

# N-acylation of L-amino acids in aqueous media: Evaluation of the catalytic performances of Streptomyces ambofaciens aminoacylases

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## N-acylation of L-amino acids in aqueous media: evaluation of the catalytic

# performances of Streptomyces ambofaciens aminoacylases

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# **Abstract:**

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N-acylated amino acids are widely used as surfactants and/or actives in cosmetics and household formulations. Their industrial production is based on the use of the Schotten-Baumann chemical and unselective reaction. Faced to the growing demand for greener production processes, selective enzymatic synthesis in more environment-friendly conditions starts to be considered as a potential alternative. This study concerns the use of the aminoacylases from Streptomyces ambofaciens to selectively catalyse aminoacid acylation reaction by fatty acids in aqueous medium. The results demonstrated that, when using undecylenoic acid as acyl donor, these aminoacylases properly catalyse the acylation of 14 of the 20 proteogenic L-amino acids tested on their α amino group with a great variability depending on the nature of the amino acid (polar or not, positively/negatively charged, aromatic or not...). More precisely, the following 9 amino acids were shown to be preferentially acylated by S. ambofaciens aminoacylases as follows: lysine > arginine > leucine > methionine > phenylalanine > valine > cysteine > isoleucine > threonine. Different fatty acids were used as acyl donors and, in most cases, the fatty acid length influenced the conversion yield. The kinetic study of  $\alpha$ -lauroy-lysine synthesis showed a positive influence of lysine concentration with Vmax and Km of 3.7 mM/h and 76 mM, respectively. Besides, the lauric acid had an inhibitory effect on the reaction with Ki of 70 mM. The addition of cobalt to the reaction medium led to a more than six-fold increase of the reaction rate. These results, achieved with the aminoacylases from S. ambofaciens represent an improved enzymebased N-acylated amino acids production in order to provide an alternative way to the Schotten-Baumann chemical reaction currently used in the industry.

### 1. Introduction:

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Amino acid surfactants (AAS) are one of the most important class of natural surface-active molecules [1]. They come from the condensation of an amino acid as acyl acceptor, and one or more non polar molecules such as long chain fatty acids as acyl donors to constitute amphiphilic molecules with high surface activity. This characteristic, in addition to their ecocompatibility, makes them molecules of choice for food, pharmaceutic and cosmetic industries [2]. Given the structural diversity of the 20 natural amino acids and the great variety of fatty acids that can be involved in acylation reaction, a wild range of surfactants with diverse properties and activities can be found. Indeed, depending on the amino acid radical and the acylated group (O-acylation or N-acylation), AAS can be classified on the basis of head group type with different commercial applications. For instance, cationic AAS like cocoyl arginine ethyl ester were described as disinfectants, antimicrobial agents and antistatic agents. Anionic AAS like N-cocoyl-L-glutamate and potassium N-cocoyl glycinate are used in cosmetics for several applications and non-ionic AAS for their good detergency and foaming properties (phenylalanine and leucine derivatives). Zwitterionic or amphoteric AAS like ε-lauroyl-lysine, are widely used in creams and hair conditioners because of their low solubility in aqueous media [3,4]. The N-acylation position in the case of lysine as an acyl acceptor plays an important role in the property and activity of the synthetized AAS. Pérez et al. (2009) studied the influence of the N-acylated group of the lysine methyl and ethyl-ester on the antimicrobial activity [5]. They demonstrated that the N-acylation on εposition decreased considerably the antimicrobial activity as compared to what was observed when acylation occurred in α-position. This was due to the important difference in pKa between amino groups. These pKa are of 8.8 and 6.2 for the amino group in position  $\varepsilon$  and  $\alpha$ , respectively. They also reported that the hemolytic activity of N-α-acyl lysine based surfactant was lower than the Ne-acyl lysine one, but both had much lower activity as

compared classical cationic surfactants to the quaternary ammonium like alkyltrimethylammonium salts [5,6]. These results showed the practical interest of N-\alphaacylation of amino acids and consequently, of a regioselective acylation when using an amino acid such as lysine, with two different amino groups. The most common way to produce those AAS is via the Schotten-Baumann chemical reaction where the amino acid reacts with a fatty acid chloride under highly alkaline condition by using NaOH or Na<sub>2</sub>CO<sub>3</sub> in TFA/water or by using solvents like trimethylamine or Nmethylmorpholine. In addition, the N-acylation of amino acids exhibiting functional groups at their side chain requires a preliminary step of protection in order to avoid side reactions [7]. Even if this chemical way presents high synthesis yields, the production at industrial scale leads to large volumes of effluents requiring a very expensive reprocessing step. Nowadays, more and more industries try to develop sustainable technologies by switching from waste remediation and pollution control by end-of-pipe solution to waste prevention, reducing the environmental issues and the reprocessing cost [8]. One of the most interesting way to meet these aims is the use of biologically mediated reactions, more generally called bioconversions. Enzymes are catalysts of choice for such reactions to avoid the formation of by-products and salts. They can make the catalysis in mild conditions (pH and temperature), are relatively substrate specific and often show chemo-, regio- and even enantioselectivity. To date, while many articles and books have been dealing with O-acylation (esterification or transesterification), mainly using lipases, only few studies have been focused on the Nacylation reaction. Few authors have evaluated the performances of enzymes for the acylation of amino acids using even free fatty acids or triglycerides like soybean and palm oil [9–15]. Lipases, especially Candida antarctica lipase B (CALB), Rhizomucor miehei lipase, and proteases are the most used biocatalysts. They are hydrolases that can perform synthesis

reaction in non-aqueous media such as organic solvents or in solvent free condition in order to

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shift the thermodynamic equilibrium towards the synthesis. However, amino acids are poorly soluble in organic solvents [14,16,17]. Attempting to solve this problem, several authors used amino acid derivatives using Cbz (carboxybenzyle) or tert-butyl ester in order to improve their solubility and to protect the carboxylic part, allowing an increase of the conversion rate [17–19]. But this method needs an additional chemical step of deprotection to give the final product [19]. Other authors used ionic liquids in order to improve the solubility of the acyl acceptor. Husson et al., 2009 were able to improve the reaction performance of CALB for the of synthesis N-ε-oleyl lysyl-serine by using 1-butyl-3-methylimidazolium hexafluorophosphate ([Bmim]+ [PF6]-) instead of 2-methyl-2-butanol as reaction medium (initial rate was found to increase from 1 to 32 mM/h) [20]. However, the large scale use of neoteric media such as ionic liquids tends to be limited by several key drawbacks: their high viscosity, the complexity of their production but also their price as well as a lack of information on their toxicity [21]. As a catalytic alternative to lipases in anhydrous or almost anhydrous solvents, the use of aminoacylases obviously presents one key advantage: they can perform N-acylation in water while being present in a wide variety of biological matrices [22,23]. Only few studies investigated N-acylation of amino acids in aqueous media. Acylases produced by microorganisms were among the most used enzymes at industrial scale for the production of biomolecules. Microbial enzymes performing the amino acids amidation in aqueous or organic-aqueous medium were described by Koreishi et al, [10,11,24,25]. Four acylases from Streptomyces mobaraensis were identified and three of them were characterized for the N-amino acid acylation reaction. The first one is a monomeric aminoacylase metalloenzyme of 100 kDa. This enzyme was able to catalyse the N-acylation of almost all the amino acids with much less specificity for glycine, cysteine, aspartic acid and histidine and no reaction with tyrosine and proline using 10 mM of ferulic acid as acyl donor in 70% glycerol reaction media. The production of N-α-feruloyl-L-lysine was improved by using 100 mM tris

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HCl buffer (pH 7.5) without glycerol at optimum conditions [24]. The second enzyme was the ε-Lysine acylase (Sm-ELA) which was identified as a monomeric metallo-enzyme of around 60 kDa. This enzyme showed a regioselectivity towards the N-acylation of the lysine side chain (\varepsilon-position). It efficiently catalysed the N-\varepsilon-acylation of lysine with fatty and aromatic acyl groups in aqueous buffer [9,10]. The third acylase (Sm-PVA) was a dimeric enzyme of 61 and 19 kDa identified as belonging to the Penicillin V acylase family. This latter was able to catalyse the N-acylation reaction of several amino acids with lauric acid showing a better activity with lysine in \(\epsilon\)-position, arginine and cysteine [11,26]. The last acylase (Sm-AA) was a monomeric protein of 55 kDa belonging to the peptidase M20 family. This enzyme showed efficient hydrolytic activity towards amino acids acylated with short or long chain fatty acids [25]. Recently, Dettori et al. (2017) described the ability of a crude extract from S. ambofaciens ATCC23877 to perform the N- acylation of lysine and peptides with oleic acid on their α position unlike the CALB. By blast analysis, and using the genes encoding Sm-AA, Sm-ELA and Sm-PVA from S. mobaraensis as baits, 3 genes encoding putative acylases and sharing 86, 80 and 69% identity, respectively, were identified in the genome of S. ambofaciens. Another gene, presenting 55% identity with the sequence of Sm-AA was also identified. By gene deletions, and just keeping Sam-AA or Sam-ELA genes, it was shown that Sam-ELA enzyme showed hydrolytic specificity for N-\varepsilon-acetyl-L-lysine and that Sam-AA enzyme had specificity towards N-α-acetyl-L-lysine, while the crude extract was unspecific. By generating a mutant without Sam-ELA and Sam-AA genes, it was also demonstrated that 85% of the N-acylation ability from the crude extract was due to Sam AA and Sam-ELA [12]. The activity of these acylases from S. ambofaciens ATCC23877 was also evaluated for the synthesis of lauroyl-lysine after enzyme immobilisation [27]. However, to date, no data has been reported regarding the substrate specificity (acyl donor and acyl acceptor) of the acylases from S. ambofaciens ATCC23877.

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The aim of this study was therefore to determine the substrate specificity of the aminoacylases from *S. ambofaciens* for the acylation reaction of the twenty proteinogenic amino acids with middle and long chain fatty acids. The influence of the operational conditions on reaction performances was also investigated using lauroyl lysine synthesis as a reference reaction.

## 2. Materials and methods:

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## 2.1. Aminoacylases of S. ambofaciens production

In order to have enough S. ambofaciens ATCC23877 spores before starting the culture steps, a sporulation step was performed. The strains spores stored as suspensions at -80°C were streaked on Soy Flour Mannitol agar (SFM) filled. This later contain 20 g/L Mannitol, 20 g/L precooked Soy flour (Celnat) and 20 g/L Agar. After minimum 7 days of culture at 28°C, the spores of S. ambofaciens were collected by adding 2 mL of 20% (v/v) glycerol and by scratching the mycelium with a sterile handle before freezing the spore suspension at -20°C. the S. In same the spores of ambofaciens way ΔSAM23877\_1485::scarΔSAM23877\_0977::scar strain was prepared. This strain was obtained by deletion of  $\Delta SAM23877\_1485$  and  $\Delta SAM23877\_0977$  genes coding for Sam-ELA and Sam-PVA respectively as described in [12]. The culture was done in 3 steps: germination, preculture and the culture. The first step was performed in the ICS medium (Insoluble complex medium). This medium contained sucrose 15 g/L, corn steep 5 g/L sterilized beforehand, (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> 10 g/L, KH<sub>2</sub>PO<sub>4</sub> 1 g/L, NaCl 3 g/L, MgSO<sub>4</sub>·7H<sub>2</sub>O 0.2 g/L and CaCO<sub>3</sub> 0.2 g/L. The medium was then sterilized at 121°C for 20 min. In a 250 mL baffled Erlenmeyer flask, 40 mL of this medium was inoculated with 100 µL of spore suspension. After the germination, 20 mL of ICS medium were transferred in 200 mL of pre-culture medium contained in 2 L Erlenmeyer flask for the pre-culture step. The preculture medium composition was: Glucose 10 g/L, white dextrin 10 g/L, yeast extract 5 g/L, casein

hydrolysate 5 g/L, and CaCO<sub>2</sub> 1 g/L. The pH of the preculture medium was set at 6.5 before sterilization at 121°C for 20 min. The two first steps were performed for 48 h at 28°C and 250 rpm orbital agitation using an INFORS AG AK 82 agitator. 30 ml of the preculture were then added in 950 ml of culture medium in 5 L Erlenmeyer flask for incubation. The culture was maintained for 7 days at 28°C and 200 rpm orbital agitation using an INFORS HT Multitron agitator. The composition of the culture medium was meat extract 40 g/L, soluble starch 40 g/L, casein hydrolysate 20 g/L, MgSO<sub>4</sub> 20 g/L, and K<sub>2</sub>HPO<sub>4</sub> 2 g/L. The pH of the culture medium was set to 7 before sterilisation at 121°C for 20 min. All the components of the culture media were purchased from Sigma-Aldrich, except sucrose and yeast extract, from Fluka, (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> (99%), NaCl, CaCO<sub>3</sub> and acetonitrile, from Carlo Erba. At the end of the main culture, the bacterial cells were disrupted by first mixing the whole suspension for around 5 min minimum with an ULTRA-TURRAX T25 (Janke & Kunkel IKA Labortechnik) followed by 2 passages in a cell disrupter (Constant Cell disruption system, Cell-D) at 2 kBars. The cellular debris were removed by centrifugation at 8600×g at 4°C for 20 min. Thereafter, the proteins including acylases were precipitated from the cell free extract by adding ammonium sulfate to the supernatant to 60% of saturation and agitation was maintained for 1 h at room temperature. The precipitate was recovered by centrifugation at 8600×g at 4°C for 20 min and was dissolved in 25 mM Tris-HCl buffer, NaCl 50mM, at pH 8. The solution was then diafiltrated and concentrated using a Cogent uScale tangential flow filtration system equipped with a 10kDa ultrafitration membrane (Merck Millipore). Five diavolumes were necessary for removing the salt. Salt removal was checked by following the conductivity evolution during the filtrations steps. The apparent specific activity of the S. ambofaciens crude extract was evaluated using lauroyl-lysine synthesis reaction. For this reaction lauric acid and L-lysine were used as substrates at equimolar concentration (0.1 M) in 25 mM Tris-HCl buffer, NaCl 50mM, at pH 8, 45°C and at 500 rpm.

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#### 2.2. Enzyme assays

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Substrate specificity of the acylases from S. ambofaciens toward acyl acceptor was evaluated by performing the condensation reaction of the 20 L- amino acids as acyl acceptors with 10undecenoic acid as acyl donor. Reactions with taurine, 6-aminohexanoic acid, 6-amino-1hexanol, ethanolamine, hydroxylamine, 4-amino-benzoic acid and carnosine as acyl acceptor and lauric acid as acyl donor were also performed. All acyl donors were from Sigma Aldrich except 4-amino-benzoic acid, which was from Merck. Substrate specificity of the acylases from S. ambofaciens towards acyl donor was performed using lysine, arginine, methionine and phenylalanine with octanoic, 10-undecenoic, lauric and oleic acid as acyl donors. The influence of the acyl donor structure on the regioselectivity was evaluated using lysine as acyl acceptor and acetic, butyric, isobutyric, valeric, octanoic, phenylpropionic, benzoic and azelaic acid as acyl donors. All acyl donors were from sigma Aldrich except butyric acid and azelaic acid which were from Rectapur and Fluka Chemika, respectively. Substrate specificity evaluation was conducted in test tubes using 100 mM of each substrate in 2 mL of 25 mM Tris-HCl, 50 mM NaCl buffer, at 45°C, pH 8 and 250 rpm magnetic agitation. All reaction media were sonicated using an ultrasonic processor in order to improve reaction medium homogeneity. The influence of substrate concentration was studied using lauroyl-lysine synthesis reaction. Concentrations from 25 to 250 mM of lysine or from 10 to 200 mM of lauric acid were used, the other substrate being fixed to 100 mM. The inhibition by the product of the reaction was investigated by adding different concentrations of a chemically produced α-lauroyl-lysine (from GeneCust, France, with 97.21% purety). The reaction was conducted in 1 mL reaction media containing 100 mM of both substrates.

The effect of metal ions was investigated using CoCl<sub>2</sub>, ZnSO<sub>4</sub> (from Sigma Aldrich) and CuCl<sub>2</sub> (from Fluka Chemika). The reactions were performed in 3 mL of 25 mM Tris-HCl, 50 mM NaCl buffer, at 45°C, pH 8 and 500 rpm magnetic agitation.

In all cases, the reaction is started after one night agitation, in order to check if the reaction occurs without enzyme, by adding 1 g/L of the produced enzyme mixture from *S. ambofaciens*. At appropriate times, 50 or 100 µL aliquots of the reaction mixture were withdrawn and diluted with 80% methanol for HPLC and LC-MS-MS analyses. For the regioselectivity evaluation, a TLC analysis was also performed. Finally, all reactions were stopped by freezing the mixture at -18°C. Each reaction was performed at least twice.

#### 2.3. Analytical methods

### 2.3.1. Qualitative analyses by TLC

Qualitative analyses of the reaction media by TLC were performed on Kieselgel G 60 Plates  $20~\text{cm} \times 20~\text{cm}$  (Merck, Darmstadt, Germany). The mobile phase was: butanol (60%)/acetic acid (20%)/water (20%) that permitted a slight but visible separation between the  $\alpha$  and the  $\epsilon$  monoacylated products. Knowing that lysine contains two amino groups allowed the detection by spraying a commercial solution of ninhydrin reagent (Sigma-Aldrich).

### 2.3.2. Quantitative analyses by HPLC

The quantification of reaction conversion yield was monitored using HPLC (LC 10 AD-VP, Shimadzu, France) equipped with a UV detector at 214 nm followed by a light-scattering low temperature evaporative detector (Shimadzu, France). Two columns were used: the first one was a C18 amide 125 x 2.1 mm (Altima®, Altech, France) maintained at 25°C. The mobile phase (0.2 mL/min flow rate) consisted of solvent A: methanol/water/TFA (60/40/0.1 v/v/v) and solvent B: methanol/TFA (100/0.07). A linear elution gradient was applied to reach 95% methanol/TFA after 6 min. This methanol concentration was maintained for 13 min and then

decreased in 1 min to reach the initial methanol/water ratio until the end of the run (35 min). However, the use of this column did not allow the separation of the product and the fatty acids when phenylalanine, leucine and isoleucine were used as acyl acceptors. In that cases a second column, Force Biphenyl 3 µm column 150 x 2.1 mm (Restek, USA), maintained at 35°C, was used. The same mobile phase composition and gradient as before were used. Calibrations were performed using standard curve fatty acids and purified product in the case of α-lauroyl-lysine. The substrate conversion yield was determined using the following equation: conversion yield (%) =  $(1 - ([fatty acid]_{mol/L} consumed / [fatty acid]_{mol/L} initial) \times$ 100. Initial reaction rates were calculated using a third order polynomial model applied to the derivate of the first experimental data points exhibited in the figures. The maximum velocity (Vmax) and the Michaelis-Menten constant (Km) for each substrate were determined using a standard Mickaelis-Menten model based on the equation of the graphic representing  $\frac{1}{V}$  =  $f\left(\frac{1}{c}\right) = \frac{1}{c} * \frac{Km}{Vmax} + \frac{1}{Vmax}$  where V is the initial rate and S is the substrate concentration. In the case of enzymatic substrate inhibition, inhibition constants (Ki) and maximum velocity (V<sub>max</sub>) can be evaluated using  $V = \frac{V max \times S}{K m + S + \frac{S^2}{K}}$  and by neglecting Michaelis Menten constant (Km is very low as compared to S). Vmax and Ki were then determined according to the equation of the graphic representing  $\frac{1}{V} = f(S) = \frac{S}{Ki * V max} + \frac{1}{V max}$ .

#### 2.3.3. Mass spectrometry analysis

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Qualitative and semi quantitative analysis of the different acyl-amino acids were carried out on an HPLC–MS-MS system (ThermoFisher Scientific, San Jose, CA, USA) consisting in a binary pump connected to a photodiode array detector (PDA) and a Linear Trap Quadrupole mass spectrometer (LTQ). The system was equipped with an atmospheric pressure ionization interface operating in positive electrospray mode (ESI+). Chromatographic separation was

performed by injecting ten microliters of the prepared samples on a C18 Altima column (Grace/Alltech, Darmstadt, Germany, 150 mm × 2.1 mm, 5 μm porosity) equipped with a C18 pre-column (Grace/Alltech Darmstadt, Germany, 7.5 mm × 2.1 mm, 5 μm porosity) at 25°C. The flow rate was set at 200 µL.min<sup>-1</sup> and mobile phases consisted of solvent A: methanol/water/TFA (80:20:0.1, v/v/v) when lauric or oleic acid was used and methanol/water (50:50, v/v) when shorter fatty acid was used and solvent B methanol/TFA (100:0.1, v/v). Acylated products were eluted using a linear gradient from 0 to 100% of B for 5 min and then an isocratic step at 100% of B for 10 min. Allure Biphenyl column 3 µm column 150-2.1 mm (Restek, USA), maintained at 35°C, was used to perform the separation of the product and the fatty acid when using phenylalanime, leucine and isoleucine following the same gradient as with the C18 column. Mass spectrometric conditions were as follows: spray voltage was set at +4.5kV; source gases were set for sheath gas, auxiliary gas and sweep gas at 30, 10 and 10, respectively (in arbitrary units min<sup>-1</sup>); capillary temperature was set at 250°C; capillary voltage was set at 48V; tube lens, split lens and front lens voltages were set at 120 V, -34 V and -4.25 V, respectively. Ion optics parameters were optimized in ESI+ mode by automatic tuning using a standard solution of oleylysine-serine at 0.1 g L<sup>-1</sup> infused in mobile phase (A/B: 50/50) at a flow rate of 5 µL.min<sup>-1</sup>. Full scan MS spectra were performed from 100 to 1000 m/z and additional MS<sup>2</sup> scans were realized to get structural information based on daughter ions elucidation. Raw data were processed using Xcalibur software program (version 2.1, http://www.thermoscientific.com). MS<sup>2</sup> fragmentation data were interpreted in the way to evaluate chimioselectivity and regioselectivity of the aminoacylases from S. ambofaciens.

## 3. Results and Discussion

#### 3.1. Enzyme production

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All the proteins used in this study were obtained from the same seven days culture of S. ambofaciens ATCC23877 after two days of germination and pre-culture steps. After this incubation time, the cells were completely disrupted and the proteins were precipitated using ammonium sulphate. The proteins were then re-suspended and ultrafiltration using a 10 kDa membrane was performed. The filtration step allowed the elimination of almost all ammonium sulphate (decrease of the conductivity from 30 to 6.5 mS / m) and 135 mL of crude extract containing  $21 \pm 0.7$  g/L protein were recovered from the 5 L culture medium. The apparent specific activity of the extracted proteins was evaluated using lauroyl-lysine synthesis reaction. Dettori et al. (2018) used this reaction for the evaluation of the activity of immobilized acylases from S. ambofaciens on mesoporous material [27]. However, in this study equimolar substrate concentrations of 0.1 M were used (the influence of the concentration of each substrate will be treated later). The specific activity of the acylases from S. ambofaciens was  $33.4 \pm 5.0 \,\mu\text{M}$  of  $\alpha$ -lauroyl-lysine / min / mg of protein. The influence of the pH when performing this reaction was firstly investigated in order to set a fixed pH range of the reaction media for the further reactions (data not shown). The results showed that the enzyme was much less active at pH lower than 6.5 and higher than 8.5 with highest activity around pH 8. So all the further reactions were conducted at pH 8.0.

#### 3.2. Substrate specificity

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#### 3.2.1. Acyl acceptor specificity

As described in a previous study [12], acylases from *S. ambofaciens* were able to perform the N-acylation of lysine and peptides with oleic acid on the amine function located on  $\alpha$ -position. The specificity towards other amino acids have never been reported. 10-undecenoic acid was used as acyl donor in this study. It is a mono-unsaturated fatty acid having antifungal, antiviral and insect-repelling activity which makes it useful to treat superficial infection [28,29]. It was also described as having neuroprotective and radical-scavenging

activities [30]. The enzymatic acylation of amino acids with this fatty acid has never been reported before. All condensation reactions were carried out with 1 g/L of acylases mixture at 45°C ± 3°C during 3 days. As shown in fig. 1, the aminoacylases from S. ambofaciens allowed the acylation of amino acids with conversion yields from 5 to 25%, except for tyrosine, tryptophane, histidine, proline, glutamic and aspartic acids for which the quantity produced was below the detection limit. The results obtained with tyrosine and tryptophane can be attributed to their very low solubility at pH 8 which is the optimal pH of these enzymes. In the case of non-polar aliphatic amino acids and also in the case of polar and noncharged amino acids with hydroxyl group, a relation between the side chain length and the conversion percentage was observed: the longer the chain length the higher the conversion yield. The best results were obtained with lysine, arginine, leucine and methionine with 24  $\pm$ 1.5,  $22 \pm 1.2$ ,  $23 \pm 0.14$  and  $20 \pm 0.27\%$  conversion yields, respectively, after 3 days. According to these results, the specificity of the aminoacylases from S. ambofaciens was very different from that described by Koreishi et al. for the aminoacylase (of 100 kDa molecular weight) and the penicillin V acylase (Sm-PVA) from S. mobarensis [11,24]. Indeed, these authors used ferulic acid (10 mM) as acyl-donor for the evaluation of the 100 kDa aminoacylase of S. mobarensis specificity towards amino acids in tris-HCl buffer pH 7.2, containing 70% glycerol, for 6 days, at 37°C. The best conversion yield was around 12% with phenylalanine and between 0 and 8% with the other amino acids. They were able to improve the conversion rate of lysine by using only tris HCl buffer (pH 7.5) [24]. The evaluation of Sm-PVA specificity was done using lauric acid as acyl-donor (7.5 mM) in tris-HCl buffer containing 78% glycerol, for 2 days, at 37°C. The best results were observed with lysine that was acylated on its ε amino group, arginine and cysteine with 44, 32 and 32% conversion yields, respectively. 0 to 10% conversion yields were obtained with the other amino acids tested [11]. Recently, Takakura et al. (2019) identified a new aminoacylase from

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Burkholderia sp. specific for the acylation of amino acids with lauric acid. The evaluation of the amino acids specificity was performed after 60 h at 25°C in 100 mM sodium borate buffer, pH 9.0, containing 100 mM sodium laurate and 200 mM amino acid. The results showed that the enzyme was able to catalyse the acylation of most of the amino acids, especially when arginine, phenylalanine, lysine or valine (89, 51, 28 and 23 % yield respectively) were used. Like with the aminoacylases from *S. ambofaciens* no product was observed when using aspartic and glutamic acid as substrates [31].

## Fig.1:

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The acylation reaction media were analysed by LC-MS-MS after 3 days of incubation in order to check the presence of the product and to evaluate the chemo and the regioselectivity of the aminoacylases from S. ambofaciens. The acylation of amino acids by 10-undecenoic acid led to the formation of mono-acylated products in almost all cases except with glutamic acid and proline where no product was detected. Probably, N-undecenoyl-proline was not synthesized because of the cyclic structure of the proline with no primary amino function available. The fragmentation of the parent ion [M + H]<sup>+</sup> of each product generated daughter ions corresponding to  $[M - COOH + H]^+$  or  $[M - COOH - H_2O + H]^+$  or  $[M - COOH - 2 H_2O +$ H]<sup>+</sup> which confirmed that the acylation took place on the amino group of the amino acids. In the case of some of the amino acids carrying a reactive side chain, daughter ions that came from the radical part of the acyl-acceptor were generated. Indeed, the fragmentation profiles of acylated serine and threonine, which present a potentially reactive hydroxyl group on their radical part, were very similar. The fragmentation of the parent ion generated daughter ions coming from the carboxyl group and hydroxyl group with no fragment coming from the amine group (Fig. 2). These fragments showed that the acylation reaction occurred only on amino acid amino groups. Cysteine also generated a daughter ion that corresponded to [M -CH<sub>2</sub> - SH]<sup>+</sup> coming from the –SH group, thus confirming the acylation of the amino acid on

its amino group (Fig. 2). However, the LC-MS-MS analysis showed the presence of a mass (241 m/z) corresponding to the cystine that came from the condensation of two cysteine molecules with disulphide bond bridge formation which has a very low solubility in the buffer [32]. In terms of regioselectivity, the results of the acylation of lysine and arginine showed that the acylation occurred on the  $\alpha$ -position. Indeed, the acylation of lysine with fatty acids has already been described in the literature. Whatever the fatty acid involved, two distinct fragmentation profiles were observed for the  $\alpha$  and the  $\epsilon$ -acyl-lysine [12,33]. The authors showed that for  $\varepsilon$  acylated lysine only one major daughter ion corresponding to  $[M - H_2O -$ COOH + H]<sup>+</sup> was found, whereas two major daughter ions corresponding to [M - 2 H<sub>2</sub>O + H]<sup>+</sup> and [M - H<sub>2</sub>O - COOH + H]<sup>+</sup> were obtained for the  $\alpha$ -acylated lysine. In the present study, the fragmentation of the parent ion of undecenoyl-lysine (m/z = 313) produced two major daughter ions corresponding to [M - 2 H<sub>2</sub>O + H]<sup>+</sup> and [M - H<sub>2</sub>O - COOH + H]<sup>+</sup>, showing that acylation took place on the  $\alpha$ -position. The fragmentation of the acylated arginine generated two major daughter ion corresponding to [M - NH2 - C - NH2 - NH] and [M - NH<sub>2</sub> - C - NH<sub>2</sub> - NH - H<sub>2</sub>O] which indicates also an acylation on the  $\alpha$ -position. This result was in perfect accordance with a previous study using oleic acid acid as acyl donor. Indeed, Dettori et al. (2017) used lysine and peptides containing lysine as acyl acceptors to show the ability of the aminoacylases from S. ambofaciens to perform the N-acylation reaction. They demonstrated that the extracted enzymes catalyse the acylation of the substrates, in alpha position without any acylation on the side chain despite the presence of Sam-ELA which is specific for the acylation of lysine epsilon amino group [9,12]. This result was related to the presence of the aminoacylase Sam-AA which is responsible of more than 85% of the  $\alpha$ -aminoacylase activity of the extracted enzymes [34].

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In order to extend the comprehension of the selectivity of the aminoacylases from *S. ambofaciens*, other molecules like taurine, hydroxylamine, 6-aminohexanoic acid, 4-amino-

benzoic acid, ethanolamine and the dipeptide carnosine were used as acyl-acceptor but no acylation reaction was observed (Tab.1). However, when 6-amino-1-hexanol was used, a small quantity of the product was detected by LC-MS-MS. This result suggested the negative influence of the carboxyl group at the end of the 6-aminohexanoic acid, which is replaced by a hydroxyl group on the 6-amino-1-hexanol. This may explain the lack of reaction when using the negatively charged amino acids (aspartic and glutamic acids). In order to understand the exact biological mechanism implicated, the structure of the aminoacylase has to be elucidated and a molecular modelling study has to be performed. Reactions using glycine ethyl-ester, methyl-ester and tert-butyl-ester as acyl-acceptor were also performed but no conversion was observed. These results showed that the aminoacylases from S. ambofaciens have a completely different acyl-acceptor selectivity as compared to a lipase such as CALB. Indeed, several author described the ability of CALB to perform the acylation of molecules such as carnosine and 6-amino-1-hexanol in organic solvent and ionic liquid [35–37]. They also compared the acylation of amino acids and amino acid derivatives like glycine and glycine tert-butyl-ester, showing an improvement of the conversion rate due to their better solubility in organic solvent. Goujard et al. (2004) evaluated the performances of CALB for the N-acylation reaction of the glycine with arachidonic acid in acetonitrile. They demonstrated that this enzyme was not efficient for the acylation of the glycine. However, the use of an activated amino acid containing tert butyl group on its carboxylic part led to a significant increase in the conversion yield [19]. Those results are completely different from the ones obtained in the present study where an activated amino acid or fatty acid doesn't allow any improvement of the activity. In order to check the enantio-selectivity of the enzyme, reactions using D-phenylalanine and L-phenylalanine were compared. No acylation of the D-phenylalanine was observed

suggesting that aminoacylases from S. ambofaciens are very specific to the natural L-amino

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acids. Koreishi et al. (2005) also showed that the 100 kDa aminoacylase from *S. mobaraensis* wasn't able to catalyse the hydrolysis of the N-acetyl-D-methionine while exhibiting a very high hydrolysis activity of the L-enantiomer, showing the same enantioselectivity as the one of the aminoacylases from *S. ambofaciens* [24].

406 Fig.2

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407 Tab.1

### 3.1.1. Acyl donor specificity

The evaluation of the specificity towards fatty acids was first performed by using lysine, arginine, methionine, leucine and phenylalanine as acyl-acceptors as these substrates presented high reactivity with different radical groups. Four different fatty acids were used in equimolar conditions (0.1 M of each substrate). The synthesis of N-acyl-amino acids with various fatty acids is reported in fig.3.A. The aminoacylases from S. ambofaciens showed different activities towards the fatty acids, depending on the amino acid used. The conversion yields obtained for N-acyl-L-phenylalanine derivatives were similar whatever the fatty acid used, with values from 10 to 15%, after 24 h. Likewise, N-acyl-L-methionine products were obtained with similar conversion yields, however a slight decrease of the conversion yield was noted when using oleic acid as acyl donor. The results obtained with arginine, lysine and leucine showed an increase of the conversion yields when increasing the fatty acid chain length from C8 to C12, and then, a decrease when oleic acid was used. This result is consistent with the work of Wada et al. (2002) who studied the substrate specificity of the acylase I from pig kidney towards different fatty acids, using arginine as acyl-acceptor. They showed nearly the same profile as the one observed with the aminoacylases from S. ambofaciens with an increase of the conversion yield from C10 to C12 fatty acid, then a significant decrease with C18 [38]. The final samples were analysed by LC-MS-MS in order

to evaluate the influence of the chain length on the regioselectivity of the aminoacylases from S. ambofaciens. The MS<sup>2</sup> spectrum after fragmentation of the parent ion of the product of each N-acyl-amino acid did not change by changing the fatty acid confirming that the ionisation of the product occurred on its amino acid part. This result confirmed that in all cases the acylation of the amino acids on the amine function was preserved. However, the structure of the lysine is particular because of the presence of two reactive amino groups. The products resulting from lysine acylation with the different fatty acids always corresponded to N- $\alpha$ -acyl-lysine derivatives except for octanoic acid that led to the synthesis of both  $\alpha$  and  $\epsilon$ acyl-lysine. As described in Fig.4, two products corresponding to the same molar mass were found with 1.5 min retention time difference. Those products presented different MS<sup>2</sup> fragmentation profiles where the highest peak corresponded to the  $\alpha$ -acylated lysine, the same as the one described previously, and the second one, with  $[M - H_2O - COOH + H]^+$  as major daughter ion, corresponded to the  $\varepsilon$ -acylated lysine. The presence of  $\varepsilon$ -octanoyl-lysine can be explained by the presence of Sam-ELA in the crude extract of S. ambofaciens which is similar to Sm-ELA from S. mobarensis described by Koreishi et al. with 80% sequence identity [10]. Sm-ELA was able to catalyse the acylation of lysine on its \epsilon-amino group with 58.7% conversion yield by using 10 mM octanoic acid, in 100 mM Tris-HCl, pH 7.0 buffer, at 45°C. However, Sm-ELA also led to 100% conversion by using lauric acid in the same conditions, which would not be the case with the corresponding enzyme from S. ambofaciens which seemed not able to catalyse the acylation of ε-amino group (fig. 1). A better acylation of the lysine ε-position was observed with lauric acid using Sm-PVA compared with the lysine αposition. No acylation occurred with octanoic acid [11]. The arginine presents also two free amine groups. But the guanidine side chain of this amino acid remains protonated under operational conditions (pH 8) preventing its acylation [6].

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In order to check if there is a structural relation between the regio-selectivity and the fatty acid chain length, different short chain fatty acids (acetic acid, butyric acid and valeric acid) were used with lysine as acyl-acceptor (Tab.2). The regioselectivity was evaluated by combining TLC and LC-MS-MS analyses (data not shown). In all reactions performed with fatty acid chains shorter than eight carbons, both  $\alpha$  and  $\epsilon$  mono-acylated products were found. However, there was a relation between the chain length and the regioselectivity since short carbon chains seemed to favour acylation on the ε-position. Also, regioselectivity towards amino group in α-position was higher with butyric acid as compared to isobutyric acid. This kind of relation has never been reported before. With molecules containing cyclic group such as benzoic acid, 4 amino benzoic and homovanillic acid, no product was detected by LC-MS-MS analysis, except for the phenylpropionic acid. This result can be explained by the short distance between the carboxylic group and the cyclic group which may lead to a steric hindrance. For the phenylpropionic acid, the LC-MS-MS analysis showed the presence of both mono-acylated products thus confirming the previous results. Even if this fatty acid contains 9 carbon atoms, the presence of the phenyl group made it similar to short length fatty acids in terms of regioselectivity. Reaction with azelaic acid (C9), which is a dicarboxylic acid widely used in cosmetics, was conducted in the same conditions. The results of the LC-MS-MS analysis showed the presence of a monoacylated product corresponding to the pseudo mass 317 m/z and a diacylated product corresponding to the pseudo mass 487 m/z. The MS<sup>2</sup> fragmentation profiles of the monoacylated product, suggested that acylation occurred on the α-amino group of the lysine. Another mass corresponding to 444 m/z formed by the condensation of two lysine units with both carboxylic parts of the azelaic acid was also found. This product represents a bolaamphiphile amino acid surfactant which belongs to a product family widely used in household products and cosmetics [3]. These results were confirmed by TLC-MS analysis which showed the three products described before. These results are of

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clear interest for further studies aiming to produce such products knowing that the enzymatic synthesis of bolaform surfactants using acylases in aqueous media has never been reported until now. Indeed, several authors used enzymes, such as lipases, in the synthesis of bolaform surfactants [39]. However, a chemical step have to be performed to recover the bolaamphiphilic surfactant. Valivety et al. (1998) performed the reaction of esterification of N-protected amino acids for the synthesis of bolaform surfactant using lipases. However, in order to recover the final product a chemical step had to be performed in order to remove the protection from the amino acids [39]. Consequently, it is the first time that the one-step enzymatic condensation on both carboxylic groups of a diacid is described.

In order to confirm the contribution of Sam-ELA to the acylation of the lysine with octanoic acid on its side chain, the gene sequence that express the production of Sam-ELA and Sam-PVA in *S. ambofaciens* was deleted. This allowed the expression of the sequence corresponding to Sam-AA, which was described as being of major importance in the N-α-acylation activity [34]. The transformed strain was cultivated and the enzyme was extract following the same steps as with the wild strain of *S. ambofaciens*. As shown in the fig.3 B, the presence of only Sam-AA allowed the N-acylation of the lysine on its α-position with all fatty acid used even with octanoic acid with nearly similar conversion rate. Knowing that Sam-PVA is similar to Sm-PVA described by Koreishi et al., 2006 as being specific to the acylation of amino acids with lauric acid only, this result confirms that the ε-acylation of lysine with octanoic acid observed with the aminoacylases from the non-transformed *S. ambofaciens* was due to the presence of Sam-ELA [11].

496 Fig.3:

- 497 Tab.2:
- 498 Fig.4:

#### 3.2. Influence of substrate concentration on the synthesis of N-α-lauroyl-L-lysine

Lauroyl-lysine is one of the most used acylated derivative of amino-acids among healthcare products due to its interesting surface active property and biological activity. Several studies reported the acylation of lysine on its ε-position by using lipases or acylases but very few enzymes were described as being able to catalyse the acylation of lysine on its  $\alpha$ -position [12,24]. In order to evaluate the influence of each substrate on the reaction rate, the concentration of one substrate was set to 100 mM while different concentrations of the substrate under study were used (from 25 to 250 mM for lysine and from 10 to 200 mM for lauric acid). The increase of lysine concentration from 25 to 150 mM led to an increase of the initial rate of the reaction. The curves of initial rates followed standard Michaelis-Menten model allowing the determination of Km and Vmax for the amino acid. According to the equation of Lineweaver and Burk, a Vmax and Ks of around 3.1 mM / h and 69 mM were found, respectively (Tab.3). However, at concentrations higher than 175 mM, a decay in the initial rate was observed. This decay might be due to an inhibition by the amino acid. The effect of the lauric acid concentration was conducted by using 100 mM lysine and concentrations from 0 to 200 mM of lauric acid. Between 10 and 50 mM lauric acid, a low increase in the conversion was observed but a decrease was observed at concentrations higher than 75 mM. This decrease can be attributed to mass transfer limitation. Wada et al., 2002 also attributed the decrease of the activity by increasing lauric acid concentration when using acylase I from Pig Kidney in buffer/glycerol mixture, to the fatty acid saturation which is reached at around 10 mM [38]The second reason could be the inactivation or the inhibition of the enzyme by the reaction product [10,15]. In this case, Vmax and Ki of around 4.4 mM h<sup>-1</sup> and 85 mM, respectively, were found (Tab.3). A Km of around 2 mM was also determined.

Tab.3:

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#### 3.3. Influence of metal ions on the reaction rate

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As described by Dettori et al., [12] the genes encoding the aminoacylases from S. ambofaciens present close similarities to the ones encoding acylases from S. mobarensis, which were described by Koreishi et al. [9,11,25]. Two of those identified acylases (Sm-AA and Sm-ELA) were described as belonging to a metallo-dependant family which can be influenced by the addition of metal ions. In order to evaluate the effect of metal ions on the activity of acylases from S. ambofaciens, Co2+, Zn2+ and Cu2+, described as the most influencing ones, were added in the reaction medium [10,11,24,25]. As shown in Fig.5A, for the same concentration of the three metal ions (0.1 mM), the addition of Co<sup>2+</sup> ion led to a three time increase of the initial rate while Zn<sup>2+</sup> and Cu<sup>2+</sup> ions only led to a low increase of the initial rate. Very low conversion yield was observed when EDTA (metal scavenger) was used, showing that the observed activity originates from a metallo-enzyme. Indeed, even if the initial rate in the presence of EDTA was similar to the one without, a decrease of 75% of the conversion rate was observed. This result can be explained by a decay of the enzymatic stability when a chelating agent is used. The same result was reported by different authors with others metallo-enzymes, where a partial decrease of the activity was registered when using EDTA [10,24,31]. However, the conversion rate at thermodynamic equilibrium decreased to 11 and 17% using Zn<sup>2+</sup> and Cu<sup>2+</sup> ions, respectively at 0.1 mM, while it remained stable with Co<sup>2+</sup>. The influence of Co<sup>2+</sup> concentration was therefore investigated (Fig.5B). The increase of the Cobalt concentration up to 0.4 mM allowed an important increase (about 6 times) of the initial rate of the reaction (Tab.4). The thermodynamic equilibrium in those conditions was reached after only 4 h reaction time instead of around 72 h without CoCl<sub>2</sub>. The LC-MS-MS analysis did not show any modification in the regioselectivity of the enzyme. Koreishi et al. 2005 also studied the influence of metal ions on the hydrolytic activity of S. mobarensis aminoacylaseS. Their results were similar to the ones of the present study with an

increase of around 3 times of the activity by adding 0.1 mM CoCl<sub>2</sub>. However, they observed a decrease of around 80% of the activity by adding Cu<sup>2+</sup> ion, which was not the case of the acylases from *S. ambofaciens* [24].

551 Fig.5:

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552 Tab.4:

## **Conclusion**

The performances of the aminoacylases from S. ambofaciens in terms of substrate specificity and regioselectivity were evaluated. This work showed for the first time a relationship between the side chain length of the amino acids, especially with non-polar aliphatic amino acids, and the conversion rate (from 6% for glycine to 23% for leucine). Furthermore, the side chain nature of the amino acid was shown to play an important role. Best results were obtained with positively charged amino acids (lysine and arginine) while no reaction was obtained with the negatively charged ones (aspartic and glutamic acids). The results concerning the acyl donor specificity showed that the regioselectivity was affected by the fatty acid chain length which was attributed to the presence of Sam-ELA in the crude extract of S. ambofaciens. Indeed, this enzyme is able to perform the acylation of the lysine on its side chain and appears to be also specific for short chain fatty acid. Moreover, for the first time, the possible synthesis of bolaamphiphile amino acid surfactant using azelaic acid and lysine was shown. The study concerning the synthesis of lauroyl lysine in various operational conditions showed an inhibition of the enzymes by the product of the reaction. In order to overcome this issue, the future development of processes using continuous flow reactors such as packed bed reactors with immobilized aminocylases from S. ambofaciens appears as a promising way. The results obtained in the presence of metal ions showed that it was possible to significantly increase the reaction by adding CoCl<sub>2</sub> that allowed to reach the

thermodynamic equilibrium in few hours. Further works are in progress to intensify the process increasing the conversion yield in order to develop a green process for the production of eco-friendly lipoaminoacids as bio-based surfactant using *S. ambofaciens* aminoacylases in aqueous media.

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#### **Figures:**

Fig.1 Conversion yields for the synthesis reaction using various amino acids (100 mM) and 10-undecenoic acid (100 mM). Reaction conditions: 3 days, at 45°C, 250 rpm agitation, in 25 mM Tris-HCl buffer, NaCl 50mM, pH 8, using 1 g/L enzyme

Fig.2 MS<sup>2</sup> spectra after fragmentation of the parent ion of (A) N-10-undecenoyl-threonine (B) N-10-undecenoyl-cysteine synthesized using aminoacylases from *S. ambofaciens* 

Fig.3 Conversion yields using various amino acids (100 mM) and fatty acids (100 mM). A. acylation of arginine, methionine, leucine and phenylalanine using the aminoacylases from *S. ambofaciens*, B. Acylation of lysine using the aminoacylases from *S. ambofaciens* and Sam-AA from transformed *S. ambofaciens*. Reaction conditions: 1 day, at 45°C, 250 rpm agitation, in 25 mM Tris-HCl buffer, NaCl 50mM pH 8, using 1 g/L enzyme

Fig.4 Mass analysis of octanoyl-lysine using aminoacylases from *S. ambofaciens* (A) single ion chromatogram of the acylated product  $[M+H]^+$  parent ion (m/z = 273); (B)  $MS^2$  spectrum after fragmentation of the parent ion at 5.79 min retention time; (C)  $MS^2$  spectrum after fragmentation of the parent ion at 7.36 min retention time.

Fig.5: Influence of metal ions on the initial rate of the α-lauroyl-lysine (C12K) synthesis reaction (0.1 M of lauric acid and lysine) using *S. ambofaciens* acylases in tris HCl 25mM NaCl 50 mM buffer pH 8 at 45°C, (A) Produced concentration of C12K during the time according to the metal ion used and the concentration of COCl<sub>2</sub> used (B)

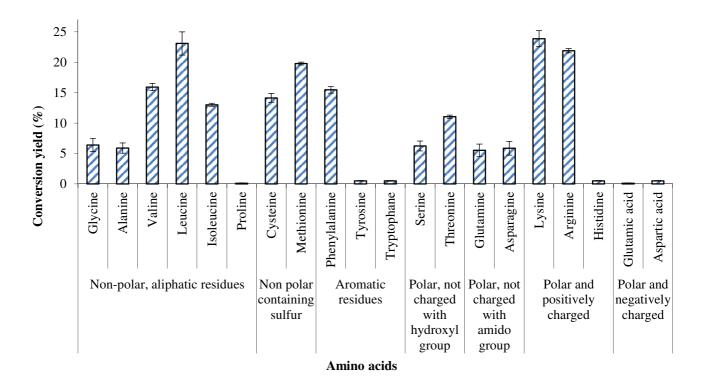
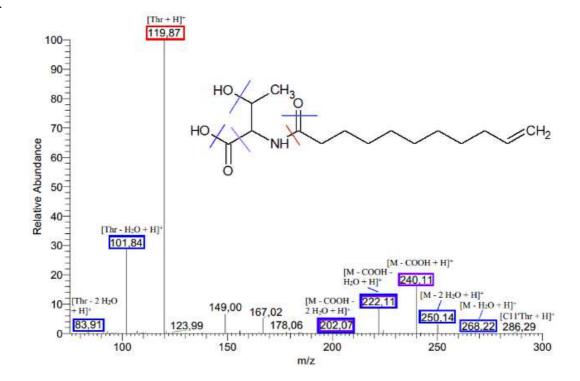


Fig.1





В

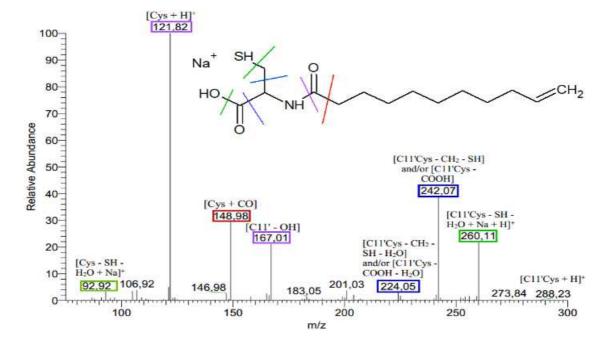
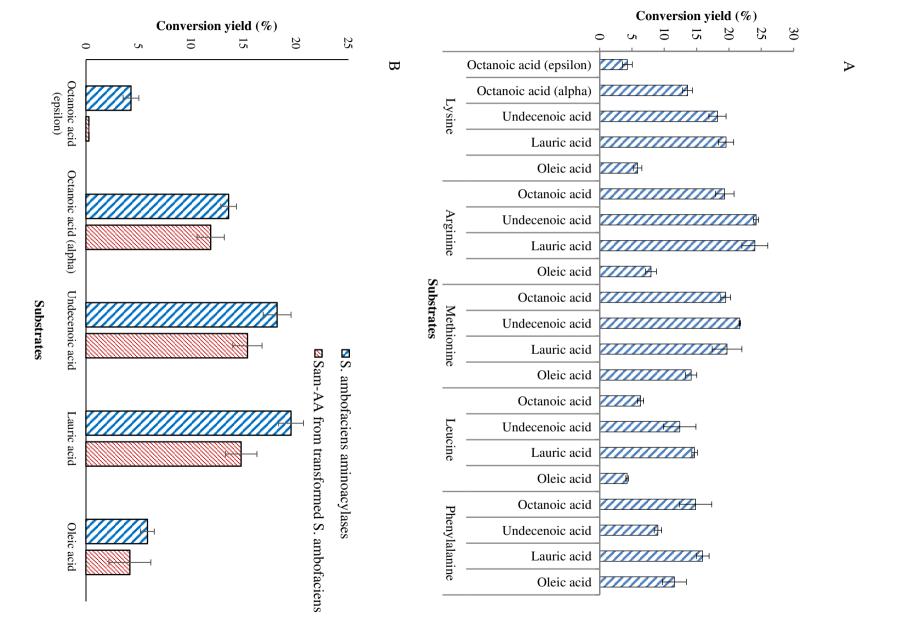
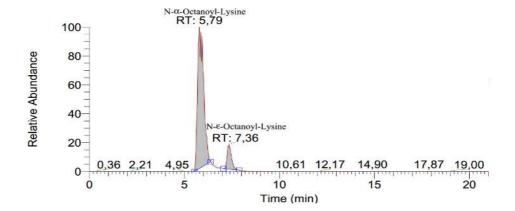
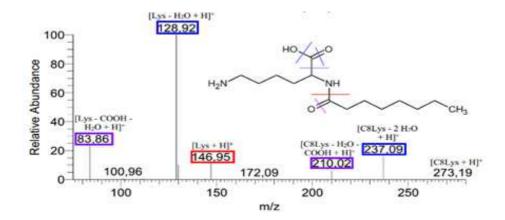


Fig.2





В



C

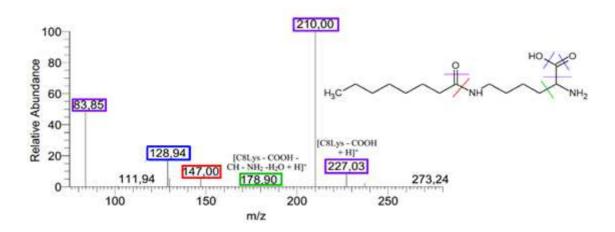
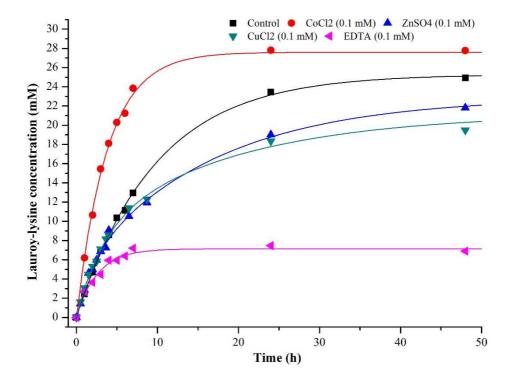


Fig.4



В

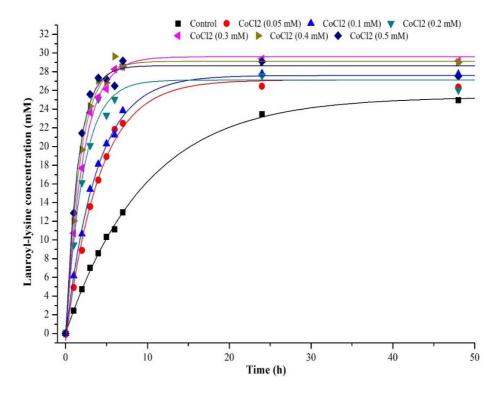


Fig.5

# Table:

Tab.1 structure of the acyl receivers used

Fatty acid	Molecular structure	Reaction
Taurine	H <sub>2</sub> N————————————————————————————————————	-
Hydroxylamine	H <sub>2</sub> N—OH	-
Ethanolamine	H <sub>2</sub> N OH	-
6-aminohexanoic acid	$H_2N$ OH	-
6-amino-1-hexanol	H <sub>2</sub> N OH	+
4-amino-benzoic acid	$H_2N$ OH	-
Carnosine (N-β alanyl, L- histidin)	O OH N NH NH	-
Glycine	H <sub>2</sub> N OH	+
Glycine methyl-ester	$H_2N$ $O$ $CH_3$	-
Glycine ethyl-ester	$H_2N$ $O$ $CH_3$	-
Glycine tert-butyl- ester	$H_2N$ $O$ $CH_3$ $CH_3$	-

Tab.2 structure of the fatty acids used

Eatty soid	Molecular structure	Acylation position		
Fatty acid	Moleculal structure	Alpha	Epsilon	
Acetic acid	H <sub>3</sub> C OH	+	+	
Isobutyric acid	H <sub>3</sub> C OH	+	+	
Butyric acid	Н <sub>3</sub> С ОН	+	+	
Valeric acid	H <sub>3</sub> C OH	+	+	
Octanoic acid	Н <sub>3</sub> С ОН	+	+	
10-undecenoic acid	H <sub>2</sub> C OH	+	-	
Lauric acid	H <sub>3</sub> C OH	+	-	
Oleic acid	H <sub>3</sub> C 7 OH	+	-	
Azelaic acid <sup>a</sup>	но	+	-	
Phenylpropionic acid	ОН	+	+	

a: acylation on both carboxylic group

Tab.3: Kinetic parameters of the substrates concentrations for  $\alpha$ -lauroyl-lysine synthesis

	Lysine	Lauric acid
Vmax (mM/h)	$3.1 \pm 0.2$	$4.4 \pm 0.2$
Km (mM)	69 ± 7	$2.1 \pm 0.3$
Ki (mM)	-	$85.5 \pm 7.8$

Tab.4 Initial rate evolution depending on the condition.

Metal ion	No metal ions	EDTA	CuCl <sub>2</sub>	ZnSO <sub>4</sub>	CoCl <sub>2</sub>	CoCl <sub>2</sub>				
Concentration (mM)		0.1	0.1	0.1	0.1	0.05	0.2	0.3	0.4	0.5
Initial rate	$2.6 \pm 0.4$	2.4 ±	2.9 ±	3.2 ±	6.3 ±	4.6 ±	12.1	14.2	14.6	15.8 ±
(mM/h)		0.2	0.4	0.0	0.6	0.1	$\pm 0.2$	$\pm 0.0$	± 1.8	2.0

# Graphical abstract

