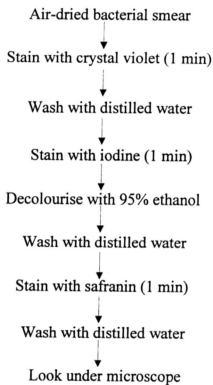


APPENDICES

APPENDIX

APPENDIX A : ANALYTICAL METHODS

1. Gram Staining (Collins and Lyne, 1984)



(Note : Gram + cells are stained purple)

Gram - cells are stained red)

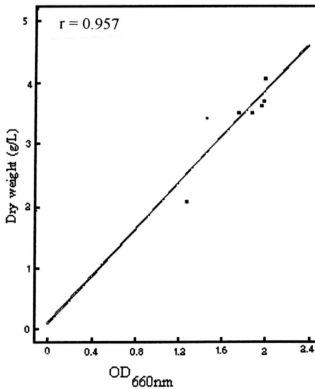
2. Cell Dry Weight

The cell dry weight was determined by filtering 10 mL of thoroughly mixed culture sample through Whatman filter paper No. 4 which was previously dried and weighed. The cell mass in the filter paper was oven-dried at 60 °C overnight, then cooled in a dessicator and finally weighed. Measurements were carried out in duplicates.

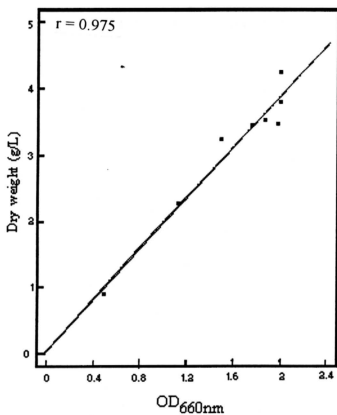
3. Cell Mass Concentration (Noparatnaraporn *et al.*, 1983)

The cell mass concentration was determined by measuring the optical density at 660 nm with a spectrophotometer (Shimadzu UV-160A). The OD_{660nm} values were then converted to cell dry weight by interpolating from the standard curve of cell dry weight (g/L) against OD_{660nm} .

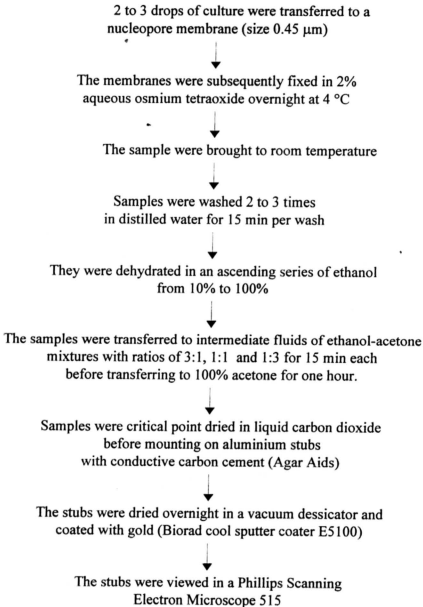
(a) Regression of *R. palustris* strain B1 dry cell weight against OD_{660nm} (Appendix C32)



- (b) Regression of *R. palustris* ATCC 17001 Dry Cell Weight Against OD_{660nm} (Appendix C33)

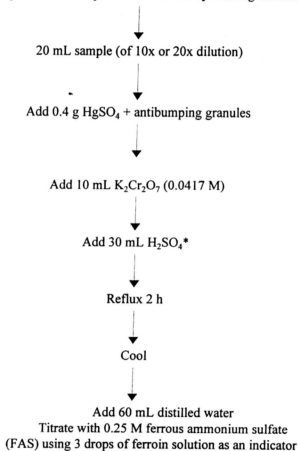


4. Electron Microscopy (Vijaya, N., Personal Communication)



5. Chemical Oxygen Demand (COD) Analysis (APHA, AWWA and WPCF, 1989)

The COD determination is a measure of the oxygen equivalent of that portion of organic matter in a sample that is susceptible to oxidation by a strong chemical oxidant.



(*Add Ag_2SO_4 to concentrated H_2SO_4)

The end-point is a sharp colour change from blue-green to reddish-brown.

The amount of oxidizable organic matter, measured as oxygen equivalent, is proportional to the potassium dichromate used.

The COD of the sample was calculated based on the formula:

$$\text{COD (mg/L)} = \frac{(A - B) \times M \times 8000}{\text{mL sample}}$$

where, A = mL FAS used for the titration of blank

B = mL FAS used for the titration of sample

M = molarity of FAS (Appendix B2)

APPENDIX B : MEDIA, REAGENTS AND BUFFERS

1. Media

a. *Glutamate-malate medium, GMM* (Noparatnaraporn, 1994)

| | | |
|---|---|---------|
| | DL - malic acid | 3.0 g |
| | <i>Sodium glutamate</i> | 2.0 g |
| | $(\text{NH}_4)_2 \text{PHO}_4$ | 1.0 g |
| A | K_2HPO_4 | 0.5 g |
| | $\text{KH}_2 \text{PO}_4$ | 0.5 g |
| | $\text{CaCl}_2 \cdot 2\text{H}_2\text{O}$ | 0.6 g |
| | $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ | 0.2 g |
| | $\text{MnSO}_4 \cdot 5\text{H}_2\text{O}$ (1 mg/mL) | 1.3 mL |
| | $\text{CoCl}_2 \cdot 6\text{H}_2\text{O}$ (1 mg/mL) | 1.0 mL |
| | Ferric citrate (10 mg/mL) in 30% NH_4OH | 0.25 mL |
| | Nicotinic acid (1 mg/mL) | 1.0 mL |
| | Thiamine - HCl (1 mg/mL) | 1.0 mL |
| | Biotin (100 mg/mL) | 1.0 mL |
| | Yeast extract | 1.0 g |

The organic substrates were added before the vitamins. The solution was made up to 1000 mL in distilled water. pH was adjusted to pH 6.8-7.2 by adding 3M NaOH prior to autoclaving. The medium may conveniently be made double-strength and stored at 4 °C with a few drops of toluene before sterilization.

b. GM-salt (1) Medium

| | |
|--|---------|
| DL - malic acid | 3.0 g |
| Sodium chloride | 5.0 g |
| <i>p</i> -Aminobenzoic acid (50 µg/mL) | 1.0 mL |
| Biotin (100µg/mL) | 1.0 mL |
| Distilled water | 1000 mL |

Add ingredients *A* and adjust pH to 7.0 prior to autoclaving.

c. Glutamate Starch (Medium)

| | |
|--|---------|
| Potato starch | 1.0 g |
| Sodium chloride | 5.0 g |
| <i>p</i> -Aminobenzoic acid (50 µg/mL) | 1.0 mL |
| Biotin (100 µg/mL) | 1.0 mL |
| Yeast extract | 1.0 g |
| Distilled water | 1000 mL |

Add ingredients *A* and adjust pH to 7.0 prior to autoclaving.

d. GM-salt (2) Medium

| | |
|---|---------|
| DL - malic acid | 3.0 g |
| Sodium chloride | 5.0 g |
| Yeast extract | 1.0 g |
| MnSO ₄ .5H ₂ O (1 mg/mL) | 1.3 mL |
| CoCl ₂ .6H ₂ O (1 mg/mL) | 1.0 mL |
| Ferric citrate (10 mg/mL) in 30% NH ₄ OH | 0.25 mL |
| <i>p</i> -Aminobenzoic acid (50 µg/mL) | 1.0 mL |
| Nicotinic acid (1 mg/mL) | 1.0 mL |
| Biotin (100 µg/mL) | 1.0 mL |
| Thiamine-HCl (1 mg/mL) | 1.0 mL |
| Distilled water | 1000mL |

Add ingredients *A* and adjust the pH was adjsuted to 7.0 prior to autoclaving.

2. Reagents

a. Gram Strain

Crystal Violet:

A - Crystal violet 2 g/ 20 mL 95% ethanol

B - Ammonium oxalate 0.8 g / 80 mL distilled water

(solution mixed, stood for 24 h and filtered)

Iodine Solution:

Iodine 1 g

Potassium iodide 2 g

Distilled water 300 mL

Safranin :

0.25 g safranin is ground in a mortar with

10 mL 95% ethanol before making it upto

100 mL with distilled water

b. Standard potassium dichromate solution, 0.0417 M

Dissolve 12.259 g $K_2Cr_2O_7$ previously dried at 103 °C for 2 h in distilled water and dilute to 1000 mL.

c. Sulfuric acid reagent ($H_2SO_4^*$)

Add 10.5 g Ag_2SO_4 to 2.5 L concentrated H_2SO_4 . Let stand for 1 to 2 days for dissolution.

d. Ferroin indicator solution

Dissolve 1.485 g 1,10-phenanthroline monohydrate and 695 mg $\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$ in distilled water and dilute to 100 mL.

e. Standard ferrous ammonium sulfate (FAS) titrant, approximately 0.25 M

Dissolve 98 g $\text{Fe}(\text{NH}_4)_2(\text{SO}_4)_2 \cdot 6\text{H}_2\text{O}$ in distilled water. Add 20 mL concentrated H_2SO_4 . Cool and dilute to 1000 mL. Standardize this solution daily against standard $\text{K}_2\text{Cr}_2\text{O}_7$ solution as follows:-

Dilute 10.0 mL standard $\text{K}_2\text{Cr}_2\text{O}_7$ to about 100 mL. Add 30 mL concentrated H_2SO_4^* and cool. Titrate with FAS titrant using 2 to 3 drops ferroin indicator.

Molarity of FAS solution:

$$M = \frac{\text{Volume 0.0417 M } \text{K}_2\text{Cr}_2\text{O}_7 \text{ solution titrated, mL}}{\text{Volume FAS used in titration, mL}} \times 0.25$$

f. Mercury sulphate, HgSO_4 , crystals or powder

3. Buffers

a. Phosphate buffer saline (pH 7.0) (Grist *et al.*, 1979)

| | |
|------------------------------------|---------|
| NaCl | 8.0 g |
| KCl | 0.2 g |
| Na ₂ HPO ₄ - | 1.15 g |
| KH ₂ PO ₄ | 0.2 g |
| Distilled water | 1000 ml |

Adjust pH to 7.0 prior to autoclaving.

APPENDIX C : EXPERIMENTAL AND STATISTICAL DATA

1. OD₆₆₀ nm and cell dry weight of *R. palustris* strain B1 grown in GM medium under anaerobic-light and aerobic-dark conditions

| Time (Day) | Anaerobic-light | | Aerobic-dark | |
|---------------|-----------------------|--------------------|-----------------------|---------------------|
| | OD ₆₆₀ nm* | Cell dry wt* (g/L) | OD ₆₆₀ nm* | Cell dry wt.* (g/L) |
| 0 | 0.496 | 1.03 | 0.602 | 0.56 |
| 1 | 1.277 | 2.05 | 1.626 | 1.37 |
| 2 | 1.884 | 3.46 | 1.518 | 1.21 |
| 3 | 2.002 | 3.66 | 1.459 | 2.13 |
| 4 | 2.009 | 4.02 | 1.229 | 2.40 |
| 5 | 1.979 | 3.59 | 1.362 | 2.23 |
| 6 | 1.775 | 3.48 | 1.359 | 1.33 |
| 7 | 1.482 | 3.40 | 1.302 | 1.23 |

*Mean of three values

2. OD₆₆₀ nm and cell dry weight of *R. palustris* ATCC 17001 grown in GM medium under anaerobic-light and aerobic-dark conditions

| Time (day) | Anaerobic-light | | Aerobic-dark | |
|---------------|----------------------|--------------------|-----------------------|--------------------|
| | CD ₆₆₀ nm | Cell dry wt. (g/L) | CD ₆₆₀ nm* | Cell dry wt. (g/L) |
| 0 | 0.490 | 0.89 | 0.602 | 0.57 |
| 1 | 1.131 | 2.26 | 1.607 | 0.77 |
| 2 | 1.871 | 3.53 | 1.581 | 1.04 |
| 3 | 2.006 | 3.79 | 1.445 | 3.00 |
| 4 | 2.009 | 4.23 | 1.229 | 2.40 |
| 5 | 1.979 | 3.59 | 1.310 | 2.31 |
| 6 | 1.780 | 3.47 | 1.419 | 1.74 |
| 7 | 1.492 | 3.24 | 1.403 | 1.69 |

*Mean of three values

3. Optical density at 660 nm after 72 h at anaerobic-light conditions of *R. palustris* strain B1 and ATCC 17001 grown on different carbon sources

| Carbon source | OD _{660nm} * | |
|-------------------|-----------------------|------------|
| | Strain B1 | ATCC 17001 |
| DL-malic acid | 1.965 | 2.020 |
| Sodium succinate | 1.967 | 1.900 |
| Sodium tartrate | 1.921 | 1.696 |
| Sodium formate | 1.724 | 1.401 |
| Sodium benzoate | 1.694 | 1.732 |
| Sodium sulfite | 1.602 | 1.148 |
| Methanol 1% | 1.598 | 1.541 |
| Pyruvic acid 0.3% | 1.719 | 1.766 |
| Starch | 1.915 | 1.325 |
| Amylopectin | 1.882 | 1.379 |
| Amylose | 0.827 | 0.793 |

*Mean of three values

4. ANOVA (Effect of Carbon Source on Day 3 Growth of Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| Carbon source | 10 | .7387727 | 38.117 | .0000* |
| Residual | 11 | .0193818 | | |
| Total | 21 | | | |

* $p < 0.05$

Multiple Range Analysis Using Tukey HSD at 95% Confidence Intervals (Strain B1)

| Carbon source | Average | Homogenous group* |
|-------------------|-----------|-------------------|
| Succinate | 3.7750000 | a |
| Malic acid | 3.7750000 | a |
| Starch | 3.6750000 | b |
| Tartrate | 3.6750000 | b |
| Amylopectin | 3.6000000 | b |
| Pyruvic acid 0.3% | 3.3500000 | b |
| Formate | 3.3500000 | b |
| Benzoate | 3.3150000 | c |
| Methanol 1% | 3.1250000 | c |
| Sulfite | 3.1250000 | c |
| Amylose | 1.6250000 | d |

*Means followed by a common letter are not significantly different at 5% level.

5. ANOVA (Effect of Carbon Source on Day 3 Growth of ATCC 17001)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| Carbon source | 10 | .9694882 | 18.185 | .0000* |
| Residual | 11 | .0533136 | | |
| Total | 21 | | | |

* $p < 0.05$

Multiple Range analysis Using Tukey HSD at 95% Confidence Intervals (ATCC 17001)

| Carbon source | Average | Homogenous group* |
|-------------------|-----------|-------------------|
| Malic acid | 4.0500000 | a |
| Succinate | 3.8500000 | a |
| Pyruric acid 0.3% | 3.5750000 | b |
| Benzoate | 3.5500000 | b |
| Tartrate | 3.4750000 | b |
| Methanol 1% | 3.1500000 | b |
| Formate | 2.8750000 | c |
| Amylopectin | 2.8400000 | c |
| Starch | 2.7300000 | c |
| Sulfite | 2.3750000 | d |
| Amylose | 1.6750000 | e |

*Means followed by a common letter are not significantly different at 5% level.

6. Optical density at 660 nm after 72 h at $30^{\circ} \pm 2^{\circ}\text{C}$, 3 klux of *R. palustris* strain B1 grown on various types of starch

| No. | Type of starch | OD _{660nm} * |
|-----|--|-----------------------|
| 1 | Potato | 1.888 |
| 2 | Starch (soluble) | 1.825 |
| 3 | Sago | 1.814 |
| 4 | Tapioca | 1.811 |
| 5 | Corn | 1.442 |
| 6 | Wheat | 1.375 |
| 7 | Glutinous rice | 1.315 |
| 8 | Rice | 1.103 |
| 9 | Starch (soluble) without the yeast extract | 0.808 |
| 10 | Raw starch (soluble) ⁺ | 0.311 |

*Mean of three values

⁺Sterilized at 120 °C for 2 h.

7. ANOVA (Effect of various types of starch on Day 3 Growth of Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| Carbon source | 9 | 2.4223569 | 999.999 | .0000* |
| Residual | 12 | .0003444 | | |
| Total | 21 | | | |

* $p < 0.05$

Multiple Range analysis Using Tukey HSD at 95% Confidence Intervals (Strain B1)

| Types of starch | Average | Homogenous group* |
|----------------------------------|-----------|-------------------|
| Potato | 3.6000000 | a |
| Sago | 3.5000000 | a |
| Soluble starch | 3.5000000 | a |
| Tapioca | 3.4500000 | a |
| Corn | 2.8000000 | b |
| Wheat | 2.7000000 | b |
| Glutinous rice | 2.5500000 | b |
| Rice | 2.1500000 | b |
| Soluble starch - YE ⁺ | 1.5666667 | c |
| Raw starch (soluble) | 0.6333333 | d |

*Means followed by a common letter are not significantly different at 5% level.

⁺Yeast extract

8. Optical density at 660 nm after 72 h at $30^{\circ} \pm 2^{\circ}\text{C}$, 3 klux of *R. palustris* strain B1 grown on various concentrations of potato starch

| No. | Potato starch concentration (%) | OD _{660nm} * |
|-----|---------------------------------|-----------------------|
| 1 | 0.3 | 1.952 |
| 2 | 0.5 | 1.979 |
| 3 | 1.0 | 2.342 |
| 4 | 2.0 | 2.356 |
| 5 | 3.0 | 2.369 |
| 6 | 4.0 | 2.371 |
| 7 | 5.0 | 2.362 |

*Mean of three values

9. ANOVA (Effect of potato starch concentrations on Day 3 Growth of Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|-----------------------------|----|-------------|---------|-------------|
| Potato starch concentration | 6 | .2647167 | 999.999 | .0000* |
| Residual | 7 | .0001286 | | |
| Total | 13 | | | |

* $p < 0.05$

Multiple Range analysis Using Tukey HSD at 95% Confidence Intervals (Strain B1)

| Potato starch concentration | Average | Homogenous group* |
|-----------------------------|-----------|-------------------|
| 4.0 | 4.5550000 | a |
| 3.0 | 4.5500000 | a |
| 5.0 | 4.5300000 | a |
| 2.0 | 4.5150000 | a |
| 1.0 | 4.4600000 | b |
| 0.5 | 3.8100000 | c |
| 0.3 | 3.7500000 | d |

*Means followed by a common letter are not significantly different at 5% level.

10. ANOVA (Effect of *p*-aminobenzoic acid (PABA) concentrations on Day 3 Growth of Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| PABA concentration | 4 | .0113400 | .344 | .8381* |
| Residual | 5 | .0329300 | | |
| Total | 9 | | | |

* $p < 0.05$

11. ANOVA (Effect of *p*-aminobenzoic acid (PABA) concentrations on Day 3 Growth of ATCC 17001)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| PABA concentration | 4 | .0041150 | .031 | .9974* |
| Residual | 5 | .1311000 | | |
| Total | 9 | | | |

* $p < 0.05$

12. Effect of temperature on the growth of *R. palustris* strain B1 and ATCC 17001

| Temperature (°C) | OD _{660nm} * | |
|------------------|-----------------------|------------|
| | Strain B1 | ATCC 17001 |
| 25 | 1.892 | 1.819 |
| 30 | 1.831 | 1.775 |
| 35 | 1.739 | 1.745 |
| 37 | 1.722 | 2.009 |
| 40 | 1.715 | 1.484 |
| 45 | 0.513 | 0.305 |
| 50 | - | - |
| 55 | - | - |

*Mean of three values

13. ANOVA (Effect of Temperature on Day 3 Growth of Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| Temperature | 7 | 5.3554286 | 181.233 | .0000* |
| Residual | 8 | .0295500 | | |
| Total | 15 | | | |

* $p < 0.05$ **Multiple Range analysis Using Tukey HSD at 95% Confidence Intervals
(Strain B1)**

| Temperature (°C) | Average | Homogenous group* |
|------------------|------------|-------------------|
| 25 | 3.6350000 | a |
| 30 | 3.5050000 | a |
| 35 | 3.37500000 | a |
| 37 | 3.3350000 | a |
| 40 | 3.3100000 | a |
| 45 | 1.0000000 | b |
| 55 | .0000000 | c |
| 50 | .0000000 | c |

*Means followed by a common letter are not significantly different at 5% level.

14. ANOVA (Effect of Temperature on Day 3 Growth of ATCC 17001)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| Temperature | 7 | 6.3389107 | 892.804 | .0000* |
| Residual | 8 | .0071000 | | |
| Total | 15 | | | |

* $p < 0.05$ **Multiple Range analysis Using Tukey HSD at 95% Confidence Intervals (ATCC 17001)**

| Temperature (°C) | Average | Homogenous group* |
|------------------|-----------|-------------------|
| 37 | 4.0700000 | a |
| 25 | 3.690000 | b |
| 30 | 3.600000 | b |
| 35 | 3.540000 | b |
| 40 | 3.050000 | c |
| 45 | .6400000 | d |
| 55 | .0000000 | e |
| 50 | .0000000 | e |

*Means followed by a common letter are not significantly different at 5% level.

15. Effect of initial pH on the growth of *R. palustris* strain B1 and ATCC 17001

| pH | OD 660 nm* | |
|-----|------------|------------|
| | Strain B1 | ATCC 17001 |
| 4.0 | 0.765 | 0.758 |
| 4.5 | 1.330 | 1.175 |
| 5.0 | 1.842 | 1.842 |
| 5.5 | 2.357 | 2.416 |
| 6.0 | 2.275 | 2.323 |
| 6.5 | 2.084 | 2.087 |
| 7.0 | 2.080 | 2.087 |
| 7.5 | 2.063 | 2.064 |
| 8.0 | 2.163 | 1.995 |
| 8.5 | 1.902 | 1.902 |

*Mean of three values

16. ANOVA (Effect of pH on Day 3 Growth of Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| pH | 9 | 1.6906689 | 31.454 | .0000* |
| Residual | 10 | .0537500 | | |
| Total | 19 | | | |

* $p < 0.05$ **Multiple Range analysis Using Tukey HSD at 95% Confidence Intervals
(Strain B1)**

| pH | Average | Homogenous group* |
|-----|----------|-------------------|
| 5.5 | 4.510000 | a |
| 6.0 | 4.360000 | a |
| 8.0 | 4.140000 | a |
| 6.5 | 4.010000 | a |
| 7.0 | 4.000000 | a |
| 7.5 | 3.965000 | a |
| 8.5 | 3.660000 | a |
| 5.0 | 3.550000 | b |
| 4.5 | 2.590000 | c |
| 4.0 | 1.495000 | d |

*Means followed by a common letter are not significantly different at 5% level.

17. ANOVA (Effect of pH on Day 3 Growth of ATCC 17001)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| pH | 9 | 2.1213894 | 16.923 | .0001* |
| Residual | 10 | .1253550 | | |
| Total | 19 | | | |

* $p < 0.05$ **Multiple Range analysis Using Tukey HSD at 95% Confidence Intervals (ATCC 17001)**

| pH | Average | Homogenous group* |
|-----|----------|-------------------|
| 5.5 | 4.875000 | a |
| 6.0 | 4.700000 | a |
| 7.0 | 4.220000 | a |
| 6.5 | 4.220000 | a |
| 7.5 | 4.185000 | a |
| 8.0 | 4.035000 | a |
| 8.5 | 3.850000 | a |
| 5.0 | 3.730000 | a |
| 4.5 | 2.400000 | b |
| 4.0 | 1.550000 | c |

*Means followed by a common letter are not significantly different at 5% level.

18. Effect of light intensity on growth of *R. palustris* strain B1 and ATCC 17001

| light intensity (klux) | OD 660 nm* | |
|---------------------------|------------|------------|
| | Strain B1 | ATCC 17001 |
| 1000 | 1.690 | 1.736 |
| 2000 | 1.709 | 1.768 |
| 3000 | 1.769 | 1.855 |
| 4000 | 1.897 | 1.906 |
| 5000 | 1.906 | 1.906 |
| 6000 | 1.906 | 1.906 |

*Mean of three values

19. ANOVA (Effect of light intensity on Day 3 Growth of Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| Light intensity | 5 | .0682133 | 1.165 | .4217* |
| Residual | 6 | .0585333 | | |
| Total | 11 | | | |

* $p < 0.05$

20. ANOVA (Effect of light intensity on Day 3 Growth of ATCC 17001)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| Light intensity | 5 | .0431083 | 22.789 | .0008* |
| Residual | 6 | .0018917 | | |
| Total | 11 | | | |

* $p < 0.05$ **Multiple Range analysis Using Newman-Keuls at 95% Confidence Intervals (ATCC 17001)**

| Light intensity (klux) | Average | Homogenous group* |
|------------------------|-----------|-------------------|
| 6 | 3.8600000 | a |
| 5 | 3.8600000 | a |
| 4 | 3.8600000 | a |
| 3 | 3.7550000 | a |
| 2 | 3.6050000 | b |
| 1 | 3.5250000 | b |

*Means followed by a common letter are not significantly different at 5% level.

21. Effect of Salinity on Growth

| % NaCl | OD 660 nm* | |
|--------|------------|------------|
| | Strain B1 | ATCC 17001 |
| 0.0 | 1.875 | 1.875 |
| 0.5 | 1.951 | 1.935 |
| 1.0 | 1.936 | 1.375 |
| 1.5 | 1.478 | 1.316 |
| 2.0 | 1.302 | 1.145 |
| 2.5 | 1.152 | 0.944 |
| 3.0 | 0.693 | 0.874 |

*Mean of three values

22. ANOVA (Effect of NaCl Concentration on Day 3 Growth of Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| NaCl concentration | 6 | 1.5952833 | 10.459 | .0034* |
| Residual | 7 | .1525286 | | |
| Total | 13 | | | |

* $p < 0.05$

Multiple Range analysis Using Newman-Keuls at 95% Confidence Intervalsn (Strain B1)

| NaCl conc. (%) | Average | Homogenous group* |
|----------------|-----------|-------------------|
| 3.0 | 1.3800000 | c |
| 2.5 | 2.2350000 | b |
| 2.0 | 2.5300000 | a |
| 1.5 | 2.8650000 | a |
| 0.0 | 3.6000000 | a |
| 1.0 | 3.7300000 | a |
| 0.5 | 3.7500000 | a |

*Means followed by a common letter are not significantly different at 5% level.

23. ANOVA (Effect of NaCl Concentration on Day 3 Growth of ATCC 17001)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| NaCl concentration | 6 | 1.4975071 | 3.233 | .0751* |
| Residual | 7 | .4632429 | | |
| Total | 13 | | | |

* $p < 0.05$ (Range analysis using 10%)

(Table 21)

| Inoculum size (%) |
|-------------------|
| 5 |
| 10 |
| 15 |
| 20 |

*Means followed by a capital letter

24. ANOVA (Effect of Inoculum Size on Day 3 Growth rate of Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|--------------|---------|-------------|
| Inoculum size | 3 | 8.05000E-005 | 7.318 | .0422* |
| Residual | 4 | 1.10000E-005 | | |
| Total | 7 | | | |

*p < 0.05

Multiple Range analysis Using Tukey HSD at 95% Confidence Intervals (Strain B1)

| Inoculum size (%) | Average | Homogenous group* |
|-------------------|----------|-------------------|
| 5 | .0560000 | a |
| 10 | .0520000 | a |
| 15 | .0500000 | a |
| 20 | .0410000 | b |

*Means followed by a common letter are not significantly different at 5% level.

25. ANOVA (Effect of Inoculum Size on Day 3 Growth rate of ATCC 17001)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|--------------|---------|-------------|
| Inoculum size | 3 | 2.74583E-005 | 12.922 | .0159* |
| Residual | 4 | 2.12500E-006 | | |
| Total | 7 | | | |

* $p < 0.05$ **Multiple Range analysis Using Tukey HSD at 95% Confidence Intervals (ATCC 17001)**

| Inoculum size (%) | Average | Homogenous group* |
|-------------------|----------|-------------------|
| 10 | .0540000 | a |
| 15 | .0505000 | a |
| 5 | .0500000 | a |
| 20 | .0450000 | b |

*Means followed by a common letter are not significantly different at 5% level.

26. ANOVA (Effect of Inoculum Age on Day 3 Growth rate of Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|--------------|---------|-------------|
| Inoculum age | 5 | 3.73333E-006 | .974 | .5012* |
| Residual | 6 | 3.83333E-006 | | |
| Total | 11 | | | |

*p < 0.05

27. ANOVA (Effect of Inoculum Age on Day 3 Growth rate of ATCC 17001)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|--------------|---------|-------------|
| Inoculum age | 5 | 1.06833E-005 | 2.981 | .1080* |
| Residual | 6 | 3.58333E-006 | | |
| Total | 11 | | | |

*p < 0.05

28. COD Removal of Settled and Unsettled Sago Effluent by Free and Immobilized Cells of *R. palustris* Strain B1

| | COD removal (%)* | | | |
|-------|------------------|----------------------|--------------------|----------------------|
| | Settled effluent | | Unsettled effluent | |
| | Free cells | Agar-entrapped cells | Free cells | Agar-entrapped cells |
| Day 1 | 34.8 | 16.6 | 73.6 | 70.8 |
| Day 2 | 58.3 | 27.0 | 75.7 | 68.2 |
| Day 3 | 60.4 | 39.5 | 86.4 | 79.5 |
| Day 4 | 64.2 | 44.3 | 85.1 | 81.8 |
| Day 5 | 70.8 | 55.2 | 74.4 | 79.0 |

*Mean of three values

29. ANOVA (Effect of Sago Effluent Concentration on COD Removal by Immobilized Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| Sago effluent conc. | 3 | 174.42917 | 10.157 | .0242* |
| Residual | 4 | 17.17252 | | |
| Total | 7 | | | |

*p < 0.05

Multiple Range analysis Using Tukey HSD at 95% Confidence Intervals

| Sago effluent conc. (%) | Average | Homogenous group* |
|-------------------------|-----------|-------------------|
| 100 | 57.590000 | a |
| 75 | 49.615000 | a |
| 50 | 39.665000 | b |
| 25 | 37.390000 | b |

*Means followed by a common letter are not significantly different at 5% level.

30. ANOVA (Effect of Mixing on the COD Removal of Sago Effluent by Immobilized Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| Effect of mixing | 3 | 128.04661 | 3.701 | .1193* |
| Residual | 4 | 34.59774 | | |
| Total | 7 | | | |

*p < 0.05

31. ANOVA (Effect of Inoculum Size on the COD Removal of Sago Effluent by Immobilized Strain B1)

| Sources of variation | df | Mean square | F-ratio | Sign. level |
|----------------------|----|-------------|---------|-------------|
| Inoculum size | 2 | .792067 | .079 | .9259* |
| Residual | 3 | 10.022800 | | |
| Total | 5 | | | |

*p < 0.05

32. Regression of *R. palustris* Strain B1 Dry Cell Weight Against OD_{660nm}

$$r = 0.957$$

| OD at 660 nm | Mean dry weight (g/L) |
|--------------|-----------------------|
| 0.496 | 1.03 |
| 1.277 | 2.05 |
| 1.482 | 3.40 |
| 1.775 | 3.48 |
| 1.884 | 3.46 |
| 1.979 | 3.59 |
| 2.002 | 3.66 |
| 2.009 | 4.02 |

33. Regression of *R. palustris* ATCC 17001 Dry Cell Weight AgainstOD_{660nm}

$$r = 0.975$$

| OD at 660 nm | Mean dry weight (g/L) |
|--------------|-----------------------|
| 0.490 | 0.89 |
| 1.131 | 2.26 |
| 1.492 | 3.24 |
| 1.780 | 3.47 |
| 1.871 | 3.53 |
| 1.979 | 3.47 |
| 2.006 | 3.79 |
| 2.009 | 4.23 |