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Production of biosurfactant using the endemic bacterial community of a PAHs contaminated soil, and its potential use for PAHs remobilization Florian CAZALS^{1, 2, 3*}, David HUGUENOT¹, Marc CRAMPON³, Stéfan COLOMBANO³, Stéphanie BETELU³, Nathalie GALOPIN², Arnault PERRAULT², Marie-Odile SIMONNOT⁴, Ioannis IGNATIADIS³, Stéphanie ROSSANO¹

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10 11

16 Abstract

17 Biosurfactants are surface-active agents produced by microorganisms whose use in soil remediation processes is increasingly discussed as a more environmentally friendly 18 alternative than chemically produced surfactants. In this work, we report the production of a 19 biosurfactant by a bacterial community extracted from a polluted soil, mainly impacted by 20 PAHs, in order to use it in a soil-washing process coupled with bioremediation. Nutrient 21 balance was a critical parameter to optimize the production. Best conditions for biosurfactant 22 production were found to be 20 g/L of glucose, 2 g/L of NH₄NO₃, and 14.2 g/L of Na₂HPO₄, 23 corresponding to a C/N/P molar ratio equal to 13/1/2. Purification of the produced 24 biosurfactant by acidification and double extraction with dichloromethane as a solvent 25 allowed measuring the Critical Micellar Concentration (CMC) as equal to 42 mg/L. The 26 capacity of the purified biosurfactant to increase the apparent solubility of four reference 27 PAHs (naphthalene, phenanthrene, pyrene and benzo[a]pyrene) was completed. The 28 29 solubilisation ratios, in mg of PAH/g of biosurfactant for phenanthrene, pyrene and benzo[a]pyrene are 0.214, 0.1204 and 0.0068, respectively. Identification of the bacteria 30 found in the colony producing the biosurfactant showed the presence of bacteria able to 31

produce biosurfactant (*Enterobacteriaceae*, *Pseudomonas*), as well as, others able to degrade
PAHs (*Microbacterium*, *Pseudomonas*, *Rhodanobacteraceae*).

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35 Keywords

Bacterial biosurfactant, Polycyclic Aromatic Hydrocarbons, desorption, solubility, Critical
Micellar Concentration

38 **1 - Introduction**

39

Surface-active agents produced by microorganisms, including bacteria, are called
biosurfactants. These compounds are able to reduce the surface tension between liquids,
solids or gases. According to this property, biosurfactants can be used to increase the
solubility of hydrophobic compounds in water (Soberón-Chávez et al., 2005).

When present in soil, Polycyclic Aromatic Hydrocarbons (PAHs), have low accessibility,
availability and solubility due to their proper physical and chemical characteristics (especially
molecular weights) and to those of the soil (Haritash and Kaushik, 2009).

Biodegradation of PAHs is then limited due to their low bioavailability, defined as "the part of 47 the compound which is freely available to cross an organism membrane from the medium the 48 organism inhabits at a given point in time" (Riding et al., 2013). Due to their hydrophobicity, 49 in soil, an important part of the PAHs, isn't bioavailable, but still bioaccessible (Riding et al., 50 2013) namely "could become bioavailable if desorbed from the solid matrix". However, 51 increasing the PAHs solubility could lead to an increase of the risk of PAHs dispersion in the 52 environment. Thus, a thorough control of the flow during their solubilisation process is 53 crucial in order to limit this risk. 54

55 Chemical surfactants are used in bioremediation to increase the apparent solubility of PAHs 56 in groundwater (Lamichhane et al., 2017; Lukić et al., 2016), but the bioavailability of PAHs 57 trapped in chemical surfactant micelles is not insured (Shao et al., 2017) and the toxicity of 58 the chemical surfactant itself could lead to a decrease in the biodegradation process. For 59 example, addition of SDS in high concentrations (100 and 500 mg/kg) increased the PAHs 60 concentration in water , but it decreased the biodegradation of PAHs due to the toxicity of the 61 SDS towards the bacterial strains (Deschênes et al., 1995; Lima et al., 2011).

To solve this problem, the use of biosurfactants for soil remediation has been increasingly 62 studied and is close to be implemented (Koch et al., 1991; Souza et al., 2014; Whang et al., 63 2008). In comparison with chemically synthetized surfactants which are often toxic and 64 persistent in the environment, biosurfactants have no proven toxicity and are more 65 biodegradable (Mulligan, 2005; Rufino et al., 2014), which is more in line with their use for 66 environmental applications. Some studies (Lima et al., 2011; Varjani and Upasani, 2017a) 67 suggest that the biodegradation of biosurfactants happens after the consumption of other 68 69 carbon sources and should not affect the solubility of entrapped pollutants.

Remediation of soils polluted by hydrophobic compounds could lead to better achievements if biosurfactants are used. This is due to the capacity of microorganisms to better degrade pollutants trapped in micelles of biosurfactant. Using a bacterial culture able to produce biosurfactant and to metabolize PAHs for bioremediation could then combine the two fold effect of soil washing and biodegradation (Ali Khan et al., 2017; Chebbi et al., 2017; Gabet, 2004; Hassanshahian, 2014; Lamichhane et al., 2017; Li et al., 2015; Portet-Koltalo et al., 2013).

Numerous studies have investigated the optimization of the production of biosurfactants
from bacteria isolated from polluted sites to this purpose (Ali Khan et al., 2017; Chebbi et al.,
2017; Hassanshahian, 2014; Koch et al., 1991; Liang et al., 2017; Pereira et al., 2013; SoberónChávez et al., 2005).

Soil and groundwater remediation of PAHs often focus on the remediation of the source zone. 81 The remediation of pollution plumes is economically limited by the cost of the remediation 82 processes, and technically by the difficulty to access the sorbed PAHs, due to their 83 hydrophobicity. The association of soil washing with biosurfactant, coupled with a 84 bioremediation process could be an innovative method for the remediation of sites impacted 85 by PAHs pollution plumes though the fate of solubilized PAHs needs a careful monitoring 86 that must control and promulgate actions to avoid their dispersion in the environment. The 87 use of biosurfactant will increase the bioavailability of the PAHs and so their biodegradability 88 89 which will limit their dispersion. Due to the biodegradability of the biosurfactant, the solubilisation of the PAHs will be time limited and the risk of dispersion in the environment 90 will decrease quickly. 91

In this present work are reported i) the extraction and isolation from a polluted soil impacted by PAHs of bacterial populations able to produce biosurfactants, and ii) the development of a method for the production and recovery of the biosurfactants. Each isolated colony was tested for the production of biosurfactant. After the selection of one colony, tests were performed to optimize the production medium based on nutrient nature and concentrations. Two types of carbon sources, soluble and insoluble, were compared. The effect of the C/N/P ratios for the production of this biosurfactant has also been studied.

99 2 - Materials and methods

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101 **2.1. Chemicals**

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Anhydrous D-glucose (99% purity), K₂HPO₄ (97% purity), NH₄NO₃ (>99% purity)
FeCl₃.6H₂O (99% purity) and Na₂HPO₄ (>99% purity) were purchased from VWR (VWR
International SAS, France).

KH₂PO₄ (analytical standard), KNO₃ (99% purity), MgSO₄.7H₂O (>98% purity), CaCl₂.2H₂O
(>99% purity), NaCl (99.5% purity) dichloromethane (>99.8% purity), naphthalene (NAPH)
(99% purity), phenanthrene (PHE) (98% purity), pyrene (PYR) (97% purity) and
benzo[*a*]pyrene (BaP) (96% purity) were purchased from Sigma-Aldrich (Merck KGaA,
Darmstadt, Germany).

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112 2.2. Site of interest and soil sampling

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114 A former coke plant located in the Northeast of France, in the region of Lorraine, was 115 selected for collecting water and soil samples and for *in-situ* implementation of the developed 116 remediation process.

117 The site is impacted by pure PAHs, BTEX and other hydrocarbons (TPH) in the source zone 118 through a layer of 4 m depth. A pollution plume due to an aquifer passing through the site at 119 a depth of 4 to 8 m affects the surroundings of the site.

To extract bacteria able to resist to PAHs, several soil samples were performed by drilling in the pollution plume area. Three samples were collected from three different depths, the first one at 2 m, the second one from 2 to 4 m depth and the last one, in the saturated zone from 4 to 8 m. Each sample was homogenized and aliquots were sent to Synlab (Gennevilliers, France) for the analysis of the main pollutants, hydrocarbons, phenols, BTEX and PAHs, in water and soil. The results showed high concentrations of PAHs, both in water and in soil.

The total BTEX concentration in water was 8400 μ g/L, total concentration of the 16 PAHs (US EPA) was about 10000 μ g/L and the concentration of the total hydrocarbons (C10 to C40) in water was 14000 μ g/L. The corresponding concentrations in soils were 33, 2164 and 2486 mg/kg, respectively. A more detailed summary of these analyses can be found in supplementary materials (SM1). Granulometric and chemical analyses of the soil sample are detailed in supplementary materials, SM2 and SM3, respectively. 132

133 2.3. Experimental design

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To achieve the production of biosurfactant and evaluate its potential use in bioremediation processes, different steps were necessary. Figure 1 presents the successive steps of the study, from the extraction of the soil bacteria to the purification and characterization of the produced biosurfactant and the PAHs desorption assays, with an important focus on the optimization of the biosurfactant production.

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141 **2.3.1.** Extraction of cultivable bacteria from contaminated soil

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Aliquots of 10g of the three contaminated soil samples were added to 40 mL solution 143 containing 0.016 g of MgSO₄.7H₂O, 0.001 g of CaCl₂.2H₂O, 0.040 g of KH₂PO₄, 0.040 g of 144 K₂HPO₄, 0.040 g of NH₄NO₃ and 0.003 g of FeCl₃.6H₂O with a pH fixed to 7, in 50 mL flasks 145 146 (adapted from Mcgrath et al., 1997). These suspensions were agitated for 48 h (25 °C, 180 rpm). Then 4 mL were taken and added to 71 mL of the same minimum medium in 100 mL 147 flasks. The suspensions were stirred for 120 h (20 °C, 180 rpm). These media were then 148 149 spiked with pure PAHs in powder (Sigma-Aldrich (Merck KGaA, Darmstadt, Germany) at concentrations above their solubility) as sole carbon source, in order to select only PAHs 150 tolerant bacteria. Colonies were obtained after plating on solid minimal medium (Bois et al., 151 152 2011), and were used for next experiments.

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2.3.2. Selection of bacteria able to produce biosurfactant

Each of the obtained colonies was cultivated in fresh media and then tested for the 156 production of biosurfactant. A first experimental medium was prepared following the 157 literature, selecting the most widespread components on biosurfactant production media to 158 increase the probability of having a positive result (Ali Khan et al., 2017; Davis et al., 1999; 159 Henkel et al., 2012; Liu et al., 2014; Mulligan et al., 1989; Ochsner et al., 1996; Pereira et al., 160 2013; Reis et al., 2010; Robert et al., 1989; Rocha e Silva et al., 2014). This first medium was 161 composed of 1 mL of colza oil, 50 mL of water solution containing 0.1 g of NH₄NO₃, 0.25 g of 162 K₂HPO₄, 0.25 g of Na₂HPO₄ and 0.25 g of NaCl in 100 mL flasks. The suspensions were 163 agitated for 7 days (25 °C, 180 rpm). After 7 d, the collapsing drop test (Jain et al., 1991) was 164 used to prove the production of biosurfactant. A drop of each suspension was placed on a 165 colza oil coated glass side. Visually, the presence of biosurfactant can be observed: the 166 collapsed drop indicates the presence of biosurfactant, when a stable drop indicates its 167 absence, or its presence at a very low concentration. 168

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2.3.3. Optimization of the biosurfactant production medium

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For the optimization of the biosurfactant production medium, two carbon sources (colza oil and glucose), and four nutrients (NH₄NO₃, Na₂HPO₄, K₂HPO₄, KNO₃), were selected in order to highlight the effect of C, N, P and K concentrations in the biosurfactant production. For each experiment, running time was 7 d, and the flasks were agitated at 180 rpm at 20 °C. In the principle of having a fast, cheap and easily reproducible biosurfactant production, the two carbon sources selected were colza oil and glucose.

177 Several tens of experiments were carried out following the scheme presented in Figure 2.

178

179 The first step of assays was achieved with the purpose of having an equal molar concentration

180 of 0.05 mol/L of K, N and P in the medium where they were present. The two carbon sources,

181 glucose and colza oil, were tested at 20 g/L.

The Collapsing Drop Test was performed daily to assess the production of biosurfactant. The positive result of the test and the speed of production were the two critical factors for selecting the best carbon source to optimize the production of biosurfactant. The objective of the second step was to evaluate the effect of the presence or absence of the three different nutrients on the biosurfactant production by the bacterial community. The same protocol was used to assess the production of biosurfactant.

After the first two steps of tests, following the tests used, only a medium with glucose as carbon source and NH_4NO_3 and Na_2HPO_4 as nutrients was selected. Next experiments were carried out with various concentrations of each component to optimize the production of biosurfactant: glucose, NH_4NO_3 and Na_2HPO_4 concentration varied from 20 to 40 g/L, 1 to 4 g/L, 7.1 to 14.2 g/L, respectively. Table 1 presents the composition of the media (named from A to I) where biosurfactant production was achieved. Each medium will be named after this table in the following of the text.

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196 **2.3.4.** Upscaling of biosurfactant production in batch bioreactor

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Once the optimization of the medium used for biosurfactant production was achieved, a scale-up process to increase the production volume started. The objective of this scale-up process was to obtain a large quantity of biosurfactant for its characterization and to use it in PAHs desorption experiments in dynamic sand and soil columns.

The production was carried out in bioreactors of 5 L and the impact of several physical and chemical conditions on biosurfactant production was tested: (i) temperature (10, 15, 20 and 40° C), (ii) pH (from 5 to 8) and (iii) water oxygenation of the batch bioreactor (bulk solution between below saturation and full saturation in O₂).

For each bioreactor, pH, redox potential (with Pt/Ag-AgCl probe), temperature, conductivity and dissolved O_2 content in the fluid were continuously monitored using probes (Mettler Toledo, Viroflay, France) immersed in the batch bioreactor and were recorded continuously
with a data acquisition device (Keithley Instruments, model 2700, Cleveland, OH, USA).

2.4.1. Extraction and purification of the biosurfactant

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211 **2.4. Characterization of the biosurfactant**

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214 To extract and purify the produced biosurfactant from batch bioreactors, the protocol was adapted from previous ones (Varjani and Upasani, 2017b). Three litres of the liquefied 215 foam obtained from three different 5 L bioreactors, were centrifuged, and the supernatant 216 filtered at 0.22 µm (MF-Millipore[™] 0.22 µm mixed cellulose esters membrane, Merck 217 Millipore), to separate the bacteria from the biosurfactant. The obtained filtrate was then 218 acidified to pH 2 with hydrochloric acid and stored at 4 °C for 48 h. The precipitate resulting 219 220 from the acidification was extracted by centrifugation and filtration at 0.22µm, and then redissolved in ultrapure water. The pH was finally stabilized at 7. The biosurfactant was then 221 extracted from water two times using dichloromethane in separating funnel. 222 223 Dichloromethane was then evaporated under a nitrogen flux during 1 h, and finally freezedried (Christ Alpha 2-4 LSCPlus freeze-dryer), to eliminate the remaining water. 224

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2.4.2. Determination of the Critical Micellar Concentration (CMC)

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To evaluate the CMC, the curve of the surface tension (mN/m) between the solution of purified biosurfactant and air, as a function of the concentration of biosurfactant in solution was made (Mahalingam and Sampath, 2014; Patowary et al., 2017; Varjani and Upasani, 2017b). Surface tensions were measured using a KRUSS DSA-100 goniometer and the method of the pending drop (using Young-Laplace fit directly from the Advance software combined to the KRUSS DSA-100). Using the tangent intersection method on the twodifferent parts of the curve, CMC was graphically evaluated and reported.

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2.4.3. PAHs solubilisation assays

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238 The ability to solubilize PAHs by the produced biosurfactant was tested. Four PAHs: NAPH (99% purity), PHE (98% purity), PYR (97% purity) and BaP (96% purity) were used as 239 240 representative of the 16 priority PAHs. The PAHs were introduced in flasks in pure acetone at concentrations equal to 10 times their solubility in water at 25 °C. On each batch, 6.4 mg of 241 NAPH (solubility at 25 °C = 32 mg/L), 0.24 mg of PHE (solubility at 25 °C = 1.2 mg/L), 0.032 242 mg of PYR (solubility at 25 °C = 0.140 mg/L) and 0.0016 mg of BaP (solubility at 25 °C = 6 243 μ g/L) were introduced. After evaporation of the acetone for 24 h (and formation of PAHs 244 crystals), the flasks were filled with 20 mL of different solutions of biosurfactant at different 245 concentrations (final concentrations of 1/2 CMC, 1 CMC, 5 CMC, 10 CMC and 20 CMC). After 246 247 24 h at 20 °C with agitation at 180 rpm, extractions by Solid Phase Extraction (SPE) were performed to evaluate the solubilized quantity of PAHs (Ma et al., 2010). Extracts were 248 analysed by Gas Chromatography coupled with Mass Spectrometry for PAHs concentrations 249 (Shimadzu QP 2020). For the extraction, SPE columns from Supelclean Envichrom (ENVI -250 Chrom P SPE Tube) were used. The extraction protocol was adapted for this specific use. The 251 columns were conditioned with 2 mL dichloromethane, followed by 2 mL MilliQ water. The 252 water samples were percolated, and the cartridges were then washed with 3 mL MilliQ water. 253 254 The cartridges were finally dried with vacuum aspiration for 2 h, and samples were eluted with 5 mL dichloromethane before analyses. Due to the low quantity of biosurfactant 255 extracted and purified, the tests could not be performed in triplicate. 256

257

258 2.5. Characterization of the biosurfactant-producing bacterial community

Firstly, DNA extractions were performed on culture from bioreactor producing 260 biosurfactant. A volume of 10 mL of culture were centrifuged and the supernatant discarded. 261 DNA extractions were carried out on the pellet using Fast Prep DNA kit for soils (MP 262 263 Biomedicals) following the manufacturer's instructions. The total extracted DNA was then quantified with a Quantus Fluorometer (Promega) and stored at -20 °C prior to further 264 analyses. To characterize the bacterial community structure and the diversity in the 265 bioreactor, Capillary Electrophoresis-Single Strand Conformational Polymorphism (CE-266 SSCP) profiles were performed based on an amplicon of the gene coding for 16S rDNA, and 267 amplified fragments of the gene coding for 16S rDNA were cloned and sequenced to identify 268 bacteria from the community. 269

For CE-SSCP fingerprints, about 200 bp of the V3 region of bacterial 16S rRNA genes were 270 271 amplified from DNA extract with the forward primer w49 (5'-ACGGTCCAGACTCCTACGGG-3'; E. coli position, 331) and the reverse primer w34 (5'-TTACCGCGGCTGCTGGCAC-3'; E. 272 coli position, 533). The 5' end was labelled with the fluorescent dye FAM, 25 cycles, 273 274 hybridization at 61 °C, and 30 s elongation at 72 °C were used. One µL of diluted (5- to 100-275 fold in nuclease-free water) PCR product was added to a mixture of 18.8 µL of deionized formamide and 0.2 µL of Genescan-600 LIZ internal standard (Applied Biosystems). To 276 277 obtain single-strand DNA, samples were heated denaturised for 5 min at 95 °C, and immediately cooled on ice. CE-SSCP analyses were performed on an ABI Prism 310 genetic 278 analyser using a 47-cm length capillary, a non-denaturating 5.6 % CAP polymer (Applied 279 Biosystems) and the following electrophoresis conditions: run temperature 32 °C, sample 280 injection for 5 s at 15 kV, data collection for 35 min at 12 kV. SSCP data was analysed using 281 BioNumerics V7.5 (Applied Maths) and lining SSCP profiles up to the internal standard and 282 283 to a common baseline.

The bacterial 16S rRNA gene sequences were then amplified from the extracted DNA. Universal bacterial primers 8F (5'-AGAGTTTGATCMTGGCTCAG-3') and 1406R (5'-ACGGGCGGTGTGTGTRC-3') were used with elongation at 55 °C and 30 cycles. Purified PCR

products (Macherey-Nagel) were used to build a gene library with the TOPO-TA Cloning Kit 287 and TOP 10 chemically competent cells (Invitrogen). Sequences were then analysed by 288 Sanger sequencing (Eurofins genomics, Germany). Consensus 16S rRNA gene sequences 289 290 were constructed and manually verified with the BioEdit program version 7.5.2 (Hall, 1999). The phylogenetic assignment was based on sequence analysis by the Ribosomal Database 291 Project version 11.5 classifier tool (Wang et al., 2007) and by the NCBI Blast tool 292 (http://www.ncbi.nlm.nih.gov/blast/Blast.cgi). Phylogenetic tree was built with the MegaX 293 software (version 10.0.1). All sequences (5 clones corresponding to 5 peaks on fingerprint 294 295 profiles) were posted on GenBank platform (NCBI) under accession numbers MN422137 to 296 MN422141.

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298 2.6. Chemical analysis methods

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To analyse the biosurfactant in cultures, a High Performance Liquid Chromatography with a fluorescence detector (HPLC-FLD, Hitachi LaChrom Elite® L-2400) was used. The media were analysed with a Reverse Phase column (5 mm, 25 cm × 4.6 mm) RP C-18 (Purospher®, Merck), and as mobile phase a mix of acidified water (pH =2.5) and acetonitrile at a ratio of 25:75 v:v (Mubarak et al., 2015). Due to absence of standard for the biosurfactant, only relative quantifications between the different experiments were possible with peak area values.

For PAH analyses, a GC-MS with a 30 m column (RXi®-5Sil MS from RESTEK, length: 30 m, thickness: 0.25 μm and diameter: 0.25 mm) was used. The oven temperature was set at 80 °C at the beginning of the analysis, and then increased up to 300 °C in 25 min and maintained at 300 °C for 10 min. Helium was used as the gas carrier, at a flow rate of 1.30 mL/min. For the mass spectrometer, the ion source temperature was set at 240 °C and the detector voltage at 1.20 kV (adapted from Crampon, 2015).

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14 2.7. Statistical analysis method

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316 Statistical analysis of the significant differences between data have been made with XLSTAT

- 317 software, version 2019.3.2 (Addinsoft).
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319 3. Results and discussion

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321 **3.1. Optimization of the production of biosurfactant**

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After plating the extraction medium, 15 different colonies were grown. These colonies 323 324 were re-inoculated in fresh medium in batches, filled with the first biosurfactant production medium (1 mL of colza oil, 50 mL of water solution containing 0.1 g of NH₄NO₃, 0.25 g of 325 K₂HPO₄, 0.25 g of Na₂HPO₄ and 0.25 g of NaCl). Only one of the batches had a positive 326 327 response on the collapsing drop test. The three replicates of the batch showed a positive result, reinforcing the reliability of the result. In addition, an emulsion was visible in the 328 batches between the two immiscible fluids, the aqueous medium and the colza oil. This batch 329 330 seems to be the only one inoculated by a colony containing bacteria able to produce biosurfactant with these conditions. The colony (containing several bacterial species), coming 331 from the 2 to 4 m depth sample, was then selected as the most suitable colony for 332 333 development and used in the following tests.

The second step of the assays, with colza oil medium and glucose medium was monitored for two weeks. After 7 days, in the medium containing 20 g/L glucose, 2 g/L NH_4NO_3 , and 7.1 g/L Na_2HPO_4 , a positive result was obtained on the collapsing drop test. It took 14 days to obtain a biosurfactant production in the medium containing 20 g/L colza oil, 2 g/L NH_4NO_3 , 7.1 g/L Na_2HPO_4 and 4.35 g/L K_2HPO_4 . No other medium allowed the production of biosurfactant. The medium containing glucose allowed a faster production of biosurfactant.
Glucose being an easily degradable carbon source, its biodegradation is faster, thus
improving the bacterial growth and so the production of the biosurfactant. In the case of any
development of a biosurfactant production process, the speed of production is a crucial
parameter. Glucose was then selected for the next experiments.

Variations of the selected medium with a lack of glucose, of NH_4NO_3 or of Na_2HPO_4 were tested, but did not show any production of biosurfactant. The presence of the two nutrients with glucose is mandatory to produce biosurfactant for the selected bacterial community.

347 The purpose of the next step of assays was to optimize the ratio of each nutrient and glucose to increase the production of biosurfactant. Each batch showed a production of biosurfactant, 348 with positive collapsing drop tests. Using HPLC-Fluo analysis on these media, it was found 349 that the medium producing more biosurfactant was medium A (Figure 3). These 350 concentrations matched with a C/N/P ratio equal to 13/1/2. It was deduced that a C/N ratio 351 higher than 10 was mandatory for the biosurfactant production. This result was consistent 352 with others biosurfactant production studies, which have proved the influence of the C/N 353 ratio on biosurfactant production (Karanth, 1989; Reis et al., 2010). Influence of phosphorus 354 had been studied (Mulligan et al., 1989; Wei et al., 2007) and differed in function of the 355 bacterial community and biosurfactant type. Tukey analysis of the significant differences 356 between the different conditions shows different groups of result. The first condition, 357 medium A with the higher peak area, is in a distinct group than the others, meaning that it is 358 significantly different from the other (p<0.05, Tukey ANOVA test), supporting the fact that 359 360 this medium is the best for the biosurfactant production.

361 This medium was then selected as biosurfactant production medium for the next362 experiments.

364 **3.2.** Monitoring of the parameters for the optimisation of biosurfactant

365 production in bioreactor

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Firstly, an open to air bioreactor with only mechanical stirring for oxygenation of the 367 solution showed no evidence of biosurfactant production. The lack of dissolved oxygen in the 368 water was a limiting parameter for the bacterial community growth, as shown below (Figure 369 4), where O_2 concentrations were close to 0 ppm during most of the time of the experiment. 370 Consumption of the O₂ by the bacteria in the bioreactor could be an explanation to these low 371 372 O_2 concentrations, with a too low input by natural dissolution of the O_2 present in the air. Two assays to inject directly O2 into the bioreactor did not allow the production of 373 biosurfactant with a surface tension remaining constant at ca 72 mN/m during the whole 374 375 experiment (Figure 6). Temperature – due to a thermal control –, pH and conductivity values 376 remained stable during the test and redox potential increased with the injection of O₂. The 377 lack of oxygen could have led to the non-consummation of the glucose, explaining the 378 stability of the physical parameters. These results seems to indicate that the bacteria 379 composing the studied community are aerobic bacteria.

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For next experiments, oxygen was supplied by direct air injection in the bioreactors, what generates a biosurfactant production after two days which was achieved and attested by collapsing drop test and measurement of the surface tension (Jain et al., 1991).

During the production of biosurfactant, the consumption of glucose by bacteria led to the production of carboxylic acids (probably puryvic acid and its conjugate base, pyruvate) (Schuit and Schuit, 2015), which decreased the pH of the solution (Figure 5). When pH was lower than 7, the production of biosurfactant was stopped and the foam on the bioreactor disappeared. Measurements of the surface tension and collapsing drop tests made at pH inferior to 6.5 proved that the biosurfactant was no longer active. The acidification of the medium could lead to the partial precipitation of the biosurfactant and then its disappearance at the interface between water and air, preventing its role in the decrease of the surface tension. It's also possible that the nature of the biosurfactant is affected by the pH as it can happen to the amphoteric surfactant and so causing the destabilization of the biosurfactant micelles (Johansson and Somasundaran, 2007).

As shown by Figure 6, surface tension stayed stable during the first day of experiment, during the decrease of the pH. When pH was adjusted to 7, surface tension values began to decrease, and foam production started again. Measurements of the surface tension between the bioreactor medium and air at different pH in the bulk, showed that when pH was lower than 7, the biosurfactant was no longer active and the surface tension was close to 72 mN/m, close to the reference value between pure water and air.

Other biosurfactants, like lipopeptides, can be produced at large pH range, from 4.5 to 10.5
(Makkar and Cameotra, 1997). However, at acidic pH, rhamnolipids and surfactin lower their
efficiency on emulsifying diesel (Whang et al., 2008). For other biosurfactants, a pH value
greater than 7 seems important for their production (Lotfabad et al., 2009; Zinjarde and
Pant, 2002) or stability (Mukherjee, 2007).

An explanation could be the change of the molecular aggregates morphologies of the
biosurfactant (Helvac et al., 2004) at acidic pH or the slowdown of the cells growth, leading
to a decrease in the production (Lotfabad et al., 2009).

To limit this phenomenon, some drops of a NaOH 2 M solution were regularly injected intothe bioreactor to keep the pH value higher than 7, explaining the curve shape (Figure 5).

Conductivity and temperature – due to a thermal control – remained stable during the test. O₂ concentration decreased quickly at the beginning of the experiment due to its consumption by bacteria. After the first drop, O₂ concentration increased and remained stable at a concentration of approximatively 9 ppm enabling enough oxygen for biosurfactant production. After an initial decrease, due to the O₂ concentration drop, redox potential increased during the experiment due to oxidation of the glucose by the bacteria. If a decrease in redox potential can be observed when 2 M NaOH was injected into the bioreactor, it
always stays positive, meaning that the producing medium is in oxidative conditions, which is
in line with the idea of oxidizing the PAHs by the bacteria (Haritash and Kaushik, 2009).

420

Four different temperatures of production were tested, 10, 15, 20 and 40 °C. No production 421 422 of biosurfactant occurred at 10 °C and 40 °C and the surface tension remained at a value around 72 mN/m during the whole experiment. Considering the 15 °C bioreactor, it took 423 424 more than 2 wk to observe a decrease in the surface tension and the decrease was lower than 425 the one in the 20 °C bioreactor. The bioreactors at 20°C showed a higher production of biosurfactant; these results illustrate the low-mesophilic nature of the bacterial strains able 426 427 to produce biosurfactant. Finally, the upscaling of the biosurfactant production was a success. 428 The continuous monitoring of the physical parameters allowed determining that the O₂ saturation of the medium, the pH of 7 and a temperature of 20 °C were the most important 429 operating parameters to succeed the biosurfactant production. This preference for a 430 temperature of 20 °C may be due to the conditions of extraction and culture of the bacterial 431 community. To maximize the number of extracted bacterial species, it was decided to make 432 an extraction medium at a temperature of 20 °C (Dobrić and Bååth, 2018). Next media of 433 culture were all carried out at 20 °C. These tests could lead to the selection of only mesophilic 434 bacteria, explaining that later 20 °C is the optimum temperature for the biosurfactant 435 production, which could be due to a highest bacterial growth. 436

Enough biosurfactant could be produced to go further by purifying and characterizing it. Inparallel, the characterization of the producing community was achieved.

439

- 440 3.3. Characterization of the biosurfactant
- 441

442 **3.3.1. Characterization of the biosurfactant properties**

From the 3 L of foam and medium extracted used for the purification assays of the biosurfactant, 80 mg of purified biosurfactant were obtained after lyophilisation of the extracted product.

446 The surface tension as a function of the biosurfactant concentration can be obtained by 447 measuring surface tension between different solutions of purified biosurfactant and air. This 448 is achieved in order to determine its critical micellar concentration (CMC).

Two different behaviours, represented by the two different lines in figure 7, were recorded with the increase of concentration. The surface tension decreased linearly with the increase in concentration, due to the increasing concentration of biosurfactant molecule at the surface of the drop medium. Then the surface tension remained stable with the increase of the concentration, due to the saturation of biosurfactant at the surface of the drop. The junction of the two lines is the point where the concentration of surfactant in solution is sufficient to start to produce naturally micelles of surfactant; it corresponds to the CMC.

The estimated CMC of the purified biosurfactant is 42 mg/L, for a maximum reduction of the
surface tension from 72 mN/m to 40 mN/m (Figure 7).

458 This value of CMC at pH 7 and 20 °C is in the same range of value compared to the CMC of other biosurfactants, like rhamnolipids - CMC = 10-400 mg/L - or surfactin - CMC = 20459 mg/L - (Lang and Wullbrandt, 1999; Liu et al., 2014). It is also lower than the CMC of 460 chemically produced surfactants, like sodium dodecyl sulfate (SDS) – CMC = 2307 mg/L –, 461 Octylphenol ethylene oxide condensate (marketed as Triton X-100[™]) – CMC = 150 mg/L –or 462 Polyoxyethylene lauryl ether (marketed as $Brij-35^{TM}$) – CMC = 108 mg/L –. These three 463 surfactants are commonly used in soil washing processes (Befkadu and Chen, 2018). This 464 465 quite low value of CMC could be advantageous in a soil washing process perspective, with micelle formation at lower concentrations, it will lead to a better efficiency for PAHs 466 desorption and a better control of the soil remediation method(Mao et al., 2015; Mulligan et 467 468 al., 2001). Moreover, another advantage is that the biosurfactant will not be regarded itself as 469 a pollutant and so it will not require recovery operations after the total removal of PAHs. 470 However, the precise monitoring of the PAHs concentration in the groundwater is necessary
471 until the end in order to control and promulgate actions to prevent the pollutant dispersion
472 in the groundwater due to the biosurfactant action.

473

474

3.3.2 PAHs solubilisation assays

The evolution of the concentration in water of each of the 4 PAHS was studied as a function 475 476 of biosurfactant concentration. For PHE, PYR and BaP, an increase in PAH concentration can be observed with the increase of the biosurfactant production, attesting the efficiency of the 477 biosurfactant to solubilize PAHs in water (Figure 8). NAPH concentration is decreasing 478 479 slightly with the increase of biosurfactant concentration. This phenomenon could be explained by the fact that the maximum solubility of naphthalene in water at 20 °C, which is 480 481 32 mg/L, was not reached. The method of PAH deposit, by solubilisation in acetone (Sawada 482 et al., 2004), injection in the dry batches and evaporation of the acetone, could have led to the partial evaporation of NAPH and then a final concentration below its solubility value. The 483 chemical formula of the produced biosurfactant could not be determined, therefore molar 484 485 mass, it was not possible to calculate the Molar Solubilisation Ratios (MSR) (Ansari et al., 2013). Then comparison with other bio-sourced and chemical surfactant of its ability to 486 solubilize PAHs was not possible. However, the solubilisation capacity of the biosurfactant 487 488 was evaluated thanks to the slope of the lines and presented as mg solubilized PAH per g biosurfactant in solution. The solubilisation ratios of PHE, PYR and BaP are respectively 489 490 0.214 mg/g, 0.1204 mg/g and 0.0068 mg/g.

These results show the ability of the produced biosurfactant to increase the solubilisation of PAHs. Concerning soil remediation, this ability is appealing and enables opportunity to combine the desorption effect of the biosurfactant with a biodegradation process. The analysis of the producing bacterial community could support this possibility by bringing to light the potential ability of the bacteria to degrade PAHs. 496

497 **3.4. Identification of the bacterial community**

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DNA extractions and analyses were performed on a sample from bioreactor in which biosurfactant production was attested and optimal. First, CE-SSCP fingerprint (Figure 9a) highlighted a relatively low biodiversity in the bacterial community, with five major peaks corresponding to five dominant bacterial species in the culture. These species could be determined by cloning and sequencing the 16S rRNA gene, and closer affiliations could be determined as listed in a phylogenetic tree (Figure 9b). Clones 1 and 5, corresponding to peaks 1 and 5 in the fingerprint profile, were found to be part of the *Pseudomonas* genus.

506

507 As Pseudomonas 16S rRNA genes are close one from each other, it was difficult here to clearly link clones 1 and 5 to a particular Pseudomonas specie, nevertheless some of the 508 species close from our clones are described as able to produce biosurfactants (Abouseoud et 509 510 al., 2008; Mellor et al., 2011) and to degrade PAHs, generally low molecular ones (Isaac et al., 511 2015). Some members of the Enterobacteriaceae family, especially Klebsiela grimonti who 512 was found to be close from clone 2, were described as able to produce biosurfactants (Jain et 513 al., 2013). The presence of these species in the consortium could then be interesting for bioremediation perspectives. 514

515 Clone 3 was found to be close to members of the *Microbacterium* genus, especially from 516 *Microbacterium maritypicum* and *Microbacterium oxydans*. *Microbacterium maritypicum* 517 is not known to produce biosurfactants, but was described as able to degrade BaP (Mansouri 518 et al., 2017). Combined with the ability of some strains in our culture to produce a 519 biosurfactant able to increase the solubilisation (and so the bioavailability) of PAHs, the 520 presence of bacteria able to degrade High Molecular Weight (HMW) PAHs in the consortium 521 is very interesting. Other members of *microbacterium* genus were described as able to degrade benzo[a]pyrene and pyrene (Qin et al., 2017). The presence of such strains in the bacterial community clearly let us think that its use in soil for PAHs remediation could be very interesting. The degradation of HMW PAHs such as BaP by some members of the community could also lead to the production of degradation by-products, easier to metabolize for other bacteria known to degrade Low Molecular Weight (LMW) PAHs, such as *pseudomonas* strains (clones 1 and 5) or Enterobacteriaceae family members (clone 2).

528 Members of the *Rhodanobacteraceae* family (close to clone 4) have also been described as 529 able to mineralize BaP (Kanaly et al., 2002), as shown by the production of ${}^{14}CO_2$ from ${}^{14}C$ 530 labelled BaP on a high-boiling-point diesel fuel distillate (HBD). BaP solubilisation into the 531 culture suspension was however an essential step for rapid mineralization. In the case of our 532 consortium, the solubilisation could occur thanks to biosurfactant producing bacteria, 533 making the degradation of BaP possible.

Finally, the community grown in our study seems to cumulate the ability of different strains 534 that could lead to an efficient PAHs biodegradation, with biosurfactant producing strains that 535 536 could increase PAHs bioavailability (Enterobacteriaceae, Pseudomonas), and PAHs degradation bacteria (Microbacterium, some Pseudomonas, Rhodanobacteraceae). The 537 more recalcitrant PAHs, such as benzo[a]pyrene, can be metabolized by some bacteria close 538 from members of our consortium, creating more easily degradable by products. LMW PAHs 539 such as NAPH or PHE can be degraded more easily by bacteria, including strains from our 540 consortium. Even if assays of remobilization and degradation in real matrices contaminated 541 with PAHs have to be performed to clearly conclude on the efficiency of the community for 542 543 bioremediation, all parameters seem to be very promising.

544

545 **4. Conclusions**

547 In this study, the extraction and isolation of a bacterial community able to produce biosurfactant from a contaminated soil was achieved. The nutritional and operating 548 conditions were optimized to maximize biosurfactant production. Four main factors were 549 identified for the biosurfactant production: (i) a C/N ratio higher than 10, with a maximum 550 551 production with C/N ratio = 13/1, (ii) the presence of phosphorus in the culture medium, with a maximum production with a C/N/P ratio = 13/1/2, (iii) a constant presence of 552 dissolved oxygen in the medium, and (iv) a pH superior to 7. Purification of the produced 553 biosurfactant allowed its characterization, with a CMC = 42 mg/L, and the determination of 554 555 the Molar Solubility Ratio for 4 PAHs with this biosurfactant, highlighting its capacity for PAHs remobilization. Determination of the bacterial community has been achieved and 556 showed the presence of bacteria (Enterobacteriaceae family, Microbacterium genus) (i) able 557 558 to produce biosurfactant, which were responsible for the production observed in our experiments, and (ii) bacteria able to metabolize both LMW and HMW PAHs. Despite the 559 proved effect of the studied biosurfactant in the PAHs desorption, due to the long and 560 expensive process of purification of biosurfactant, and the relatively small quantity obtained, 561 562 the use of purified biosurfactant in a real scale soil washing process seems difficult. However, the fast and relatively not expensive production of biosurfactant in batch bioreactor, and its 563 564 low CMC value make possible the use of non-purified biosurfactant rather than the purified product. When using raw bioreactor solution, the presence of biosurfactant combined with 565 bacteria, which are able to degrade both LMW and HMW PAHs, could lead to a combined 566 567 effect of desorption and biodegradation and so a reduction of the remediation cost, by only using easily available nutrients to both help the bacterial growth and the biosurfactant 568 production. Furthermore, the biosurfactant is certainly biodegradable and non-toxic, which 569 facilitates the post-treatment acceptance on completion of the works, with or without 570 obligation of monitoring. 571

572 Due to these benefits, it seems that the consortium used in our study, and its produced 573 biosurfactant, is a good candidate for bioremediation tests, and future studies will determine 574 its possible use for increasing PAHs bioavailability and biodegradation in contaminated soils.

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591 **References**

- Abouseoud, M., Maachi, R., Amrane, A., Boudergua, S., Nabi, A., 2008. Evaluation of different carbon and nitrogen sources
 in production of biosurfactant by Pseudomonas fluorescens. Desalination 223, 143–151.
- 595 https://doi.org/10.1016/j.desal.2007.01.198
- Ali Khan, A.H., Tanveer, S., Alia, S., Anees, M., Sultan, A., Iqbal, M., Yousaf, S., 2017. Role of nutrients in bacterial
- 597 biosurfactant production and effect of biosurfactant production on petroleum hydrocarbon biodegradation. Ecol.
- 598 Eng. 104, 158–164. https://doi.org/10.1016/j.ecoleng.2017.04.023
- 599 Ansari, W.H., Fatma, N., Panda, M., Kabir-Ud-Din, 2013. Solubilization of polycyclic aromatic hydrocarbons by novel
- 600 biodegradable cationic gemini surfactant ethane-1,2-diyl bis(N,N-dimethyl-N- hexadecylammoniumacetoxy)
- 601 dichloride and its binary mixtures with conventional surfactants. Soft Matter 9, 1478–1487.

602 https://doi.org/10.1039/c2sm26926k

- BEFKADU, A.A., CHEN, Q., 2018. Surfactant-Enhanced Soil Washing for Removal of Petroleum Hydrocarbons from
 Contaminated Soils: A Review. Pedosphere 28, 383–410. https://doi.org/10.1016/S1002-0160(18)60027-X
- Bois, P., Huguenot, D., Norini, M.P., Haque, M.F.U., Vuilleumier, S., Lebeau, T., 2011. Herbicide degradation and copper
- 606 complexation by bacterial mixed cultures from a vineyard stormwater basin. J. Soils Sediments 11, 860–873.
- 607 https://doi.org/10.1007/s11368-011-0354-3
- 608 Chebbi, A., Hentati, D., Zaghden, H., Baccar, N., Rezgui, F., Chalbi, M., Sayadi, S., Chamkha, M., 2017. Polycyclic aromatic
 609 hydrocarbon degradation and biosurfactant production by a newly isolated Pseudomonas sp. strain from used motor
 610 oil-contaminated soil. Int. Biodeterior. Biodegrad. 122, 128–140. https://doi.org/10.1016/j.ibiod.2017.05.006
- 611 Crampon, M., 2015. Influence des facteurs biogéochimiques et de l'ajout de biosurfactant sur la biodégradation des HAP
 612 dans des sols contaminés.
- 613 Davis, D.A., Lynch, H.C., Varley, J., 1999. The production of Surfactin in batch culture by Bacillus subtilis ATCC 21332 is
- 614 strongly influenced by the conditions of nitrogen metabolism. Enzyme Microb. Technol. 25, 322–329.
- 615 https://doi.org/10.1016/S0141-0229(99)00048-4
- 616 Deschênes, L., Lafrance, P., Villeneuve, J.P., Samson, R., 1995. The effect of an anionic surfactant on the mobilization and
- 617 biodegradation of PAHs in a creosote-contaminated soil. Hydrol. Sci. J. 40, 471–484.
- 618 https://doi.org/10.1080/02626669509491433
- 619 Dobrić, S., Bååth, E., 2018. Temperature affects lag period and growth of bacteria in soil according to a Ratkowsky (square
- 620 root) model after a drying/rewetting episode. Soil Biol. Biochem. https://doi.org/10.1016/j.soilbio.2018.05.023
- Gabet, S., 2004. Remobilisation d'Hydrocarbures Aromatiques Polycycliques (HAP) présents dans les sols contaminés à
 l'aide d'un tensioactif d'origine biologique. Thèse Dr. Univ. LIMOGES 177.
- Hall, T.A., 1999. BIOEDIT: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/
- 624 NT. Nucleic Acids Symp. Ser.
- Haritash, A.K., Kaushik, C.P., 2009. Biodegradation aspects of Polycyclic Aromatic Hydrocarbons (PAHs): A review. J. Hazard.
 Mater. 169, 1–15. https://doi.org/10.1016/j.jhazmat.2009.03.137
- 627 Hassanshahian, M., 2014. Isolation and characterization of biosurfactant producing bacteria from Persian Gulf (Bushehr
- 628 provenance). Mar. Pollut. Bull. 86, 361–366. https://doi.org/10.1016/j.marpolbul.2014.06.043

- 629 Helvac, S.S., Peker, S., Özdemir, G., 2004. Effect of electrolytes on the surface behavior of rhamnolipids R1 and R2. Colloids
- 630 Surfaces B Biointerfaces 35, 225–233. https://doi.org/10.1016/j.colsurfb.2004.01.001
- Henkel, M., Müller, M.M., Kügler, J.H., Lovaglio, R.B., Contiero, J., Syldatk, C., Hausmann, R., 2012. Rhamnolipids as
- 632 biosurfactants from renewable resources: Concepts for next-generation rhamnolipid production. Process Biochem.
- 633 47, 1207–1219. https://doi.org/10.1016/j.procbio.2012.04.018
- 634 Isaac, P., Lozada, M., Dionisi, H.M., Estévez, M.C., Ferrero, M.A., 2015. Differential expression of the catabolic nahAc gene
- 635 and its effect on PAH degradation in Pseudomonas strains isolated from contaminated Patagonian coasts. Int.
- 636 Biodeterior. Biodegrad. 105, 1–6. https://doi.org/10.1016/j.ibiod.2015.08.011
- 637 Jain, D.K., Collins-Thompson, D.L., Lee, H., Trevors, J.T., 1991. A drop-collapsing test for screening surfactant-producing
- 638 microorganisms. J. Microbiol. Methods 13, 271–279. https://doi.org/10.1016/0167-7012(91)90064-W
- 639 Jain, R.M., Mody, K., Joshi, N., Mishra, A., Jha, B., 2013. Production and structural characterization of biosurfactant
- 640 produced by an alkaliphilic bacterium, Klebsiella sp.: Evaluation of different carbon sources. Colloids Surfaces B
- 641 Biointerfaces 108, 199–204. https://doi.org/10.1016/j.colsurfb.2013.03.002
- Johansson, I., Somasundaran, P., 2007. Handbook for cleaning/decontamination of surfaces, Handbook for
 Cleaning/Decontamination of Surfaces.
- 644 Kanaly, R.A., Harayama, S., Watanabe, K., 2002. Rhodanobacter sp. strain BPC1 in a benzo[a]pyrene-mineralizing bacterial
- 645 consortium. Appl. Environ. Microbiol. 68, 5826–5833. https://doi.org/10.1128/AEM.68.12.5826-5833.2002
- 646 Karanth, V.R. &N. G., 1989. Factors Affecting Biosurfactant Production Using 249–257.
- Koch, A.K., Kappeli, O., Fiechter, A., Reiser, J., 1991. Hydrocarbon assimilation and biosurfactant production in
 Pseudomonas aeruginosa mutants. J. Bacteriol. 173, 4212–4219. https://doi.org/134212-08\$02.00/0
- 649 Lamichhane, S., Bal Krishna, K.C., Sarukkalige, R., 2017. Surfactant-enhanced remediation of polycyclic aromatic
- 650 hydrocarbons: A review. J. Environ. Manage. 199, 46–61. https://doi.org/10.1016/j.jenvman.2017.05.037
- 651 Lang, S., Wullbrandt, D., 1999. Rhamnose lipids--biosynthesis, microbial production and application potential. Appl.
- 652 Microbiol. Biotechnol. 51, 22–32. https://doi.org/10.1007/s002530051358
- Li, S., Pi, Y., Bao, M., Zhang, C., Zhao, D., Li, Y., Sun, P., Lu, J., 2015. Effect of rhamnolipid biosurfactant on solubilization of
- 654 polycyclic aromatic hydrocarbons. Mar. Pollut. Bull. 101, 219–225. https://doi.org/10.1016/j.marpolbul.2015.09.059
- Liang, X., Guo, C., Liao, C., Liu, S., Wick, L.Y., Peng, D., Yi, X., Lu, G., Yin, H., Lin, Z., Dang, Z., 2017. Drivers and applications of

- 656 integrated clean-up technologies for surfactant-enhanced remediation of environments contaminated with polycyclic
- aromatic hydrocarbons (PAHs). Environ. Pollut. 225, 129–140. https://doi.org/10.1016/j.envpol.2017.03.045
- Lima, T.M.S., Procópio, L.C., Brandão, F.D., Carvalho, A.M.X., Tótola, M.R., Borges, A.C., 2011. Biodegradability of bacterial
 surfactants. Biodegradation 22, 585–592. https://doi.org/10.1007/s10532-010-9431-3
- Liu, Q., Lin, J., Wang, W., Huang, H., Li, S., 2014. Production of surfactin isoforms by Bacillus subtilis BS-37 and its
- applicability to enhanced oil recovery under laboratory conditions. Biochem. Eng. J. 93, 31–37.
- 662 https://doi.org/10.1016/j.bej.2014.08.023
- 663 Lotfabad, T.B., Shourian, M., Roostaazad, R., Najafabadi, A.R., Adelzadeh, M.R., Noghabi, K.A., 2009. An efficient
- biosurfactant-producing bacterium Pseudomonas aeruginosa MR01, isolated from oil excavation areas in south of
- 665 Iran. Colloids Surfaces B Biointerfaces 69, 183–193. https://doi.org/10.1016/j.colsurfb.2008.11.018
- Lukić, B., Panico, A., Huguenot, D., Fabbricino, M., van Hullebusch, E.D., Esposito, G., 2016. Evaluation of PAH removal
- 667 efficiency in an artificial soil amended with different types of organic wastes. Euro-Mediterranean J. Environ. Integr.
- 668 1, 5. https://doi.org/10.1007/s41207-016-0001-x
- 669 Ma, J., Xiao, R., Li, J., Yu, J., Zhang, Y., Chen, L., 2010. Determination of 16 polycyclic aromatic hydrocarbons in
- 670 environmental water samples by solid-phase extraction using multi-walled carbon nanotubes as adsorbent coupled
- 671 with gas chromatography-mass spectrometry. J. Chromatogr. A. https://doi.org/10.1016/j.chroma.2010.06.060
- 672 Mahalingam, P.U., Sampath, N., 2014. Isolation, characterization and identification of bacterial biosurfactant. Eur. J. Exp.
- 673 Biol. 4, 59–64.
- Makkar, R.S., Cameotra, S.S., 1997. Biosurfactant production by a thermophilic Bacillus subtilis strain. J. Ind. Microbiol.
 Biotechnol. 18, 37–42. https://doi.org/10.1038/sj.jim.2900349
- 676 Mansouri, A., Abbes, C., Landoulsi, A., 2017. Combined intervention of static magnetic field and growth rate of
- 677 Microbacterium maritypicum CB7 for Benzo(a)Pyrene biodegradation. Microb. Pathog. 113, 40–44.
- 678 https://doi.org/10.1016/j.micpath.2017.10.008
- Mao, X., Jiang, R., Xiao, W., Yu, J., 2015. Use of surfactants for the remediation of contaminated soils: A review. J. Hazard.
 Mater. https://doi.org/10.1016/j.jhazmat.2014.12.009
- 681 Mcgrath, S.P., Shen, Z.G., Zhao, F.J., 1997. Heavy metal uptake and chemical changes in the rhizosphere. Plant Soil 188,
- 682 153–159. https://doi.org/10.1023/A:1004248123948
- 683 Mellor, G.E., Bentley, J.A., Dykes, G.A., 2011. Evidence for a role of biosurfactants produced by Pseudomonas fluorescens in

- 684 the spoilage of fresh aerobically stored chicken meat. Food Microbiol. 28, 1101–1104.
- 685 https://doi.org/10.1016/j.fm.2011.02.003
- Mubarak, M.Q.E., Hassan, A.R., Hamid, A.A., Khalil, S., Isa, M.H.M., 2015. A simple and effective isocratic HPLC method for
 fast identification and quantification of surfactin. Sains Malaysiana 44, 115–120.
- 688 Mukherjee, A.K., 2007. Potential application of cyclic lipopeptide biosurfactants produced by Bacillus subtilis strains in
- 689 laundry detergent formulations. Lett. Appl. Microbiol. 45, 330–335. https://doi.org/10.1111/j.1472-
- 690 765X.2007.02197.x
- Mulligan, C.N., 2005. Environmental applications for biosurfactants. Environ. Pollut. 133, 183–198.
 https://doi.org/10.1016/j.envpol.2004.06.009
- 693 Mulligan, C.N., Mahmourides, G., Gibbs, B.F., 1989. The influence of phosphate metabolism on biosurfactant production by
- 694 Pseudomonas aeruginosa. J. Biotechnol. 12, 199–210. https://doi.org/10.1016/0168-1656(89)90041-2
- 695 Mulligan, C.N., Yong, R.N., Gibbs, B.F., 2001. Surfactant-enhanced remediation of contaminated soil: A review. Eng. Geol.

696 60, 371–380. https://doi.org/10.1016/S0013-7952(00)00117-4

- 697 Ochsner, U.A., Hembach, T., Fiechter, A., 1996. Production of rhamnolipid biosurfactants. Adv. Biochem. Eng. Biotechnol.
 698 53, 89–118. https://doi.org/10.1007/BFb0102326
- Patowary, K., Patowary, R., Kalita, M.C., Deka, S., 2017. Characterization of biosurfactant produced during degradation of
- 700 hydrocarbons using crude oil as sole source of carbon. Front. Microbiol. 8, 1–14.
- 701 https://doi.org/10.3389/fmicb.2017.00279
- 702 Pereira, J.F.B., Gudiña, E.J., Costa, R., Vitorino, R., Teixeira, J.A., Coutinho, J.A.P., Rodrigues, L.R., 2013. Optimization and
- 703 characterization of biosurfactant production by Bacillus subtilis isolates towards microbial enhanced oil recovery

704 applications. Fuel 111, 259–268. https://doi.org/10.1016/j.fuel.2013.04.040

- 705 Portet-Koltalo, F., Ammami, M.T., Benamar, A., Wang, H., Le Derf, F., Duclairoir-Poc, C., 2013. Investigation of the release of
- 706 PAHs from artificially contaminated sediments using cyclolipopeptidic biosurfactants. J. Hazard. Mater. 261, 593–601.
- 707 https://doi.org/10.1016/j.jhazmat.2013.07.062
- 708 Qin, W., Zhu, Y., Fan, F., Wang, Y., Liu, X., Ding, A., Dou, J., 2017. Biodegradation of benzo(a)pyrene by Microbacterium sp.
- 709 strain under denitrification: Degradation pathway and effects of limiting electron acceptors or carbon source.
- 710 Biochem. Eng. J. 121, 131–138. https://doi.org/10.1016/j.bej.2017.02.001
- 711 Reis, R.S., da Rocha, S.L.G., Chapeaurouge, D.A., Domont, G.B., Santa Anna, L.M.M., Freire, D.M.G., Perales, J., 2010. Effects

- 712 of carbon and nitrogen sources on the proteome of Pseudomonas aeruginosa PA1 during rhamnolipid production.
- 713 Process Biochem. 45, 1504–1510. https://doi.org/10.1016/j.procbio.2010.05.032
- 714 Riding, M.J., Doick, K.J., Martin, F.L., Jones, K.C., Semple, K.T., 2013. Chemical measures of bioavailability/bioaccessibility of
- 715 PAHs in soil: Fundamentals to application. J. Hazard. Mater. 261, 687–700.
- 716 https://doi.org/10.1016/j.jhazmat.2013.03.033
- 717 Robert, M., Mercadé, M.E., Bosch, M.P., Parra, J.L., Espuny, M.J., Manresa, M.A., Guinea, J., 1989. Effect of the carbon
- 718 source on biosurfactant production by Pseudomonas aeruginosa 44T1. Biotechnol. Lett. 11, 871–874.
- 719 https://doi.org/10.1007/BF01026843
- 720 Rocha e Silva, N.M.P., Rufino, R.D., Luna, J.M., Santos, V.A., Sarubbo, L.A., 2014. Screening of Pseudomonas species for
- 721 biosurfactant production using low-cost substrates. Biocatal. Agric. Biotechnol. 3, 132–139.
- 722 https://doi.org/10.1016/j.bcab.2013.09.005
- 723 Rufino, R.D., de Luna, J.M., de Campos Takaki, G.M., Sarubbo, L.A., 2014. Characterization and properties of the
- biosurfactant produced by Candida lipolytica UCP 0988. Electron. J. Biotechnol. 17, 34–38.
- 725 https://doi.org/10.1016/j.ejbt.2013.12.006
- Sawada, A., Kanai, K., Fukushima, M., 2004. Preparation of Artificially Spiked Soil with Polycyclic Aromatic Hydrocarbons for
 Soil Pollution Analysis. Anal. Sci. 20, 239–241. https://doi.org/10.2116/analsci.20.239
- 728 Schuit, F.C., Schuit, F.C., 2015. Glycolyse, in: Leerboek Metabolisme. https://doi.org/10.1007/978-90-368-0620-6_3
- 729 Shao, B., Liu, Z., Zhong, H., Zeng, G., Liu, G., Yu, M., Liu, Y., Yang, X., Li, Z., Fang, Z., Zhang, J., Zhao, C., 2017. Effects of
- rhamnolipids on microorganism characteristics and applications in composting: A review. Microbiol. Res. 200, 33–44.
 https://doi.org/10.1016/j.micres.2017.04.005
- Soberón-Chávez, G., Lépine, F., Déziel, E., 2005. Production of rhamnolipids by Pseudomonas aeruginosa. Appl. Microbiol.
 Biotechnol. 68, 718–725. https://doi.org/10.1007/s00253-005-0150-3
- Souza, E.C., Vessoni-Penna, T.C., De Souza Oliveira, R.P., 2014. Biosurfactant-enhanced hydrocarbon bioremediation: An
 overview. Int. Biodeterior. Biodegrad. 89, 88–94. https://doi.org/10.1016/j.ibiod.2014.01.007
- 736 Varjani, S.J., Upasani, V.N., 2017a. Critical Review on Biosurfactant Analysis, Purification and Characterization Using
- 737 Rhamnolipid as A Model Biosurfactant. Bioresour. Technol. https://doi.org/10.1016/j.biortech.2017.02.047
- 738 Varjani, S.J., Upasani, V.N., 2017b. Critical Review on Biosurfactant Analysis, Purification and Characterization Using
- 739 Rhamnolipid as A Model Biosurfactant. Bioresour. Technol. 232, 389–397.

- 740 https://doi.org/10.1016/j.biortech.2017.02.047
- Wang, Q., Garrity, G.M., Tiedje, J.M., Cole, J.R., 2007. Naïve Bayesian classifier for rapid assignment of rRNA sequences into
 the new bacterial taxonomy. Appl. Environ. Microbiol. 73, 5261–5267. https://doi.org/10.1128/AEM.00062-07
- 743 Wei, Y.H., Lai, C.C., Chang, J.S., 2007. Using Taguchi experimental design methods to optimize trace element composition
- for enhanced surfactin production by Bacillus subtilis ATCC 21332. Process Biochem. 42, 40–45.
- 745 https://doi.org/10.1016/j.procbio.2006.07.025
- 746 Whang, L.M., Liu, P.W.G., Ma, C.C., Cheng, S.S., 2008. Application of biosurfactants, rhamnolipid, and surfactin, for
- 747 enhanced biodegradation of diesel-contaminated water and soil. J. Hazard. Mater. 151, 155–163.
- 748 https://doi.org/10.1016/j.jhazmat.2007.05.063
- 749 Zinjarde, S.S., Pant, A., 2002. Emulsifier from a tropical marine yeast, Yarrowia lipolytica NCIM 3589. J. Basic Microbiol. 42,
- 750 67-73. https://doi.org/10.1002/1521-4028(200203)42:1<67::AID-JOBM67>3.0.CO;2-M

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753 Supplementary materials

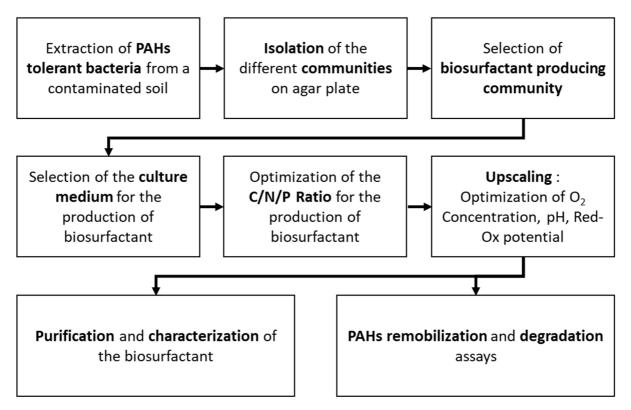


Figure 1: Experimental scheme showing all the successive steps from extraction of PAHs tolerant bacterial strains from contaminated soil to PAHs remobilization and degradation assays.

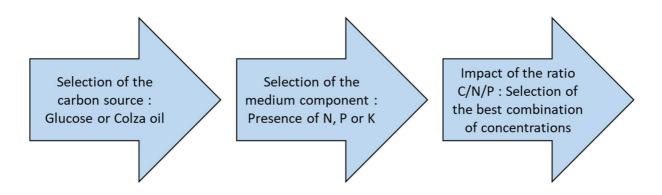


Figure 2: Experimental scheme showing the three successive steps from the selection of carbon source nutrients to the selection of the most optimized combination of concentrations in N, P and K as nutrients for the selected population of bacteria, tolerant to PAHs and producing biosurfactants.

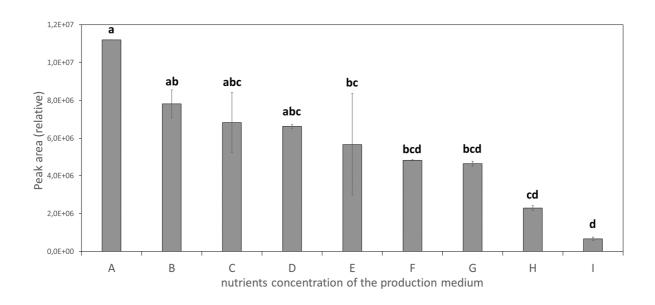


Figure 3: Relative biosurfactant concentration as shown by HPLC-FLD peak areas in the 9 media tested, where biosurfactant production was achieved, groups a to d represent significants differences between the conditions, analyzed by Tukey test with multiple pairwise comparisons from an ANOVA analysis of the results

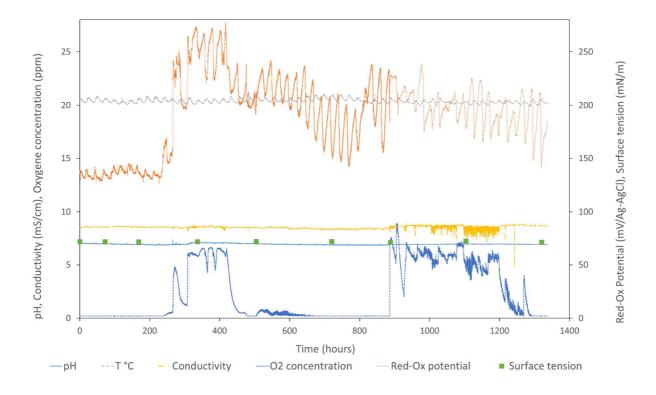


Figure 4: Monitoring of the main physical and chemical parameters of the solution (pH, redox potential, conductivity, dissolved oxygen concentration and surface tension between air and solution), on the first bioreactor

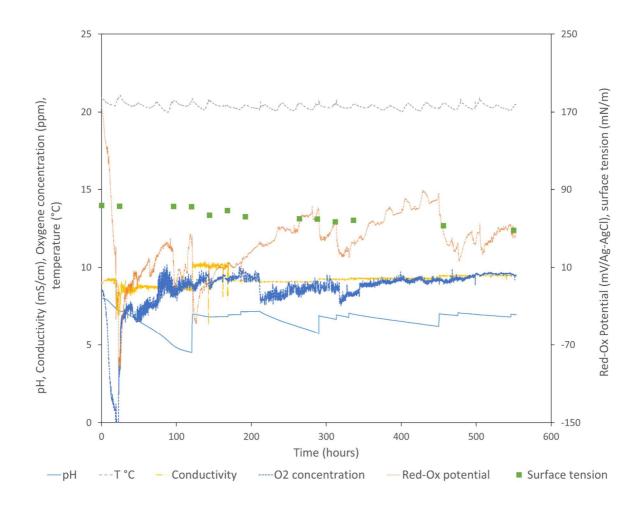


Figure 5: Monitoring of the main physical and chemical parameters of the solution (pH, redox potential, conductivity, dissolved oxygen concentration and surface tension between solution and air), on the second bioreactor.

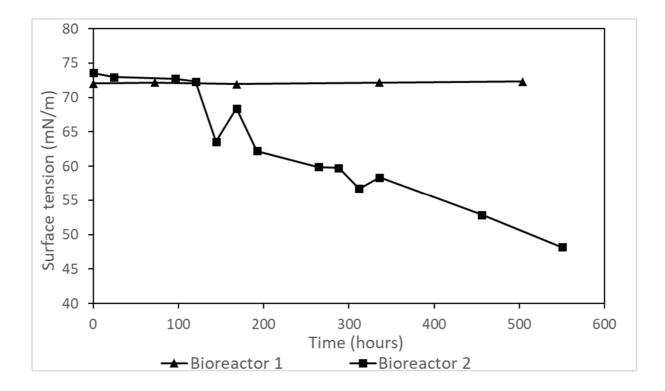


Figure 6: Evolution of the surface tension between solution and air in the solution of the two bioreactors

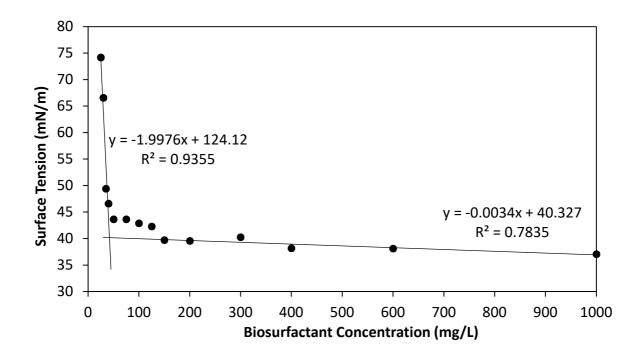


Figure 7: Surface tension between different purified biosurfactant solutions and air as a function of the purified biosurfactant concentration (in mg/L). The estimated CMC of the biosurfactant is equal to 42 mg/L, for a surface tension of 40 mN/m.

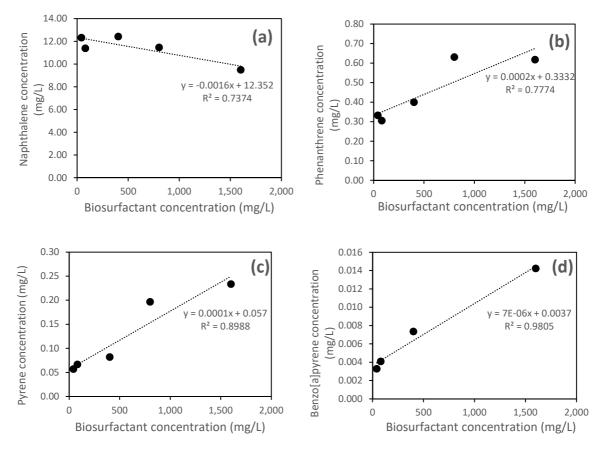


Figure 8: Molar solubilisation Ratio (MSR) in water of the four studied PAHs, (a) naphthalene, (b) phenanthrene, (c) pyrene, (d) benzo[a]pyrene, as a function of the biosurfactant concentration

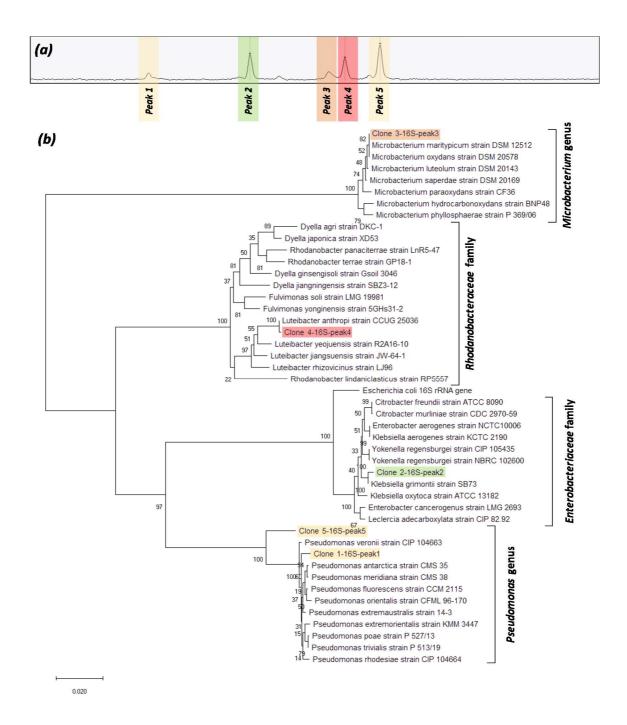


Figure 9 : (a) CE-SSCP fingerprint analysis showing the biodiversity of the bacterial community and (b) Evolutionary relationships of taxa inferred using the Neighbor-Joining method [1]. The optimal tree with the sum of branch length = 0.62696946 is shown. The percentage of replicate trees in which the associated taxa clustered together in the bootstrap test (1000 replicates) are shown next to the branches [2]. The evolutionary distances were computed using the Jukes-Cantor method [3]. The analysis involved 46 nucleotide sequences, with a total of 1341 positions in the final dataset. Evolutionary analyses were conducted in MEGA X [4].

Table 1: Composition of the different media tested for the biosurfactant production

Medium name	Α	В	С	D	Ε	F	G	Н	Ι
Glucose concentration (g/L)	20	20	40	20	40	20	40	40	20
NH4NO3 concentration (g/L)	2	4	2	2	4	4	4	2	1
Na2HPO4 concentration (g/L)	14.2	14.2	14.2	7.1	7.1	7.1	14.2	7.1	7.1
C/N/P ratio	13/1/2	13/2/2	26/1/2	13/1/1	26/2/1	13/2/1	13/1/1	26/1/1	26/1/2

Graphical abstract

