



# New insight into the role of oxygen supply for surfactin production in bench-scale bioreactors using induced surface aeration

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## Abstract

Surfactin biosurfactant produced by *Bacillus* sp. has been studied, because it has enormous potential in several applications in the oil and cosmetics industry. The cultivation conditions for obtaining this bioproduct, however, still require attention, as, for example, parameters related to oxygen supply and consumption. In this study, different volumetric oxygen transfer coefficient ( $K_L a$ ) levels (0–11.56 h<sup>-1</sup>) were tested in bench-scale bioreactor for surfactin biosurfactant production by *Bacillus velezensis* H2O-1, using induced surface aeration. While conditions close to anaerobiosis showed insignificant production of surfactin, an intermediated  $K_L a$  condition (4.24 h<sup>-1</sup>) generated the best surfactin concentration (579.6 mg L<sup>-1</sup>), with a volumetric productivity of 11.9 mg L<sup>-1</sup> h<sup>-1</sup>. These results showed that the oxygen demand to produce surfactin is not high, being possible to use induced surface aeration strategy in bioreactors, minimizing foam formation. In addition, in all  $K_L a$  conditions tested, surfactin homologues C<sub>14</sub> and C<sub>15</sub> had higher relative abundance. Nevertheless, the  $K_L a$  parameter seems to have had minimal influence on affecting the relative abundances of surfactin homologues produced. Particularly noteworthy in this study is the possibility of producing surfactin using a low-cost and scale-up feasible aeration strategy, unlike the foam collection strategies developed in other studies to obtain this bioproduct.

**Keywords** Surfactin · Induced surface aeration · Bioprocess scale-up

## Introduction

Biosurfactants are amphiphilic molecules with surface-active properties produced by biological systems, mainly microorganisms. When compared to synthetic surfactants, biosurfactants have several advantages, such as low toxicity, biodegradability and the possibility of being synthesized from renewable raw material [1]. On the other hand, the main disadvantages of biosurfactants in comparison with synthetic surfactants is the high production costs and scale-up difficulties [2].

Among the different types of biosurfactants, cyclic lipopeptides (CLPs) have interesting characteristics, such as the fact that they present incredibly low values of critical micellar concentration (CMC) and the fact that they are generally

produced by *Bacillus* sp., non-pathogenic microorganisms [3]. Structurally, CLPs molecules have a fatty acid in combination with a peptide moiety. Several CLPs families and homologues are normally produced by the same strain [4]. These CLPs differ by the amino acid composition of the peptide moiety, the length of the fatty acid chain, and the link between the two parts [5]. Among others, surfactin, iturin, fengicin, kurstakin and polymixin can be mentioned among Biosurfactant LPs families [5]–[7], with the surfactin family being the most studied [8].

Despite the great interfacial properties and potential applications of surfactin that have been recognized in the literature for some time—this molecule was discovered in 1969 [9], this bioproduct production in industrial scale is not yet a reality. This is mainly due to the difficulty of supplying oxygen to the surfactin production bioprocess, avoiding excessive foam formation. Several studies report the use of foam collection systems on bioreactors [10]–[14]. Some authors suggest the use of aeration through membranes [15]–[17]. These technologies, despite being indeed

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successful in controlling excessive foaming, have high costs and scale-up complexity.

In addition, the difficulty in supplying oxygen to the bioprocess to obtain surfactin has led the scientific community to a lack of information regarding the real oxygen demand for the formation of this bioproduct. Several works about culture medium optimization have already been published [5, 18, 19], but few have studied the role of oxygenation in the production of CLPs. Jokari et al. (2012) [20] studied the effect of aeration rate on surfactin production in shaken flasks with specific designs, reporting a surfactin volumetric productivity of  $49 \text{ mg L}^{-1} \text{ h}^{-1}$  using a  $K_L a$  of  $259.56 \text{ h}^{-1}$ . Yeh et al. (2006) [10] reported a foam collection and cell recycling system, capable of producing  $106 \text{ mg L}^{-1} \text{ h}^{-1}$  of surfactin at an optimized  $K_L a$  of  $47.52 \text{ h}^{-1}$  reaching a maximum bioproduct concentration of  $6.45 \text{ g L}^{-1}$  on the collected foam. Nazareth et al. (2021) [12] reported a volumetric productivity of  $9.98 \text{ mg L}^{-1} \text{ h}^{-1}$  for an optimized  $K_L a$  of  $45.41 \text{ h}^{-1}$ , tested in a benchtop bioreactor coupled to a foam trap, reaching a surfactin concentration of  $239.74 \text{ mg L}^{-1}$  considering the total volume of the test ( $2159.84 \text{ mg L}^{-1}$  of surfactin in the collected foam). Interestingly, due to the operational difficulty of supplying oxygen without foam excessive generation, Hoffmann et al. (2020) [21] studied the anaerobic production of surfactin in bioreactors by nitrate respiration, reaching  $190 \text{ mg L}^{-1}$  of surfactin. However, under anaerobic culture conditions, a higher production of acetate has been reported, impairing cell growth. Hoffmann et al. (2020) [21] conducted a specific experiment to evaluate *Bacillus subtilis* cellular growth with different initial concentrations of acetate added artificially. While in the control condition (no acetate added) the  $\mu_{\text{max}}$  was  $0.123 \text{ h}^{-1}$ , with an initial concentration of  $0.5 \text{ g L}^{-1}$  acetate the  $\mu_{\text{max}}$  dropped to  $0.095 \text{ h}^{-1}$ . Thus, the search for alternative strategies to reduce foam formation allows the reduction of expenses with defoamer and with the foam separation step, in addition to enabling the use of conventional bioreactors to produce this biomolecule.

Another issue to be considered is whether the supply of oxygen to the bioprocess may be altering the relative quantity of surfactin homologues or other CLPs produced. Many published works report the variety of CLPs produced by new *Bacillus sp.* strains discovered [22]–[24]. Some authors also describe changes in the relative quantities of CLP families produced according to culture media composition variation [18, 19]. More deeply, the peptide biodiversity of the surfactin biomolecules may be related to the peptide's synthesis by multienzymatic proteins called nonribosomal peptide synthetases. This biodiversity also depends on the fatty acid moiety, which length can vary from 12 to 17 carbons atoms, with fifteen different isoforms, enlarging the surfactin structures possibility [9]. However, no studies were found correlating different bioprocess oxygenation levels with changes

in the production of different surfactin homologues or other CLPs.

It is interesting to note that, if hypothetically the oxygen demand to produce surfactin is not high, it is possible to use induced surface aeration [25] strategies in bioreactors, injecting air into the bioreactor headspace instead of using the spargers and also adjusting an impeller near the liquid surface to increase oxygenation. Naturally, it is not possible to obtain a high  $K_L a$  value with this aeration method, but it is a simple, inexpensive and easily scalable technique.

The aim of this study, therefore, was to evaluate the surfactin production by *Bacillus velezensis* H2O-1 in bench-scale bioreactors using induced surface aeration, at different oxygenation levels, making it possible to correlate different  $K_L a$  with overall surfactin productivity and with the relative abundance of different surfactin and other CLPs homologues.

## Materials and methods

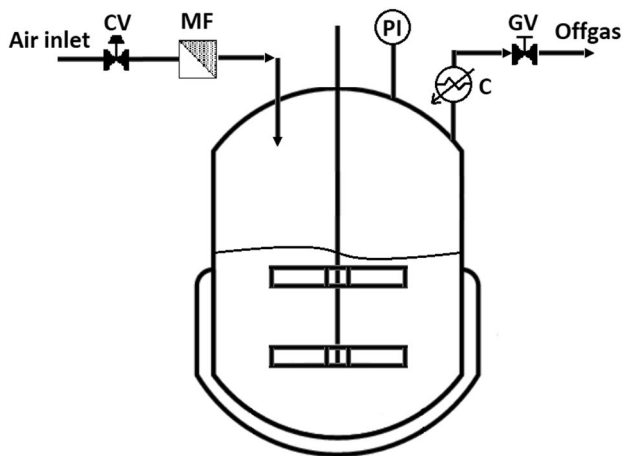
### Microorganism and culture medium

The surfactin producer strain *Bacillus velezensis* H2O-1, originally isolated from an oil reservoir in Brazil and previously described by Korenblum et al. [26, 27] was used in bioreactor experiments. The microorganism was kept in a freezer at  $-80^\circ \text{C}$  in Luria–Bertani Miller (LB) broth, added with 20% glycerol.

The culture medium previously optimized for surfactin production in shake flasks by *Bacillus velezensis* H2O-1 was used in this work [28, 29], having the following composition (% w/v): glucose 1.0;  $\text{Na}_2\text{HPO}_4$  0.5;  $\text{KH}_2\text{PO}_4$  0.2;  $(\text{NH}_4)_2\text{SO}_4$  0.1. In addition, the biomass propagation medium was supplemented with yeast extract (0.05% w/v).

### Bioreactor design and culture conditions

As represented in Fig. 1, an induced surface aeration system was used in the surfactin production experiments. A 2 L New Brunswick Bioflo 310 benchtop bioreactor was used. The vessel had a height-to-diameter ratio  $H/D = 2$ , allowing a large headspace surface area. The agitation system had two flat blade impellers, with a diameter one-half of the vessel diameter. One of the impellers was set near the liquid surface to increase oxygenation from induced surface aeration [25]. The vessel also had four equally spaced baffles to improve mixing and reduce vortex formation. A controlled air flow was fed directly into the bioreactor headspace, passing through a control valve (CV) and a  $0.22 \mu\text{m}$  membrane filtration (MF). Exhausted air was removed by passing through a condenser (C) and a manually controlled globe valve (GV). Headspace pressure



**Fig. 1** Bioreactor setup diagrammatic representation for induced surface aeration experiments. CV stands for control valve; MF for membrane filtration; PI for pressure indicator; C for condenser and GV for globe valve

was measured by a pressure indicator (PI). The fact that no air bubbles into the liquid prevents most of the excessive foaming. The agitation rate was controlled by a motor driven from the top of the vessel. The vessel was equipped with probes for temperature control and dissolved oxygen (DO) measurement (not represented in Fig. 1). Samples were taken via a sample valve (not shown in Fig. 1) for offline analysis of substrate and product concentrations.

The medium culture volume inside the vessel was 0.9 L and the bioreactor was inoculated with 100 mL of a preinoculum previously grown at 30 °C and 170 rpm for 16 h. The temperature was maintained at 30 °C for every condition tested. Each condition was performed in duplicate and ended with 48 cultivation hours. The volumetric mass transfer coefficient  $K_La$ , that describes the efficiency with which oxygen can be delivered to a bioreactor for a given set of operating conditions, tested in each condition ranged between  $0.97 \text{ h}^{-1}$  and  $11.56 \text{ h}^{-1}$  by varying aeration rate (into the bioreactor headspace) and stirring speed (Table 1). A condition close to anaerobiosis was also tested (condition A), where there was no air supply throughout the process and the culture medium dissolved oxygen was set to zero by bubbling  $\text{N}_2$  prior to the grown pre-inoculum addition. No antifoaming agent was added in any experiment.

$K_La$  determination was performed by a static gassing out technique [30], prior to the surfactin production experiments. In a cell-free culture medium, DO concentration is lowered by gassing the liquid out with nitrogen gas. Aeration is then initiated at a constant stirring speed and air flow rate (according to Table 1) and the DO concentration increase is monitored using dissolved  $\text{O}_2$  electrode. The plot of  $\ln(C_L^* - C)$  versus time will produce a straight line, where the

**Table 1** Aeration conditions tested, varying aeration rate (into the headspace) and bioreactor stirring speed

| Condition | Agitation rate (RPM) | Air flow (L/min) | $K_La$ ( $\text{h}^{-1}$ ) |
|-----------|----------------------|------------------|----------------------------|
| A         | 150                  | 0                | 0                          |
| B         | 150                  | 0.5              | 0.97                       |
| C         | 200                  | 1                | 2.07                       |
| D         | 400                  | 1                | 4.24                       |
| E         | 500                  | 2                | 5.89                       |
| F         | 600                  | 3                | 11.56                      |

Condition A had no air supply throughout the process and the culture medium dissolved oxygen was set to zero by bubbling  $\text{N}_2$  prior to the pre-inoculum

slope is equal to  $K_La$ . ( $C$  is the concentration of DO in the culture medium,  $C_L^*$  is the saturated DO concentration).

The oxygen transfer rate (OTR) was calculated as  $\text{OTR} = K_La(C_L^* - C)$ . Oxygen uptake rate (OUR) was calculated using Eq. (1):

$$\text{OUR} = \text{OTR} - \frac{dC}{dt} \quad (1)$$

The specific oxygen uptake rate (SOUR) was calculated by dividing the OUR by the biomass concentration as shown in Eq. (2). Both OUR and SOUR were calculated in the first hours of the stationary phase of microbial growth, where surfactin production is most pronounced:

$$\text{SOUR} = \frac{\text{OUR}}{X} \quad (2)$$

### Surfactin, glucose, and biomass quantification

Surfactin production was quantified as previously described by Guimarães et al. (2019) [28]. Bioreactor samples were centrifuged at  $12,500 \times g$  at 4 °C for 20 min. The supernatant was diluted (2 ×) in acetonitrile, filtered through a 0.22  $\mu\text{m}$  membrane and injected into a C18 column (150 × 4.6 mm, 5  $\mu\text{m}$  particle size; ZORBAX) in an Agilent 1200 high-performance liquid chromatography system (HPLC) to separate the surfactin homologues according to Sousa et al. (2012) [31]. The mobile phase used was 20% trifluoroacetic acid (3.8 mM) and 80% acetonitrile, the flow rate was set at  $1.0 \text{ mL min}^{-1}$ , and the system temperature was 30 °C. Detection was performed at 210 nm and injection volume was 20  $\mu\text{L}$ . The relative peaks of the different surfactin isoforms were integrated and quantified according to the calibration curve obtained with standard surfactin (Sigma-Aldrich, Ref: S3523).

Glucose consumption was analyzed directly by HPLC (HPLC Agilent Technologies) equipped with HPX-87H

column (BioRad—300 mm × 7.8 mm) and refractive index detector. H<sub>2</sub>SO<sub>4</sub> 5 mM was used as mobile phase at a flow rate of 0.6 mL min<sup>-1</sup>. Column temperature was set at 40 °C. Quantification was based on a calibration curve obtained with standard glucose (Sigma-Aldrich, Ref: G6918).

Cell growth was assessed by measuring absorbance at 600 nm. The absorbance value was converted to dry cell weight (DCW) concentration using a standard curve [DCW (g L<sup>-1</sup>) = 0.715 × ABS; R<sup>2</sup> = 0.988].

### Surface tension and critical micellar concentration measurement

Surface tension (ST) and critical micellar concentration (CMC) were analyzed using the pendant drop method (Krüss DSA100 goniometer, Model OF 3210) according to Song and Springer (1996) [32]. Pendant drops ST measurements were performed at 23 °C, with the relative humidity maintained at 55%. The same method was used to determine the CMC by serial dilutions of the biosurfactant according to Sheppard and Mulligan (1987) [33].

### Surfactin relative abundance

The detection of CLPs in the supernatant samples was performed according to a methodology similar to that presented by Liu et. al (2007) [34]. Aliquots of the cell-free supernatant were diluted 3 × in a 1:1 (v/v) methanol/acetonitrile solution containing 0.05% formic acid (v/v) and subjected to direct injection (3 µL/min flow) in the Bruker Maxis Impact spectrometer with ESI-Q-TOF-MS (electrospray ionization quadrupole-time-of-flight mass spectrometry) configuration. As electrospray ionization source a capillary voltage of 4500 V, 180 °C, was used and an acquisition rate of 1 Hz and a scanning range of 100 to 2000 m/z. Detection occurred using the positive ion mode. Data-dependent acquisition (DDA/AutoMS) was used as the data acquisition mode, with collision energy from 30 to 100 eV.

For CLP biosurfactant identification and relative abundance determination, the peak quantification was performed using the mMass software [35] from a CLP biosurfactant mass library adapted from Price et. al (2007), Sarwar et al. (2017), Liu et al. (2007), Gu. et Al. (2017) and Kaisermann (2017) [23, 34, 36]–38 and presented in Table 2.

### Data analysis

A spreadsheet software (Microsoft® Excel®) was used to analyze the obtained data. Means, standard deviations (SD) and variance values were calculated. Single factor analysis of variance (ANOVA) was used to determine significant differences between treatments at  $p < 0.05$ .

**Table 2** Molecular masses considered for cyclic lipopeptides detection by mass spectrometry

| Cyclic Lipopeptide             |  | Mass (m/z)                   |
|--------------------------------|--|------------------------------|
| Family                         | Homologue  |                              |
| Surfactin                      | Surfactin Leu/Ile-7 C <sub>12</sub>  | 993.64                       |
|                                | Surfactin Leu/Ile-7 C <sub>13</sub>  | 1007.65                      |
|                                | Surfactin Leu/Ile-7 C <sub>14</sub>  | 1021.66                      |
|                                | Surfactin Leu/Ile-7 C <sub>15</sub>  | 1035.68                      |
|                                | Surfactin Leu/Ile-7 C <sub>16</sub>  | 1049.70                      |
|                                | Iturin   | Iturin Asn-1 C <sub>10</sub> |
| Iturin Asn-1 C <sub>12</sub>   |  | 1014.51                      |
| Iturin Asn-1 C <sub>14</sub>   |  | 1042.54                      |
| Iturin Asn-1 C <sub>15</sub>   |  | 1056.56                      |
| Bacillomycin D C <sub>14</sub> |  | 1030.51                      |
| Bacillomycin D C <sub>15</sub> |  | 1044.51                      |
| Fengycin                       | Fengycin Val-7 C <sub>16</sub>   | 1462.80                      |
|                                | Fengycin Val-7 C <sub>17</sub>   | 1476.82                      |
|                                | Fengycin Val-7 C <sub>18</sub>   | 1490.84                      |
| Kurstakin                      | Kurstakin 1 (C <sub>39</sub> H <sub>64</sub> N <sub>11</sub> O <sub>12</sub> )   | 878.47                       |
|                                | Kurstakin 2/3 (C <sub>40</sub> H <sub>66</sub> N <sub>11</sub> O <sub>12</sub> ) | 892.41                       |
|                                | Kurstakin 4 (C <sub>41</sub> H <sub>68</sub> N <sub>11</sub> O <sub>12</sub> )   | 906.5                        |

[M+H]<sup>+</sup>, [M+Na]<sup>+</sup> and [M+K]<sup>+</sup> ions were considered at 1.008, 22.9898 and 38.9637 above the molecular masses, respectively

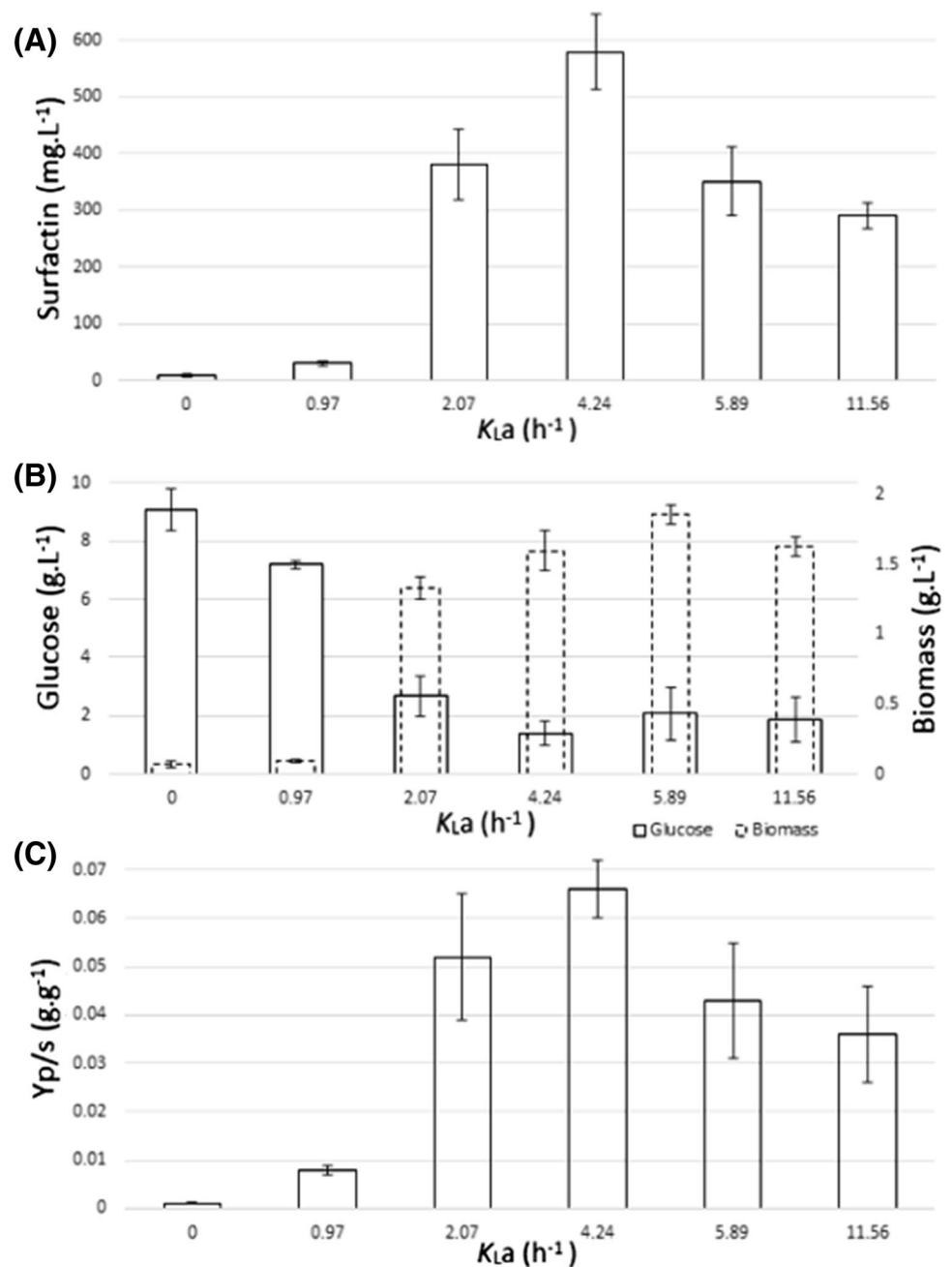
## Results and discussion

### K<sub>L</sub>a effect on surfactin production

Figure 2A, B and C illustrates the obtained results of surfactin final concentration, biomass and glucose final concentration, and Y<sub>P/S</sub>, respectively, for each K<sub>L</sub>a value tested. All results presented in Fig. 2 were obtained with a cultivation time of 48 h. Summary of each process parameters obtained are shown in Table 3. ST and CMC results obtained for each K<sub>L</sub>a tested are presented in Table 4. The highest surfactin concentration (579.6 mg L<sup>-1</sup>) and volumetric productivity (11.90 mg L<sup>-1</sup> h<sup>-1</sup>) were obtained in the K<sub>L</sub>a of 4.24 h<sup>-1</sup>. Interestingly, at the highest tested K<sub>L</sub>a level of 11.56 h<sup>-1</sup>, a lower production of surfactin was observed: 290.96 mg L<sup>-1</sup> final surfactin concentration, and a volumetric productivity of 5.88 mg L<sup>-1</sup> h<sup>-1</sup>. On the other hand, obtained values for the yield factor Y<sub>P/S</sub> showed that there was no statistically significant difference ( $p < 0.05$ ) for the conditions of K<sub>L</sub>a 2.07 to 11.56 h<sup>-1</sup>, indicating similarity in the glucose uptake for surfactin formation.

Except for the condition close to anaerobiosis (K<sub>L</sub>a = 0 h<sup>-1</sup>), no significant differences ( $p < 0.05$ ) were observed in the measured physicochemical parameter CMC (Table 4), being the reported CMC values of 15–19 mg.L<sup>-1</sup> consistent with those commonly described in the literature [28, 39]. As for the TS measured values, K<sub>L</sub>a conditions

**Fig. 2** **A** Final surfactin concentration for each  $K_La$  condition tested in a bioreactor, with 48 h of cultivation time. **B** Final biomass and glucose concentration for each  $K_La$  condition tested in a bioreactor, with 48 h of cultivation time.; **C** Final  $Y_{p/s}$  for each  $K_La$  condition tested in a bioreactor, with 48 h of cultivation time



**Table 3** Summary of process parameters obtained for each  $K_La$  tested

| $K_La$ (h <sup>-1</sup> ) | SOUR (mgO <sub>2</sub> gX <sup>-1</sup> h <sup>-1</sup> ) | $P_f$ (mg L <sup>-1</sup> )   | $Q_p$ (mg L <sup>-1</sup> h <sup>-1</sup> ) | $Y_{p/s}$ (g g <sup>-1</sup> ) | $Y_{p/x}$ (g g <sup>-1</sup> ) |
|---------------------------|---|-------------------------------|---|--------------------------------|--------------------------------|
| 0                         | –   | 9.31 ± 1.18 <sup>(a)</sup>    | 0.02 ± 0.01 <sup>(a)</sup>                  | 0.001 ± 0.0001 <sup>(a)</sup>  | –                              |
| 0.97                      | –   | 31.14 ± 3.91 <sup>(b)</sup>   | 0.47 ± 0.08 <sup>(b)</sup>                  | 0.008 ± 0.001 <sup>(b)</sup>   | 0.281 ± 0.002 <sup>(a)</sup>   |
| 2.07                      | 7.10 ± 0.13 <sup>(a)</sup>                                | 381.18 ± 62.13 <sup>(c)</sup> | 7.76 ± 1.29 <sup>(c)</sup>                  | 0.052 ± 0.013 <sup>(c)</sup>   | 0.317 ± 0.030 <sup>(b)</sup>   |
| 4.24                      | 7.35 ± 0.18 <sup>(a)</sup>                                | 579.60 ± 66.53 <sup>(d)</sup> | 11.9 ± 1.59 <sup>(d)</sup>                  | 0.066 ± 0.006 <sup>(c)</sup>   | 0.373 ± 0.098 <sup>(b)</sup>   |
| 5.89                      | 7.58 ± 0.19 <sup>(a)</sup>                                | 350.91 ± 59.64 <sup>(c)</sup> | 7.13 ± 1.18 <sup>(c)</sup>                  | 0.043 ± 0.014 <sup>(c)</sup>   | 0.187 ± 0.011 <sup>(c)</sup>   |
| 11.56                     | 13.72 ± 0.21 <sup>(b)</sup>                               | 290.96 ± 22.89 <sup>(c)</sup> | 5.88 ± 0.47 <sup>(c)</sup>                  | 0.036 ± 0.011 <sup>(c)</sup>   | 0.176 ± 0.007 <sup>(c)</sup>   |

± indicates SD. Values in the same column that are followed by the same letter are not significantly different ( $p < 0.05$ )

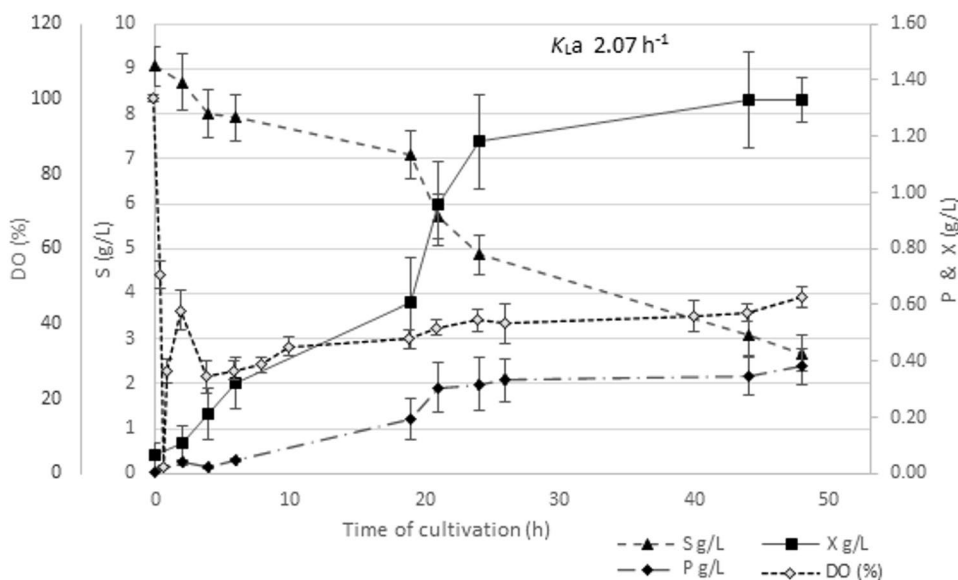
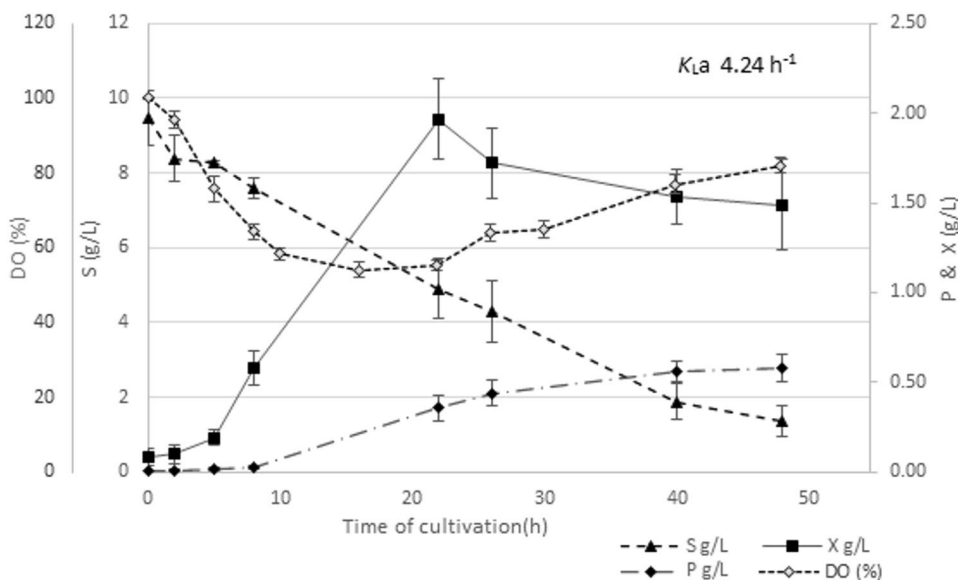
**Table 4** ST and CMC results obtained for each  $K_L a$  tested

| $K_L a$ ( $\text{h}^{-1}$ ) | ST ( $\text{mN m}^{-1}$ ) | CMC ( $\text{mg L}^{-1}$ ) |
|-----------------------------|---------------------------|----------------------------|
| 0                           | $47.7 \pm 0.13^{(a)}$     | –                          |
| 0.97                        | $29.1 \pm 0.02^{(b)}$     | $15.57 \pm 1.95^{(a)}$     |
| 2.07                        | $26.94 \pm 0.41^{(c)}$    | $15.52 \pm 1.90^{(a)}$     |
| 4.24                        | $28.3 \pm 0.28^{(b)}$     | $19.30 \pm 2.55^{(a)}$     |
| 5.89                        | $29.21 \pm 0.37^{(b)}$    | $17.55 \pm 2.10^{(a)}$     |
| 11.56                       | $27.2 \pm 0.42^{(c)}$     | $14.55 \pm 1.14^{(a)}$     |

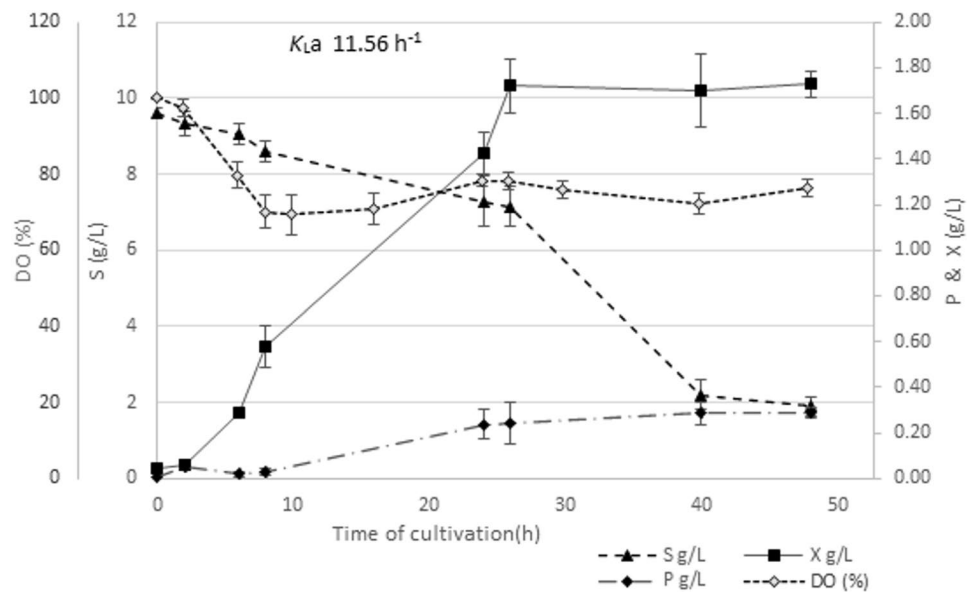
$\pm$  indicates SD. Values in the same column that are followed by the same letter are not significantly different ( $p < 0.05$ )

$2.07\text{--}11.56 \text{ h}^{-1}$  were statistically equal ( $p < 0.05$ ), and lower than the TS values obtained for the other  $K_L a$  conditions. Interestingly, even at the lowest surfactin concentration ( $9.31 \text{ mg L}^{-1}$ ), the surface tension lowering was considerably effective ( $47.7 \text{ mN m}^{-1}$ ) compared with others biosurfactants, as rhamnolipids, for example, that needs higher concentrations to promote this effect. The literature reports that the uniqueness of the peptide loop present in surfactin provides to it an excellent molecular action at the interfaces [9].

Figures 3, 4 and 5 show the kinetics of surfactin (P) production, glucose (S) consumption and biomass (X) generation for  $K_L a$  condition of 2.07, 4.24 and  $11.56 \text{ h}^{-1}$ , respectively. In the kinetics presented, it is remarkable to notice that the oxygen consumption is in fact low in the bioprocess.

**Fig. 3** Kinetics of surfactin (P) production, glucose (S) consumption and biomass (X) generation for the  $K_L a$   $2.07 \text{ h}^{-1}$  condition**Fig. 4** Kinetics of surfactin (P) production, glucose (S) consumption and biomass (X) generation for the  $K_L a$   $4.24 \text{ h}^{-1}$  condition

**Fig. 5** Kinetics of surfactin (P) production, glucose (S) consumption and biomass (X) generation for the  $K_L a$  11.56  $\text{h}^{-1}$  condition



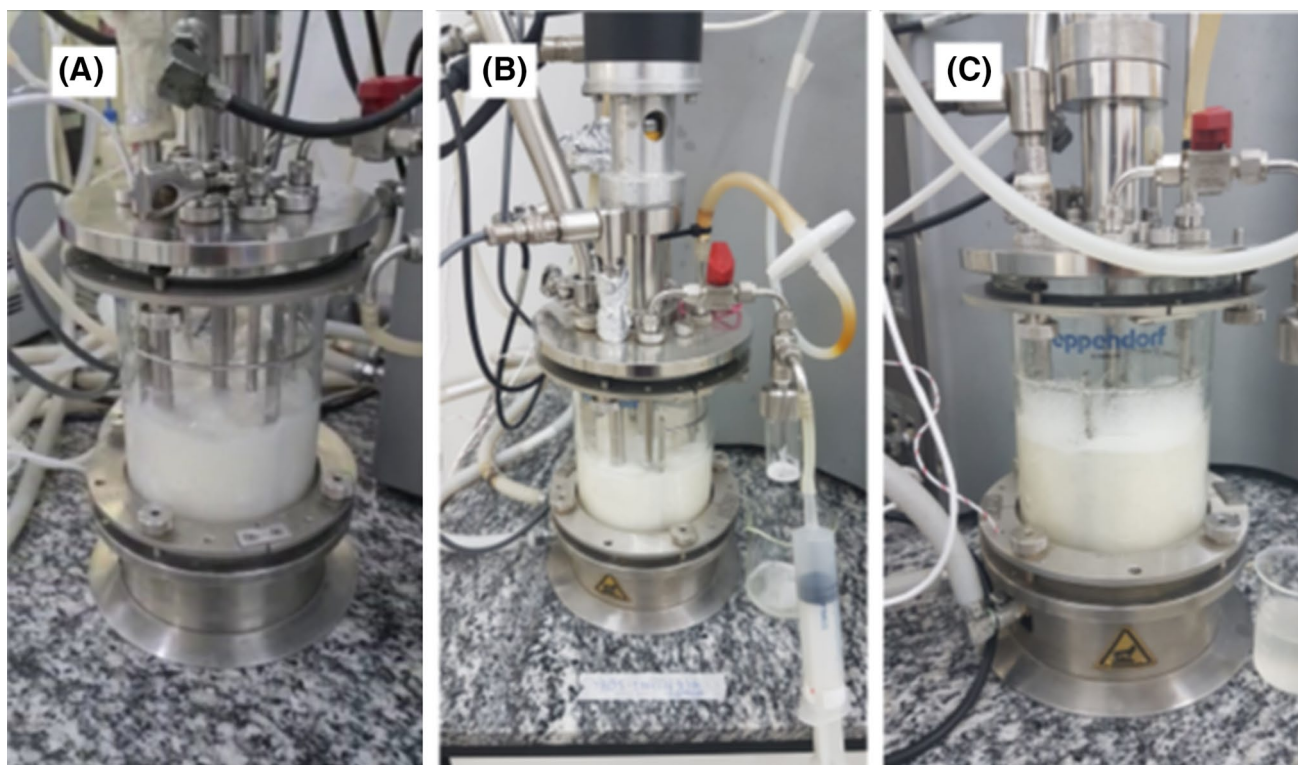
Only in the kinetics of the  $K_L a$  2.07  $\text{h}^{-1}$  condition did the DO value reach zero during the exponential phase of cell growth, remaining around 40% in the stationary phase of cell growth. For the conditions of  $K_L a$  4.24 and 11.56  $\text{h}^{-1}$ , the DO did not even reach zero in the exponential phase of cell growth, and remained around 64% in the stationary phase of growth for the condition of  $K_L a$  4.24  $\text{h}^{-1}$  and 75% in the  $K_L a$  condition 11.56  $\text{h}^{-1}$ . These results are strong indications that, in fact, the oxygen demand for conducting this type of bioprocess is relatively low, making an induced surface aeration strategy viable.

Although the calculated SOUR values were similar for  $K_L a$ 's of 2.07, 4.24 and 5.89  $\text{h}^{-1}$ , there was a significant increase for the condition of  $K_L a$  11.56  $\text{h}^{-1}$  (Table 3), which in fact was accompanied by a decrease in surfactin production. SOUR is routinely used to indicate metabolic shifts in cells [37]. These results may be related to the Leucine metabolism, the main amino acid present in the surfactin peptide moiety (4 of 7 residues). Coutte et. al (2015) [38] reported significant surfactin yield improvement by leucine supplementation. Leucine is known to be an amino acid synthesized via pyruvate,  $\alpha$ -acetolactate and  $\alpha$ -ketoisovalerate. When the oxygen consumption is high, the amount of pyruvate oxidized in the TCA cycle may increase, resulting in low leucine formation [39] and consequent low surfactin formation. On the other hand, if the oxygen supply is extremely low or non-existent, pyruvate will be directed into lactate fermentation [39], also resulting in low surfactin formation.

In many surfactin production studies in benchtop bioreactors available in the literature, the surfactin concentration is measured in the collected foam, which means a bioproduct concentration up to 10x [10]. Interestingly, studies that related  $K_L a$  to surfactin production described much higher  $K_L a$  values as optimal, around 45–48  $\text{h}^{-1}$ , using foam collection [8, 10].

Yeh et al. (2006) [8] reported optimal surfactin production at a  $K_L a$  value of 47.52  $\text{h}^{-1}$ , reaching surfactin concentrations of 6.45  $\text{g L}^{-1}$  in the foam. Nazareth et al. (2021) [10] reported a volumetric productivity of 9.98  $\text{mg L}^{-1} \text{h}^{-1}$  for an optimized  $K_L a$  of 45.41  $\text{h}^{-1}$ , reaching a surfactin concentration of 239.74  $\text{mg L}^{-1}$  in the main bioreactor and 2159.84  $\text{mg L}^{-1}$  of surfactin in the collected foam. Possibly, in aeration system using bubbling and foam collection, it is necessary to work with higher  $K_L a$  values to compensate for the loss of bacterial cells entrapped in the foam and carried out into the foam collector. In other words, for the cell density within the bulk liquid to remain at adequate levels for surfactin production, considering the biomass loss carried out in the collected foam, it is likely that higher  $K_L a$  values must be used, due to the oxygen consumption required for cell multiplication.

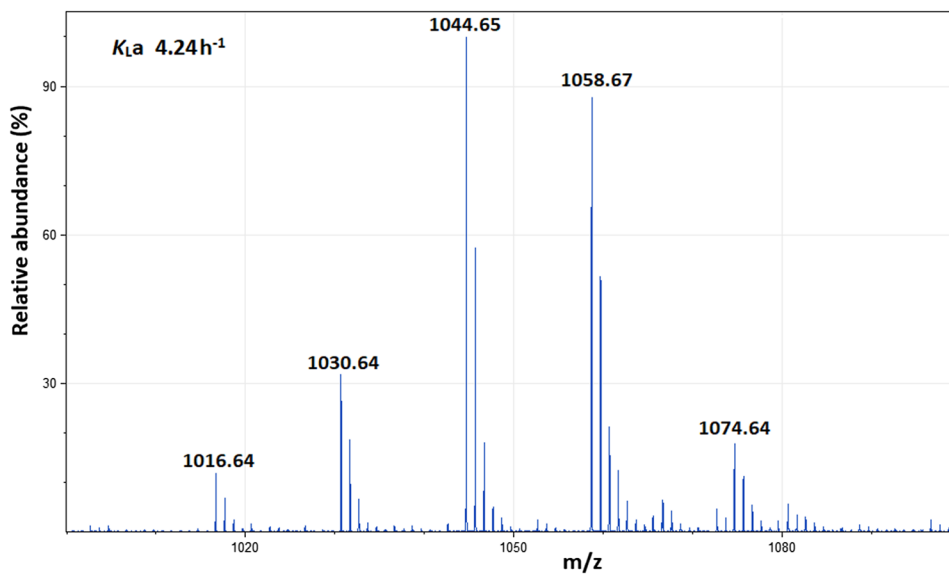
In this study, in none of the aeration conditions tested, excessive foam formation was observed, as depicted in Fig. 6. At the highest  $K_L a$  tested of 11.56  $\text{h}^{-1}$ , in fact, a slightly thicker layer of foam was observed (roughly 2–4 cm of foam thickness), probably caused by the higher stirring speed with an impeller close to the liquid surface (characteristic of the induced surface aeration strategy). However, even in this condition, the foam remained stable and far from reaching the top of the bioreactor, which would cause filters fouling and other operational difficulties. This result indicates that induced surface aeration is a promising technology for surfactin production, reducing the expenses and operational complexity with the foam collection step. In addition, because there was no irregular loss of culture volume or cells through the foam in the tests performed, it was possible, in an unprecedented way in the literature, to make a comparison of different  $K_L a$ 's in the bioprocess for surfactin production.



**Fig. 6** Bioreactor photos under different  $K_L a$  conditions tested, near the end of the bioprocess with 48 h. **A**  $2.07 \text{ h}^{-1}$ ; **B**  $4.24 \text{ h}^{-1}$ ; **C**  $11.56 \text{ h}^{-1}$ . There was no excessive foaming under any tested aera-

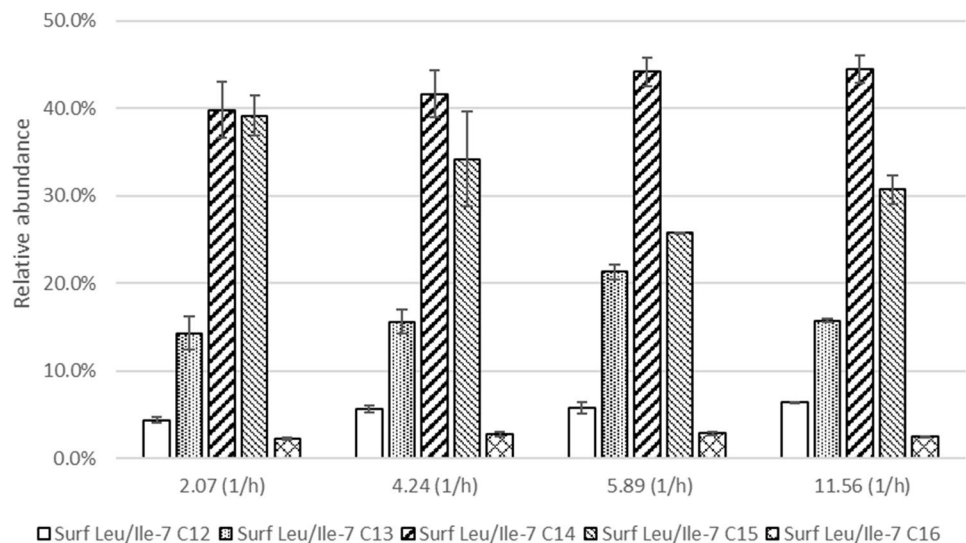
tion conditions. In the condition of  $K_L a 11.56 \text{ h}^{-1}$ , in fact, a slightly thicker layer of foam was observed, probably caused by the higher agitation speed with an impeller close to the surface

**Fig. 7** ESI-Q-TOF-MS mass spectra of  $K_L a 4.04 \text{ h}^{-1}$  sample. Main surfactin homologues molecular ions are marked at  $m/z = 1044.65$  (surfactin Leu/Ile-7  $C_{14} [M + Na]^+$ ),  $m/z = 1058.67$  (surfactin Leu/Ile-7  $C_{15} [M + Na]^+$ ),  $m/z = 1030.64$  (surfactin Leu/Ile  $C_{13} [M + Na]^+$ ),  $m/z = 1016.64$  (surfactin Leu/Ile-7  $C_{12} [M + Na]^+$ ),  $m/z = 1074.64$  (surfactin Leu/Ile-7  $C_{15} [M + K]^+$ )





**Fig. 8** Relative abundance of each surfactin homologue Surfactin Leu/Ile-7 C<sub>12</sub>–C<sub>16</sub> for 48 h samples of each tested  $K_L a$  value



### $K_L a$ effect on surfactin homologues relative abundance

Figure 7 shows the final supernatant sample mass spectrum of  $K_L a = 4.24 \text{ h}^{-1}$  condition, more productive in terms of surfactin concentration. The highest peak in spectra at  $m/z = 1044.65$  refers to molecular ion surfactin Leu/Ile-7 C<sub>14</sub> [M+Na]<sup>+</sup>. Naturally, mass spectrometry results do not provide the differentiation of compounds having the same mass (such as Leu and Ile, for example), nor the type of fatty acid chain (linear, iso or anteiso), but provides the global mass and the peptide moiety primary sequence. Figure 8 depicts the relative abundance of the identified surfactin homologues for each aeration condition. All supernatant samples showed that the surfactin homologue present in the highest relative abundance was Surfactin Leu/Ile-7 C<sub>14</sub>, followed by Surfactin Leu/Ile-7 C<sub>15</sub>, Surfactin Leu/Ile-7 C<sub>13</sub>, Surfactin Leu/Ile-7 C<sub>12</sub> and by Surfactin Leu/Ile-7 C<sub>16</sub>. This result is in agreement with what Kaisermann (2017) [38] reported for the analysis of Sigma's surfactin standard by mass spectrometry, where it was observed that the main homologue present is indeed Surfactin Leu/Ile-7 C<sub>14</sub>. In addition, Iturin Asn-1 C<sub>14</sub>, Iturin Asn-1 C<sub>15</sub>, Bacillomycin D C<sub>14</sub> and Bacillomycin D C<sub>15</sub> peaks were identified in all samples, but in very low relative amounts, approximately 1–2% of the relative intensity of Surfactin Leu/Ile-7 C<sub>14</sub> (major homologue). Species of fengycin, kurstakin or polymyxin families were not detected in any of the samples.

The increase in  $K_L a$  was accompanied by a small tendency to decrease the relative abundance of the surfactin leu/Ile-7 C<sub>15</sub> homologue, from  $K_L a$  2.07–5.89  $\text{h}^{-1}$  (Fig. 8). Likewise, there was a slight tendency to increase in the relative abundance of the Leu/Ile-7 C<sub>14</sub> surfactin homologue with increasing  $K_L a$ . Intracellular fatty acids metabolism is divided into straight chain fatty acid synthesis and branched-chain fatty

acid synthesis. Acetyl-CoA is the precursor of straight-chain fatty acids, while the precursors of branched-chain fatty acids are catabolic products of Val, Leu, and Ile [40]. More specifically, branched-chain fatty acid synthesis from Leu and Ile generates uneven chains of fatty acids [40]. In fact, Liu et al. (2012) [41] experimentally demonstrated that, in a culture for surfactin production using *B. subtilis* TD7, Leu and Ile supplementation significantly enhanced the proportion of surfactins with odd  $\beta$ -hydroxy fatty acid. It is to be expected, therefore, that the generation of C<sub>15</sub> fatty acids is to some extent a competitive metabolic pathway for the total surfactin production, which has Leu in large amounts in its peptide portion. This would mean a decreasing trend, therefore, for the relative abundance of C<sub>15</sub> surfactin homologues under conditions, where the total surfactin concentration was higher. Observing the results of the present study, this trend was observed to a certain extent, comparing the results of  $K_L a$  5.89–11.56  $\text{h}^{-1}$ , where in  $K_L a$  11.56  $\text{h}^{-1}$  a lower total concentration of surfactin was obtained and an increase in the C<sub>15</sub> homologue was observed (Fig. 8). However, this same behavior was not observed when comparing the results of  $K_L a$  4.24–5.89  $\text{h}^{-1}$ , since in  $K_L a$  4.24  $\text{h}^{-1}$  a higher total concentration of surfactin and a higher concentration of the C<sub>15</sub> homologue were observed at the same time.

In addition, in all  $K_L a$  conditions tested, surfactin homologues C<sub>14</sub> and C<sub>15</sub> had higher relative abundance. Interestingly, these are the surfactin homologues reported in the literature with the greatest disease-suppression capacity [4], antiviral activity [42], insecticidal activity [43] and as activators of several plant-defense mechanisms [44]. This relationship is a strong indication that the surfactin production profile of the strain tested in this study *Bacillus velezensis* H2O-1 has great application potential in plant disease biocontrol.

## Conclusions

The studies in this work indicate that the oxygen consumption for surfactin production is relatively low (SOUR values of approx.  $7.3 \text{ mgO}_2 \text{ gX}^{-1} \text{ h}^{-1}$ ), allowing the use of induced surface aeration and evidencing that complex operational strategies such as foam collection or culture medium composition strategies to induce nitrate respiration are not necessary. It must be considered that the induced surface aeration system described in this study is considerably simpler operationally and scalable when compared to foam collection strategies. Due to the tendency of cells to be entrained in the generated foam, problems such as loss of cells from the main bioreactor and clogging of the foam collection system are widely reported in the literature of foam collection strategies [8, 12]. The operation of a bioprocess where the bioreactor volume is erratically lost and that requires the collected cells separation and recycling will be more difficult to scale-up and operate, in addition to entailing higher fixed and operational costs.

Concerning the different surfactin homologues produced at the tested  $K_L a$  levels,  $C_{14}$  and  $C_{15}$  homologues were the most abundant in all conditions tested. Surfactin  $C_{14}$  and  $C_{15}$  homologues are often described as more efficient for many applications [4, 39, 40]. Nevertheless, the  $K_L a$  parameter seems to have had minimal influence on affecting the relative abundances of surfactin homologues produced.

In addition, this study highlights the indication that high  $K_L a$  values can be deleterious to the total surfactin production (above  $11.56 \text{ h}^{-1}$ ), which makes sense when considering the biosynthesis of the main amino acid present in the peptide portion of the surfactin molecule, leucine. As pyruvate is one of its main precursors, it can be expected that high oxygen availability will increase the amount of pyruvate being oxidized in the TCA cycle, resulting in insufficient leucine formation and consequent low surfactin production. On the other hand, conditions close to anaerobiosis ( $K_L a$  values close to zero), in fact, showed almost zero production of surfactin and little cell growth.

Finally, because there was no irregular loss of culture volume or cells through the foam in the tests performed with this study, it was possible to carry out, in an unprecedented way in the literature, a benchtop bioreactor system study comparing different  $K_L a$ 's in the bioprocess for surfactin production, obtaining surfactin volumetric productivity values close to  $11.9 \text{ mg L}^{-1} \text{ h}^{-1}$  in  $K_L a$  of  $4.24 \text{ h}^{-1}$ .

**Data Availability** The data that support the findings of this study are available from the corresponding author, upon reasonable request.

## Declarations

**Conflict of interest** The authors declare that they have no conflicts of interest in the publication.

**Ethics approval** Not applicable.

**Consent to participate** All authors consent to participate.

**Consent for publication** All authors consent for publication.

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