

Query

- Q1. Please confirm if the forename and surname of all the authors are presented appropriately in the author byline here.
- Q2. Please provide expansion for abbreviations “HPLC and MALDI-TOF” both in abstract and in the text.
- Q3. References “Shimoni et al. (1998), Todokoro et al. (2015), and Lee and Chen (1997)” are cited in the text but not provided in the list. Please check and provide the references in the list or delete the citation.
- Q4. As per journal specification, numbered reference style is followed. So the Name-and-Date reference citations “Meiwes et al. (1990), Nakouti and Hobbs (2014), Tóth et al. (2009), Emri et al. (2013), and Szigeti et al. (2014)” have been replaced with numbered reference citations to match with reference list. Please check and confirm.
- Q5. Please check whether “variance analysis” can be changed to “analysis of variance” in the sentence that begins with “The effects of the carbon...”
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- Q7. Figures 2 and 3 have been changed to Figures 1 and 2 and also the citations “Figures 2 and 1” have been changed to “Figures 1 and 2” in the text. Please check and amend if necessary.
- Q8. The value “ $715 \pm 34.99 \pm 2$ mg/l” seems incorrect in the sentence that begins with “In MOPS buffered media...” Please check and amend if necessary.
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- Q10. Please provide expansion for “DF” in Table IV.
- Q11. Please check the suggested placement of footnote “a” in Table IV, amend if necessary.
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OPTIMIZATION OF DESFERRIOXAMINE E PRODUCTION BY *STREPTOMYCES PARVULUS*

TAMÁS GÁLL^{1*}, GÁBOR LEHOCZKI², GYÖNGYI GYÉMÁNT², TAMÁS EMRI³,
ZSUZSA M. SZIGETI³, GYÖRGY BALLA^{1,4}, JÓZSEF BALLA⁵ and ISTVÁN PÓCSI³

¹MTA-DE Vascular Biology, Thrombosis and Hemostasis Research Group, Hungarian
Academy of Sciences, Debrecen, Hungary

²Department of Inorganic and Analytical Chemistry, Faculty of Science and Technology,
University of Debrecen, Debrecen, Hungary

³Department of Microbiology and Biotechnology, Faculty of Science and Technology,
University of Debrecen, Debrecen, Hungary

⁴Department of Pediatrics, Faculty of Medicine, University of Debrecen, Debrecen,
Hungary

⁵Department of Internal Medicine, Faculty of Medicine, University of Debrecen,
Debrecen, Hungary

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Siderophores are produced by a number of microbes to capture iron with outstandingly high affinity, which property also generates biomedical and industrial interests. Desferrioxamine E (DFO-E) secreted by streptomyces bacteria can be an ideal candidate for iron chelation therapy, which necessitates its cost-effective production for *in vitro* and animal studies. This study focused on the optimization of DFO-E production by *Streptomyces parvulus* CBS548.68. Different combinations of various carbon and nitrogen sources as well as the addition of 3-morpholinopropane-1-sulfonic acid (MOPS) markedly affected DFO-E yields, which were attributed, at least in part, to the higher biomass productions found in MOPS-supplemented cultures. In MOPS-supplemented glucose and sodium glutamate medium, DFO-E productions as high as $2,009 \pm 90$ mg/l of culture medium were reached. HPLC analysis demonstrated that a simple two-step purification process yielded DFO-E preparations with purities of ~97%. MALDI-TOF mass spectrometry analysis showed that purified DFO-E always contained traces of desferrioxamine D2.

Keywords: *Streptomyces parvulus*, desferrioxamine E, MOPS, Na-glutamate

*Corresponding author; E-mail: tamasgallmsc@gmail.com

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Introduction

33 Siderophores are iron chelators produced by many microorganisms to
34 sequester iron from the environment [1]. These compounds bind iron with
35 remarkably high affinity and, hence, they are pivotally important in supporting
36 the growth of microbes and are also required in the invasion of host organisms by
37 many pathogens [2]. Although iron is an abundant element on Earth, its bioavail-
38 ability is considerably limited in the environment by the low solubility of Fe^{3+}
39 ($\sim 10^{-24}$ mol/l) [3–5].

40 Some soil bacteria like the Gram-positive *Streptomyces* species produce
41 commercially important antibiotics and other metabolites like iron-chelator side-
42 rophores, e.g., desferrioxamine B (DFO-B), a product of *Streptomyces pilosus*,
43 which is employed to treat iron overload in humans [6]. Iron chelators are
44 routinely used for the medication of the patients with iron overloads like in the
45 case of β -thalassemia or sickle cell disease [7, 8]. Several studies have demon-
46 strated that DFO-B is able to mitigate oxidative cell damages in various biological
47 systems through disrupting iron-mediated redox cycling reactions [9–11]. Hartley
48 et al. [11] also showed that DFO-B itself can act as a radical scavenger by donating
49 an electron or hydrogen atom from the hydroxamate center. Therefore, side-
50 rophores of both bacterial and fungal origins are considered in the prevention and
51 treatment of oxidative-stress-mediated diseases, e.g., ischemia–reperfusion injury
52 diabetes, inflammation, and atherosclerosis [12–16]. Not surprisingly, there is also
53 an increasing body of evidence that these siderophores are able to target cancer
54 cells selectively [17, 18].

55 In a recent study published by Todokoro et al. (2015), a promising
56 new application was described for the natural hexadentate iron chelator
57 deferriferriochrysin, fermented by the rice-saccharifying fungus *Aspergillus*
Q3 58 *oryzae* as a food-grade antioxidant [19]. Furthermore, Shimoni et al. (1998)
59 demonstrated that the hexadentate cyclic siderophore desferrioxamine E (DFO-E,
60 also produced by various streptomycetes) possessed significant antioxidant
61 effects in linoleic acid and also in combined linoleic acid– β -carotene
62 emulsions.

63 These results suggest that various siderophores of microbial origin, includ-
64 ing DFO-E, could be candidate drugs to medicate oxidative-stress-or iron-excess-
65 mediated diseases in humans. Because today's knowledge on the optimization of
66 microbial siderophore productions is still rather uneven, further studies are
67 definitely needed to introduce new fermentation processes for the more cost-
68 effective production of microbial siderophores like DFO-E, e.g., for *in vitro* and
Q4 69 animal studies [20, 21].

Materials and Methods

Chemicals

D-Glucose monohydrate, agar, K_2HPO_4 , NaCl, methanol, chloroform, acetonitrile (HiPerSolv CHROMANORM, isocratic grade for HPLC), and glycerol were purchased from VWR BDH Prolabo (VWR, Radnor, PA, USA); yeast extract and $FeCl_3 \cdot 6H_2O$ from Alfa Aesar (Ward Hill, MA, USA); L-aspartic acid (L-Asp), Amberlite XAD-2, Na-L-glutamate (monosodium salt monohydrate), 3-morpholinopropane-1-sulfonic acid (MOPS), and $ZnSO_4 \cdot 7H_2O$ from Sigma-Aldrich (St. Louis, MO, USA); anhydrous $MgSO_4$, Silica Gel 60, and soluble starch from Merck (Darmstadt, Germany); $CaCO_3$ and K_2SO_4 from Reanal (Budapest, Hungary); and malt extract from Oxoid (Basingstoke, UK), while anhydrous $CaCl_2$ was bought from Scharlau (Barcelona, Spain).

Production and harvesting of spores

S. parvulus CBS 548.68 was cultured on glucose–yeast extract–malt extract (GYM) *Streptomyces* medium (pH 7.2), containing 4.0 g/l glucose, 4 g/l yeast extract, 10 g/l malt extract, 2.0 g/l $CaCO_3$, and 12 g/l agar. After incubation at 28 °C for 9 days, spores were harvested by agitation in sterile 0.9% NaCl solution supplemented with 0.1% Tween-80. Spore suspensions were passed through sterile cotton wool and centrifuged at $12,000 \times g$ for 2 min at room temperature, washed with distilled water, centrifuged again, and stored in 20% glycerol at –20 °C. Colony-forming units (CFU) were determined by streaking a limited dilution of spores on GYM *Streptomyces* medium and incubated at 28 °C for 5 days.

Siderophore production

All glasswares were rinsed with 36% HCl and were washed subsequently with distilled water to remove any residual iron. Siderophore production was carried out in glass Erlenmeyer flasks (250 ml) containing 50 ml aliquots of production medium, which were inoculated with 1.9×10^8 CFU spores from glycerol stock, and were incubated at 28 °C and at 4.2-Hz shaking frequency. The basic siderophore production medium (pH 7.5) contained 5 g/l L-Asp or Na-L-glutamate monosodium salt monohydrate (Na-L-Glu), 5 g/l L-lysine monohydrochloride (L-Lys), 2 g/l K_2HPO_4 , 2 g/l K_2SO_4 , 1 g/l NaCl, 0.1 g/l anhydrous $CaCl_2$, 0.2 g/l anhydrous $MgSO_4$, 0.1 g/l $ZnSO_4 \cdot 7H_2O$, and 2.5%

102 carbon source (glucose, glycerol, or soluble starch). The buffered siderophore
103 production medium contained the same ingredients as the basic medium, but it was
104 also supplemented with 21 g/l MOPS and autoclaved for 15 min at 121 °C. Carbon
105 sources were autoclaved separately as 25% stock solutions. In order to determine
106 the siderophore productions by different cultures, 1.0 ml of production medium
107 samples were taken at 72-, 96-, and 120-h incubation times after inoculation.
108 Samples were centrifuged at $14,000 \times g$ for 5 min at room temperature, and then,
109 supernatants from each flasks (500 μ l) were transferred to 1.5-ml Eppendorf tubes
110 and supplemented with 20 μ l of 10% $\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$ to saturate all produced
111 siderophores with iron before HPLC analysis. The effects of the carbon source,
Q5 112 nitrogen source, and MOPS were evaluated with variance analysis (ANOVA), and
113 Tukey's difference test using the R 2.3.1 software (The R Development Core
114 Team; <http://www.r-project.org/>) was adapted to describe the response surface
115 (Lee and Chen 1997).

116 *HPLC analysis*

117 HPLC measurements were carried out with a HP 1090 liquid chromato-
118 graph, equipped with autosampler and diode array detector. Ferrioxamines were
119 separated on a mixed-phase column (Luknova Hyper SCX 4.6 mm \times 100 mm,
120 5 μ m), in which both the cation exchange and hydrophobic interactions contrib-
121 ute to the separation. The 10- μ l injected sample was separated by isocratic elution
122 with 1:4 acetonitrile: 0.5 M NaCl solvent mixture at 0.5 ml/min flow rate.
123 Siderophores were detected at $\lambda = 435$ nm, and Agilent ChemStation was used to
124 evaluate the chromatograms. Commercially available, ferrated DFO-E was used
125 as standard.

126 *Purification of siderophores with column chromatography*

127 Culture media of *S. parvulus* ($V = 1,000$ ml, incubation time = 96 h) were
128 centrifuged at $4,000 \times g$ for 20 min at 4 °C, and the supernatants were
129 supplemented with 2 g/l of $\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$ to reach iron complexation of all side-
130 rophores. Ferrated siderophores were isolated first using Amberlite XAD2 resin
131 and were purified further with Silica Gel 60 chromatography with a chloroform:
132 methanol:water (17.5:6:1) solvent mixture as the mobile phase. Organic solvents
133 were evaporated with a Buchi Rotavapor R-3000 equipment. Following that, the
134 ferrated DFO-E was dissolved in distilled water and analyzed with HPLC, and the
135 two fractions detected with HPLC were analyzed with MALDI-TOF mass
136 spectrometry (MS).

137 *MALDI-TOF MS*

138 HPLC fractions of the purified siderophores were analyzed with MALDI-
139 TOF MS in positive-ion mode using a Bruker Biflex MALDI-TOF mass
140 spectrometer equipped with delayed-ion extraction. Desorption/ionization of
141 the sample molecules was performed with a $\lambda = 337$ nm nitrogen laser, and
142 2,5-dihydroxy benzoic acid was used as matrix. Spectra from multiple (at least
143 100) laser shots were summarized using 19-kV accelerating and 20-kV
144 reflectron voltage. External calibration was applied using the $[M + Na]^+$
Q6 145 peaks of cyclodextrins DP 6–8 with m/z : 995 Da, m/z : 1,157 Da, and m/z :
146 1,319 Da, respectively.

147 *Response surface methodology (RSM)*

148 The optimal medium composition for DFO-E production was determined
149 using the RSM with two selected variables (Na-L-Glu and L-Lys concentra-
150 tions) with the central composite design and full-factorial design consisting of
151 two-factor-two-level pattern with 11 design points (nine combinations with
152 three replications of the center points). In accordance with our preliminary
153 experiments, DFO-E yields depended highly on L-Lys and Na-L-Glu concen-
154 trations. Two plausible variables, glucose and MOPS concentrations, were
155 omitted from the analysis since neither increasing glucose concentration from
156 25 g/l to 50 g/l nor decreasing to 10 g/l, or decreasing MOPS concentration
157 from 21 g/l to 15 g/l or 10 g/l did not affect the DFO-E yields considerably,
158 while decreasing either L-Lys or Na-L-Glu from 5 g/l to 2.5 g/l reduced DFO-E
159 yields by 20%–40% (data not shown). A full second-order polynomial model
160 was obtained by a multiple regression technique for two factors using the R
161 2.3.1 software.

162 **Results**163 *Identification of DFO-E produced by S. parvulus*

164 When culture media supplemented with Fe(III) were subjected to HPLC
165 analysis, four clearly distinguishable peaks were identified in each fermentation
166 medium. Two of them, observed at ~3.5-min and ~4.4-min retention times, had
167 the same retention values as the DFO-B and DFO-E standards, respectively (data
168 not shown).

169 HPLC and MALDI-TOF MS analysis of the purified DFO-E

170 The purity of DFO-E prepared from the culture supernatant of *S. parvulus*
171 after Amberlite XAD and Silica Gel purification was checked with HPLC, and
172 approximately 97% of the siderophore content was DFO-E (~4.5-min retention
173 time, Peak 2 in Figure 1A). A minor siderophore peak always appearing at 4.1-min
174 retention time in all preparations (Peak 1 in Figure 1A) was identical to
175 desferrioxamine D2 (DFO-D2). The chemical structures of DFO-E and DFO-
176 D2 were verified by MALDI-TOF MS analyses (Figure 1B and C).

177 Influence of MOPS as well as various carbon and nitrogen sources on biomass 178 and siderophore yields

179 In unbuffered culture media, DFO-E production reached its maximum at 96-h
180 incubation time after inoculation, and it remained constant or even slightly
181 decreased following that (Figure 2). The highest DFO-E production was detected
182 in the glucose + Na-L-Glu medium followed by glucose + L-Asp and glycerol +
183 Na-L-Glu media, while the glycerol + L-Asp medium showed slightly decreased
184 yields. Starch was not a favored carbon source either in L-Asp- (96 ± 3 mg/l) or in
185 Na-L-Glu- (99 ± 2 mg/l) supplemented media, which may indicate that the positive
186 effect of Na-L-Glu is dependent on the nature of carbon source (Figure 2). On the
187 other hand, MOPS supplementation markedly increased DFO-E yields in the
188 presence of all carbon and nitrogen sources and their combinations tested
189 (Figure 2).

190 In MOPS-buffered media, DFO-E production reached its peak at 120 h after
191 inoculation, and the most beneficial effects of MOPS on DFO-E yields were
192 recorded in the presence of starch, where 7- to 9-fold increases in DFO-E
193 productions were recorded when compared with unbuffered systems (from 96
Q8 194 ± 3 mg/l to 882 ± 179 mg/l in starch + L-Asp and from 99 ± 2 mg/l to 715 ± 34 99
195 ± 2 mg/l in starch + Na-L-Glu media). The maximum DFO-E yield ($1,390 \pm$
196 65 mg/l) was detected in MOPS-buffered glucose + Na-L-Glu medium (Figure 2).

197 The final pH of the fermentation media for all tested combinations (includ-
198 ing media with MOPS) was slightly acidic (pH 5.3–6.2) except starch + L-Asp
199 (pH 8.1) and starch + Na-L-Glu (pH 7.1). Dry cell masses correlated well with
200 yields with the exception of unbuffered starch + L-Asp and starch + Na-L-Glu
201 media (Figure 2 and Table I). Surprisingly, the final biomass gains found in
202 MOPS-buffered starch + L-Asp and MOPS-buffered glucose-Na-L-Glu media
203 were comparable (7.9 g/l and 8.9 g/l, respectively), but the DFO-E yields recorded
204 in these systems differed considerably. In MOPS-buffered Na-L-Glu supplemented

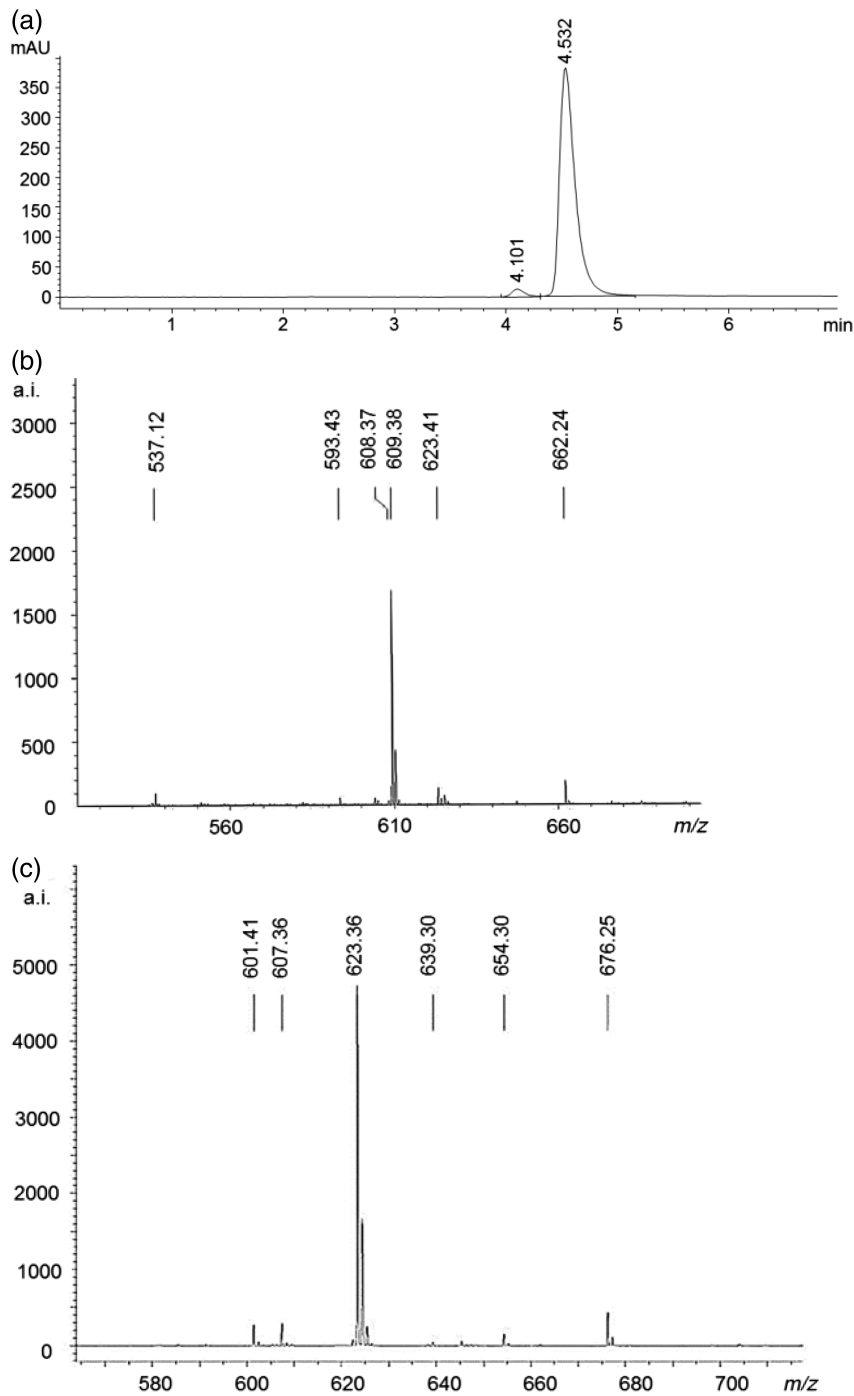


Figure 1. HPLC and MALDI-TOF MS analysis of the purified DFO-E

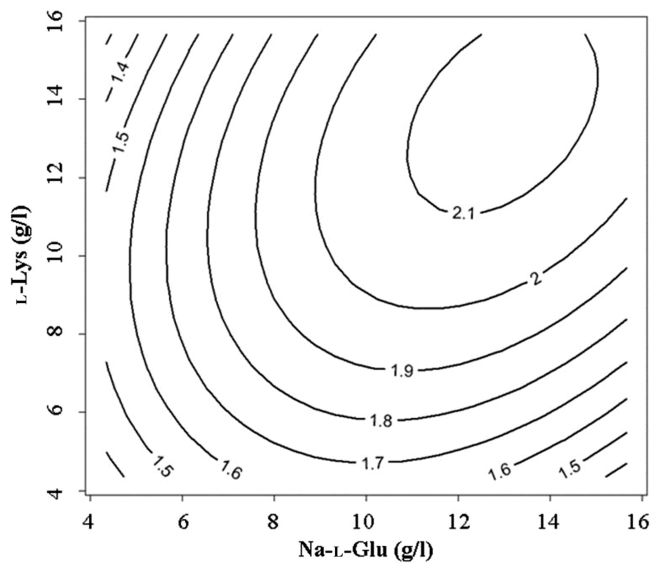


Figure 2. Fitted regression surface showing Na-L-Glu and L-Lys dependence of DFO-E production

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Table I. Effect of MOPS, carbon source, and nitrogen source on dry cell mass. MOPS supplementation markedly increased dry cell mass compared with the unbuffered media

| DFO-E production medium | Dry cell mass at 72-h incubation (g/l) | Dry cell mass at 96-h incubation (g/l) | Dry cell mass at 120-h incubation (g/l) |
|---------------------------------|--|--|---|
| Glucose + aspartic acid | 2.2 ± 0.2 | 2.3 ± 0.2 | 2.4 ± 0.2 |
| Glucose + aspartic acid + MOPS | 4.2 ± 0.1 | 4.7 ± 0.2 | 4.7 ± 0.2 |
| Glycerol + aspartic acid | 2.6 ± 0.1 | 2.6 ± 0.1 | 2.7 ± 0.1 |
| Glycerol + aspartic acid + MOPS | 5.4 ± 0.5 | 6.2 ± 0.6 | 6.4 ± 0.4 |
| Starch + aspartic acid | 2.2 ± 0.1 | 2.8 ± 0.1 | 2.8 ± 0.1 |
| Starch + aspartic acid + MOPS | 4.7 ± 0.2 | 6.3 ± 0.5 | 7.9 ± 0.3 |
| Glucose + glutamic acid | 5.2 ± 0.1 | 5.3 ± 0.1 | 5.5 ± 0.1 |
| Glucose + glutamic acid + MOPS | 5.5 ± 0.3 | 8.1 ± 0.2 | 8.9 ± 0.1 |
| Glycerol + glutamic acid | 3.5 ± 0.3 | 3.5 ± 0.2 | 3.7 ± 0.2 |
| Glycerol + glutamic acid + MOPS | 5.8 ± 0.5 | 8.1 ± 0.2 | 8.4 ± 0.4 |
| Starch + glutamic acid | 2.3 ± 0.1 | 2.3 ± 0.1 | 2.4 ± 0.1 |
| Starch + glutamic acid + MOPS | 6.0 ± 0.3 | 6.2 ± 0.3 | 7.9 ± 0.2 |

Note: Results are shown as mean values of three independent experiments with standard deviations.

Table II. ANOVA of DFO-E production data

| Variable | Difference in DFO-E production (mg/l) | Adjusted <i>p</i> values |
|----------------------------------|---------------------------------------|--------------------------|
| Effect of carbon source | | |
| Glycerol–glucose | –175.75 | <0.000001 |
| Starch–glucose | –372.99 | <0.000001 |
| Starch–glycerol | –197.24 | <0.000001 |
| Effect of nitrogen source | | |
| Glutamic acid–aspartic acid | 202.18 | <0.000001 |
| Effect of MOPS | | |
| Without MOPS–with MOPS | –441.32 | <0.000001 |
| Effect of sample taking | | |
| 72–120 h | –178.13 | <0.000001 |
| 96–120 h | –35.49 | 0.26 |
| 96–72 h | 142.64 | <0.000001 |

Note: According to ANOVA, the effects of carbon source, nitrogen source, and the presence of MOPS were significant. There was no significant difference between 96-h and 120-h sample taking, but at 72 h, the DFO-E concentrations were significantly lower than at 120 h or at 96 h.

media, all the glucose, glycerol, and starch carbon sources resulted in high growth rates, but siderophore productions varied again within a relatively wide range (Figure 2 and Table I).

According to ANOVA, the effects of carbon sources, nitrogen sources, and the presence of MOPS on DFO-E productions were significant (Table II). On the other hand, there was no significant difference between taking samples at 96-h or 120-h incubation times, but DFO-E concentrations recorded at 72-h incubation time were significantly lower than those found at 120-h or 96-h incubations. According to the Tukey's difference test, the DFO-E contents of media containing glucose, glutamic acid, and MOPS incubated for 120 h were significantly higher (adjusted $p < 0.05$) than those found in other cultures at any incubation times tested, except the glucose, glutamic acid, and MOPS containing cultures when incubated for 96 h (Supplementary Material).

Optimization of Na-L-Glu and L-Lys concentrations in DFO-E production using RSM

To describe the optimal medium for DFO-E production with *S. parvulus*, the response surface was studied as a function of Na-L-Glu and L-Lys concentrations as described earlier [22]. Glucose and MOPS levels in the fermentation media were set to the concentrations (25 g/l for glucose and 21 g/l for MOPS) used before for the optimization of the carbon and nitrogen sources. The levels for initial Na-L-Glu and L-Asp concentrations as well as experimental data for the

Table III. Experimental data of the central composite and full-factorial design

| Trial | Na-L-Glu (g/l) | L-Lys (g/l) | Coded values ^a | | DFO-E (g/l) |
|-------|----------------|-------------|---------------------------|--------|-------------|
| | X_1 | X_2 | Z_1 | Z_2 | Y |
| 1 | 6 | 6 | −1 | −1 | 1.7 |
| 2 | 6 | 14 | −1 | 1 | 1.6 |
| 3 | 14 | 6 | 1 | −1 | 1.7 |
| 4 | 14 | 14 | 1 | 1 | 2.0 |
| 5 | 15.66 | 10 | 1.414 | 0 | 2.0 |
| 6 | 4.34 | 10 | −1.414 | 0 | 1.5 |
| 7 | 10 | 15.66 | 0 | 1.414 | 2.1 |
| 8 | 10 | 4.34 | 0 | −1.414 | 1.6 |
| 9 | 10 | 10 | 0 | 0 | 2.1 |
| 10 | 10 | 10 | 0 | 0 | 2.0 |
| 11 | 10 | 10 | 0 | 0 | 2.0 |

Note: The glucose content of the medium was set to 25 g/l, while MOPS to 21 g/l, and DFO-E production was quantified after 120 h of cultivation.

^a $Z_i = (X_i - 10)/4$.

226 two-factor-two-level response surface analysis are summarized in Table III. Three
227 repeats were included at the center of the design, and the total number of the test
228 runs was 11. Regression coefficients, their *t* values, and determination coefficient
229 (R^2) for the full second-order polynomial model of DFO-E production are
230 presented in Table IV. The regression model for DFO-E production was signifi-
231 cant ($p = 0.01912$) with a satisfactory value of R^2 (0.89). The estimated DFO-E
232 production at optimal Na-L-Glu (12.96 g/l) and L-Lys (13.60 g/l) concentrations
233 was 2,130 mg/l, which corresponded well with the measured $2,009 \pm 90$ mg/l
234 DFO-E production ($n = 3$) at 13 g/l Na-L-Glu and 14 g/l L-Lys concentrations.

Table IV. Regression of coefficients, *p* values, determination coefficient (R^2), and *F*-statistic of a full second-order polynomial model for DFO-E production

| Term ^a | Coefficient | <i>p</i> value |
|-------------------|-------------|------------------------|
| b0 | 2.033 | 4.524×10^{-7} |
| b1 | 0.138 | 0.01383 |
| b2 | 0.113 | 0.02871 |
| b12 | 0.100 | 0.11637 |
| b11 | −0.154 | 0.01777 |
| b22 | −0.104 | 0.06578 |

$R^2 = 0.89$
F-statistic: 8.124 (DF 5 and 5), *p* value = 0.01912

^a $Y = b_0 + b_1Z_1 + b_2Z_2 + b_{11}Z_{12} + b_{12}Z_1Z_2 + b_{22}Z_{22}$, where *Y* is DFO-E (g/l) production, and Z_1 and Z_2 are Na-L-Glu and L-Lys content of the media (given in coded values), respectively.

Discussion

There is a plethora of evidence that the types of carbon sources and nitrogen sources as well as other culture media supplements (salts and precursors) greatly influence secondary metabolite biosyntheses, e.g., siderophore [20, 23–25]. In good accordance with this, different combinations of carbon sources, nitrogen sources, and MOPS resulted in various DFO-E yields during the production of this hexadentate cyclic siderophore by *S. parvulus*.

In unbuffered systems, i.e., when MOPS was not supplemented, the gains in biomasses recorded in starch + L-Asp and starch + Na-L-Glu media were similar to those found in glucose + L-Asp and glycerol + L-Asp media, but DFO-E yields were considerably higher in the latter cases (Table I). The lower DFO-E yields detected in unbuffered starch systems may be explained by the neutral or even slightly alkaline pH of the fermentation media (pH 7.1 for the starch + Na-L-Glu and pH 8.1 for the starch + L-Asp media). It is worth noting that even in the presence of MOPS, the final pH values measured in most tested culture media were slightly acidic (pH 5.3–6.2), and it is well established that hydroxamate-type siderophores are most stable under acidic conditions [26]. Szigeti et al. [27] also found that triacetylfusarinin C production by the opportunistic human pathogenic fungus *Aspergillus fumigatus* also favored acidic conditions. Because culture media underwent acidification even in the presence of MOPS and DFO-E yields were not affected by decreasing MOPS concentration from 21 g/l to 10 g/l, we can conclude that the positive effects of MOPS on DFO-E yields cannot be attributed solely to its buffering effects in the culture media.

The addition of MOPS is likely to facilitate DFO-E productions through supporting higher biomass productions (Table I) because dry cell masses correlated well with DFO-E yields when siderophore productions in MOPS-buffered and not buffered media were compared (Figure 2 and Table I). On the other hand, the improved yields could not be explained exclusively with the increased biomass gains because various DFO-E productions were observed in MOPS-buffered Na-L-Glu media supplemented with different carbon sources (glucose, glycerol, and starch), meanwhile the biomasses measured at 120-h incubation times were quite similar (Table I). Hence, further experiments are needed to shed light on the mechanism(s) through which MOPS stimulates DFO-E production in *S. parvulus*.

Considering the beneficial effects of the nitrogen sources on siderophore yields, the positive effect of L-Lys on DFO production is well known because L-Lys, the precursor of DFOs, greatly enhances their productions [20, 21]. As shown above, Na-L-Glu also stimulated the biomass production and the biosynthesis of

273 DFO-E by *S. parvulus* in both MOPS-buffered and unbuffered systems with the
274 exception of media containing starch as carbon source (Figure 2 and Table I). In the
275 absence of MOPS, starch-containing media supplemented with either L-Lys or Na-L-
276 Glu as nitrogen sources provided us with comparable biomass and DFO-E yields
277 (Figure 2 and Table I). These data indicated that both siderophore yields and biomass
278 gains were influenced by the carbon source as well, and the beneficial effect of Na-L-
279 Glu was carbon source dependent. Similar phenomena were also observed when
280 MOPS was also added to the culture media (Figure 2).

281 After optimizing the Na-L-Glu (14 g/l) and L-Lys (13 g/l) concentrations in
282 glucose (25 g/l) and MOPS (21 g/l) containing fermentation medium, *S. parvulus*
283 CBS548.69 provided us with DFO-E in considerable yields (~2 g/l). Following
284 that high purity grade (~97%), DFO-E preparations were obtained in a simple
285 two-step purification scheme relying on the capture of DFO-E by XAD resin and
286 silica gel chromatography (Figure 1A–C). MALDI-TOF MS demonstrated that
287 purified DFO-E always contained traces of DFO-D2 (Peak 1 in Figure 1A).

288 There are plenty of experimental data available on the optimization of the
289 production of hydroxamate-type hexadentate siderophores of microbial origin,
290 including compounds produced by either fungi (coprogen, ferrichrysin, and
291 triacetylfusarinin C) [22, 27, 29] or bacteria [23, 30]. Considering yields,
292 maximum coprogen productions by the filamentous fungus *Neurospora crassa*
293 varied between 200 mg/l and 250 mg/l, meanwhile triacetylfusarinin C production
294 by *A. fumigatus* reached as high as 2.9 g/l [22, 27, 29]. Todokoro et al. [19]
295 demonstrated that ferrichrysin production by *A. oryzae* can even reach 2.8 g/l.
Q12 296 Furthermore, Mortazavi and Akbarzadeh reported on almost 2 g/l DFO-B yields
297 with *S. pilosus* ATCC 19797, and Chiani et al. [25] published a 5 g/l DFO-B
298 production with the same streptomycete by adding Na₂HPO₄·12H₂O, NaH₂PO₄,
299 MgSO₄·7H₂O, ZnSO₄·7H₂O, FeSO₄·7H₂O, CaCl₂·2H₂O, NaCl, MnSO₄, NH₄Cl,
300 KH₂PO₄, and K₂HPO₄ to the medium.

301 Not surprisingly, although production data for DFO-B [25, 28, 31] are
302 abundant because this siderophore is widely used for the treatment of iron
303 overload in humans, similar data for the optimization of DFO-E production are
304 scarcely available [20, 21]. Nevertheless, the highest DFO-E yield was reported by
305 Meiwes et al. [21], which reached 12 g/l in a fed-batch fermentation system with
306 *Streptomyces olivaceus* Tü 2718. Although this yield is six times higher than the
307 highest DFO-E production by *S. parvulus* recorded in this study, there are
308 arguments for considering *S. parvulus*, when future DFO-E production technolo-
309 gies will be developed, at least at laboratory scale. For example, the *S. parvulus*
310 CBS 548.68 strain used in this study is commercially available and the shaking
311 flask method reported here may be scaled up simply in bench-top bioreactors [22].
312 Importantly, the favorable effects of MOPS on DFO-E production can be tested

313 and, hopefully, also exploited in any other chemically defined fermentation media
314 currently applied in siderophore productions [22, 27, 29].

315 We hope that our results will enable us to carry out further experiments to
316 reveal the complexometric properties, cytotoxicity, and physiological/pharmaco-
317 logical characteristics of this DFO-E. Large-scale production of various side-
318 rophores would make many potential applications of these iron chelators possible
319 in the future, e.g., in managing human diseases like atherosclerosis, imaging
320 fungal infections, or preparing siderophore–antibiotic conjugates [16, 32, 33].
321 Other possible applications of siderophores include the prevention of post-harvest
322 diseases in stored crops, fruits, and vegetables or using siderophores as food-grade
323 antioxidants or in heavy metal bioremediation technologies [19, 34–36]. There-
324 fore, we need further studies to set up efficient and profitable technologies for the
325 bulk production of these promising compounds.

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330 Conflict of Interest

331 The authors declare that they have no conflict of interest.

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