



Removal of ammonia and orthophosphate from domestic wastewater using marine actinomycetes

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ABSTRACT

The present study aims to experiment the efficiency of marine actinomycetes in ammonia and orthophosphates removing from domestic wastewaters collected from El-Gona Wastewater Treatment Station, Hurgada, Red Sea. Ten marine actinomycetes isolates were isolated from sediments in the Suez Gulf and identified as: *Kocuria palustris*, *Streptomyces parvus*, *Streptomyces griseorubens*, *Streptomyces rochei*, *Streptomyces albidoflavus*, *Streptomyces griseus* and four belonged to *Streptomyces* sp based on their 16S rRNA gene sequence analysis. The results demonstrated that, the experimented actinomycetes have differential effectiveness in the domestic wastewater treatment and have significant abilities to remove ammonia and orthophosphates. *Streptomyces griseorubens* and *Streptomyces griseus* are more efficient organisms in ammonia and orthophosphates removal (77.35, 79.02%, respectively) from the raw wastewater relative to the other strains.

INTRODUCTION

The shortage of freshwater resources has become a globally serious problem where the growing population, advanced agricultural practices, industrialization, urbanization and the multiple uses of freshwater have increased the demand for freshwater. Sewage after appropriate treatment and the reuse of the recycled wastewater have become the consensus solution all-over the world (Qin *et al.* 2013).

Actinomycetes have many benefits as a wastewater treatment tool for many purposes ; first, they can use several growth substrates ranging from sugars to polysaccharides, proteins and aromatic compounds of elevated molecular weight (Lemmer and Kroppensted, 1984; Lemmer, 1986), secondly the high resistances of these actinomycetes to desiccation and ultraviolet irradiation and thirdly, their storage capacity for polyphosphates and poly- β -hydroxybutyric acid (Lemmer and Baumann, 1988)

Actinomycetes have recently become the focus of studies in activated sludge and sewage because they are thought to play a significant part in activated sludge bulking and foaming. (Davenport *et al.* 2000; Madoni *et al.* 2000; Heard *et al.* 2008).

Nitrogen and phosphorus removal from the wastewater is the main purpose for the wastewater treatment. Nitrogen is removed by energy and time consuming procedure, which is based on the combination of anaerobic or anoxic denitrification and autotrophic nitrification (Khardenavis *et al.* 2007; Khin and Annachhatre, 2004).

Denitrification is known to proceed as conversion of nitrates to nitrites and subsequent conversion of nitrites to nitric oxide, nitrous oxide and nitrogen gas. Nitrogen appears in wastewater as; ammonia, nitrite, nitrate and organic nitrogen. Part of the organic nitrogen is decomposed to ammonia, which is assimilated to the bacterial cells leading to growth and the other part is oxidized to nitrite and nitrate. Latterly, nitrate is converted to gaseous nitrogen and then removed from the wastewater.

Phosphorus in wastewater was found as orthophosphate, polyphosphate and organic phosphorus, the last two components representing up to 70% of the influent phosphorus. Microbes utilize phosphorus during cell synthesis and energy transport. About 10% to 30% of the influent phosphorus was found to be removed during traditional mechanical and biological treatments (Wenzel and Ekama, 1997; Mulder and Rensink, 1987; Metcalf and Eddy, 1991; Henze, 1996; Sedlak, 1991).

The aim of the present study to experiment the efficiency of actinomycetes in ammonia and orthophosphates removing from domestic wastewaters collected from El-Gona Wastewater Treatment Station, Hurghada, Red Sea.

MATERIALS AND METHODS

Sampling of isolated actinomycetes

Sediments samples were collected according to (Norouzi *et al.* 2018) from four sites along Suez gulf, Egypt using piston corer in clean plastic bags. The samples were immediately transported for bacteriological analysis to the laboratory in an ice box that was always completed within 24 hours.

Isolation of actinomycetes

Approximately 5 g of each sediment sample was added to 15 ml of sterilized seawater, then shaken for 25 min and thermally treated to decrease the amount of non-cell bacteria in favor of actinomycetes. Starch nitrate agar medium plates (Starch, 20; KNO₃, 1; K₂HPO₄, 0.5; MgSO₄.7H₂O, 0.5; FeSO₄.7H₂O, 0.01; agar 20) were inoculated with 1 ml of the samples (Mohanraj and Sekar, 2013). To minimize fungal contamination, the media was supplemented with 75 and 25 ug ml⁻¹ filter sterilized cycloheximide and nystatin (Shrivastava *et al.* 2015). 10 isolates were selected from the plates after incubation and repeatedly streaked to the same medium to obtain pure cultures. The pure isolates were preserved in 20% glycerol at -20 ° C as inventory suspensions. Then 10 isolates were identified as morphologically distinct as; M5, M8, M9, M13, M15, M23, M38, M50, MN and MP then were used for morphological and biochemical identifications.

Morphological and physiological characteristics of isolated actinomycetes:

The chosen prospective actinomycetes isolates have been explored according to morphological, cultural, physiological and biochemical properties (Bensultana *et al.* 2010). Under light microscope, the color of the spore mass was examined and estimated by color chart. Cultural characteristics of actinomycetes isolates were examined by the eye of 14-day-old culture grown on Starch nitrate agar media. The inclined cover slip technique on starch casein agar after 7 days of incubating at 30 ° C observed micro morphology and sporulation under a light microscope. Eyes examined the colors of aerial and substratum mycelia (Li *et al.* 2016). Unless otherwise specified, all physiological tests were performed at 30 ° C and pH 7.0. The efficacy of different temperatures and pH conditions on growth and tolerance of actinomycetes isolates to salt was determined by the use of plates produced with altered starch nitrate agar medium (Akond *et al.* 2016). Utilization of carbon source

(Starch, Lactose, Dextrose, Maltose, glucose, and Glycerol) was examined using mineral salts agar medium containing $g\ l^{-1}$: $(NH_4)_2SO_4$, 2.64; KH_2PO_4 , 2.38; $K_2HPO_4 \cdot 3H_2O$, 5.65; $MgSO_4 \cdot 7H_2O$, 1 and agar, 15, pH adjusted to 7.0. Various biochemical tests performed for the identification of the actinomycetes isolates are as follows: production of Protease, Lipase, Urease, Catalase, Chitinase, Oxidase, Sulphide production and Melanin production (Yu *et al.* 2015).

DNA extraction and 16S rRNA sequencing

The molecular identification of the isolates used was carried out by extracting genomic DNA using the Gene Jet genomic DNA purification kit (Fermentas) genomic DNA extraction protocol. Using Maxima Hot Start PCR Master Mix (Fermentas) Polymerase Chain Reaction (PCR) was used. The PCR thermocycler was programmed as follows: 95 ° C for initial denaturation for 5 min, 30 cycles for 1 min at 95 ° C, 55 ° C for 1 min, 72 ° C for 2 min and a final extension for 10 min at 72 ° C. The PCR mixture contained 25 pmol of each primer, 10 ng chromosomal DNA, 200 mmol/LdNTPs and 2.5 U of Taq Polymerase in 50 μ L of Taq polymerase buffer 10X Standard Taq Reaction Buffer. The PCR Clean-Up of the PCR product was performed using Gene JET™ PCR Purification Kit (Fermentas) at "Sigma Scientific Services Company, Egypt". The GATC Company used ABI 3730xl DNA sequencer with universal primers to sequence the PCR product (16S 27F and 16S 1492R), (5'AGAGTTTGATCCTGGCTCAG-3' and 5'-GGTTACCTTGTTACGACTT-3'). Genotypical characterization was carried out using sequence analysis of 16S. For comparison, sequences of rRNA genes were obtained from the database of the National Biotechnology Information Center (NCBI).

Effect of the actinomycete isolates on removal of ammonia and phosphate from the wastewater

The raw wastewater samples were collected from El Gona wastewater station on the northern Egyptian Red Sea coast. The samples were filtered through 45 μ filter membrane to remove the suspended particles and any materials ($\geq 45\mu$), then sterilized in the autoclave at 121° C for 20 minutes to avoid any contaminations by the microbial organisms (fungal and bacterial). Physical parameter (pH, TDS Dissolved oxygen and salinity) were measured using Inst. Model (YSI 5913). The dissolved ammonia and orthophosphates in the wastewater were measured using (JENWAY-6800UV/VIS) spectrophotometer at 630nm, and 880nm (APHA, 1995) before the treatment process.

Experimental applications

The pre-sterilized wastewater sample was divided into 10 flasks (about 250 ml for each) then inoculated with a young culture of the selected actinomycete isolates. The flasks were shaking in incubator at 30°C for 7 days. After incubation, 100 ml of each cultured sample was filtered through bacterial filter paper then the filtrates were taken for measuring the physicochemical parameters (dissolved oxygen, TDS, salinity and pH) and the nutrient salts; ammonia and orthophosphate. The percentage of nutrients removal was calculated according to the equation:

$$\text{Removal Efficiency (\%)} = \frac{(\text{Initial concentration} - \text{Final concentration})}{(\text{Initial concentration})} \times 100$$

RESULTS AND DISCUSSION

The selected marine actinomycetes isolates were distinguished by their differential morphological and chemical characteristics (Table 1). The substrate mycelium color of four isolates of them was white, three isolates were creamy color,

two isolates were grey and the last one was brown color and the most of them have grey to white aerial mycelia. The majority of isolates is producing beige diffusible pigments and only one produces violet pigment. The isolates were growing under the conditions; 25-38°C and pH 7–9. Starch and glycerol were utilized by all isolates as carbon sources, the majority utilized lactose, dextrose, maltose and D-glucose. All the used isolates were growing in presence of 6% NaCl, three isolates grew in presence of 9% NaCl and three isolates grew in presence of 10% NaCl.

Table 1: Phenotypic characteristics of actinomycetes isolates under investigation

Character	Isolates									
	M5	M 8	M 9	M 13	M 15	M 23	M 38	M 50	MN	MP
<u>Substrate mycelium</u>										
Grey	+	-	+	-	-	-	-	-	-	-
Brown	-	-	-	-	+	-	-	-	-	-
Cream	-	+	-	+	-	-	-	-	+	-
Beige	-	-	-	-	-	-	-	-	-	-
White	-	-	-	-	-	+	+	+	-	+
<u>Aerial mycelium</u>										
Yellow	-	+	-	-	-	-	-	-	-	-
White	-	-	-	-	-	-	-	-	-	-
Cream	-	-	-	-	-	+	-	-	+	-
Grey	+	-	+	+	-	-	+	+	-	+
Brown	-	-	-	-	+	-	-	-	-	-
<u>Diffusible pigments</u>										
Beige	+	+	-	+	+	-	+	+	+	-
Violet	-	-	+	-	-	-	-	-	-	-
<u>Growth at (°C)</u>										
25-35	+	+	+	+	+	+	+	+	+	+
38	+	+	+	+	-	-	+	-	+	+
45	-	-	-	-	-	-	-	-	-	-
<u>Growth at pH</u>										
5-6	-	-	-	-	-	-	-	-	-	-
7	+	-	-	+	+	+	+	+	+	+
8	+	+	+	+	+	+	+	+	+	+
9	-	-	-	+	+	+	+	+	+	+
<u>Utilization of</u>										
Starch	+	+	+	+	+	+	+	+	+	+
Lactose	+	+	+	+	-	+	+	+	+	+
Dextrose	+	+	+	+	-	+	+	+	+	+
Maltose	+	-	+	+	+	+	+	-	+	+
Glucose	+	+	+	+	-	+	+	+	-	-
Glycerol	+	+	+	+	+	+	+	+	+	+
<u>NaCl (%)</u>										
6	+	+	+	+	+	+	+	+	+	+
7-9	-	-	-	+	-	-	+	+	-	+
10	-	-	-	+	-	-	+	+	-	-
<u>Biochemical tests</u>										
Protease	+	-	+	+	-	+	-	+	+	+
Lipase	+	+	-	+	+	+	+	+	+	-
Urease	+	-	-	+	-	+	-	+	+	+
Catalase	+	+	+	+	+	+	+	+	+	+
Chitinase	-	-	-	-	-	-	-	-	-	-
Oxidase	-	-	-	-	-	-	-	-	-	-
Sulphide production	-	-	-	-	-	-	-	-	-	-
Melanin production	-	-	-	-	-	-	-	-	-	-
<u>Degradation of</u>										
Cellulose	+	+	+	+	+	+	+	+	+	+
Gelatin	-	-	-	-	-	-	-	-	-	-

Actinomycete numbers and kinds vary with environmental circumstances such as geographical area, pH, temperature, humidity, nutrients and climate (Hong *et al.* 2009 and Arifuzzaman *et al.* 2010). Rifaat (2003) it has been indicated that

actinomycetes are commonly distributed in different water bodies and attached as biofilms to any difficult substrate where they play a major part in the carbon cycle owing to their capacity to grow at low carbon content and degrade the remaining organic matter. All the cultured isolates were produced catalase, meanwhile the majority were also produced lipase and urease. Six isolates were produced protease and only one produced chitinase and oxidase. No precipitated sulphite and melanin were detected. Cellulose was hydrolyzed but no gelatin liquefaction occurred.

The BOX-PCR method can be a useful identifying material for isolates in collection with a extensive database or 16S rRNA gene sequencing and phylogenetic analysis. The amplification of bacterial genomic DNA as a biotechnological instrument plays a significant role in isolating improvements in a variety of pharmaceutical and production applications, including antibiotic biosynthesis, bioconversion and poisonous compound devolution (Li *et al.* 2017). Phenotypic characterization and molecular phylogenetic analysis recognized the actinomycete isolates used for treatment of wastewater. The actinomycetes isolate genomic DNA was prepared and the 16S rRNA gene coding was partly amplified using the universal primers (16S 27F and 16S 1492R), (5'AGAGTTTGCTGCTGCAG-3' and 5'-GGTTACCTTGACTGACTT-3'). The sequencing data utilizing of the strategy (ABI 3730xl) was 1500 base pair. Sequencing data were aligned against the 16S rRNA sequences of (<http://blast.ncbi.nlm.nih.gov/Blast.cgi>). The amplicons produced by the actinomycetes isolates were identified using electrophoresis of agarose gel as shown in (Figure 1). These sequences were compared with those gave the high esthomology using Blast Search Computer Based Program and the results reported in (Table 2). The ten actinomycete isolates were then tested for biological treatment and the removal of ammonia and orthophosphate from the raw wastewater.

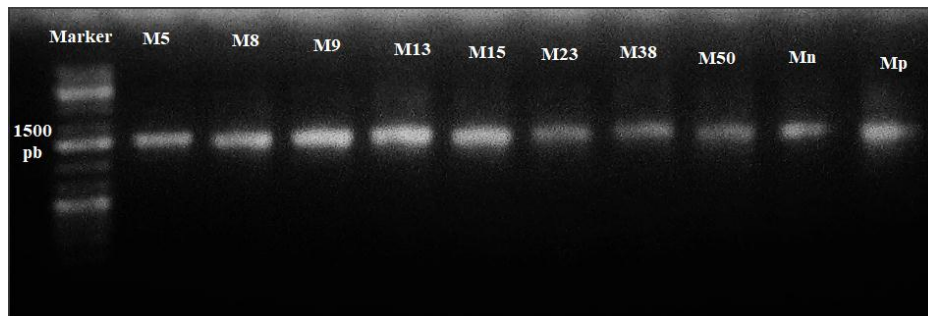


Fig. 1: Agarose gel electrophoreses of the amplified 16S rRNA gene of the actinomycetes isolates

Table 2: The similarity percentages of the most related actinomycetes species

Actinomycetes isolates no.	Most related Species	Similarity (%)
M 5	<i>Kocuriapalustris</i> strain TH126	95
M 8	<i>Streptomyces parvus</i> strain J59	96
M 9	<i>Streptomyces griseorubens</i> strain 173477	96
M 13	<i>Streptomyces rochei</i> strain R1-2A/D104	99
M 15	<i>Streptomyces albidoflavus</i> strain DT-A40	98
M 23	<i>Streptomyces sp.</i> strain L116	99
M 38	<i>Streptomyces sp.</i> strain A300Ydz-QZ	91
M 50	<i>Streptomyces sp.</i> Strain I08A-00376	95
MN	<i>Streptomyces griseus</i> strain SYA.E50	99
Mp	<i>Streptomyces sp.</i> strain 66P31-1	99

Capabilities of the actinomycete strains for wastewater treatment

The physico-chemical parameters of the raw sewage are as follows: dark brown color, with un-acceptable odors, alkaline (pH~9.32) and depleted dissolved oxygen content (0.6mg/l). The measured total dissolved salts (TDS) and salinity (‰) recorded 1508mg/l and 1.17‰ respectively. The nutrient salts of raw wastewater before bacterial treatment showed significantly high contents for ammonia and orthophosphate; 139.55, and 22.38mg/l respectively (Table 3).

Table 3: physicochemical parameters of raw wastewater before the biological treatment

Parameter	Concentration
color	Dark brown
odor	Un acceptable
pH	9.32
TDS	1508mg/l
Salinity	1.17‰
D.O	0.6mg/l
Ammonia	139.55±9.31mg/l
Phosphate	22.38±0.53 mg/l

Wastewater with high organic load causes many ecological problems (Manh, 2008). It shows adverse effects on both flora and fauna; its discharge to the land alters physical and chemical properties of the soil, thus reducing the fertility of land for crop production and its discharge to the water bodies may results in eutrophication, affecting the aquatic life and making water unfit for drinking (Manu *et al.* 2011). Hence, in recent years, the biological treatment system has become popular, more efficient with low cost in waste treatment (Vishakha *et al.* 2013).

Dissolved oxygen is considered an important pollution indicator parameter to examine the water quality. After treatment and as shown in Table (3), a slight increasing in dissolved oxygen ranging from 0.7 ± 0 to 0.967 ± 0.289 mg/l was observed with isolates M8, M9, M13, M15, M23 and M38, while MP, M50, MN and M5 isolates recorded significant high increasing in the dissolved oxygen (4.233 ± 0.289 , 3.33 ± 0.702 , 2.3 ± 0.1 and 1.967 ± 0.153 mg/l, respectively). The recorded high dissolved oxygen mean that there is plenty of organic matter present that provided energy for some actinomycete organisms (MP, M50, MN and M5) and they have ability to make biodegradation to most of the organic compounds, meanwhile the other actinomycetes isolates haven't this ability.

The recorded alkaline value (pH~9.32) in the raw wastewater indicated that the actinomycetes have the ability to survive under the elevated pH conditions. It is known that actinomycetes grow well and survive in neutral and slightly alkaline media (Trenozhnikova and Azizan, 2018). During the period of the experiment, pH values at all the cultured isolates were significantly declined from 9.32 to 8.283.

It is very important to be existed a sufficient alkalinity in the wastewater to balance the acid produced by bacterial nitrification process, where alterations in pH could have an adverse effect on nitrification. About 7.14 mg of alkalinity (as CaCO₃) are consumed to oxidize mg NH₃-N. (Sedlak, 1991). This consumption of alkalinity throughout nitrification is responsible for the depressing effect on the pH value.

In the same manner, slight decreasing in the total dissolved salts (TDS) was observed in the used wastewater flasks during the period of the experiment. The TDS removal efficiency of the experimented species was ranged from 6.6 to 9.6% for all isolates. This would be explained by the consuming these salts by experimented

isolates which used them as source of nourishment. Andriambelason *et al.* (2017) were succeeded to decrease the TDS from wastewater collected from oilfield using the actinomycetes strains; *Streptomyces albidoflavus* DSM40455T and *Streptomyces antibioticus* NBRC12838T. Shruthi *et al.* (2012) found 68.8% reduction in TDS using *Pseudomonas sp.* in rubber processing. Gaikwad *et al.* (2014) recorded maximum decrease in TDS (74.36%) using microbial consortia of different bacterial strains; *Pseudomonas*, *Actinomycetes*, *Bacillus*, *Staphylococcus* and *Streptomyces*.

Autotrophic and heterotrophic bacteria can remove ammonia through two processes; nitrification that converting ammonia to nitrite and nitrate in two sequential steps and de-nitrification. These bacteria can tolerate nitrogen and organic carbon loads (Joo *et al.* 2005). The *Thiobacillus denitrificans* autotrophic organism and the heterotrophic organisms *Micrococcus denitrificans*, *Pseudomonas* and *Archomobacter* are accountable for this phase. (Torrento *et al.* 2010 and Kim *et al.* 2008).

Nitrification takes place in two phases: first, ammonia is converted to nitrite by ammonia oxidizing bacteria and this nitrite is converted to nitrate by nitrite oxidizing bacteria and actinomycetes isolates use ammonia or nitrite as an electron-acceptor and carbon dioxide as a source of carbon (Laanbroek *et al.* 2002). The de-nitrification liberates N₂ gas from nitrates. Some microorganisms can transform nitrate into nitrous oxide and nitrogen gas under anaerobic conditions through a method called denitrification that results in a net loss of nitrogen from the soil. Recent progress in identifying heterotrophic ammonia remediation bacteria has revolutionized ammonia extraction techniques for sewage treatment procedures. Bacteria obtain their source of elemental nitrogen through the assimilation of ammonia, but can also use nitrites and nitrates as well as gaseous nitrogen under certain conditions. Microorganism assimilation of ammonia is regarded one of the significant measures to remove ammonia from the manure treatment systems. The metabolism of actinomycetes has been observed as a notable source of enzymes and bioactive products and therefore an significant problem is the function of actinomycetes with respect to the removal of ammonia during the composting process. (Sasaki *et al.* 2005).

Table (3) shows the capacity of the chosen actinomycete isolates for ammonia removal from wastewater samples. The efficiency of the selected isolates in ammonia removal was varied obviously. The ammonia removal efficiency was ranged between 23.96 and 77.35%. The highest removal of ammonia is achieved by isolate M9 followed by isolates M15 and M23 with the removal efficiencies of; 77.35%, 67.03% and 62.58% respectively). On the other hand, isolates M5 and M8 recorded the lowest removal efficiencies of ammonia (29.77 and 23.96 %).

By assimilating soluble orthophosphate (PO₄), bacteria obtain their source of elemental phosphorus. Phosphorus ion is the final breakdown of the organic phosphorus process and is the most easily accessible type of phosphorus for biological uses. (Ahemad *et al.* 2009). Throughout the period of the experiment, the concentration of orthophosphate was decreased from 22.38±0.54 to 7.00±0.00 mg/l with removal efficiencies ranging from 58.86% to 79.02%. The isolates; M5, M13, M38, MN and MP recorded orthophosphate removal efficiency more than 70%, while the rest isolates achieved removal efficiency exceeded 63% except isolate M50 which recorded 58.86% removal efficiency. This could be because of the differential abilities of these isolates to produce acids and enzymes that responsible for breakdown the orthophosphate compounds (De Souza *et al.* 2000 and Seshadri *et al.* 2002).

Table 4: Physicochemical parameters of wastewater after the treatment using the selected actinomycetes strains

Bacterial name	pH	D.O	TDS	Salinity
IS/5	8.603±0.025	1.967±0.153	1408.33±27.06	1.093±0.020
IS/8	8.5±0.052	0.767±0.058	138.6	1.086
IS/9	8.477±0.032	0.7±0	1391±39.53	1.08±0.03
IS/13	8.48±0.046	0.7±0	1362.83±32.06	1.06±0.02
IS/15	8.507±0.047	0.7±0	1375.16±11.02	1.07±0.01
IS/23	8.453±0.032	0.7±0	1378±28.33	1.07±0.02
Is/38	8.42±0.026	0.967±0.289	1384.5±6.5	1.076±0.005
IS/50	8.443±0.032	3.333±0.702	1391±6.5	1.083±0.005
IS/MN	8.377±0.067	2.3±0.1	1388.83±3.75	1.08±0
IS/NP	8.283±0.064	4.233±0.289	1369.16±44.52	1.036±0.037

Table 5: Removal percentage (%) of nutrients (ammonia and orthophosphate) in wastewaters after treatment using actinomycetes strains

Bacterial strains	Ammonia removal (%)	Ammonia removal (mg/L)	orthophosphate removal (%)	Phosphate removal (mg/L)
IS/5	29.77	41.55	72.04	24.05
IS/8	23.96	33.44	63.45	21.18
IS/9	77.35	107.95	64.94	21.68
IS/13	59.37	82.85	73.30	24.47
IS/15	67.03	93.54	68.99	23.03
IS/23	62.58	87.33	68.82	22.974
Is/38	46.96	65.54	70.24	23.447
IS/50	47.39	66.14	58.86	19.65
IS/MN	33.63	46.94	79.02	26.38
IS/NP	55.83	77.91	76.73	25.6134

CONCLUSION

Wastewater treatment is one of the extreme problems for both developed and developing countries. Addendum for that, globalization placing defy for waste management within communities. Wastewater treatment is very essential to achieve sustainable expansion in economic issues, health concerns, and environmental problems and the role of microorganisms in wastewater treatment helps to treat and purify wastewater and make it less harmful to the environment. In this study, ten marine actinomycetes isolates were isolated from Suez gulf, Egypt using for biological treatment to samples of raw sewage. All of isolates have ability to treatment of domestic wastewater in an acceptable manner especially strains *Streptomyces griseorubens* and *Streptomyces griseus*.

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