# Biodegradation and microbial CO<sub>2</sub> evolution of blended polymers in food-waste and coffee supplemented disposal environments

by

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#### Abstract

The dependency of many industries on single-use materials, has led to an accumulation of plastic waste and subsequent harm to the planet. Since avoidance of plastic is often not viable and the current processes of recycling have their own environmental impact, many have looked toward development of sustainable replacements for plastics. Homogenous blended polymers made from varied ratios of polycaprolactone (PCL), starch, and biochar were made in the Rochester Institute of Technology (RIT) Packaging Science Department, tested for percent weight loss and for carbon dioxide (CO<sub>2</sub>) evolution in soils. Testing was done across three burial matrices: soil amended with compost starter, soil with 30% pulverized food waste, and soil with 30% spent coffee grounds. Percent weight loss derived from burial experiments food waste and soil with compost starter show a positive correlation between material degradation and starch content. However, burial experiments with a coffee amended environment did not lead to as high of percent weight loss. CO2 evolution showed samples of 45% starch/ 45% PCL/ 10% biochar producing the highest cumulative amount (346.9mg) over 120 days. Fungal and bacterial isolates from the burial experiments show the most diversity in food and soil environments, as well as more species of interest (used in bioremediation or found to degrade polymers) compared to the coffee environment.

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#### **Chapter 1. Introduction**

#### 1.1. Overview

The increased demand for plastics in industrial and domestic applications has amplified the accumulation of plastic waste over time[1], [2]. Fifteen to thirty million tons of plastic waste is generated annually in the United States and Western Europe [3] and fifty-seven million tons is generated globally [1]. The presence of these discarded non-biodegradable polymeric materials in terrestrial and aquatic habitats poses a substantial threat to the various species that reside in these affected ecosystems, as well as overall ecosystem function [4]. In addition to directly impacting ecosystem health after consumer use, plastic manufacturing is solely petroleum based which leads to greenhouse gas emissions. These emissions are problematic when considering the drastic greenhouse gas reductions needed to combat climate change [5]. Current manufacturing, use, and disposal of plastic material is consistently unsustainable, and to lessen further anthropogenic harm to the planet there needs to be a disruption in our current plastic usage.

To lessen the accumulation of plastic waste, we want to think beyond current methods. Reduce, reuse, and recycle is a widely known slogan meant to promote sustainable navigation through plastic consumption [6]. However, reducing plastic consumption is often unavoidable because of how deeply plastic is engrained in modern society. Reuse of plastic materials may extend the lifetime of a single-use plastic, but disposal is inevitable. Recycling can reutilize nonvirgin plastics, but within the recycling process are steps that are environmentally detrimental. Also, the potential impact of recycling is often greater in the minds of consumers versus what is achieved or possible.

The recycling rate of plastic is low compared to the high rate of plastic use. Also, of the small percent of plastics sent to be recycled, some materials are unable to be recycled and then discarded. The processing and chemical treatment of recyclables creates its own environmental impact. Fillers and additives routinely used within mechanical recycling processes are often pollutants, eventually making their way into water supplies [6]. Public ideations of recycled plastic has the same potential as virgin plastic. However, this is not the case. Plastic materials are often "downcycled" into materials that are not able to be recycled again, therefore virgin plastic is still needed to maintain the anthropogenic demand of plastic products [7]. As for recycling rates, from 2015 to 2017 total plastic recycling fell from 9.1% to 8.4% of total plastic waste in the U.S., according to the EPA [8].

At our current rate of plastic use and inevitable disposal, we need to make more substantial changes to how we interact and produce plastic [9]. Instead of looking at ways we can sustainably clean up conventional plastic waste, we should be considering how we can change plastic itself to be more environmentally friendly [9]. Development of biodegradable plastic alternatives sourced partially or fully from renewable resources has the potential to change the environmental impact of single and short-term use packaging.

When developing biodegradable packaging, the many industries that utilize plastic material need to be considered. To maximize the environmental benefits gained from the adoption of

biodegradable plastics, it's necessary to identify a target industry that heavily utilizes plastics and has widespread impact. The food industry, specifically food packaging, is an ideal target for implementation of biodegradable packaging material [10]. This industry is convenience-driven with individually portioned and ready-to-eat food packaging used daily by a large population of individuals. If the plastic utilized for food packaging were more sustainable, the positive environmental impact of this change could be massive.



Figure 1. Cumulative plastic waste generation and disposal. Graph source: [10]

#### **1.2. Materials of interest**

#### 1.2.1. Polycaprolactone (PCL)

The Packaging Science Department at R.I.T. has identified polycaprolactone (PCL) and starch based blended polymers as having high potential for utilization as biodegradable packaging. Polycaprolactone is a hetero-chain biodegradable polymer, once primarily utilized for medical applications like drug delivery and tissue engineering [11], [12]. Biodegradation is an enzymatic process driven by microbial activity. The inclusion of oxygen atoms in the molecular backbone of PCL characterizes the aliphatic polyester as a hetero-chain biodegradable polymer. Unlike conventional plastic which is comprised solely of carbon-carbon bonds, PCL contains an ester functional group, making the material more susceptible to enzymatic microbial processes. Polycaprolactone has also been shown to biodegrade in sea water via hydrolysis and enzymatic microbial activity. The degradation of PCL in salt water is also greatly influenced by mechanical

stress and light exposure present in ocean environments [13]. The presence of the ester functional group within the PCL backbone also results in polymer flexibility which is necessary for the polymer to fit correctly into the enzymatic active site allowing for microbial attack of the polymer [14], [15]. However, PCL has a slow rate of hydrolytic degradation, ranging from two to three years [16]. To successfully utilize PCL as a biodegradable packaging material in efforts to reduce packaging waste buildup, the rate of degradation as a function of time must be increased. [17]



#### Figure 2. Polycaprolactone (PCL).

Chemical structures generated using ACD/ChemSketch [18].

#### 1.2.2. Starch

Starch is a natural polymer that is highly biodegradable, however the hydrophilicity of starch proves problematic when trying to develop a mechanically sound packaging material. Incorporation of starch with PCL in a homogenous blended polymer may have the potential to yield a blended polymer with the benefits of PCL's mechanical function, and increased degradation rates due to starch content. The inclusion of starch with more mechanically sound polymers has been identified as having potential for development of promising biodegradable blended polymers [19], [20]. Starch is comprised of two isomers: amylose and amylopectin. The specific ratio of these two components depend on the botanical source of the starch [21].



Figure 3. Structure of the starch isomer: amylose.

Chemical structures generated using ACD/ChemSketch [18].



Figure 4. Structure of the starch isomer: amylopectin.

Chemical structures generated using ACD/ChemSketch [18].

#### 1.2.3. Biochar

The Intergovernmental Panel on Climate Change (IPCC) has identified biochar as having potential for carbon sequestration, making it a desirable material to utilize for sustainable practices [22] Biochar is a high carbon content material produced via pyrolysis of various organic feedstocks under anaerobic conditions, molecular morphology of biochar can vary depending on feedstock and pyrolysis methods [23]. Many potential feedstocks of biochar are food waste based. Pyrolysis of food waste to form a new material would divert food waste from landfills, thereby reducing methane emissions. High porosity makes biochar an attractive candidate for biodegradable material synthesis because microbial surface colonization is needed for microbial decomposition. The use of biochar based degradable materials may benefit the burial environment post-material degradation, since biochar has been shown to improve soil quality [24], [25]





# 1.3. Potential within food industry

Targeting the food industry for the adoption of biodegradable packaging material poses a unique opportunity to find a multi-dimensional solution to two major sustainability issues in the industry: plastic waste and food waste [26]. When looking to implement a biodegradable material into consumer use, it is vital to determine if the potential disposal environment of the material will facilitate microbial decomposition. Decomposition is a microbial process, so if the disposal environment is not hospitable for the necessary or ideal microbial community the materials may not degrade. Food waste can serve as a powerful agricultural fertilizer due to microbial richness [27]. If biodegradable materials were adopted within the food industry, could food waste be utilized as a disposal environment to help degradation? If an end-of-life plan was utilized for the disposal of a biodegradable food packaging that incorporated food waste, would the rate of decomposition increase over time? This would avoid a shortfall seen in many currently available biodegradable materials where their disposal environment post-consumer use does not facilitate decomposition. Consequently, a slow rate of degradation would lead to a buildup of packaging waste.



Figure 6. Product lifetime distributions. Graph source: [28]

## **Chapter 2**. Materials and methods

#### 2.1. Polymers

Material name (used in this paper)	Material composition	Material preparation
PCL60	60% polycaprolactone (PCL) 40% starch	Pressed homogenous polymer ~0.5mm
ST60	60% starch 40% PCL	Pressed homogenous polymer ~0.5mm
BC	45% starch 45% PCL 10% biochar	Pressed homogenous polymer ~0.5mm
CELL	100% cellulose	Paper (<0.5mm)
LDPE	low density polyethylene	<1.0mm
PET	Polyethylene terephthalate	Film
PP	Polypropylene	Film
BOPP	Biaxially Oriented Polypropylene	Film
PLA	Polylactic Acid	Film
BIOBAGS	MATER-BI	Film

Table 1. Overview of all materials tested throughout study.

## 2.2. Methods

Three primary experimental methods were performed: burial experiment, carbon dioxide capture, and microbial culture and identification. The primary polymer blends being tested are designated by: PCL60, ST60, BC, and CELL (*table 1*). Other polymers (PP, PET, BOPP, PLA, BIOBAGS) were tested for percent weight loss in a soil environment to serve as reference to currently used packaging materials. BIOBAGS and PLA are both marketed as biodegradable. However, products marketed as biodegradable may require specific post-consumer treatment and burial matrix to uphold their marketed biodegradability. These necessary treatments (such as shredding) of biodegradable material create a false consumer illusion of the end-of-life fate when a biodegradable material is disposed of.

Comparison of degradation rates was made between polymer type, as well as burial matrices. Three burial matrices, or disposal environments, were applied across all testing of the four main polymers: PCL60, ST60, BC, CELL. These matrices are designated in this paper as: soil, coffee, and food. Their specific composition is detailed in *table 2*.

Burial Matrix Designation	Matrix composition
Soil 100% Miracle-Grow® All Purpose Pottin	
	Soil
	Amended with Espoma® compost starter (35-
	40 grams per bin)
Food Waste	30% pulverized food waste (64% potato peels
	and 36% tomato scraps (by weight))

	70% Miracle-Grow® All Purpose Potting Soil	
	(compost started omitted)	
Coffee	30% spent coffee grounds (caffeine extracted)	
	70% Miracle-Grow® All Purpose Potting Soil	
	(compost starter omitted)	

Table 2. Composition of burial matrices used across all methods of testing.

# 2.2.1. Initial Burial Experiment

The percent weight loss of polymers was measured in soil environments over 122 days following the methods of similar burial studies [29]. Two novel polymer blends were tested: PCL60 and ST60. These blended polymer samples were supplied by the Diaz-Acosta lab group from the Packaging Science Department at RIT and were prepared as thin malleable pressed polymers (~0.5mm). In addition, BIOBAGS and CELL were tested to provide comparison. CELL is a known biodegradable material, and BIOBAGS are a commercially available product marketed as fully compostable. Ten samples of PCL60 and ST60, and 6 samples each of BIOBAG and CELL were tested for percent weight loss via decomposition within a burial environment. All samples were cut into 3cm<sup>2</sup> pieces.

Miracle-Grow® All Purpose Potting Soil was used as a burial matrix. Espoma® Compost Starter was used as a microbial source for the soil. The compost starter was added according to the manufacturers' recommendations with 35-40grams being added based on the surface area of the bins. Bins were filled with the soil and compost mixture approximately 11.5cm deep. This blended burial matrix of potting soil and compost starter will be designated as a 'soil' burial environment as detailed in *table 2*. All bins were watered routinely and monitored with a moisture meter throughout the experiment. Bins were also given drainage holes and lined with mesh to avoid oversaturation of soil. Arrangement of samples within bins is shown in *figure 7*. with sample buried approximately 6cm beneath the soil surface.

The burial experiment ran for 122 days. At days 0, 10, 20, 30, 42, 55, 65, 77, 88, 98, 112, and 122 each sample was weighed. At these checkpoints, samples were cleaned of debris, washed with alcohol, and weighed to track weight loss via decomposition. After weight assessment, samples were placed back into soil as they previously were. Photos were also taken to visually monitor the breakdown of materials. Routine water application to bins was done to maintain a consistent soil moisture content, without saturation of soil. At the conclusion of the experiment, weights were analyzed and percent weight loss was calculated to easily compare degradation as a function of weight loss, since weight of initial samples varied due to material specific density.



Figure 7. Initial burial experiment layout.

Shown in *figure 7*, samples were labeled with an alpha-numeric system for easier data collection to follow the decomposition of individual samples. Soil was uniform across all bins and each bin contained duplicates of the same sample type. Sample groups A, B, E, F, K, L, M, and N were in bins the larger bins (45.7cm  $\times$  38cm) and sample groups C, D, G, and H were in the smaller bins (39.4cm  $\times$  28cm).

# 2.2.2. Revised Burial Experiments

Methods were slightly revised for the remaining burial experiments. Revisions were made to improve and further standardize the general methods used to collect preliminary burial data. The table below shows the experimental variables for all revised burial experiments performed:

Exp	Burial	Tested	Duration	Data Sampling	Polymer size
#	Matrix	Polymers			
2	Soil	PCL60, ST60, BC, CELL, LDPE	120d	Initial weight as average of samples, experimental weight as individual sample not reused for later checkpoints	2.5cm <sup>2</sup> – PLC60, ST60, BC, LDPE 3.8cm <sup>2</sup> -CELL
3	Coffee	PCL60, ST60, BC,	120d	Initial weight as average of samples, experimental weight as	2.5cm <sup>2</sup> – PLC60, ST60,

		CELL, LDPE		individual sample not reused for later checkpoints	BC, LDPE 3.8cm <sup>2</sup> -CELL
4	Food	PCL60, ST60, BC, CELL	120d	Initial weight as each individual samples' initial weight, experimental weight as individual sample not reused for later checkpoint	7.6cm <sup>2</sup> - All samples
5	Soil	PP, PET, BOPP, PLA	120d	Initial weight as average of samples, experimental weight as individual sample not reused for later checkpoints	7.6cm <sup>2</sup> - All samples

Table 3. Overview of revised burial experiment setups.

The percent weight loss of PCL60, ST60, CELL, and BC was monitored in three burial environments: soil, coffee, and food waste, over 120 days. The placement and sample collection of the three revised burial experiments differed compared to the initial burial experiment, as illustrated in *figure 7*. To better regulate data collected, triplicates of each sample type was placed into each bin. Shown in *figure 8* for coffee and soil environments, each bin represented a different timepoint within each experiment. All bins were watered routinely and monitored with a moisture meter throughout the experiment. At each timepoint samples within one bin are unearthed, cleaned, dried in an oven at 60 °C for 4 hours, cleaned again, and weighed. No samples were placed back into the burial environments for further data collection. This allowed for the samples to be completely untouched until their respective checkpoint day, unlike how sample data was collected within the first burial experiment.

Percent weight loss was calculated with the same formula in the initial and revised burial experiment:

Weight Loss (%) = 
$$\frac{w_i - w_e}{w_i} \times 100$$

Where  $w_i$  is the initial dried weight of sample and  $w_e$  is the experimental weight of sample after being cleaned and dried.



Figure 8. Revised burial experiment layout

*Figure 8* illustrates the revised burial experimental setup for two checkpoints, t=10 and t=20, for two burial environments: coffee and soil. For all revised burial experiments, the number of bins was reflective of the number of checkpoints. At each timepoint, the samples within one bin for both environments was unearthed. The food burial experiment was setup in the same manner as shown in the figure.

#### 2.2.3. Carbon Dioxide Evolution

Biometer flasks were used to measure the microbially driven  $CO_2$  evolution of polymers in each burial environment: soil, food, and coffee. This experimental protocol to measure  $CO_2$ production is outlined by EPA guidelines [30]. A 2.5cm<sup>2</sup> sample of polymer in 50g of burial matrix (soil, food, or coffee) was placed in the Erlenmeyer portion of a biometer flask. The sidearm contained 10mL KOH, the concentration of KOH used varied due to differing  $CO_2$ production across experiments. An Ascarite tower was used to absorb atmospheric  $CO_2$  when opening the system to remove the spent KOH at 10-day intervals. Polymers tested across the three burial matrices include: PCL60, ST60, BC, and CELL. The described biometer flask setup was done in duplicate for each polymer type, as well as the respective burial matrix alone. The burial matrix-only setups were done to obtain approximate measures of endogenous microbial activity of each disposal environment. Flasks were kept at room temperature and incubated for 120 days. Data was collected in duplicate, with two flasks labeled A and B for each polymer type plus the control setup.

The 10mL KOH was drawn from the side arm via syringe for titration and replenished every 10 days. Titration of KOH was done using 0.1mL of a 1% phenolphthalein solution as a pH indicator and HCl as the acid. A sample of control KOH was also neutralized each week to get the  $V_c$  value. Molarity of KOH and HCl was adjusted based on absorption of CO<sub>2</sub>. Changes in reagent molarity is accounted for within the equation by the *CF*. The equation below was used to determine the gross amount of CO<sub>2</sub> production per flask [31] [32]:

$$Mg \ CO_{2} = (V_{c} - V_{E}) {\binom{M_{CO_{2}}}{2}} (M_{HCl}) (CF)$$

$$Correction \ Factor = CF = \frac{M_{HCl}}{M_{KOH}}$$

$$V_{c} = vol \ of \ HCl \ to \ neutralize \ control$$

$$V_{F} = vol \ of \ HCl \ to \ neutralize \ experimental$$

Calculations were done to measure the  $CO_2$  production for both experimental setups and control setups (burial matrix only). To get the amount of  $CO_2$  produced by the microbial breakdown of materials, the  $CO_2$  production of the soil-only controls were subtracted to account for endogenous microbial activity within the soil.



Figure 9. Experimental setup of biometer flasks used to collect CO<sub>2</sub> evolution data.

# 2.2.4. Microbial Culturing and Identification

Soil samples were taken off the surface of polymers: PCL60, ST60, CELL, and BC for burial experiments 2, 3, and 4. Samples were also taken off LDPE, which was tested in soil and coffee environments. One-gram samples of the burial matrix was scraped off the surface of each polymer type, then diluted in a 0.9% NaCl solution for serial dilutions and plating. Potato dextrose agar (PDA) containing 100ug/ml streptomycin was used for fungal cultures and plate count agar (PCA) was used for bacterial cultures. The dilutions of fungal cultures were plated from  $10^{-2}$ - $10^{-5}$  and the dilutions of bacterial cultures were plated from  $10^{-3}$ - $10^{-6}$ . Plate counts were done at day 3 of incubation at room temperature. Day 3 was chosen because it showed the best distribution of countable colonies (30-300) for cultures across all environments. The soil samples were collected at day 100 to select from the most prevalent colonies remaining at the end of the experiment. Plate counts were done using an image mapping software (ImageJ)[33].

DNA sequencing of bacterial isolations was done at the genomics lab at RIT, under the supervision of Dr. Andre Hudson. Following dilution, plating, and incubation of samples, bacterial colonies of interest were isolated for purity and the 16S rRNA segment was amplified via polymerase chain reactions (PCR). PCR is used to amplify the genomic area of interest, which for bacterial samples is the 16S rRNA segment. For each isolated bacterial colony sample, a PCR tube was inoculated with the sample,  $1\mu$ L forward and reverse V3/V4 primers,  $12.5\mu$ L master mix, and  $10.5\mu$ L nuclease-free H<sub>2</sub>O. Samples are then centrifuged and processed in the thermo cycler. Gels were made with 50mL TAE 1X and 1.2g agarose, heated until fully dissolved. After cooling slightly,  $3\mu$ L of ETBR is added prior to pouring in the gel box for solidification. Larger gels were made with triple the amounts given. After the gel is solidified and the comb removed, TAE 1X buffer is added to the box to completely cover the gel, channels are filled with  $5\mu$ L PCR samples, and one lane is designated to hold the DNA ladder as reference. Once the gel box is connected to a current with the channels closest to the negative node, the gel electrophoresis is run until the visible dye travels past the end of where the ladder extends.

PCR samples that successfully show bands at the 16S point on the DNA ladder are then cleaned up and sent to GENEWIZ for analysis. Bands present at the 16S point in the DNA ladder show amplification of the necessary 500bp section of DNA needed for identification. Samples that show replication of the 16S DNA segment are then tested for DNA concentration (ng/ $\mu$ L). Based on DNA concentrations, amounts totaling at least 20ng DNA are combined with 1ng Forward primer and brought to 15 $\mu$ L with nuclease-free H<sub>2</sub>O.

Sequencing data was then run through the BLAST NCIB to analyze for similarity against the current database of bacterial sequences. [34] Specifically nucleotide BLAST was used to find percent match of 16S rRNA sequences. After species were identified, a cladogram of species found was created with MUSCLE and NCIB Genomic Workbench programs shown in *figure A*. *1*. in the appendix [35][36].

Genomic sequencing was not possible for fungal specimens, due to the necessary primers not being available. So, identification of fungal colonies was done based on colony morphology and microscopy staining. Lactophenol cotton blue was used to stain fungal slides. This allowed visualization of hyphae variations (septate or non-septate) as well as spore morphology. [37]

# 2.2.5. SEM imaging

Samples of polymers retrieved at days 40 and 100 of burial experiments 2, 4, and 5 were analyzed via SEM imaging. Scanning electron microscope (SEM) images were taken off the surface of polymers to visualize the progression of material decomposition at a microscopic level. Prior to taking the images, samples roughly 5mm<sup>2</sup> were sputter coated with a metal on their

surface. Sputter coating is a necessary pre-treatment when SEM imaging surfaces. The thin layer of a conducting material allows the electrons that are bombarding the material during imaging to evenly disperse over the entire surface- resulting in an accurate SEM depiction of the material surface[38]. Images were taken at 500X and 2000X. These magnifications were chosen based on related literature [39]. Images were also taken on the polymer samples in their initial state to provide a starting point comparison.

## Chapter 3. Results: Burial Decomposition

#### 3.1. Overview

Multiple burial experiments were performed to follow the progression of blended polymer decomposition with variant soil environments. Soil amended with food waste, coffee grounds, or soil amended with compost starter (details of burial matrix composition in *table 2*) were used as variable environments for the novel blended polymer samples that were tested for decomposition within soil.

## 3.2. Soil burial experiment

Exp	Burial	Tested	Duration	Data Sampling	Polymer size
#	Matrix	Polymers			
1	Soil	PCL60, ST60, CELL, BIOBAGS	122d	Removal, clean of debris, weigh, then re-bury for addition data collection	2.5cm <sup>2</sup>

Table 4. Overview of experimental details for burial experiment 1.





The starch-heavy blended polymer ST60 degraded at a faster rate compared to PCL60, . Cellulose paper degraded as expected, functioning well as a positive control. On day 42, no distinguishable pieces of the cellulose material were found in the soil or able to be weighed. BioBag samples proved difficult to accurately weigh due to their low weight, leading to their data collection to be stopped on day 55.

When collecting data for BIOBAG decomposition, the low initial weight of the material (<0.1g) led to difficulties weighing the material as the experiment progressed. The light-weight material was not able to be cleaned of soil adhering its' surface without damaging the sample. Weights measured after day 20 showed innacuracies because the weight of soil unabled to be cleaned off heavily impacted the weighed mass of the BioBag material. This led to data collection on BioBag decomposition to be haulted after day 55, shown in *figure 10*. However, it should be noted that visually there was no major physical deterioration of the material within the 55 days data was collected.

Exp	Burial	Tested	Duration	Data Sampling	Polymer size
#	Matrix	Polymers			
2	Soil	PCL60, ST60, BC, CELL, LDPE	120d	Initial weight as average of samples, experimental weight as individual sample not reused for later checkpoints	2.5cm <sup>2</sup> – PLC60, ST60, BC, LDPE 3.8cm <sup>2</sup> -CELL

## 3.3. Revised soil burial experiment

Table 5. Overview of experimental details for burial experiment 2



Figure 11. Percent weight loss of polymers in soil burial experiment 2.

Prior to day 60, bins of samples were kept in a temperature-controlled lab at RIT, however due to COVID-19 school closures in March of 2020, the bins were relocated to my apartment for the remainder of the experiment. The temperature variance from an unregulated residential heating system may be to blame for the irregularities in data after day 70.

Decomposition of materials followed similar trends displayed by the preliminary burial experiments. Days 0-120 showed samples of ST60 had higher degradation trend compared to PCL60. In this time span the ST60 samples also showed a slightly higher degradation trend when compared with BC. The three experimental blended polymer samples showed a positive correlation between degradation and starch content for days 0-120. For days 0-70, prior to possible temperature variations, CELL functioned as a positive control with the highest degradation trend line, and LDPE functioned as a negative control with the lowest degradation rate.

Exp #	Burial Matrix	Tested Polymers	Duration	Data Sampling	Polymer size
3	Coffee	PCL60, ST60, BC, CELL, LDPE	120d	Initial weight as average of samples, experimental weight as individual sample not reused for later checkpoints	2.5cm <sup>2</sup> – PLC60, ST60, BC, LDPE 3.8cm <sup>2</sup> -CELL

#### 3.4. Coffee burial experiment

Table 6. Overview of experimental details for burial experiment 3.

As stated earlier for experiment 2, burial bins for experiment 3 were also relocated after COVID closures in March 2020. Data preceding day 50 was kept under temperature conditions aligning with the other experimental setups. However past day 50, the watering schedule of the bins did not maintain soil moisture at the higher room temperature they were being kept at. Despite the shortened period of time where temperature and moisture levels were uniform, data collected prior to day 50 showed low rates (<10%) of percent weight loss for the three novel polymers besides BC at day 10 and the control: CELL. The sporadic jumps in CELL percent weight loss, paired with the generally low percent weight loss for the remaining polymers do not support coffee amended soil as an ideal or improved degradation environment for the polymers being tested. Although generally coffee grounds are accepted as a food waste type in composting, this may negatively impact the microbial community needed for degradation of materials. One study found that amending plant material-based composts with coffee grounds led to compost acidification and reduced biological activity. Reduced biological activity as defined by reduced germination capacity of seeds compared to control compost.[40]





Exp #	Burial Matrix	Tested Polymers	Duration	Data Sampling	Polymer size
4	Food	PCL60, ST60, BC, CELL	120d	Initial weight as each individual samples' initial weight, experimental weight as individual sample not reused for later checkpoint	7.6cm <sup>2</sup> - All samples

3.5. Food burial experiment

Table 7. Overview of experimental details for burial experiment 4.

In burial experiment 4, CELL was fully degraded by day 40, and continued to be fully degraded at all check points following day 40. Similar to burial experiments 1 and 2 in soil, ST60 showed higher percent weight loss overtime when compared to the other novel polymer blends. Interestingly, at day 80 BC and PCL60 followed similar downward trends, while ST60 peaked in degradation amount. This could indicate microbial preference to the ST60 material within the day 80 bin. The ending percent weight loss for the novel polymers were also within 10% of one another. When thinking about curating the ideal decomposition environment, having samples clustered closely in an upward degradation trend is ideal.



Exp	Burial	Tested	Duration	Data Sampling	Polymer size
#	Matrix	Polymers			
5	Soil	PP, PET, BOPP, PLA	120d	Initial weight as average of samples, experimental weight as individual sample not reused for later checkpoints	7.6cm <sup>2</sup> - All samples

3.6. Soil burial experiment with PP, PET, BOPP, and PLA

Table 8. Overview of experimental details for burial experiment 5

Decomposition of commercially available films over 120 days are shown in *figure 14*. BOPP samples were found to average 31% weight loss by day 120. Film samples tested showed soil adhesion and weathering as the burial experiment progressed, which lead to the increasing and decreasing levels of decomposition seen at the progressing time points. Compared to the novel polymers (PCL60, ST60, and BC) tested in the same burial environment, PP, PET, PLA, and BOPP did not show a general upward trend in percent degradation.





# 3.7. SEM imaging



Figure 15. SEM images of polymers post-burial at 2000X magnification.



Figure 16. SEM images of polymers post-burial at 2000X magnification.

2000X magnification scanning electron microscope (SEM) imaging of polymer samples from burial experiments 3, 4, and 5 at days 40 and 100. From left to right: CELL, LDPE. BC, ST60, PCL60. Images of untreated polymers included for reference (T=0). 'X' represents complete degradation of material, and the black boxes are for polymers that were not tested in all environments.

#### 3.8. Burial experiment statistics

Percent degradation data gathered from burial experiments 2, 3, and 4 were compared at days 40 and 100. A two-way Analysis of variance (ANOVA) was performed on each set of data: T=40 and T=100 using the JMP program[41]. The ANOVA method was used to test if the null hypotheses was supported or rejected by the data. Meaning, if the null hypothesis is accepted,

there is no difference in means. This hypothesis testing looks at comparisons withing polymer type, burial environment, and the interactions between those two factors. For both T=40 and T=100 all null hypotheses were rejected.

# 3.8.1. T=40 statistics

Source	DF	Sum of	Mean	F Ratio
		Squares	Square	
Model	11	2.5642384	0.233113	9.1525
Error	24	0.6112736	0.025470	Prob > F
C. Total	35	3.1755120		<.0001*

Table 9. Analysis of variance for percent weight loss data from burial experiments 2, 3, and 4 at day 40.

Source	Nparm	DF	Sum of Squares	F Ratio	<b>Prob</b> > <b>F</b>
POLYMER	3	3	1.0275907	13.4485	<.0001*
ENVIRONMENT	2	2	0.6299486	12.3666	0.0002*
POLYMER*ENVIRONMENT	6	6	0.9066991	5.9332	0.0007*

Table 10. Effects Tests for percent weight loss data from burial experiments 2, 3, and 4 at day 40.



Figure 17. Least square means plot for T=40 burial experiment data.

# 3.8.2. T=100 statistics

Source	D	Sum of	Mean	F Ratio
	F	Squares	Square	
Model	11	3.5624167	0.323856	25.6498
Error	24	0.3030261	0.012626	<b>Prob</b> > <b>F</b>
C. Total	35	3.8654428	_	<.0001*

Table 11. Analysis of variance for percent weight loss data from burial experiments 2, 3, and 4 at day 100.

Nparm	DF	Sum of Squares	F Ratio	<b>Prob</b> > <b>F</b>
3	3	2.9491721	77.8592	<.0001*
2	2	0.2733563	10.8251	0.0004*
6	6	0.3398883	4.4866	0.0035*
	<b>Nparm</b> 3 2 6	Nparm         DF           3         3           2         2           6         6	NparmDFSum of Squares332.9491721220.2733563660.3398883	NparmDFSum of SquaresF Ratio332.949172177.8592220.273356310.8251660.33988834.4866

Table 12. Effects Tests for percent weight loss data from burial experiments 2, 3, and 4 at day 100.



Figure 18. Least square means plot for T=100 burial experiment data.

# Chapter 4. Results: Carbon Dioxide Evolution

## 4.1. Overview

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Carbon dioxide production was determined by the pH change of KOH within biometer flasks, and the subsequent titration every 10 days of the experimental KOH. Mg CO<sub>2</sub> shown on
the *figures 17, 18*, and *19* below are the net totals  $CO_2$ , where the each set of data is averaged and the average of the control flask containing only the respective burial matrix is deducted. The plotted lines represent the cumulative totals and reflect the total of the data gathered at and prior to the timepoint given.

# 4.2. CO<sub>2</sub> evolution in soil environment

 $CO_2$  evolution in soil burial matrix mostly followed the ranking of percent weight loss of the same samples in the soil burial experiment, minus CELL. Plateaus of  $CO_2$  production were seen for all materials except BC by day 110. In the soil biometer flasks, polymers with higher starch content produced the most  $CO_2$ . Over the 120 days, the net production of  $CO_2$  for ST60 was 99.5mg, BC evolved 93.5mg, and PCL60 produced 43mg. CELL produced 29.8mg and the lowest production was seen from LDPE (22.6mg). There was also an initial plateau in net  $CO_2$ production seen for all polymers until day 30.

# 4.3. CO<sub>2</sub> evolution in coffee environment

Unlike the percent weight loss data from the coffee burial experiment, the CO<sub>2</sub> evolution of polymers all followed a positive trend in coffee, and by day 120 ST60 produced more CO<sub>2</sub> (119.6mg) versus CO<sub>2</sub> evolution seen with soil. PCL60 produced 60.2mg CO<sub>2</sub> in coffee, almost 1.5 times the amount seen in soil. CELL also produced almost twice as much CO<sub>2</sub> in coffee (57.8mg) compared to soil. BC produced the least CO<sub>2</sub> (25.3mg) versus other polymers in coffee. Both BC and PCL60 saw plateaus of net production from days 80 to 120.

# 4.4. CO<sub>2</sub> evolution in food environment

 $CO_2$  evolution in the food burial matrix showed BC, CELL and PCL60 with higher production when compared to the other  $CO_2$  experiments: coffee and soil. BC produced the highest average of  $CO_2$  across all burial matrices (346.9mg) which was over three times the amount of the second highest  $CO_2$  evolution across all experiments: ST60 in coffee (119.6mg) shown in *figure 20*. ST60 only produced an average of 25.0 mg CO<sub>2</sub> across 120 days, the lowest  $CO_2$  evolution of ST60 across all burial matrices. Fungal growth in individual flasks of CELL and PCL60 was seen to increase  $CO_2$  production, but was not seen throughout all flasks.

#### 4.5. CO<sub>2</sub> statistics

A two-way ANOVA was done on the  $T=120 \text{ CO}_2$  evolution data for each biometer flask within all burial matrices using JMP[41]. Based on the F ratio and P>F value, there was no significant difference between means between polymer type, burial environment, and their interactions. Although no significant statistical difference was found, the trends seen from the averaged duplicates will be further discussed. CO<sub>2</sub> experimental setups were done in duplicate due to availability of biometer flasks, but to obtain better statistical results methods should be increased to triplicate or more.

Source	DF	Sum of	Mean	F Ratio
		Squares	Square	
Model	11	170826.13	15529.6	0.7009
Error	12	265875.05	22156.3	<b>Prob</b> > <b>F</b>
C. Total	23	436701.18		0.7182

Table 13. Analysis of variance for carbon dioxide evolution experiments.

Source	Nparm	DF	Sum of Squares	F Ratio	Prob > F
POLYMER	3	3	19457.54	0.2927	0.8299
ENVIRONMENT	2	2	50528.95	1.1403	0.3521
POLYMER*ENVIRONMENT	6	6	100839.65	0.7585	0.6155



Table 14. Effects test for carbon dioxide evolution experiments.

Table 15. Least square means plot for carbon dioxide evolution experiments.









# **Chapter 5 . Results: Microbial Communities**

# 5.1. Bacterial

Polymer		Burial Environment	
Туре			
PCL60	FOOD	SOIL	COFFEE
	Microbacterium sp.	Ochrobactrum sp	unable to sequence
	Gram-positive, non-spore	Gram-negative bacillus,	
	forming rod shaped.	oxidase producing, non-	
	Isolated from soil, insects,	lactose fermenting with	
	human specimens, dairy	environmental origins that	
	productions and more	has been shown to be	
	[41] Has been found to	pathogenic in	
	produce enzymes that	immunocompromised	
	degrade Polystryene (PS)	people. [43]	
	films when buried in soil	Ochrobactrum anthropi	
	[42].	was found in high	
		frequency in activated	
	Rhodococcus globerulus	sludge. [44] Microbes that	
	Present in soils, some	exist within	
	being pathogenic, some	anthropogenic-derived	
	harmless. Well known	waste conditions are often	
	taxa for their	considered for	
	biodegradation and	bioremediation potential	
	bioremediation [43]. Has	[45].	
	been shown to form		
	biofilms along with		
	Aeromonas sp. in study		
	investigating the		
	biodegradability of films		
	(including PCL) in		
	activated sludge (a		
	wastewater treatment		
	process)[44].		
	Chryseobacterium		
	cucumeris		
	This genus is rod shaped,		
	occurring in soil, water,		
	rhizospheres, chicken,		
	fish, and raw milk. Many		
	isolated from plants have		
	part in the plants defense		
	of pathogens[42].		

ST60
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# FOOD **Pedobacter sp.** Rapidly growing genus. One species, *P. heparinus*, produces enzymes that degrade mucopolysaccharides.[46]

# Stenotrophomonas maltophilia

an environmentally ubiquitous bacteria used in bioremediation that has also been found to be an opportunistic pathogen to immuno-compromised persons [47].

SOIL Enterobacter cloacae Gram-negative, rodshaped, facultatively anaerobic bacterium of clinical significance. Research using E. cloacae in microbial fuel cells showed its degradation of cellulose. Highlighting the bacterium's cellulolytic and exoelectrogenic activity [48] E. cloacae strain AKS7 has been shown to develop biofilm over LDPE – leading to enhanced degradation of the plastic [49].

# Enterobacter sp.

Common gram negative bacteria, with some species acting as opportunistic pathogens [50] Found in water, soil, sewage, and intestinal tracts of animals. Capable of nitrogen fixation and are have been isolated from the rhizospheres of crops like wheat and rice. [51]. COFFEE Bacillus subtilis A rhizobacterium species that is non-pathogenic, gram-positive, rod shaped bacteria, that feeds on decayed organic matter that can form biofilms. B. subtilis is also endospore forming [52], [53].

COFFEE
Pseudomonas sp.
Broad genus of gram-
negative bacterium. Many
species are used for
bioremediation of various
anthropogenic pollutants,
due to their metabolic
diversity [57].

Brucella melitensis Soil microbe that causes reproductive losses and illness in some ruminant species [55].

COFFEE

species.[55]

Brucella melitensis

Soil microbe that causes

reproductive losses and

illness in some ruminant

*Pseudomonas moorei* Specific strain (KB4) of this species was shown to degrade paracetamol (an emerging medicinederived pollutant). [56]

SOIL

# Enterobacter cloacae

gram-negative, rodshaped, facultatively anaerobic bacterium of clinical significance. Research using E. cloacae in microbial fuel cells showed its degradation of cellulose. Highlighting the bacterium's cellulolytic and exoelectrogenic activity [48] E. cloacae strain AKS7 has been shown to develop biofilm over LDPE - leading to enhanced degradation of the plastic [49].

#### FOOD Bacillus megaterium Gram positive, endospore forming rod shaped bacteria, aerobic. Large variety of enzymes produced-leading to its definition

produced-leading to its use in bioremediation[54].

# Brucella melitensis

Soil microbe that causes reproductive losses and illness in some ruminant species [55].

# CELL

FOOD *Pseudomonas sp.* Broad genus of gramnegative bacterium. Many species are used for bioremediation of various anthropogenic pollutants, due to their metabolic diverstiy [57].

# Paenibacillus sp.

Rhizobacterium species with at least 16 strains known to be nitrogenfixing. Other bacterial characteristics to enhance plant health and soil health are seen within this species [58]. Paenarthrobacter ureafaciens a gram-positive aerobic bacterium shown to degrade sulfamethazine[59]

SOIL

# Sphingobacterium faecium

A species of Sphingobacterium isolated from cattle feces. The genus is comprised of gram-negative bacilli that are positive for catalase and oxidase. Commonly

# isolated from soil and compost [60].

# \_\_\_\_\_

LDPE	FOOD SOIL	COFFEE
	Rhodococcus sp.	Pedobacter ginsengisoli
	Environmental bacterium	This species was first
	commonly found in soils.	isolated from a ginseng
	Large group of diverse	field soil sample in South
	bacteria that have shown	Korea. Members of the
	great degradation	Pedobacter genus are
	properties. This is due to	obligate aerobes, gram-
	their ability to amass	negative, and positive for
	many catabolic genes [61]	oxidase, catalase, and
		heparinase.[46], [64]
	Pseudoxanthomonas sp.	
	Gram-negative bacteria	
	with some specific sub-	
	species showing potential	
	for the bioremediation of	
	diesel oil [62] and	
	antibiotics in the	
	environment [63]	

Table 16. Bacterial species identified from burial experiment soil samples via 16S rRNA genomic sequencing.



Figure 22. log(CFU/g) of bacterial colonies isolated from polymer surfaces at T=100 of burial experiments 2,3 and 4

5.2. Fungal

Polymer Type		<b>Burial Environment</b>	
PCL60	FOOD	SOIL	COFFEE
	Trichophyton	unidentified	Trichoderma
	Genus of fungi that		Used for industrial
	can be pathogenic		production of
	(ex. ringworm).		cellulase, specifically
	Present in natural		β-glucosidases which
	environments. Some		break down cellulose
	species are wood		into monomers of
	degrading and have		glucose. [66]
	been shown to be		have also been shown
	successful for the		to degrade PLA [67]
	bioremediation of		
	textile azo dyes		
	through fungal		
	biodegradation and		
	bioadsorption [65]		
ST60	FOOD	SOIL	COFFEE
	Deuteromycetes	Trichoderma	Trichoderma
	Grouping of mold	Used for industrial	Used for industrial
	fungi. Broad	production of	production of
	category, but	cellulase, specifically	cellulase, specifically
	different species	$\beta$ -glucosidases which	β-glucosidases which
	strains have been	break down cellulose	break down cellulose
	shown to degrade	into monomers of	into monomers of
	lignin in composts	Trichoderma species	glucose. [66]
	[68]. As well as	have also been shown	Trichoderma species
	Polycyclic aromatic	to degrade PLA [67]	have also been shown
	hydrocarbons or		to degrade PLA [67]
	PAHs which		
	naturally occur in		
	crude oil and gasoline		** unidentified yeast
	[69]		species also present
	Fusarium		
	Common soil fungi		
	that is also found in		
	mycorrhizae of		
	plants. Strain F.		
	proliferatum CF2 has		

COFFI	SOII	been shown to degrade the pesticide allethrin.[70] Fusarium sp were also in a fungal community that led to depredation of polythene[71] FOOD
		1000
**unidentified yeas specie <i>Aspergillus</i> Aspergillus strain were found in abundance on the surface of polymer (PCL, PHB, PLA and PBS) in biodegradation study[72	<i>Fusarium</i> Common soil fungi that is also found in mycorrhizae of plants. Strain <i>F.</i> <i>proliferatum</i> CF2 has been shown to degrade the pesticide allethrin.[70] <i>Fusarium sp</i> were also in a fungal community that led to depredation of polythene[71]	Deuteromycetes Grouping of mold fungi. Broad category, but different species strains have been shown to degrade lignin in composts [68]. As well as Polycyclic aromatic hydrocarbons or PAHs which naturally occur in crude oil and gasoline

CELL

FOOD	SOIL	COFFEE
Trichoderma	Trichoderma	Trichoderma
Used for industrial	Used for industrial	Used for industrial
production of	production of	production of
cellulase, specifically	cellulase, specifically	cellulase, specifically
β-glucosidases which	β-glucosidases which	β-glucosidases which
break down cellulose	break down cellulose	break down cellulose
into monomers of	into monomers of	into monomers of
glucose. [66]	glucose. [66]	glucose. [66]
Trichoderma species	Trichoderma species	Trichoderma species
have also been shown to degrade PLA [67]	have also been shown to degrade PLA [67]	have also been shown to degrade PLA [67]

LDPE	FOOD	SOIL	COFFEE
		Trichophyton	Aspergillus
		Genus of fungi that	Aspergillus strains
		can be pathogenic	were found in
		(ex. ringworm).	abundance on the
		Present in natural	surface of polymers
		environments. Some	(PCL, PHB, PLA,
		species are wood	and PBS) in a
		degrading and have	biodegradation
		been shown to be	study[72]
		successful for the	-
		bioremediation of	
		textile azo dyes	
		through fungal	
		biodegradation and	
		bioadsorption [65]	
T-11. 17 E	· · · · · · · · · · · · · · · · · · ·	· · · · · · · · · · · · · · · · · · ·	1

Table 17. Fungal isolates from burial experiment soil samples identified by colony morphology and microscopy.

Soil samples were collected off the surface of polymers at T=100 from burial experiments 2, 3, and 4. Identification of fungal colonies was based off lactophenol cotton blue staining and colony morphology.



Figure 23. log(CFU/g) of fungal colonies isolated from polymer surfaces at T=100 of burial experiments 2,3 and 4

# **Chapter 6**. Discussion

#### **6.1. Burial Experiments**

As supported by the ANOVA findings, the inclusion of food waste shows a favorable increase in percent weight loss for ST60, PCL60, and partially BC. The soil environment showed higher rates of BC percent weight loss (61%) versus food burial BC (47%) only at T=100. Prior to T=100, the BC buried with food waste outperformed the soil-only environment. Across all novel blended polymers (PCL60, ST60, and BC) the coffee burial environment resulted in lower percent weight loss. Cellulose paper functioned well as a control in all environments, with all three reaching 100% degradation by day 60. However, within the coffee burial environment, there was a drop below 30% degradation which may be attributed to the temperature variations that were previously mentioned.

Utilization of food and soil burial environments both resulted in polymers with higher starch content (ST60 and BC) reaching higher percent weight loss that PCL60. Higher percent inclusion of starch was shown to reflect more degradation, as measured by percent weigh loss, in food and soil burial environments. These findings show that PCL and starch blended polymers should be considered as having potential for use as biodegradable materials. In addition, increasing starch content is shown to increase percent weight loss in burial environments with favorable microbial communities. Also, the inclusion of biochar was not inhibitory to degradation.



Figure 24. Percent weight loss of PCL60 in three burial environments over 100 days.



Figure 25. Percent weight loss of ST60 in three burial environments over 100 days.



Figure 26. Percent weight loss of BC in three burial environments over 100 days.



Figure 27. Percent weight loss of CELL in three burial environments over 100 days.

# 6.1.1. SEM imaging

SEM imaging done at 500X and 2000X magnification showed progression of polymer surface degradation in all three burial environments for days 0, 40 and 100. Comparison between polymer types showed materials with higher starch content ST60 (60% starch) and BC (45% starch) showed more pore formation compared to PCL60 and LDPE.

Looking between burial environments, samples unearthed from coffee show the least amount of surface breakdown for both timepoints 40 and 100. However, the control CELL was seen to fully degrade by day 100. Soil burial environment showed breakdown of CELL and minimal pore formation for BC at day 40. At day 100 in soil there was noticeable degradation for BC, ST60, and PCL60, as well as full CELL degradation. The food burial environment showed the most noticeable surface breakdown for BC, ST60, and PCL60 at both timepoints. Also, in the food burial matrix was the only CELL reached full degradation by day 40.

# 6.2. Microbial Identification

Both bacterial and fungal species identified were chosen due to their majority colonization on either PCA or PDA with streptomycin plates. The microbial source of the plated

cultures was the polymer surfaces at T=100 from the three burial environments. Although the microbial source are the polymer surfaces, it is necessary to note that selection of the best growing colony on plates may favor species that simply grow well on the selected medias. Although general growth medias were selected to best grow all fungal or bacterial species present in soil samples, it cannot be assumed that the colonization of plate growth is perfectly reflective of polymer surface colonization.

# 6.2.1. Bacterial 16S rRNA sequencing

Nine unique species were identified across all polymers in both soil and food environments. Within the coffee environment, four unique species were identified. Food and soil burial environments saw the most diversity of isolates, there was also more bacteria of interest for these burial matrices. These bacteria of interest are the same species or closely related to specific strains that have shown material degradation properties in literature cited. Isolated from PCL60 in the food burial environment, *Rhodococcus globerulus* is a bacterial species that has been shown by other research to form biofilms with *Aermonas sp.* and degrade PCL films. [73]. In soil, *Enterobacter cloacae* was isolated off ST60. Interestingly, a specific strain (AKS7) of *E*, *cloacae* has been shown to enhance the degradation of LDPE via the formation of a biofilm [49]

Beyond bacteria that has been researched specifically for material degradation properties, many species isolated have been cited for having potential bioremediation applications. *Bacillus megaterium, Pseudomonas sp.,Pseudomonas moorei, Ochrobactrum sp., Stenotrophomonas maltophilia, Rhodococcus globerulus,* and *Pseudoxanthomonas sp.* are all bacterium cited within literature for their potential use for bioremediation [39] [43] [45] [52] [54] [55] [60]. Out of the bacterium of interest for bioremediation, only *Pseudomonas sp.* were isolated from coffee environments.

# 6.2.2. Fungal identification

Fundal identification was done based off isolated colony morphology and staining with lactophenol cotton blue (LPCB). Between disposal environments, there were four uniquely identified fungal species or groups in the food burial matrix: *Trichophyton, Deuteromycetes, Fusarium* and *Trichoderma*. Three unique classifications for soil environments: *Trichoderma, Fusarium*, and *Trichophyton*. And in the coffee burial experiment only *Aspergillus* and *Trichoderma* species were found. The only sample without an identified fungal species was PCL60 in soil burial. The LPCB stain for this unidentified species is shown in *figure 28*.

In both soil and coffee environments, *Trichoderma* was isolated off ST60 samples. In literature cited, *Trichoderma* was shown to break down cellulose and PLA. This is of interest because PLA has an ester functional group, just like PCL. Also, cellulose is similar structurally to starch due their shared repeating monomer unit: glucose. Though a *Trichoderma* species was not isolated from the food environments: a *Deuteromycetes* species was which is the group of fungi *Trichoderma* belong to. *Trichoderma* was also seen on CELL samples across all environments.



Figure 28. Lactophenol cotton blue stain of unidentified fungi.



Figure 29. Lactophenol cotton blue stain of Aspergillus.



Figure 30. Lactophenol cotton blue stain of *Trichoderma*.



Figure 31. Lactophenol cotton blue stain of *Trichophyton*.



Figure 32. Lactophenol cotton blue stain of Fusarium.

# 6.3. CO<sub>2</sub> Evolution

 $CO_2$  evolution data across all three burial environments for polymers PCL60, ST60, BC, and CELL were not shown to have statistically significant differences. This may be attributed to only duplicates being used to collect data. Any replication of these methods are recommended to use triplicates for data collection. Although shown to be statistically insignificant, observations from the  $CO_2$  experiment will be discussed further. Comparisons of cumulative  $CO_2$  production of polymers within the three environments are shown in *figures 33, 34*, and *35*.

From the CO<sub>2</sub> data gathered from all burial matrices, BC in the food burial environment/matrix showed the highest evolution of CO<sub>2</sub> (346.9mg) over 120 days amongst all polymer types. Polymers PCL60 and CELL showed higher CO<sub>2</sub> emissions in the food burial environment compared to coffee and soil. However, ST60, which showed higher percent weight loss for 3 out of 4 burial experiments (1,2,4) of the novel blended polymers produced the least amount of CO<sub>2</sub> in the food burial environment.

It is important to note that beyond the time when the ascarite tower is being vented on the biometer flasks and the KOH is taken via syringe from the side arm, the system is closed. This lack of oxygen influx during  $CO_2$  capture may have proven problematic when using a burial matrix with high endogenous microbial activity, like food waste. It has also been cited that soils rich in nitrogen may turn biometer flask systems anaerobic [75].

All  $CO_2$  evolution shown is theoretically sourced only from the breakdown of polymer samples, since the  $CO_2$  evolution of the respective burial matrix is deducted from the experimental  $CO_2$  evolution that involves both the burial matrix and the polymer. When

comparing the  $CO_2$  evolution of polymers in varied burial matrices, nutrient availability could dictate the microbial breakdown of materials. Microbial favored burial environments may favor breaking down the burial material over the polymer material in a closed system like a biometer flask. Conversely, a microbially unfavorable burial environment may lead microbial breakdown of the polymer. Because  $CO_2$  evolution is microbially driven, and we are purely monitoring the output of  $CO_2$  based on pH change and titrations, it is necessary to acknowledge the complexities of microbial activity and consider all potential feedstocks that are present and may be used by both fungi and bacteria to produce  $CO_2$ .



Figure 33. CO<sub>2</sub> evolution of PCL60 polymer samples in food, coffee, and soil burial matrices.



Figure 34. CO2 evolution of ST60 polymer samples in food, coffee, and soil burial matrices.



Figure 35. CO2 evolution of BC polymer samples in food, coffee, and soil burial matrices.



Figure 36. CO2 evolution of CELL polymer samples in food, coffee, and soil burial matrices.

# **Chapter 7**. Future research

To successfully implement biodegradable materials that rely on enzymatic activity for breakdown, the microbiology of disposal environments needs to be understood. However, soilbased disposal environments with or without food waste amendments, are extraordinarily complex. The interactions within the microbial flora, available nutrients, and polymers of interest cannot be oversimplified. Based on findings in this research, future research should address:

- Impact of nitrogen-rich disposal environments on CO<sub>2</sub> evolution within biometer flasks.
- Does inclusion of specific food waste impact polymer degradation? For example, did the starch heavy polymers degrade with the food waste used because it contained starchy food waste (potato peels)?
- Do microscopic fragments of PCL, biochar, or starch remain in the disposal environment, or are they fully utilized and broken down

by microbes? Would these materials be considered fully compostable?

• How would a disposal environment with both food waste and coffee interact with polymer degradation?

Blended polymers made from PCL and starch show promise as potential biodegradable polymers for utilization in food packaging. Inclusion of biochar has been shown to not inhibit biodegradation and should also be considered in further studies of biodegradable packaging. Inclusion of food waste generally led to a more diverse and numerous microbial community and resulted in more desirable percent weight loss trends seen in the food burial environment.

# **Chapter 8 . References**

- T. Ahmed *et al.*, "Biodegradation of plastics: current scenario and future prospects for environmental safety," *Environ. Sci. Pollut. Res.*, vol. 25, no. 8, pp. 7287–7298, Mar. 2018, doi: 10.1007/s11356-018-1234-9.
- [2] P. A. Popescu, M. E. Popa, A. Mitelut, E. E. Tanase, M. Geicu-Cristea, and M. Draghici, "Screening of different methods to establish the biodegradability of packaging materials - a useful tool in environmental risk assessment approach," *Adv. Environ. Sci. Cluj-Napoca*, vol. 9, no. 1, pp. 30–36, Apr. 2017.
- [3] D. Jayanth, P. S. Kumar, G. C. Nayak, J. S. Kumar, S. K. Pal, and R. Rajasekar, "A Review on Biodegradable Polymeric Materials Striving Towards the Attainment of Green Environment," *J. Polym. Environ.*, vol. 26, no. 2, pp. 838–865, Feb. 2018, doi: 10.1007/s10924-017-0985-6.
- [4] L. Van Cauwenberghe and C. R. Janssen, "Microplastics in bivalves cultured for human consumption," *Environ. Pollut.*, vol. 193, pp. 65–70, Oct. 2014, doi: 10.1016/j.envpol.2014.06.010.
- [5] J. C. Philp, A. Bartsev, R. J. Ritchie, M.-A. Baucher, and K. Guy, "Bioplastics science from a policy vantage point," *New Biotechnol.*, vol. 30, no. 6, pp. 635–646, Sep. 2013, doi: 10.1016/j.nbt.2012.11.021.
- [6] F. Gu, J. Guo, W. Zhang, P. A. Summers, and P. Hall, "From waste plastics to industrial raw materials: A life cycle assessment of mechanical plastic recycling practice based on a real-world case study," *Sci. Total Environ.*, vol. 601–602, pp. 1192–1207, Dec. 2017, doi: 10.1016/j.scitotenv.2017.05.278.
- B. Geyer, G. Lorenz, and A. Kandelbauer, "Recycling of poly(ethylene terephthalate) A review focusing on chemical methods," *Express Polym. Lett.*, vol. 10, no. 7, pp. 559–586, Jul. 2016, doi: http://dx.doi.org/10.3144/expresspolymlett.2016.53.
- [8] O. US EPA, "National Overview: Facts and Figures on Materials, Wastes and Recycling," *US EPA*, Oct. 02, 2017. https://www.epa.gov/facts-and-figures-about-materials-waste-and-recycling/national-overview-facts-and-figures-materials (accessed Nov. 02, 2020).
- [9] A. K. Urbanek, W. Rymowicz, and A. M. Mirończuk, "Degradation of plastics and plasticdegrading bacteria in cold marine habitats," *Appl. Microbiol. Biotechnol.*, vol. 102, no. 18, pp. 7669–7678, Sep. 2018, doi: 10.1007/s00253-018-9195-y.
- [10] Z. A. Nur Hanani, Y. H. Roos, and J. P. Kerry, "Use and application of gelatin as potential biodegradable packaging materials for food products," *Int. J. Biol. Macromol.*, vol. 71, pp. 94–102, 2014, doi: 10.1016/j.ijbiomac.2014.04.027.
- [11] L. S. Nair and C. T. Laurencin, "Biodegradable polymers as biomaterials," *Prog. Polym. Sci. Oxf.*, vol. 32, no. 8–9, pp. 762–798, 2007, doi: 10.1016/j.progpolymsci.2007.05.017.
- [12] M. Abedalwafa, F. Wang, L. Wang, and C. Li, "Biodegradable PCL for tissue engineering applications: A review," *Rev. Adv. Mater. Sci.*, vol. 34, no. DECEMBER 2012, pp. 123– 140, 2013.
- [13] M. Rutkowska, M. Jastrzębska, and H. Janik, "Biodegradation of polycaprolactone in sea water," *React. Funct. Polym.*, vol. 38, no. 1, pp. 27–30, 1998, doi: 10.1016/S1381-5148(98)00029-7.
- [14] S. J. Huang, "Biodegradation," in Comprehensive Polymer Science and Supplements, Elsevier, 1989, pp. 597–606. doi: 10.1016/B978-0-08-096701-1.00201-9.
- [15] A. Gijpferich, "Mechanisms of polymer degradation and erosion," 1996.

- [16] B. C. Almeida, P. Figueiredo, and A. T. P. Carvalho, "Polycaprolactone Enzymatic Hydrolysis: A Mechanistic Study," ACS Omega, vol. 4, no. 4, pp. 6769–6774, 2019, doi: 10.1021/acsomega.9b00345.
- [17] G. Kale, T. Kijchavengkul, R. Auras, M. Rubino, S. E. Selke, and S. P. Singh, "Compostability of bioplastic packaging materials: An overview," *Macromolecular Bioscience*, vol. 7, no. 3. John Wiley & Sons, Ltd, pp. 255–277, Mar. 08, 2007. doi: 10.1002/mabi.200600168.
- [18] ACD/ChemSketch.
- [19] M. P. Guaras, V. A. Alvarez, and L. N. Luduena, "Biodegradable nanocomposites based on starch/polycaprolactone/compatibilizer ternary blends reinforced with natural and organomodified montmorillonite," *J. Appl. Polym. Sci.*, vol. 133, no. 44, p. 10, 2016, doi: 10.1002/app.44163.
- [20] F. Zia, K. M. Zia, M. Zuber, S. Kamal, and N. Aslam, "Starch based polyurethanes: A critical review updating recent literature," *Carbohydr. Polym.*, vol. 134, pp. 784–798, Dec. 2015, doi: 10.1016/J.CARBPOL.2015.08.034.
- [21] C. L. Luchese, P. Benelli, J. C. Spada, and I. C. Tessaro, "Impact of the starch source on the physicochemical properties and biodegradability of different starch-based films," *J. Appl. Polym. Sci.*, vol. 135, no. 33, p. 46564, 2018, doi: 10.1002/app.46564.
- [22] "Chapter 4 Global Warming of 1.5 °C." https://www.ipcc.ch/sr15/chapter/chapter-4/ (accessed Jul. 05, 2021).
- [23] R. Janu *et al.*, "Biochar surface functional groups as affected by biomass feedstock, biochar composition and pyrolysis temperature," *Carbon Resour. Convers.*, vol. 4, pp. 36–46, Jan. 2021, doi: 10.1016/j.crcon.2021.01.003.
- [24] G. Agegnehu, A. M. Bass, P. N. Nelson, and M. I. Bird, "Benefits of biochar, compost and biochar–compost for soil quality, maize yield and greenhouse gas emissions in a tropical agricultural soil," *Sci. Total Environ.*, vol. 543, pp. 295–306, Feb. 2016, doi: 10.1016/j.scitotenv.2015.11.054.
- [25] J. S. Singh and C. Singh, Eds., Biochar Applications in Agriculture and Environment Management. Cham: Springer International Publishing, 2020. doi: 10.1007/978-3-030-40997-5.
- [26] T. Aarnio and A. Hämäläinen, "Challenges in packaging waste management in the fast food industry," *Resour. Conserv. Recycl.*, vol. 52, no. 4, pp. 612–621, Feb. 2008, doi: 10.1016/j.resconrec.2007.08.002.
- [27] "Multifunctional food waste fertilizer having the capability of Fusarium-growth inhibition and phosphate solubility: A new horizon of food waste recycle using microorganisms | Elsevier Enhanced Reader." https://reader.elsevier.com/reader/sd/pii/S0956053X1930354X?token=99B17B67BC1C4A 1C1D031C20A5DF3E712BC565E96866402DE34291F68A234D6A491A58664663A662C 6454CC311BC54D0&originRegion=us-east-1&originCreation=20210716145647 (accessed Jul. 16, 2021).
- [28] R. Geyer, J. R. Jambeck, and K. L. Law, "Production, use, and fate of all plastics ever made," *Sci. Adv.*, vol. 3, no. 7, p. e1700782, Jul. 2017, doi: 10.1126/sciadv.1700782.
- [29] C. L. Luchese, P. Benelli, J. C. Spada, and I. C. Tessaro, "Impact of the starch source on the physicochemical properties and biodegradability of different starch-based films," *J. Appl. Polym. Sci.*, vol. 135, no. 33, p. 46564, 2018, doi: 10.1002/app.46564.

- [30] O. US EPA, "Series 835 Fate, Transport and Transformation Test Guidelines," *US EPA*, May 28, 2015. https://www.epa.gov/test-guidelines-pesticides-and-toxic-substances/series-835-fate-transport-and-transformation-test (accessed Jun. 25, 2020).
- [31] "Fate, Transport and Transformation Test Guidelines," United States Environmental Protection Agency, OPPTS 835.3300, Jan. 1998.
- [32] L. Rosario and E. Dell, "AC 2010-593: BIODEGRADABILITY OF PLASTICS TESTING IN AN UNDERGRADUATE MATERIALS LABORATORY COURSE," Rochester Institute of Technology, 2010.
- [33] J. O'Brien, H. Hayder, and C. Peng, "Automated Quantification and Analysis of Cell Counting Procedures Using ImageJ Plugins," *JoVE J. Vis. Exp.*, no. 117, p. e54719, Nov. 2016, doi: 10.3791/54719.
- [34] S. F. Altschul, W. Gish, W. Miller, E. W. Myers, and D. J. Lipman, "Basic local alignment search tool," *J. Mol. Biol.*, vol. 215, no. 3, pp. 403–410, Oct. 1990, doi: 10.1016/S0022-2836(05)80360-2.
- [35] E. Robert C., "MUSCLE: multiple sequence alignment with high accuracy and high throughput," *Nucleic Acids Res.*, vol. 32, no. 5, Mar. 2004, doi: 10.1093/nar/gkh340.
- [36] A. Kuznetsov and C. J. Bollin, "NCBI Genome Workbench: Desktop Software for Comparative Genomics, Visualization, and GenBank Data Submission," in *Multiple Sequence Alignment: Methods and Protocols*, K. Katoh, Ed. New York, NY: Springer US, 2021, pp. 261–295. doi: 10.1007/978-1-0716-1036-7\_16.
- [37] I. L. Pepper and C. P. Gerba, *Environmental Microbiology*, 2nd ed. Elsevier Academic Press, 2005.
- [38] P. Echlin, "Sample Stabilization for Imaging in the SEM," in *Handbook of Sample Preparation for Scanning Electron Microscopy and X-Ray Microanalysis*, P. Echlin, Ed. Boston, MA: Springer US, 2009, pp. 137–183. doi: 10.1007/978-0-387-85731-2\_8.
- [39] D. Adamcová *et al.*, "SEM Analysis and Degradation Behavior of Conventional and Bio-Based Plastics During Composting," *Acta Univ. Agric. Silvic. Mendel. Brun.*, vol. 66, no. 2, pp. 349–356, May 2018, doi: 10.11118/actaun201866020349.
- [40] M. Kopeć, M. Kopeć, A. Baran, and A. Baran, "Effect of the Addition of Biochar and Coffee Grounds on the Biological Properties and Ecotoxicity of Composts," *Waste Biomass Valorization*, vol. 9, no. 8, pp. 1389–1398, Aug. 2018, doi: 10.1007/s12649-017-9916-y.
- [41] P. Goos and D. Meintrup, *Statistics with JMP: Hypothesis Tests, ANOVA and Regression*. New York, UNITED KINGDOM: John Wiley & Sons, Incorporated, 2016. Accessed: Aug. 02, 2021. [Online]. Available: http://ebookcentral.proquest.com/lib/rit/detail.action?docID=4413728
- [42] S. G. Kim, S. S. Giri, S. W. Kim, J. Kwon, S. B. Lee, and S. C. Park, "First Isolation and Characterization of Chryseobacterium cucumeris SKNUCL01, Isolated from Diseased Pond loach (Misgurnus anguillicaudatus) in Korea," *Pathogens*, vol. 9, no. 5, Art. no. 5, May 2020, doi: 10.3390/pathogens9050397.
- [43] N. Jelveh and B. A. Cunha, "Ochrobactrum anthropi bacteremia," *Heart Lung*, vol. 28, no. 2, pp. 145–146, Mar. 1999, doi: 10.1053/hl.1999.v28.a94602.
- [44] "Effect of Different Nanomaterials on the Metabolic Activity and Bacterial Flora of Activated Sludge Medium - Ergön-Can - 2016 - CLEAN – Soil, Air, Water - Wiley Online Library." https://onlinelibrary-wileycom.ezproxy.rit.edu/doi/full/10.1002/clen.201500628#clen201500628-bib-0034 (accessed Jul. 04, 2021).

- [45] V. K. Nguyen, D. D. Nguyen, M.-G. Ha, and H. Y. Kang, "Potential of versatile bacteria isolated from activated sludge for the bioremediation of arsenic and antimony," *J. Water Process Eng.*, vol. 39, p. 101890, Feb. 2021, doi: 10.1016/j.jwpe.2020.101890.
- [46] C. Han *et al.*, "Complete genome sequence of Pedobacter heparinus type strain (HIM 762-3T)," *Stand. Genomic Sci.*, vol. 1, no. 1, pp. 54–62, Jul. 2009, doi: 10.4056/sigs.22138.
- [47] G. Berg, N. Roskot, and K. Smalla, "Genotypic and Phenotypic Relationships between Clinical and Environmental Isolates of Stenotrophomonas maltophilia," J. Clin. Microbiol., vol. 37, no. 11, pp. 3594–3600, Nov. 1999.
- [48] F. Rezaei, D. Xing, R. Wagner, J. M. Regan, T. L. Richard, and B. E. Logan, "Simultaneous Cellulose Degradation and Electricity Production by Enterobacter cloacae in a Microbial Fuel Cell," *Appl. Environ. Microbiol.*, vol. 75, no. 11, pp. 3673–3678, Jun. 2009, doi: 10.1128/AEM.02600-08.
- [49] R. K. Sarker, P. Chakraborty, P. Paul, A. Chatterjee, and P. Tribedi, "Degradation of lowdensity poly ethylene (LDPE) by Enterobacter cloacae AKS7: a potential step towards sustainable environmental remediation," *Arch. Microbiol.*, vol. 202, no. 8, pp. 2117–2125, Oct. 2020, doi: 10.1007/s00203-020-01926-8.
- [50] M. Adeolu, S. Alnajar, S. Naushad, and R. 2016 S. Gupta, "Genome-based phylogeny and taxonomy of the 'Enterobacteriales': proposal for Enterobacterales ord. nov. divided into the families Enterobacteriaceae, Erwiniaceae fam. nov., Pectobacteriaceae fam. nov., Yersiniaceae fam. nov., Hafniaceae fam. nov., Morganellaceae fam. nov., and Budviciaceae fam. nov.," *Int. J. Syst. Evol. Microbiol.*, vol. 66, no. 12, pp. 5575–5599, doi: 10.1099/ijsem.0.001485.
- [51] "Enterobacter -- Britannica Academic." https://academic-ebcom.ezproxy.rit.edu/levels/collegiate/article/Enterobacter/487686 (accessed Jul. 04, 2021).
- [52] Y. Chai, F. Chu, R. Kolter, and R. Losick, "Bistability and biofilm formation in Bacillus subtilis," *Mol. Microbiol.*, vol. 67, no. 2, pp. 254–263, 2008, doi: 10.1111/j.1365-2958.2007.06040.x.
- [53] T. Stein, "Bacillus subtilis antibiotics: structures, syntheses and specific functions," *Mol. Microbiol.*, vol. 56, no. 4, pp. 845–857, 2005, doi: 10.1111/j.1365-2958.2005.04587.x.
- [54] P. Vary *et al.*, "Bacillus megaterium from Simple Soil Bacterium to Industrial Protein Production Host," *Appl. Microbiol. Biotechnol.*, vol. 76, pp. 957–67, Nov. 2007, doi: 10.1007/s00253-007-1089-3.
- [55] A. R. Spickler, "Brucellosis: Brucella melitensis," p. 11.
- [56] J. Żur, D. Wojcieszyńska, K. Hupert-Kocurek, A. Marchlewicz, and U. Guzik, "Paracetamol – toxicity and microbial utilization. Pseudomonas moorei KB4 as a case study for exploring degradation pathway," *Chemosphere*, vol. 206, pp. 192–202, Sep. 2018, doi: 10.1016/j.chemosphere.2018.04.179.
- [57] S. Wasi, S. Tabrez, and M. Ahmad, "Use of Pseudomonas spp. for the bioremediation of environmental pollutants: a review," *Environ. Monit. Assess.*, vol. 185, no. 10, pp. 8147– 8155, Oct. 2013, doi: 10.1007/s10661-013-3163-x.
- [58] L. Seldin, "Paenibacillus, Nitrogen Fixation and Soil Fertility," in *Endospore-forming Soil Bacteria*, N. A. Logan and P. Vos, Eds. Berlin, Heidelberg: Springer, 2011, pp. 287–307. doi: 10.1007/978-3-642-19577-8\_15.
- [59] L. Yu, Y. Wang, X. Su, Y. Fu, F. Ma, and H. Guo, "Biodiversity, isolation and genome analysis of sulfamethazine-degrading bacteria using high-throughput analysis," *Bioprocess Biosyst. Eng.*, vol. 43, no. 8, pp. 1521–1531, Aug. 2020, doi: 10.1007/s00449-020-02345-1.

- [60] A. Lambiase, "The Family Sphingobacteriaceae," in *The Prokaryotes: Other Major Lineages of Bacteria and The Archaea*, E. Rosenberg, E. F. DeLong, S. Lory, E. Stackebrandt, and F. Thompson, Eds. Berlin, Heidelberg: Springer, 2014, pp. 907–914. doi: 10.1007/978-3-642-38954-2\_136.
- [61] V. Gürtler, B. C. Mayall, and R. Seviour, "Can whole genome analysis refine the taxonomy of the genus Rhodococcus?," *FEMS Microbiol. Rev.*, vol. 28, no. 3, pp. 377–403, Jun. 2004, doi: 10.1016/j.femsre.2004.01.001.
- [62] W. Nopcharoenkul, P. Netsakulnee, and O. Pinyakong, "Diesel oil removal by immobilized Pseudoxanthomonas sp. RN402," *Biodegradation*, vol. 24, no. 3, pp. 387–397, Jun. 2013, doi: 10.1007/s10532-012-9596-z.
- [63] Z. Lu, W. Sun, C. Li, X. Ao, C. Yang, and S. Li, "Bioremoval of non-steroidal antiinflammatory drugs by Pseudoxanthomonas sp. DIN-3 isolated from biological activated carbon process," *Water Res.*, vol. 161, pp. 459–472, Sep. 2019, doi: 10.1016/j.watres.2019.05.065.
- [64] L. N. Ten, Q.-M. Liu, W.-T. Im, M. Lee, D.-C. Yang, and S.-T. 2006 Lee, "Pedobacter ginsengisoli sp. nov., a DNase-producing bacterium isolated from soil of a ginseng field in South Korea," *Int. J. Syst. Evol. Microbiol.*, vol. 56, no. 11, pp. 2565–2570, doi: 10.1099/ijs.0.64414-0.
- [65] S. K. Yesiladalı, G. Pekin, H. Bermek, İ. Arslan-Alaton, D. Orhon, and C. Tamerler, "Bioremediation of Textile Azo Dyes by Trichophyton rubrum LSK-27," *World J. Microbiol. Biotechnol.*, vol. 22, no. 10, pp. 1027–1031, Oct. 2006, doi: 10.1007/s11274-005-3207-7.
- [66] P. Tiwari, B. N. Misra, and N. S. Sangwan, "β-Glucosidases from the Fungus Trichoderma: An Efficient Cellulase Machinery in Biotechnological Applications," *BioMed Res. Int.*, vol. 2013, p. 203735, 2013, doi: 10.1155/2013/203735.
- [67] R. Lipsa, N. Tudorachi, R. N. Darie-Nita, L. Oprică, C. Vasile, and A. Chiriac, "Biodegradation of poly(lactic acid) and some of its based systems with Trichoderma viride," *Int. J. Biol. Macromol.*, vol. 88, pp. 515–526, Jul. 2016, doi: 10.1016/j.ijbiomac.2016.04.017.
- [68] B. Kluczek-Turpeinen, M. Tuomela, A. Hatakka, and M. Hofrichter, "Lignin degradation in a compost environment by the deuteromycete Paecilomyces inflatus," *Appl. Microbiol. Biotechnol.*, vol. 61, no. 4, pp. 374–379, May 2003, doi: 10.1007/s00253-003-1272-0.
- [69] A. R. Clemente and L. R. Durrant, "Biodegradation of PAHs in Soil by Two Deuteromycete Fungi," in *Contaminated Soils, Sediments and Water: Science in the Real World Volume 9*, E. J. Calabrese, P. T. Kostecki, and J. Dragun, Eds. Boston, MA: Springer US, 2005, pp. 13–20. doi: 10.1007/0-387-23079-3\_2.
- [70] P. Bhatt *et al.*, "Biodegradation of Allethrin by a Novel Fungus Fusarium proliferatum Strain CF2, Isolated from Contaminated Soils," *Microorganisms*, vol. 8, no. 4, p. 593, 2020, doi: http://dx.doi.org/10.3390/microorganisms8040593.
- [71] M. K. Sangale, M. Shahnawaz, this link will open in a new window Link to external site, and A. B. Ade, "Potential of fungi isolated from the dumping sites mangrove rhizosphere soil to degrade polythene," *Sci. Rep. Nat. Publ. Group*, vol. 9, pp. 1–11, Mar. 2019, doi: http://dx.doi.org/10.1038/s41598-019-41448-y.
- [72] A. S. Al Hosni, J. K. Pittman, and G. D. Robson, "Microbial degradation of four biodegradable polymers in soil and compost demonstrating polycaprolactone as an ideal

compostable plastic," *Waste Manag.*, vol. 97, pp. 105–114, Sep. 2019, doi: 10.1016/j.wasman.2019.07.042.

- [73] M. Swiontek Brzezinska *et al.*, "Biofilm formation during biodegradation of polylactide, poly (3,4 hydroxybutyrate) and poly(ε-caprolactone) in activated sludge," *Int. J. Biol. Macromol.*, vol. 159, pp. 539–546, Sep. 2020, doi: 10.1016/j.ijbiomac.2020.05.107.
- [74] H. M. Alvarez, Ed., *Biology of Rhodococcus*, vol. 16. Berlin, Heidelberg: Springer Berlin Heidelberg, 2010. doi: 10.1007/978-3-642-12937-7.
- [75] H. D. Skipper, I. Arthur G. Wollum, R. F. Turco, and D. C. Wolf, "Microbiological Aspects of Environmental Fate Studies of Pesticides," *Weed Technol.*, vol. 10, no. 1, pp. 174–190, 1996.
- [76] "Polypropylene (PP) Plastic: Types, Properties, Uses & Structure Info." https://omnexus.specialchem.com/selection-guide/polypropylene-pp-plastic (accessed Jul. 18, 2021).
- [77] "PET Plastic (Polyethylene Terephthalate): Uses, Properties & Structure." https://omnexus.specialchem.com/selection-guide/polyethylene-terephthalate-pet-plastic (accessed Jul. 18, 2021).
- [78] "Polylactic Acid or Polylactide, PLA Plastic, Lactic Acid Polymer Guide." https://omnexus.specialchem.com/selection-guide/polylactide-pla-bioplastic (accessed Jul. 20, 2021).

**Appendix- Supplemental figures** 



Figure A. 1. Structure of polypropylene (PP) [76]



Figure A. 2. Structure of repeating monomer unit of polyethylene terephthalate (PET) [77]



Figure A. 3. Structure of repeating monomer unit for polylactic acid (PLA) [78]



Figure A. 4. SEM imaging at 500X of polymers from burial experiments.



Figure A. 6. Image of gel after gel electrophoresis used for 16S rRNA DNA sequencing.



Figure A. 5. Dilution process used for plating and culturing of bacteria and fungi.



Figure A. 7. ImageJ processed image of colonies formed on a plate [33].


0.02	



Figure A. 9. Cladogram of sample source (material-environment) of bacterial species shown in figure A. 8

0.02	