

1 **Review Article**

2 **Methods of purification and characterization of biosurfactants: an overview**

4 **Abstract**

5 Biosurfactants are amphiphilic molecules produced by several microorganisms including
6 bacteria, filamentous fungi and yeasts. They are classified according to their chemical
7 composition (glycolipids, lipopeptides, glycoproteins, glycolipopeptides, phospholipids) and
8 their molecular weight (low and high molecular weight). Biosurfactants are currently used in
9 several industrial fields (food industry, pharmacy, medicine, detergents, agriculture,
10 cosmetics, oil recovery and bioremediation) owing to some of their specific properties such
11 as their low toxicity, biocompatibility, biodegradability, environmentally-friendly, stability
12 under extreme conditions (temperature, pH and salinity), structural diversity, production using
13 renewable low-cost substrates, agreement with green chemistry and sustainability. Despite the
14 properties of biosurfactants, there are few studies leading to their structural characterization.
15 The most reported structures are from biosurfactants produced by *Bacillus* spp. (surfactins,
16 fengycins, lichenysines) and *Pseudomonas* spp. (rhamnolipids). The majority of researches
17 performed on biosurfactants emphasized their functional groups characterization. These
18 biosurfactants have yet to be fully characterized at structural level. Hence, in this review, we
19 highlight the different purification and structural characterization techniques which can be
20 combined to provide information on the structure of the biosurfactants independently of its
21 complexity. This will lead to enhance their application in some highly pointed industrial fields
22 such as medicine and pharmacy.

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27 **Keywords:** Biosurfactants, Microorganisms, Purification, Structural characterization.

28 **Introduction**

29 Biosurfactants are amphiphilic surface-active compounds produced by microorganisms from
30 diverse sources (terrestrial, marine environments, sludge, etc.) including bacteria, molds and
31 yeasts (Nataraj *et al.*, 2021a; Mouafo *et al.*, 2022). They are either excreted extracellularly or
32 remain bound to the producing cells (Vallejo *et al.*, 2021). They are synthesized by
33 microorganisms through different metabolism routes depending of the nature and the
34 composition of the substrates (Figure 1). Once the substrates are intake, some microorganisms
35 produced enzymes that might combine the hydrophilic and hydrophobic components of the
36 substrates leading to the formation of biosurfactants. The substrates can also be metabolized
37 into coenzyme into acetyl coenzyme A, that will be used in *de novo* pathway as the precursor
38 of the synthesis of sugars, lipids and proteins (Tokumoto *et al.*, 2009). Hence, the enzymatic
39 combination of these hydrophilic (sugar, proteins) and hydrophobic (lipids) components
40 synthesized by the microorganisms will lead to the biosurfactants. In some case, the
41 hydrophilic (sugar, proteins) or hydrophobic (lipids) components synthesized by the
42 microorganisms through *de novo* pathway can be combined with carbohydrates, proteins or
43 lipids' components intake from substrates (Sydatk and Wagner, 1987). According to Desai
44 and Banat (1997), microorganisms often used water insoluble substrates for the synthesis of
45 the hydrophobic moiety of the biosurfactants and the water-soluble ones for the synthesis of
46 the hydrophilic moiety. Biosurfactants are classified according to their chemical composition
47 and their molecular weight. With regards to the chemical composition, they can be glycolipids

48 (rhamnolipids, Mannosylerythritol lipids, trehalolipids, xylolipids), lipopeptides (surfactins,
49 viscosines, fengycins, lichenysines), glycolipopeptides, glycoproteins, phospholipids, neutral
50 lipids, polymeric biosurfactants (emulsan, alasan, biodispersan), and particular biosurfactants
51 (protein–sugar–lipid complex molecules) (Bergevin *et al.*, 2017; Nitschke and Silva, 2018;
52 Mouafo *et al.*, 2022). Considering the molecular weight, biosurfactants are grouped as low
53 molecular weight (glycolipids, lipopeptides, phospholipids) and high molecular weight
54 (particular biosurfactants, polymeric biosurfactants) compounds (Smyth *et al.*, 2014).

55 Several investigations on the potential replacements of synthetic surfactants by biosurfactants
56 have been conducted this last decade. The main reasons that have impelled these
57 investigations are their low toxicity, biodegradability, environmentally friendly,
58 biocompatibility, structural diversity, stability under extreme conditions of temperature,
59 salinity and pH, production using renewable low-cost substrates, agreement with green
60 chemistry and sustainability (Marchant and Banat, 2012a; Marchant and Banat, 2012b;
61 Mouafo *et al.*, 2018a; Nitschke and Silva, 2018). These advantages of biosurfactants
62 associated with their diverse properties (emulsifying, de-emulsifying, foaming, dispersing,
63 wetting, solubilizing, antioxidant, antiadhesive, antimicrobial and antibiofilm agents) confer
64 to these latter's, the ability to be applied in many industrial fields including, cosmetics,
65 detergents, oil recovery and bioremediation, agriculture, pesticides,
66 pharmaceuticals/medicines, and food industry (Nitschke and Silva, 2018; Silva *et al.*, 2020;
67 Patel *et al.*, 2021; Mouafo and Somashekar, 2020; Mouafo *et al.*, 2022). As examples, Wattoo
68 *et al.* (2023) highlighted the application of biosurfactants as emulsifiers and antimicrobial
69 agents in the stabilization and preservation of juices. Reshmy *et al.* (2023) reported the
70 exploitation of the antioxidant properties of biosurfactants in reducing lipids oxidation and
71 thus, improving the preservation of lipids-containing foods. The improvement in the shelf life
72 of raw ground goat meat using biosurfactants was reported by Mouafo *et al.* (2020b). The

73 authors noticed a significant inhibition of microbial proliferation, lipids oxidation, proteins
74 degradation, and color changes. Fookao *et al.* (2022) demonstrated the ability of
75 biosurfactants to improve the texture and dough stability of milk bread. The beneficial effects
76 of biosurfactants in enhancing the texture profile (Santoso *et al.*, 2023; Sharma and Arya,
77 2023) and sensory attributes (Rahaman *et al.*, 2023) of starchy foods were recently noticed.
78 Numerous other recent applications of the different properties of biosurfactants in the food
79 industry were summarized in the book of Inamuddin and Adetunji (2023). In view of
80 improving the oral bioavailability of hydrophobic drugs at the target sites, the emulsification
81 properties of biosurfactants were successfully exploited as drug delivery system (Kurozuka *et*
82 *al.*, 2017; Ohadi *et al.*, 2020; Bjerk *et al.*, 2021). For different administration routes of drugs
83 (intravenous, oral, ocular, nasal and topical), there is nowadays the formulation of
84 microemulsion drug delivery systems with biosurfactants that insures their delivery to targets
85 sites and significantly enhance their bioavailability (Ohadi *et al.*, 2020). Besides, there are
86 also interesting reports highlighting the successful applications of biosurfactants for the
87 bioremediation of heavy metals (Cd, Cu, and Pb) in contaminated soils (Sun *et al.*, 2021). All
88 these industrial exploitations of biosurfactants justify the increasing demand of biosurfactants
89 for which the markets reached 2210.5 million dollars by 2018 (Transparency Market
90 Research, 2013) and the high number of companies (approximately 17) around the world,
91 which currently produced biosurfactants (Randhawa and Rahman, 2014; Sekon and Rahman,
92 2014).

93 The activity and structure of biosurfactants which derived from the secondary metabolism of
94 substrates by microorganisms, strictly depend on the producing strain, the medium
95 composition, and the culture conditions (Mulligan *et al.*, 2014; Liu *et al.*, 2015). Although in
96 the literature there are some studies on the full structural characterization of biosurfactants,
97 the majority of these researches concerned lipopeptides' biosurfactants derived from *Bacillus*

98 spp. (surfactins, fengycins, utirins, lichenysins) and glycolipid biosurfactants derived from
99 *Pseudomonas* spp. (rhamnolipids) (Joshi *et al.*, 2016; Sharma *et al.*, 2016; Abdalsadiq *et al.*,
100 2018). For the other microorganisms, almost researches performed on biosurfactants
101 emphasized their functional groups characterization (Lara *et al.*, 2020; Nataraj *et al.*, 2021a,
102 Nataraj *et al.*, 2021b; Sakr *et al.*, 2021; Vallejo *et al.*, 2021; Mouafo *et al.*, 2022). These
103 biosurfactants have yet to be fully characterized at structural level. The principal hurdles are
104 the complex nature of some biosurfactants particularly those derived from lactic acid bacteria
105 (Mouafo *et al.*, 2022). Hence, several purification and characterization techniques are needed
106 to obtain structural information on the molecule. In this review, we highlight the different
107 purification techniques reported in the literature as well as the different colorimetric,
108 spectrophotometric, chromatographic and spectrometric methods which can be combined to
109 obtain the structure of the biosurfactants independently of its complexity. Relevant data used
110 in this review were from several original articles, review articles, book chapters and books.
111 They are were obtained from PubMed, EMBASE, Google Scholar, Scopus, Google, Web of
112 Science and Google. The search terms were the combination of these four keywords:
113 “biosurfactants”, “techniques”, “purification” and “structural characterization”.

114 **1. Purification techniques of biosurfactants**

115 Several techniques were reported in the literature concerning the purification process of
116 biosurfactants. These techniques included dialysis, membrane filtration, centrifugation, acid
117 precipitation, thin layer chromatography and column chromatography. Currently, no defined
118 commercial method is available for the purification of the biosurfactants. Table 1 summarizes
119 the different methods of purification of biosurfactants according to the producing strain et the
120 nature of the biosurfactant.

121 *1.1. Centrifugation and ultracentrifugation*

122 Biosurfactants extracted for culture media are complex mixture of several compounds. The
123 centrifugation allows the separation of these compounds based on their density even tiny the
124 difference can be. According to the size of biosurfactants, higher gravitational-force
125 instruments or ultra-centrifuges are very often used. That technique has been widely used for
126 purification of biosurfactants. A centrifugation at high speed (13,000×g, 4°C, 20 min) was
127 used by Mouafo *et al.* (2020a) to separate the biosurfactants produced by *L. casei* subsp. *casei*
128 TM1B from contaminants. A centrifugation at 10,000 rpm, 4 °C, 15 min was applied by
129 Madhu and Paprulla (2013) in the purification process of biosurfactants. It is important to
130 mention that centrifugation and ultracentrifugation are not used as lone purification
131 techniques of biosurfactants. They are always used as a unit operation of a process that
132 included several other unit operations

133 *1.2. Precipitation*

134 Almost biosurfactants are known for their low solubility under acidic conditions (Nelson et
135 al., 2020). That property was exploited by several authors to purify biosurfactants from other
136 contaminants. Madhu and Paprulla (2013) acidified the solution of crude biosurfactants to pH
137 2 for 18 h, and used acidic water (pH 2.0) to wash the precipitate in order to remove possible
138 contaminants. The precipitation of biosurfactants was also achieved using ice-cold acetone
139 (Bhosale *et al.*, 2019), chilled ethanol (Fookao *et al.*, 2021; Devale *et al.*, 2023) and
140 ammonium sulphate (Vigneshwaran *et al.*, 2021).

141 *1.3. Chromatography*

142 Generally, biosurfactants are produced by microorganisms in association with other
143 molecules. These molecules can be extracted together with biosurfactants following the
144 extraction method chosen. Hence there is a need for purification and chromatography appears
145 a one of the main methods. By the means of the mobile phase the mixture of compounds is
146 flowing through the stationary phase and they are separated. The separation process is based

147 on the differential partitioning of compounds between the mobile and stationary phases. The
148 most reported chromatographic methods for biosurfactants purification are column
149 chromatography, thin layer chromatography (TLC) and reverse phase chromatography
150 (Rodrigues *et al.*, 2006; Sauvageau *et al.*, 2012; Mouafo *et al.*, 2020a).

151 1.3.1. Thin layer chromatography

152 Based on the properties of the biosurfactants present in the mixture, a solvent is chosen. That
153 solvent into which biosurfactants should be dissolved will move up the plate the spot of the
154 compounds deposited at the bottom of the plate. Depending on the physical properties and the
155 structure of compounds from the mixture, they will move up the plate or stays behind. Then,
156 spots are visualized in UV light or after chemical treatment. The solvent systems generally
157 used for purification of biosurfactants on TLC plates are: petroleum ether/diethyl ether/acetic
158 acid, acetone/acetic acid/water, chloroform/methanol/water, ethyl acetate/methanol/water,
159 chloroform/methanol/water/acetic acid (Sauvageau *et al.*, 2012; Banerjee and Ghosh, 2021;
160 Meena *et al.*, 2021, Dabaghi *et al.*, 2023; Fernandes *et al.*, 2023).

161 1.3.2. Column chromatography

162 That technique separates a complex mixture of biosurfactants based on their molecular
163 weight, size and their affinity with the stationary phase. The stationary phases used are
164 generally either silica gel (60–120 mesh) or sephadex, while mobile phase consists of a
165 mixture of solvents with different polarities. Rodrigues *et al.* (2006) used an octyl Sepharose
166 4 FF Prep hydrophobic interaction column and a linear gradient of PBS buffer containing 1.0
167 M $(\text{NH}_4)_2\text{SO}_4$ for elution of the complex mixture of biosurfactants from *Streptococcus*
168 *thermophilus* A. Thavasi *et al.* (2011) and Sharma and Saharan (2016) used silica a gel (60–
169 120 mesh) column with a gradient solvent system (chloroform/methanol) starting from 20:1 to
170 2:1 (v/v). Reverse phase HPLC using a C-18 column with a gradient solvent system
171 (trifluoroacetic acid/millipore water, 0.05:99.95 and TFA/millipore water/acetonitrile,

172 0.05:19.95:80) was applied by for purification of biosurfactants from *L. casei* subsp. *casei*
173 TM1B (Mouafo *et al.*, 2020a). A silica gel column (60 Mesh) with chloroform–methanol–
174 water (65:25:4) as mobile phase was successfully used by Banerjee and Ghosh (2021) to
175 purify biosurfactants from *Bacillus oceanisediminis* H2. A Sephadex G-25 column with the
176 sodium phosphate buffer (20 mM; pH 7.5) as elution solvent was used by to purify
177 biosurfactants from *B. velezensis* KLP2016 (Meena *et al.*, 2021). Biosurfactants from *B.*
178 *subtilis* LSFM-05 was purified on a silica gel column (0.03–0.07 mm, 60 Å) using the
179 following solvent system at different polarities: chloroform, methanol, and an aqueous
180 solution of 28% (v/v) ammonium hydroxide (de Faria *et al.*, 2021). Vigneshwaran *et al.*
181 (2021) purified biosurfactant from *Brevibacillus* sp. AVN13 using fast protein liquid
182 chromatography.

183 However, in almost reported studies, the collected active fractions following column
184 chromatography were pooled by the authors (Thavasi *et al.*, 2011; Sharma and Saharan,
185 2016). This might reduce the probability to discovery new biosurfactants.

186 1.4. Filtration

187 This method uses the difference in pressure between two sides of a special membrane to
188 separate biosurfactants from other molecules according to their size. Depending on the
189 membrane pores' size and the pressure, it can be microfiltration, ultrafiltration or
190 nanofiltration (Satpute *et al.*, 2010). The crude biosurfactants derived from *L. acidophilus*
191 NCIM 2903 and *L. casei* subsp. *casei* TM1B was purified from contaminants using filtration
192 through a 0.22 µm membrane pore size (Satpute *et al.*, 2019; Mouafo *et al.*, 2020). It is
193 important to highlight that, filtration is always used as a unit operation included in a
194 purification process that contains other purification techniques like TLC, column
195 chromatography, etc.

196 1.5. Dialysis

197 As a good method for sensitive compounds, dialysis enable separation of biosurfactants from
198 accompanying substances simultaneous extracted with biosurfactants such as salts. In the
199 dialysis process, molecules driven by a differential concentration gradient, are transported
200 through a semipermeable membrane based on their size. Hence, contaminants present in the
201 crude biosurfactants will pass through the semipermeable membrane leading to the
202 purification of this latter. In the purification process of biosurfactants, Vecino *et al.* (2017),
203 Ghasemi *et al.* (2018), Satpute *et al.* (2019) and Vigneshwaran *et al.* (2021) used a membrane
204 of molecular weight cutoff 6000–8000 Da to perform dialysis against double demineralized
205 water. The process has led to the obtention of pure biosurfactants.

206 Globally, the choice of the purification techniques of biosurfactants depends on the producing
207 strain, the culture media and the extraction method. There are some microorganisms that
208 produce only one molecule of biosurfactant as major metabolism compound (*Bacillus* spp.,
209 *Pseudomonas* spp.). Depending on the production mode (extracellular or cell-bound), the
210 biosurfactants is extracted from the culture media. The great challenge in the purification
211 process is to separate the biosurfactants from the culture media or cells' components which
212 can be extracted together with the biosurfactants. However, other microorganisms (lactic acid
213 bacteria) produced complex mixture of molecules with different features as biosurfactants. In
214 this case, the choice of the purification technique becomes difficult and requests that several
215 techniques should be used with special care to the screening of desired activity during the
216 process. Hence, at least three different techniques should be used in combination with TLC or
217 column chromatography to strengthen the purity of the biosurfactants (Mouafo *et al.*, 2022).

218 **2. Physicochemical characterization of biosurfactants**

219 *2.1. Identification of ionic properties of biosurfactants*

220 The different types of biosurfactants vary according to their biochemical composition. Based
221 on their large structural diversity, the biosurfactants might be charged or not. This has led to
222 the classification of biosurfactants depending on their ionic properties. Hence, the different
223 classes are: anionic, cationic, amphoteric and non-ionic. The CTAB agar method was
224 developed and was generally used to detect the production of biosurfactants of glycolipidic
225 nature (Satpute *et al.*, 2008). The method is based on the fact that the biosurfactants can form
226 an insoluble ion pair (precipitation lines) with the cationic cetyl trimethyl ammonium bromide
227 (CTAB) or anionic sodium dodecyl sulfate (SDS). Methylene blue is sometimes added in the
228 culture media to ease the observation of the precipitation line materialized as a blue halo
229 around well containing culture (Satpute *et al.*, 2008). This method was successfully used to
230 confirm the anionic nature of biosurfactants derived from *Lactococcus lactis* (Saravanakumari
231 and Mani 2010) *E. faecium* (Sharma *et al.*, 2015), and *Pseudomonas guguanensis* strain Iraqi
232 ZG.K.M (Faisal *et al.*, 2023). However, the method is not suitable for other types of
233 biosurfactants such as lipopeptides or glycoproteins.

234 2.2. Color of biosurfactants

235 The color of biosurfactants generally varies according to the producing strain, the culture
236 media composition and the extraction technique. The colors reported in the literature were
237 either whitish, yellowish or brownish (Faisal *et al.*, 2023). Fig. 2 presents the color of
238 biosurfactants produced by some lactobacilli strains (Mouafo, 2019).

239 2.3. Identification of chemical nature

240 2.3.1. Screening of proteins

241 The reaction of ninhydrin and amino acids is used to screen the presence of proteins in
242 biosurfactants. After TLC plates development, plates are dried, sprayed with ninhydrin
243 solution and kept at 90°C for 30 min. Appearance of pink or red spots indicates the presence

244 of amino acids and thus reveals the proteinaceous nature of the biosurfactants (Banerjee and
245 Ghosh, 2021; Faisal *et al.*, 2023).

246 2.3.2. Screening of sugars

247 At high temperature and in presence of α -naphthol and sulfuric acid, carbohydrates reacted,
248 leading to the formation of a pink complex. That reaction was exploited to screen the presence
249 of carbohydrates in biosurfactants. Plates obtained following TLC development of complex
250 mixture of biosurfactants are sprayed with α -naphthol solution and concentrated sulfuric acid
251 and heated at 100°C for 5 min. Appearance of pink spots indicates the presence of
252 carbohydrates in biosurfactants (Satpute *et al.*, 2010). That method was applied directly on a
253 solution of pure biosurfactants by Atta *et al.* (2009). The authors confirmed the presence of
254 carbohydrates in the biosurfactants after the apparition of a violet or purple colour between
255 the two layers formed by the solution of biosurfactants and the reagents. Sen *et al.* (2017)
256 reported that the TLC plates can be sprayed with anthrone reagent for detection of sugars
257 while Satpute *et al.* (2019) noticed that sugars detection can be assessed through TLC plates
258 spraying with diphenylamine. In the studies of Reddy *et al.* (2016), orcinol reagent was
259 sprayed on TLC plates to detect the presence of sugars in biosurfactants.

260 2.3.3. Screening of lipids

261 The presence of lipids in biosurfactants is generally assessed on TLC plates. The principle is
262 based on the reaction between iodine vapors and lipids in a close chamber that leads to the
263 development of a yellow color (Sen *et al.*, 2017; Faisal *et al.*, 2023). The method is not
264 specific as a positive result might indicate the presence of neutral or polar lipids (Atta *et al.*,
265 2009). According to de Faria *et al.* (2021) and Meena *et al.* (2021), the presence of lipids in
266 biosurfactants can be assessed through the spraying of TLC plates with water followed with

267 drying. Appearance of white spots on the TLC plates indicates the lipophilic nature of the
268 biosurfactants.

269 2.3.4. Screening of phosphate groups

270 Some biosurfactants were reported as containing phosphate groups in their constitution
271 (Rodrigues *et al.*, 2006). A rapid method to assess phospholipids in biosurfactants was
272 developed by Okpokwasili and Ibiene (2006). In that method, a solution of biosurfactants is
273 mixed with nitric acid 6M, the mixture is heated at 70°C for 30 min and a solution of
274 ammonium molybdate 5% (w/v) is added drop by drop. Apparition of a yellow color and
275 formation of a yellow precipitate at the bottom of the tube indicate the presence of
276 phospholipids.

277 2.4. Identification of the chemical composition

278 2.4.1. Protein content

279 In the literature, the protein content of biosurfactants was quantified through different
280 methods. The dye-binding method of Bradford (1976) was used by Ghasemi *et al.* (2019) and
281 Behzadnia *et al.* (2020) to assess the protein content of biosurfactants. In that method, the
282 Coomassie G-250 dye is bound to proteins leading to the formation of a blue complex that
283 absorbs at 595 nm. The dye-binding method of Bradford is relatively rapid and suitable for
284 determination of protein content of biosurfactants with high molecular weight (Smyth *et al.*,
285 2010). In the study of Morais *et al.* (2017), the method of Lowry *et al.* (1951) was used to
286 quantify protein in biosurfactants. That method combines the Biuret reagent (used to assess
287 protein through the presence of peptide bonds) and the Folin-Ciocalteu phenol reagent (used
288 to assess the residues of tryptophan and tyrosine). The intensity of the blue colored complex
289 developed is measured spectrophotometrically at 660 nm. The Kjeldahl method (AOAC,
290 1990) was used by Mouafo *et al.* (2018b) to assess the protein content of biosurfactants

291 derived from three lactobacilli strains. In that method, biosurfactant was digested with a
292 strong acid leading to the release of nitrogen which was titrated. The nitrogen content was
293 then converted to protein content using an appropriate conversion factor.

294 2.4.2. Lipid content

295 Lipids are one of the major constituents of biosurfactants. They are mostly responsible for the
296 amphiphilic nature of biosurfactants. The lipid contents of biosurfactants were assessed using
297 the method of Folch *et al.* (1957) by Ferreira *et al.* (2017) and Ghasemi *et al.* (2019) [57].
298 That method based on the solubility of lipids in solvents of different polarities. When solvent
299 is heated, it evaporated, passed through the sample, extracted lipids and carried these later into
300 the flask where they can be quantified after solvent evaporation. However, as lipids are
301 always complexed in biosurfactants with proteins (lipoproteins), carbohydrates (glycolipids)
302 or the association of proteins and carbohydrates (glycolipoproteins) (Mouafo *et al.*, 2022), the
303 most suitable method to assess their content required to break the bond that hold lipids prior
304 solvent extraction. Generally, hydrolysis with HCl 3N at 100°C for 1 h is used to release these
305 bound lipids (Morais *et al.*, 2017).

306 2.4.3. Sugar content

307 Sugars are important elements for the composition of some types of biosurfactants such as
308 glycolipids, glycoproteins, and glycolipoproteins. They can be simple sugars,
309 oligosaccharides, or polysaccharides. The most reliable method widely used in the literature
310 to assess the sugar content of biosurfactants is the phenol–sulfuric acid of Dubois *et al.*
311 (1956). In that method, hydrolyzed saccharides of biosurfactants are dehydrated into furfural
312 derivatives while reacting with concentrated sulfuric acid. The furfural derivatives obtained
313 reacted with phenol to form a colored complex that absorb light at 490 nm. The phenol–
314 sulfuric acid was successfully used by Ferreira *et al.* (2017), Mouafo *et al.* (2018a), Mouafo *et*

315 *al.* (2018b), Behzadnia *et al.* (2020), Mouafo *et al.* (2020a) and Devale *et al.* (2023) while
316 identifying the chemical nature of biosurfactants.

317 2.5. Elemental composition of biosurfactants

318 Element analysis provides information that can be used to determine the chemical nature and
319 structure of biosurfactants. The elemental composition of biosurfactants was assessed in the
320 literature with different equipment. Rodrigues *et al.* (2006), used X-ray photoelectron
321 spectroscopy (XPS) to determine the percentage of C, N, O and P in biosurfactants from
322 *Streptococcus thermophilus* A. Based on the higher amount of C and O, the authors concluded
323 on the glycolipid nature of the biosurfactants. The energy dispersive X-ray was successfully
324 used by Habib *et al.* (2023) to assess the elemental composition of biosurfactants from *P.*
325 *benzoelyticum* Pb4, *Bacillus albus* S2i and *Proteus mirabilis* Th1. Ferreira *et al.* (2017), and
326 Vecino *et al.* (2017), used a Carlo Erba EA-1108CHNS-O element analyzer to determine the
327 percentage of C, N, H and S in the biosurfactants from *L. paracasei*. The authors reported
328 respectively the glycoprotein and glycolipopeptide natures of biosurfactants based on their C,
329 N, H and S contents. The glycolipoprotein nature of biosurfactants derived from *L. paracasei*
330 subsp. *tolerans* N2 was identified by Mouafo *et al.* (2018b) using a Vario EL III elemental
331 analyzer associated with biochemical analyses.

332 **3. Identification of functional groups of biosurfactants**

333 3.1. FTIR spectroscopy

334 Fourier transform infrared spectroscopy (FTIR) is a method widely used for the
335 characterization of biosurfactants. It generally leads to determine the chemical nature of
336 biosurfactants based on the functional groups and chemical bonds that are present in the
337 biosurfactants (Jemil *et al.*, 2017). In the method, biosurfactants are submitted to radiation
338 ranging from 400 cm^{-1} to 4000 cm^{-1} . When the radiation frequency is close to the resonance

339 frequency of the biosurfactants that is analyzed, there is absorption of the luminous energy. A
340 decrease of the transmitted energy is then observed. The absorption bands obtained that vary
341 according to chemical bonds and functional groups of constituent elements of the
342 biosurfactants are characteristic of the studied biosurfactants. FTIR was used by several
343 authors to determine the chemical nature of biosurfactants (Morais *et al.*, 2017; Vecino *et al.*,
344 2017; Mouafo *et al.*, 2018b; Mouafo *et al.*, 2020a; Banerjee and Ghosh, 2021; Dabaghi *et al.*,
345 2023; Faisal *et al.*, 2023; Habib *et al.*, 2023). It is generally performed by 32 scans of the
346 compounds at a spectral resolution of 4 cm⁻¹ and a wave number accuracy between 400 and
347 4,000 cm⁻¹. Background reference used is always potassium bromide pellets.

348 *3.2. Ultraviolet spectroscopy*

349 Several microorganisms produced biosurfactants thus leading to a broad structural diversity.
350 They are mixture of lipids (saturated or unsaturated fatty acids), proteins and sugars that
351 possess several kinds of bonds which can absorb ultraviolet (UV) light in the range of 100 to
352 800 nm (Wilcox and Wilcox, 1995; Sakr *et al.*, 2021). The ultraviolet spectroscopy method
353 was used by Dehghan-Noudeh *et al.* (2005) and Ismail *et al.* (2013) to identify the
354 lipopeptidic nature of biosurfactants produced by *Bacillus* sp. Authors reported peaks at 215
355 nm corresponding to α and β -unsaturated ketones, peaks at 260 nm corresponding to aromatic
356 rings thus indicating the presence of aromatic amino acids. Based on these information's, they
357 concluded on the lipopeptidic nature of the biosurfactants. Sakr *et al.* (2021) reported that
358 biosurfactants from *L. plantarum* 60FHE, *L. paracasei* 75FHE, and *L. paracasei* 77FHE
359 absorbed in the far UV region (270-277 nm) and were all glycolipopeptides.

360 *3.3. Fatty acids profile of biosurfactants*

361 The structural diversity of biosurfactants leads to a great variation in their lipid moiety. To
362 obtain detailed structural information on that lipid moiety, the different fatty acid entering in

363 their constitution should be identified. The most suitable methods for that are GC-FID (gas
364 chromatography coupled with flame ionization detection) and GC-MS (gas chromatography
365 coupled mass spectrometry). In these methods, the lipid moiety of biosurfactants is prior
366 separated from the other moiety through a hydrolytic process that will cleave the link (Smyth
367 *et al.*, 2010). The fatty acids obtained are transformed into a volatile derivative (fatty acid
368 methyl esters) which will be easily analyzed in GC-FID using standards or in GC-MS. Fatty
369 acid methyl esters are separated according to their chain length, and their molecular mass
370 registered provide information on their structures. The fatty acids profile of biosurfactants
371 were determined by several authors in the literature using GC-MS (Sharma *et al.*, 2015;
372 Morais *et al.*, 2017; Vecino *et al.*, 2017; Behzadnia *et al.*, 2020; De Gregorio *et al.*, 2020;
373 Mouafo *et al.*, 2020a ; Faisal *et al.*, 2023).

374 *3.4. Monosaccharide's profile of biosurfactants*

375 The sugar moiety of biosurfactants is always made of monosaccharides bound through
376 glycosidic links. To identify these monosaccharides, the first step is the separation of the
377 sugar moiety of the biosurfactants from the other moiety which could be lipids, proteins, or
378 both depending on the nature of the biosurfactants. That separation is generally carried out
379 through hydrolysis with trifluoroacetic acid in a sealed tube at 120°C for 4-6 h. Then, the mixture is
380 neutralized with 2 M NH₄OH, reduced with an aqueous solution of sodium borohydride
381 (NaBH₄) and then, the sugars are transformed into volatile derivatives (N-
382 trimethylsilylimidazole, alditol acetates). The obtained derivatives are injected in GC-MS which
383 will provide structural information. The sugar moieties of biosurfactants were successfully
384 identified by Sauvageau *et al.* (2012), Morais *et al.* (2017) and Mouafo *et al.* (2020a) using
385 GC-MS.

386 *3.5. Amino acids profile of biosurfactants*

387 3.5.1. Amino acids profile

388 Amino acids are constitutive units of peptides that entering into the composition of several
389 class of biosurfactants such as lipopeptides, glycopeptides and glycolipopeptides. The peptide
390 moiety of biosurfactants is linked to other compounds through amide and lactone bonds (for
391 fatty acids chain) or N-glycosidic and O-glycosidic bonds (for sugars) (Joshi *et al.*, 2016).
392 Prior identification of the amino acids of the peptide moiety of the biosurfactants, these bonds
393 should be cloven. That cleavage generally occurs after hydrolysis at high temperature (110°C)
394 with concentrated HCl (6 N) for approximately 24 h in sealed tubes. The amino acid obtained
395 can be analyzed by GC-MS after conversion into volatile derivatives (trimethylsilylation).
396 That method provides information on the amino acid composition of the biosurfactants based
397 on the retention times and masses (m/z). GC-MS was successfully used by You *et al.* (2015)
398 to identify four amino acids (Val, Leu, Asp and Glu) in the lipopeptide from *Enterobacter* sp.
399 N18 after acid hydrolysis and trimethylsilylation. However, that method does not provide
400 information on the sequence of amino acids.

401 3.5.2. Amino acid sequence

402 Amino acids sequence of biosurfactants can be assessed using Edman degradation or
403 Quadruple-time-of-flight tandem mass spectrum (Q-TOF MS/MS). Edman degradation is the
404 earlier method reported in the literature to elucidate the composition and sequence of amino
405 acids of biosurfactants (Smyth *et al.*, 2014). The method was mostly applied for lipopeptides.
406 In that method, the biosurfactant is hydrolyzed and the smaller cleaved peptides are submitted
407 to Edman degradation. Mild alkaline hydrolysis is generally applied to open the ring of
408 lipopeptides. The Edman degradation method required peptides from the biosurfactants to be
409 purified to homogeneity (Smyth *et al.*, 2014). The Edman degradation process is conducted in
410 an automated sequencers and the cleaved amino acids are submitted to chromatography. The

411 retention time of the cleaved amino acid is compared with the one of amino acid standards for
412 identification (Zachara and Gooley, 2000).

413 Opposite to the Edman degradation technique, Q-TOF MS/MS is applied directly on the
414 entire biosurfactants without a preliminary hydrolysis step (You *et al.*, 2015). In that method,
415 the initial mass (m/z) of the biosurfactants is determined. Then, amino acids from
416 biosurfactants are lost sequentially leading to a decrease of its m/z after MS/MS
417 fragmentation. The m/z lost corresponds to the mass of the amino acid eliminated. The C-term
418 amino acid is often identified using the double hydrogen transfer mechanism of Yang *et al.*
419 (2006). In that mechanism, the m/z of the C-term amino acid is increased of m/z value of 18
420 which correspond to the m/z of water. Direct infusion in the mass spectrometer or application
421 of liquid chromatography (LC, HPLC, UPLC) leading to separation of individual peptides
422 before mass spectrometer analysis can be used. The method is less time consuming and
423 required small volume of samples (Smyth *et al.*, 2014). Korenblum *et al.* (2012) used Q-TOF
424 MS/MS to identify Glu-Leu-Leu-Val-Asp-Leu-Leu as the amino acid sequence of
425 biosurfactants produced by *Bacillus* sp. H2O-1. The peptide sequence Gly-Ser-Thr-Leu-
426 Leu-Ser-Leu-Leu was identified from the biosurfactants produced by *P. fluorescens*. BD5
427 using MALDI TOF/TOF mass spectrometry and MS/MS fragmentation (Janek *et al.*, 2010).
428 The same method was used by Pereira *et al.* (2013) to identify the sequence of the
429 heptapeptide moiety (L-Glu-L-Leu-D-Leu-L-Val-L-Asp-D-Leu-L-Leu) of the biosurfactants
430 from three strains of *B. subtilis*. Fragments indicating the sequential losses of the amino acid
431 residues Leu/Leu/Asp/Val/Leu/Leu/Glu-OMe was identified in the biosurfactants from *B.*
432 *subtilis* LSFM-05 by de Faria *et al.* (2011) while using Fourier transform ion cyclotron
433 resonance mass spectrometry with electrospray ionization (ESI-FTICR-MS).

434 **4. Molecular weight of biosurfactants**

435 The method commonly used to assess the molecular weight of biosurfactants is Sodium
436 Dodecyl Sulphate Polyacrylamide Gel Electrophoresis (SDS-PAGE). The method is
437 recommended for high molecular weight biosurfactants (Smyth *et al.*, 2010). In that method,
438 biosurfactants are separated in a gel load with a reducing buffer and SDS. They will be
439 separated according to their molecular mass (Toren *et al.*, 2001). Fouad *et al.* (2010) have
440 successfully determined the molecular weight of surlactin (a glycolipid biosurfactants)
441 produced by *L. acidophilus* using SDS-PAGE. The molecular weight of the protein fraction of
442 biosurfactants derived *L. acidophilus* NCIM 2903 was also determined by Satpute *et al.*
443 (2019) using SDS-PAGE. Tricine-SDS-PAGE was performed by Vigneshwaran *et al.* (2021)
444 to identify the molecular mass the biosurfactant *Brevibacillus* sp. AVN13

445 **5. Mass spectrum of biosurfactants**

446 Mass spectrometers provide information on the structure of biosurfactants through an
447 identification of all constitutive elements of the molecule. The method identifies both the
448 moieties of the biosurfactants without a preliminary hydrolysis. The MS spectra provides
449 information on the mass (m/z) of the protonated molecules $[M+H]^+$ of and their sodium
450 $[M+Na]^+$ or potassium $[M+K]^+$ adducts. During MS/MS fragmentation, constitutive elements
451 of the biosurfactants are released and identified based on the m/z values. The direct infusion
452 to the mass spectrometer and injection into liquid chromatography (LC) connected to a MS
453 detector system, so as 90% of each fraction is collected and 10% sent to MS detector, are
454 generally used. The only condition is the purity of the biosurfactants. For this purpose, it is
455 always recommended to perform and HLPC-UV or HPLC-DAD to confirm the purify of the
456 compound with the appearance of a well-constructed and differentiated peaks. These peaks
457 correspond to pure fractions. HPLC-ESI-MS, LC-MS/MS, Tandem MS/MS, HRMS and other
458 mass spectrometry techniques were used by several authors to determine the structure of

459 biosurfactants (de Faria *et al.*, 2011; Janek *et al.*, 2013; Pereira *et al.*, 2013; You *et al.*, 2015;
460 Oluwaseun *et al.*, 2017; Abdalsadiq *et al.*, 2018; Adu *et al.*, 2023).

461 **6. Nuclear magnetic resonance of biosurfactants**

462 NMR is an analytic technique that consists to measure the absorption of radio frequencies by
463 atoms of biosurfactants submitted to a magnetic field. That magnetic field allows the
464 resonance of atoms present in the molecule. The different resonance frequencies of the atoms
465 are consigned on a graph which permits to determine the structure of the molecule based on
466 chemical shifts (Abdel-Mawgoud *et al.*, 2011). Briefly, NMR required a principal static
467 magnetic field and a magnetic field that oscillates in the field of radio frequencies (10^6 - 10^7
468 Hz). Sample is dissolved in solvents (deuterated chloroform, methanol pyridine, acetic acid,
469 dimethyl sulfoxide acetone, benzene) and introduced into an induction coil that generates an
470 oscillating magnetic field. Depending on the nucleus present in the sample and the value of
471 the principal static magnetic field, the sample will absorb and reemit energy in a particular
472 way and at a precise frequency named resonance. Some recent NMR spectrometers are
473 equipped with triple resonance helium-cooled TCI cryoprobe (^1H , ^{13}C and ^{15}N or ^{31}P) and
474 provide information on the chemical environment of atoms of H, C, N, and P within the
475 biosurfactants (Hamerly *et al.*, 2015).

476 Proton and carbon NMR are amongst the most used techniques for structure elucidation of
477 biosurfactants. It was used by several authors independently of the biosurfactants producing
478 strain (Monteiro *et al.*, 2007; Sauvageau *et al.*, 2012; Pereira *et al.*, 2013; Sharma *et al.*,
479 2015 ; Oluwaseun *et al.*, 2017; Mouafo *et al.*, 2020 ; Dabaghi *et al.*, 2023). In the process of
480 identifying the exact position of constitutive atoms (carbons, hydrogen, nitrogen, phosphorus)
481 of the biosurfactants with respect of their chemical environment, two dimensions NMR
482 (COSY, HMBC, ROSY, and HSQC) is often used to strengthen information provided by one
483 dimension ^1H and ^{13}C NMR (Sauvageau *et al.*, 2012).

484 7. Successful structure elucidation of biosurfactants.

485 Successful characterization leading to the proposition of a chemical structure to biosurfactants
486 was achieved by some authors in the literature. Globally, the combination of at least two
487 techniques depending on the biosurfactants and the producing strains, was required.

488 Nelson *et al.* (2020) used HPLC coupled to high-resolution electrospray ionization mass
489 spectrometry to characterize the biosurfactants from *L. sakei* JN-185, *L. fermentum* JN-119
490 and *L. plantarum* JN-141. They identified several compounds including glycosyldiglycerides,
491 surfactin C13, iturin A8, octapeptin D, plantaricin A, lichenysin A, sakacin-A, glyserin,
492 plusbacin A3 and laterocin. The biosurfactants from *Pseudomonas putida* BD2 was identified
493 by Janek *et al.* (2013) using TLC and UPLC/ESI-MS/MS. They found that the biosurfactants
494 was a mixture of di-rhamnolipid (Rha-Rha-C10-C10) and phosphatidylethanolamines PE
495 (32:1), PE (33:1). You *et al.* (2015) used the combination of GC-MS, ESI-MS and Q-TOF
496 MS/MS to identify the biosurfactants from *Enterobacter* sp. N18. They reported the presence
497 of surfactin homologues such as n-C12, iso-C13, anteiso-C13, iso-C14, n-C14, iso-C15,
498 anteiso-C15, iso-C16, n-C16 and iso-C17 b-OH fatty acids. Korenblum *et al.* (2012)
499 identified surfactin analogues as biosurfactants produced by the *Bacillus* sp. H2O-1 using
500 TLC, GC-MS, ESI-MS and Tandem-MS. These compounds contained a similar heptapeptide
501 chain of surfactin (Glu-Leu-Leu-Val-Asp-Leu-Leu) as the hydrophilic moiety, and C13, C14,
502 C15, C16 β -hydroxy-fatty acids as the hydrophobic moiety. TLC, GC-MS and MALDI TOF-
503 MS/MS was used to identify biosurfactants derived from *P. fluorescens* BD5 (Janek *et al.*,
504 2010). The authors reported two cyclic peptides namely pseudofactin I (palmitoyl-Gly-Ser-
505 Thr-Leu-Leu-Ser-Leu-Val-O-) and pseudofactin II (palmitoyl-Gly-Ser-Thr-Leu-Leu-Ser-
506 Leu-Leu-O-).

507 The biosurfactants from *B. subtilis* 309, *B. subtilis* 311 and *B. subtilis* 573 were identified by
508 Pereira *et al.* (2013) using FTIR-ATR, ^1H NMR and MALDI-TOF MS/MS. They found that

509 the biosurfactants were made of a heptapeptide (L-Glu-L-Leu-D-Leu-L-Val-L-Asp-D-Leu-L-
510 Leu) linked to a C13, C14 or C15 hydroxy fatty acid chain. Lichenysin-A, is a compound
511 made of a small peptide (Gln, Leu, Leu, Val, Asp, Leu, and Ile) that was linked to 3-hydroxy
512 fatty acid residue (3-hydroxylated tri, tetra, penta, or hexadecanoic acids) with amide and
513 lactone bonds forming a cyclic structure, was identified as the biosurfactants produced by
514 *Bacillus licheniformis* W16 while using FTIR, HPTLC-ESI-MS and MALDI-TOF-MS, ¹H
515 and ¹³C NMR as analytical techniques. In the study conducted by Oluwaseun *et al.* (2017), L-
516 rhamnosyl-L-rhamnosyl-3-b-hydroxydodecenoate was identified as the biosurfactants from
517 *Pseudomonas aeruginosa* C1501 using LC-ESI-MS and ¹H and ¹³C NMR.

518 Saravanakumari and Mani (2010) used GC-MS and ¹H NMR to identify O-methyl-β-D-
519 xylopyranoside and octadecanoic acid was the major constitutive elements of the
520 biosurfactants from *L. lactis*. The final structure proposed by the authors was 2-methyl-O-
521 methyl-β-D-xylopyranosyl octadecanoic acid. The combination of TLC, GC-MS, ESI-MS and
522 NMR (¹H and ¹³C) at one and two dimensions (COSY, HMBC and HSQC) was used in the
523 characterization process of biosurfactants from *L. plantarum* (Sauvageau *et al.*, 2012). The
524 authors identified four fractions including α-D-Glucopyranosyl-diglyceride (GL1), α-D-
525 Galactopyranosyl-(1→2)-α-D-Glucopyranosyl-diglyceride (GL2a), β-D-Glucopyranosyl-
526 (1→6)-β-D-Galactopyranosyl-(1→2)-6-O-acyl-α-D-Glucopyranosyl-diglyceride (GL2b) and
527 β-D-Glucopyranosyl-(1→6)-α-D-Galactopyranosyl-(1→2)-α-D-Glucopyranosyl-diglyceride
528 (GL3). Xylopyranosyl linked to octadecanoic acid was identified as the biosurfactants from *L.*
529 *helveticus* MRTL91 while using TLC, UPLC-ESI-MS, GC-MS, FTIR and NMR (¹H and ¹³C)
530 (Sharma *et al.*, 2014). Using the same techniques latter, the authors identified xylopyranosyl
531 β-hydroxydecanoic acid as the biosurfactants produced by *E. faecium* MRTL9 (Sharma *et al.*,
532 2015). Mouafo *et al.* (2020a) identified 2,5-O-methyl-rhamnofuranosyl-palmitate in

533 biosurfactants from *L. casei* TM1B using elemental analysis, FTIR, GC-MS, and NMR (¹H
534 and ¹³C).

535 **Conclusion**

536 This review presents the different techniques of purification of biosurfactants. It suggests that,
537 the choice of the purification techniques of biosurfactants depends on the producing strain, the
538 culture media composition and the extraction method. Chromatographic, spectrometric and
539 mass spectrometry methods are suitable for fully characterization at structural level of
540 biosurfactants. However, biosurfactants which are known as complex mixture of compounds
541 must be fractionated before being analyzed. They are in the food industry to enhance the shelf
542 live of food, remove biofilms from food processing surfaces, stabilize emulsions, improve
543 dough stability and texture of bakery products. In agricultural field, they are used as
544 biopesticides. In environmental field, they are used for bioremediation, to enhance oil
545 recovery and remove heavy metals from contaminated soils. They are also used in cosmetic,
546 detergency and painting. The full structural characterization of biosurfactants will improve
547 their added values and their application in some highly pointed industrial fields such as
548 medicine and pharmacy.

549

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551 Conceptualization, A.M. and H.T.M.; methodology, investigation, data curation, writing—
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560 The authors declare no conflict of interest.

561

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Table 1. Purification methods and nature of biosurfactants produced by some microorganisms

Microorganisms	Method of purification	Nature of biosurfactants	References
<i>L. casei</i> subsp. <i>casei</i> TM1B	- Centrifugation (13,000×g, 4°C, 20 min) - Filtration (0.22 µm) - RP-HPLC	Glycolipid	(Mouafo <i>et al.</i> , 2020)
<i>L. plantarum</i> CFR 2194	- Centrifugation (10,000×g, 4 °C, 15 min) - Precipitation (pH 2)	Glycoprotein	(Nelson <i>et al.</i> , 2020)
<i>B. velezensis</i> KLP2016	- TLC - Size exclusion chromatography - Ion exchange chromatography	Lipopeptide	(Dabaghi <i>et al.</i> , 2023)
<i>B. oceanisediminis</i> H2	- TLC - Column chromatography	Lipopeptide	(Meena <i>et al.</i> , 2021)
<i>B. subtilis</i> LSFM-05	- Column chromatography	Lipopeptide	(Satpute <i>et al.</i> , 2010)
<i>S. thermophilus</i> A	- Filtration (0.22 µm) - Dialysis (6000–8000 Da) - Hydrophobic interaction chromatography	Glycolipid	(Sauvageau <i>et al.</i> , 2012)
<i>L. plantarum</i> IRL- 560	- TLC - Column chromatography - Size exclusion chromatography	Glycolipid	(Banerjee and Ghosh, 2021)
<i>L. delbrueckii</i>	- Column chromatography - Dialysis (6000–8000 Da)	Glycolipid	(Sharma and Saharan, 2016)
<i>L. helveticus</i> MRTL91	- Dialysis (6000–8000 Da) - Column chromatography	Glycolipid	(de Faria <i>et al.</i> , 2021)
<i>L. lactis</i>	- Precipitation (pH 2) - Centrifugation (6000×g, 15 min, 4°C)	Glycolipid	(Sharma <i>et al.</i> , 2015)
<i>L. rhamnosus</i> PTCC 1637	- Filtration (0.22 µm) - Dialysis (6000–8000 Da)	Phosphoglycoprotein	(Satpute <i>et al.</i> , 2008)
<i>L. paracasei</i> ssp. <i>paracasei</i> A20	- Filtration (0.22 µm) - Dialysis (6000–8000 Da)	Glycolipoprotein	(Ghasemi <i>et al.</i> , 2018)
<i>L. acidophilus</i> NCIM 2903	- Filtration (0.22 µm) - Dialysis (6000–8000 Da)	Glycolipoprotein	(Vecino <i>et al.</i> , 2017)

List of Figures

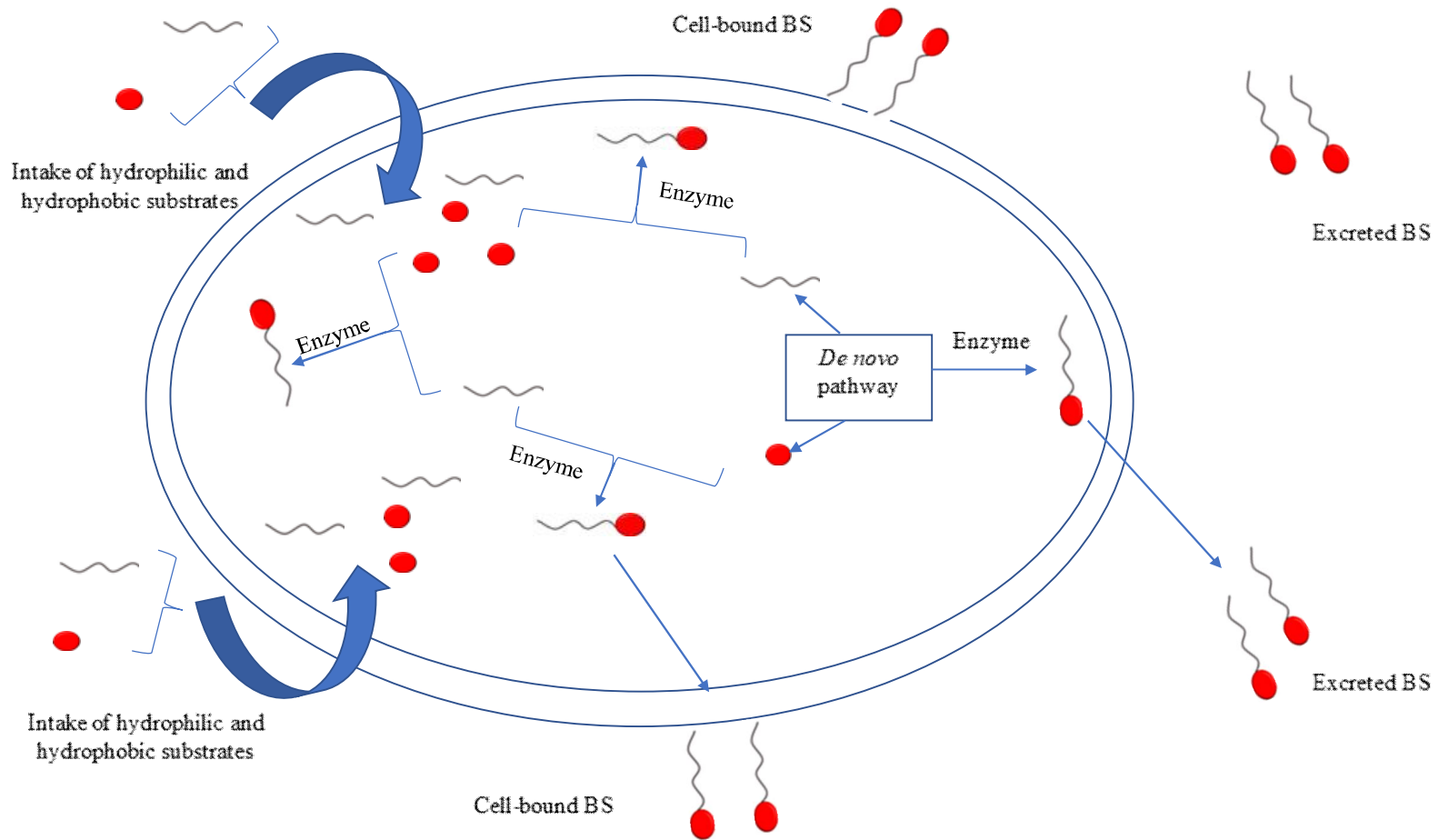


Fig. 1. The different metabolism routes of biosurfactants by microorganisms.

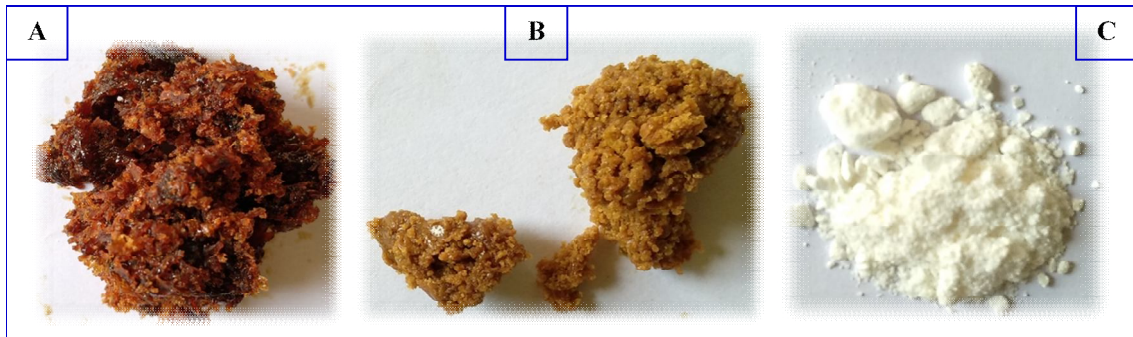


Fig. 2. Pictures showing color of biosurfactants produced by *Lactobacillus paracasei* subsp. tolerans N2 with sugar cane molasses as substrate (A), *Lactobacillus casei* subsp. casei TM1B with sugar cane molasses as substrate (B) and *Lactobacillus rhamnosus* G88 with glycerol as substrate (C) (Mouafo, 2019).