

## **Increased production of carrageenase by *Pseudomonas aeruginosa* ZSL-2 using Taguchi experimental design**

**M. Ziyoddin<sup>1,\*</sup>, Junna Lalitha<sup>1</sup>, Manohar Shinde<sup>2</sup>**

<sup>1</sup>Department of Biochemistry, Gulbarga University, Gulbarga - 585106, Karnataka, India

<sup>2</sup>Department of Studies and Research in Biochemistry, Tumkur University, Tumkur - 572103, Karnataka, India

\*E-mail address: ziyagug@gmail.com

### **ABSTRACT**

The culture conditions for the production of carrageenase were optimized using one-factor-at-a-time method combined with orthogonal array design. With the one-factor-at-a-time method revealed optimal conditions for carrageenase production were 24 h of fermentation period, 28 °C incubation temperature at pH 8.0 with NaNO<sub>3</sub> as nitrogen source and carrageenan as carbon source in MMS media. Further optimization of carrageenase production by using orthogonal experimental design L<sub>9</sub> (3<sup>4</sup>) with four factors, temperature, pH, NH<sub>4</sub>NO<sub>3</sub> and carrageenan with their relevant levels revealed optimised conditions for carrageenase production were temperature of 28 °C, pH 8.0, 2 g L<sup>-1</sup> NaNO<sub>3</sub> and 2 g L<sup>-1</sup> carrageenan. The order of the factors affecting the fermentation process was found to be temperature > pH > NaNO<sub>3</sub> > carrageenan. The temperature played a significant role on the carrageenase production. Higher carrageenase yield with activity of 0.542 ± 0.045 U ml<sup>-1</sup> was obtained in the optimised medium when compared to those of basal medium. Carrageenase hydrolysed products of carrageenan were identified by LC-ESI-MS as neocarrabiose, neocarrabiose-4 sulfate, neocarratetraose, neocarratetraose-4 sulfate, anhydrogalactose, galactose, galactose-4 sulphate and sulphate.

**Keywords:** Carrageenase; *Pseudomonas aeruginosa* ZSL-2; L<sub>9</sub> orthogonal array design; LC-ESI-MS

### **1. INTRODUCTION**

The algal polysaccharides, also known as phycocolloids, constitute a crucial carbon source for many marine bacteria which degrade the cell wall of algae by secreting specific glycoside hydrolases [1]. Carrageenan, one of the phycocolloids is a sulfated galactan made up of linear chains of galactose and 3,6-anhydrogalactose with alternating  $\alpha$ -(1→3) and  $\beta$ -(1→4) linkages and further classified based on the number and the position of sulfated ester(s);  $\kappa$ -,  $\iota$ - and  $\lambda$ -carrageenan [2]. Microbial enzymes which hydrolyse phycocolloids have drawn considerable interest. The biotechnology of enzymes and enzyme degraded products of

carrageenan is still in infancy compared to that of agar, alginate, starch [3] or pectin [4]. Carrageenases which hydrolyse 1,4 linkages in carrageenan to a series of homologous, even numbered oligosaccharides [5] are useful tool for the structural analysis of the cell walls and protoplast isolation from red algae [6,7]. The sulfated carrageeno-oligosaccharides have also drawn considerable interest [8] owing to their diverse biological and physiological activities including anticoagulation [9], anti-inflammation [10], anti-thrombosis [11], antitumor activity [12] and viral inactivation [13], which depend on structural parameters such as carbohydrate structure, molecular mass, degree of sulfate esterification, and the linking position of sulpho groups [14].

Several bacterial genera including *Pseudomonas carrageenovora* [8,9], *Cytophaga* [10,11], *Alteromonas carrageenovora* [12,13], *Vibrio sp.* [15] and *Zobellia galactanovorans* [16] have been reported carrageenase production one-factor-at-a-time method. Improvement of productivity of microbial metabolite by the organisms was done by manipulating the nutritional parameters, physical parameters and strain improvement [16]. Medium optimization by single dimensional search was laborious and time consuming and requires a large number of experiments to determine optimum levels, which are unreliable. Optimizing all the effecting parameters can eliminate these limitations of a single factor optimization process collectively by statistical experimental design. Several statistical factorial designs ranging from two-factorial to multi-factorial designs are available for optimization of process parameters [17,18]. Taguchi method is a popular experimental design currently applied in industries for optimization serves as screening filters that examine the effects of many process variables and identify those factors that have major effects on process by using a few experiments.

Previously we have reported the optimization of production conditions for carrageenase by solid state fermentation of *Pseudomonas aeruginosa* ZSL-2 [19]. In present investigation we report establishment of culture conditions for the production of extracellular carrageenase by SmF of *Pseudomonas aeruginosa* ZSL-2 using one-factor-at-a-time method by changing one of the independent variables, temperature, pH etc. while fixing the others at certain levels and  $L_9 (3^4)$  design was applied to screen the significant factors according to preliminary experiments. The analysis of carrageenan degraded products by LC-ESI-MS is also carried out.

## 2. MATERIALS AND METHODS

### 2. 1. Bacterial strain and culture conditions

One of the agarolytic bacteria isolated from marine water *Pseudomonas aeruginosa* ZSL-2 capable of utilizing both agar and carrageenan [20] was selected for further studies and stored at 4 °C on minimal mineral salts (MMS)-agar slants. The bacterium was routinely grown on MMS medium with agar as energy source at 37 °C. Seed culture was prepared from 24 h grown cultures on MMS liquid broth at 180 rpm and 37 °C.

### 2. 2. Culture conditions for acclimatization to carrageenan

The fermentation medium composed  $g.L^{-1}$ ;  $K_2HPO_4$ , 0.38;  $MgSO_4$ , 0.20;  $FeCl_3$ , 0.05;  $NH_4NO_3$ , 1.0; pH 8.0 was supplemented with carrageenan (0.3 %) as the only source of carbon. The bacterial seed culture was used for inoculation. The cultures were maintained on minimal salts agar (1.5 %) and carrageenan (0.5 %) plate or slants, or LB agar (1.5 %) and carrageenan (0.5 %) plate or slants and sub cultured fortnightly.

### 2. 3. Assay of carrageenase activity

The carrageenase activity was determined by neocuproine method as described by Dygert et al [21]. The assay mixture 1 ml contained 20 mM Tris-HCl buffer (pH 8.0), 0.5 ml of carrageenan (0.1 % w/v in Tris-HCl buffer) and 50  $\mu$ l suitably diluted enzyme at 40 °C. For carrageenase assay it was necessary to add ethanol (50 %, v/v) to reaction mixture before measuring the absorbance.

One unit of the enzyme activity was defined as the amount which liberates 1  $\mu$ mol galactose equivalent per minute under assay conditions. The protein content of the enzyme solution was determined according to Lowry et al [22].

### 2. 4. Optimization of culture conditions for carrageenase production

The optimization of fermentation conditions and medium compositions were carried out based on the stepwise modification of the governing parameters for carrageenase production. They include fermentation period, pH, temperature, NaCl, carbon source and nitrogen sources. The optimum culture conditions obtained in previous step were used in next step.

The *P. aeruginosa* ZSL-2 was inoculated into the MMS medium supplemented with carrageenan (0.3 % w/v) and incubated. A 1 ml exponential growth phase culture was inoculated into a fresh 250 ml Erlenmeyer flask containing 50 ml MMS medium supplemented with carrageenan (0.3 % w/v).

The growth of the bacterium and carrageenase production in culture supernatants were analysed at different incubation periods. The effect of incubation temperature was determined by incubating the bacterium at different temperature ranging from 20 to 40 °C, at pH 8.0 after 24 h. Similarly, influence of medium pH was determined from pH 5 to 10 at 28 °C. The effect of NaCl on the growth and enzyme production was determined by incubating at 28 °C for 24 h with NaCl in MMS medium at pH 8.0.

To study the effect of different nitrogen sources on carrageenase production,  $\text{NH}_4\text{NO}_3$  in the medium was replaced with different organic nitrogen sources (0.1%, w/v) such as, peptone, yeast extract, beef extract, and inorganic nitrogen sources (1 %, w/v) such as,  $\text{KNO}_3$ ,  $\text{NH}_4\text{Cl}$ ,  $\text{NaNO}_3$ ,  $(\text{NH}_4)_2\text{SO}_4$  and fermentation was carried out as described earlier.

The influence of carbon source on the carrageenase production was analysed by growing *P. aeruginosa* ZSL-2 in the MMS medium supplemented with various carbon sources (0.2 % to 0.5 %, w/v) such as glucose, galactose, fructose, lactose, sucrose, mannose, maltose and carrageenan alone or the said carbon sources were co-supplemented with carrageenan.

### 2. 5. Orthogonal test (Taguchi method)

Based on the one-factor-at-a-time method, an  $L_9 (3^4)$  orthogonal array method was used to determine the optimal medium conditions for carrageenase production. This enables to determine the process variables affecting the response.

The level-setting values of the factors used in the orthogonal array design are shown in Table 4. The four variables, carrageenan,  $\text{NaNO}_3$ , temperature and pH with three concentration levels were used. The experiments were carried out in 250 ml Erlenmeyer flasks containing 50 ml MMS medium. The carrageenase activity of each flask was assayed after 24 h of incubation and analysed statistically by ANOVA.

## 2. 6. LC-ESI-MS analysis of carrageenan oligosaccharides

The hydrolysed products of carrageenan by *P. aeruginosa* ZSL-2 supplemented in the growth medium were analyzed by LC-ESI-MS as reported by Baswaraj et al., [23]. Briefly; a 1 ml of the culture medium was withdrawn, cooled to 4 °C and centrifuged for 15 min at 14,000 rpm at 4 °C. A 500 µl of clarified culture supernatant was extracted by 80 % methanol (-20 °C) and evaporated to dryness by a vacuum manifold.

The residue was then re-dissolved in 100 µl of 80 % methanol and was filtered through a 5 kDa centricon filter and the filtrate obtained was transferred to an autosampler vial. Ten microliters of culture supernatant filtered through 0.22 µ syringe filter was injected through auto sampler into a 150 mm × 4 mm i.d, 5 µ phenomenon, C-18, RP-column (thermostated at 40 °C) at flow rate 1 ml min<sup>-1</sup>.

LC is synchronized with mass Q TRAP 4000 MS-MS (Applied Biosystems MDS-SCIX). LC-ESI-MS data acquisition was carried out at 22 min. EMS spectra were generated from the TIC. Spectra were recorded in negative mode between m/z 50 and 1200. Identification was carried out with AB's Analyst 1.2.2 software.

## 3. RESULTS

### 3. 1. Optimization of culture parameters for carrageenase production from *Pseudomonas aeruginosa* ZSL-2 by “one-factor-at-a-time” (OFAT)

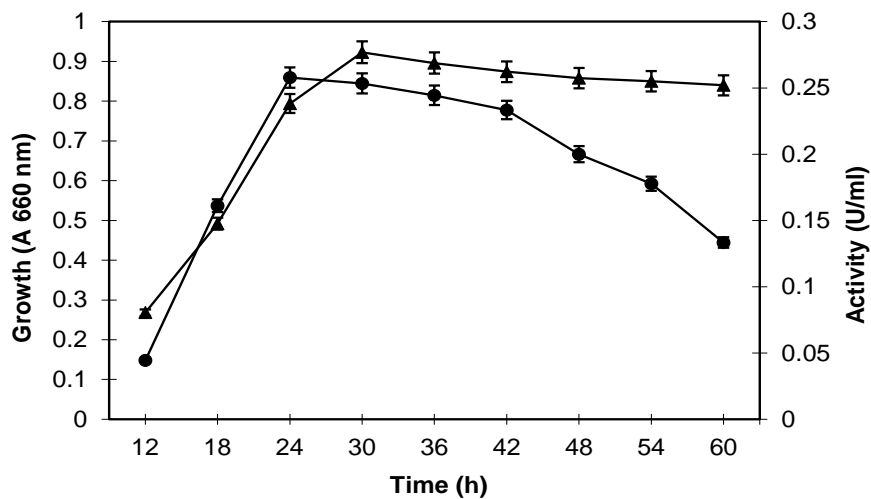
The effect of different culture parameters viz., fermentation period, temperature, optimum pH of fermentation medium, ionic strength and carbon and nitrogen sources, on the growth of *P. aeruginosa* ZSL-2, and production of extracellular carrageenase was investigated.

The growth of the bacterium at different incubation periods was determined by monitoring the change in the absorbance at 660 nm and the carrageenase activity in the fermentation broth was assayed simultaneously. It was observed that, the growth of the bacterium was increased with the increase in the incubation time. The carrageenase production by *P. aeruginosa* ZSL-2, was observed after 12 h of incubation, which increased with an increase in the incubation period.

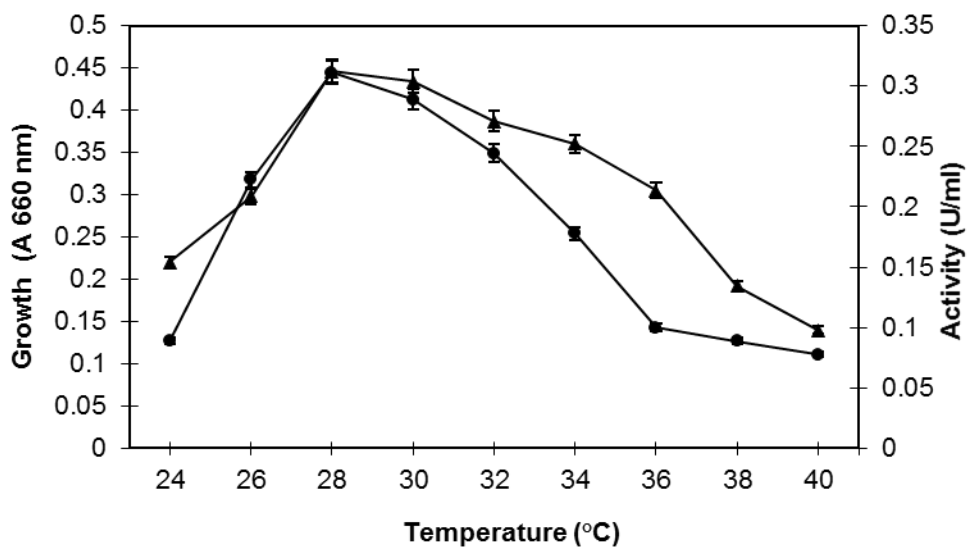
Maximum carrageenase activity (0.257 U/ml) was observed in late log phase at 24 h incubation, thereafter the carrageenase production decreased (Fig. 1). The bacterium was able to grow at temperatures ranging from 25 to 40 °C, maximum carrageenase production (0.311 U/ml) was observed at 28 °C (Fig. 2). *P. aeruginosa* ZSL-2 grew in a broad range of pH 5.0-11.0. showed good growth at pH range from 7.0-10.0; however, the maximal carrageenase production (0.355 U/ml) was observed when grown at pH 8.0 (Fig. 3).

The effect of sodium chloride on the growth of the microorganism in the fermentation medium was studied with the amendment of carrageenan (0.3 %, w/v) at pH 8, temperature 28 °C It is evident that the microorganism grew in presence of sodium chloride (0 to 5%). The maximal growth appeared at 1 % NaCl.

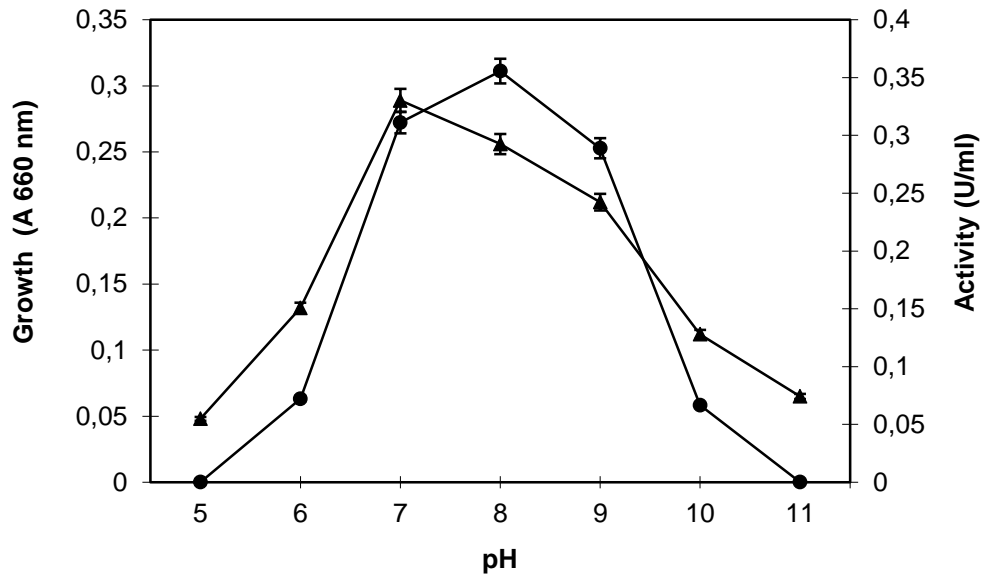
The organism did not show a specific requirement of sodium chloride and grew even at 0-2 % levels of sodium chloride. The carrageenase production was found to be slightly increased with increase in the concentration of NaCl up to 1 % (w/v), beyond which there was sharp decline in the secretion. At 4 % NaCl, the carrageenase activity was not detected though the organism was able to grow (Fig. 4).



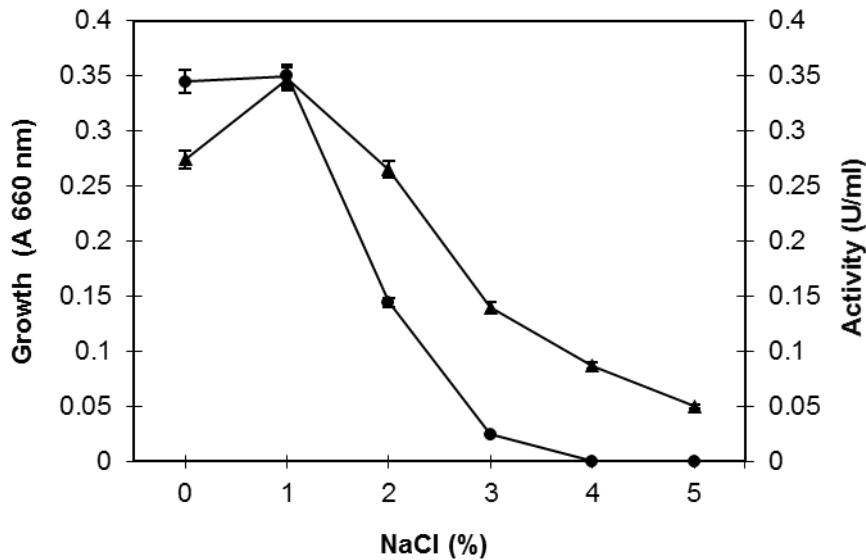
**Fig. 1.** Effect of incubation time on the growth of *Pseudomonas aeruginosa* ZSL-2 (▲) and the production of extracellular carrageenase (●).



**Fig. 2.** Effect of temperature on the growth of *Pseudomonas aeruginosa* ZSL-2 (▲) and the activity of of extracellular carrageenase (●).



**Fig. 3.** Effect of media pH on growth (▲) and production (●) of extracellular carrageenase by *Pseudomonas aeruginosa* ZSL-2.



**Fig. 4.** Effect of NaCl on growth (▲) and production (●) of extracellular carrageenase by *Pseudomonas aeruginosa* ZSL-2.

The growth of the bacterium was found to be optimum in the medium supplemented with  $\text{NaNO}_3$  or  $\text{NH}_4\text{NO}_3$  as nitrogen source and carrageenan as the carbon source. The production of carrageenase was found to maximum (0.393 U/ml) in the medium, which had received  $\text{NaNO}_3$  as nitrogen source. The production of carrageenase was found to be less in the fermentation medium, containing organic nitrogen sources.

However, peptone served as a good source of nitrogen and supported luxuriant growth however carrageenase production was found to be less (Table 1).

**Table 1.** Effect of nitrogen source on the growth and production of carrageenase by *Pseudomonas aeruginosa* ZSL-2.

Nitrogen source*	Growth (A <sub>660nm</sub> )	Carrageenase activity (U/ml)
NH <sub>4</sub> NO <sub>3</sub>	0.317	0.288
KNO <sub>3</sub>	0.284	0.222
NH <sub>4</sub> Cl	0.275	0.094
NaNO <sub>3</sub>	0.297	0.393
(NH <sub>4</sub> ) <sub>2</sub> SO <sub>4</sub>	0.206	0.166
Peptone	0.273	0.177
Yeast extract	1.111	0.100
Beef extract	0.902	0.088

\* Fermentation conditions: MMS media containing one of the nitrogen source, supplemented with carrageenan at pH 8, temperature 28 °C after 24h of incubation

The *P. aeruginosa* ZSL-2 utilized various other organic chemicals as carbon source. carrageenase production was observed in the fermentation media supplemented with carrageenan. The organism grew well utilizing other simple sugars, while carrageenase production was not observed (Table 2). On the other hand carrageenase production was observed when simple sugars were co-supplemented to carrageenan containing fermentation media. The results revealed that carrageenase production was inducible only in presence of carrageenan and co-supplementation of other simple sugars found to decrease the production slightly.

**Table 2.** Effect of carbon sources on growth and production of extracellular carrageenase by *Pseudomonas aeruginosa* ZSL-2.

Carbon source	Growth (A <sub>660nm</sub> )	Carrageenase production (U/ml)	Co-Supplementation of sugars *	Growth (A <sub>660nm</sub> )	Carrageenase production (U/ ml)
Carrageenan	0.382	0.3444	C alone	0.342	0.322
Fructose	1.02	0.0174	C + Fructose	0.402	0.277
Glucose	1.175	ND	C + Glucose	0.475	0.305
Galactose	1.035	ND	C + Lactose	0.711	0.233
Lactose	0.911	0.0054	C + Sucrose	0.236	0.222
Sucrose	0.836	0.003	C + Mannose	0.298	0.244





4	2(30 °C)	1(pH 7.5)	2	3	0.244	± 0.007
5	2(30 °C)	2(pH 8.0)	3	1	0.210	± 0.006
6	2(30 °C)	3(pH 8.5)	1	2	0.156	± 0.004
7	3(32 °C)	1(pH 7.5)	3	2	0.128	± 0.003
8	3(32 °C)	2(pH 8.0)	1	3	0.132	± 0.003
9	3(32 °C)	3(pH 8.5)	2	1	0.068	± 0.002
<sup>a</sup> K 1	1.239	0.714	0.63	0.62		
K 2	0.61	0.862	0.832	0.804		
K 3	0.328	0.601	0.715	0.753		
<sup>b</sup> k 1	<b>0.413</b>	0.238	0.21	0.206		
k 2	0.203	<b>0.287</b>	<b>0.277</b>	<b>0.268</b>		
k 3	0.109	0.200	0.238	0.251		
<sup>c</sup> Δ	0.303	0.087	0.067	0.061		
R	1	2	3	4		
Opt	A <sub>1</sub>	B <sub>2</sub>	C <sub>2</sub>	D <sub>2</sub>		

The assignments of column A, B, C, and D were performed by orthogonal array consisted of nine experiments corresponding to the nine rows and four columns.

<sup>a</sup>  $k_i = \sum k$  of all experiment at the same factor level.

<sup>b</sup> Average of  $k_i$ .

<sup>c</sup>  $\Delta = \max \{ \text{average of } k_i \} - \min \{ \text{average of } k_i \}$ .

Values are mean of triple determinations with standard deviation ( $\pm$ ).

Opt – optimization

**Table 5.** ANOVA of  $L_9(3^4)$  Orthogonal array testing strategy (OATS) for optimization of fermentation parameters for carrageenase production by submerged fermentation of *P. aeruginosa* ZSL-2.

	Sum of squares	df	Mean of square	F value	Critical value	Significance
Temperature	0.0483	2	0.0241	3.4249	$F_{0.10}(2,8) = 3.113$	*
pH	0.0038	2	0.0019	0.2701		
Nitrogen	0.0022	2	0.0011	0.1623		
Carbon	0.0020	2	0.0010	0.1425		
Error	0.0564	8	0.0070			

df – Degree of freedom F value -  $F_{0.10}$

\* Significant terms.

### 3. 3. LC-ESI-MS analysis of carrageenan hydrolysed products

The presence of oligosaccharides ranging from mono- to hexa- carrageenan were identified based on signals observed at m/z 161, 179 and 79, which corresponds to the anhydrogalactose, desulfated galactose and sulfate, respectively (Table 6). The signals observed at m/z 259.1, 323, 403, 629.9 and 788.8 were corresponding to galactose-4 sulphate, neocarrabiose, neocarrabiose-4 sulfate, neocarratetraose and neocarratetraose-4 sulfate, respectively. The carrageenase secreted by *P. aeruginosa* ZSL-2, thus acted as endo-enzymes and capable of hydrolysing carrageenan into mainly tetrose, biose and then to monomers.

**Table 6.** LC-ESI-MS spectrometry analysis of carrageenase hydrolyzed products of carrageenan in culture supernatant by *P. aeruginosa* ZSL-2 grown for 24 h.

Carrageenana oligosaccharide	m/z
[G <sub>4</sub> S]	259.1
[DA-G]	323.0
[DA-G <sub>4</sub> S]	403.0
[DA-G] <sub>2</sub>	629.9
[DA-G <sub>4</sub> S] <sub>2</sub>	788.8
[SO <sub>3</sub> <sup>-</sup> ]	79
[DA]	161.1
[G]	179.1

DA - anhydrogalactose, G - Galactose, G<sub>4</sub>S - Galactose 4 sulphate, DA-G - neocarrabiose, DA-G<sub>4</sub>S - neocarrabiose 4 sulfate, [DA-G]<sub>2</sub> - neocarratetraose, [DA-G<sub>4</sub>S]<sub>2</sub> - neocarratetraose 4 sulfate, and SO<sub>3</sub><sup>-</sup> sulphate ions.

## 4. DISCUSSION

The optimum pH and temperature for carrageenase production were around pH 8 and 28 °C, respectively, and this is because the isolated strain had adapted to the alkaline and low temperature marine environment for the growth. The pH of the culture strongly affects enzymatic processes and transport of compounds across the cell membrane. Several carrageenan degrading bacteria *Pseudomonas elongata* [24], *Cytophaga* sp. [25], *Cytophaga* like bacterium [26-28] were found to produce carrageenases at pH range of 7-8 and few at different temperatures, ranging from 25 °C [15,29] to 37 °C [24,27] are reported. The *P. aeruginosa* ZSL-2 grows better, when the growth medium was supplemented with 1 % sodium chloride. It grew nearly optimal even in the absence of sodium chloride. It is commonly assumed that marine bacteria live in the sea, most of them are salt tolerant. Most carrageenan utilizing bacteria like *P. elongata* [24], *P. alteromonas*-like bacterium WZUC10

[30], *P. carrageenovora* [7,31], *Vibrio sp.* CA-1004 [15] and *Cytophaga sp* [26] isolated from marine sources are reported to have a specific requirement of sodium chloride for their growth. The *P. aeruginosa* ZSL-2 grows better at 1 % NaCl concentration and unlike other carrageenolytic marine bacterium *P. aeruginosa* ZSL-2 do not demand for sodium chloride as a specific requirement, more over, even in its absence, the organism could grow well and produces extracellular carrageenase. An understanding of the influence of nitrogen source on the growth of *P. aeruginosa* ZSL-2 was derived by growing the later in the fermentation medium containing one of the nitrogen sources and supplemented with carrageenan (0.3 %, w/v) as sole carbon source. The observed results indicated that  $\text{NaNO}_3$  as a sole nitrogen source supported the optimum growth and maximum enzyme production, thus nitrogen was found to be an essential element for the production of the carrageenase.

All the simple sugars tested were utilized as carbon sources for the growth of the *P. aeruginosa* ZSL-2. Carrageenase production was observed when it was grown on the carrageenan as carbon source and also in co-supplemented with other carbon sources. When simple sugars alone were included in the medium, although good growth was observed, carrageenase production was not occurred in the medium. This explains that the carrageenase production was inducible only in presence of carrageenan and co-supplementation of other simple sugars found to decrease the production slightly. Catabolite repression of carrageenase production was observed in *P. aeruginosa* ZSL-2, in which carrageenase production was repressed when other carbon sources were supplemented to carrageenan. Similar observations were reported in agarase production by *P. aeruginosa* AG-LSL11 [32], *Alteromonas Sp.* (049/1) and *Cytophaga saccharophila* (024), in which the enzyme production was repressed by glucose, co-supplemented with agar [33]. While increased growth was observed in the medium co-supplemented with glucose, fructose or maltose. This may be due to rapid utilization of these simple sugars by the bacterium resulting in an increase in cell mass by vigorous growth. However, carrageenase production was found to be reduced.

The results with the one-factor-at-a-time method revealed that optimal conditions for the production of carrageenase were 24 h of fermentation period, 28 °C incubation temperature at pH 8.0 with  $\text{NaNO}_3$  as nitrogen source and carrageenan as carbon source in MMS media. Among these four factors, temperature, pH,  $\text{NH}_4\text{NO}_3$  and carrageenan with their relevant levels are selected for further optimization of carrageenase production by using orthogonal experimental design  $L_9 (3^4)$ . The one-factor-at-a-time (single factor) experiments enable us to determine which process variables affect the response. A logical next step is to determine the point in the important factors that leads to the best possible response with maximum production of the enzyme [34,35]. The Latin square used to optimize the nutritional factors was  $L_9 (3^4)$ , where  $L_9$  indicates a Latin square with nine combinations of variables and  $(3^4)$  denotes four factors with three levels. Appropriately selecting the factor one can improve the quality of the product.

The optimum culture conditions obtained from the statistical analysis were at temperature of 28 °C, pH 8.0,  $2 \text{ gL}^{-1} \text{ NaNO}_3$  and carrageenan  $2 \text{ gL}^{-1}$ . The order of effect of four factors on carrageenase production was temperature > pH >  $\text{NaNO}_3$  > carrageenan. According to the interaction among the tested factors, the temperature had significant effect on the carrageenase production. The validation experiments were conducted to obtain the maximum yields of carrageenase using the above optimal culture conditions. The carrageenase activity, 0.542 U/ml was obtained which was much higher than those in basal culture media, suggests that the carrageenase production has been further improved with orthogonal experiment design. The orthogonal array design technique that has been successfully applied for the improvement of culture conditions of fermentation process to

provide the relationships among various factors, and the order of significant factors for the optimum results [4,19,36,37].

The electrospray ionization mass spectrometry (ESI-MS) technique was employed for the analysis of carrageenan hydrolysed oligosaccharides samples. The ESI-MS analysis of the culture supernatants of *P. aeruginosa* ZSL-2 revealed that commercial carrageenan was hydrolyzed initially by the action of extracellular carrageenase to oligosaccharides ranging from mono- to tetra-carrageenan oligosaccharides. ESI-MS creates multiple charged ions making the interpretation of a mixture of highly charged oligosaccharides through direct ESI-MS analysis a rather complex task [38,39]. This has been demonstrated for the first time for the characterization of enzymatically digested correspondent  $\kappa$ - and  $\iota$ -carrageenan using LC/ESI-MS method, which is based on ion pair liquid chromatography coupled with ESI-MS in the negative-ion mode. This method can directly determine and verify the exact number of sulphate groups, as well as the molecular weight of the primary backbone structure of each oligosaccharide in the mixture.

## 5. CONCLUSION

Using the one-factor-at-a-time method and orthogonal matrix design, it was possible to determine optimal conditions to obtain maximum yield of carrageenase. Two optimization techniques used in this work can be widely applied to other processes for optimization of submerged culture conditions for production of other enzymes. The *P. aeruginosa* ZLS-2 could be used for the production of biotechnologically important, commercial, alkali active enzymes for their industrial applications.

## ACKNOWLEDGEMENTS

This work was supported in part by research grant from Department of Science and Technology (DST), New Delhi, India (Project No. 100/IFD/5186/2007-2008 dated 6/11/2007). M. Ziyoddin is grateful to UGC New Delhi, India for research fellowship (MANF JRF (No.F.40-49(M)/2009(SA-III/MANF) dated 08 Jan 2011). The authors thank Gulbarga University, Gulbarga for laboratory facility.

## References

- [1] Michel G., Nyval-Collen P., Barbeyron T., Czjzek M., Helbert W., *Appl Microbiol Biotechnol.* 71 (2006) 23-33.
- [2] Knutsen S., Myslabodski D., Larsen B., Usov A., *Bot Mar.* 37 (1994) 163-169.
- [3] Guzman-Maldonado H., Paredes-Lopez O., *Critical Review in Food Science Nutrition* 35 (1995) 373-403.
- [4] Beldman G, Mutter M, Searle-van Leeuwen MJF, van den Broek LAM, Schols HA, Voragen AGJ. Visser J, & Voragen, A.G.J. Amsterdam: Elsevier; 1996.
- [5] Weigl J., Yaphe W., *Can J of Microbiol.* 12 (1966) 939-947.
- [6] Le Gall Y., Braud J. P., Kloareg B., *Plant Cell Rep* 8 (1990) 582-585.

- [7] Ostgaard K., Wangen B. F., Knutsen S. H., Aasen I. M., *Enzy Microbial Technol.* 5 (1993) 326-333.
- [8] Mou H., Jiang X., Guan H., *J of Appl Phycol.* 15( 2003) 297-303.
- [9] Alban S., Schauerte A., Franz G., *Carbohydrte Polymer* 47 (2002) 267-276.
- [10] Arfors K. E., Ley K., *J Lab Clinical Medicine* 121 (1993) 201-202.
- [11] Suzuki N, Kitazato K, Takamatsu J, Saito H. Thrombosis and Haemostasis. 65 (1991)369-373.
- [12] Hiroishi S, Sugie K, Yoshida T, Morimoto J, Taniguchi Y, Imai S, Kurebayashi J. Cancer Letters. 167 (2001) 167:145-150.
- [13] Caceres P. J., Carlucci M. J., Damonte E. B., Matsuhira B., Zuniga E. A., *Phytochemistry* 53 (2000) 81-86.
- [14] Knutsen S., Myslabodski D., Larsen B., *Usov A Bot Mar* 37 (1994) 163-169.
- [15] Araki T., Higashimoto Y., Morishita T., *Fisheries Science.* 65 (1999) 937-942.
- [16] Greasham R. L., Biotechnology. In: Rehm HJ, Read G, Puhler A, Stagler P (Eds.), *Bioprocessing*, vol. 3. VCH Publishers, Inc., New York; 1983.
- [17] Thomas D. M., *J of Chem Engi.* 6 (1977) 180.
- [18] Deming S. N., Morgan SL., Elsevier, Oxford; 1987.
- [19] Ziyaddin M., Shinde M., Lalitha J., *The Bioscan.* 4 (2012) 096-095.
- [20] Ziyaddin M., Manohar S, Lalitha J., *The Bioscan.* 5 (2010) 279-283.
- [21] Dygert S., Li L. H., Florida D., Thoma J. A., *Anal Biochem.* 13 (1965) 367-374.
- [22] Lowry O. H., Rosebrough N. J., Farr A.L., Randall R. J., *J of Biol Chem.* 193 (1951) 265-275.
- [23] Basawaraj A. K., Manohar S., Lalitha J. *Biotechnology and Bioprocess Engineering* 18 (2013) 333-341.
- [24] Khambhaty Y, Mody K, Jha B., *Biotechnol Bioproc Engi.* 12 (2007) 668-675.
- [25] Sarwar G., Matayoshi S., Oda H., *Microbiol Immunol.* 31 (1987) 869-877.
- [26] Potin P., Sanseau A., Le Gall Y., Rochas C., Kloareg B., *Europ J of Biochem.* 201 (1991) 241-247.
- [27] Mou H. J., Jiang X. L., Jiang X., Guan H. S., *J Fish Sci China* 9 (2002) 251-254.
- [28] Yaphe W., Baxter B., *Appl Microbiol* 3 (1955) 380-383
- [29] Sarwar G., Sakata T., Kakimoto D., *Bulletin of the Japanese Society of Scientific Fisheries* 49 (1983) 1689-1694.
- [30] Mao-hong Zhou, Jian-she Ma, Jun Li, Hai-ren Ye, Ke-xin Huang, Xiao-wei Zhao, *Biotechnol Bioproc Engi* 13 (2008) 545-551.
- [31] McLean M. M., Williamson F. B., *Europ J of Biochem.* 93 (1979) 553-558.
- [32] Lakshmikanth M., Manohar S., Patnakar J., Vaishampayan P., Shouche Y., Lalitha J., *World J of Microbiol and Biotechnol.* 22 (2006) 531-537.

- [33] Agbo J., Moss M., *Canadian J. Microbiol.* 115 (1979) 355-368.
- [34] Li Y., Chen J., Lun S. Y., Rui X. S., *Appl Microbiol and Biotechnol.* 55 (2001) 680-685.
- [35] Tarnng Y. S., Juang S. C., Chang C. H., *J of Materials Processing Technology* 128 (2002) 1-6.
- [36] Krishna P. K., Venkata M. S., Sreenivas R. R., Bikas R. P., Sarma P. N., *Biochem Engi J.* 24 (2005) 17-26.
- [37] Sreenivas R. R., Prakasham R. S., Krishna P. K., Rajesham S., Sarma P. N., Venkateswar R. L., *Proc Biochem.* 39 (2004) 951-956.
- [38] Antonopoulos A., Favetta P., Helbert W., Lafosse M., *Anal Chem.* 77 (2005) 4125-4136.
- [39] Aristotelis A., Hardouin J., Favetta P., Helbert W., Lafosse M., *Rapid Communications In Mass Spectrometry* 19 (2005) 2217-2226.

( Received 22 June 2014; accepted 30 June 2014 )