

1 Original Research Article
2 Antibacterial Screening and Analysis of
3 *Streptomyces coelicolor* Secondary Metabolites

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Abstract

Aims: The widespread usage of commercially accessible antibacterial agents has resulted in the emergence of multidrug-resistant bacterial pathogens across the world. Since Streptomyces produce a variety of antibacterial secondary metabolites, the aim of this study was to look into the antibacterial potential of *Streptomyces coelicolor* against some pathogenic bacteria.

Study design: The bacterial strains of *S. coelicolor* by conducting a study of phenotyping and genotyping for these strains, then extracting the crude extract and actinorhodin using Thin Layer Chromatography technique, and antibacterial assessments.

Place and Duration of Study: Department Medical Microbiology, Science and Health research center, University of Koya, between April 2009 and December 2020.

Methodology: All genomic DNA of strains were extracted and genotypic confirmed by using specific primers, then extracted of crude extract and purified actinorhodin by Thin Layer Chromatography (TLC) and ethyl acetate of Trypticase soy agar plates contained *S. coelicolor*. Crude extract and purified actinorhodin were tested against standard strains of *Staphylococcus aureus* (ATCC 25923), *Streptococcus pyogenes* (ATCC 19165), *E. coli* (ATCC 25218), and *Salmonella typhi* (ATCC 14028), also disc diffusion and minimum inhibitory concentration (MIC) techniques were used to serve this purpose.

Results: The crude extract and actinorhodin have an effect as an inhibitor against *S. pyogenes* & *S. aureus* that rang between (10-27 mm), (9-24 mm), while less effect against *E. coli* & *S. typhi*. The crude extract and actinorhodin exhibited potent activity against *S. aureus* & *S. pyogenes* that rang MIC values of (8-32µg/ml), while it showed less activity against *S. typhi* & *E. coli* with rang of (64-128 µg/ml). The mean growth inhibition values were most significant with (P=.05) outcome.

Conclusion: According to the findings of this study, the extracts be a viable option for generating novel antibiotics to eliminate gram-positive pathogenic bacteria.

8 *Keywords:* [crude extract; *Streptomyces coelicolor*; TLC; actinorhodin; MIC]

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10 **1. Introduction**

11 The *Streptomycetaceae* family which belongs to the *Actinobacteria* phylum has shown to
12 be a great source of bioactive small molecules and medicines [1], it is the most abundant
13 drug-producing family in its phylum [2]. In this family, the *Streptomyces* genus has

14 demonstrated a remarkable ability to produce secondary metabolites, especially antibiotics
15 that have therapeutic potential in humans. Its species parvome has been linked to a variety
16 of biological activities such as inhibiting other microorganisms' growth and/ or killing them
17 [3]. Among these species, the model organism *S. coelicolor* has attracted the interest of
18 researchers due to its ability to produce 20 secondary metabolites, 30% of the compounds
19 belongs to one of the well characterised natural substance. The remaining 70% that called
20 cryptic compounds are not produced in laboratory conditions [4]. *Streptomyces* goes through
21 a cellular development process that is similar to that of fungi [5]. Their development begins
22 with spores that germinate and evolve into a vegetative mycelium of branching hyphae. Most
23 secondary metabolites are generated during the subsequent formation of aerial hyphae,
24 which is thought to be a cell response to nutrition deficiency [6].

25 Most of the recognized antibiotics in today's arsenal were discovered between 1940 and
26 1980 including tetracyclines, chloramphenicol, vancomycin, daptomycin, and many more [7].
27 These chemical signalling molecules/ antibiotics were identified through huge screens of
28 tens of thousands of *Streptomyces* strains undertaken either in academia or industrial [8-9].
29 The majority of this research was done on culture supernatants to see if they might suppress
30 the development of pathogenic bacteria such as *Staphylococcus aureus* and *Mycobacterium*
31 *tuberculosis*.

32 The emergence of multi-drug resistance among human pathogens has pushed the mining
33 for novel antibiotics from existent well-known ancient microorganisms which is *Streptomyces*
34 [10]. *S. coelicolor* can serve this purpose based on its ability to produce two different
35 pigmented secondary metabolites which are actinorhodin with blue pigment and
36 undecylprodigiosin with red pigment in addition to a wide variety of therapeutic compounds
37 such as anti-viral, anti-cancer, immune sensitive modulators, herbicides, insecticides, and
38 anti-parasitic [11-12-13].

39 The production of this individual derivative is encoded by a cluster with one or more
40 transcription-regulating genes, in the case of actinorhodin (ACT) biosynthesis, AtrA has been
41 identified as ACT activator beside several other proposed transcription factors that can bind
42 to the *actII-ORF4* promoter region and regulate ACT biosynthesis [14]. Since the most
43 promising types of antibiotics in the future tend to be microbial natural metabolites, the main
44 goal of this work is to investigate the antimicrobial activity of secondary metabolite extracted
45 from *S. coelicolor* strain L646 that has an extra copy of the global regulon *atrA* against some
46 pathogenic bacteria include *Staphylococcus aureus* (ATCC 25923), *Streptococcus pyogenes*
47 (ATCC 19165), *Escherichia coli* (ATCC 25218) and *Salmonella typhi* (ATCC 14028).

48 49 **2. Materials and Methods**

50 **2.1 Bacterial strains and culture conditions**

51 Bacterial strains that were used in this study are shown in Table 1. *Staphylococcus*
52 *aureus*, *Streptococcus pyogenes*, *Escherichia coli*, *Salmonella typhi* and *Pseudomonas*
53 *aeruginosa* that were used for susceptibility testing were part of a culture collection
54 belonging to the Faculty of Science and Health Research Centre (FSHRC) at Koya
55 University. *Streptomyces* strains were kindly provided by Dr Kenneth MacDowall (University
56 of Leeds/ UK). The bacterial strains, unless otherwise stated, were cultured in/ on Mueller
57 Hinton broth/ agar (MHB or MHA) (Oxoid Ltd, Cambridge, UK) for 24 h at 37°C for antibiotic
58 sensitivity and antibacterial activity tests. All *S. coelicolor* strains were cultured in a 2XYT
59 medium, incubated at 30°C for 6 days [15]. Always *Streptomyces* strains broth cultures were
60 incubated with shaking at 220 rpm in 250 mL Erlenmeyer flasks containing 50 ml of media
61 and fitted with a spring baffle to aid dispersed growth of the mycelia.

62
63 **Table 1: Bacterial strains used in this study**

Strain	Description	Reference/ Source
<i>Staphylococcus aureus</i>	ATCC: 25923	FSHRC collection
<i>Streptococcus pyogenes</i>	ATCC: 19165	FSHRC collection
<i>Escherichia coli</i>	ATCC: 25218	FSHRC collection
<i>Salmonella typhi</i>	ATCC: 14 028	FSHRC collection
<i>Streptomyces coelicolor</i> M145	SCP1-, SCP2-. Contains a mutation of the <i>sre-I</i> gene. Wild type	[15]
<i>S. coelicolor</i> L646	<i>S. coelicolor</i> M145 contains an integrating plasmid overexpressing wild-type <i>atrA</i> , which leads to overproduction of actinorhodin.	[16]
<i>S. coelicolor</i> L645	<i>S. coelicolor</i> M145 disrupted <i>atrA</i>	[17]
<i>S. coelicolor</i> M511	it is $\Delta actII-ORF4$	[18]
<i>S. coelicolor</i> M1145	it is Δact , Δred , Δcpk , and Δcda	[19]

64

65 2.2 Preparation of *S. coelicolor* spore stocks

66 *S. coelicolor* spore stocks were generated according to Jalal [20].

67 2.3 *Streptomyces coelicolor* strains integrity

68

69 2.3.1 Phenotyping

70 To investigate the cultural characteristics of the *S. coelicolor* strains, 1×10^5 spore
71 suspensions from M145, L646, L645, M511, and M1146 strains were spotted on tryptone
72 soya agar (TSA) plates, then kept at 28°C for 6 days, and the colour changes were observed
73 during that time. When the bacterial strains were grown as circular patches (14-16 mm
74 diameter) on TSA media, their phenotypes were compared to each other. Starting on the
75 second day of incubation, the plates were imaged every 24 hours.

76

77 2.3.2 Genotyping

78 2.3.2.1 Preparation of Genomic DNA and purification

79 Genomic DNA from *S. coelicolor* strains was extracted after 5 days of incubation in a
80 YEME medium containing 10 % sucrose according to [21]. The quantity and quality of the
81 extracted DNA were measured by Nanodrop (Thermo Scientific NanoDrop 2000. SN. 6113)
82 then stored at -20°C until the time of use.

83 2.3.2.2 Polymerase Chain Reaction (PCR)

84 To confirm *S. coelicolor* strains' identity, three different sets of primers were designed to
85 target a unique specific region of the certain genomic strain (Table 2). PCR reactions were
86 carried out in a T100™ Thermal Cycler, BIO-RAD (621BR11592, Singapore) using Q5® High-
87 Fidelity 2X Master Mix PCR Kit. The thermal cycling conditions consisted of initial
88 denaturation at 98°C for 5 min followed by 30 cycles of denaturation at 98°C for 10 seconds,

89 annealing for 30 seconds at various temperatures, and extension at 72°C for 30 seconds.

90 Then the final extension was performed at 72°C for 5 minutes.

91 **Table 2. Represents sequences, and size of primes used in this study and the name**
92 **of their target genes.**

93

Primer Name	Sequence 5' >>>>>> 3'	Target gene	Product Size (bp)	Description
atrA –ACT gene-for	GGACTCCTCGGACACCAG	<i>atrA</i>	281 bp	differentiate L645 and M1145 strains from the others
atrA –ACT gene-rev	GATAGCCGAGGAGGAGACG	(SCO4118)		
red-prodigiosin-for	AGTTCCTTCGACCGACGTTTC	<i>redZ</i>	248 bp	differentiate M1145 strain from the others
red-prodigiosin-rev	ACGACATGAAAGTGCAGGTG	(SCO5881)		
<i>actII</i> /ORF4 - for	GATTCAACTTATTGGGACGTG	<i>actII</i> -ORF4	746 bp	differentiate M511 strain from the others
<i>actII</i> /ORF4 - rev	CCGTTGAGAATTTCCATGTG	(SCO5085)		

94

95 2.4 Crude Extraction

96 To evaluate the antibacterial activities of *S. coelicolor* crude extract, spores of M145 (wild
97 type), L645 ($\Delta atrA$, the *actII*-ORF4 regulon), and L646 (has an additional copy of *atrA*)
98 strains were spread on TSA plates separately and incubated at 28°C for 6 days that is when
99 the colour of media turned into blue which is an indicator of actinorhodin production. The
100 agar was collected and mixed with 2.5 volumes of dH₂O before being incubated at room
101 temperature for 1 hour with 200 rpm shaking. The supernatant containing diffusible
102 actinorhodin was collected, and acetic acid was added to lower the pH, causing
103 actinorhodin's colour to shift from blue to red.

104 The supernatant was extracted with an equal volume of absolute ethyl acetate (Fisher
105 Scientific) and was separated from the aqueous phase by incubating the mixture for 15
106 minutes at room temperature in a glass separatory funnel. A rotary evaporator (Rotavapor,
107 RE Buchi) was used to evaporate the ethyl acetate, and a parallel evaporator was used to
108 dry the sample (EZ-2 personal evaporator, Genevac) [16]. The crude extract was weighed,
109 then dissolved in 100% (v/v) methanol to a final concentration of 10 mg/mL and kept at 4°C.
110 The crude extract activity was checked as mentioned in sections 2.6.1 and 2.6.2.

111

112 2.5 Actinorhodin (ACT) purification

113 Using a capillary glass tube, aliquots of crude extract dissolved in methanol from strain
114 L646 were spotted on a pencil line about 1 cm from the bottom of a TLC silica gel plate (DC-
115 Alufolien, Kieselgel 60/Kieselgur F254, Merck). Just behind the line where the crude extract
116 loaded, the plate was placed into a TLC tank and submerged in toluene: glacial acetic acid
117 (4:1 [v/v]). The plate was removed once the solvent front reached about 1/3 of the plate, and
118 the solvent front was indicated with a pencil line. The top red band with retention factor (Rf)
119 0.45 mm was taken out and eluted with ethyl acetate after the plate was dry. The ethyl
120 acetate extract was weighed after drying then methanol was used to dissolve the resultant
121 substance. The Rf value was calculated by dividing the distance travelled by the pigment by
122 the distance travelled by the solvent front. ACT activity was verified as indicated in sections
123 2.6.1 and 2.6.2 [22-23].

124 **2.6 Antibacterial activity assessment of the crude extract and ACT**

125 Two alternative methods were used to examine the bioactivity of the crude extract and
126 purified ACT.

127

128 **2.6.1 Disc diffusion assay**

129 Disc diffusion experiment was used to evaluate the antimicrobial activity of dried crude
130 extracts from M145, L646, and L645 strains, as well as purified actinorhodin from the L646
131 strain [24-25]. Prior to the experiment, 6.0 mm sterile Whatman No. 1 discs were prepared
132 then impregnated with the crude extract or purified actinorhodin at the following
133 concentrations: 100g/ml, 200g/ml, 400g/ml, 600g/ml, and 800g/ml. The discs were placed on
134 overnight Muller-Hinton agar (MHA) (Oxiod) plates that were previously inoculated with 0.1
135 ml of each of the following standardized test organisms: *Staphylococcus aureus* ATCC
136 25923, *Streptococcus pyogenes* ATCC 19165, *E. coli* ATCC: 25218, and *Salmonella typhi*
137 ATCC 14028.

138 Standard antibiotic discs were used to serve as positive controls, which were
139 azithromycin (10 µg/ml) and ciprofloxacin (15 µg/ml). Blank discs impregnated with ethanol
140 solvent were used as a negative control. This experiment was done in triplicate. All the
141 inoculated plates were kept in a refrigerator at 4°C for 1 h to allow the diffusion of the
142 extracts in the media then incubated at 37°C for 24 h to detect and measure the inhibition
143 diameter zone around the discs.

144

145 **2.6.2 Standard susceptibility test**

146 The Clinical and Laboratory Standards Institute (CLSI) criteria were used to calculate the
147 Minimum Inhibitory Concentrations (MICs) of L646 crude extracts and proposed actinorhodin
148 against the same set of pathogenic bacteria from section 2.6.1 [26]. MICs were read after
149 18-24 hours of incubation at 37°C. Following incubation, the MIC was defined as the lowest
150 concentration of antibiotic that inhibited all visible growth.

151 MIC determination with appropriate comparator antibiotics was undertaken to permit
152 comparison of the antibacterial activity of the investigated natural products with existing
153 clinical agents against the tester strains. Positive control (growth control) and solvent control
154 (MHB) were included in each experiment. A susceptibility test was conducted on a minimum
155 of three independent occasions to ensure reproducibility. MIC results were read with high
156 accuracy after incubation, for both groups of crude extract and actinorhodin using an ELISA
157 device (ELX800 ABSORBANCE MICROPLATE READER, BioTek, Germany), and with a
158 Micro-plate (96 wells), at a wavelength of 540 nm.

159

160 **2.6.3 Data analysis**

161 All data analysis was conducted by Statistical Package for Graphpad prism version 8 in
162 terms of the mean of the growth inhibition zone value obtained from each of the four
163 bacterial pathogens and two standard antibiotics. The data on L646 crude extract and
164 purified actinorhodin extract were analyzed by comparing the mean growth inhibition zone
165 value in disc diffusion test through analysis of variances (one-sample t-test) while in MIC test
166 was analysed by comparing the mean growth inhibition value (Two-way ANOVA). Results
167 with ($P = .05$) and (P -value summary = ****) were considered as most significant outcome.

168

169 **3. Results and Discussion**

170

171 3.1 Streptomyces coelicolor identification approaches

172 The integrity of *S. coelicolor* strains M145 (the wild type), L645 ($\Delta atrA$), M511
173 ($\Delta actIIOF4$), M1164 (Δact , Δred , Δcpk , and Δcda), and L646 (extra copy of *atrA*) were
174 verified using the classical culture method and at the molecular level as follows:

175 3.1.1 Phenotypic Verification

176 The phenotype of *S. coelicolor* strains was compared to each other, after 6 days of
177 incubation at 28°C on TSA plates. The representative results of a triplicate set are given in
178 figure 1 as patches with 12-15 mm diameter. The first strain that started actinorhodin
179 production (blue colour) on day 3 was L646 as it contains an extra copy of the regulon *atrA*
180 that regulates *actI*-ORF4, which is in turn synthesis the activator of the cluster-situated act
181 biosynthesis to produce actinorhodin [16]. M145 followed L646 in extracellular production of
182 actinorhodin as it contains a single copy of *atrA* [17]. The actinorhodin blue pigment was not
183 seen in the L645 strain that lacks the *atrA* gene. However, it started prodigiosin production
184 on day 3 [18]. Strains M511 and M1146 did not express any colour changing during the six
185 days of incubation that was expected due to lack of *actI*-ORF4 gene and red cluster in both
186 strains [18-19].

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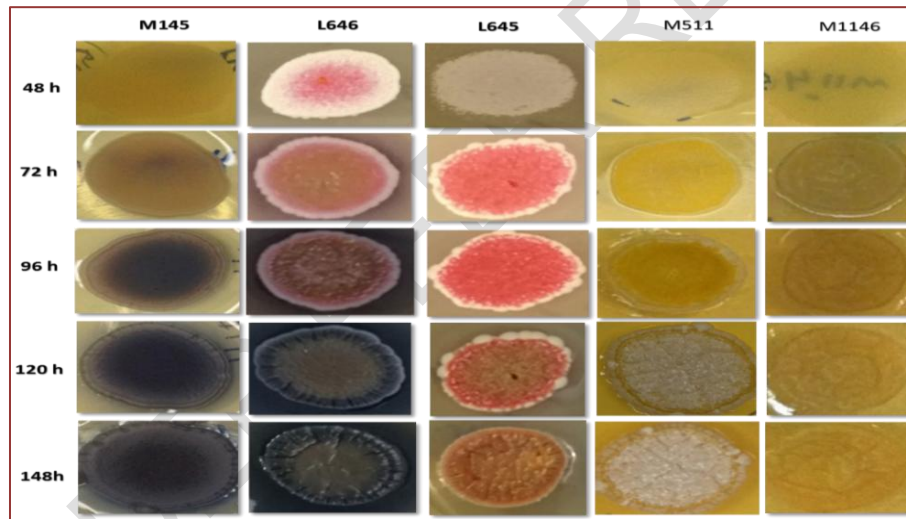
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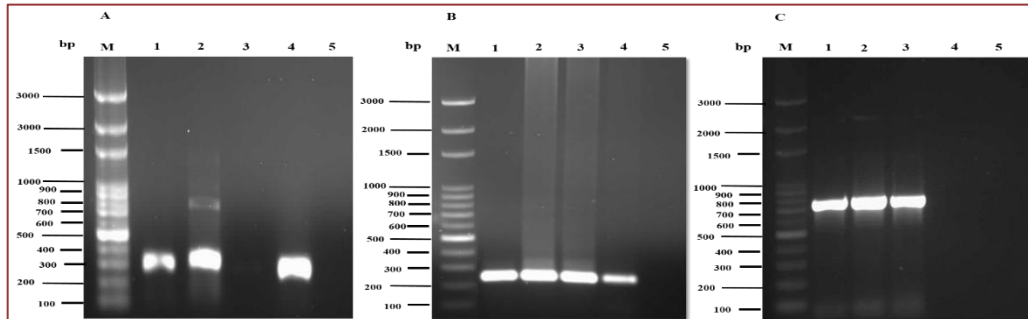
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Figure 1. Phenotypic characteristic of *S. coelicolor* strains on TSA. The strains are identified by the top labelling. The hour's period on the left indicates how long the patches had been incubating before being photographed. At least three separate replicates were used to create these images.

211 3.1.2 Genotypic Confirmation

212 The molecular method of *S. coelicolor* strains identification started with mycelia collection
213 from the cultured strains then subjected to total DNA extraction. The isolated DNA was
214 analysed using PCR to confirm the identity of the strains. The existence of *atrA* was
215 confirmed using the *atrA* –*ACT* gene-for and *atrA* –*ACT* gene-rev primers that target the *atrA*
216 gene in the genome. The expected amplicon of 281 bp was produced for M145, L646, and
217 M511, but not for L645 and M1146 (Figure 2, panel A). The deletion of undecylprodigiosin
218 biosynthetic cluster (RED) was confirmed using *red*-prodigiosin-for and *red*-prodigiosin-rev
219 primers that bind to the chromosomal *redZ* (SCO5881) that activates transcription of *redD*,

220 the final regulator of the RED biosynthetic cluster. An expected 248 bp PCR amplicon was
221 found for all the strains except M1146 (Figure 2, panel A, lane 5). The presence of the *actII*-
222 ORF4 (SCO5085) gene in the *act* cluster was investigated by targeting it using *actII*ORF4 –
223 for and *actII*ORF4 – rev primers. A 746 bp of PCR amplicon was detected for M145, L646,
224 and L645 but not for M511 and M1146 (Figure 2, panel C).
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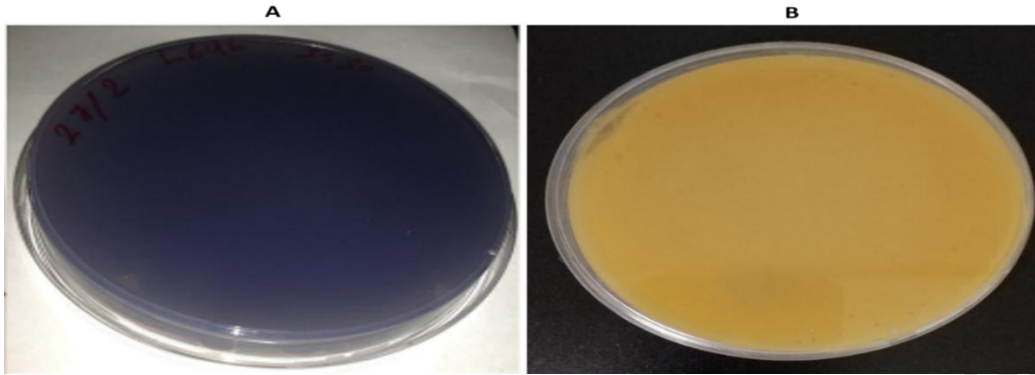
228 **Figure 2. PCR verification of M145, L646, L645, M511, and M1146 strains using (A) *atrA***
229 **–*ACT* gene-for and *atrA* –*ACT* gene-rev primers that target *atrA* gene (SCO4118), the**
230 **strains that contain a copy of *atrA*, a 281 bp amplicon was detected, no PCR product**
231 **was gained for L645 and M1146 strains. (B) *red*-prodigiosin-for and *red*-prodigiosin-rev**
232 **primers that target *redZ* (SCO5881), a 248 bp amplicon was detected for all strains**
233 **except M1146, lane 5. (C) *actII*ORF4 – for and *actII*ORF4 – rev primers that target *actII*-**
234 **ORF4 gene (SCO5085), a 746 bp PCR product was visualised for M145, L646, and**
235 **L645, but not for M511 and M1146. In all panels M: contains 100 bp DNA marker**
236 **(Promega Corporation, USA), lanes 1-5 contains PCR amplicon using a DNA template**
237 **from M145, L646, L645, M511, M1146 DNA, respectively. A 1.4% agarose gel in 1X TAE**
238 **was used.**
239

240 3.2 Chemical molecules Extraction

241 In order to investigate the ability of *S. coelicolor* to produce biologically active
242 compounds, small molecules were extracted from L646 and L645 strains (as discussed
243 above), then tested against some pathogenic bacteria include *Staphylococcus aureus*
244 (ATCC 25923), *Streptococcus pyogenes* (ATCC19165), *Escherichia coli* (ATCC 25218) and
245 *Salmonella typhi* (ATCC14028).

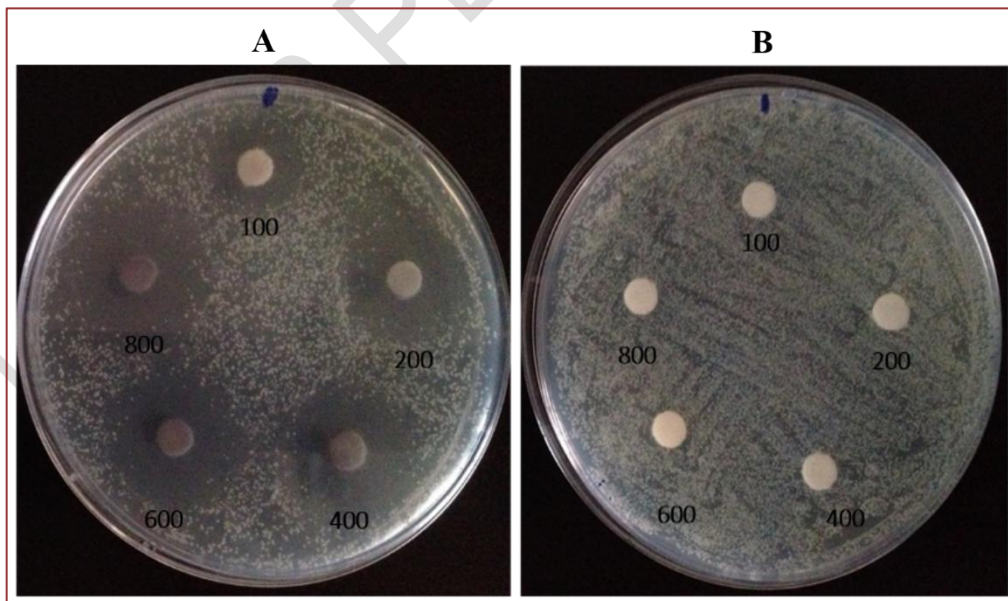
246 3.2.1 Chemical Extraction from TS Agar

247 Chemical extraction contains actinorhodin was isolated from the L646 strain, according to
248 [16] (See section 2.6). L646 strain overproduces actinorhodin as a result of constitutive
249 expression of *actII*-ORF4 from a strong promoter *ermE**p, and ribosome-binding site *tuf* [17].
250 The extraction proceeded after 6 days of incubation at 28°C when the colour of the TSA
251 plates turned blue, which is an indicator of actinorhodin production (Figure 3, panel A). Same
252 culturing conditions and crude extraction were applied to the *S. coelicolor* L645 strain to be
253 used as a control. Since the L645 strain lacks the *actII*-ORF4 gene, no detectable blue
254 pigment was observed (Figure 3, panel B).
255



256
 257 **Figure 3. TSA agar plates were inoculated with L646 and L645 strains. Panel A shows**
 258 **the production of blue pigment by L646. Panel B shows L645 culture with no change**
 259 **in the colour as an indicator of no actinorhodin production. The plates were incubated**
 260 **for 6 days at 28°C.**
 261

262 The extracted chemicals were concentrated, dried, weighed, and then re-suspended in
 263 methanol at a known concentration. A pilot study was set to evaluate the validity of the
 264 extracted chemicals before proceeding to any further investigation. Different quantities of the
 265 crude extract were prepared, dried and suspended with methanol then spotted at certain
 266 concentrations onto 6 mm diameter filter papers, which in turn they placed on the surface of
 267 MHA agar plates that have been cultured with *Staphylococcus aureus* (ATCC 25923) and
 268 incubated at 37°C for 24 h. The results confirmed that the chemical extract that contains
 269 actinorhodin from L646 has the ability to inhibit *Staphylococcus aureus* by generating a clear
 270 zone ranging between (14-25mm) around the discs at 100µg/ ml (Figure 4, panel A). This
 271 activity was absent when a crude extract from the L645 strain was used in the experiment
 272 (Figure 4, panel B) since L645 does not possess the *atrA* gene that activates actinorhodin
 273 production.
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 277 **Figure 4. Shows the impact of crude extract from L646 and L645 against**
 278 ***Staphylococcus aureus* (ATCC 25923). Panel (A) represents the effect of the crude**

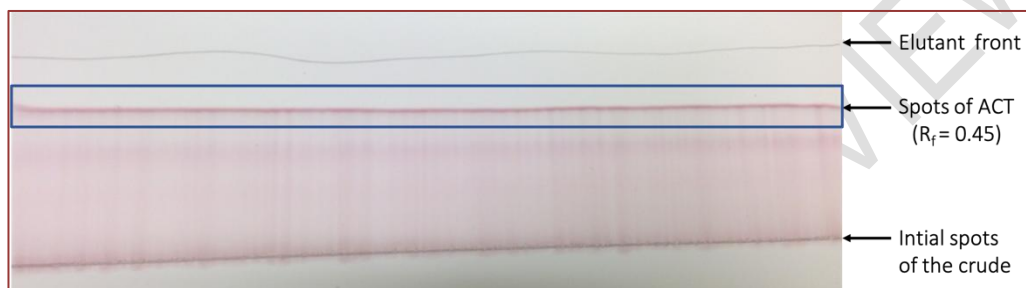
279 compound from L646 that shows a clear zone around the tested discs. Panel (B)
280 represents a crude compound from *S. coelicolor* L645 with no effect. 100, 200, 400,
281 600, 800 refer to the used concentrations as $\mu\text{g/ml}$ or what) of the crude extract on
282 MHA medium. All the plates were incubated overnight at 37°C .

283

284 **3.2.2 Thin-layer chromatography (TLC) for actinorhodin (ACT) purification**

285 As an insight was obtained from the previous experiment (Section 3.2.1), which was the
286 inhibition ability of actinorhodin containing crude, we attended to prepare quantities of the
287 active compound that we believe is actinorhodin based on a previous study [27]. Thin-layer
288 chromatography (TLC) plate with toluene: acetic acid [v / v (4: 1)] as a solvent was used for
289 this purpose (see Section 2.5). The proposed actinorhodin was collected at a retention factor
290 (R_f) of 0.45 mm [28] (Figure 5). The proposed actinorhodin was concentrated, air-dried,
291 weighed, and then re-suspended in methanol at a known concentration.
292

292



293

294 **Figure 5. Represents the position of the actinorhodin band that has been removed**
295 **from the TLC plate. A blue box indicates the actinorhodin band that has been**
296 **removed from the TLC plate. Toluene: acetic acid [v/v (4:1)] was used as a solvent.**
297 **The R_f value is (0.45) mm.**

298

299 **3.3 Antibacterial activity of the crude extract and proposed actinorhodin**

300 The crude extract and suggested actinorhodin (ACT) from the L646 strain were used
301 against the mentioned set of pathogenic gram-positive and gram-negative bacteria at
302 different concentrations by two methods, which were disc diffusion assay and minimum
303 inhibitory concentration assay.

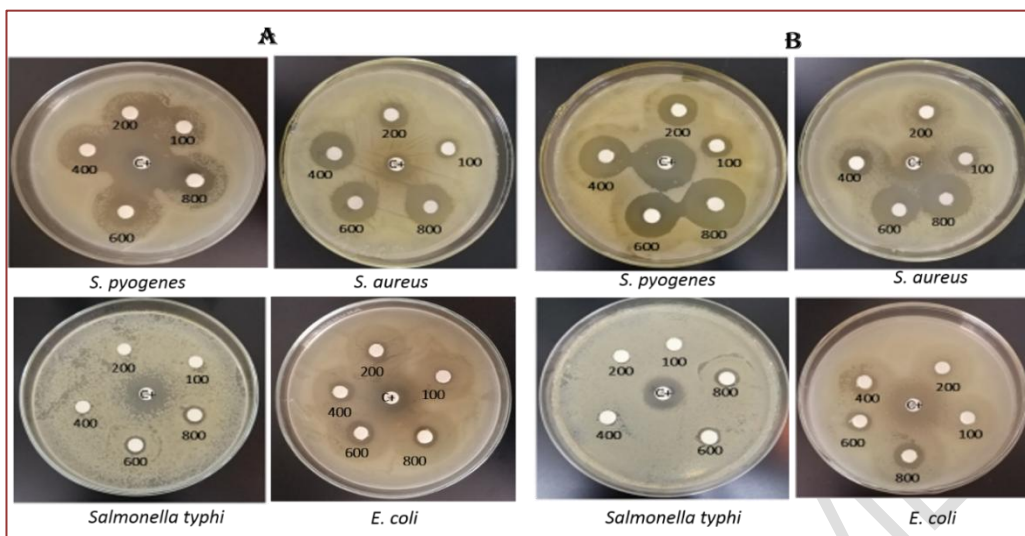
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305 **3.3.1 Disc Diffusion**

306 Disc diffusion assay of crude extract and proposed actinorhodin showed significant
307 differences against some important pathogens such as *Staphylococcus aureus* ATCC
308 25923, *Streptococcus pyogenes* ATCC: 19165, *E. coli* ATCC: 25218, and *Salmonella typhi*
309 ATCC: 14028, on MHA medium. The inhibition zone size around the discs was used as an
310 indication of the efficacy of the tested extracts on bacterial growth. Ciprofloxacin (CIP) and
311 azithromycin (AZM) were used as positive controls at $10 \mu\text{g/ml}$ and $15 \mu\text{g/ml}$, respectively.

312 It was noted that the crude extract has an effect as an inhibitor against *Streptococcus*
313 *pyogenes* and *Staphylococcus aureus* in standard conditions with diameter zone inhibition of
314 (11-23 mm) and (9-20 mm), respectively, while fewer inhibition activities were shown against
315 *E. coli* with diameter zone inhibition of (8-16 mm) and no effect was detected against *S. typhi*
316 (figure 6 panel A). However, actinorhodin showed a reasonable wider effect against them
317 with diameter zone inhibition of (10-27 mm) against *S. pyogenes* and (10-24 mm) against *S.*
318 *aureus*; while less effect against *E. coli* with diameter zone inhibition of (8-17 mm) was
319 detected and no effect was determined against *S. typhi* (figure 6, panel B). In all cases, the
320 proposed actinorhodin was shown considerably more activity against the tested pathogens
321 compared to the crude extract.
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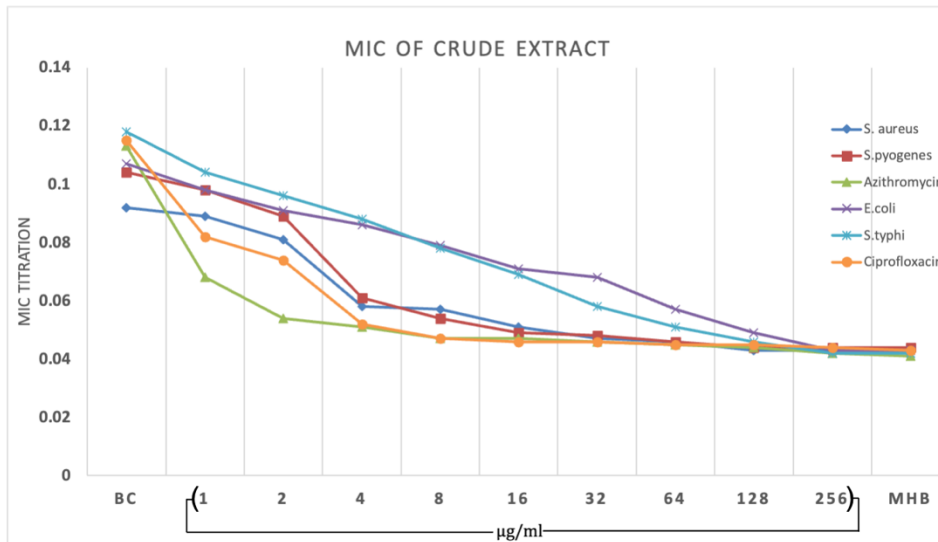
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 324 **Figure 6. Antibacterial activity of the crude extract and purified actinorhodin from *S.***
 325 ***coelicolor* L646 by disc diffusion test. Panel (A) shows the use of crude extract**
 326 **against four pathogenic bacteria at different concentrations (100, 200, 400, 600, 800**
 327 **µg/ ml). In parallel, actinorhodin was used in panel (B) in the same manner and**
 328 **conditions as in panel A. All the tests were performed on MHA medium and the**
 329 **results were taken after 24 h of incubation at 37°C. C⁺ refers to a positive control**
 330 **(ciprofloxacin and Azithromycin).**
 331

332 **3.3.2 Minimum inhibitory concentrations MICs**

333 MICs of the crude extract and purified actinorhodin were determined for several clinically
 334 important Gram-positive and Gram-negative bacteria which included *Staphylococcus*
 335 *aureus*, *Streptococcus pyogenes*, *E. coli*, and *Salmonella typhi* using the standard CLSI
 336 broth microdilution method. The crude extract exhibited potent antibacterial activity against
 337 *Staphylococcus aureus* and *Streptococcus pyogenes*, giving MIC values of 16 µg/ml.
 338 However, it showed less antibacterial activity against *Salmonella typhi* and *E. coli* with MICs
 339 values of (64 and 128 µg/ml), respectively (Figure 7). The results fall within the comparable
 340 range for systemically administrated antibacterial drugs in current clinical use such as
 341 azithromycin and ciprofloxacin, which were given a MIC value of 4 µg/ml against *S. aureus*
 342 and *E. coli*, respectively.

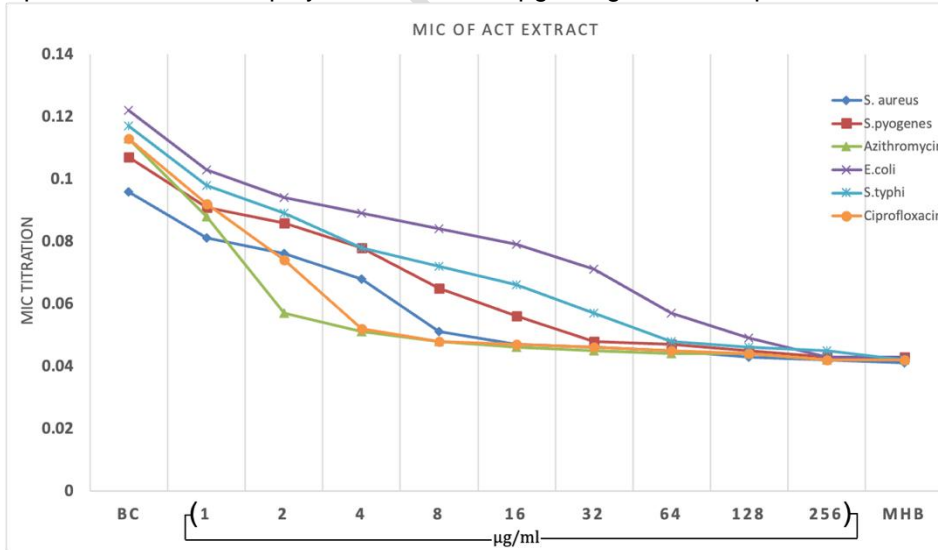


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345 **Figure 7. A statistical chart representing minimum inhibitory concentration (MIC) of**
346 **the crude extract from *S. coelicolor* L646 against *S. aureus*, *S. pyogenes*, *S. typhi* and**
347 ***E. coli* after 18-24 h of incubation at 37°C in MH broth. BC stands for the antibiotic-free**
348 **bacterial culture that is set as positive control and MHB stands for bacterial free**
349 **Mueller Hinton Broth that has been established as a negative control.**

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On the other hand, MICs were determined for the purified actinorhodin against the same group of bacteria following the same conditions and procedures. Actinorhodin expressed notable antibacterial activity against *S. aureus*, *S. pyogenes* and *S. typhi* with MICs of (8, 16 and 32 µg/ml), respectively. However, it gave a MIC value of 128 µg/ml against *E. coli* as shown in figure (8). The above results fall within the comparable range for systemically administrated antibacterial drugs in current clinical use such as chloramphenicol and ciprofloxacin which display MIC values of 4 µg/ml against susceptible *S. aureus* strains.



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359

360 **Figure 8. A statistical chart representing minimum inhibitory concentration (MIC) of**
361 **the purified actinorhodin isolated from *Streptomyces coelicolor* L646 against *S.***
362 ***aureus*, *S. pyogenes*, *S. typhi* and *E. coli* after 18-24 h. BC represents antibiotic-free**

363 **bacterial culture, which was used as positive control and MHB represents bacterial**
364 **free Mueller Hinton Broth, which was used as a negative control.**
365

366 Although secondary metabolite synthesis is not required for *Streptomyces* to survive,
367 these compounds provide an evolutionary advantage over other occupiers since they may
368 be employed as a chemical agent to positively influence other microorganisms. Since
369 *Streptomyces* genus has been discovered to be a reliable source of bioactive compounds
370 with human health applications, much effort is being put into isolating novel
371 pharmacologically active compounds from this genus [29]. As a result, we focused our work
372 on whether *Streptomyces coelicolor* can produce any antibacterial active molecules through
373 their biosynthetic pathways.

374 Our data indicated that the crude extract from *Streptomyces coelicolor* possesses
375 significant antibacterial activity against representative gram-positive bacteria, with MIC
376 values of 16 µg/ml. However, it lacked useful antibacterial efficacy against gram-negative
377 pathogens, with MICs of >128 µg/ml against the tested gram-negative bacteria. It was
378 expected from previous pilot studies by our group that actinorhodin is the active compound
379 that expresses this antibacterial activity. So, we attempted to examine its biological activity
380 after isolating it from the crude extract of *S. coelicolor* strain L646 that overexpresses *atrA*,
381 which results in an increase in actinorhodin production. A prominent, well-separated purple
382 band with R_f 0.45 was obtained when the crude chromatographed on TLC (Figure 5). This
383 proposed actinorhodin showed good efficacy against *S. aureus* with a MIC of 8 µg/ml based
384 on the MIC value for the clinically useful antibiotic azithromycin against the same bacterial
385 genus with a MIC of 4 µg/ml [27]. Against *S. pyogenes* and *E. coli*, the actinorhodin had the
386 same effect as the crude extract. The ACT and the crude extract lacked efficacy against *S.*
387 *typhi* with MICs between 32-64 µg/ml, respectively and *E. coli* with MICs of >128 µg/ml.

388 Compared to the Gram-positive pathogens, both extracts showed less effectiveness against
389 the representative of Gram-negative pathogens that could be due to the architecture
390 differences in their cell wall structure, particularly the peptidoglycan content. Furthermore,
391 gram-negative bacteria have a lipopolysaccharide membrane on the exterior of their cell wall
392 that makes it impenetrable [30-31]. According to [32], actinorhodin's poor efficacy against
393 Gram-negative bacteria is due to its restricted ability to traverse the outer membrane and
394 reach its target site, or it could be due to the action of the existence of substrate specificity
395 range efflux pumps such as AcrAB-TolC [27-33].

396 Our findings are in line with those of several previous studies, such as one by [28], in which
397 they partially purified 1 mg of crude from *Streptomyces ruber* EKH2 using a silica gel plate
398 by TLC technique. Different fractions were obtained with various (R_f) values, however, the
399 one with R_f of 0.45 was more effective against a broad range of medically important bacteria.
400 Another study by [34-35] supported our finding by confirming that the purified actinorhodin
401 from *Streptomyces coeruleorubidus* BTSS-301 by TLC method showed high effectiveness
402 against gram-negative tested pathogens including *Staphylococcus aureus* and *E. coli* using
403 minimum inhibitor concentration (MIC) of the partially purified compound.

404 The disc diffusion experiment results of the crude extract and suggested actinorhodin
405 confirms the MIC findings, which is backed by [36], which found that the extracted and
406 purified material by TLC was effective against the organisms tested in both the disk diffusion
407 and MIC assays equally. The antistaphylococcal activity of actinorhodin demonstrated here
408 is substantially higher than that previously reported for actinorhodin, the original work
409 reported antibacterial activity against *Staphylococcus* with a MIC of 25-30 µg/ml using the
410 agar diffusion method [37]. This obvious difference might be explained by a number of
411 factors. The first factor can be that the MIC value was calculated in the original study using
412 the agar plug diffusion technique. Briefly, the actinorhodin producer bacteria were grown on
413 a complete medium (CM) for 48 h, and then 6 mm diameter colony centred plugs of CM agar
414 were cut and moved into a Petri dish filled with nutrient agar containing *S. aureus*. In
415 contrast, we used broth culture to estimate the MICs values using ELISA. So, the

416 considerable disparity in the MICs achieved is most likely due to the choice of culture
417 medium. The second factor can be returned to the differences in actinorhodin analogues that
418 can express various levels of bioactivity against *S. aureus*.

419 This study demonstrates the value of revisiting an old mine to re-evaluate an excitant natural
420 product (e.g., actinorhodin) in the quest for novel antibacterial reagents. The finding of this
421 study supports such action through proposing actinorhodin as an active future antibiotic
422 against staphylococcal infection, which we believe acts on the bacterial membrane
423 differently from other membrane-perturbing agents used in clinical practice [38].

424 **4. Conclusion**

425 In conclusion, this work has examined the antibacterial activity of *S. coelicolor* secondary
426 metabolites. The remarkable result that we obtained from studying small molecules was the
427 ability of proposed actinorhodin from the L646 strain (figure 8), in which an extra *atrA* gene
428 has been added, to inhibit *S. aureus* and *S. pyogenes* growth with a MIC value of 8 µg/ml.
429 However as yet, the small molecule(s) exact structure is unknown. The next step would be to
430 fractionate a crude extract from the L646 strain and screen the fractions against some
431 human pathogenic bacteria to identify the fraction containing the inhibition activity. Analytical
432 LC-MS may be used to produce a high-quality molecular weight from the most active
433 fraction, which could then be utilized to create a formula. Finally, NMR might be carried out
434 to identify the small molecule's physical and chemical characteristics.

435

436

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438

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444

445 **COMPETING INTERESTS**

446 Authors have declared that no competing interests exist. The products used for this research
447 are commonly and predominantly used products in our area of research and country. There
448 is absolutely no conflict of interest between the authors and producers of the products
449 because we do not intend to use these products as an avenue for any litigation but for the
450 advancement of knowledge. Also, the research was not funded by the producing company
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452

453 **AUTHORS' CONTRIBUTIONS**

454 Author A' and B' designed the study, performed the statistical analysis, wrote the protocol,
455 and wrote the first draft of the manuscript. 'Author A' managed the analyses of the study.
456 'Author B' managed the literature searches... All authors read and approved the final
457 manuscript.

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459 **REFERENCES**

460

- 461 1. Barka EA, Vatsa P, Sanchez L, Gaveau-Vaillant N, Jacquard C, Klenk H-P, Clément C,
462 Ouhdouch Y, van Wezel GP. Taxonomy, physiology, and natural products of
463 Actinobacteria. Microbiol. Mol. Biol. Rev. 2016; 80:1-43.
- 464 2. Challis GL, Hopwood DA. Synergy and contingency as driving forces for the evolution of
465 multiple secondary metabolite production by *Streptomyces* species. Proc. Natl. Acad. Sci.
466 USA 2003; 100:14555-14561.

- 467 3. Metsä-Ketelä M, Salo V, Halo L, Hautala A, Hakala J, Mäntsälä P, Ylihonko K. An efficient
468 approach for screening minimal PKS genes from *Streptomyces*. *FEMS Microbiol. Lett.*
469 1999; 180:1-6. doi: 10.1111/j.1574-6968.1999.tb08770.x.
- 470 4. van Keulen G, Dyson PJ. Production of specialized metabolites by *Streptomyces coelicolor*
471 A3(2). *Adv Appl Microbiol.* 2014; 89: 217-66. doi: 10.1016/B978-0-12-800259-9.00006-8.
- 472 5. Seipke RF, Kaltentpoth M, Hutchings MI. *Streptomyces* as symbionts: an emerging and
473 widespread theme? *FEMS Microbiol Rev.* 2012; 36(4):862-76. doi: 10.1111/j.1574-
474 6976.2011.00313.x.
- 475 6. Sello JK, Buttner MJ. The gene encoding RNase III in *Streptomyces coelicolor* is transcribed
476 during exponential phase and is required for antibiotic production and for proper
477 sporulation. *J Bacteriol.* 2008;190(11):4079-83. doi: 10.1128/JB.01889-07.
- 478 7. Watve MG, Tickoo R, Jog MM, Bhole BD. How many antibiotics are produced by the genus
479 *Streptomyces*? *Arch Microbiol.* 2001;176(5):386-90. doi: 10.1007/s002030100345.
- 480 8. Baltz RH. Marcel Faber Roundtable: is our antibiotic pipeline unproductive because of
481 starvation, constipation or lack of inspiration? *J Ind Microbiol Biotechnol.* 2006; 33(7):507-
482 13. doi: 10.1007/s10295-005-0077-9.
- 483 9. Baltz RH. Renaissance in antibacterial discovery from actinomycetes. *Curr Opin Pharmacol*
484 2008 Oct;8(5):557-63. doi: 10.1016/j.coph.2008.04.008.
- 485 10. Mohammadipanah F; Wink J. Actinobacteria from arid and desert habitats: Diversity and
486 biological activity. *Front. Microbiol.* 2016; 6:1541.
- 487 11. Hopwood DA, Chater KF, Bibb MJ. Genetics of antibiotic production in *Streptomyces*
488 *coelicolor* A3(2), a model streptomycete. *Biotechnology.*1995; 28:65-102.
489 doi:10.1016/b978-0-7506-9095-9.50009-5.
- 490 12. Thompson CJ, Fink D, Nguyen LD. Principles of microbial alchemy: insights from
491 the *Streptomyces coelicolor* genome sequence. *Genome Biol rev.* 2002; 3:1020.
492 doi.org/10.1186/gb-2002-3-7.
- 493 13. Wohlleben W, Mast Y, Stegmann E, Ziemert N. Antibiotic drug discovery. *Microb*
494 *Biotechnol.* 2016; 9(5):541-8. doi: 10.1111/1751-7915.12388.
- 495 14. Bednarz B, Kotowska M, Pawlik KJ. Multi-level regulation of coelimycin synthesis
496 in *Streptomyces coelicolor* A3(2). *Appl Microbiol Biotechnol.* 2019;103: 6423–6434.
497 doi.org/10.1007/s00253-019-09975-w.
- 498 15. Kieser T, Bibb MJ, Buttner MJ, Chater KF, Hopwood DA. *Practical Streptomyces Genetics.*
499 The John Innes Foundation Norwich, 2000.
- 500 16. Towle JE. AtrA-mediated transcriptional regulation in *Streptomyces* secondary metabolite
501 production and development. In: Faculty of Biological Sciences. Leeds: University of Leeds.
502 2007, pp. 208.
- 503 17. Uguru GC, Stephens KE, Stead JA, Towle JE, Baumberg S, McDowall KJ. Transcriptional
504 activation of the pathway-specific regulator of the actinorhodin biosynthetic genes in
505 *Streptomyces coelicolor*. *Mol Microbiol.* 2005; 58(1):131-50. doi: 10.1111/j.1365-2958.
- 506 18. Floriano B, Bibb M. *afsR* is a pleiotropic but conditionally required regulatory gene for
507 antibiotic production in *Streptomyces coelicolor* A3(2). *Mol Microbiol.* 1996;21(2):385-96.
508 doi: 10.1046/j.1365-2958.
- 509 19. Gomez-Escribano JP, Bibb MJ. Engineering *Streptomyces coelicolor* for heterologous
510 expression of secondary metabolite gene clusters. *Microb Biotechnol.* 2011; 4(2):207-15.
511 doi: 10.1111/j.1751-7915.
- 512 20. Jalal BK, Hasan AH. Molecular and Phenotypic Characterization of Novel *Streptomyces*
513 Species Isolated from Kurdistan Soil and its Antibacterial Activity Against Human
514 Pathogens. *Jordan J. Biological Sciences.* 2021; 14(3): 441-451.
- 515 21. Kim SH, Traag BA, Hasan AH, McDowall KJ, Kim BG, van Wezel GP. Transcriptional
516 analysis of the cell division-related *ssg* genes in *Streptomyces coelicolor* reveals direct
517 control of *ssgR* by AtrA. *Antonie Van Leeuwenhoek.* 2015;108(1): 201-13. doi:
518 10.1007/s10482-015-0479-2.

- 519 22. Fessenden JR. *Organic Laboratory Techniques*. Books/Cole Publishing Company Pacific
520 Grove, California. 1993; pp.189-240.
- 521 23. Selvameenal L, Radhakrishnan M, Balagurunathan R. Antibiotic pigment from desert soil
522 actinomycetes; biological activity, purification and chemical screening. *Indian J Pharm Sci.*
523 2009; 71(5):499-504. doi: 10.4103/0250-474X.58174.
- 524 24. Hassan MA, El-Naggat MY, Said WY. Physiological factors affecting the production of an
525 antimicrobial substance by *Streptomyces violatus* in batch cultures. *Egyptian Journal of*
526 *Biology*. 2001; 3:1-10.
- 527 25. Ningthoujam D S, Sanasam S, Nimaichand S. Screening of Actinomycete Isolates from
528 Niche Habitats in Manipur for Antibiotic Activity. *American Journal of Biochemistry and*
529 *Biotechnology*. 2009; 5(4): 221-225. doi.org/10.3844/ajbbbsp.2009.221.225
- 530 26. Clinical and Laboratory Standards Institute (CLSI). M07-A10. Methods for dilution
531 antimicrobial susceptibility test for bacteria that grow aerobically; approved standard, 10th
532 ed. CLSI 2015; Wayne PA.
- 533 27. Nass NM, Farooque S, Hind C, Wand ME, Randall CP, Sutton JM, Seipke RF, Rayner CM,
534 O'Neill AJ. Revisiting unexploited antibiotics in search of new antibacterial drug candidates:
535 the case of γ -actinorhodin. *Sci Rep*. 2017; 12, 7(1):17419. doi: [10.1038/s41598-017-17232-](https://doi.org/10.1038/s41598-017-17232-1)
536 [1](https://doi.org/10.1038/s41598-017-17232-1)
- 537 28. Barakat KM, Beltagy EA. Bioactive phthalate from marine *Streptomyces ruber* EKH2
538 against virulent fish pathogens. *Egyptian Journal of Aquatic Research*. 2015; 41: 49-56.
539 DOI:[10.1016/j.ejar.2015.03.006](https://doi.org/10.1016/j.ejar.2015.03.006)
- 540 29. Parte AC. LPSN-List of prokaryotic names with standing in nomenclature (bacterio.net) 20
541 years on. *Int J. Syst. Evol. Microbiol*. 2018; 68: 825-1829.
- 542 30. Nikaido H, Vaara M. Molecular basis of bacterial outer membrane permeability. *Microbiol*
543 *Rev*.1985; 49(1):1-32. doi: 10.1128/mr.49.1.1-32.
- 544 31. Chen L, Todd R, Kiehlbauch J, Walters M, Kallen A. Notes from the Field: Pan-Resistant
545 New Delhi Metallo-Beta-Lactamase-Producing *Klebsiella pneumoniae* - Washoe County,
546 Nevada, 2016. *MMWR Morb Mortal Wkly Rep*. 2017; 13, 66 (1):33. doi:
547 [10.15585/mmwr.mm6601a7](https://doi.org/10.15585/mmwr.mm6601a7).
- 548 32. Sahlan AZ, Dixon RA. Role of the cell envelope in the antibacterial activities of polymyxin B
549 and polymyxin B nonapeptide against *Escherichia coli*. *International Journal of Antimicrobial*
550 *Agents*. 2008; 31(3): 224-227.
- 551 33. Poole K. Outer membranes and efflux: the path to multidrug resistance in Gram-negative
552 bacteria. *Curr Pharm Biotechnol*. 2002; 3(2):77-98. doi: 10.2174/1389201023378454.
- 553 34. Siva KK, Haritha R, Jagan-Mohan YSYV, Ramana T. Screening of Marine Actinobacteria
554 for Antimicrobial Compounds. *Research Journal of Microbiology*. 2011; 6: 385-
555 393. [10.3923/jm.2011.385.393](https://doi.org/10.3923/jm.2011.385.393).
- 556 35. Siva KK, Ramana T. Production, purification and characterization of an antimicrobial
557 compound from marine *Streptomyces coeruleorubidus* BTSS-301. *Research Journal of*
558 *Microbiology*. 2013; 7397 e403.
- 559 36. Houssam M A. Biochemical studies on antibiotic production from *Streptomyces* sp.:
560 Taxonomy fermentation, isolation and biological properties. *Journal Saudi chemical society*,
561 2015;19:12-22. doi.org/10.1016/j.jscs.2011.12.011.
- 562 37. Wright LF, Hopwood DA. Actinorhodin is a chromosomally-determined antibiotic in
563 *Streptomyces coelicolor* A3(2). *J Gen Microbiol*. 1976; 96(2): 289-97. doi:
564 [10.1099/00221287-96-2-289](https://doi.org/10.1099/00221287-96-2-289).
- 565 38. Mak S., Nodwell J.R. Actinorhodin is a redox-active antibiotic with a complex mode of action
566 against Gram-positive cells. *Mol Microbiol*. 2017;106(4): 597-613. doi: 10.1111/mmi.13837.