

## IDENTIFICATION OF EIGHT HALOTOLERANT STREPTOMYCETE ISOLATES USING A SUGGESTED NUMERICAL TAXONOMY

# [42]

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### ABSTRACT

A numerical method was suggested for the identification of eight local halotolerant streptomycete isolates. Cultural, morphological, physiological and antagonistic characteristics of those isolates were determined. Arbitrary scoring of these characters for the eight unknown isolates and those of the more related known *Streptomyces* species in the key of **Pridham and Tresner (1974)** was given and resulted 58 characteristic units. The degree of similarity was determined using the Dice coefficient method and clustering was achieved using the unweighted pair group method average (UPGMA) algorithm. Using this numerical method, six out of the eight *Streptomyces* isolates, i.e., ST08, ST09, ST11, ST13, ST14 and ST15 were identified as *S. longisporus*, *S. janthinus*, *S. griseochromogenes*, *S. antibioticus*, *S. baarnensis* and *S. albolongus* with 100, 95.8, 95.0, 92.8, 95.0 and 91.1%, respectively. Isolates ST10 and ST12 were duplicate of *S. echinatus* with similarities of 93.3 and 94.6%, respectively. Application of the suggested numerical taxonomy on the 14 known *Streptomyces* species revealed that these species fell into three major clusters based on their color of aerial mycelia.

**Key words:** *Streptomyces*, Numerical identification, Halotolerant, Characteristic units, Taxonomy

### INTRODUCTION

Genus *Streptomyces* comprises, by far, the largest number of species of actinomycetes now known to occur in nature (**Williams et al 1989**). Many investigators throughout the world are isolating cultures of streptomycetes from soils (**Goodfellow et al 1987; Srinivasan**

**et al 1991 and Mohamed et al 2001**) and other substrates (**Saleh et al 1990 and Mohamed et al 2000**) and studying their cultural, physiological and biochemical activities (**Abdel-Fattah, 2005**).

**Goodfellow (1967)** described a numerical taxonomy method and cultures accordingly. Numerical taxonomy methods were thereafter applied to the

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genus *Streptomyces* by **Paszkiewicz (1972)**; **Kurylowicz et al (1975)** and **Goodfellow et al (1992)**. **Goodfellow et al (1979)** used a numerical system to classify 156 *Actinomadura* strains, and found marker strains of related taxa, and related isolates from bagasse and fodder via the numerical phenetic analyses using 90 unit characters. **Williams et al (1983)** suggested a standard numerical classification of 475 strains, of which 394 type cultures of *Streptomyces*, and 14 other actinomycete genera.

The present study suggests a numerical taxonomy method for eight *Streptomyces* isolates by comparing their phylophenetic characters with their corresponding strains in the eighth edition of Bergey's Manual of Determinative Bacteriology (**Pridham and Tresner, 1974**).

## MATERIAL AND METHODS

### *Streptomyces* isolates

Eight-halotolerant streptomycete isolates belonging to white (ST08, ST14 and ST15); red (ST09) and gray (ST10, ST11, ST12, and ST13) series were kindly provided from the Department of Agricultural Microbiology, Institute of Soil, Water and Environment Research, ARC, Giza, Egypt.

### Characterization of *Streptomyces* isolates

Cultural and morphological characteristics of *Streptomyces* isolates under investigation were determined as proposed by **Pridham and Tresner (1974)** using the media and methods of The International *Streptomyces* Project (ISP) as described by **Shirling and Gottlieb (1966)**. In addition, antagonistic activities of the tested *Streptomyces* isolates against 2 fungi, 2 yeasts and 5 bacteria (kindly provided by Cairo MIRCEN, Faculty of Agriculture, Ain Shams University) were determined as described by **Mohamed et al (2001)**. Salt tolerance range was also studied using four NaCl concentrations, i.e., 3.5, 7.0, 10 and 13% as mentioned by **Mohamed et al (2000)**. Chain type and spore surface of the *Streptomyces* isolates were determined as recommended by **Pridham and Tresner (1974)** using the light and electron microscopy, respectively. The abilities of the isolates to produce melanoid pigment, growth on Czapek's medium, to produce diffusible pigments, to tolerate streptomycin ( $4\mu\text{g ml}^{-1}$ ) and to utilize nine carbon compounds were studied as described by **Saleh et al (1990)**.

### Numerical identification of *Streptomyces*

Characters of *Streptomyces* reported in the key of **Pridham and Tresner (1974)** in the eighth edition of Bergey's Manual of Determinative Bacteriology

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were divided into two parts. Part I represents the main taxonomical characters that included color of aerial mycelium, spore-chain, melanoid pigment, spore surface and utilization of carbon compounds. Part II includes the other assisted characters for streptomycetes classification, i.e., growth on Czapek's medium, anti-bacterial and -

fungal activities, sensitivity to streptomycin, color of substrate mycelium, diffusible pigments and NaCl tolerance.

Suggested scoring of each of the previous characters (main and assisted) was given 1 unit if present or 0 unit if absent (**Table 1**). Relative importances for each of these characters were arbitrary



## Identification of streptomycete isolates by numerical taxonomy

Table 1. Cont.

Characters for streptomycetes identification	Present	Absent
II- Other assisted characters		
f- Growth on Czapek's medium (5 units)		
Fair	1 (1)	0
Poor	1 (2)	0
Moderate	1 (3)	0
Good	1 (4)	0
Excellent	1 (5)	0
g- Antibacterial activity (2 units)	1 (2)	0
h- Antifungal activity (2 units)	1 (2)	0
<b>Subtotal</b>	<b>58 (1)</b>	<b>0</b>
i- Sensitivity to streptomycin (1 unit)	1 (1)	0
j- Color of substrate mycelium (1 unit)	1 (1)	0
k- Diffusible pigments (1 unit)	1 (1)	0
l- NaCl tolerance (4 units)		
0-7%	1 (2)	0
0-10%	1 (3)	0
> 10%	1 (4)	0
<b>Total units</b>	<b>1 (65)</b>	

suggested according to their importance in the identification of streptomycetes. Accordingly, color of aerial mycelium was given a maximum of **22** units, type of spore chain **4** units, melanoid pigments **1** unit and spore surface **5** units. Utilization of carbon compounds was given **17** units, growth on Czapek's medium **5** units, antibacterial activity **2** units, antifungal activity **2** units, sensitivity to streptomycin **1** unit, color of substrate mycelium **1** unit, diffusible pigments **1** unit and NaCl tolerance **4** units. Within

each character, its units were distributed according to frequency distribution of sub-characters. For example, gray, red, white, blue, yellow, green, and violet aerial mycelia were given 22, 16, 13, 10, 7, 4 and 1 units, respectively.

Cultural, morphological and assisted characters of the eight *Streptomyces* isolates were compared with those of the most similar 14 *Streptomyces* species in the key of **Pridham and Tresner (1974)**. The similarity matrix between the experimental and identified species, in

**Pridham and Tresner (1974)**, was determined by Dice Coefficient method. In addition clustering of all characters was determined by the unweighted pair group method with average (UPGMA) algorithm (**Sneath and Sokal, 1973**).

Analyses were done using the Diversity Data base™ Version 2.0 from Bio-Rad.

Suggested numerical identification was based on the four characteristics, which are used for streptomycetes identification in **Pridham and Tresner (1974)**. In addition, some of the assisted characters, i.e., growth on Czapek's medium, anti- bacterial and -fungal activities due to their use in the identification of more than 60% of *Streptomyces* species. The unit character for the main characters was 49, and those of selected assisted characters were 9 units making a total of 58 unit characters. The rest of assisted characters, namely, sensitivity to streptomycin, color of substrate mycelium, diffusible pigments, and NaCl tolerance are not usually reported for identification of the majority of *Streptomyces* species in the key of **Pridham and Tresner (1974)**.

## RESULTS AND DISCUSSION

### Characterization of the *Streptomyces* isolates

Characterization of white, red and gray streptomycete isolates are given in

Table (2), Figures (1), (2) and could be represented as follows:

### White *Streptomyces* isolates

The two white *Streptomyces* isolates, namely, ST14 and ST15 were characterized with RF chain spores and smooth surface spores, while the third (ST08) had RA chain spores and spiny spore surface. Both of ST08 and ST15 were able to produce melanoid pigment while, ST14 did not. Regarding the utilization of carbon compounds, ST08 utilized all the carbon compounds as sole sources of carbon. On the other hand, ST14 and ST15 varied in their utilization of raffinose and sucrose, and did not utilize mannitol. The three isolates showed good or excellent growth on Czapek's medium. The ST14 and ST15 isolates showed white aerial mycelium and only showed antibacterial activities. On the other hand, ST08 showed colorless aerial mycelium and no antagonistic activities against the bacterial or fungal test organisms. Variation was also observed in the sensitivity of these isolates to streptomycin.

### Red *Streptomyces* isolates

The red *Streptomyces* isolate, namely, ST09 was characterized with RA or spiny chains spore with smooth surface, producing melanoid pigments, utilizing of

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all carbon compounds as sole source of carbon, good growth on Czapek's medium, yellow substrate mycelium, not sensitive to streptomycin, antagonistic activity against the bacterial test organisms.

**Gray *Streptomyces* isolates**

Three of the four gray *Streptomyces* isolates, namely, ST10, ST11 and ST12 were characterized with RA or spiny chain spores, while the fourth (ST13) showed RF chain spores with smooth spores, and all of them produced melanoid pigment. The color of their aerial mycelia varied between yellow to grayish yellow. They showed variation in

Table 2. Taxonomical characters of eight halotolerant streptomycete isolates according to the key of **Pridham and Tresner (1974)** .

<i>Streptomyces</i> isolates	I- Main taxonomical characters							
	a-Color of aerial mycelium		b-Spore-chain*		c-Melanoid pigment		d-Spore surface	
ST08	white		RA		+		spiny	
ST09	red		RA-S		+		spiny	
ST10	gray		RA		+		spiny	
ST11	gray		RA-S		+		spiny	
ST12	gray		RA		+		spiny	
ST13	gray		RF		+		smooth	
ST14	white		RF		-		smooth	
ST15	white		RF		+		smooth	
e-Utilization of carbon compounds								
Carbon sources	<i>Streptomyces</i> isolates							
	ST08	ST09	ST10	ST11	ST12	ST13	ST14	ST15
No carbon	-	-	-	-	-	-	-	-
D-Glucose	+	+	+	+	+	+	+	+
D-Xylose	+	+	+	+	+	+	+	+
L-Arabinose	+	+	+	+	+	+	+	+
L-Rhamnose	+	+	+	-	+	+	+	+
D-Fructose	+	+	+	+	+	-	+	+
Raffinose	+	+	-	+	-	+	-	+
D-Mannitol	+	+	+	+	+	+	-	-
i-Inositol	+	+	+	+	+	+	+	+
Sucrose	+	+	-	+	-	-	+	-
II- Other assisted characters								
Isolates	Growth on Czapek's medium	Antagonistic activity	Sensitivity to streptomycin	Color of substrate mycelium	Diffusible pigments	NaCl tolerance (Up to 10%)		
ST08	Excellent	-	-	Colorless	-	+		
ST09	Good	Anti-bacterial	-	Yellow	-	+		
ST10	Moderate	-	+	Grayish yellow	-	+		
ST11	Excellent	Anti-bacterial	-	Yellow	-	+		
ST12	Good	Anti-bacterial	+	Grayish yellow	-	+		

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\*RA: spore chain in the form of open loops, hooks or greatly extended coils of wide. RF: spores in straight (R) or flexuous (F) chains. S: spira; spore chain in form of hooks, open loops and coils

Figure 1. Microphotographs of spore chain of some *Streptomyces* isolates (x-400).

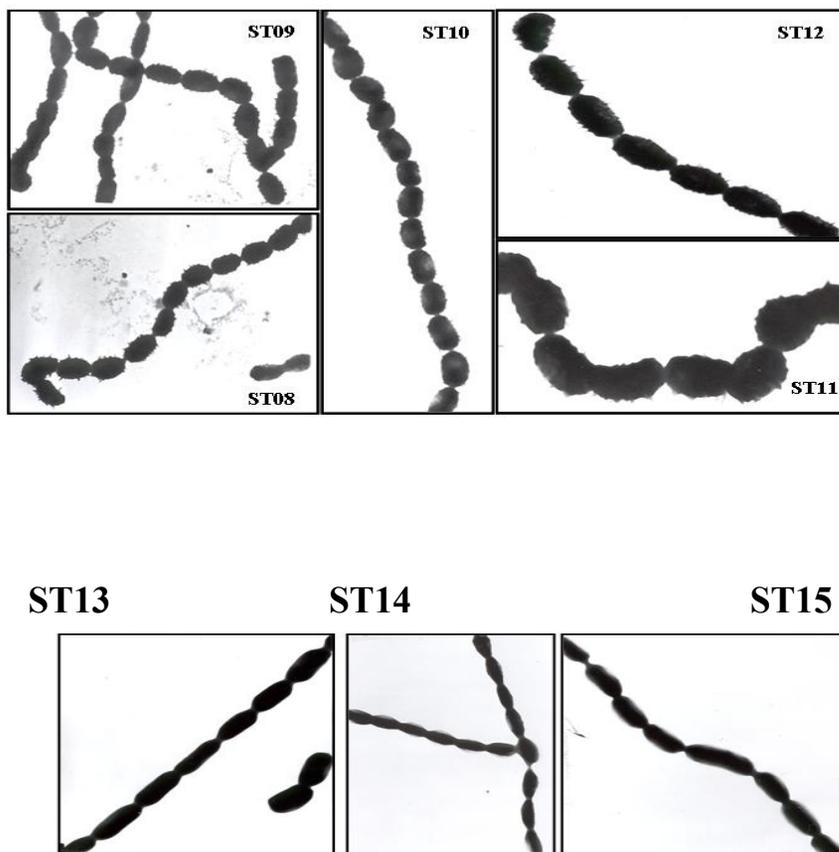


Figure 2. Electron micrographs (X-10000) of spore surface of some *Streptomyces* isolates.

the utilization of sucrose, raffinose, rhamnose and fructose as sole carbon sources. Growth on Czapek's medium was moderate (ST10 and ST13), good (ST12) or excellent (ST11). They showed antagonistic activities against the bacterial test organisms except for ST10. They were also sensitive to streptomycin except for ST11.

All the isolates under investigation were able to grow in the presence of 10% NaCl in the growth medium and could not produce diffusible pigments.

#### **Numerical identification of the *Streptomyces* isolates**

According to the proposed numerical taxonomy in this study, cultural, morphological, physiological and antagonistic characteristics of the eight *Streptomyces* isolates and related *Streptomyces* species recorded in the proposed key of **Pridham and Tresner (1974)** were given 1 unit if present or 0 unit if absent. Scoring results for all characters of the eight *Streptomyces* isolates are given in **Table (3)**.

Accordingly, isolates numbers ST13, ST11, ST09, ST12, ST10, ST14, ST15 and ST08 had the following scoring in descending order, being 50, 50, 45, 45, 42, 41, 40 and 40 units, respectively.

Clustering of all scoring units was determined (**Sneath and Sokal, 1973**) and the results as phylogenetic tree are given in **Figure (3)**. The data reveal the

presence of two major related clusters, one includes, ST08, ST09, ST14 and ST15 and the second includes ST10, ST11, ST12 and ST13. It was also found that each major cluster contained two subclusters. Subcluster A contained ST14 and ST15; subcluster B contained ST08 and ST09, subcluster C contained ST11 and ST13 and finally subcluster D contained ST10 and ST12.

#### **Numerical identification of white series isolates**

Characters of the three white *Streptomyces* isolates, namely, ST08, ST14 and ST15 and most similar *Streptomyces* species, i.e., *S. albolongus*, *S. alboniger*, *S. baarnensis*, *S. longisporus* and *S. viridaris* in the key proposed by **Pridham and Tresner (1974)** were scored (**Tables 4 and 5**).

The analysis of the scored data showed that ST08, ST14 and ST15 isolates were most related to *S. longisporus*, *S. baarnensis* and *S. albolongus*. Results in **Table (6)** and **Figure (4)** showed that ST08 & *S. longisporus*; ST14 & *S. baarnensis* and ST15 & *S. albolongus* fell in three subclusters with similarities of 100, 95 and 91.1%, respectively. However, slight differences were found between ST14 and *S. baarnensis* in the utilization of raffinose and antibacterial activities.

#### **Identification of red series isolates**

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The scored characters of the red ST09 *Streptomyces* isolate under investigation was compared with the scored characters of three red *Streptomyces* species, namely, *S. purpurascens*, *S. yokosukanensis* and *S. janthinus* in **Pridham and Tresner (1974)** key (**Table 7**). Result in **Table (8)** and **Figure**

**(5)** show that ST09 could be identified as a strain of *S. janthinus* with similarity of 95.8 %. The 4.2% differences between them could be due to the type of spore-chain, growth on Czapek's medium, antagonistic activities and production of diffusible pigments as shown in **Table (7)**.

Table 3. Scoring of the characters of the eight *Streptomyces* isolates under investigation.

Characters	<i>Streptomyces</i> isolates								
	ST08	ST09	ST10	ST11	ST12	ST13	ST14	ST15	
a- Color of aerial mycelium (22)	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	1	1	1	1	1	1	1	1	
	0	1	1	1	1	1	1	0	0
	0	1	1	1	1	1	1	0	0
	0	1	1	1	1	1	1	0	0
	0	0	1	1	1	1	1	0	0
	0	0	1	1	1	1	1	0	0
	0	0	1	1	1	1	1	0	0
	0	0	1	1	1	1	1	0	0
	0	0	1	1	1	1	1	0	0
	b-Spore-chain (4)	1	1	1	1	1	1	1	1
1		1	1	1	1	1	1	1	
0		1	0	1	0	1	1	1	
0		0	0	0	0	1	1	1	
c-Melanoid pigment (1)	1	1	1	1	1	1	0	1	

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Table 3. Cont.

Characters	<i>Streptomyces</i> isolates							
	ST08	ST09	ST10	ST11	ST12	ST13	ST14	ST15
e-Utilization of carbon compounds (17)								
No carbon	0	0	0	0	0	0	0	0
D-Glucose	1	1	1	1	1	1	1	1
D-Xylose	1	1	1	1	1	1	1	1
L-Arabinose	1	1	1	1	1	1	1	1
L-Rhamnose	1	1	1	1	0	1	1	1
	1	1	1	1	0	1	1	1
D-Fructose	1	1	1	0	1	0	1	1
	1	1	1	0	1	0	1	1
Raffinose	1	1	0	1	1	1	0	1
	1	1	0	1	1	1	0	1
D-Mannitol	1	1	1	1	1	1	0	0
	1	1	1	1	1	1	0	0
i-Inositol	1	1	1	1	1	1	1	1
	1	1	1	1	1	1	1	1
Sucrose	1	1	0	1	0	0	1	0
	1	1	0	1	0	0	1	0
	1	1	0	1	0	0	1	0
f- Growth on Czapek's medium (5)	1	1	1	1	1	1	1	1
	1	1	1	1	1	1	1	1
	1	1	0	1	1	0	1	1
	1	0	0	1	0	0	1	0
g- Anti-bacterial activity (2)	0	1	0	1	1	1	1	1
	0	1	0	1	1	1	1	1
h- Anti-fungal activity (2)	0	0	0	0	0	1	0	0
	0	0	0	0	0	1	0	0
<b>Total (58 Units)</b>	<b>40</b>	<b>45</b>	<b>42</b>	<b>50</b>	<b>45</b>	<b>50</b>	<b>41</b>	<b>40</b>

1: Present. 0: Absent.

Figure 3. Phylogenetic tree of eight *Streptomyces* isolates based on analysis of their selected characters.

Table 4. Scoring of the characters of ST08 isolate compared with those of related species in **Pridham and Tresner (1974)** key.

Characters	ST08 isolate	<i>Streptomyces</i> species in <b>Pridham and Tresner (1974)</b> key		
		<i>S. alboniger</i>	<i>S. longisporus</i>	<i>S. viridaris</i>
<b>a- Color of aerial mycelium (22)</b>	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	0	0	0	0
	0	0	0	0
	0	0	0	0
	0	0	0	0
	0	0	0	0
	0	0	0	0
	0	0	0	0
	<b>b-Spore-chain (4)</b>	1	1	1
1		1	1	1
0		1	0	1
0		1	0	1
<b>c-Melanoid pigment (1)</b>	1	0	1	1
<b>d-Spore surface (5)</b>	1	1	1	1
	1	1	1	1
	1	1	1	1
	0	1	0	1
	0	1	0	1

Table 4. Cont.

Characters	ST08 isolate	<i>Streptomyces</i> species in <b>Pridham and Tresner (1974)</b> key		
		<i>S. alboniger</i>	<i>S. longisporus</i>	<i>S. viridaris</i>
<b>e- Utilization of carbon compounds (17)</b>				
No carbon	0	0	0	0
D-Glucose	1	1	1	0
D-Xylose	1	1	1	1
L-Arabinose	1	1	1	0
L-Rhamnose	1	0	1	1
	1	0	1	1
D-Fructose	1	1	1	0
	1	1	1	0
Raffinose	1	0	1	1
	1	0	1	1
D-Mannitol	1	1	1	0
	1	1	1	0
i-Inositol	1	1	1	0
	1	1	1	0
Sucrose	1	0	1	1
	1	0	1	1
	1	0	1	1
<b>f- Growth on Czapek's meium (5)</b>	1	0	1	1
	1	0	1	1
	1	0	1	1
	1	0	1	1
	1	0	1	1
<b>g- Anti-bacterial activity (2)</b>	0	1	0	1
	0	1	0	1
<b>h- Anti-fungal activity (2)</b>	0	0	0	0
	0	0	0	0
<b>Total (58 Units)</b>	<b>40</b>	<b>33</b>	<b>40</b>	<b>38</b>

1: Present.0: Absent.

Table 5. Scoring of the characters of white ST14 and ST15 isolates compared with those of related species in **Pridham and Tresner (1974)** key.

Characters	ST14	<i>Streptomyces</i> species in		ST15
		<b>Pridham and Tresner (1974)</b>		
		key		
	<i>S. albolongus</i>	<i>S. baarnensis</i>		
<b>a- Color of aerial mycelium (22)</b>	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	0	0	0	0
	0	0	0	0
	0	0	0	0
	0	0	0	0
	0	0	0	0
	0	0	0	0
	0	0	0	0
	<b>b-Spore-chain (4)</b>	1	1	1
1		1	1	1
1		1	1	1
1		1	1	1
<b>c-Melanoid pigment (1)</b>	0	1	0	1
<b>d-Spore surface (5)</b>	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1



Table 5. Cont.

Characters	<i>Streptomyces</i> species in			
	ST14	Pridham and Tresner (1974)		ST15
		key		
	<i>S. albolongus</i>	<i>S. baarnensis</i>		
<b>e- Utilization of carbon compounds (17)</b>				
No carbon	0	0	0	0
D-Glucose	1	1	1	1
D-Xylose	1	1	1	1
L-Arabinose	1	1	1	1
L-Rhamnose	1	0	1	1
	1	0	1	1
D-Fructose	1	1	1	1
	1	1	1	1
Raffinose	1	0	0	1
	1	0	0	1
D-Mannitol	1	1	1	0
	1	1	1	0
i-Inositol	1	1	1	1
	1	1	1	1
Sucrose	0	0	0	0
	0	0	0	0
	0	0	0	0
<b>f- Growth on Czapek's medium (5)</b>	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	0
<b>g- Anti-bacterial activity (2)</b>	1	1	0	1
	1	1	0	1
<b>h- Anti-fungal activity (2)</b>	0	0	0	0
	0	0	0	0
<b>Total (58 Units)</b>	<b>42</b>	<b>39</b>	<b>38</b>	<b>40</b>

1: Present. 0: Absent.

Table 6. Similarities between the white *Streptomyces* isolates and related species in **Pridham and Tresner (1974)** key.

<i>Streptomyces</i> isolates	Related white <i>Streptomyces</i> species in <b>Pridham and Tresner (1974)</b> key				
	<i>S. longisporus</i>	<i>S. baarnensis</i>	<i>S. albolongus</i>	<i>S. viridaris</i>	<i>S. alboniger</i>
ST08	<b>100</b>	87.2	83.5	82.1	74.0
ST14	87.8	<b>95.0</b>	93.8	85.0	88.0
ST15	85.0	89.7	<b>91.1</b>	87.2	84.9

Bold number represents the most similar species.

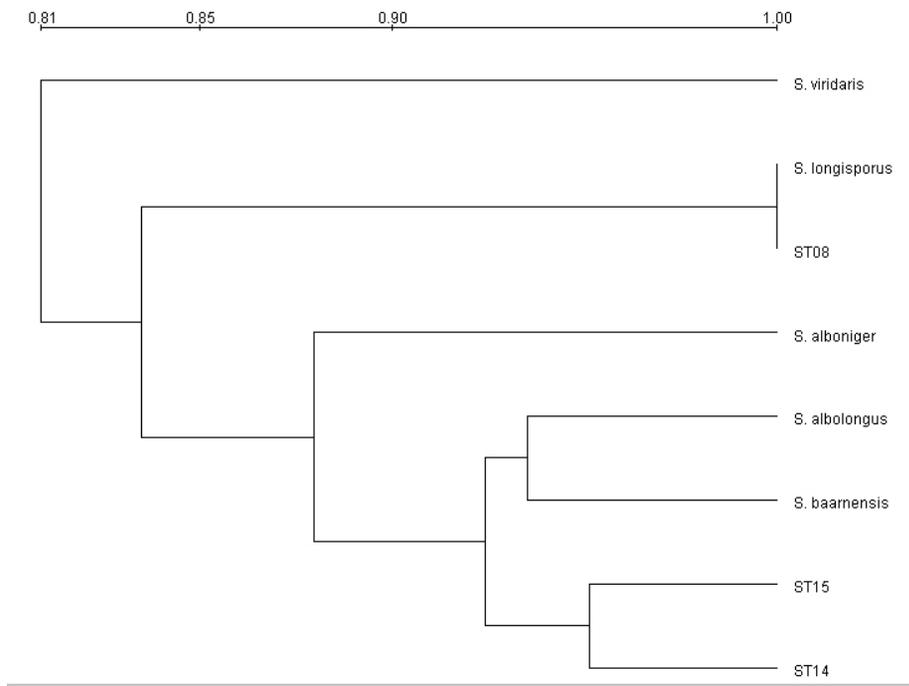


Figure 4. Phylogenetic tree of three *Streptomyces* isolates belonging to white series and related species in **Pridham and Tresner (1974)**.



## Identification of streptomycete isolates by numerical taxonomy

0	0	0	0
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Table 7. Cont.

Characters	ST09 isolate	<i>Streptomyces</i> species in <b>Pridham and Tresner (1974)</b> key		
		<i>S. janthinus</i>	<i>S. purpurascens</i>	<i>S. yokosukanensis</i>
e- Utilization of carbon compounds (17)				
No carbon	0	0	0	0
D-Glucose	1	1	1	1
D-Xylose	1	1	1	1
L-Arabinose	1	1	1	1
L-Rhamnose	1	1	1	1
D-Fructose	1	1	1	1
Raffinose	1	1	1	1
D-Mannitol	1	1	1	1
i-Inositol	1	1	1	1
Sucrose	1	1	1	1
f- Growth on Czapek's medium (5)	1	1	0	1
	1	1	0	1
	1	1	0	0
	1	1	0	0
g- Anti-bacterial activity	1	1	1	1
	1	1	1	1
h- Anti-fungal activity	0	1	0	1
	0	1	0	1
<b>Total (58 Units)</b>	<b>45</b>	<b>47</b>	<b>40</b>	<b>44</b>

1: Present. 0: Absent.

Table 8. Similarities between the red *Streptomyces* isolate and those related species in **Pridham and Tresner (1974)** key.

<i>Streptomyces</i> Isolate	Related red <i>Streptomyces</i> species in <b>Pridham and Tresner (1974)</b> key		
	<i>S. janthinus</i>	<i>S. yokosukanensis</i>	<i>S. purpurascens</i>
ST09	<b>95.8</b>	94.6	94.4

### Identification of streptomycete isolates by numerical taxonomy

Bold number represents the most similar species.

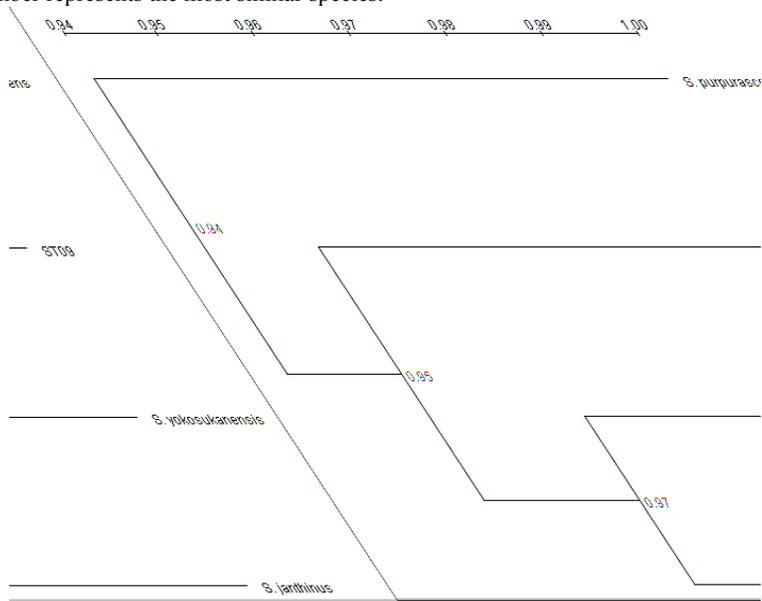


Figure 5. Phylogenetic tree of ST09 isolate belonging to red series and related species in Pridham and Tresner (1974).

### Identification of gray series isolates

Characters of ten gray streptomycetes, i.e., the four isolates under investigation (ST10, ST11, ST12 and ST13) and most related six species in the key of Pridham and Tresner (1974) (*S. durhamensis*, *S. filipinensis*, *S. griseochromogenes*, *S. chromofuscus*, *S. echinatus* and *S. antibioticus*) were scored (Tables 9 and 10). Clustering (Figure 6) and similarities (Table 11) analyses showed that ST11 and ST13 isolates were strains of *S. griseochromogenes*, and *S. antibioticus* with similarities of 95.0 and 92.8%, respectively. Isolates ST10 and ST12 were duplicate strains of *S. echinatus* with similarities of 93.3 and 94.6%, respectively. This result confirmed the previous results in Figure

(3). As these two *Streptomyces* isolates were fell in one subcluster with a similarity of 92%.

*Streptomyces* isolate ST11 differed from *S. griseochromogenes* in the utilization of L-rhamnose and fructose as sole carbon sources and color of substrate mycelium. ST11 tolerated the presence of NaCl up to 10% in the growth medium, while, *S. griseochromogenes* did not. There were some differences between ST13 and *S. antibioticus*, in utilization of raffinose, D-fructose, growth on Czapek's medium, antagonistic activities and salt tolerance (Table 9).

### Application of the suggested numerical taxonomy on some known *Streptomyces* species

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The suggested numerical taxonomy under investigation was applied for

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0	0	0	0
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Table 9. Cont.

Characters	ST11 isolate	<i>Streptomyces</i> species in <b>Pridham and Tresner (1974)</b> key		
		<i>S. durhamensis</i>	<i>S. filipinensis</i>	<i>S. griseochromogenes</i>
e- Utilization of carbon compounds (17)				
No carbon	0	0	0	0
D-Glucose	1	1	1	1
D-Xylose	1	1	1	1
L-Arabinose	1	1	1	1
L-Rhamnose	1	0	0	0
	1	0	0	0
D-Fructose	0	1	1	1
	0	1	1	1
Raffinose	1	1	1	1
	1	1	1	1
D-Mannitol	1	1	1	1
	1	1	1	1
i-Inositol	1	1	1	1
	1	1	1	1
Sucrose	1	0	1	1
	1	0	1	1
	1	0	1	1
F- Growth on Czapek's medium (5)	1	1	1	1
	1	1	1	1
	1	1	1	1
	1	1	1	1
g- Anti-bacterial activity (2)	1	0	0	1
	1	0	0	1
h- Anti-fungal activity (2)	0	1	1	0
	0	1	1	0
<b>Total (58 Units)</b>	<b>49</b>	<b>47</b>	<b>51</b>	<b>50</b>

Table 10. Scoring of the characters of ST10, ST12 and ST13 isolates compared with those of related species in **Pridham and Tresner (1974)** key.

Characters	ST10 isolate	Streptomyces species in <b>Pridham and Tresner (1974)</b> key			ST13 isolate	ST12 isolate	
		S.	S.	S.			
		<i>chromofuscus</i>	<i>echinatus</i>	<i>antibioticus</i>			
a- Color of aerial mycelium (22)	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	1	1	1	1	1	1	
	b-Spore-chain (4)	1	1	1	1	1	1
		1	1	1	1	1	1
0		0	1	1	1	1	
0		0	1	1	1	1	
c-Melanoid pigments (1)	1	1	1	1	1	1	
d-Spore surface (5)	1	1	1	1	1	1	
	1	1	1	1	1	1	
	0	0	0	1	1	0	
	0	0	0	1	1	0	

## Identification of streptomycete isolates by numerical taxonomy

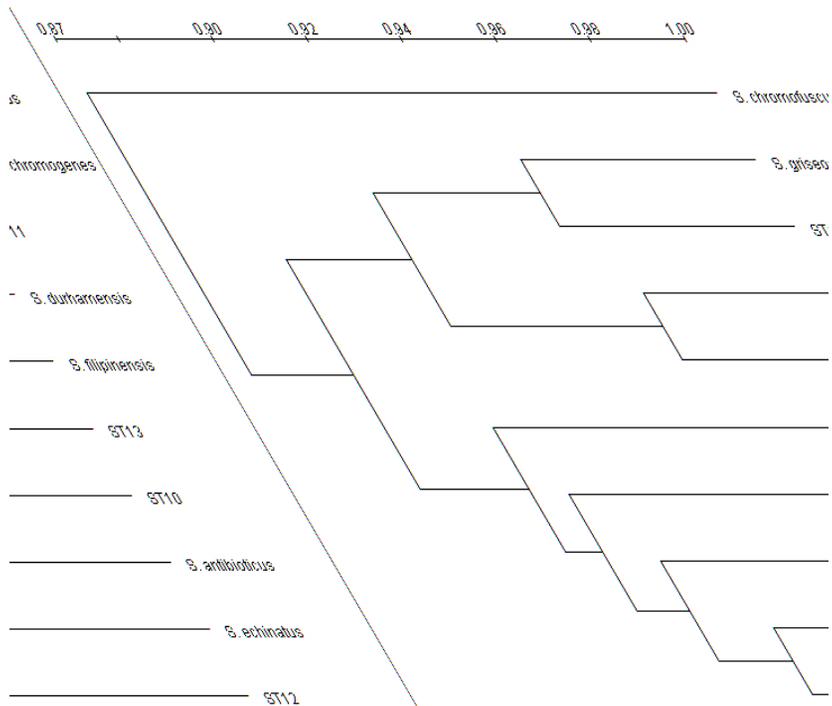
Table 10. Cont.

Characters	ST10	Streptomyces species in Pridham and			ST13	ST12
	isolate	Tresner (1974) key			isolate	isolate
		S.	S.	S.		
		<i>chromofuscus</i>	<i>echinatus</i>	<i>antibioticus</i>		
e- Utilization of carbon compounds (17)						
No carbon	0	0	0	0	0	0
D-Glucose	1	1	1	1	1	1
D-Xylose	1	1	1	1	1	1
L-Arabinose	1	1	1	1	1	1
L-Rhamnose	1	1	1	1	1	0
	1	1	1	1	1	0
D-Fructose	1	1	1	1	0	1
	1	1	1	1	0	1
Raffinose	1	0	1	0	1	1
	1	0	1	0	1	1
D-Mannitol	1	1	1	1	1	1
	1	1	1	1	1	1
i-Inositol	1	1	1	1	1	1
	1	1	1	1	1	1
Sucrose	0	1	0	0	0	0
	0	1	0	0	0	0
	0	1	0	0	0	0
f- Growth on Czapek's medium (5)	1	0	1	1	1	1
	1	0	0	0	1	1
	0	0	0	0	0	1
	0	0	0	0	0	0
g- Anti-bacterial activity (2)	0	1	1	1	1	1
	0	1	1	1	1	1
h- Anti-fungal activity (2)	0	0	0	0	1	0
	0	0	0	0	1	0
<b>Total (58 Units)</b>	<b>44</b>	<b>44</b>	<b>46</b>	<b>48</b>	<b>50</b>	<b>46</b>

Table 11. Similarities between the gray *Streptomyces* isolates and related species in Pridham and Tresner (1974) key.

<i>Streptomyces</i> isolates	Related gray <i>Streptomyces</i> species in Pridham and Tresner (1974) key					
	<i>S.</i> <i>griseochrom</i> <i>ogenes</i>	<i>S.</i> <i>echinatus</i>	<i>S.</i> <i>filipinensi</i> <i>s</i>	<i>S.</i> <i>chromofus</i> <i>cus</i>	<i>S.</i> <i>durhamen</i> <i>sis</i>	<i>S.</i> <i>antibioticus</i>
ST10	88.4	<b>93.3</b>	87.5	88.6	91.3	90.1
ST11	<b>95.0</b>	90.2	89.6	89.4	87.8	86.6
ST12	93.9	<b>94.6</b>	90.9	85.7	92.6	91.5
ST13	87.1	91.7	88.2	83.0	89.8	<b>92.8</b>

Bold number represents the most similar species.



## Identification of streptomycete isolates by numerical taxonomy

Figure 6. Phylogenetic tree of four isolates belonging to gray series and related species in **Pridham and Tresner (1974)**.

determination of the relationship between the 14 known *Streptomyces* species used in the present study. Data presented in the phylogenetic tree (**Figure 9**) reveal that the *Streptomyces* species fell into three major clusters based on their color of aerial mycelia as follows:

**First:** includes white series species, i.e., *S. viridaris*, *S. alboniger*, *S. baarnensis*, *S. albolongus* and *S. longisporus*.

**Second:** includes red series species, i.e., *S. purpurascens*, *S. yokosukanensis* and *S. janthinus*

**Third:** includes gray series species, i.e., *S. chromofuscus*, *S. antibioticus*, *S. echinatus*, *S. griseochromogenes*, *S. durhamensis* and *S. filipinensis*.

Our results reveal that the suggested numerical taxonomy proved valuable as a base for the identification of *Streptomyces*. However, further studies are needed for more evaluation of this method by its application on the all known *Streptomyces* species presented in **Pridham and Tresner (1974)** key.

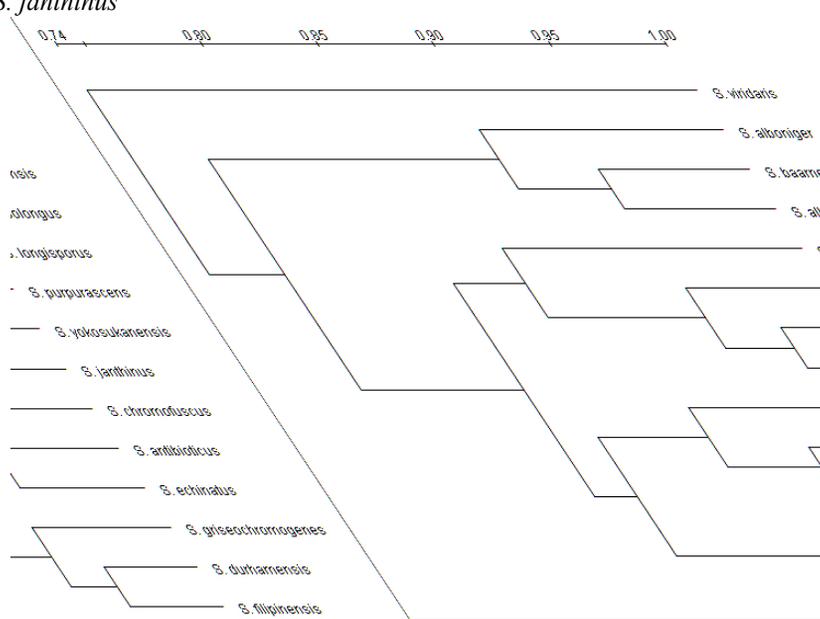


Figure 7. Phylogenetic tree of 14 *Streptomyces* species in the key of **Pridham and Tresner (1974)** achieved using suggested numerical taxonomy.

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