

FTIR Analysis of Biodegradation of Polystyrene by Intestinal Bacteria Isolated from *Zophobas Morio* and *Tenebrio Molitor*

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Received 02 April 2020; received in revised form 03 September 2020; accepted 06 November 2020

DOI: <https://doi.org/10.46604/peti.2021.5450>

Abstract

Since the 1970s, polystyrene (PS) plastic marine pollution has become a global concern. *Tenebrio molitor* and *Zophobas morio* beetle larva actively respond to a diet of primarily polystyrene. The tantalizing evidence indicates that the gut bacteria of these beetle larva enables them not only to consume polystyrene, but to successfully biodegrade polystyrene. Heretofore, data collection to verify polystyrene degradation by the gut bacteria of these larva has taken up to six months per test. Our laboratory created a platform by dissolving PS into a liquid mineral culture medium to examine PS degradation by the gut bacteria. Under investigation, PS works as main carbon source supporting the growth of gut bacteria. Fourier-transform Infrared spectroscopy (FTIR) is an effective method which can provide relevant information on chemical changes. This study presents a methodology using FTIR and Visible Light Spectrometry as an efficient methodology to verify the physiological degradation of polystyrene.

Keywords: polystyrene, *tenebrio molitor*, *zophobas morio*, FTIR

1. Introduction

Polystyrene (PS) marine pollution had intensively increased since the 1970s [1]. Plastic contributes to 60-90% of marine debris and 50% of debris belongs to the single-use plastics (SUPs), including polystyrene [2]. Recent publications suggest that weathering processes promote further fragmentation of PS into nano-plastics, and show a metabolic toxicity which can change the behavior of marine creatures, or damage human health [3-4].

Previous research successfully isolated two bacterial strains from the intestines of *Tenebrio molitor* and *Zophobas morio* larval worms. Turbidity assay evidence suggested that these bacteria have the ability to digest polystyrene and successfully survive [5]. With two potentially PS-degrading enzymes identified in the foregoing research, the efficacy and enzymatic catalyzing mechanism were still under investigation. That is the research that this paper takes up.

Our laboratory adopted the naming conventions of “ZM1” for the bacterial strains from *Zophobas morio* and “TM1” for the strains taken from *Tenebrio molitor*. 16sRNAs sequence of each revealed the taxonomic status of TM1 to be genus *Pantoea* and that of ZM1 to be genus *Klebsiella*. ZM1 has a 99% genetic similarity to *Klebsiella pneumoniae* DSM 30104; TM1 has a 96% genetic similarity to *Pantoea agglomerans* JCM 1236 [5].

With these potentially PS-degrading strains identified, they were cultured in a medium with polystyrene (PS) as the sole carbon source, making the TM1 and ZM1 no option for sustenance other than Polystyrene. According to our data, TM1 and ZM1 began to synthesize bacterial enzymes that seem capable to successfully degrading the PS-medium [5]. Fourier-transform

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Infrared Spectroscopy (FTIR) has been commonly applied in the field of biochemistry as a powerful technique to measure structural changes of organic and inorganic molecules and their functional groups [6-10].

In order to digest polystyrene as a carbon source, a secretion of the specific enzymes that can bind a polystyrene substrate which is required for the survival of the intestinal bacteria. Therefore, if any enzymatic catalyzation of the carbon bonds in the PS-medium occurred, FTIR could be a potential tool for measuring these molecular changes and demonstrating the sites where enzymes tend to activate [6].

2. Materials and Methods

2.1. Isolating polystyrene-degrading microbes

The experiments began by acclimating the *Tenebrio molitor* and *Zophobas morio* larval worms to a polystyrene diet for a period of 21 days (Fig. 1). PS-degrading bacterial strains TM1 and ZM1 were then isolated from the intestines by selective PS agar plates [5].



Fig. 1 Two larval worms' species subsisting on Polystyrene

2.2. Bacterial cultivation and PS-biodegradation experiments

The intestinal bacterial strains ZM1 and TM1 were inoculated and incubated within the PS-medium. A customized platform for polystyrene biodegradation testing of TM1 and ZM1 from the previously cited research was used [5]. The PS-medium was 50 ml, with 45 ml of inorganic salt basal buffer, 5 ml of polystyrene emulsion, and 0.05 ml of 38% yeast extract. Chloroform is used to dissolve polystyrene into emulsion, but after that it will be vaporized, so the final product doesn't contain chloroform.

At this stage, three tubes of PS-medium were loaded in 50 ml centrifuge tubes; two of them were inoculated with bacterial strains ZM1 and TM1. Each was cultivated for further FTIR testing. The remaining tube of medium was designated as the un-inoculated control. The FTIR cultivation courses were operated under a stable temperature at 37°C, 144 hours (6 days).

2.3. Turbidity analysis by absorbance and transparency

First, quantification and comparison of absorbance and transparency in PS-medium during cultivation course was conducted. Second, the absorbance and transparency levels of the PS-medium were used as a measure of turbidity. Turbidity is considered a quantitative indicator for PS degrading efficiency; analysis of this kind is also referred to as a Turbidity Assay [5, 11].

The absorbance of PS-medium was measured by microplate spectrophotometer (Thermo Scientific Multiskan GO), and 200 μ l of sample were loaded for each measurement: from 230-900 nm. The OD value was measured every 1 nm of wavelength. Transparency was determined by spectrophotometer (Metertech SP-830), and 1 ml of sample was loaded for each

test: from 400-700 nm. The percentage of transparency was measured at every 10 nm of wavelength. All measurements were performed with a time lapse at 0 hr, 24 hr and 48 hr.

2.4. FTIR analysis

FTIR spectroscopy was applied to the PS-medium after the cultivation of two strains to determine if enzymatic changes consistent with degradation of carbon bonds took place [12]. To avoid the interference of non-specific organic molecules within the microbes' outcome spectra, the samples (two inoculums and one control) were first centrifuged (4000 rpm, 25°C, 10 min) to remove debris and particles.

After centrifugation, 3 ml of supernatant of the medium was dried in a drying oven. The dried sample was then mixed with KBr salt powder (0.5 mg of sample plus 20 mg of KBr) and hand-pulverized by mortar; 10 mg of mixture were tableted for FTIR test afterward. FTIR spectroscopy (Perkin-Elmer FT-IR RX1) at 4000-400 cm^{-1} was performed and the samples were collected at day 0 and day 6 [13].

3. Result and Discussion

3.1. Absorbance and transparency determination

The turbidity of PS-medium during cultivation was measured based on absorbance and transparency. The absorbance levels of the experimental groups gradually decreased during the cultivation time lapse (Fig. 2), and the transparency rose (Fig. 3), while the control remained stable. This data suggests that the activation of TM1 and ZM1 caused PS to degrade in the medium.

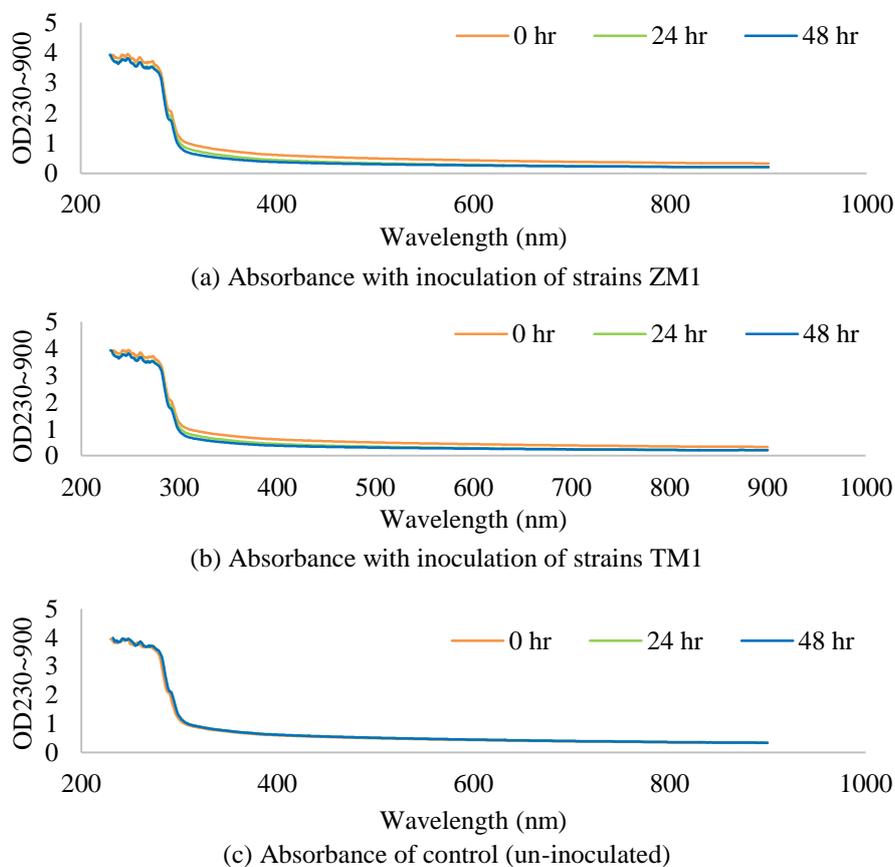
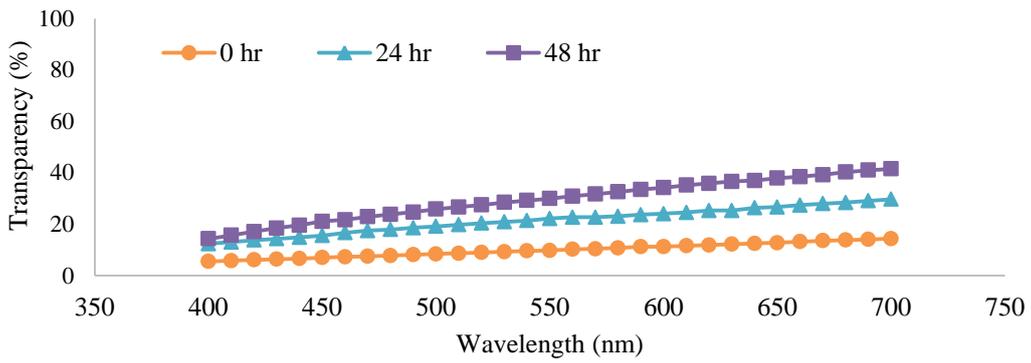


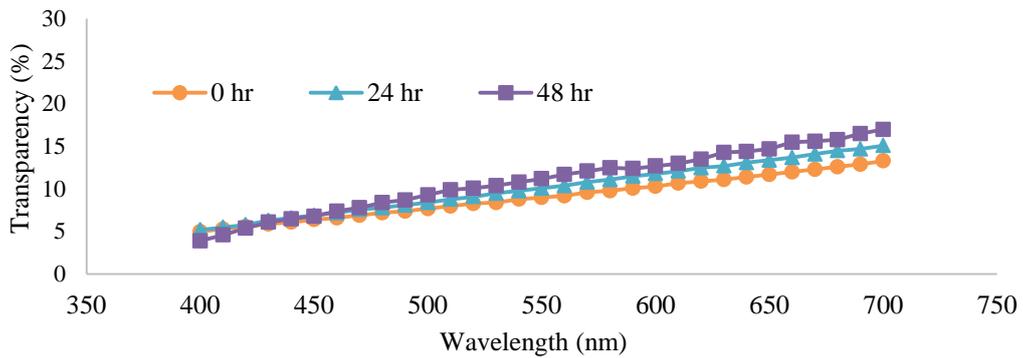
Fig. 2 Turbidity assay based on absorbance

To demonstrate biological degradation of polystyrene by the gut bacteria of *Tenebrio molitor* and *Zophobas morio* beetle larva this study performed turbidity assays based on absorbance and transparency on inoculated TM1 and ZM1 gut bacteria

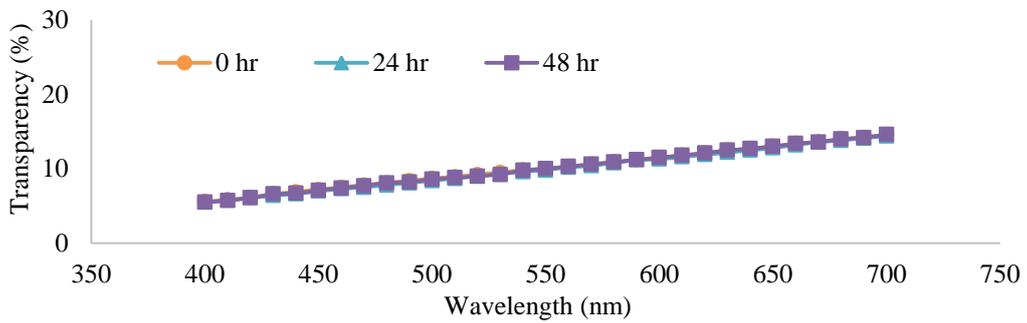
strains. The assay data reveal declining turbidity in the PS-medium across the full visible light spectrum, not only at the 600 nm point as the previous study had described [5].



(a) Transparency with inoculation of strains ZM1



(b) Transparency with inoculation of strains TM1



(c) Transparency of control (un-inoculated)

Fig. 3 Turbidity assay based on transparency

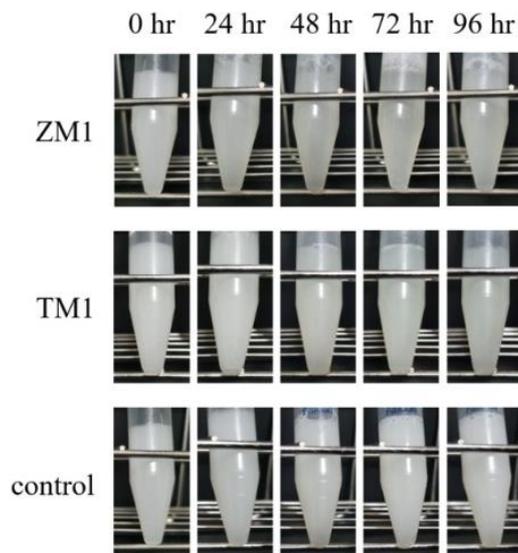


Fig. 4 Visualization of turbidity

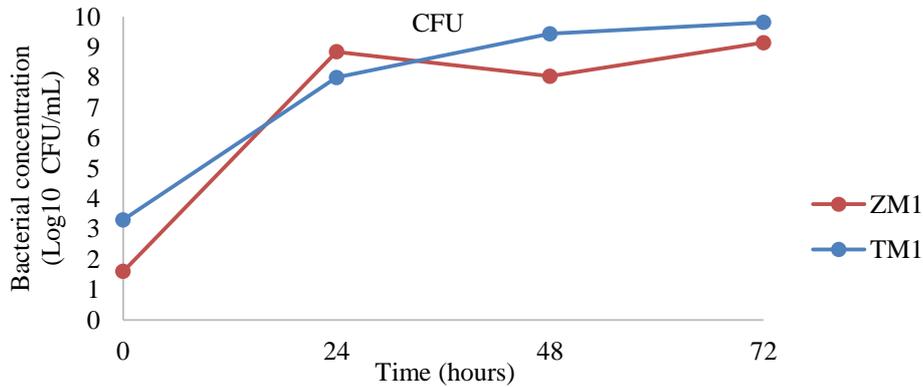


Fig. 5 Bacterial CFU (measured by MacConkey agar plate)

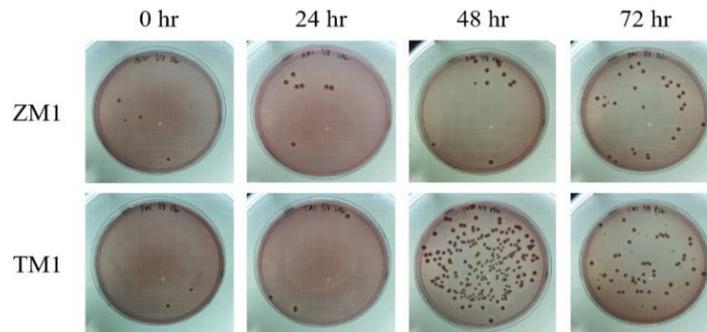


Fig. 6 Bacterial growth curve

A series of photos of strain TM1 and ZM1 growing in PS-medium also shows the changes of turbidity. PS-medium inoculated with TM1 and ZM1 become increasingly transparent from time at 0 hr to 96 hr, while the control group remains its originally opaque and turbid (Fig. 4). Figs. 5-6 shows the CFU data (measured by MacConkey agar plate) represent bacterial concentration and growth. In CFU test, the dilution factor of ZM1 at time 0, 24, 48 and 72 hr were 1, 10^7 , 10^6 and 5×10^6 , respectively; TM1 at time 0, 24, 48 and 72 hr were 10^2 , 2.5×10^6 , 10^6 and 10^7 , respectively. All colonies were yielded by plating 0.1 ml of diluted sample (Fig. 5).

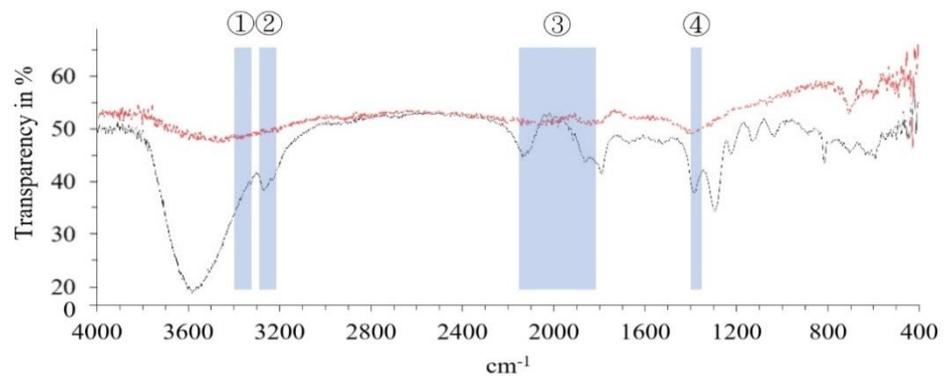
3.2. FTIR analysis

Fig. 7 shows an FTIR spectrum of the PS medium which is inoculated with TM1 and ZM1. Fig. 7(c) shows the un-inoculated control group for comparison. Both the test and the control samples contained the same polystyrene and inorganic salts medium. Hence, overlapping of peaks across these multiple salts and polystyrene substrate would be possible.

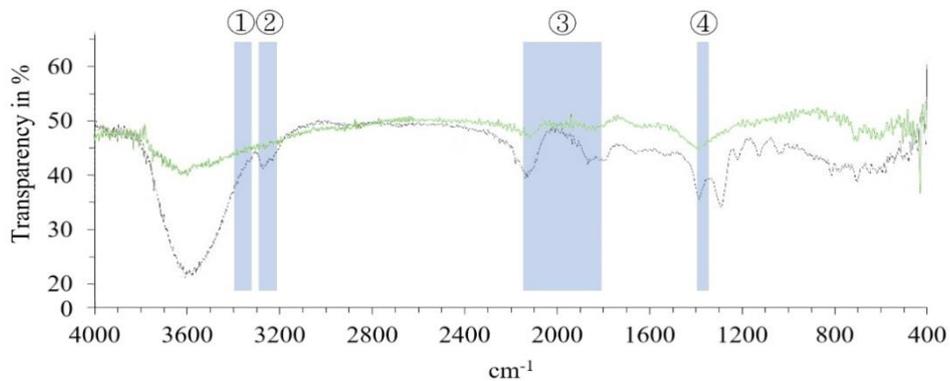
According to the spectrum at day 0, the absorption peaks at 3400 cm^{-1} , 1044 cm^{-1} , 994 cm^{-1} , 926 cm^{-1} , and 856 cm^{-1} have appeared with initial inoculation of ZM1 and TM1 (Figs. 7(a) and 7(b)). By comparing the spectral data onto Spectral Database for Organic Compounds (SDBS), these peaks were regarded as signal of glycerol, the remaining solution from ZM1 and TM1 bacterial stock (glycerol was used to preserve bacterial strain).

At the initial stage, the medium of inoculated groups share the same constituent of the control group, except glycerol from the bacterial stock solution. After 6 days of the experiment, weakening on multiple peaks occurs to the inoculated test groups compared to the control group. The reduced peaks seen in the range of 3000 cm^{-1} to 3100 cm^{-1} (Fig. 7) represent organic molecule functional groups including the band attributed to the aromatic C–H stretch. Another absorption peak change in the test group occurred in the range of 2800 cm^{-1} to 3000 cm^{-1} , which is assigned to the aliphatic C–H stretch. Additionally, changes seen occurring at 1654 cm^{-1} , 1494 cm^{-1} , and 1453 cm^{-1} are attributed to aromatic C = C stretch. And a C–H in-plane bending peak change occurred at 1109 cm^{-1} in the test group. Aromatic rings and aliphatic C–H-rich backbones are two of the typical molecular features of polystyrene.

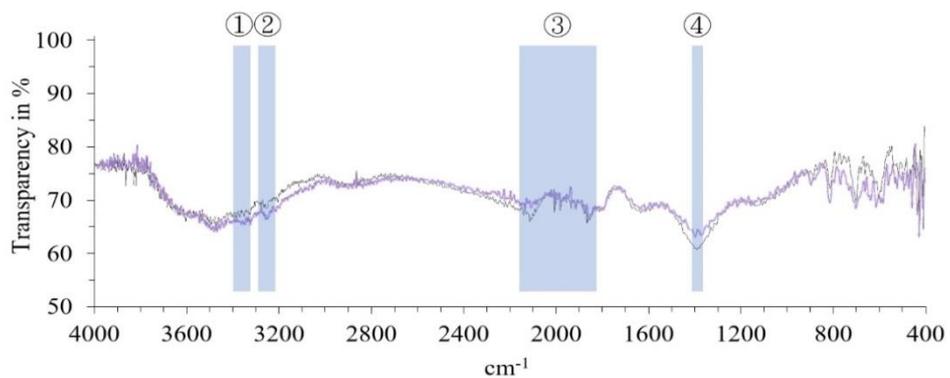
Compared with the spectra of the un-inoculated control, the FTIR peak reductions seen in the carbon chain functional groups indicates that ZM1 and TM1 have the ability to degrade carbon in Polystyrene. For those polymers with long hydrocarbon backbones, it is possible that enzymes produced by TM1 and ZM1 are degrading polystyrene from the side chains of the molecular backbone itself (Fig. 8).



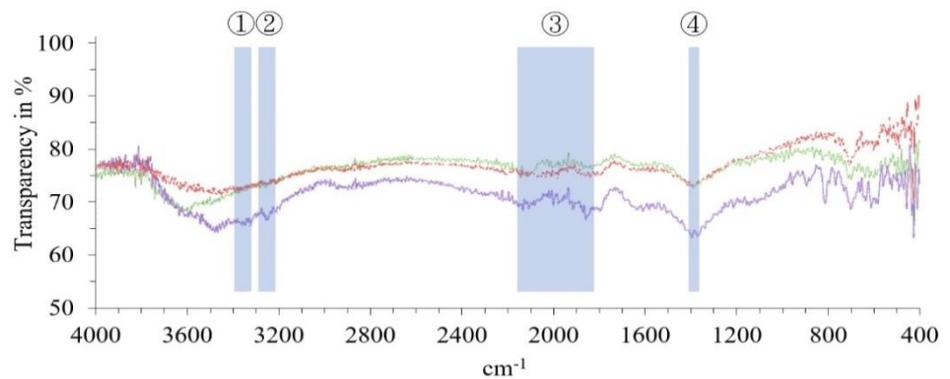
(a) Spectra with inoculation of strains ZM1 (day 0: black; day 6: red)



(b) Spectra with inoculation of strains TM1 (day 0: black; day 6: green)



(c) Spectra of un-inoculated control (day 0: black; day 6: violet)



(d) Spectra of all three groups at day 6 (ZM1: red; TM1: green; control: violet)

Fig. 7 FTIR spectrum within the frequency range of 4000–400 cm^{-1} ; ① refers to aromatic C–H stretch; ② refers to aliphatic C–H stretch; ③ refers to aromatic C = C stretch; ④ refers to C–H in-plane bending

The alteration or replacement of these side groups could create a new functional group, which would be feasible for other enzymes to break down along the main hydrocarbon chain. For example, research indicates that benzene 1,2-dioxygenase can degrade benzene (an aromatic derivative).

Considering that the aromatic rings of PS were also degraded in our study, TM1 and ZM1 may be shown to share a similar mechanism with benzene 1,2-dioxygenase degrading PS, by altering the benzene ring side chain.

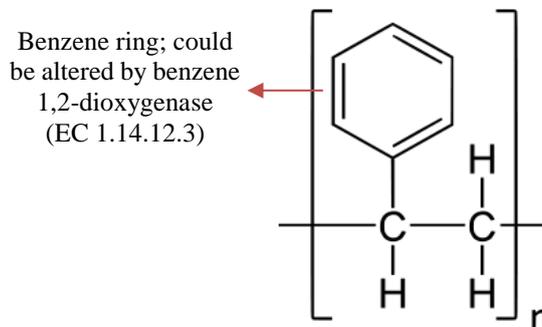


Fig. 8 molecular structure of polystyrene (PS), with a benzene side chain and C–H rich backbone

4. Conclusions

To exclude the possibility of bacteria death causing the turbidity decrease, CFU tests were conducted and these showed no morbidity in the bacteria. With bacterial colony stable and turbidity decreasing, the inference is that the decrease in PS-medium turbidity was the result of digestion of PS emulsion. Turbidity assay as a means of verifying plastic biodegradation, as well as a means to quantify the efficiency of plastic decomposition, has been cited in other studies [11, 14-15].

Through FTIR analysis, multiple carbon bonding changes were revealed. These changes at the aromatic C–H stretch; the aliphatic C–H stretch; the aromatic C=C stretch and the C–H in-plane bending offer further evidence to suggest that it is the digestive processes of the bacteria strains TM1 and ZM1 in the PS-medium which were responsible for the alternations of the absorption peaks levels related to organic function seen [16-17].

Other research has discovered a potential manner for enzymatic degradation of polystyrene, utilizing hydroquinone peroxidase produced by *Azotobacter beijerinckii* HM121 [18]. Hydroquinone peroxidase is an enzyme which can oxidatively catalyze phenols such as hydroquinone, a derivative of benzene rings [19]. As a polymer molecular structure, it is notable that polystyrene is a rich in benzene rings as well [20]. Other studies [21-22] have mentioned the degradation ability of certain beetle larvae including the *Tenebrio molitor*, however the identification of the specific microbes and their enzymatic degradation mechanisms were not yet be described in detail.

In conclusion, our laboratory utilized the entire range of visible light wavelengths to scan turbidity via absorption and transparency modes to explore the optimal condition for future experiments. Through FTIR analysis Our laboratory revealed structural changes of the polystyrene at the sites above as the TM1 and ZM1 incubations which indicate biodegradation. Further studies offers researchers the chances to isolate, identify and potentially clone the specific intestinal enzymes that enable bacterial strains such as TM1 and ZM1 to break down polystyrene and expanded genetic engineering targets for this biodegrading enzymes.

Conflicts of Interest

The authors declare no conflict of interest.

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