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Production of Carotenoids by a Newly Isolated Marine *Micrococcus* sp.

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Abstract: A total of 75 bacterial strains were isolated from different sediment samples. Of the 75 isolates, 4 strains were found to be carotenoids producers, termed (PAH81, PHAH82, PAH83 and PAH84). The isolates were identified by complete 16S rRNA gene sequence. Analysis of the 16S rDNA sequence of strain PAH81 showed 99% similarity with *Halomonas* sp., PAH82 (98%) similarity with *Bacillus aquimaris*. PAH83 showed highest similarity with *Micrococcus* sp., (98%), PAH84 showed 99% with *Micrococcus luteus*. Strain PAH83 was selected for further studies. The major carotenoids of the marine *Micrococcus* sp. PAH83 were identified by LC/MS, UV spectra and proton NMR to be decaprenoxanthin diglucoside, decaproxanthin monoglucosides and decaproxanthin with relative molecular mass (m/z) of 1028, 866 and 706 with the absorption maxima of 418, 440 and 470 nm, respectively. Production of total carotenoids by the marine *Micrococcus* sp. PAH83 was investigated. The carotenoid production was increased by aeration up to 300 rpm, maximum production was seen in medium with initial pH of 6. While nutrient rich medium supported the cell biomass, maximum carotenoids production was supported with nutrient poor medium. The carotenoids production rate by the marine *Micrococcus* sp. PAH83 was increased about five fold by varying the nutritional and incubation conditions 92.73 and 430 mg g L⁻¹, respectively

Key words: Carotenoid, decaprenoxanthin, marine, 16S rDNA, micrococcus, optimization

INTRODUCTION

Carotenoids are yellow to red colored pigments which originate from the terpenoid biosynthetic pathway. They are synthesized by plants, algae and by some fungi and bacteria. They are involved in photosynthesis as accessory pigments, functioning as antioxidants, light protection pigments and membrane stabilizers. Over 600 different carotenoids are known at present (Krubasik *et al.*, 2001). Most consist of 40 carbon atoms. The enzymatic reactions leading to their formation are well understood in plants (Sandmann, 2001) and bacteria (Armstrong, 1997). The characterization of the enzymes catalyzing their formation was facilitated after the corresponding genes had been cloned and heterologously expressed. Several groups of Gram-positive bacteria, including species from the genera *Micrococcus*, *Corynebacterium* and *Flavobacterium*, synthesize cyclic and acyclic C₄₅ and C₅₀ carotenoids (Krubasik *et al.*, 2001). The branchpoint from the typical C₄₀ pathway to the elongated carotenoids is lycopene, but very little is known about the details of the subsequent reactions. The C₅₀ carotenoid decaprenoxanthin, a cyclic carotenoid with two substituted ϵ -rings, was first identified in *Flavobacterium dehydrogenans* (Miki *et al.*, 1994). Later it was found in a few other bacteria including

Corynebacterium glutamicum (Krubasik *et al.*, 2001). The most important biological function of carotenoids is as antioxidants owing to their potential to inactivate singlet oxygen and to quench carboxy radicals (Britton, 1995; Van den Berg *et al.*, 2000). More and more evidences are being accumulated to show that carotenoids play an important role in human health. A number of epidemiological studies have revealed that an increased consumption of a diet rich in carotenoids is correlated with a diminished risk for various types of cancer, cardiovascular or ophthalmological diseases (Mayne, 1996; Sandmann, 2001). Carotenoids with unsubstituted β -ionone end groups are precursors of vitamin A. Vitamin A may inhibit tumor promotion, carotenoids are also protective because of their antioxidant effects to prevent cells and tissues from oxidative damage (Stahl and Sies, 2003). Carotenoids also influence cellular signaling and may trigger redox-sensitive regulatory pathways (Stahl *et al.*, 2002). Their value in medical and pharmaceuticals has triggered an increased interest in the synthesis of new carotenoid structures and the economic production of compounds in engineered cells. The goals of the present research are to isolate carotenoids producing bacteria from marine source, structure elucidation and optimization of the culture conditions for carotenoids production.

MATERIALS AND METHODS

Isolation and enumeration of coral-associated bacteria:

Sediment samples were collected in sterile bag under water by SCUBA diving in 2006 from different sites in Bimini, Russel Cay, Eleuthera (south end), Sweetings Cay, Cross Harbour Sandi point and Peterson Cay around The Bahamas. For bacterial isolation, the different sediment samples were suspended in sterile seawater (1 g/100 mL) and stirred for one hour. Ten-fold serial dilutions of the suspension were prepared to a dilution of 10⁻⁴ and 100 µL of each dilution spread-plated in triplicate on nutrient rich agar media, Marine Agar 2216 (Difco Laboratories, Detroit) and Nutrient Agar in seawater (Difco) and nutrient poor media, 1/10 strength Marine agar and 1/10 strength Nutrient Agar in seawater (Difco). Plates were incubated at 30°C for 3-4 weeks. Representatives of each colony morphotype were serially streak-plated onto fresh marine agar media to obtain pure cultures. All pure isolates were grown in Marine broth media overnight and stored in 20% glycerol at -80°C.

Identification of the carotenoid producing strains by 16S

rDNA sequencing: The bacterial strains were identified by 16S rRNA gene sequence analysis. The bacterial isolates were grown overnight in 2 mL Marine Broth. Total DNA was extracted with the UltraClean Microbial DNA Isolation Kit (Mo Bio Laboratories Inc., Carlsbad, CA) according to the manufacturer's procedure. Eubacterial-specific primers, forward primer 16F27 (5'-AGA GTT TGA TCC TGG CTC AG-3') and reverse primer 16R1525 (5'-AAG GAG GTG ATC CAG CCG CA-3') derived from *E. coli* 16S-rDNA sequence (Lane, 1991) were used to amplify 16S rDNA gene. The reaction mixture of 50 µL contained at least 100 ng of genomic DNA (in 10 mM Tris-HCl, pH 8), 0.2 µM of each primer and PCR Supermix High fidelity (Taq and Go, Promega, CA). PCR fragments were purified using a QIAquick Gel Extraction Kit (Qiagen, Valencia, CA) and sequenced on an ABI 377 automated sequencer using the PRISM Ready Reaction Kit (Applied BioSystems, Foster City, CA). Sequence data were analyzed by comparison with 16S rRNA genes in the GenBank database. The nearest relatives of each organism were obtained by BLAST searches (Altschul *et al.*, 1997).

Fermentation conditions and optimization of carotenoid production:

The microorganism was grown on sterilized marine broth medium (5 mL) and incubated overnight at 30°C and 100 rpm. One milliliter of the inoculum (2%) was transferred to 250 Erlenmeyer flask containing 50 mL marine broth medium and incubated at 30°C. For the optimization of the carotenoid production, several

nutritional and incubation conditions were studied including medium initial pH, incubation temperature, agitation rate, carbon source, poor and rich media and effect of some additives.

Analytical methods: The carotenoid was analysed by a modification of the method reported by Aksu and Eren (2007). Ten milliliters sample was taken from each flask at definite time intervals. Great care was taken to protect the samples from light. For the measurement of bacterial growth, dry biomass concentration was determined by measuring turbidity of the diluted samples at 600 nm using a standard curve of absorbance against dry cell mass weight. Absorbance measurements were carried out by using a Nanodrop ND-1000 spectrophotometer. The medium without cell growth was used as the blank. The total carotenoids content was also determined spectrophotometrically. For this purpose the sample was centrifuged to remove supernatant and the cell pellet was washed three times with deionized water. Cells were resuspended in methanol and blended to prevent clotting. Samples were then wrapped with aluminum foil to protect them from light and samples were vortexed until the methanol layer turned yellow (within 15 min). The methanol extract was purified from cell debris by further centrifugation at 8,000 g for 10 min, extraction were done several time until no yellow colour was extracted from the cells. Methanol solution was collected and absorbance was measured at 440 nm.

HPLC analysis: Methanol extract was evaporated to dryness under vacuum at 30°C in a Buchi rotavapor and analysis of carotenoids was carried out without saponification. Dry pigments were dissolved in 1 mL methanol and injected (50 µL) onto a µBondapak C18 column (300×4.6 mm, 5 µm particle size, RCM type; Water, USA). All other HPLC apparatus (Series 200 pump, 785A UV/Vis detector at 440 nm) was from Perkin Elmer. Separation was achieved using reverse phase HPLC at a flow rate of 1.0 mL min⁻¹. Solvents and conditions for separation were as follows: 0 to 10 min, 90% methanol: H₂O; 10-20 min, 90% methanol: H₂O to 100% methanol; 20-40 min, 100% methanol (Sander *et al.*, 1994).

LC-MS: For analysis of carotenoids by HPLC-APCI-MS, an Accela instrument (Thermo Electron Corporation) equipped with automatic sample injector, photodiode-array detector (PDA) and Finnigan LXQ series APCI mass selectivity detector was used. The mass spectra were recorded in the positive ion mode in the mass range from m/z 50 to 1500. The voltage of the corona needle was +3.9 kV, cone voltage of +17 V, probe temperature of 300°C and

source temperature of 200°C. Separation was performed on 50×2.1 mm ×1.9 µm Hypersil Gold C18 column using the gradient solvent system as follows: 0-1 min, 70% methanol; H₂O; 1-10 min, 100 % MeOH; 10-15 min, 100 % MeOH. The flow rate was set to 0.4 mL min⁻¹ and injection volume was 1 µL. The PDA was operated at 200-800 nm.

RESULTS AND DISCUSSION

In the present study, a total of 75 bacterial strains were isolated from different sediment samples. Total bacterial count ranged from a highest of 3×10⁵ to a lowest of 3×10⁷ CFU per gram of sediment. Of the 75 isolates, 4 strains were found to be carotenoid producers, termed (PAH81, PHAH82, PAH83 and PAH84). The isolates were identified by complete 16S rRNA gene sequence. Analysis of the 16S rDNA sequence of strain PAH81 showed 99% similarity with *Halomonas* sp., PAH82 (98%) similarity with *Bacillus aquimaris*. PAH83 showed highest similarity with *Micrococcus* sp. (98%), PAH84 showed 99% with *Micrococcus luteus*. Strain PAH83 was selected for further studies. The physiological and biochemical features of strain PAH83 is shown in Table 1.

Identification of the major carotenoids produced by isolate PAH83: Carotenoids of the cells of the isolated *Micrococcus* sp. PAH83 were extracted and analyzed by HPLC (Fig. 1A). The elution profile indicated the presence of three major carotenoid peaks, with the same absorption spectra. LC-MS analysis of the major peaks indicated their relative molecular mass (m/z) to be 1028, 866 and 706 with the absorption maxima of 418, 440 and 470 nm (Fig. 1B). This data with proton NMR (data not shown) indicated that the structure of the major carotenoids to be decaprenoxanthin diglucoside, decaproxanthin monoglucosides and decaproxanthin, respectively. The deccaprenoxanthin biosynthesis pathway of by *Micrococcus luteus* has been identified by Sandmann (2001).

Production and optimization of carotenoids production by isolate PAH83: An experiment on the growth pattern and the time course of carotenoid production by *Micrococcus* sp. using marine broth was carried out. The carotenoids production was increased during cultivation and reached maximum level at the end of the stationary phase (30 h). The effects of aeration, initial pH, temperature, initial sugar and incubation period on the growth and carotenoids biosynthesis of the marine *Micrococcus* sp. were investigated.

Table 1: Physiological and biochemical characteristics of the marine isolate PAH83

Gram staining	Gr+
Shape and growth	Cocci
Motility stab	Non-motile
Growth on MacConkey plates	+
Tellurite reduction	
Oxidase production	
Catalase production	+
Indole production	-
O/F test	
ONPG Hydrolysis	
H ₂ S production	-
Growth at 4 °C	
Growth at 40 °C	
Production of extra-cellular enzymes	
Amylase	
Gelatinase	+
Cellulase	
Urease	-
Caesinase	
Acid from	
D-Arabinose	-
D-Inositol	-
D-Lactose	-
D-Maltose	+
D-Rhamnose	-
D-Sorbitol	-
D-Glucose	
D-Mannose	-
D-Trehalose	+
D-Xylose	-/+
Maltotriose	+
Acetate	+
Lactic acid	+
Malic acid	+
Propionic acid	+
Pyruvic acid	+
Glycerol	+
Alpha Hydroxybutyrate	+
Beta Hydroxybutyrate	+
Serine	+
Glutamate	+
Sorbitol	-
Glucose	+
Growth in NaCl	
0%	+
0.5%	+
1%	+
Tween 40	+
Tween 80	+
0%	+
0.5%	+
1%	+

Effect of initial aeration on the growth and production of carotenoids: If the microorganism requires oxygen, aerating the growth medium is very important for the successful progress of the fermentation. As *Micrococcus* sp. is an aerobic microorganism, the effect of the aeration rate on the growth and total carotenoids formation of the bacterium was examined and the results obtained were shown in Table 2. Both the growth and total carotenoids formation changed significantly with increasing the aeration rate from 0-300 rpm. Maximum carotenoids production rate (188.31 mg g L⁻¹) was shown

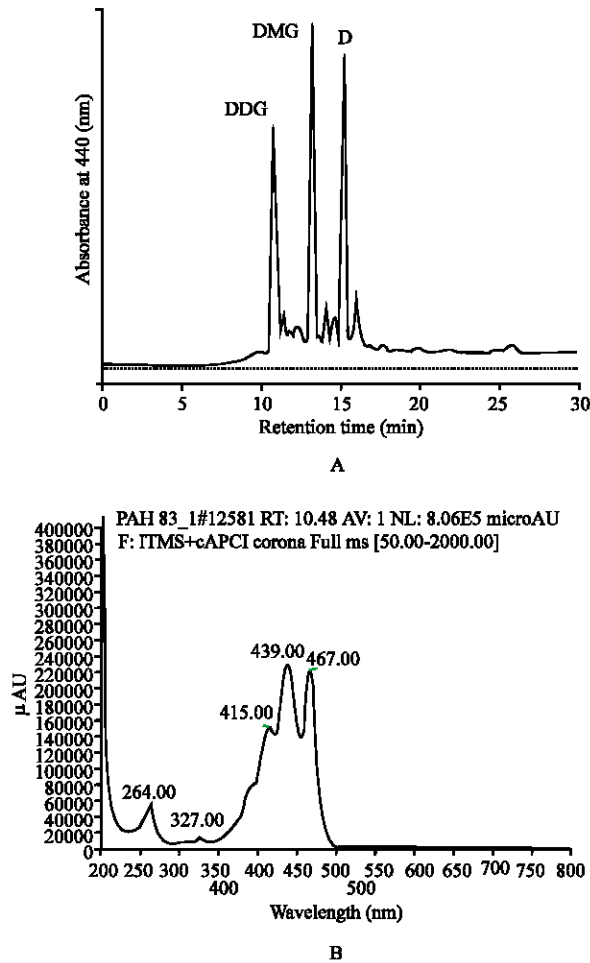


Fig. 1: A: Separation of carotenoid of *Micrococcus* sp. PHA83 on μ Bondapak C18 column. DDG: Decaprenoxanthin diglucoside, DMG: Decapromanthin monoglucosides and D: Decaprotanthin. B: Absorption spectra of DDG, DMG and D with maxima at 418, 440 and 470 nm

Table 2: Effect of aeration (rpm) on the cells dry weight, total carotenoid and carotenoid production rate by the marine *Micrococcus* sp.

Aeration (rpm)	X_M (g L ⁻¹)	P_M (mg L ⁻¹)	Y_{PX} (mg L ⁻¹)
0	0.009	0.85	92.73
100	0.027	5.10	188.31
150	0.034	5.20	152.20
250	0.035	5.50	157.14
300	0.036	5.40	150.00

X_M : cell dry weight (g L⁻¹); P_M : Total carotenoid; Y_{PX} : Carotenoid production rate Standard deviation is 2-4%

to be at rpm of 100, which was two fold more that obtained in anaerated culture (90.73 mg g⁻¹). The maximum dry cell biomass was shown at 250 rpm. These results are consistent with that reported by Aksu and Eren (2007) where the carotenoid production by *Rhodotorula glutinis* was increased by increasing the aeration level. Aeration

Table 3: Effect of initial pH on the cells dry weight, total carotenoid and carotenoid production rate by the marine *Micrococcus* sp.

pH	X_M (g L ⁻¹)	P_M (mg L ⁻¹)	Y_{PX} (mg g L ⁻¹)
4	0.001	0.00	0.00
5	0.023	4.16	178.37
6	0.026	5.85	222.86
7	0.030	5.70	190.00
8	0.048	5.34	110.44
9	0.025	2.60	104.00

X_M : cell dry weight (g L⁻¹); P_M : Total carotenoid; Y_{PX} : Carotenoid production rate. Standard deviation is 4-6%

Table 4: Effect of incubation temperature on the cells dry weight, total carotenoid and carotenoid production rate by the marine *Micrococcus* sp.

T (°C)	X_M (g L ⁻¹)	P_M (mg L ⁻¹)	Y_{PX} (mg g L ⁻¹)
25	0.017	2.80	164.71
30	0.026	5.74	220.77
37	0.033	5.20	157.58
45	0.008	1.00	120.00

X_M : cell dry weight (g L⁻¹); P_M : Total carotenoid. Y_{PX} : Carotenoid production rate. Standard deviation is 2-6%

could be beneficial to the growth and performance of microbial cells by improving the mass transfer characteristics with respect to substrate, product and oxygen (Davoli *et al.*, 2004).

Effect of initial pH on the growth and production of carotenoid:

The pH value of growth medium affects not only biosynthesis activity of culture, but also culture growth rate. Table 3 shows that with ranging from 3-8. With raising the pH, carotenoid production rates and growth increased and reached to a maximum level at pH 6 and 8, respectively. On the other hand, a further increase of the pH resulted in a reduction of both rates. Although the optimal pH of 6 yielded the highest values of total carotenoids concentration and product yields, maximum biomass concentration was obtained at an initial pH value of 8. It has been reported that the optimal pH value of carotenoid production was different of that of growth in carotenoids production by *Rhodotorula glutinis* (Aksu and Eren, 2007).

Effect of temperature on the growth and production of carotenoid:

Temperature is another important parameter affecting the performance of cells and product formation. The temperature of the growth medium had also a considerable effect on both the growth and carotenoid production of the marine *Micrococcus* sp. the cell biomass increased notably with raising the temperature with up to 37°C and lessened sharply at 45°C due to the denaturation of the enzyme system of microorganism at higher temperatures (Table 4). The total carotenoids production rate was also affected by the variation of fermentation temperature; enhanced strictly up to 30°C (220.77 mg g L⁻¹). Maximum cell biomass where found to be at 37°C (0.31 mg L⁻¹).

Effect of different media on the growth and production of carotenoids:

The effect of nutrient rich (Marine broth and Nutrient broth media) and nutrient poor media (1/10 strength marine broth and 1/10 strength Nutrient broth) was investigated. The results presented in Table 5 showed that nutrient poor media showed less cell biomass but more carotenoids production rate, by the marine *Micrococcus* sp., than the nutrient rich media with diluted marine broth as the best one. It was found that using diluted marine broth medium resulted in increase of the total carotenoids production rate by about 1.5 fold more than marine broth medium, 331.58 and 220.77 mg g L⁻¹, respectively. However maximum cell biomass was found to be in culture using marine broth medium.

Effect of different sugars on the growth and production of carotenoid:

The effect of addition different sugars to the diluted Marine broth media, triose, tetrose, pentose, monosaccharide, disaccharide and polysaccharides, was investigated. Glucose was found to enhance the cell

Table 5: Effect of different media on the cells dry weight, total carotenoid and carotenoid production rate by the marine *Micrococcus* sp.

Media	X _M (g L ⁻¹)	P _M (mg L ⁻¹)	Y _{PR} (mg L ⁻¹)
MB	0.026	5.72	220.00
MB 1/2	0.023	6.10	265.22
MB 1/10	0.019	6.30	331.58
NB	0.022	5.10	231.82
NB 1/2	0.019	4.90	257.89
NB 1/10	0.016	4.30	268.75

MB: Marine broth media, MB ½: half strength Marine broth media, MB 1/10: 1/10 strength Marine broth media NB: Nutrient broth media, NB ½: half strength Nutrient broth media, NB1/10: 1/10 strength Nutrient broth media. X_M: cell dry weight (g L⁻¹); P_M: Total carotenoid; Y_{PR}: Carotenoid production rate. Standard deviation is 3-6%

Table 6: Effect of different sugars on the cells dry weight and carotenoid production rate by the marine *Micrococcus* sp.

Sugars	X _M (g L ⁻¹)	P _M (mg L ⁻¹)	Y _{PR} (mg L ⁻¹)
MB 1/10	0.019	6.25	328.95
Glycerol	0.016	4.90	301.54
Arabinose	0.021	6.10	292.80
Xylose	0.044	7.10	160.75
1% Glucose	0.027	10.25	379.63
Maltose	0.022	6.50	295.45
Starch	0.019	6.00	315.79

X_M: cell dry weight (g L⁻¹); P_M: Total carotenoid; Y_{PR}: Carotenoid production. Standard deviation is 2-6%

Table 7: Effect of different glucose concentrations on the cells dry weight and carotenoid production rate by the marine *Micrococcus* sp.

Glucose conc. (%)	X _M (g L ⁻¹)	P _M (mg L ⁻¹)	Y _{PR} (mg L ⁻¹)
0.25	0.02	6.50	295.45
0.01	0.02	8.00	333.33
0.01	0.03	10.20	377.78
1.50	0.03	12.50	431.03
2.00	0.03	11.50	410.71
2.50	0.03	10.67	395.19
0.03	0.03	7.25	290.00

X_M: cell dry weight (g L⁻¹); P_M: Total carotenoid; Y_{PR}: Carotenoid production. Standard deviation is 2-4%

biomass, total carotenoids and carotenoids production rate (Table 6). It was found that the addition of xylose to the diluted marine broth medium reduce the carotenoid cell productivity to about 50% of the control. Addition of 1% glucose increased the total carotenoids slightly. The effect of different glucose concentration on the growth and carotenoid production was investigated (Table 7). Both cell biomass and carotenoids production was increased by increasing the glucose concentration up to 1.5% with cell biomass and carotenoid production rate of 0.029 g L⁻¹ and 430 mg g L⁻¹, respectively.

In summary, the carotenoids production rate by the marine *Micrococcus* sp. PAH83 was increased about five fold by varying the nutritional and incubation conditions 92.73 and 430 mg g L⁻¹, respectively.

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