification processes that result in maximum biosurfactant yield (Nitschke and Silva 2018). Biosurfactant production is dependent on the producer strain, physicochemical conditions (temperature, pH, agitation and aeration) and medium composition (carbon source, nitrogen source and salinity). Table 1 indicates various biosurfactant-producing *Serratia* species, the type of biosurfactant produced by each strain and the media and culture conditions used for small-scale biosurfactant production (excluding production in bioreactors).

The production of biosurfactants by *Serratia* species occurs during the late log and early stationary phase of growth indicating that the biosurfactants are secondary metabolites (Pruthi and Cameotra 2000; Bidlan et al. 2007; Dusane et al. 2011). It is noteworthy that the production of secondary metabolites, including biosurfactants and prodigiosin, by *Serratia* species is temperature-dependent (Matsuyama et al. 2011; Eckelmann et al. 2018). Matsuyama et al. (1986, 1990) investigated the production of secondary metabolites by *Serratia* species at varying temperatures and found that while the bacterial strains grew well at both 30 °C and 37 °C, serrawettin (W1, W2 and W3), rubiwettin (R1 and RG1) and the pigment (prodigiosin) were produced at 30 °C and were significantly reduced or absent when the bacterial suspension was grown at 37 °C. Numerous studies have subsequently utilised 30 °C as the optimum growth temperature for the production of biosurfactants by *Serratia* species (Pruthi and Cameotra 2000; Anyanwu et al. 2010; Dusane et al. 2011; Kadouri and Shanks 2013; Su et al. 2016; Almansoori et al. 2017).

Another physicochemical condition that influences production is oxygen transfer during cell growth. Several factors may contribute to the transfer of oxygen from the gas phase to the

aqueous phase within the growth medium, such as agitation speed and aeration, amongst others, thereby affecting cell growth and, ultimately, biosurfactant production (Fakruddin 2012). Various agitation speeds have been used during the growth and production of biosurfactants by *Serratia* species (Table 1). In addition, the media composition, pH values (5 to 9) and cultivation times (24 to 168 h) similarly influence the quality and quantity of the biosurfactants produced. Almansoori et al. (2017) investigated the effects of varying cultivation times (24 to 168 h), agitation speeds (100, 125, 150, 180 or 200 rpm) and pH (5 to 9) for the production of biosurfactants by a *Serratia* strain. Results indicated that the optimum cultivation time, agitation speed and pH for maximum biosurfactant production and surface tension reduction were after 120 h at 200 rpm with the pH of the growth medium at 8.0. However, for the small-scale production of biosurfactants by *Serratia* species, an agitation speed of 200 rpm, pH of 7.2 and a cultivation time of 72 h are extensively used (Table 1). The salt concentration within the growth medium was also hypothesised to influence microbial growth and biosurfactant production by *Serratia* species (Almansoori et al. 2017). Almansoori et al. (2017) further indicated that the growth and production of a biosurfactant by an *S. marcescens* strain was reduced in the absence of salt. The authors additionally tested varying salt concentrations (ranging from 1 to 5%), with 1% recorded as optimum for biosurfactant production. The production of biosurfactants can also be improved by the presence of different carbon and nitrogen sources, as these factors strongly influence cell growth and the accumulation of metabolic products (Santos et al. 2016). Moreover, research has indicated that the concentration and type of biosurfactant compounds synthesised by the producer strain are influenced by the type of carbon substrate (Rahman and Gakpe 2008; Ndlovu et al. 2017).

Water miscible substrates, such as glucose, sucrose, fructose, mannitol and glycerol, and water immiscible substrates, such as mahua oil and olive oil, have been used for biosurfactant production by *Serratia* species (Table 1; Almansoori et al. 2017). At low nitrogen levels, bacterial growth is also limited, which favours cell metabolism towards the production of secondary metabolites (Santos et al. 2016). Numerous nitrogen sources have subsequently been used for the production of biosurfactants by *Serratia* species, including peptone, yeast extract, tryptone, ammonium sulphate, ammonium nitrate and casamino acids. Of the various carbon and nitrogen sources utilised for the production of biosurfactants by *Serratia* species, the most widely used are glycerol and peptone (Table 1), respectively. Almansoori et al. (2017) also investigated the effects of different carbon (glycerol, olive oil, glucose, sucrose and fructose) and nitrogen sources (ammonium sulphate, yeast extract, peptone and combinations of these three nitrogen sources) on lipopeptide production by *S. marcescens* by measuring the biosurfactant yield and the

**Table 1** The culture conditions and media composition for the production of biosurfactants produced by *Serratia* spp.

<i>Serratia</i> species	Strain	Biosurfactant	Carbon source	Nitrogen source	Cultivation time	Agitation speed	pH	Reference
<i>S. marcescens</i>	ATCC 13880, NS38, 274	Serrawettin W1	Glycerol	Peptone	72 h	n/a	7.2	Matsuyama et al. (1985, 1986)
<i>S. marcescens</i>	SHHRE645	Serratamolides A–F	Glucose	Tryptone and Soytone	120 h	n.s	7.0	Dwivedi et al. (2008)
<i>S. marcescens</i>	H30	Serrawettin W1	Sucrose	Yeast extract	24 h	200 rpm	7.2	Zhang et al. (2010)
<i>S. marcescens</i>	MSRBB2	Serratamolides	Glucose	Peptone and yeast extract	168 h	n.s	n.s	Eckelmann et al. (2018)
<i>S. marcescens</i>	S823	Serratamolides A and G	n.s	Yeast extract	48 h	180 rpm	7.0–7.2	Zhu et al. (2018)
<i>S. marcescens</i>	DSM12481	Serrawettin W1	Luria broth (LB) media	LB media	24 h	150 rpm; 130 rpm	n.s	Theis et al. (2014); Hage-Hülsmann et al. (2018)
<i>S. marcescens</i>	NS 25	Serrawettin W2	Glycerol	Peptone	72 h	n/a	7.2	Matsuyama et al. (1986, 1992)
<i>S. marcescens</i>	n.s	Serrawettin W2	Glycerol	Ammonium chloride	168 h	135 rpm	7.0	Motley et al. (2016)
<i>S. liquefaciens</i>	MG1	Serrawettin W2	Glucose	Casamino acids	48 h	n/a	n.s	Lindum et al. (1998)
<i>S. surfactantifaciens</i>	YD25 <sup>T</sup>	Serrawettin W2	Glycerol	Protease peptone	72 h	n.s	7.2	Su et al. (2016)
<i>S. marcescens</i>	NS 45	Serrawettin W3	Glycerol	Peptone	72 h	n/a	7.2	Matsuyama et al. (1986)
<i>S. marcescens</i>	HDB	Lipopeptide	Glycerol*	Ammonium sulphate and peptone*	120 h	100–200 rpm	5.0–9.0	Almansoori et al. (2017)
<i>S. marcescens</i>	NSK-1	Lipopeptide	Glycerol	Ammonium sulphate	72 h	180 rpm	7.0	Anyanwu et al. (2010)
<i>S. marcescens</i>	n.s	Lipopeptide	n.s	Ammonium nitrate	168 h	n.s	7.2	Ibrahim et al. (2013)
<i>Serratia</i> sp.	n.s	Stephensiolides	LB media	LB media	24 h	250 rpm	7.0	Ganley et al. (2018)
<i>S. marcescens</i>	MTCC 86	Glycolipid (sucrose lipid)	Sucrose	Ammonium sulphate	24 h	200 rpm	7.2	Pruthi and Cameotra (2000)
<i>S. marcescens</i>	n.s	Glycolipid	n.s	Peptic digest of animal tissue and yeast extract	48 h	120 rpm	n.s	Dusane et al. (2011)
<i>S. rubidaea</i>	SNAU02	Rhamnolipid	Mannitol	Yeast extract	72 h	200 rpm	6.97	Nalini and Parthasarathi (2013)
<i>S. rubidaea</i>	SNAU02	Rhamnolipid	Mahua oil	Ammonium nitrate	168 h	n.s	7.0	Nalini and Parthasarathi (2014)
<i>S. rubidaea</i>	ATCC 27593	Rubiwettin R1 and RG1	Glycerol	Peptone	72 h	n/a	7.2	Matsuyama et al. (1990)

n/a not applicable (agar used), n.s not specified

\*Various carbon and nitrogen sources tested



reduction of surface tension. Results indicated that of the five carbon sources used, glycerol resulted in the highest yield of 1.05 g/L and the lowest surface tension with a value of 30.4 mN/m recorded. Furthermore, the combination of ammonium sulphate and peptone (shown in Table 1) as nitrogen sources resulted in the highest yield of the biosurfactant compounds (1.33 g/L) and the lowest reduction in surface tension with a value of 29.9 mN/m recorded. Although biosurfactant production by *Serratia* species is strongly dependent on the producer strain, in summary the most widely used growth conditions and media composition for the small-scale production include an incubation temperature of 30 °C in a medium containing glycerol as a carbon source and peptone as a nitrogen source at a pH of 7.2 with agitation at 200 rpm.

In addition to the small-scale production of biosurfactants, studies have evaluated the use of bioreactors for the large-scale production of biosurfactants by *Serratia* species. A study by Granada et al. (2018) investigated the effects of dissolved oxygen on the large-scale production of bioactive metabolites, including serratamolide A, prodigiosin and haterumalide NC (a cytotoxic molecule), by *Serratia* sp. ARP5.1 using a 7-L stirred tank bioreactor. The strain was inoculated into 4 L of mineral medium (with glucose as a carbon source) at 28 °C for 96 h. Additionally, three agitation speeds (150, 300, 450 rpm) and three aeration rates (0.5, 1.0 and 1.5 vvm) were tested for optimal production. It was found that oxygen was a crucial factor for the biosynthesis of these secondary metabolites in a bioreactor, with the best combination of agitation speed and aeration observed at 450 rpm and 1.5 vvm, respectively. It was therefore recommended that dissolved oxygen be included as a parameter for the large-scale production of secondary metabolites by *Serratia* species. In addition, Roldán-Carrillo et al. (2011) utilised a Box-Behnken experimental design to evaluate the effect of nutrient ratios (C/N, C/Mg and C/Fe) on biosurfactant production by a *S. marcescens* strain. The results indicated that a nutrient ratio of C/N = 5, C/Mg = 30 and C/Fe = 26,000 was optimal for biosurfactant production. This media composition was then utilised for large-scale production in a 3-L bioreactor. The large-scale production of the biosurfactant by this strain was conducted in a volume of 1.5 L of the growth medium within the 3-L bioreactor. After 48 h, the biosurfactant was extracted using two volumes of ethanol after acid precipitation from the cell-free broth and the crude extract was freeze-dried and weighed. It was thus found that the nutrient ratios optimised by the Box-Behnken experimental design for biosurfactant production by this strain successfully yielded 21.6 g/L of crude extract after 48 h.

Although studies have used various methods to recover biosurfactants, further optimisation of the large-scale production process and downstream recovery of biosurfactants produced by *Serratia* species is still required. Furthermore, biosurfactants can be used to replace chemically synthesised surfactants in various industries as they exhibit a low toxicity,

high biodegradability, can be produced from cost-effective materials and they are stable at extreme temperatures, pH and salinity (Satpute et al. 2010; Santos et al. 2016).

## Applications of biosurfactant compounds produced by *Serratia* species

Biosurfactants have several advantages over chemical surfactants as they exhibit a low toxicity, high biodegradability, can be produced from cost-effective materials and exhibit stability at extreme temperature, pH and salinity (Satpute et al. 2010; Santos et al. 2016). Thus, due to their diverse chemical properties and biological activity, biosurfactants have the potential to replace their chemical counterparts in a number of industries, such as the petroleum, medical, pharmaceutical, food, agriculture, beverage, cosmetics, textiles and mining industries as well as in bioremediation strategies. Furthermore, biosurfactants produced by *Serratia* species have the potential to be applied as antimicrobial compounds, antifouling agents and antitumour compounds and as emulsifying agents of hydrocarbons.

### Medical and pharmaceutical industries

#### Biosurfactant compounds as antimicrobial agents

Although the exact mode of action of biosurfactant compounds has yet to be elucidated, both lipopeptides (including serrawettin W1, serrawettin W2 and stephensiolides) and glycolipids (rhamnolipid and glucose-palmitic acid glycolipid) produced by *Serratia* species have been reported to display antimicrobial activity (Dwivedi et al. 2008; Dusane et al. 2011; Kadouri and Shanks 2013; Nalini and Parthasarathi 2014; Su et al. 2016; Eckelmann et al. 2018; Ganley et al. 2018). Dwivedi et al. (2008) purified serratamolide A (serrawettin W1) and homologues (serratamolide B to F) and tested these compounds for antimicrobial activity against *Mycobacterium* species. It was found that serratamolide A and all the homologues exhibited antibacterial activity against *M. diernhoferi* at a minimum inhibitory concentration of 0.18 mM. A recent study by Su et al. (2016) also investigated the antimicrobial activity of secondary metabolites produced by *S. surfactantifaciens* sp. YD25<sup>T</sup>. The secondary metabolite was identified as serrawettin W2 and was found to exhibit inhibitory activity against *Staphylococcus aureus*, *Pseudomonas aeruginosa* and *Shigella dysenteriae*, amongst other bacterial pathogens, at a concentration of 300 µg/mL (Su et al. 2016). Hage-Hülsmann et al. (2018) further investigated the synergistic antibiotic effects of prodigiosin and biosurfactants produced by *S. marcescens* DSM12481 strain against a soil bacterium, *Corynebacterium glutamicum*. As results indicated that the combination of prodigiosin and

serrawettin W1 generated a larger zone of inhibition compared to the individual compounds, employing a combination of biomolecules may be a useful strategy for future antimicrobial formulations.

### Biosurfactant compounds as antifouling agents

Biosurfactants have been reported to effectively inhibit biofilm formation on various surfaces (Epstein et al. 2011). When a surface is conditioned with a layer of a biosurfactant, it becomes more hydrophilic and is thus expected to have reduced microbial attachment (Gudiña et al. 2010; Zeraik and Nitschke 2010). In addition, many biosurfactant compounds disrupt preformed biofilms. McLandsborough et al. (2006) hypothesised that a biosurfactant has to penetrate the biofilms' extracellular matrix (possibly through the water channels) and adhere to the interface, thereby reducing the interfacial tension between the substratum surface and the biofilm. The interactions involved in bacterial adhesion are also reduced, ultimately leading to biofilm removal. Dusane et al. (2011) investigated the antifouling activity of a glycolipid, composed of palmitic acid esterified to glucose, produced by a *S. marcescens* strain to disrupt a preformed biofilm or prevent biofilm formation on polystyrene microtitre plates. Results indicated that 100 µg/mL of the glycolipid prevented the attachment of 94%, 88% and 82% of *Bacillus pumilus* TiO1, *P. aeruginosa* PAO1 and *Candida albicans* BH cells, respectively, to polystyrene microtitre plate surfaces. A glycolipid concentration of 50 µg/mL also resulted in a reduction of up to 55% *C. albicans*, 62% *P. aeruginosa* and 55% *B. pumilus* reduction of preformed biofilms when compared to untreated control biofilms grown on polystyrene microtitre plates (Dusane et al. 2011). A similar study conducted by Motley et al. (2016) investigated the ability of serrawettin W2 produced by a *Serratia* strain to inhibit the microbial adhesion of *C. albicans* to a 96-microwell plate. This was conducted by incubating the test compound with the inoculum in the wells of the 96-microwell plate and determining half the inhibitory concentration (IC<sub>50</sub>) value using a tetrazolium salt (XTT) reduction assay. An IC<sub>50</sub> value of 7.7 ± 0.7 µM for serrawettin W2 inhibited the biofilm formation of *C. albicans* and it was concluded that the cyclic lipodepsipeptides produced by the *Serratia* strain may be used to control *C. albicans* infections associated with biofilm modulation.

### Other medical and pharmaceutical applications

Kadouri and Shanks (2013) suggested that *Serratia* species could serve as a potential source of antibiotics to combat multidrug-resistant opportunistic pathogens such as methicillin-resistant *S. aureus* (MRSA) and *Staphylococcus epidermis*. Despite their antimicrobial properties, some compounds such as serrawettin W1 have

cytotoxic activity and are therefore unlikely to be used directly as a systemic antibiotic. In addition to antimicrobial activity, Tomas et al. (2005) filed a patent for the use of a serratamolide (serrawettin W1) produced by a *S. marcescens* 2170 strain as a chemotherapeutic agent. The serratamolide was found to reduce the cell viability (induce apoptosis) of various cancer cell lines (Jurkat, Molt-4, NSO, HGT-1, HT-29 and GLC-4S), while displaying no effect on non-malignant cell lines (NIH-3T3, NRK-49F and IEC-18). In addition to serratamolides (serrawettin W1), serrawettin W2 produced by *S. surfactantfaciens* sp. YD25<sup>T</sup> was also shown to exhibit anticancer activity by suppressing the growth of cancer cell lines (HeLa and Caco-2 cell lines), while not significantly affecting the viability of non-malignant cells (Vero and HEK293 cell lines) (Su et al. 2016). Based on this research, serrawettins have the potential to be used as anticancer chemotherapeutic agents against leukaemia, lymphoma, myeloma, carcinoma, melanoma and sarcoma (Tomas et al. 2005).

### Agricultural industry

Although biosurfactants produced by *Serratia* species have been shown to display emulsification activity against hydrocarbons, such as diesel and kerosene (Wei et al. 2004), limited research on the use of lipopeptides and glycolipids produced by *Serratia* species in the food industry has been conducted. Research has however indicated that biosurfactants produced by *Serratia* species can be used in the agricultural industry, due to their antifungal and plant-protecting properties. A patent was filed by Strobel et al. (2005) for the use of serratamolides produced by a *S. marcescens* strain in the agricultural industry, due to its anti-mycotic activity against oomycete pathogens, which has the potential to be applied for crop protection. Similarly, a study by Nalini and Parthasarathi (2014) investigated the antifungal activity of a rhamnolipid produced by *S. rubidaea* SNAU02. The study found that the rhamnolipid exhibited antifungal activity against *Fusarium oxysporum* and *Colletotrichum gloeosporioides* (plant pathogens) at a concentration of 100 µg/mL. In addition, the rhamnolipid (at a concentration of 100 µg/mL) did not exhibit toxicity towards the seeds of *Brassica oleracea* and *Artemia salina* employed as bio-indicators. The biosurfactant produced by *S. rubidaea* SNAU02 thus exhibits potential as a biocontrol agent against plant pathogens (Nalini and Parthasarathi 2014). Similarly, Granada et al. (2018) investigated the use of crude extracts obtained from a *Serratia* strain as a biological control agent of avocado pathogens. The study identified serratamolide A, prodigiosin and haterumalide NC in the crude extract and found that the crude extract displayed high activity against the pathogens *Phytophthora cinnamomi* and *C. gloeosporioides*. It was

therefore concluded that the secondary metabolites produced by this strain may be used for crop protection or to maintain plant health.

### Bioremediation and petroleum industries

Chemically synthesised surfactants have been used for the bioremediation of oil-contaminated sites as well as to enhance the recovery of oil from oil reservoirs in the petroleum industry (Banat 1995). However, as synthetic surfactants are not biodegradable and can be toxic to the environment, biosurfactants provide an environmentally friendly alternative and have been shown to exhibit equivalent emulsification properties. In addition to emulsifying hydrocarbons, studies have indicated that biosurfactants produced by *Serratia* species have the potential for application as microbial enhanced oil recovery (MEOR) agents (Pruthi and Cameotra 2000; Ibrahim et al. 2013; Nalini and Parthasarathi 2013). MEOR is considered a tertiary recovery method that could recover the residual oil using microorganisms or their products (biosurfactants) and can be evaluated using a sandpack technique (sand previously saturated with oil in a column) (Pruthi and Cameotra 2000; Amani et al. 2010). A study by Ibrahim et al. (2013) isolated a *S. marcescens* strain that was found to produce a lipopeptide and investigated the effects of the lipopeptide to enhance the recovery of crude oil from a sandpack column. The lipopeptide was able to recover 76% of the crude oil from the column and was earmarked as a potential enhanced oil recovery agent. In addition, a sucrose lipid (glycolipid) produced by a *S. marcescens* strain was found to exhibit excellent emulsification activity against a wide range of hydrocarbons and effectively recovered 90% of residual oil from an oil-saturated sandpack column (Pruthi and Cameotra 2000). Pruthi and Cameotra (2000) further indicated that the glycolipid displayed a strong ability to recover oil from the walls of containers and therefore could possibly be applied in cleaning operations. Similarly, Nalini and Parthasarathi (2013) demonstrated that a concentration of 0.05% rhamnolipid solution produced by a *S. rubidaea* SNAU02 strain was able to recover 92% of used engine oil previously absorbed to a sand sample, while the commercially available surfactant (sodium dodecyl sulphate) only removed 60% of the oil from the contaminated sand sample.

### Conclusions and future perspectives

The review provided insight into the classification and structure of biosurfactants produced by *Serratia* species that have

been characterised and identified as serrawettins, stephensioides, rubiwettins and rhamnolipids. Furthermore, the biosynthetic pathways of three lipopeptides synthesised by NRPS enzymes, including serrawettin W1, serrawettin W2 and stephensioides, encoded by *swrW*, *swrA* and *sphA*, respectively, were discussed. However, further research is still required to elucidate the chemical structure of serrawettin W3 and rubiwettin R1 as well as the biosynthetic pathway of glycolipids and serrawettin W3 produced by *Serratia* species. While the small-scale production of biosurfactants produced by *Serratia* species has been extensively applied, the large-scale production and downstream recovery of these compounds still require optimisation. Furthermore, a few biosurfactants produced by *Serratia* species have been patented for use in industries such as the agricultural sector, due to their antifungal and plant-protecting properties, and numerous studies have demonstrated that these compounds have the potential to be applied as hydrocarbon emulsifiers for enhanced oil recovery. The potential of these compounds as antitumor agents is also being explored and while biosurfactants produced by *Serratia* species display antibacterial and antifungal activity, the mode of action of these compounds still needs to be extensively investigated. Biosurfactants produced by *Serratia* species are thus a vital source of antimicrobial, antifouling, surface active (emulsifier) and antitumor compounds and should be considered one of the major candidates for application in the medical, agricultural and petroleum industries.

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### Compliance with ethical standards

**Conflict of interest** The authors declare that they have no competing interests.

**Ethical approval** This article does not contain any studies with human participants or animals performed by any of the authors.

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