Standardized method for controlled modification of poly (ethylene terephthalate) (PET) crystallinity for assaying PET degrading enzymes

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**A B S T R A C T**

Poly(ethylene terephthalate) (PET) is a polyester plastic, which is widely used, notably as a material for single-use plastic bottles. Its accumulation in the environment now poses a global pollution threat. A number of enzymes are active on PET providing new options for industrial biorecycling of PET materials. The enzyme activity is strongly affected by the degree of PET crystallinity ($X_C$), and the $X_C$ is therefore a relevant factor to consider in enzyme catalyzed PET recycling. Here, we present a new experimental methodology, based on systematic thermal annealing for controlled preparation of PET disks having different $X_C$, to allow systematic quantitative evaluation of the efficiency of PET degrading enzymes at different degrees of PET substrate crystallinity. We discuss the theory of PET crystallinity and compare PET crystallinity data measured by differential scanning calorimetry and attenuated Fourier transform infrared spectroscopy.

- This study introduces a simple method for controlling the crystallinity of PET samples via annealing in a heat block.
- The present methodology is not limited to the analytical methods included in the methods details.

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**A R T I C L E  I N F O**

Method name: Standardized method for controlled modification of poly(ethylene terephthalate) (PET) crystallinity for assaying PET degrading enzymes

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Specifications table

| Subject Area:       | Chemical Engineering |
|---------------------|----------------------|
| More specific subject area: | Enzyme Technology |
| Method name:        | Standardized method for controlled modification of poly(ethylene terephthalate) (PET) crystallinity for assaying PET degrading enzymes |
| Name and reference of original method: | N/A |
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Method details

Background: Polyethylene terephthalate as an enzyme substrate

Poly(ethylene terephthalate) (PET) is a synthetic polyester used in packaging materials and plastic bottles. PET accounts for ~7% of global plastic usage [1], and is a key contributor to the increasing plastic pollution [2]. The polymeric chains of PET consist of repeating units of ethylene glycol and terephthalic acid linked together by ester bonds. These polymer chains are either in an amorphous state (absence of long-range molecular order) or in a crystal structure (highly ordered) [3].

The degree of crystallinity ($X_C$) of a PET sample refers to the fraction of the total polymer chains being in the crystal structure state. The $X_C$ is a result of the PET process history, as crystallization is induced either thermally, at temperatures above the glass transition temperature $T_g$ ($T_g$ is 76°C for amorphous PET), or during certain mechanical operations [4,5]. The amorphous fraction of PET is in fact comprised of two fractions, namely a rigid amorphous fraction (RAF) and a mobile amorphous fraction (MAF); the ratio of each is written as $X_{RAF}$ and $X_{MAF}$, respectively. The RAF is present at the interface between the crystal and the amorphous regions as an immobilized phase caused by the crystallization, and the mobility of RAF is more restrained than the completely amorphous region MAF. The $X_{RAF}$ of a PET sample increases with the $X_C$, while $X_{MAF}$ decreases [6,7].

Although PET is a synthetic polymer, recent data have shown that certain esterolytic enzymes (EC 3.1.1.x), including the PET hydrolase from the bacterium Ideonella sakainese 201-F6, originally specified as a PETase (or IsPETase) [8], now classified as EC 3.1.1.101, are capable of catalyzing hydrolysis of the ester bonds within PET. The activity of these enzymes are, however, limited on PET samples with a high $X_C$ [9–12]. $X_C$ of PET in water bottles ranges from 21–31% [13] making the significance of $X_C$ on PET degrading enzyme activity of critical importance in biobased industrial PET recycling.

Here we present the practical details and the theory behind an experimental methodology that we have developed for preparing PET samples of different degrees of crystallinity, $X_C$, to allow systematic evaluation of the efficiency of PET degrading enzymes in response to the $X_C$ of PET [11]. We introduce annealing involving controlled heating and isothermal cold crystallization to control the $X_C$ of standardized PET sheets (Fig. 1).

Substrate preparation

Preparation of PET material

Amorphous or low crystalline PET films or sheets (e.g. 1 mm thick amorphous PET cat. No. ES30301 from Goodfellow Cambridge Ltd, Huntingdon, UK) are used as starting material to obtain PET materials having different $X_C$ via annealing and isothermal cold crystallization. The PET material is cut into uniform disks using a generic hole punch (Ø 6mm), prior to annealing. Although the starting materials are categorized as amorphous PET materials, it is important to note that the crystallinity of the PET films and sheets differ, and observe that they are not completely amorphous (see Method Validation, below).

Annealing and isothermal cold crystallization of PET

The $X_C$ of the PET discs is systematically altered via annealing followed by cold crystallization as follows: A PET disk is transferred into a 2 mL Eppendorf tube and annealed in a heating block at
115°C for a defined amount of time (minutes); up to three disks can be added per Eppendorf tube, still achieving the same crystallization result as if only one disk is added at a time. The crystallization of the annealed PET sample is quenched by immediately transferring the Eppendorf tube into an ice water bath. It is kept on ice for at least 30 sec to ensure that the crystallized sample has been sufficiently cooled. “Untreated” samples to be used for comparison should be annealed at 85°C for 5 min to remove the enthalpy relaxation caused by the aging of the polymer [14].

Substrate characterization

Quantification of \( X_{\text{MAF}} \) by ATR-FTIR

Attenuated total reflection Fourier-transform infrared spectroscopy (ATR-FTIR) does not require any substrate preparation and can thus be performed directly on the PET disks after annealing. As it is a non-destructive method, the samples analyzed by ATR-FTIR can be used for further analysis, such as differential scanning calorimetry (DSC) or enzymatic treatment.

The peaks of interests described in this paper are at 973 cm\(^{-1}\) and 898 cm\(^{-1}\). These two peaks are associated with the trans (973 cm\(^{-1}\)) and gauche (898 cm\(^{-1}\)) conformer of ethylene glycol [14,15]. While the gauche conformer is only associated with amorphous PET the trans conformer is present
in both the crystalline and amorphous regions [14]. The ratio of the absorbances at these two peaks \( (A_{973}/A_{898}) \) is proportional with the \( X_{MAF} \) of the sample.

**Quantification of the \( X_C \) and \( X_{MAF} \) by DSC**

DSC can be used to quantify the \( X_C \) of a sample annealed for a specific period of time, by analyzing a few samples \((n=3)\) from the batch. A temperature range from 20°C to 270°C is sufficient to determine the thermal features required to estimate \( X_C \) and \( X_{MAF} \) of PET samples. A constant heating rate of 10°C min\(^{-1}\) is recommended, as some thermal features are dependent on the heating rate \([9,11,16]\).

The \( X_C \) of a PET sample, measured by DSC, is then calculated according to Eq. (1):

\[
X_C = \frac{\Delta H_m - \Delta H_{cc}}{\Delta H_m^0} \times 100\%
\]

(1)

Here, \( \Delta H_{cc} \) is the cold crystallization enthalpy (numerical value) of the sample, \( \Delta H_m \) is the heat of melting of the sample, and \( \Delta H_m^0 \) is the heat of melting of a pure crystalline sample. According to literature \( \Delta H_m^0 \) is 140 J g\(^{-1}\) \([17]\).

The \( X_{MAF} \) is calculated according to Eq. (2):

\[
X_{MAF} = \frac{\Delta C_P(m)}{\Delta C_P(a)}
\]

(2)

Where \( \Delta C_P(m) \) is the change in heat capacity of the sample at \( T_g \) and \( \Delta C_P(a) \) is the change in heat capacity at \( T_g \) of a completely amorphous sample. By extrapolation from a linear regression curve of measured \( \Delta C_P(m) \) values and \( X_C \) of PET disk samples we have previously estimated \( \Delta C_P(a) \) of an amorphous PET disk sample to be approximately 0.47 J g\(^{-1}\) K\(^{-1}\) \([11]\).

**Enzyme assay**

The effect of the increased substrate crystallinity on the particular enzyme activity is evaluated on PET disks of different \( X_C \) by measuring the concentration of soluble products formed during enzymatic reaction. Products are measured in terms of bis(2-hydroxyethyl)-terephthalic acid (BHET) equivalents via absorbance measurements of the reaction at 240 nm \([18]\). The rate of the product formation is then plotted against \( X_C \) of the PET disks used.

The enzyme reactions are performed in Eppendorf tubes using 1 disk and a reaction volume of 1 mL in buffer. During the enzymatic treatment, 10μL of the reaction is sampled at various time points. The concentration of soluble product should be normalized with respect to the starting volume \( V_0 \) according to Eq. (3):

\[
C_i = \frac{C_s,i \cdot V_i + \sum_{n=1}^{n} (C_s,n \cdot V_{s,n})}{V_0}
\]

(3)

Where \( C_i \) and \( V_i \) is the normalized concentration of soluble products and the reaction volume at time point \( i \), \( C_s,i \), \( C_s,n \), and \( V_{s,n} \) is the measured concentration of soluble products in the sample measured at time point \( i \) or \( n \) and the sampling volume at time point \( n \).

The product release rate on each PET sample is then calculated from the linear regions of the progress curves. The rate data are then plotted against the \( X_C \) or \( X_{MAF} \) quantified by either DSC or ATR-FTIR.

**Method validation (Case study)**

**Materials and instrumentation**

**PET material**

1 mm thick amorphous PET sheets (Cat. No. ES303010) denoted as PET-S. 250 μm thick amorphous PET film (Cat. No. ES301445) denoted as PET-F, and semi-crystalline PET particles, \( \varnothing < 300 \mu m \) (Cat. No. ES306031), denoted as PET-P were all from Goodfellow (Cambridge Ltd, Huntingdon, UK). The PET-P particles were cast into an amorphous PET sheet by melting 0.5 g of particles in an aluminum dish
the absorbance \(F\) directly increased X PET-C, which observed at 115°C for 5, 6, 7, 8, 9, 10, 12, 15 min (PET-S, PET-F, and for 30 min (PET-S, PET-F, and PET-C) in a heating block. At the specific time points the samples were quenched (cold crystallized), by transferring the annealed samples into ice water for 30 sec. “Untreated” PET samples were annealed at 85°C for 5 min, and subsequently quenched in ice water for 30 sec.

**Isothermal crystallization**

PET-S, PET-F, and PET-C were all cut into disks using a generic hole punch (Ø 6mm). Samples were annealed at 115°C for 5, 6, 7, 8, 9, 10, 12, 15 min (PET-S, PET-F), and for 30 min (PET-S, PET-F, and PET-C) in a heating block. At the specific time points the samples were quenched (cold crystallized), by transferring the annealed samples into ice water for 30 sec. “Untreated” PET samples were annealed at 85°C for 5 min, and subsequently quenched in ice water for 30 sec.

**ATR-FTIR analysis**

ATR-FTIR was performed directly on the PET disks (n=3) using a Spectrum 100 FTIR spectrometer (PerkinElmer, MA, USA). Samples were monitored from 4000 to 650 cm\(^{-1}\) with a resolution of 4 cm\(^{-1}\). Each spectrum consisted of an average of 4 scans.

**DSC analysis**

The \(X_C\) of the PET samples was quantified using DSC measurements (n=3) on a Pyris 1 Calorimeter (Perkin Elmer, Waltham, Massachusetts, USA). A constant heating rate of 10°C min\(^{-1}\) was applied on samples weighing 8.5 ± 0.5 mg for PET-P and PET-S or 6.5 ± 0.5 mg for PET-F between temperatures of 20°C and 270°C. \(X_C\) and \(X_{MAF}\) was calculated according to Eqs. (1) and (2), respectively.

**Enzymes**

\(\text{LCC}_{1\text{CGC}}\) was expressed recombinantly in *E. coli* Shuffle T7 and purified as described in [11] (Expression also works in *E. coli* BL21 (DE3) cells).

**Enzyme activity assay**

1 disk of PET was treated with 150 nM \(\text{LCC}_{1\text{CGC}}\) in 1 mL 0.5M glycine-NaOH buffer pH 9 at 70°C during shaking in a thermomixer (Eppendorf, Hamburg, Germany) for up to 8 h. 10 \(\mu\)L were sampled at specific time points during the reaction. The product formation, at the time points, was quantified by measuring the absorbance at 240 nm using a Synergy HT\textsuperscript{TM} plate reader (BioTek, Winooski, Vermont, United States) as described in [18]. A sample containing 32.6 mg/mL PET-P, corresponding to the average weight of a PET-S disk, was included as a benchmark and assayed under the same conditions as PET-S. All reactions were performed in triplicate.

**Results validation**

The increase in \(X_C\) caused by the annealing followed the characteristic Avrami equation (Fig. 2A), which is associated with crystal growth during annealing [19,20] and the \(X_C\) data obtained per annealing time were higher for the PET-S (sheet) that the PET-F (film) samples (Fig. 2A).

As a supplement to the DSC measurements the ATR-FTIR spectra of the untreated PET samples and extensively annealed PET samples (heated for 30 min at 115°C), respectively, were obtained for the PET-C, PET-F, and PET-S. From the ATR-FTIR spectra, it was observed that the peak at 973 cm\(^{-1}\) increased upon annealing while the absorbance at 898 cm\(^{-1}\) decreased for both the PET-S and the PET-C samples. A non-linear relationship was observed between the \(X_C\) and the \(A_{973}/A_{898}\) ratio of PET-S (linear model \(r^2 = 0.96\)) (Fig. 2C), but a linear correlation, \(r^2 = 0.99\), was observed between the \(X_{MAF}\) and \(A_{973}/A_{898}\) ratio (Fig. 2D). These observations are in agreement with the gauche conformation only being present in \(X_{MAF}\). For PET-S, ATR-FTIR can thereby be used as a non-destructive method to directly quantify the \(X_{MAF}\) of thermally annealed samples before enzymatic treatment.

It is important to note that no changes were observed on the ATR-FTIR spectrum of the PET-F samples. Moreover, both the untreated and annealed PET-F samples produced a broad peak with absorbance maximum at 955 cm\(^{-1}\). We attribute these data to the possible presence of impurities in the original PET film. For this reason, PET-S is recommended as a substrate for the method presented here for quantification of the effect of \(X_C\) on the product release rate of PET degrading enzymes.
Fig. 2. Characterization of PET samples (n=3) annealed at 115°C. (A) Change in the degree of crystallinity (X\textsubscript{C}), measured by DSC of amorphous PET-P (light blue dots) and PET-S (black dots) as a function of annealing time at 115°C. The bold lines represent a nonlinear curve fit to a modified version of the Avrami equation. (B) ATR-FTIR spectrum of PET-C (light red), PET-F (light blue), and PET-S (black lines) which are either untreated (solid lines) or cold crystallized (dashed lines) at 115°C for 30 min. (C) Correlation (non-linear) between the X\textsubscript{C} measured by DSC and the ATR-FTIR ratio \(\frac{A_{973}}{A_{898}}\); the dotted line is included only as an aid for the eye. (D) Linear correlation between the X\textsubscript{MAF} measured by DSC and the ATR-FTIR ratio \(\frac{A_{973}}{A_{898}}\).

From the enzymatic reaction progress curves it was evident that the product release rate catalyzed by LCC\textsubscript{ICCG} on the untreated sample (X\textsubscript{C}=8.2%) was constant throughout the 8 h of the enzymatic treatment (Fig. 3A). A lag phase was evident for the samples with higher X\textsubscript{C} [11]. The substrate X\textsubscript{C}, induced by annealing affected the length of the observed lag phases, which increased with an increasing substrate crystallinity. The samples with a low X\textsubscript{C} (9.3 and 13.8%) thus also displayed lag phases, although the product release rate did not differ once a steady-state reaction rate had been reached. Once a steady-state was reached for the samples with 9.3 or 13.8% X\textsubscript{C} (determined by the linear region of the progress curves displayed in Fig. 3A), it was observed that the product release rate of these samples had reached the same level as the untreated sample. In contrast, the samples with a X\textsubscript{C} of 18.8 % or higher displayed significantly lower product release rates.

The correlation between the product release rate and the X\textsubscript{C} followed a sigmoidal trend (Fig. 3B). This indicates that the negative effect on the LCC\textsubscript{ICCG} catalyzed product release rate caused by increasing substrate crystallinity, reached a critical point at a X\textsubscript{C} between 13.8 and 18.8%, as the reaction rate decreased ~10 fold within this range. A similar observation was made for the *Ideonella sakaiensis* PETase, IsPETase, using PET-F samples crystallized via non-isothermal crystallization from melt [10].
Important considerations

The annealing temperature affects the crystallization rates, $X_c$ at saturation/metastable state, and the crystal thickness [7]. The annealing temperature of 115°C was chosen as $X_c$ saturation was reached within a reasonable amount of time [11]. The enzyme response data (Fig. 3B) underline that it is particularly important to evaluate the reaction rates based on progress curves rather than on end-point measurements. Although PET hydrolase activity as EC 3.1.1.101 has now been defined as producing mono(2-hydroxyethy) terephthalate (MHET) and (ethylene terephthalate)$_n$ from (ethylene terephthalate)$_n$ (www.BRENDA-enzymes.org), it is known that many PET degrading (poly)esterases, including the *I. sakaiensis* PETase, produce some BHET in addition to MHET [18,21] measureable by HPLC. In the present method, we employed direct spectrophotometry of supernatant samples at 240 nm, principally as described in [18]. This quantification of the products as BHET$_{eq}$ is thus mainly a method for enzyme screening or for assessing e.g. the significance of systematic substrate alterations.

A sample containing 32.6 mg of PET-P, corresponding to the average mass of a PET-S disk, was included as a reference. The $X_c$ of PET-P was very high ($37.5 \pm 0.7 \%$), while the total surface area of the PET-P sample was several fold larger than that of a PET-S disc (according to our calculations minimum 6 times larger considering a particle diameter of 300 μm of each PET-P particle). During the initial 1.5 h of incubation, the initial product release rate on the PET-P was significantly higher than the low crystalline PET-S sample. After this point (after an extent of reaction of approximately 4%) the product release rate decreased 14 fold, from 7 ± 0.9 mM BHET$_{eq}$, h$^{-1}$ to 0.5 ± 0.03 mM BHET$_{eq}$, h$^{-1}$. This observed leveling off with time agrees with other studies and appears to be a common trait of enzymatic PET degradation [9,11,12]. Based on recording a large variation in the enzymatic degradability of PET chips from different parts of postconsumer packaging material, it has been suggested [9], that differences in degradability may be due to differences in the distribution of the crystalline microstructures within heterogeneous PET samples. Our data on PET-P versus PET-S (Fig. 3) agree with this interpretation. Taken together, the results underline the importance of annealing the PET disks in a controlled manner to examine the significance of crystallinity on PET hydrolase degradation rates.

Conclusion

In this study, we present a new methodology used to quantitatively evaluate the influence of PET substrate crystallinity, $X_c$, on the action of PET degrading enzymes. The method involves the
preparation of PET samples having different levels $X_C$, by annealing PET samples above their $T_g$ using commercially available PET sheets as starting material.

As a validation of the methodology the effect of the $X_C$, on the product release rate of the gold standard thermostable PET-hydrolyzing enzyme LCCCG was investigated. The findings showed that the enzymatic rate was heavily dependent on the $X_C$. Especially between a $X_C$ of 13.8 and 18.8%, at which the reaction rate decreased $\sim$10 fold. Furthermore, it was observed that the product release rates were not constant during the reaction using PET at a $X_C > 8.2\%$. The data thus emphasize the importance of using progress curves rather than end-point measures when evaluating the catalytic efficiency of PET degrading enzymes on PET, and notably on PET substrate samples of different $X_C$.

It is suggested that the methodology presented in this paper is used as a standardized methodology for addressing the influence of $X_C$ on PET degrading enzymes to allow a direct comparison between different studies.

**Perspectives**

As a final remark, it is important to state that this methodology is not limited to the characterization of the influence of $X_C$ on the enzyme activity. An alternate annealing temperature (above $T_g$) may be used. This will, however, result in different crystallization rates, $X_C$ at saturation/metastable state, and crystal thickness. The thermally annealed PET samples may also be used in other enzyme assays, which are addressing different properties than the product release rate.

**Declaration of Competing Interest**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

**Data availability**

Data will be made available on request.

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