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Streptomyces: isolation, optimization of culture conditions and extraction of secondary metabolites

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ABSTRACT

The objectives of this study were to isolate and identify *Streptomyces* from soil sediments as well as to optimize cultural growth conditions for maximum antibacterial productivity. A total of fifty soil sediments were collected from Red Sea, Sudan. The soil sediments were pretreated and cultivated on agar medium. Promising *Streptomyces* spp. were isolated by agar overlay method using indicator organisms. Optimization of chemical and physical culture conditions was carried out. The later was judged by assessment of antibacterial activity. Ethyl acetate was used to extract the secondary metabolite compounds. The separation of the active ingredients was performed using both thin layer chromatography (TLC) and gas chromatography-mass spectrometer (GC-MS). The results revealed nine strains of *Streptomyces*. Of them two (PS1 and PS28) isolates exhibited high activity against pathogenic bacteria. The optimum growth conditions were pH 7.5, temperature at 30°C, soyabean concentration 2.5 g/l, incubation period in 7 days, MgSO₄·7H₂O conc. 1g/l and K₂HPO₄ conc. 2.5g/l. TLC test showed three and two fragments from metabolites of PS1 and PS28 respectively, while the GC-MS analysis revealed eight and eleven compounds with antibacterial activity of PS1 and PS28 respectively. It is concluded that marine is promising source of secondary metabolites.

Key Words: *Streptomyces*, Red Sea, antibacterial activity, GC-MS, Sudan.

INTRODUCTION

The increase of bacterial resistance to antibiotics is due to the concurrent usage of existing antibiotics, the search for new potent antibiotic to control these resistant pathogens, new antibiotics are in need and should be developed (Naine *et al.*, 2015). The discovery of antibacterial metabolites from marine sources with a potent activity has been proved. *Streptomyces* spp. have the ability to produce many different biologically active secondary metabolites such as antibacterial (Nandhini and Selvam, 2013). Marine *Streptomyces* have a unique secondary metabolites variety by producing new natural antimicrobials. About 2/3 of the famous antibiotics was produced by the genus *Streptomyces*. And about 75% of useful antibiotics are produced by the *Streptomyces* species (Bhavana *et al.*, 2014). There are about 23,000 recognized secondary metabolites and around 80% of which are produced by *Streptomyces* species. Many pathways are related with secondary metabolites produced by the genus *Streptomyces*, these include antimicrobial, antitumor and enzyme inhibitors, compared to terrestrial species, marine *Streptomyces* are essential sources of unique antibiotics. Therefore, the marine *Streptomyces* are investigated to extract secondary metabolites and there are many secondary metabolites have been screened recently (Vijayakumar *et al.*, 2012). It is commonly acknowledged that new antibiotics are urgently required, and that the most promising sources are natural habitats. The marine environment is largely unexploited source for new antibiotics, in a view of the enormous diversity of microorganisms-producing secondary metabolites (Bull

and Stach, 2007). The production of secondary metabolites from the genus *Streptomyces* can be influenced by optimization of the nutritional requirements and cultural conditions. These conditions play an important role in the production of these secondary metabolites (Al-Hulu, 2013).

MATERIALS AND METHODS

Isolation of *Streptomyces*

Fifty marine soil samples were collected from Port Sudan (Falamango coast area) in the depth vary from 0.5 to 20 meter (Average depth 10.5 meters). The samples were air dried to reduce bacterial and fungal contaminants, and stored at 4°C (Kathiresan *et al.*, 2005).

One gram of soil was suspended in 4 ml of sterile seawater. The suspension was heated for 6 min at 55°C to reduce non-spore forming bacteria. The contents were diluted (1:4) in sterile seawater. Aliquots of 1 ml of the sample was spread onto the isolation media (Starch casein Agar (SCA) and International Streptomyces Project (ISP) No.7 media) were prepared and sterilized at 121°C in 15 lbs pressure for 15 min (Jensen *et al.*, 2005). Rifampicin 5µg/ml and Nystatin 25µg/ml were added to prevent bacterial and fungal growth respectively. Plates were incubated at 30°C for 7 days (Panchagnula and Terli, 2011). Antibacterial activities of the obtained colonies were detected using agar overlay technique (Srinu, 2013).

Identification of *Streptomyces* strain

The DNA was extracted using Jena Bioscience bacterial DNA isolation and purification kit (Jena Bioscience Laboratories, GmbH, Germany) according to the manufacturer instructions, and the 16S rDNA was amplified by PCR using a Maxime PCR PreMix kit (iNtRON, Korea). The primers and the PCR conditions were done according to Malinova (2014).

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Primary Screening of Antibacterial Activity

Antimicrobial activity was determined using agar well diffusion method (Augustine *et al.*, 2005), with the test pathogenic bacteria *Staphylococcus aureus* ATCC 25923 and *Escherichia coli* ATCC 25922. The inhibition zones were recorded and only isolated strains with the broad spectrum activity were selected for more investigations.

Optimization of growth and antimicrobial compounds production

To test the growth and antimicrobial compounds production, (PS1 And PS28) isolates were inoculated into a 250 ml Erlenmeyer flask containing 100 ml Soyabean Casein Digest Medium (Tryptone Soya Broth), and the filtrate was detected tested for antibacterial activity by agar well diffusion method (Jeffrey and Halizah, 2014; Bhavana *et al.*, 2014).

Effect of nitrogen source

Different soybean meal concentration (1.5, 2.5, 3.5 and 5 g/L) were added to the culture medium and incubated for 7 days at 30°C.

Effect of pH

To determine the optimal pH for antimicrobial compounds production, flasks containing 50 ml of broth Soyabean Casein Digest Medium (Tryptone Soya Broth) adjusted to pH values of (6.5, 7, 7.5 and 8). The broth medium was inoculated with *Streptomyces* isolates and the flasks were incubated at 30°C for 7 days on the shaker, and then the supernatants were tested for antimicrobial activity.

Effect of Temperature

Streptomyces isolates were inoculated in optimized medium and incubation at different temperatures ranging from 20, 25, 30 and 40°C for 7 days.

Optimum incubation period

To study the influence of incubation period, the culture was inoculated in growth medium for 10 days. The broth was sequentially tested for antimicrobial disc sensitivity test in 4, 5, 6, 7, 8, 9 and 10 days. Inhibition zones were measured and the optimum period was fixed for other experiments.

Effect of Dark and Light Conditions

Two flasks containing culture media were incubated for 7 days in dark and light conditions.

Effect of MgSO₄·7H₂O concentration

Different concentration of MgSO₄·7H₂O (0.5, 1, 1.5 and 2 g/L) were tested for optimum concentration in culture media.

Effect of K₂HPO₄ concentration

Different concentration of K₂HPO₄ (1, 1.5, 2.5 and 4 g/L) were tested for optimum concentration in culture media.

Fermentation

The two selected strains *Streptomyces* (SP 1 and SP 28) isolates were inoculated into Tryptone soya broth, and incubated at 30°C in a shaker for seven days.

After fermentation filtrate was separated by centrifugation at 5000 rpm for 10 min. and the supernatant was filtrated through Millipore filter (Millipore Millex-HV Hydrophilic PVDF 0.45 µm) (Remya and Vijayakumar,

2008). The filtrate was transferred aseptically into a conical flask and stored at 4°C for further assay.

Extraction

The culture filtrate (200 ml) was extracted 3 times with solvent ethyl acetate. The solvent was added to the filtrate in the ratio of 1:1(v/v) and shaken vigorously for 20 minutes. The ethyl acetate phase that contains antibiotic was separated from the aqueous phase using separating funnel. Ethyl acetate layer was concentrated by evaporating to dryness at 40°C and residue obtained was purified using methanol to give (0.8 g) of brown crude extract (Ahmed 2007). This obtained compound was used to determine antibacterial activity against test pathogenic bacteria by agar-well diffusion method (Rabah *et al.*, 2007).

Secondary Screening of Antibacterial Activity

The crude extract was tested for antimicrobial activity using agar-well diffusion method by spreading 25µl of *S. aureus* and *E. coli* on Muller-Hinton agar, the wells were dug by the help of sterile 6 mm cork-borer and loaded with the 100 µl of crude metabolite.

The plates were left at 4°C for 12-16 hours then incubated overnight at 37°C. Inhibition zone of test microorganisms around the wells were measured (Augustine *et al.*, 2005, Khan and Patel, 2011).

Thin layer chromatography (TLC)

To visualize the number of compounds present in the extract of isolates (PS1 and PS28), thin layer chromatography (TLC) was performed. Aluminum plates pre-coated with silica gel (20×20 cm, 0.25 mm Alugram® SIL G/UV 254, Macherey and Nagel, Duren) and two mobile phases [ethyl acetate : methanol (6:4) and Petroleum ether : Chloroform (1:1)] (Attimarad *et al.*, 2012).

Chromatograms were observed under UV light, fractions of the isolates retention factor R_f value were measured and the ratio calculated.

Identification of bioactive compounds

The antibacterial compounds were identified by using Gas Chromatography-Mass Spectrometer technique (GC-MS) (Joel and Bhimba, 2012). The mass spectrum was recorded by using GC-MS-Shimadzu QP2010 Ultra, Japan.

RESULTS

Among fifty isolates obtained from marine soil sediment an agar overlay technique revealed that, 21 (42%) showed antibacterial activities against both Gram-positive and Gram-negative bacteria. Polymerase chain reaction (PCR) was done to determine the identities of the 21 promising isolates revealed that nine (n=9) (18%) were identified as *Streptomyces* species (PS1, PS5, PS10, PS13, PS20, PS21, PS23, PS24, and PS28). The antibacterial activity of the nine *Streptomyces* revealed that the strains PS1 and PS28 have the highly activity against pathogenic test bacteria. Maximum antibacterial production was observed among several concentration of soyabean meal (figure 1). The antibacterial activity was increasing continuously from 0.5g/L to 2.5g/L of soyabean meal concentration. Further increase in the soyabean meal concentration showed a gradual decrease in the production of antimicrobial compounds. The strains of *Streptomyces* showed high activity at pH with value between 7-7.5 (figure 2), the optimum activity of strains showed highly activity at 30°C, while the activity was decreased at lower temperature (figure 3), the activity of strains against pathogenic

microorganisms was at high peak in the range of 5-7 days of incubation, while it is decreased with other periods (figure 4.), the activity of antibacterial compounds exhibited there is no important difference between dark and light environment (figure 5), optimum concentration of $MgSO_4 \cdot 7H_2O$ required for the production of antimicrobial compounds was 1g/L. Further increase in $MgSO_4 \cdot 7H_2O$ concentration showed a gradual decrease in the production of antimicrobial compounds (figure 6); optimum K_2HPO_4 concentration required for the production of antimicrobial compounds was 2.0g/L. Further increase in the K_2HPO_4 concentration showed a gradual decrease in the production of antimicrobial compounds (figure 7). The two strains (PS1 and PS 28) were selected for the extraction of crude compound by ethyl acetate extraction method. Gummy brown crude extract was obtained. The antibacterial activity against *S. aureus* and *E. coli* by agar well diffusion method of crude extracted is shown in table 1.

The crude extracts by ethyl acetate were separated by TLC plates for the strain PS1 yielded three fractions (A, B and C), while the strain PS28 yielded two fractions (A and B) (figure 8.). The Rf value for the strain PS1=0.7cm, while the value for PS28=0.6cm.

The GC-MS analysis showed the chemical composition of extract of isolate PS1 were 54 compounds, while the isolate PS28 were 96 compounds. From these compounds the isolate PS1 eight bioactive compounds is identify with antimicrobial activity (table 1), while the isolate PS28 eleven compounds identify with antimicrobial activity (table 2).

DISCUSSION

The resistance of numerous pathogenic bacteria to antibiotics necessitate search for new antibacterial agents to fight these pathogens. Secondary metabolites produced by bacteria still interested, due to their complicated chemical structures and highly specific antimicrobial activities. The soil bacteria resembling to the genus *Streptomyces* are rich sources of large number of bioactive natural products; they are widely used as antimicrobials. *Streptomyces* species produce about 75% of useful antibiotics (Kumar *et al.*, 2014). Al-Zahrani, (2007) reported that incubation conditions influence the qualitative production of secondary metabolites by *Streptomyces* species. Kathiresan *et al.*, (2005) reported that ability of bacteria to produce antimicrobial agents can be increased or lost under different culture conditions. Maximum antibiotic production was observed with soyabean meal as a source of nitrogen compared to other compounds. The present study confirmed this fact as soyabean meal was used in this study to influence the production of antibacterial compounds; the pH values affect the cellular metabolisms and biosynthesis of secondary metabolites in *Streptomyces* species (Bhavana *et al.*, 2014). In this study, the pH was found suitable for maximum production of metabolites near the neutral value. Ripa *et al.*, (2009) reported that extreme pH is unfavorable for production of secondary metabolites. Kokare *et al.*, (2007) showed that 28°C is the optimum temperature, while 30°C was the optimum temperature for maximum production of metabolites in this study. Our study showed that incubation period was 7 days. Kathiresan *et al.*, (2005) reported that 5 days of incubation gave the broad activity of secondary metabolites. Ethyl acetate was used extensively to extract secondary metabolites. Ghadin *et al.*, (2008) reported that the extraction of secondary metabolites by ethyl acetate

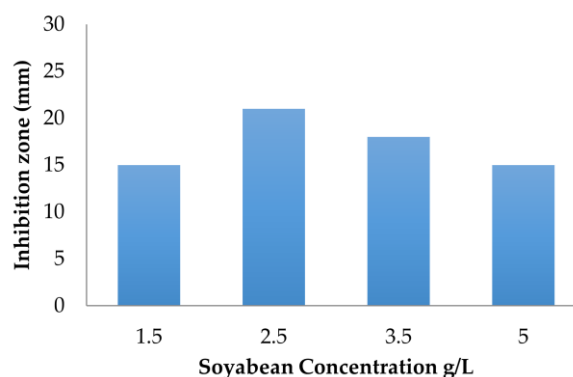


Figure 1: Effect of nitrogen source.

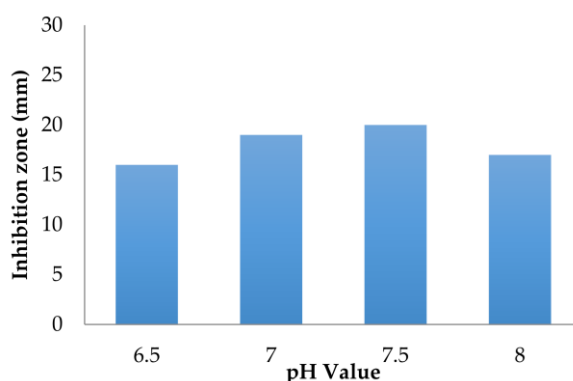


Figure 2: Effect of pH value.

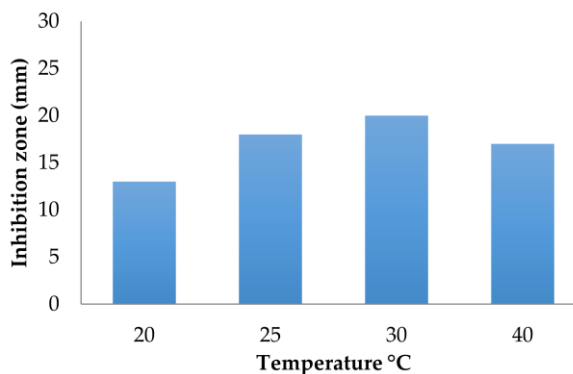


Figure 3: Effect of temperature.

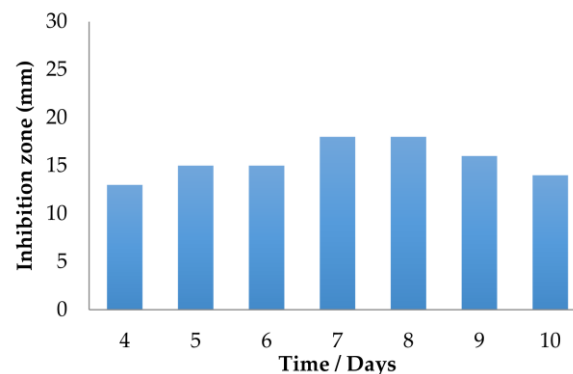


Figure 4: Effect of incubation period.

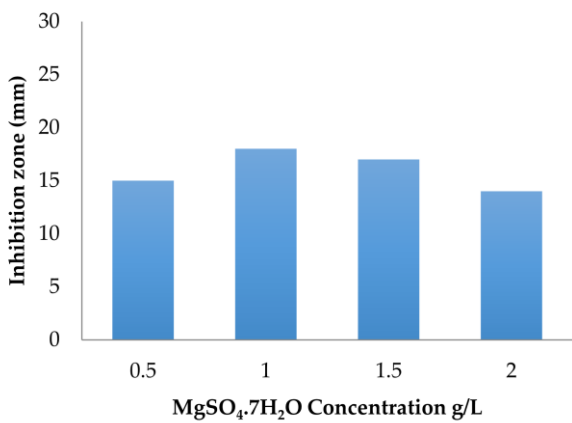


Figure 5: Effect of MgSO₄.7H₂O concentration.

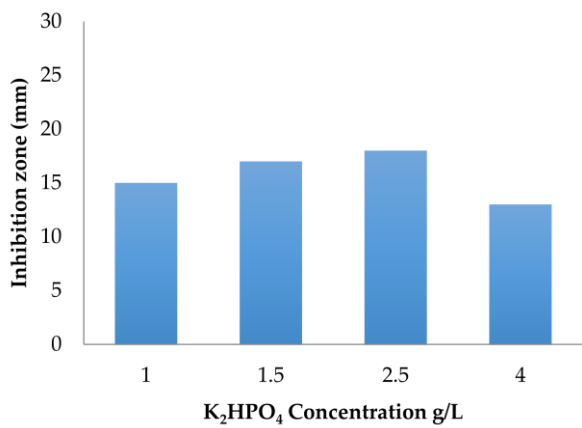


Figure 6: Effect of K₂HPO₄ concentration.

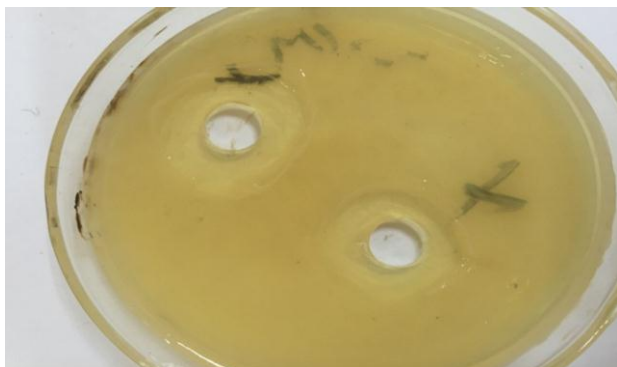


Figure 7: The inhibition zone (mm) of PS1 and PS28 against *S. aureus*.

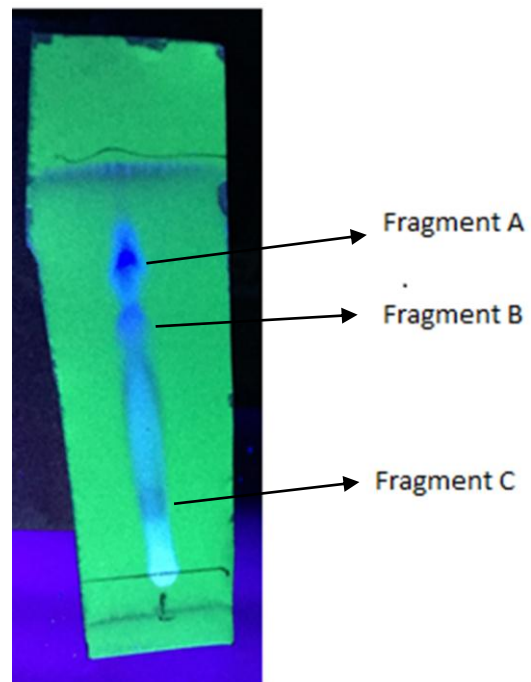


Figure 8: TLC plate of PS1 showing fragments.

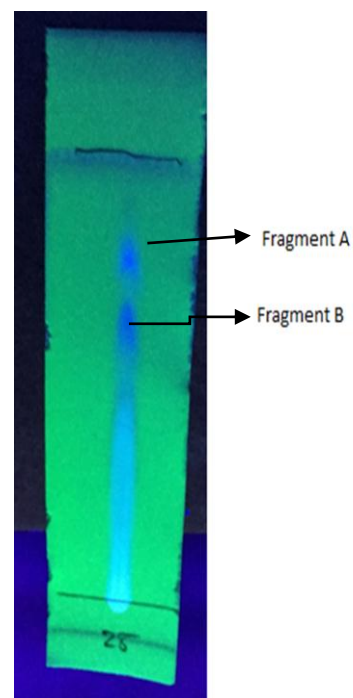


Figure 9: TLC plate of PS28 showing fragments.

Table1: Compounds identify with antimicrobial activities of PS1 isolate.

Compound No.	Compound name	R-Time	Similarity %	Area %	Compound formula	Molecular weight
I	4-Vinyl-imidazole	6.727	96	2.52	C ₅ H ₆ N ₂	94
II	Ornithine	8.595	95	5.2	C ₅ H ₁₂ N ₂ O ₂	132
III	4-Methyleneproline	13.055	96	15.02	C ₆ H ₉ NO ₂	127
IV	Pyrimidinone, 6-amino-2-methyl	15.712	95	0.82	C ₅ H ₆ N ₄ O ₂	154
V	2,6-Dibuty1-4-methylpiperidine	15.879	95	2.70	C ₁₄ H ₂₉ N	211
VI	Actinomycin C2	15.809	99	0.46	C ₆₃ H ₈₈ N ₁₂ O ₁₆	1268
VII	n-3-imidazol-1-yl-propyl-n-4-isopropyl-phenyl-oxalamide	20.024	98	2.14	C ₁₇ H ₂₂ N ₄ O ₂	314
VIII	5-nitroso-2,4,6-triaminopyrimidine	22.817	98	5.79	C ₄ H ₆ N ₆ O	154

Table 2: Compounds identify with antimicrobial activities of PS28 isolate.

Compound No.	Compound name	R-Time	Similarity %	Area %	Compound formula	Molecular weight
IX	pyrrolidine n 3-methyl-3- butenyl	3.870	97	0.67	C ₉ H ₁₇ N	139
X	Piperidine, 2,3-dimethyl	4.875	96	0.14	C ₇ H ₁₅ N	113
XI	Cyclohexanamine, N-2-propenyl	7.793	96	0.17	C ₉ H ₁₇ N	139
XII	alpha-campholenal	8.329	96	1.0	C ₁₀ H ₁₆ O	152
XIII	Norvaline	15.072	99	6.07	C ₂₆ H ₄₇ NO ₄	437
XIV	2-Piperidinone	15.415	98	0.51	C ₁₀ H ₁₆ N ₂ O	180
XV	l-leucine, n-cyclopropylcarbonyl- pentadecyl ester	15.783	99	5.76	C ₂₅ H ₄₇ NO ₃	409
XVI	5,10-diethoxy-2,3,7,8-tetrahydro-1h,6h-dipyrrolo[1,2-a;1,2-d]pyrazine	16.059	97	6.21	C ₁₄ H ₂₂ N ₂ O ₂	250
XVII	n(1) 3-methyl-1,2,4-oxadiazole	17.737	95	0.55	C ₉ H ₁₅ N ₅ O	209
VII	n-3-imidazol-1-yl-propyl-n-4-isopropyl-phenyl-oxalamide	20.062	99	5.65	C ₁₇ H ₂₂ N ₄ O ₂	314
VIII	5-nitroso-2,4,6-triaminopyrimidine	22.914	98	7.23	C ₄ H ₆ N ₆ O	154

showed the obvious activity against pathogenic bacteria than other solvent. Similar result was obtained in this study when used in ratio 1:1. In recent years GC-MS studies have been increasingly applied for the analysis medical used components (Nandagopalan *et al.*, 2015).

In this study the GC-MS analysis showed a variety of compounds from the extract of PS1 and PS28, Jalaluldeen *et al.*, (2015) revealed that 77 compounds were gain from GC-MS analysis, this is higher than PS1 strain compounds and lower than PS28 strain compounds. Many of these compounds are responsible for the antibacterial activity that detected in this study.

CONCLUSION

The study concluded that marine is promising habitat *Streptomyces* spp. and optimizing of culture conditions may influence the production of secondary metabolites. The GC-MS as a new technology is useful for separation components that exist in secondary metabolites.

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