

1 **Antifungal polycyclic tetramate macrolactam HSAF is a novel oxidative stress**
2 **modulator in *Lysobacter enzymogenes***

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4 Running title: HSAF as a novel oxidative stress modulator

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23 **ABSTRACT**

24

25 Polycyclic tetramate macrolactams (PoTeM) are a fast-growing family of antibiotic natural
26 products found in phylogenetically diverse microorganisms. Surprisingly, none of the PoTeM
27 had been investigated for potential physiological functions in their producers. Here, we used
28 HSAF (heat-stable antifungal factor), an antifungal PoTeM from *Lysobacter enzymogenes*, as
29 a model to show that PoTeM forms complexes with iron ion, with a K_a of 2.71×10^6 . The *in*
30 *vivo* and *in vitro* data showed formation of 2:1 and 3:1 complexes between HSAF and iron
31 ions, which were confirmed by molecular mechanical and quantum mechanical calculations.
32 HSAF protected DNA from degradation in high concentrations of iron and H_2O_2 or under UV
33 radiation. HSAF mutants of *L. enzymogenes* barely survived under oxidative stresses and
34 markedly increased the production of reactive oxygen species (ROS). Exogenous addition of
35 HSAF into the mutants significantly prevented ROS production and rescued the mutants to
36 normal growth under the oxidative stresses. The results reveal that the function of HSAF is to
37 protect the producer microorganism from oxidative damages, rather than as an
38 iron-acquisition siderophore. The characteristic structure of PoTeM,
39 2,4-pyrrolidinedione-embedded macrolactam, may represent a new iron-chelating scaffold of
40 microbial metabolites. Together, the study demonstrated a previously unrecognized strategy
41 for microorganisms to modulate oxidative damages to the cells.

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43

44 **Importance**

45 Polycyclic tetramate macrolactams (PoTeM) are a family of structurally distinct metabolites
46 that have been found in a large number of bacteria. Although PoTeM exhibit diverse
47 therapeutic properties, the physiological function of PoTeM in the producer microorganisms
48 had not been investigated. HSAF from *Lysobacter enzymogenes* is an antifungal PoTeM that
49 has been subjected to extensive studies for mechanism of biosynthesis, regulation and the
50 antifungal activity. Using HSAF as a model system, we here showed that the characteristic
51 structure of PoTeM, 2,4-pyrrolidinedione-embedded macrolactam, may represent a new
52 iron-chelating scaffold of microbial metabolites. In *L. enzymogenes*, HSAF functions as a
53 small molecule modulator for oxidative damages caused by iron, H₂O₂ and UV light.
54 Together, the study demonstrated a previously unrecognized strategy for microorganisms to
55 modulate oxidative damages to the cells. HSAF represents the first member of the fast
56 growing PoTeM family of microbial metabolites whose potential biological function has been
57 studied.

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59 **Key Words:** natural products, polycyclic tetramate macrolactams, *Lysobacter enzymogenes*,
60 oxidative damage, iron binding

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63 Introduction

64

65 Polycyclic tetramate macrolactams (PoTeM) are a family of natural products with diverse
66 therapeutic properties, including antibacterial, antifungal, anti-protozoa, and anticancer (1-9).
67 Their structures share a characteristic 2,4-pyrrolidinedione (tetramate)-containing
68 macrolactam and have been found in phylogenetically diverse bacteria. For example,
69 ikarugamycin, frontalamides, clifednamides, pactamides, capsimycins, and carbamides were
70 isolated from various species of *Streptomyces* (1, 3-5, 8-10). HSAF and several alteramides
71 were reported from several *Lysobacter* strains (2, 11-15). Maltophilin and xanthobaccin were
72 isolated from *Stenotrophomonas* strains (16, 17). Discoderamide and cylindramide were from
73 marine sponges (18, 19). Umezawamides were from a combined-culture of *Umezawaea* sp.
74 and mycolic-acid containing bacterium *Tsukamurella pulmonis* (20).

75 The biosynthetic gene cluster (BGC) for several PoTeM have been reported (2, 3, 5,
76 10-14). Although the chemical structures are complex, the BGC exhibits a relative simplicity
77 and a conserved organization. In the center of the BGC is always a single-module PKS-NRPS
78 hybrid gene, which is sufficient to construct the scaffold of PoTeM (10-12, 14, 21-23).
79 Flanking the PKS-NRPS gene are 2-6 accessory genes, which are responsible for the
80 structural diversity of PoTeM (14). Cryptic BGCs with this unique organization are present in
81 numerous genome sequences in the databases, implying that there is an immense reservoir of
82 PoTeM type of natural products yet to be discovered from the vast number of microorganisms
83 (3, 9, 10). The therapeutic properties, structural novelty, diverse bioactivities, and distinct
84 biosynthetic mechanism have attracted a lot of research interests in the recent years. However,

85 essentially nothing is known about PoTeM's physiological functions in their producer
86 organisms.

87 Many antibiotic metabolites are produced by microorganisms inhabiting in diverse
88 environments. In their native environments, the metabolites are typically not to function as
89 antibiotics to kill or inhibit other microorganisms because the producers rarely produce
90 inhibitory concentrations of the metabolites in the environments (24). Many factors in the
91 environments could affect the metabolite production and stress response in microorganisms.
92 For example, reactive oxygen species (ROS) are stimulated in microorganisms when growing
93 in a high iron environment or other stressed environments. Bacteria have evolved several
94 strategies to modulate the oxidative stress induced by a high ROS level. The thioredoxin (Trx)
95 system (NADPH, thioredoxin reductase and thioredoxin) is a crucial antioxidant system in
96 bacteria. The system removes ROS through providing electrons to thiol-dependent
97 peroxidases. In most Gram-negative bacteria, glutaredoxin system (Grx) and catalase
98 provide a strong backup for the Trx system (25). Some catalase-negative bacteria such as
99 *Streptococcus pyogenes* mainly utilize the thiol-dependent peroxidase system in defense
100 against oxidative stress although both Trx and Grx exist (26). Besides, carotenoids and the
101 aryl polyene type bacterial pigments are proved to protect bacteria from ROS, which is
102 related to their conjugation double bond systems (27-29). Recently, the H₂S-mediated
103 mechanism was found in protection against oxidative stress in *Escherichia coli* (30). The
104 endogenous H₂S produced by 3-mercaptopyruvate sulfurtransferase sequesters free ion,
105 which is necessary for the genotoxic Fenton reaction (30).

106 In this study, we have used the small molecule metabolite, HSAF (heat-stable antifungal

107 factor), from *L. enzymogenes*, as a model PoTeM to explore its potential physiological
108 functions. HSAF and alteramides (Fig. 1) isolated from *L. enzymogenes* are arguably the most
109 extensively investigated PoTeM in terms of structural diversity, bioactivities and modes of
110 action, and molecular mechanisms for biosynthesis (2, 6, 7, 11-14, 31). Here, our results
111 showed that the characteristic structure of PoTeM, 2,4-pyrrolidinedione-embedded
112 macrolactam, can act as a new iron-chelating natural product scaffold. HSAF functions as a
113 small molecule modulator for oxidative damages caused by iron, H₂O₂ and UV light in *L.*
114 *enzymogenes*. Together, the study demonstrated a previously unrecognized strategy for
115 microorganisms to modulate oxidative damages to the cells.

116

117 **RESULTS**

118 *Formation of brown-orange complexes between HSAF and iron*

119 During the study of *L. enzymogenes* OH11, we serendipitously found that adding iron salts
120 into minimal culture media could make OH11 grow more robustly (Fig. S1a). When we
121 investigated the effect of different concentrations of FeSO₄ (0, 1, 10, 100 and 500 μM) on the
122 growth of OH11 in a modified minimal medium (M813m) (Figure S1b), we observed
123 formation of brown-orange substances in both the cultures and the HSAF extracts (Fig. 2a-b).
124 This color was intensified with the increase of the FeSO₄ concentration and was absent in the
125 culture or extracts from HSAF mutant (Δ HSAF) (32), even when grown in M813m
126 containing 500 μM FeSO₄. The result suggested that HSAF and iron ions might be able to
127 interact with one another and form certain pigment complexes in the culture of OH11.
128 Moreover, we found FeSO₄ could significantly boost the production of HSAF and alteramides

129 (Fig. S2). With 1-10 μM FeSO_4 in M813m, the amount of free HSAF increased about 10-14
130 folds, compared to the medium without FeSO_4 .

131 To verify the HSAF-iron complexes, we extracted the total PoTeM mixture, containing
132 both HSAF and its analogs (alteramides), from the cultures. When the mixture was added into
133 an aqueous solution of FeSO_4 , the solution turned to the brown-orange color, with a gradually
134 increased intensity following the increase of the PoTeM mixture, while the controls remained
135 colorless (Fig. 2c). Furthermore, the same color could be developed in the PoTeM mixture
136 when added with other iron salts, such as $\text{Fe}(\text{NH}_4)_2(\text{SO}_4)_2$, FeCl_3 and $\text{Fe}(\text{NO}_3)_3$ (Fig. 2d).
137 HPLC analysis of the mixtures showed that the PoTeM peaks significantly decreased or
138 disappeared when any of the iron salts was added to the solutions (Fig. 2e). The interaction
139 between PoTeM and iron ions appeared to be specific, because HPLC showed that the PoTeM
140 peaks remained in the solutions, if the mixture was added with other metal ions (Na^+ , Mg^{2+} ,
141 K^+ , Ca^{2+} , Zn^{2+}), although there might have been some interactions between PoTeMs and Cu^{2+}
142 (Fig. S3).

143

144 *Mass spectrometry of HSAF-Fe complexes*

145 The above observations indicated that HSAF and its analogs could form complexes with iron
146 ions. To obtain direct evidence, we purified HSAF from the OH11 culture and treated HSAF
147 (10 mM) with the same concentration of aqueous FeSO_4 , $\text{Fe}(\text{NH}_4)_2(\text{SO}_4)_2$, FeCl_3 , or $\text{Fe}(\text{NO}_3)_3$.
148 The products were analyzed by mass spectrometry (MS). Without the iron ion, all MS gave
149 two main peaks, m/z 513 for $[\text{HSAF}+\text{H}]^+$ and m/z 1025 for $[2\text{HSAF}+\text{H}]^+$, which are
150 expected for standard HSAF (Fig. 3). Upon treatment with the iron ion, the HSAF peaks

151 markedly decreased (when added with ferrous ion, FeSO_4 or $\text{Fe}(\text{NH}_4)_2(\text{SO}_4)_2$) or disappeared
152 (when added with ferric ion, FeCl_3 or $\text{Fe}(\text{NO}_3)_3$). Meantime, two new peaks appeared (m/z
153 1079 and 1591) when treated with FeSO_4 or $\text{Fe}(\text{NH}_4)_2(\text{SO}_4)_2$ (Fig. 3a-b), or just one new peak
154 appeared (m/z 1079) when treated with FeCl_3 or $\text{Fe}(\text{NO}_3)_3$ (Fig. 3c-d). The peak at m/z 1079
155 is coincident with $[\text{2HSAF-H+Fe}]^+$, whereas the peak at m/z 1591 is coincident with
156 $[\text{3HSAF-H+Fe}]^+$. The data showed that two or three HSAF molecules could coordinate with
157 one iron ion, to form stable HSAF-Fe complexes that displayed the observed orange-brown
158 color. The data also suggested that all HSAF chelated with Fe when a ferric salt was used, as
159 seen in FeCl_3 and $\text{Fe}(\text{NO}_3)_3$, but only a portion of HSAF chelated with iron when a ferrous
160 salt was used, as seen in FeSO_4 and $\text{Fe}(\text{NH}_4)_2(\text{SO}_4)_2$ (Fig. 3). Since ferrous iron can gradually
161 be oxidized to ferric ion in the atmosphere, it is likely that the observed partial chelation in
162 FeSO_4 and $\text{Fe}(\text{NH}_4)_2(\text{SO}_4)_2$ was due to the oxidized iron (ferric). Besides, both 2HSAF-Fe
163 and 3HSAF-Fe were observed when ferrous salts were used, whereas only 2HSAF-Fe was
164 detected when ferric salts were used. This also supports that ferric ion is the preferred iron for
165 HSAF chelation, because the concentration of ferric ion in solution would be lower when
166 FeSO_4 and $\text{Fe}(\text{NH}_4)_2(\text{SO}_4)_2$ were used than that when FeCl_3 and $\text{Fe}(\text{NO}_3)_3$ were used, and thus
167 HSAF concentration was relatively high and two or three HSAF molecules were available to
168 chelate one ferric ion in FeSO_4 and $\text{Fe}(\text{NH}_4)_2(\text{SO}_4)_2$ solutions. To further confirm the
169 formation of HSAF-Fe complexes, EDTA, a strong chelator for metal ions, was added into
170 the mixtures of HSAF and iron salts. MS clearly showed that the HSAF-Fe complexes (m/z
171 1079 and 1591) were abolished and HSAF (m/z 513 and 1025) was restored (Fig. 3).

172

173 *Absorbance spectra of HSAF-Fe complexes*

174 Next, we analyzed the absorbance spectra of the HSAF-Fe complexes (Fig. S4). Standard
175 HSAF gave a maximum peak at ~323 nm, which shifted to ~310 nm upon addition of any of
176 the iron salts, FeSO₄, Fe(NH₄)₂(SO₄)₂, FeCl₃, or Fe(NO₃)₃. While HSAF or the iron salts
177 barely had any absorption in the visible range, the HSAF-Fe complexes gave clear
178 absorptions at ~410-600 nm, which apparently contributed to the orange-brown appearance
179 of the mixtures. Furthermore, the maximal absorption shifted back to 323 nm from 310 nm
180 and the absorption at ~410-600 nm disappeared, upon addition of EDTA (Fig. S4). The
181 absorption spectroscopic data are in accordance with that of MS analysis. Using the
182 UV-visible titration of HSAF with Fe(NO₃)₃ and nonlinear curve-fitting at 470 nm, we
183 obtained the association constant (K_a) of HSAF-Fe to be 2.71×10^6 (Fig. S5). The K_a value is
184 much smaller than that for recognized siderophores (33), indicating that, rather than function
185 as a siderophore for iron acquisition from the environment, HSAF in *L. enzymogenes* may
186 play a new function during the interaction with iron ion.

187

188 *Antioxidant activity of HSAF and protection of DNA degradation in vitro*

189 To understand possible roles of HSAF in *L. enzymogenes*, we explored the potential
190 involvement of HSAF in modulation of oxidative stress, because it is well known that the
191 cellular iron could generate reactive oxygen species (ROS) due to the Fenton reaction, which
192 can lead to cell death (34). The *in vivo* and *in vitro* data described above showed the
193 formation of HSAF-Fe complexes, which might contribute to maintaining a proper free iron
194 concentration important to redox homeostasis of the bacterial cells.

195 To test this hypothesis, we used the deoxyribose degradation assay to determine HSAF's
196 antioxidant activity (35). HSAF showed a dose-dependent antioxidant activity and behaved as
197 a strong antioxidant when the concentration reached 80-160 μM (Fig. 4a). The reaction
198 system for the assay contained deoxy-D-ribose, H_2O_2 , Fe^{3+} , ascorbic acid, and purified HSAF.
199 To exclude possible direct interactions between HSAF and H_2O_2 , ascorbic acid, or
200 deoxy-D-ribose, we analyzed the mixtures using HPLC. As expected, HSAF was not affected
201 by any of the factors (H_2O_2 , ascorbic acid, deoxy-D-ribose) (Fig. S6). The results indicated
202 that the antioxidant activity of HSAF resulted from the chelation with iron ion. To confirm
203 the antioxidant function of HSAF, we carried out *in vitro* DNA fragment degradation caused
204 by Fenton reaction due to production of the radical species (36). A DNA fragment with the
205 length of 1 Kb was amplified by PCR using the genome of *L. enzymogenes* as template. The
206 DNA fragment was completely degraded in the presence of Fe^{3+} and H_2O_2 , while the DNA
207 fragment remained intact in the controls. However, the addition of purified HSAF inhibited
208 the DNA degradation, in a dose-dependent manner (Fig. 4b). When HSAF reached to 80 μM ,
209 the DNA fragment was fully protected from the Fenton reaction-caused degradation, which is
210 in good agreement with the antioxidative activity assay (Fig. 4a). To learn whether this
211 protective effect of HSAF is specific to certain DNA fragments, we tested similar length
212 DNA fragments from *Lysobacter* 3655, *Lysobacter antibioticus* OH13, and *Escherichia coli*.
213 HSAF exhibited the similar protective effect against the Fenton reaction-caused damage on
214 these DNA fragments, showing a general antioxidative effect of HSAF (Fig. S7).
215
216 *Protection of L. enzymogenes OH11 from high H₂O₂ stress by HSAF in vivo*

217 The observation that HSAF can protect *in vitro* DNA degradation by H₂O₂ inspired us to
218 investigate HSAF's function *in vivo*. The WT and Δ HSAF strains were treated with different
219 concentrations of H₂O₂ in M813m medium. The results clearly showed that the growth rate of
220 the WT and Δ HSAF was similar when culture medium contained a low concentration of H₂O₂
221 (80 μ M), but in the medium containing a high concentration of H₂O₂ (800 μ M), the growth
222 rate of Δ HSAF strain significantly decreased when compared to that of the WT (Fig. 4c-4e).
223 Actually, Δ HSAF strain did not grow in M813m containing 800 μ M H₂O₂ in the first 48 h and
224 started a slow growth only at 72 h, while the WT grown in M813m containing 800 μ M H₂O₂
225 could reach the similar OD₆₀₀ values as the WT without H₂O₂ at 72-96 h. We also analyzed
226 the HSAF level in cultures containing 0, 80 and 800 μ M H₂O₂. It showed that the production
227 of HSAF/alteramides in the WT with 800 μ M H₂O₂ decreased by 30% when compared to that
228 of the WT with 0 or 80 μ M H₂O₂, implying a consumption of HSAF for formation of
229 HSAF-Fe complexes, to protect the cells grown in a high concentration of H₂O₂ (Fig. S8). To
230 exclude the possibility that HSAF in WT could directly degrade H₂O₂ so that WT could grow
231 in a high concentration of H₂O₂, we tested the ability of HSAF and HSAF-Fe complexes to
232 degrade H₂O₂ *in vitro* (Fig. S9). The results show that H₂O₂ was not degraded by either
233 HSAF or HSAF-Fe complexes. Furthermore, MS analysis of the ethyl acetate extract from
234 the WT culture clearly detected the peaks at *m/z* 1079 for [2HSAF-H+Fe]⁺ and 1591 for
235 [3HSAF-H+Fe]⁺, in addition to *m/z* 513 for [HSAF+H]⁺ (Fig. S10). This showed HSAF-Fe
236 complexes were formed *in vivo*.

237

238 *Protection of L. enzymogenes OH11 from high iron stress in vivo*

239 Next, we looked into the effect of HSAF on the growth yield of *L. enzymogenes* OH11 under
240 different concentrations of iron ion. The WT and Δ HSAF strains in regular M813m medium
241 containing 10 μ M iron ion exhibited a similar growth yield to that in M813m medium
242 without iron ion (Fig. S11a-b). However, in M813m containing a high concentration (500 μ M)
243 of iron ion, the growth yield of Δ HSAF strain was significantly lower than that of the WT
244 after 72-120 h growth, although both strains exhibited a similar growth yield in the first 48 h
245 (Fig. S11c). The addition of HSAF to the cultures after 48 h growth restored the growth yield
246 of Δ HSAF strain to the WT level in the following 72-120 h growth, even when the iron
247 concentration was as high as 500 μ M (Fig. S11d). The results show that HSAF can protect
248 OH11 cells from high iron stress *in vivo*. Besides, the exogenously added HSAF-Fe
249 complexes, but not HSAF alone, could promote the growth of WT strain in M813m medium
250 without supplemented FeSO₄ (Fig. S12a). However, neither HSAF nor HSAF-Fe complexes
251 affect the growth of WT strain in regular M813m medium containing FeSO₄ (Fig. S12b). The
252 results suggested that iron ion could be released from HSAF-Fe complexes and then up-taken
253 by the cells to support the observed growth promotion.

254

255 *Protection of L. enzymogenes OH11 from UV radiation by HSAF*

256 In light of HSAF's protection of *L. enzymogenes* from oxidative damages induced by high
257 concentrations of iron and H₂O₂, we looked into the protective role of HSAF when the cells
258 were exposed to UV radiation, because UV exposure can also lead to a variety of ROS
259 through various mechanisms (37). When the WT and Δ HSAF strains were exposed to the UV
260 light for a short time (10 s), the survival rates of the WT and Δ HSAF cells were similar.

261 However, the survival rate of Δ HSAF cells was significantly decreased when the UV
262 exposure time increased to 30 s, and there was nearly zero colony on the plate with 60 s of
263 UV exposure, while the WT still had a large number of colonies (Fig. 5a, S13a). On the other
264 hand, the exogenous addition of purified HSAF to the Δ HSAF culture before the UV
265 exposure could restore the growth. The rescue of the Δ HSAF cells by HSAF showed a clear
266 dose-dependent manner, and when the HSAF concentration reached 160 μ M, the survival rate
267 of Δ HSAF was even higher than that of the WT (Fig. 5b, S12b). The results unequivocally
268 showed the protective effect of HSAF on cells with UV radiation.

269

270 *Repression of ROS accumulation in L. enzymogenes OH11 by HSAF*

271 Next, we tested the ROS formation in the strains using the method of H2DCFDA
272 (2',7'-dichlorodihydrofluorescein diacetate), which is a cell-permeant indicator that generates
273 fluorescence after oxidation by ROS (38). The results showed almost no detectable ROS in
274 the WT and Δ HSAF strains when cultured in the minimal medium without iron ion, and a low
275 level of ROS production similarly in the WT and Δ HSAF strains in the modified medium
276 containing a low level of iron (M813m containing 10 μ M FeSO₄) (Fig. 5c). In a high iron
277 medium (M813m containing 500 μ M FeSO₄), Δ HSAF strain produced nearly a double
278 amount of ROS than the WT, although the ROS level increased significantly in both the WT
279 and Δ HSAF. Remarkably, when exogenous HSAF was added to the Δ HSAF strain grown in
280 the 500 μ M FeSO₄ medium, the ROS level returned to the WT level. The results clearly
281 showed that HSAF is able to repress the high iron-caused ROS production (Fig. 5c).
282 Moreover, the ROS level in Δ HSAF was about 2 fold higher than that in the WT, when

283 treated with H₂O₂ or UV lights, and the exogenous HSAF restored the WT level of ROS in
284 ΔHSAF strain (Fig. 5d-e). Next, we directly measured the ROS scavenging activity of HSAF
285 and HSAF-Fe complexes using DPPH (2,2-Diphenyl-1-picrylhydrazyl), which is a stable
286 radical and has the maximum absorption at 520 nm. The results showed that HSAF and
287 HSAF-Fe complexes exhibited low ROS scavenging activity when the incubation time was
288 0.5 h, while with the extension of incubation time (24 h, 72 h and 120 h), the ROS
289 scavenging activity of both HSAF and HSAF-Fe complexes increased (Fig. S14a-b). As
290 expected, ascorbic acid showed strong ROS scavenging activity (Fig. S14c). This is
291 consistent with the observation in the UV irradiation assay, where the cultural time was 72 h
292 and HSAF exhibited ROS scavenging activity. The data clearly showed the ROS modulating
293 ability of HSAF in *L. enzymogenes*.

294

295 *Molecular structure of HSAF-Fe complexes*

296 To obtain further evidence for formation of the HSAF-Fe complexes, we performed
297 molecular mechanical and quantum mechanical calculations to determine the possible
298 molecular structures of the HSAF-Fe complexes. Using a molecular mechanical force field
299 method, a global search suggested that both Fe³⁺ and Fe²⁺ ions can be chelated by 2 or 3
300 HSAF neutral molecules. Quantum mechanical method was used to refine the molecular
301 geometries (Fig. 6). When 2 molecules of HSAF bind to an iron ion, the 3 carbonyl oxygen
302 atoms (at C7, C25, C27, see Fig. 1) of each of the two HSAF molecules form 3 coordinate
303 bonds to the iron. When 3 molecules of HSAF bind to an iron ion, two HSAF molecules
304 provide the oxygen atoms at C7 and C27, and the third HSAF molecule provides the oxygen

305 atoms at C25 and C27. Together, the three HSAF molecules form 6 coordinate bonds to the
306 iron. Due to steric factors, it would be impossible for 4 molecules of HSAF to bind to one
307 iron. The carbonyl oxygen atoms at C7, C25 and C27 are absolutely conserved in all PoTeMs,
308 suggesting that formation of such iron complexes are general for all PoTeMs. The chelation
309 status is similar to that of *Pseudomonas* quinolone signal (PQS) with iron ion, in which two
310 or three PQS molecules chelated one iron ion (39).

311

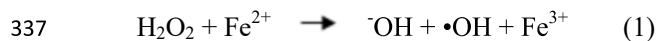
312 **DISCUSSION**

313 Since the isolation of HSAF and its analogs from *Lysobacter enzymogenes*, the research has
314 focused on their antifungal activity, as well as the molecular mechanism for their biosynthesis
315 and regulation. The work presented here is the first attempt to address the role of these
316 complex molecules in their producer organism. *Lysobacter* species are emerging as a rich
317 source of bioactive natural products. During our efforts to activate silent biosynthetic gene
318 clusters in the genomes of *Lysobacter* species, we serendipitously found that the addition of
319 iron salts enhanced HSAF production. Meantime, we observed the formation of a
320 brown-orange color when iron salts were added to the cultures that produced HSAF and
321 analogs, but not in the biosynthetic mutant. *In vitro* studies using the crude extracts and
322 purified HSAF confirmed that the color was due to formation of complexes between these
323 compounds and iron. The results also showed that the production of HSAF and analogs is
324 essential for *L. enzymogenes* to survive under oxidative conditions that are known to generate
325 reactive oxygen species (ROS).

326 When *L. enzymogenes* was exposed to a high concentration of iron, H₂O₂, or UV

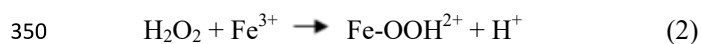
327 radiation, the HSAF biosynthetic mutant was barely able to survive, while the wild type and
328 the mutant supplemented with HSAF exogenously could grow normally. In bacteria, ROS are
329 induced by many stresses including presence of a high iron concentration, H₂O₂ treatment,
330 and UV radiation (40). Indeed, we observed a significantly higher level of ROS production in
331 the HSAF mutant than in the wild type. The supplement of HSAF into the mutant reduced the
332 ROS level to that of the wild type.

333 We also observed that HSAF protects DNA from degradation in the presence of iron and
334 H₂O₂, probably due to hydroxyl radical ([•]OH), generated by Fenton reaction [Eq. (1)].
335 Hydroxyl radical is highly reactive ROS and able to oxidize practically every molecule in the
336 cell (41, 42).



338 In both prokaryotes and eukaryotes, the oxidative DNA damage caused by hydroxyl
339 radical is the primary cause of cell death under oxidative stress conditions (43-45). We thus
340 hypothesized that HSAF may be involved in Fenton reaction and affect the redox
341 homeostasis of *L. enzymogenes*. In bacteria, both O₂^{•-} and H₂O₂ are primarily produced by
342 the accidental autoxidation of non-respiratory flavoproteins which are univalent electron
343 donors giving electrons to oxygen (46). Besides, a high iron concentration also could promote
344 the H₂O₂ generation. In deoxyribose degradation assay, ascorbic acid initializes the Fenton
345 reaction by reducing Fe³⁺ to Fe²⁺, and Fe²⁺ in turn reacts with H₂O₂ to generate ROS ([•]OH)
346 [Eq. (1)] (35). In the *in vitro* assay of DNA fragment degradation, Fe³⁺ could be reduced to
347 Fe²⁺ through reacting with H₂O₂, by following the two-step reactions [Eq. (2)-(3)] (47). The
348 reactions generate Fe²⁺ and then ROS by the reaction between Fe²⁺ and H₂O₂ as shown in Eq.

349 (1).



352 As shown in Fig. 2, HSAF prefers ferric ion over ferrous ion in the chelation. The
353 HSAF-Fe³⁺ chelation would prevent the reduction of Fe³⁺ to Fe²⁺ in the two-step reactions
354 [Eq. (2)-(3)]. Consequently, the sequestration of Fe³⁺ ion by HSAF would result in a reduced
355 amount of free Fe²⁺ in the cells and in turn a reduced ROS from the Fenton reaction. The *K_a*
356 value of HSAF-Fe complexes is significantly smaller than that of typical siderophores. This
357 suggests that a relatively high concentration (μM level) of HSAF would be needed in order to
358 form stable complexes with iron. HSAF and alteramides are the predominant secondary
359 metabolites in *L. Enzymogenes* (2). We were able to obtain up to 50 mg HSAF from 1 L
360 culture, suggesting that the concentration of HSAF in the cells would be higher than 100 μM.
361 Thus, the concentration at which HSAF exhibited iron-chelation and strong antioxidant
362 activity *in vitro* could be readily achievable *in vivo*.

363 Moreover, we showed that the exogenous addition of HSAF into the culture media could
364 restore the ΔHSAF mutant's growth under high iron concentrations, H₂O₂, and UV radiation;
365 we also showed that the exogenous HSAF was able to make the ROS level in the ΔHSAF
366 mutant return to the WT level, even when grown in the 500 μM FeSO₄ medium. Together,
367 these observations implied that extracellular HSAF was able to enter the cells. However, the
368 exact mechanism of HSAF transportation is not totally clear at the moment. Beyond chelating
369 iron, HSAF can also scavenge ROS directly as seen in Fig. 5c-e and Fig. S14a-b. These
370 explain the protective effect of HSAF on *Lysobacter* with UV exposure (Fig. 5a-b).

371 Although HSAF is able to chelate iron, the HSAF-Fe complexes form only when the iron
372 concentration is sufficiently high (above μM level). This is in contrast to siderophores whose
373 primary function is to grab metal ions from the environments where the concentration of the
374 metals can be extremely low. HSAF functions as a modulator for oxidative stresses only
375 when the cells are exposed to an environment with a high concentration of iron, H_2O_2 , or
376 exposed to UV light, all of which can lead to ROS generation that damages DNA and cell
377 survival.

378 Moreover, our study using methods in molecular mechanical force field and quantum
379 mechanical indicated that the carbonyl groups at C7, C25 and C27 of HSAF structure are
380 involved in formation of the HSAF-Fe complexes. These carbonyl groups are absolutely
381 conserved in all PoTeMs. This finding is significant because it suggests that formation of
382 such iron complexes could be general for all PoTeMs. On the other hand, there is a clear
383 structural diversity among the PoTeM family, which is derived from the polycyclic system
384 (14). While HSAF and alteramides are known for antifungal activity, other members of the
385 PoTeM family exhibit antitumor, antiprotozoal, cytotoxic, and antiviral activities. The
386 structural diversity of PoTeMs may be associated with these activities and may also confer
387 survival advantage to their producers in various habitats. For example, *L. enzymogenes* OH11
388 was originally isolated from the rhizosphere of a pepper plant (32). Whether HSAF and
389 alteramides play a role in plant root colonization is worth a further investigation, as resistance
390 to oxidative stress is important for the survival of bacteria during their interaction with plants.
391 It is not very clear what structural features in these compounds are associated with the
392 antifungal activity, although we found the carbonyl oxygen atoms at C7, C25 and C27 are

393 involved in the iron chelation to form HSAF-Fe complexes. *L. enzymogenes* can produce
394 HSAF and analogs without iron and with iron in the minimal media (up to 500 μ M). In reality,
395 it is unlikely that the iron concentration in the natural environment of *L. enzymogenes* would
396 be higher than what we have tested. Several iron chelators have exhibited antifungal activity,
397 such as siderophores produced by *Azospirillum brasilense* could inhibit the growth of
398 *Colletotrichum acutatum* and the siderophore oxachelin from *Streptomyces* sp. GW9/1258
399 showed strong antimicrobial activity against several fungi and Gram-positive bacteria (48,
400 49). Further studies are needed in order to answer whether the antifungal activity and the
401 antioxidant activity are related to each other or exclude each other. Nevertheless, it seems
402 reasonable to assume that the antioxidant activity of HSAF and analogs could enhance the
403 survival rate of the producer microorganism through the enhanced resistance to oxidative
404 stress and thus play a role in plant root colonization during their interaction with plants.
405 Several strategies are evolved in bacteria to modulate the oxidative stress (25-30).

406 In summary, HSAF has been recognized as an antifungal antibiotic with a fascinating
407 chemical structure, new mode of action, and distinct mechanism for biosynthesis (2, 6, 7, 31).
408 Here, we presented evidence to support that HSAF and its analogous compounds may
409 represent a new strategy for microorganisms to modulate the oxidative stress. These
410 “secondary metabolites” confer *L. enzymogenes* to survive in the environment with a high
411 concentration of iron, H₂O₂ or UV radiation. HSAF is the first member of the fast growing
412 PoTeM family of natural products whose potential biological function has been investigated.
413 The genome mining efforts have shown a strikingly conserved organization for PoTeM
414 biosynthetic clusters, which are present in a large number of unexplored genomes of

415 phylogenetically diverse bacteria ranging from proteobacteria to actinomycetes (3, 10, 14).
416 This means there are a large number of new PoTeMs yet to be explored. The understanding
417 the biological function of HSAF shed new lights into the critical role of “secondary
418 metabolites” in the survival of microorganisms in complex ecosystems, and the results will
419 also facilitate the future efforts in new PoTeM exploitation.

420

421 MATERIAL AND METHODS

422

423 *Bacterial cultures, primers and PCR*

424 The wild type strain *Lysobacter enzymogenes* OH11 (CGMCC No. 1978) and the HSAF
425 biosynthetic mutant Δ HSAF (Table 1) were cultured in Luria-Bertani (LB) medium (32). For
426 the production of HSAF and its analogs, the strains were cultured in M813 modified medium
427 (4 g Glucose, 3 g K_2HPO_4 , 1.2 g NaH_2PO_4 , 1 g NH_4Cl , 0.3 g $MgSO_4$, 0.15 g KCl , 10 mg
428 $CaCl_2$, 2.8 mg $FeSO_4 \cdot 7H_2O$, per liter) (50). In the initial experiments, the strains were also
429 cultured in MM2 medium (4 g Glucose, 15 g KH_2PO_4 , 34 g Na_2HPO_4 , 5.4 g NH_4Cl , 2.5 g
430 $NaCl$, 0.3 g $MgSO_4$, 10 mg $CaCl_2$, per liter). Table 2 listed the primers used in this study.
431 Phusion High-Fidelity DNA polymerase (Thermo Scientific) was used as the amplification
432 enzyme. The PCR started from an initial denaturation at 98°C for 30 s followed by 30 cycles
433 of amplification (98°C for 10 s, 60°C for 15 s, 72°C for 1 min), and completed with
434 additional 5 min at 72°C. Depending on the DNA templates and primers, the annealing
435 temperature and the elongation time were adjusted in some case

436

437 *Extraction and HPLC analysis of HSAF and its analogs*

438 *L. enzymogenes* OH11 wild type (WT) and HSAF deficiency strain (Δ HSAF), in which a part

439 of *pks-nrps* gene (from +232 bp to +1356 bp with relative to the start codon) of HSAF

440 biosynthetic gene cluster was deleted (32), were incubated into 1 ml LB at 30°C with shaking

441 of 200 rpm for overnight. An aliquot (1%) of the cultures was transferred to 25 ml M813

442 modified (M813m) medium with variable concentrations of FeSO₄ (final concentration of 0,

443 1, 10, 100 and 500 μ M), 30°C with shaking at 200 rpm for 48 h. The whole cultures (cells

444 and medium) were treated with 75 μ l TFA and 25 ml ethyl acetate. The organic phase was

445 dried with the air flow, and the residues were re-dissolved in 200 μ l methanol. A 2 μ l aliquot

446 of each extract was analyzed by HPLC (Agilent, 1220 Infinity LC). Water/0.05% FA (solvent

447 A) and acetonitrile/0.05% FA (solvent B) were used as the mobile phases with a flow rate of

448 1.0 ml/min. The HPLC program was as follows: 5–25% B in 0–5 min, 25%–80% B in 5–25

449 min, 80–100% B in 25–26 min, maintained to 28 min, back to 5% B at 29 min and

450 maintained to 30 min. HSAF and its analogs were detected at 318 nm on a UV detector. For

451 purification of HSAF, WT was incubated into 10 ml LB at 30°C with shaking at 200 rpm for

452 overnight. An aliquot (1%) of the cell cultures was transferred to 1 L M813m medium and grew

453 at 30°C with shaking at 200 rpm for 48 h. The culture broth was adjusted to pH 2.5 with 37%

454 HCl. The culture was added with the same volume of ethyl acetate, and HSAF was extracted

455 into the organic phase for three times. The ethyl acetate phase was dried using a rotavapor, and

456 HSAF was separated from other metabolites in the extract (850 mg) on a C18 reverse-phase

457 column, eluted with different concentrations of methanol (10%, 30%, 50%, 70%, and 100%).

458 The fraction (552 mg) of 100% methanol was used to purify HSAF (~50 mg) by HPLC.

459

460 *Chelation of HSAF with metal ions*

461 For the chelation with FeSO₄, an increased volume (1, 10, 50 µl, in methanol) of the total
462 PoTeM mixture (2 mg/ml) was mixed with 100 µl aqueous solution of FeSO₄ (10 mM). For
463 the chelation with other iron salts, the total PoTeM mixture (50 µl, 2 mg/ml) was incubated
464 with 50 µl aqueous solution of Fe(NH₄)₂(SO₄)₂, FeCl₃, and Fe(NO₃)₃ (each 10 mM). For the
465 chelation with other metals, the total PoTeM mixture (50 µl, 0.5 mg/ml) was incubated with
466 50 µl aqueous solution of Na₂SO₄, MgSO₄, K₂SO₄, Ca(NO₃)₂, CuSO₄, and ZnSO₄ (each 10
467 mM). Methanol was used as control. For HPLC analysis, the above mixed solutions were
468 dried and then resuspended in 100 µl methanol. A 20 µl aliquot of each of the solutions was
469 analyzed by HPLC (Agilent, 1220 Infinity LC). The HPLC program was as follows:
470 Water/0.05% FA (solvent A) and acetonitrile/0.05% FA (solvent B) were used as the mobile
471 phases with a flow rate of 1.0 ml/min. The program was as follows: 5%-60% B in 0–5 min,
472 60%-100% B in 5–20 min, maintained to 23 min, back to 5% B at 28 min, and maintained to
473 30 min. The metabolites were detected at 230 nm on a UV detector. The experiments were
474 repeated for three times.

475

476 *MS analysis and UV-Visible absorbance spectra of the HSAF-Fe complexes*

477 To prepare the complexes, purified HSAF (20 µl, 10 mM) was mixed with an equal volume of
478 each of aqueous FeSO₄ (10 mM), Fe(NH₄)₂(SO₄)₂ (10 mM), FeCl₃ (10 mM), and Fe(NO₃)₃
479 (10 mM). Each of the mixtures was dried and re-suspended in 200 µl ddH₂O. Ethyl acetate
480 (200 µl, containing 0.05% TFA) was added to the suspension to extract the HSAF-Fe

481 complexes, and the organic phase was collected and dried and re-dissolved in 1 ml methanol.
482 A 20 μ l aliquot of each of the samples was analyzed by MS and the remaining fraction of the
483 samples was used to determine the UV-visible absorbance spectrum by a spectrophotometer
484 (Shimadzu UV-Vis 2501). After the spectra were taken, 50 μ l Na₂EDTA (100 mM, pH 8.0)
485 was added into each of the samples, and the samples were dried and re-suspended in 250 μ l
486 ddH₂O. Similar to the above procedure, the HSAF-Fe complexes were extracted with 250 μ l
487 ethyl acetate (containing 0.05% TFA), the organic phase was collected and dried and
488 re-dissolved in 1 ml methanol. Then samples then were similarly analyzed by MS and
489 spectrophotometer again. The experiments were repeated for three times.

490

491 *UV-Visible absorbance titration and association constant determination*

492 To determine the association constant, absorbance spectra were recorded with a
493 spectrophotometer (Shimadzu UV-Vis 2501). The UV-Visible titration assay was performed
494 by using a constant host concentration of HSAF (0.1 mM) and variable concentrations of
495 Fe(NO₃)₃ at 25°C. Association constant (K_a) was calculated using Eq. (4) by applying a
496 nonlinear curve-fitting method (51) on Program of origin 9.0 to changes in absorbance (Δ Abs)
497 at 470 nm.

$$498 \quad \Delta\text{Abs} = (L(1 + K_a X + K_a A) - (L^2(K_a X + K_a A + 1)^2 - 4K_a^2 A X L^2)^{0.5}) / 2K_a A \quad (4)$$

499 Where X and A were the total concentration of the guest and the host, respectively, and L
500 and K_a were treated as parameters. The experiments were repeated for three times.

501

502 *Assay for deoxyribose degradation*

503 This assay was used to determine the antioxidative activity. In the assay, hydroxyl radicals
504 generated by the Fenton reaction would degrade deoxy-D-ribose into malonyldialdehyde
505 (MDA) (35). MDA then would react with 2-thiobarbituric acid to produce a pink pigment,
506 which was determined photometrically at 532 nm (34). To perform the assay, 227.5 μ l
507 variable concentration of purified HSAF (final concentration is 0, 5, 10, 20, 40, 80, and 160
508 μ M) were added into the reactant mixture containing 45.5 μ l Deoxy-D-ribose (52 mM, in 50
509 mM KH_2PO_4 , pH 7.4), 91 μ l FeCl_3 (4 μ M, in water), 91 μ l KH_2PO_4 buffer (pH 7.4), 45.5 μ l
510 H_2O_2 (10 mM, in water), and 45.5 μ l ascorbic acid (1 mM, in 50 mM KH_2PO_4 , pH 7.4). The
511 samples were mixed and incubated at 30°C for 60 min. After that, 455 μ l 2-thiobarbituric acid
512 (1%, in 3% trichloroacetic acid) was added into each of the samples, and the mixtures were
513 incubated at 85°C for 30 min. The supernatant of each of the mixtures was collected
514 following centrifugation (12,000 rpm, 1 min), and the OD_{532} value of the supernatants was
515 determined by spectrophotometer (Shimadzu UV-Vis 2501). The experiments were repeated
516 for three times.

517

518 *Assay for in vitro DNA degradation*

519 The assay followed a previously described method with some modifications (36). The DNA
520 fragments were obtained by PCR using primers listed in Table S2 with a template from the
521 genomic DNA of *L. enzymogenes* OH11, *Lysobacter* 3655, *L. antibioticus* OH13, or *E. coli*.
522 Each of the degradation mixtures contained 100 ng DNA fragment, 20 mM H_2O_2 , 150 μ M
523 FeCl_3 and various concentrations of purified HSAF (0.3125, 0.625, 1.25, 2.5, 5, 10, 20, 40
524 and 80 μ M) in 50 mM KH_2PO_4 buffer (pH 7.4). After incubating the mixtures at 37°C for 60

525 min, the samples were applied to 0.8 % agarose gels in TAE buffer. The electrophoresis was
526 performed at 140 V for 15 min, and the bands were visualized in a UV transilluminator
527 (Universal Hood, Bio-Rad). The experiments were repeated for three times.

528

529 *OH11 growth under oxidative stress induced by H₂O₂*

530 WT and Δ HSAF strains were incubated into 1 ml LB at 30°C with shaking of 200 rpm for
531 overnight. A fraction (1%) of the cultures was transferred to 25 ml M813m medium at 30°C
532 with shaking at 200 rpm for 48 h. After the OD₆₀₀ value of each of the cultures was
533 determined, a fraction (1%) of the cultures was added into 25 ml M813m medium containing
534 0, 80, or 800 μ M H₂O₂, and the cultures were incubated at 30°C with shaking at 200 rpm for
535 96 h. The OD₆₀₀ values of WT and Δ HSAF strains were recorded every 24 h. The
536 experiments were repeated for three times.

537

538 *Assay for in vitro H₂O₂ degradation*

539 The Hydrogen Peroxide (H₂O₂) Colorimetric Assay Kit (Elabscience, China) was used to
540 detect the concentration of H₂O₂ which could react with ammonium molybdate and produced
541 a yellow complex with the maximum absorption at 405 nm. The reaction system contained 2
542 ml reaction buffer, 100 μ l H₂O₂ (60 mM), and 100 μ l purified HSAF (4 mM) or 100 μ l
543 purified HSAF-Fe complexes (4 mM). Methanol was used as negative control. The reaction
544 system was incubated at 30°C for 30 min, then the OD₄₀₅ value was determined by
545 spectrophotometer.

546

547 *OH11 growth under oxidative stress induced by UV light*

548 For the WT and Δ HSAF of *L. enzymogenes* OH11, the strains were incubated into 1 ml LB at
549 30°C with shaking at 200 rpm for overnight. A fraction (1%) of the cultures was transferred
550 to 25 ml M813m medium at 30°C with shaking at 200 rpm for 48 h. The cultures were
551 adjusted with the medium to the same OD₆₀₀ of 1.5 and a fraction of 10 ml of each of the
552 cultures was spread on a petri dish (9.0 cm, external diameter) and exposed to a UV light
553 source (253.7 nm, Model TUV 30W T8, 102 Volts, 0.37 AMPS, 30 Watts), at a distance of 30
554 cm between the light and the cells for 0 s, 10 s, 30 s or 60 s. For the complementary assay,
555 purified HSAF with a various concentration (final concentration of 0, 20, 80 and 160 μ M)
556 was added into the culture of Δ HSAF strain before exposure to UV for 60 s, and methanol
557 was used as control. The cultures were serially diluted and spread on fresh LB plates. The
558 numbers of colonies on each plate were counted after 72 h of incubation at 30°C. The
559 experiments were repeated for three times.

560

561 *ROS detection*

562 The production of ROS in WT and Δ HSAF strains were detected using a previous method
563 with some modifications (38). WT and Δ HSAF strains were incubated into 1 ml LB, 30°C
564 with shaking of 200 rpm for overnight. A fraction (1%) of the cultures was transferred into 25
565 ml M813m medium with various concentrations of FeSO₄ (0, 10, 500 μ M). After 72 h of
566 growth, the cultures were diluted 30-fold with the same medium in a 96-well plate. Then
567 H2DCFDA was added to the wells with a final concentration of 10 μ M. The incubation of the
568 plate continued in the dark at 30°C with shaking at 60 rpm for 6 h. Fluorescence was

569 measured in a BioTek Synergy H1 plate reader (excitation, 495 nm; emission, 527 nm). In
570 addition, the cultures of WT and Δ HSAF from regular M813m medium containing 10 μ M
571 FeSO_4 were treated with UV for 60 s without or with purified HSAF (final concentration 160
572 μ M), or treated with H_2O_2 (final concentration 40 mM) before fluorescence detection. The
573 experiments were repeated for three times.

574

575 *Assay for ROS scavenging activity of HSAF and HSAF-Fe complexes*

576 HSAF and HSAF-Fe complexes were tested for *in vitro* ROS scavenging activity using
577 DPPH (2,2-Diphenyl-1-picrylhydrazyl) which is a stable radical and has the maximum
578 absorption at 520 nm. In reaction system, 1 ml DPPH (5 mg/ml, dissolved in ethanol) was
579 mixed with 200 μ l various concentration of purified HSAF or HSAF-Fe complexes (final
580 concentration was 0 μ M, 20 μ M, 40 μ M, 80 μ M and 160 μ M, dissolved in ethanol), and then
581 incubated at room temperature for 0.5 h, 24 h, 72 h and 120 h. The OD_{520} value of samples
582 was determined by spectrophotometer, which was used to calculate the remaining DPPH, by
583 following the formula: $\text{DPPH (\%)} = A/A_0 \times 100\%$, A_0 represents the OD_{520} value of 0 μ M.
584 Ascorbic acid was used as positive control and the OD_{520} value was determined immediately.

585

586 *Extraction of the HSAF-Fe complexes in vivo*

587 WT was incubated into 1 ml LB, 30°C with shaking at 200 rpm for overnight. A fraction (1%)
588 of the cultures was transferred into 50 ml M813m medium containing 500 μ M FeSO_4 , and
589 incubated at 30°C with shaking at 200 rpm for 48 h. After centrifugation (12,000 rpm, 5 min),
590 the precipitate presented as two layers, and the upper layer was collected and extracted with

591 50 ml ethyl acetate (containing 0.3% TFA). The organic phase was dried and re-dissolved in
592 200 μ l methanol. The methanol solutions were used for MS analysis. The experiments were
593 repeated three times.

594

595 *Molecular mechanical and quantum mechanical calculations*

596 The calculations were performed with the quantum chemistry polarizable force field
597 (QuanPol) (52) program and the General Atomic and Molecular Electronic Structure System
598 [GAMESS (53, 54)] package. The MMFF94 force field (55-58) was used in the global search
599 of the most stable molecular structures. In the global search, one million steps (time step size
600 = 1 fs) of molecular dynamic simulation were performed at 700 K, with a geometry
601 optimization at every 1000 steps. Using the MMFF94 identified minimum structures,
602 quantum mechanical density functional theory method B3LYP (59, 60) [with Grimme's
603 empirical dispersion correction version III (61)] was used to refine the molecular geometries.
604 The 6-31G* basis set (62) was used.

605

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613

614 **Conflict of Interest:** the authors have no conflict of interest to declare.

615

616

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- 781
- 782

783 **Table 1.** Bacterial strains used in this study.

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Bacterial strains	Relevant characteristics ^a	Source/reference
<i>Lysobacter</i>		
<i>L. enzymogenes</i> OH11	Wild-type, Kan ^r	CGMCC No. 1978
ΔHSAF	HSAF deficiency strain	(32)

785 ^aKan^r, kanamycin resistant.

786

787 **Table 2.** Primers used in this study.

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Primers	Sequence(5'-3')
OH11-F	CGGGGCCCCATTGGAACGACAGCCTCTT
OH11-R	CCGCTCGAGCGGCAAGACAGGGAAGA
3655-F	CGGGGCCCTTTGGTTGTTCCATCCGA
3655-R	CGGGATCCATCGAGGAGCACGGCATC
OH13-F	CTGCAGGACTTCGAACACA
OH13-R	GATTGACTCCTTGGTGCTC
E-F	ATAACGGAGAACGGAATCG
E-R	ACGCATTACTATCTGACCAA

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794 **Figure Legends**

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796 **Figure 1.** Chemical structure of HSAF and analogs (alteramides) isolated from *Lysobacter*
797 *enzymogenes*.

798 **Figure 2.** Formation of PoTeM-Fe complexes. a) Appearance of cultures of *L. enzymogenes*
799 OH11 wild type (WT) and HSAF non-producer mutant (Δ HSAF) grown in M813m medium
800 containing a different concentration of FeSO₄. b) Appearance of the total crude extract of
801 PoTeM (200 μ l) from cultures (25 ml) of WT and Δ HSAF grown in M813m containing a
802 different concentration of FeSO₄. c) Color change in the solution of FeSO₄ (10 mM, 100 μ l)
803 when added with the total crude extract of PoTeM (1-50 μ l, 2 μ g/ μ l) from WT cultured in
804 M813m containing 10 μ M FeSO₄, with methanol as negative control. I, positive control
805 (crude extract of PoTeM from WT grown in M813m containing 500 μ M FeSO₄). d) Color
806 change in the solution of various iron salts (10 mM, 50 μ l), without (0) or with 50 μ l
807 methanol (II) or with 50 μ l (2 μ g/ μ l) of the total crude extract of PoTeM (III). e) HPLC
808 analysis of the isolated PoTeMs, with or without iron salts added.

809 **Figure 3.** MS analysis of the products of HSAF with FeSO₄ (a), Fe(NH₄)₂(SO₄)₂ (b), FeCl₃ (c)
810 and Fe(NO₃)₃ (d), in absence or presence of the metal chelator EDTA. Standard HSAF gave
811 m/z 513 for [HSAF+H]⁺ and m/z 1025 for [2HSAF+H]⁺. In the mixtures of HSAF and iron
812 salts, the peak at m/z 1079 was coincident with [2HSAF-H+Fe]⁺, whereas the peak at m/z
813 1591 was coincident with [3HSAF-H+Fe]⁺.

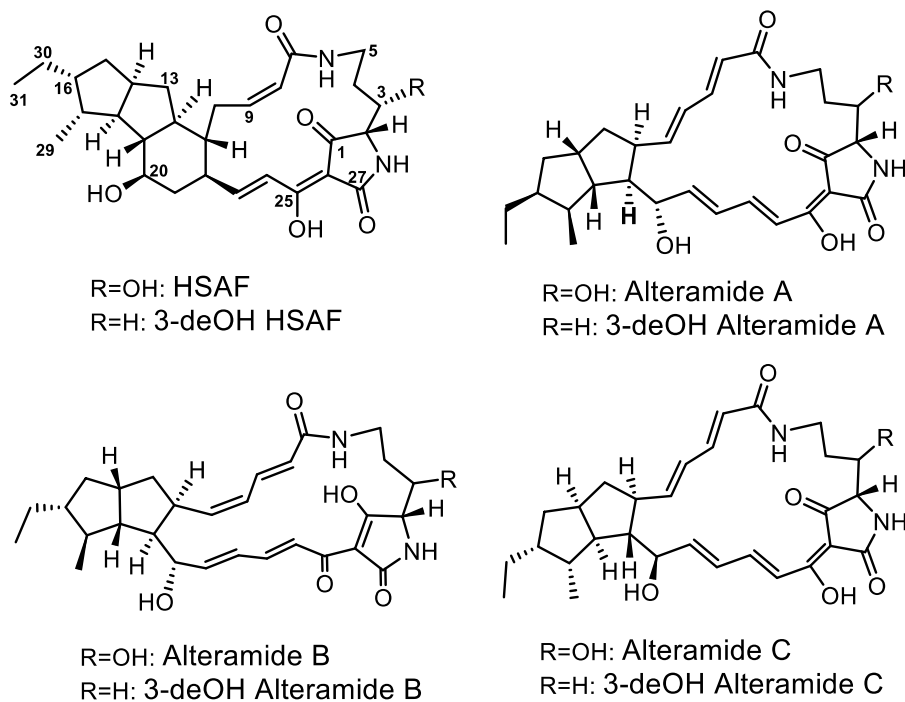
814 **Figure 4.** The *in vitro* antioxidant activity and the *in vivo* protective effect of HSAF for *L.*
815 *enzymogenes* grown in the presence of H₂O₂. a) *In vitro* deoxy-D-ribose degradation assay for

816 the antioxidant activity of HSAF. The activity is presented as thiobarbituric acid reactive
817 species (TBARS) levels relative to the control (100% = TBARS of the control reaction
818 mixture without HSAF). b) *In vitro* assay of the Fenton reaction-caused DNA fragment
819 degradation and the protective effect of HSAF. c-e) *In vivo* H₂O₂ sensitive assay of the WT
820 and Δ HSAF strains cultured in M813m medium containing 0 (c), 80 (d), or 800 μ M (e) H₂O₂,
821 and the OD₆₀₀ value was determined every 24 h. Data are presented as averages of three
822 independent experiments each conducted in triplicate. *, $P < 0.05$; **, $P < 0.01$.

823 **Figure 5.** The protective effect of HSAF for *L. enzymogenes* exposed to UV light and ROS
824 level in *L. enzymogenes* under oxidative stress. a) The WT and Δ HSAF strains were exposed
825 to UV light for 0, 10, 30, and 60 s, and the numbers of colonies on each plate were counted
826 after 72 h of incubation at 30°C. b) Rescue of the UV-light sensitivity of Δ HSAF strain by
827 exogenous HSAF. Δ HSAF strain was added with a variable amount of HSAF (0, 20, 80, 160
828 μ M) and then exposed to UV light for 60 s. c) ROS level in the WT and Δ HSAF strains
829 cultured in different media. No Fe, M813 minimal medium without FeSO₄; M813m, M813
830 modified medium containing 10 μ M FeSO₄; High Fe, M813 modified medium containing
831 500 μ M FeSO₄. d) ROS level in the WT and Δ HSAF strains treated with 40 mM H₂O₂. e)
832 ROS level in the WT and Δ HSAF strains treated with UV light for 60 s. Methanol was used
833 as control. Data are presented as averages of three independent experiments each conducted
834 in triplicate. *, $P < 0.05$; **, $P < 0.01$.

835 **Figure 6.** Molecular structure of HSAF-Fe complexes obtained from the molecular
836 mechanical force field method and quantum mechanical method. Two molecules (a) or three
837 molecules (b) of HSAF can chelate one iron ion.

838 **Figure 1.**



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841 **Figure 1.** Chemical structure of HSAF and analogs (alteramides) isolated from *Lysobacter*

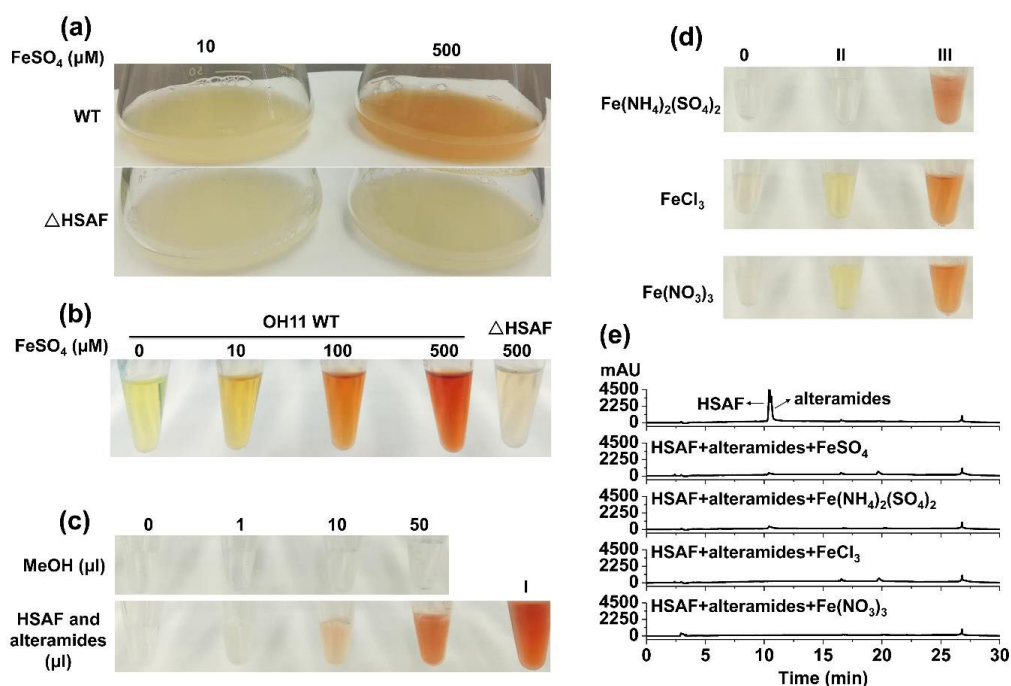
842 *enzymogenes*.

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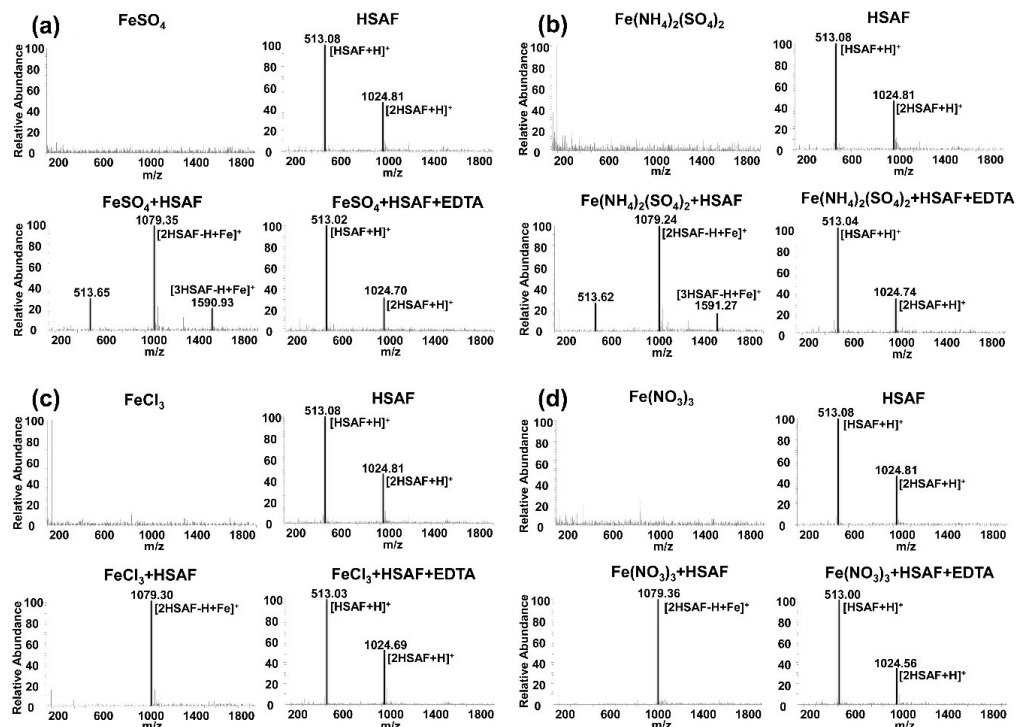
847 **Figure 2.**

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849 **Figure 2.** Formation of PoTeM-Fe complexes. a) Appearance of cultures of *L. enzymogenes*850 OH11 wild type (WT) and HSAF non-producer mutant (Δ HSAF) grown in M813m medium851 containing a different concentration of FeSO_4 . b) Appearance of the total crude extract of852 PoTeM (200 μ l) from cultures (25 ml) of WT and Δ HSAF grown in M813m containing a853 different concentration of FeSO_4 . c) Color change in the solution of FeSO_4 (10 mM, 100 μ l)854 when added with the total crude extract of PoTeM (1-50 μ l, 2 μ g/ μ l) from WT cultured in855 M813m containing 10 μ M FeSO_4 , with methanol as negative control. I, positive control856 (crude extract of PoTeM from the WT grown in M813m containing 500 μ M FeSO_4). d) Color857 change in the solution of various iron salts (10 mM, 50 μ l), without (0) or with 50 μ l858 methanol (II) or with 50 μ l (2 μ g/ μ l) of the total crude extract of PoTeM (III). e) HPLC

859 analysis of the isolated PoTeMs, with or without iron salts added.

860

861 **Figure 3.**

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863

864 **Figure 3.** MS analysis of the products of HSAF with FeSO₄ (a), Fe(NH₄)₂(SO₄)₂ (b), FeCl₃ (c)865 and Fe(NO₃)₃ (d), in absence or presence of the metal chelator EDTA. Standard HSAF gave866 *m/z* 513 for [HSAF+H]⁺ and *m/z* 1025 for [2HSAF+H]⁺. In the mixtures of HSAF and iron867 salts, the peak at *m/z* 1079 was coincident with [2HSAF-H+Fe]⁺, whereas the peak at *m/z*868 1591 was coincident with [3HSAF-H+Fe]⁺.

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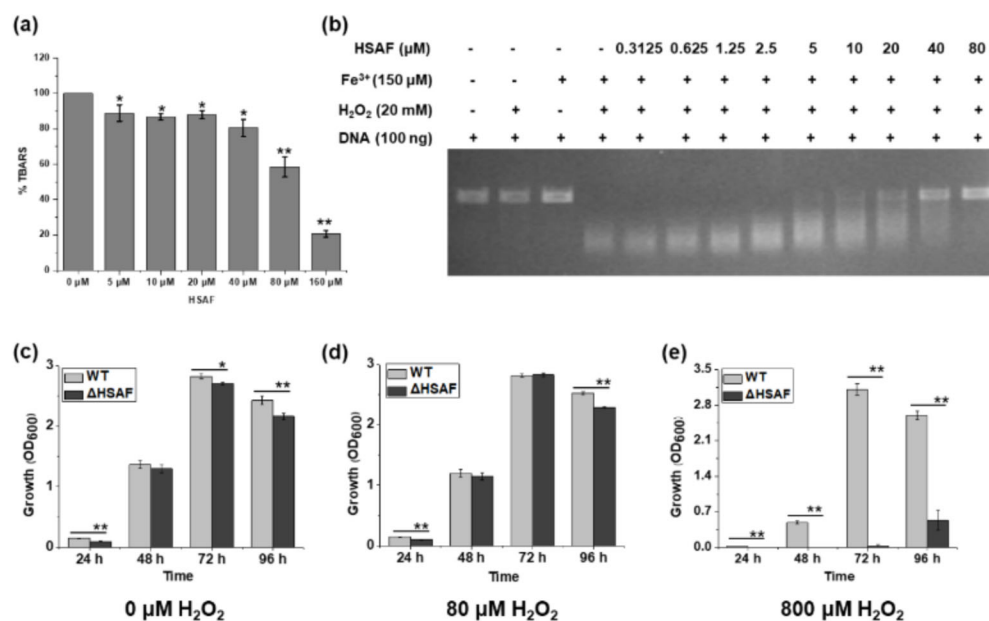
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875 **Figure 4.**

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878 **Figure 4.** The *in vitro* antioxidant activity and the *in vivo* protective effect of HSAF for *L.*879 *enzymogenes* grown in the presence of H₂O₂. a) *In vitro* deoxy-D-ribose degradation assay for

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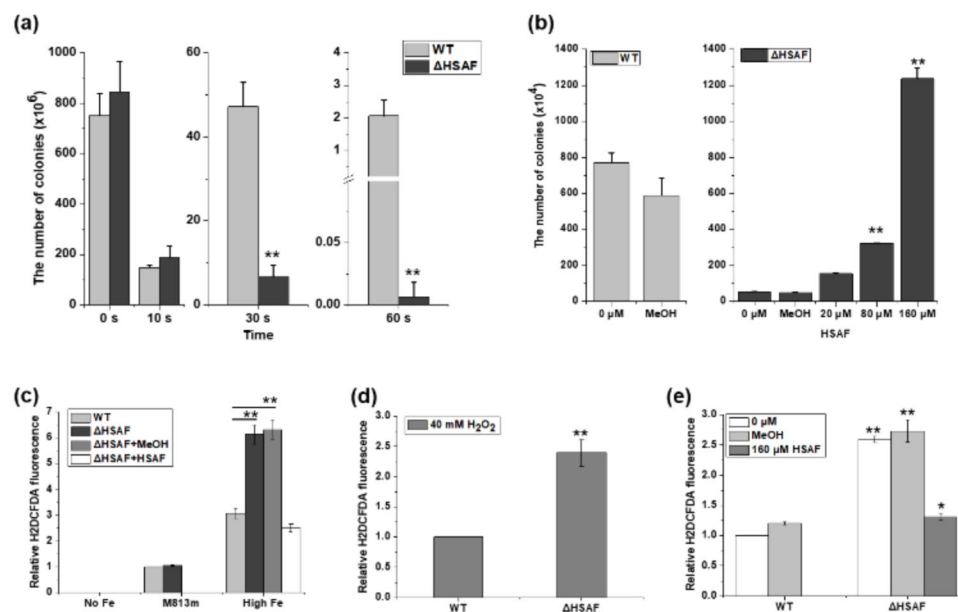
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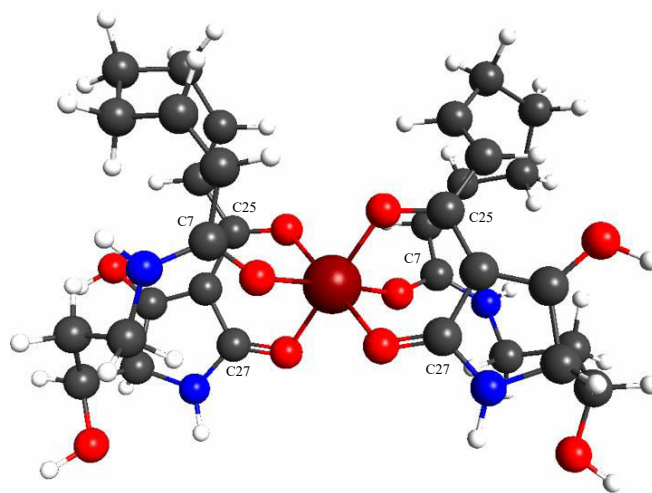
889 **Figure 5**
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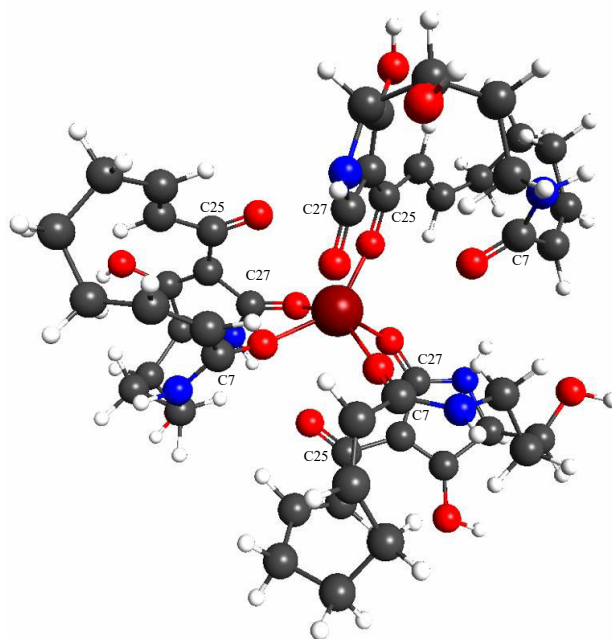
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902 ROS level in the WT and Δ HSAF strains treated with UV light for 60 s. Methanol was used
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905 **Figure 6.**
906 a)



907
908 b)



909
910 **Figure 6.** Molecular structure of HSAF-Fe complexes obtained from the molecular
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