Biodegradation of Starch Blended High Density Polyethylene using Marine Bacteria Associated with Biofilm Formation and its Isolation Characterization

Thangavelu Muthukumar1,3, Adithan Aravinthana1, R Dineshram2, Ramasamy Venkatesan1 and Mukesh Doble1**

1Bioengineering and Drug Design Lab, Department of Biotechnology, Indian Institute of Technology Madras, Chennai, India
2OSTI, National Institute of Ocean Technology, Chennai, India
3Bio-products Laboratory, Central Leather Research Institute, Adyar, Chennai, India

Abstract

The biofouling and biodegradation of starch blended high density polyethylene (HDPE) was studied by immersing it in Bay of Bengal (India) for a period of six months. A positive correlation was observed amongst the various constituents in the biofilm. FTIR spectrum showed the formation of C=O stretching band and decrease in ester and keto carbonyl bands indicating biodegradation. 17% weight loss was observed, while the polymer surface turned hydrophilic. Twenty two bacterial strains were isolated from the biofilm and biochemically characterized. 16sRNA sequence analysis was performed for three strains. In vitro biodegradation of the polymer exposed to sunlight for 150 days incubated with the isolated pure strain (Exiguobacterium) and combination of two strains (Exiguobacterium and B. subtilis) for 75 days, showed a gravimetric weight loss of 4.7 and 12.1% respectively indicating synergy between the two organisms. The current study indicated, the isolated microorganisms could degrade starch blended HDPE.

Keywords: Biodegradation; Biofilm, Starch blended HDPE, Bacillus subtilis; Exiguobacterium sp., ATP

Introduction

High density polyethylene (HDPE), particularly as a thin film, is widely used in food, clothing, shelter, transportation, construction, medical, and recreation industries. Its disposal after use causes several environmental issues because of its recalcitrant nature. The rate of biodegradation of this polymer dumped in the open environment is very slow due to the strong intermolecular bond that does not allow the microbes to access them easily. This polymer accumulates at a rate of 25 million ton per year in the terrestrial and in the marine environment [1]. Also the dumping of the waste material in the sea is the causative agent of major marine pollution. It has been estimated that more than a million of marine animals are being killed every year either by choking on floating plastic items or by becoming entangled in plastic debris. Fishing and shipping industries are a major cause of plastic litter. The two major problems which hinder the biodegradation of polypropylene are its high hydrophobicity and molecular weight. The only way to solve this problem is to develop biodegradable polymers [2]. Many conventional techniques are adopted to dispose plastics which include incineration, pyrolysis, landfill, and recycling which involves many technical and economical problems. Synthetic plastics mixed with natural polymers including starch or alginate degrades when dumped in open environment. This degradation leads to changes in their physical and mechanical properties. This degradation can be enhanced by mild physico-chemical pretreatment including thermal or UV treatment [3]. The biodegradability of high density polyethylene has been enhanced by introducing additives to it, including starch and pro-oxidants [4,5].

Ocean contains a large number of micro and macro-organisms, which could involve in the degradation of the polymer. Attachment of microorganisms to form a biofilm on the surface of the polymer is the first step in the process of biodegradation. It begins with the reversible attachment followed by an irreversible adhesion of microorganisms to the substratum [6]. The initially attached cells can be easily removed by a gentle wash since it is held by physical forces whereas the later attachment is mediated by hydrogen bonding, cation bridging, and receptor ligand interaction [7]. During the biofilm formation on the surface bacteria overproduces extracellular polymeric substances (EPS) [8]. EPS build up the biofilm matrix, serving as a multipurpose functional element for adhesion, protection, immobilization of cells and facilitating spatial arrangement of different species [9]. Rhodococcus ruber and B. borstelensis [10], have been reported to colonise polyethylene by forming biofilm and later degrade it. Only 0.75% weight loss was observed when pure HDPE was immersed in sea water for a period of six months [11].

The present study aimed to understand the biofouling and biodegradation of starch blended high density polyethylene (HDPE) immersed in sea water and isolating microbial species that can able to degrade starch blended HDPE in minimum time.

Materials and Methods

Experimental materials

Commercially available 5% (w/w) starch blended HDPE (Bio Bags, Chennai, India) film of size 10 × 7.5 cm and 0.23 mm thickness was used in this present study. The samples were washed with 70% ethanol, followed by distilled water and dried in a hot air oven for 5 h at 40°C. Later the samples were brought to room temperature and kept for...
24 h before deployment in the ocean. Firefly extract was used for the estimation of adenosine triphosphate (ATP) (Sigma Aldrich, India). All the other chemicals were purchased from Himedia, India.

**Deployment of the polymer**

The films were immersed in the surface waters of Bay of Bengal at Ennore Port Limited (EPL) Chennai, India (Latitude 13°15’ 48” N, Longitude 80° 20’ 28” E) at a depth of 1 m. The coupons were tied vertically to a floating raft made of high density polyethylene. This study was conducted for six months. The polymer films in triplicate were retrieved after every two months interval up to 6 months.

**Seawater parameters**

During the sampling period, temperature, salinity and dissolved oxygen of the sea water were measured using a pre-calibrated water quality monitoring system (Hydrolab, Quanta Instruments, USA).

**Biological parameters**

The films were retrieved from the sampling location and immersed in a bottle containing 500 ml of filtered (0.22 µm, Millipore) and sterilized seawater. The attached biofilm (attached biomass) on the polymer film was scraped using a sterile nylon brush into the container containing filtered sterilized water [12]. The sample in the container was mixed thoroughly. Then this 1 ml of sample was transferred to an Eppendorf tube and was serially diluted prior to further experiments. It was also plated on Zobell Marine Agar as well as on Pseudomonas agar (Himedia, India). After 24 h of incubation at 28 ± 2°C [13], Total viable count (TVC) were counted manually.

The total suspended solids (TSS), total carbohydrate, total protein, chlorophyll a, adenosine triphosphate (ATP), Pseudomonas and total viable count in the biofilm formed on the polymer were measured once every two months. The above parameters play a vital role in determining the extent of biofouling on a substrate. The total suspended solids in the biofilm were estimated based on the method reported by Parson et al [14]. A sample of 10 ml was filtered through a pre-weighed (Sartorius CP64, 0.1 mg accuracy) GF/C filter (47 mm, 1.25 µm, Whatman), dried at 100°C for 1 h and reweighed to get the dry weight of the residual biomass. The total carbohydrate and protein content in the biofilm were estimated using phenol - sulphuric acid method [15,16], and Lowry’s method [17] respectively.

Chlorophyll a content (chl a) indicates the amount of algal biomass present on the polymer coupons. The filtrate with the filter paper was placed in a screw capped centrifuge tube containing 10 ml of 90% acetone and was stored at -20°C until further analysis [18]. The chlorophyll a pigment was measured with a spectrophotometer (Perkin Elmer, Lambda 35, USA). For estimation of ATP, the filter paper was kept in 3 ml of 0.02 M Tris (pH 7.5) buffer and boiled for 10 min and stored at -20°C until the analysis was carried out using the method reported by Hamilton et al [19]. All the measurements were performed in triplicates. Correlation coefficients between various parameters were determined using SPSS software Version 15.0.

**Isolation and identification of bacteria from biofilm**

Twenty two bacterial colonies were isolated and they were named serially from NRBMD 1 to NRBMD 22 and their morphological, biochemical and molecular characteristics were determined. The bacteria with distinct morphology and which showed positive results on the starch utilization test were purified and sub cultured using respective agar medium. From this group three bacterial isolates were selected and were maintained at 4°C for further characterization studies. The morphologies of these organisms were also observed by growing them for 24 h at room temperature in nutrient agar medium.

These three strains were identified based on their 16S rDNA sequence. The genomic DNA from these bacteria was extracted using phenol chloroform method after incubating them in Zobell Marine Broth for 24 h [20]. PCR amplification of almost the full length of the 16S rDNA was performed with the template DNA using the universal forward and reverse primers 5’GAGTTTGATCCTGGCTCA3’ and 5’ACGGCTAACTTGTTACGACT3’ respectively [21]. The conditions used were as follows, The initial incubation was performed at 95°C for 5 min, later denaturation at 94°C for 1 min, annealing at 55°C for 1 min, extension at 72°C for 2 min. The same conditions were repeated for 30 cycles and finally a 7 min extension at 72°C [21]. The PCR product was confirmed by 1% agarose gel and it was sent for sequencing to an external agency (Anshul Biotechnologies, Hyderabad, India).

**EPS producing ability**

Bacterial strains were grown in Zobell Marine Agar (Himedia, India) and then they were transferred to test tubes containing Zobell Marine Broth and incubated for 24 h at 28 ± 2°C. A day prior to the experiments, 200 µl of the inoculum was transferred from the broth aseptically to a 250 ml conical flask containing 50 ml of modified basal salt solution [22]. The culture broth was inoculated and incubated at 28 ± 2°C for 5 d at 180 rpm. The bacterial cells were removed from the culture broth by centrifugation at 10,000g for 20 min [7]. The supernatant was mixed with 3 volumes of ethanol and kept at 4°C for 24 h. Then it was centrifuged (10,000g, 4°C, 20 min), the EPS was precipitated and maintained at 80°C for 3 days. The amount of EPS produced by each strain was measured using the method suggested by Kwon et al [23]. The procedure makes use of phenol sulphuric acid [15,24]. Briefly, 2 ml of the culture broth was centrifuged and filtered through 0.22 µm (Millipore) filter, transferred to a dialysis bag and was dialyzed against distilled water for 48 h at 4°C. After the dialysis, the samples were made up to a volume of 3 ml. From this, 500 µl of the content was taken for the estimation of EPS, using glucose as the standard and the absorbance was read at 490 nm.

**Bacterial adhesion to hydrocarbon (BATH)**

BATH assay was performed to determine the hydrophobicity of the surface of the isolated bacterial cultures [24]. All the organisms were inoculated in a modified basal salt solution and incubated at 26 ± 2°C for 18 h to get mid log phase culture. They were centrifuged and washed with phosphate buffered saline twice and the optical density of the cells was adjusted to 0.2 at 540 nm with the same buffer. To 3 ml of bacterial cell suspension, 100 µl of n-hexadecane (Himedia, India) was added. Then the suspension was vortexed for 1 min and the two phases were allowed to separate for 15 min. The absorbance of the aqueous phase was measured at 400 nm using a spectrophotometer (Perkin Elmer, Lambda 35, USA). Absorbance of the bacterial suspension without the addition of hexadecane was used as the control. The percentage of the cells bound to n-hexadecane was calculated using the following formula:

\[
\% \text{ of bacterial cell adhered to hexadecane} = \frac{(1-A)}{A_0} \times 100
\]

Where \(A_0\) and \(A\) are the absorbance at 400nm of the bacterial suspension without and with n-hexadecane respectively.
**Physicochemical properties of the polymer**

Hydrophobicity of the polymer surface is determined by measuring its contact angle using sessile drop method with easy drop contact angle measuring Goniometer (Kruess, Germany). Millipore grade distilled water is used as the wetting liquid. An average of five measurements is reported here [25,26]. Contact angle is directly proportional to the hydrophobicity of the surface.

Gravimetric weight loss is a simple and quick way to determine the degradation of the polymer. The macrofoulants from the polymer surface are removed by immersing the scraped films in 10% hydrochloric acid for few minutes, then washing them with distilled water and later drying them in a hot air oven at 40°C for 4 h [27]. Samples are weighed with an accurate four digit balance (Sartorius CP64, 0.1 mg accuracy).

Scanning Electron Microscopy (SEM) and Energy Dispersive Analysis of X-rays (EDAX) of the HDPE samples retrieved from the site were performed to detect the changes on the surface morphology and elemental composition due to degradation respectively with a JEOL JSM-6500F, (Tokyo, Japan). The latter was performed at an acceleration voltage of 10 kV resulting in a lateral and depth compositional resolution of approximately 5 µm.

Chemical changes which might have occurred in the polymer due to deployment were measured using Fourier transform infrared (FTIR) Jasco N4200 (Japan) Spectrometer. The spectra were recorded as an average of 32 scans at a resolution of 4 cm⁻¹ in the frequency range of 4000-500 cm⁻¹. The C-O peak, CH₂ deformation, CH₂ bending (symmetrical), CH₂ bending (asymmetrical) and CH stretching were compared with the control sample as a reference. The control samples were those which were kept in the laboratory at room temperature for the same study period. The keto carbonyl, ester carbonyl, and vinyl indices for the polymer were calculated using the following formula with respect to the intensity of methylene group.

\[
\text{Keto carbonyl bond index } = \frac{I_{1715}}{I_{1465}}
\]

\[
\text{Ester carbonyl bond index } = \frac{I_{1740}}{I_{1465}}
\]

\[
\text{Vinyl bond index } = \frac{I_{1640}}{I_{1465}}
\]

Where relative intensities at 1740, 1715, 1640 and 1465 cm⁻¹ correspond to ester and keto carbonyls, double bond (vinyl index) and methylene band respectively [28].

Thermograms of the polymer samples were recorded using a Differential Scanning Calorimeter (DSC) and Thermo Gravimetric/ Differential Thermal Analyzer (TG/DTA) (Model 204, Netzsch, Germany). Scans were performed under nitrogen atmosphere in the temperature range of -50 to 450°C and -28 to 600°C respectively, at a heating rate of 10°C/min. The melting temperature (Tm) and percentage crystallinity were calculated from the thermograms [29]. The enthalpy of 100% crystalline HDPE (ΔH°m) is 288 J/g and is taken as a reference [30].

\[
\% \text{ Crystallinity } = 100 \times \frac{\Delta H_m}{\Delta H^\circ \text{m}}
\]

**In vitro biodegradation studies**

The samples were exposed to sunlight for 150 days and their biodegradation were monitored in vitro for 75 days with the three bacteria isolated from the biofilm obtained during the first part of our study formed on the polymers when they were deployed in sea. Samples unexposed to sunlight were also subjected to similar biodegradation with the bacterial isolates. Pretreated films (80 mm × 25 mm) were taken in conical flask and 100 ml aliquots of filtered (0.22 µm Millipore filter paper) and autoclaved seawater was added to each of them. Log phase seed culture (1% of overnight grown culture containing 107 cells) was added to these flask and they were kept under shaking at 180 rpm at 35-37°C (Scigenics Biotech, Chennai, India). Sea water containing polymer without the microorganism was maintained as positive control and organism without the polymer served as a negative control [2,31].

**Viability of the bacterial biofilm**

Viability of the bacterial biofilm (live and dead cells) present on the polymer sample was determined by using Live/Dead BacLight™ Bacterial viability Kit (Invitrogen, Germany), according to the manufacturer’s Instructions. The kit has two dyes SYTO9 (green colour) stains the live cells and propidium iodide (red colour) stains only the dead cells. The polymers were removed at the end of in vitro study from the conical flask and were stained with Bac light, incubated for 10-15 min in the dark and then the images were captured with a blue filter at an excitation of 475 nm under a fluorescence microscope (Leica DM5000, Germany).

**Statistical analysis**

Results are reported as mean ± S.D (n = 3). Pearson correlation coefficient between various parameters was estimated using SPSS software Version 15.0. The observed differences were considered statistically significant when p < 0.05.

**Results and Discussions**

**Biological characterization of biofilms and total suspended solids (TSS)**

The average temperature, pH, salinity and dissolved oxygen over the six month study period of the sea water at Ennore port were 28 ± 1°C, 8.18 ± 0.06, 33.67 ± 2 (ppt) and 4.57 ± 0.31 (mg/l) respectively. The extent of biofouling on the starch blended HDPE polymer deployed for a period of six months is seen in photographs (Figure 1a-1c). The images were taken as soon as the films were retrieved from the ocean.

Season influences the accumulation of total suspended solids and macrofoulants on the polymer surface (Figure 2a and 2b). Maximum fouling load is observed during the month of September, which is manifested in large values in all the measured biomass parameters (TSS, protein, carbohydrate, chlorophyll a, and ATP). It reduced in the
There is a strong positive correlation between the various biomass and biofouling parameters (Table 1). ATP is a measure of the amount of viable microorganisms present on the polymer. It shows a strong positive correlation (p<0.01) with pseudomonas count (r = 0.89), protein (r = 0.88) and carbohydrate (r = 0.95). However this is to be expected since all these parameters are the direct measure of the biofouling, hence a positive correlation is observed amongst each other.

### Identification of biofilm forming bacteria

In the present study, the biofilm samples were collected from the HDPE starch blended polymer and were plated on pseudomonas and Zobell marine agar media. The morphological, biochemical and molecular characteristics of a total of three abundantly found isolates were investigated. The staining report showed that all the three isolated strains were Gram positive, and found to hydrolyze starch (starch utilization test) as well as catalase positive. Whereas they showed negative results for indole, methyl red, and citrate utilization tests.

The three abundantly found isolates had different morphology, size and growth pattern. The nucleotide sequences obtained for the three bacteria were compared with NCBI (http://www.ncbi.nlm.nih.gov/) database. The 16s rDNA sequences of these microorganisms (NRBMD 14, NRBMD 16, and NRBMD 21) were determined and were deposited in GenBank, under the accession numbers, GQ357643, GQ357644 and GQ357645 respectively, NRBMD14 aggregates and forms microcolonies on the ZMA plate. NRBMD16 grows as colonies on the Nutrient Agar and appears orange in color. NRBMD21 spreads on the surface of the ZMA plate. Phylogenetic trees were constructed for all these three isolated microorganisms using the Neighbor-joining tree method using CLUSTAL W (http://align.genome.jp/) and compared with closely related species from the NCBI database (Figure 3a and 3b). Their evolutionary distances are also presented in the figures. Strains NRBMD14 and NRBMD21, are closely related to *Bacillus subtilis*. NRBMD16 is closely related to *Exiguobacterium*.

NRBMD14 shares 98% identity with other *Bacillus subtilis* available in the NCBI database, whereas NRBMD21 shares 99% identity with *Bacillus subtilis strain ZB5*. NRBMD 16 shares 87% identity with other *Exiguobacterium* species for query coverage of 97%. It is known that *Bacillus* spp adapts to diverse habitats which vary widely in space and environmental conditions. *Exiguobacterium* is also reported to adapt to a wide range of environments with temperature ranging from -12 to 20°C. This is also reported to adapt to a wide range of environments with temperature ranging from -12 to 20°C.

### Table 1: Correlation coefficient between various microfouling products on HDPE immersed for 6 month at Ennore port limited (TSS-Total Suspended Solids, Chl a-Chlorophyll a, ATP-AdenosineTriphosphate and PS-Pseudomonas).

<table>
<thead>
<tr>
<th>Parameters</th>
<th>TSS</th>
<th>Protein</th>
<th>Carbohydrate</th>
<th>Chl a</th>
<th>ATP</th>
<th>PS</th>
</tr>
</thead>
<tbody>
<tr>
<td>TSS</td>
<td>1</td>
<td>0.77*</td>
<td>0.65*</td>
<td>0.97*</td>
<td>0.76*</td>
<td>0.37</td>
</tr>
<tr>
<td>Protein</td>
<td>1</td>
<td>0.99*</td>
<td>0.91*</td>
<td>0.99*</td>
<td>0.88*</td>
<td>-</td>
</tr>
<tr>
<td>Carbohydrate</td>
<td>1</td>
<td>0.82*</td>
<td>0.99*</td>
<td>0.95*</td>
<td>1</td>
<td>0.89*</td>
</tr>
<tr>
<td>Chl a</td>
<td>1</td>
<td>0.97*</td>
<td>1</td>
<td>0.99*</td>
<td>0.95*</td>
<td>1</td>
</tr>
<tr>
<td>ATP</td>
<td>1</td>
<td>0.89*</td>
<td>1</td>
<td>0.99*</td>
<td>0.95*</td>
<td>1</td>
</tr>
<tr>
<td>PS</td>
<td>1</td>
<td>0.89*</td>
<td>1</td>
<td>0.99*</td>
<td>0.95*</td>
<td>1</td>
</tr>
</tbody>
</table>

*Correlation is significant at p≤ 0.01 level (2-tailed).

There is a strong positive correlation between the various biomass and biofouling parameters (Table 1). ATP is a measure of the amount of viable microfouling present on the polymer. It shows a strong positive correlation (p<0.01) with pseudomonas count (r = 0.89), protein (r = 0.88) and carbohydrate (r = 0.95). However this is to be expected since all these parameters are the direct measure of the biofouling, hence a positive correlation is observed amongst each other.
A gravimetric weight loss of 17% was observed, so the unaccounted 8.5% commercial HDPE constitute 9.5% by weight. At the end of 6 months, weight loss was observed in starch blended HDPE after six months (Table 2). The additives, fillers, softeners, antioxidants and starch in the polymer lost its weight. A strong positive correlation (r = 0.97; p<0.01) was observed between weight loss and contact angle (r = -0.99; p<0.05), indicating that the surface had turned hydrophilic as the polymer lost its weight. A strong positive correlation (r = 0.97; p<0.01) was also observed between TSS and weight loss.

**FTIR and DSC**

The FTIR spectra (Figure 5) of the starch blended HDPE deployed for six months shows absorption frequencies corresponding to CH stretching at 3369 cm\(^{-1}\), CH\(_2\) deformation at 1640 cm\(^{-1}\), CH\(_2\) bending (asymmetrical) at 1466 cm\(^{-1}\) and CH\(_2\) bending (symmetrical) at 1352 cm\(^{-1}\). The weight loss can be attributed to the degradation of HDPE. Studies from our lab have reported 2.5% weight loss in unblended HDPE deployed in marine water for twelve months [27]. In vitro degradation of starch blended LDPE with a marine organism, *B. ceres*, showed 25% weight loss in a period of 12 months [26]. In this study a strong negative correlation was observed between weight loss and contact angle (r = -0.99; p<0.05), indicating that the surface had turned hydrophilic as the polymer lost its weight. A strong positive correlation (r = 0.97; p<0.01) was also observed between TSS and weight loss.

**Polymer surface analysis**

The initial contact angle of HDPE starch blended polymer before deploying it in the ocean was 83 ± 0.76° and it decreased to 70 ± 2.02° at the end of six months (Table 2). This indicated that the polymer surface became relatively hydrophilic with increasing immersion period, which was also reported earlier by several authors [13,26].

SEM focuses on the surface topology of the HDPE film. After 6 months in marine environment surface irregularities (Figure 4b), cracks and cavities are clearly seen when compared to the control (Figure 4a). SEM images clearly show that the surface of the polymer has been affected due to micro and macro foulants under the marine environment, leading to degradation and deterioration. The non-biological oxidation of polyethylene is a well-known process involving radicals attacking the polymer chain, resulting in the formation of simple compounds like water, acetone, and acetic and formic acids. Elemental composition as measured by EDAX indicates that percentage of carbon has reduced from 90 to 87.4 suggesting the microbes present in the environment have utilized the polymer as one of their carbon source [35]. The increasing percentage of oxygen from 1.45 to 4.17 indicates that the polymer has undergone oxidation. The starch in the polymer could have been consumed by the microorganisms or deteriorated and could have released degraded products leading to loss of its integrity [36]. The presence of elemental silicon in EDAX of the six months old sample must be due to the biofouling by the diatoms and ocean soil since it is not present in the original polymer sample. Calcium was also detected in both the control and 6 months immersed polymer sample which might have been incorporated as an inorganic additive to improve its mechanical properties and reduces its brittleness while casting.

**Weight loss**

Biofilm (including solid biomass and microorganism) and macrofoulants on the polymer surface not only change its surface property but also lead to its degradation. Formation of the biofilm on the surface of the polymer leads to loss of its integrity [4,10]. Biodegradation is usually initiated at the surface of the polymer by the exo enzymes produced by the bacteria and hence weight loss is proportional to the surface area. Primarily, the change in the polymer morphology is due to photodegradation and surface oxidation. This is followed by biodegradation influenced by microbes and macrofoulants. In sea water polymer degradation is influenced by several factors. 17% Weight loss was observed in starch blended HDPE after six months (Table 2). The additives, fillers, softeners, antioxidants and starch in the commercial HDPE constitute 9.5% by weight. At the end of 6 months a gravimetric weight loss of 17% was observed, so the unaccounted 8.5%
The formation of a new C-O stretching frequency at 1078 cm\(^{-1}\) and a broad absorption peak at 3400 cm\(^{-1}\) assigned to stretching vibration of –OH are not seen in the control. The former band indicates incorporation of oxygen in the polymer chain. The decrease in Keto carbonyl index from 0.156 ± 0.003 to 0.148 ± 0.001, (p<0.05) ester index from 0.146 ± 0.001 to 0.128 ± 0.002 (p<0.05) and vinyl index from 0.241 ± 0.001 to 0.211 ± 0.002 (p<0.05) of the 6 months old sample when compared to the original sample indicates biotic degradation. It has been reported that the decrease in the carbonyl indices with polymer which were deployed for a long period in the ocean water can be due to the formation of esters by Norrish type mechanism [11,28]. DSC of the control and sample retrieved from ocean after 6 months show melting peaks at 130.1 and 129.3°C respectively. The corresponding heats of enthalpies are 176 and 196 J/g respectively. So it could be concluded that the % crystallinity increased from 61.11 to 68.05. This increase in crystallinity is possibly due to the removal of starch from the polymer or the degradation of the amorphous region in the polymer. Similar increase in crystallinity has been reported earlier [4], during \textit{in vitro} degradation of polypropylene.

\textbf{In \textit{in vitro} studies}

NRBMD 14, NRBMD 16 and NRBMD 21 produced 3.23 ± 0.12, 3.04 ± 0.11 and 2.75 ± 0.13 mg/l of EPS (calculated by measuring dried EPS weight/dried cell weight) respectively indicating the amount and quality of the biofilm formed on the polymer surfaces. EPS producing ability of the bacteria is measured in terms of water soluble fraction. The amounts of EPS produced by all the three strains were found to be higher when compared with other reported strains [23]. Even though both NRBMD14 (GQ357643) and NRBMD21 (GQ357645) are the same species (\textit{Bacillus subtilis}) they showed considerable differences in their EPS producing ability. The type of EPS produced is reported to be specific to certain genus or species [37]. In our study we found that higher the cell biomass higher is the EPS amount indicating that, the cell growth and the EPS produced are positively correlated with each other (p<0.01).

BATH test indicated that the percentages of bacteria in the n-hexadecane phase were 3.45 (NRBMD14), 11.55 (NRBMD16) and 6.22 (NRBMD21), indicating that NRBMD16 was the most hydrophobic and NRBMD14 was the least hydrophobic amongst these three isolates. All the three strains are found to be highly hydrophilic with less than 12% absorbance in the hexadecane phase, which was isolated from starch blended HDPE, implying that hydrophobic substrate favored the attachment of hydrophobic bacteria [13]. The viable bacteria on the polymer surface after 75 days of \textit{in vitro} biodegradation experiments were characterized and identified. In both the experiments more living bacterial cells (green colour) were observed on the surface of starch blended HDPE incubated with \textit{Exiguobacterium} and mixture of \textit{Exiguobacterium} and \textit{Bacillus subtilis} (Figure S2).

Table 3 summarizes the effect of incubation of starch blended HDPE with \textit{Exiguobacterium} and mixture of \textit{Exiguobacterium} and \textit{Bacillus subtilis} for 75 days after being exposed to sunlight for 150 days. Exposure to sunlight, leads to the formation of radicals due to the bombardment of UV radiation. Higher biodegradation is observed with UV exposed polymer than unexposed. Higher biodegradation is observed with the mixed culture (12.1%) than with the individual culture (4.75%), indicating possible synergy between the two organisms. The contact angle of the incubated samples decreased considerably indicating that the surface had turned relatively hydrophilic. Three bacterial strains isolated from the biofilms attached to the polymer initiated degradation. \textit{In vitro} studies with multiple strains is found to biodegrade the UV exposed polymer, much more than the degradation observed with the individual organism, indicating synergy.

\textbf{Conclusions}

Starch blended HDPE exposed to sea water for six months experienced biodegradation, significantly altering its physical and chemical properties. Starch blended polymer though cannot be a solution to tackle pollution, will invariably minimize the amount of non-degradable HDPE in the environment. \textit{In vitro} studies using isolated multiple strains indicated that they could biodegrade UV exposed polymer if the right microorganism is isolated. The isolated organism are able to form a biofilm on the HDPE surface which were also found to grow on the minimal medium containing only starch blended HDPE even without nitrogen source. These organisms could be further exploited through molecular biology techniques to enhance biodegradation of such recalcitrant polymers. Hence, further studies are necessary to adequately assess the environmental decomposition of biodegradable polymers, particularly in sea water. This study is the first step in understanding the biofilm diversity and their usefulness in handling recalcitrant marine pollutants.

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