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Conformational fitting of a flexible oligomeric substrate does not explain the enzymatic PET degradation

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Joo et al.1 have recently reported a crystal structure of a polyethylene terephthalate (PET) hydrolyzing enzyme (IsPETase) from Ideonella sakaiensis which has been described able to metabolize PET at 30 °C2. They proposed a PET degradation mechanism solely based on covalent computational docking of an oligomeric substrate—2-hydroxyethyl-(mono-hydroxyethyl terephthalate)4—into the substrate binding cleft of IsPETase without considering the motions and conformations of the PET polymer chain. Here we present a solid-state nuclear magnetic resonance (NMR) analysis of amorphous PET at the degradation temperature used by Joo et al.1, indicating that the highly stiff polymer chain can hardly resemble the suggested docking conformation of 2-HE(MHET)4. In correlation with the PET degradation performance obtained at the same temperature, IsPETase is unlikely to follow the catalytic mechanism proposed by Joo et al.1, which requires simultaneous binding and interaction of all four MHET substrate moieties with the binding site of the enzyme.

While IsPETase can hydrolyze amorphous PET, it showed almost no activity against the crystalline PET polymer2. The OC–CO torsion angle \( \Psi \) in the ethylene glycol (EG) units of amorphous and crystalline PET reveals distinct probability distributions3. According to the 2-HE(MHET)4 docking conformation described by Joo et al.1, a trans \( (t; \Psi \approx 180^\circ) \) content higher than 25% was obtained, considerably higher than the literature value of 14 ± 5% obtained for amorphous PET at ambient temperature3. In our study here, we determined the probability distribution of the OC–CO torsion angle \( \Psi \) in a commercially available amorphous PET material and obtained a trans to gauche \( (\Psi \approx \pm 70^\circ) \) ratio of 9:91 at 30 °C (Fig. 1a), the temperature at which Joo et al.1 performed their PET degradation experiments1. This value is in good agreement with the literature3 but significantly lower than the \( t/g \) ratio obtained in the 2-HE(MHET)4 docking conformation, suggesting that the latter conformation is rarely present in amorphous PET polymer chains. As a consequence, even if the target ester bond in 2-HE(MHET)4 between subsite I and subsite Ia can be correctly accessed by the catalytic triad of the IsPETase (Fig. 2b in Joo et al.1), residues in distal subsites Ib and Ic are unlikely to interact with the other two MHET moieties due to a more biased presence of gauche contents in the PET polymer. In a later publication dealing with the IsPETase structure4, Austin et al.4 have also shown induced fit docking conformations for a PET tetramer, which is equivalent to 2-HE(MHET)4 used by Joo et al.1. For the wildtype IsPETase and a double mutant (S238F/W159H) with increased PET hydrolyzing activity, different energetically most-favored conformations of the PET tetramer were demonstrated by Austin et al.4: a neartrans conformer for the EG unit directly flanking the target ester bond in the wildtype enzyme and a gauche conformer in the double mutant. Nevertheless, in contrast to Joo et al.1, Austin et al.4 did not report on the questionable interactions of distal PET repeating units with surrounding residues in their hypothesis of substrate binding and catalysis.

Moreover, we examined the localized and large-scale cooperative main-chain motions in amorphous PET by magic-angle-spinning (MAS) NMR methods. The \(^{1}H–^{13}C\) dipolar coupling order parameters \( S \) measured using 2D dipolar chemical shift correlation (DIPSHIFT) experiments were used to quantify the site-specific motions on a sub-microsecond timescale to describe the relative rigidity in a value range of 0 to 1.5. Experimental data obtained with crystalline bis(2-hydroxyethyl) terephthalate (BHET) were used to define the rigid limit corresponding to \( S = 1 \). At 30 °C, the dominant gauche conformer in EG units of amorphous PET showed an order parameter of \( S_{34} = 0.96 \), indicating that the transition from gauche to trans conformation was highly restricted (Fig. 1b). Therefore, conformational changes in PET segments in order to perfectly fit the substrate binding cleft of IsPETase as described by Joo et al.1 cannot occur at 30 °C.

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Nevertheless, we determined a more than 1% weight loss of the same amorphous PET film as a result of IsPETase-catalyzed hydrolysis following 24 h of incubation at 30 °C (Fig. 2). Based on the molecular dynamics simulations shown in another study, amino acid residues and backbones involved in the substrate binding cleft of IsPETase are not allowed to move freely to fit any arbitrary rigid conformation of a PET polymer segment. Thus, IsPETase is likely to follow other binding and degradation mechanisms with regard to the PET hydrolysis at 30 °C than those proposed by Joo et al.1.

PET has a glass transition temperature above 70 °C7 which has been suggested as a more favorable reaction condition for enzymatic PET hydrolysis8,9. Yoshida et al.2 verified this by demonstrating a more than 100-fold higher release of UV-absorbing degradation products using LC-cutinase, a thermophilic homologous counterpart of IsPETase, against an amorphous PET sample than using IsPETase after an incubation time of 1 h at their individual optimal reaction conditions2. Similarly, we compared the degradation performance of both enzymes against the PET material used in this study and showed a more than 40-fold higher weight loss obtained by LC-cutinase after an incubation time of 24 h (Fig. 2). These differences in their PET degradation performance were considered to be strongly dependent on the polymer chain mobility and accessibility at different temperatures, as suggested previously also in another publication5. As shown in Fig. 1, we observed slightly larger amplitudes of motions in EG units at 70 °C indicated by the lower order parameters $S_{\text{CH}}$ of 0.86 (trans) and 0.88 (gauche). As a consequence, the transition between trans and gauche conformers was allowed resulting in a $\alpha/\gamma$ ratio of 56:44, a significantly higher value than that obtained at 30 °C. This indicated a transition tendency of PET polymers from a less-ordered state to a more-ordered one as a result of physical aging during the incubation at 70 °C, which is consistent with our recent publication10. Interestingly, the aromatic phenylene rings in amorphous PET exhibited lower order parameters with $S_{\text{CH}} = 0.90$ and 0.78 at 30 °C and 70 °C, respectively, indicating that the phenylene units in PET are more prone to motions compared with the flanking rigid EG units at both temperatures. This is in agreement with a previous publication reporting a phenylene ring flip motion in amorphous PET on a millisecond-to-second timescale11. Therefore, we hypothesize that the weak interactions between the aromatic phenylene units and the surrounding hydrophobic amino acid residues are more likely to facilitate the substrate binding requested for the subsequent enzymatic hydrolysis of PET rather than the perfect accommodation of a certain conformation of a polymer segment, especially at an ambient temperature around 30 °C.

Methods

Solid-state NMR analysis of amorphous PET powder. Amorphous PET films (product no. ES301445) were purchased from Goodfellow Cambridge Ltd. (Huntingdon, UK) and ground in a cryomill in the presence of liquid nitrogen. Particles with diameters of less than 0.25 mm were obtained by sieving and then used in the solid-state NMR analysis. All CP/MAS (cross-polarization magic-angle spinning) NMR experiments were performed at 9.4 T with a Bruker AVANCE III NMR spectrometer equipped with a 4-mm double-resonance MAS probe (Rheinstetten,
A MAS rate of 8 kHz was maintained throughout the experiments. The 207Pb NMR resonance of lead nitrate was used as the thermometer to calibrate the temperature over the sample volume of the in-situ MAS probe at 8 kHz12. The frequency two-pulse phase-modulation (SWf-TPPM) heteronuclear decoupling14 was employed for the 2D DIPSHIFT experiments5. The experiments were conducted with 2512 scans and a relaxation delay of 4 s, with optimized spin-lock pulses to satisfy the standard deviation of triplet measurements.

**Fig. 2** Weight loss of an amorphous PET film (~45 mg) as a result of enzymatic hydrolysis. Incubation with IsPEtase and LC-cutinase was carried out under agitation for 24 h at 30 °C and 70 °C, respectively, i.e., at their optimal reaction conditions for PET degradation. Error bars indicate the standard deviation of triplicate measurements.

Enzymatic hydrolysis of amorphous PET films. A codon-optimized synthetic gene encoding the IsPEtase lacking the N-terminal signal sequence containing 27 residues was ordered from Genescript (Piscataway, USA) and subcloned into the pET-21b vector (Novagen, San Diego, USA). The recombinant IsPEtase containing a C-terminal His-tag was expressed in Escherichia coli Shuffke T7 Express (New England Biolabs GmbH, Frankfurt am Main, Germany). Briefly, E. coli cells were grown to an optical density (OD600 nm of 1, followed by induction in the presence of 0.1 mM IPTG at 16 °C for more than 12 h, and then purified by immobilized metal ion chromatography (IMAC) using TALON metal affinity resin (Takara Bio Europe, Saint-Germain-en-Laye, France) to homogeneity. LC-cutinase was recombinantly expressed in E. coli BL21(DE3) and purified by IMAC to homogeneity using Ni-NTA (Qiagen, Hilden, Germany) as previously described16. 5 µg of IsPEtase and 50 µg of LC-cutinase were used to degrade a piece of amorphous PET film (Goodfellow Cambridge Ltd., 3 × 0.5 cm², ~45 mg) using the optimal degradation conditions for the respective enzymes16. Briefly, IsPEtase required 50 mM Na2HPO4-HCl at pH 7 while LC-cutinase required 1 M K2HPO4/KH2PO4 at pH 8. Degradation was performed by shaking the reaction vials on a thermoshaker T51 (Biometa, Göttingen, Germany) at 1000 rpm for 24 h at 30 °C for IsPEtase and at 70 °C for LC-cutinase. The reaction was stopped by cooling the samples on ice. PET films were washed sequentially with 0.1% aqueous SDS, ethanol and ultrapure water and then dried at 50 °C for 48 h before subjected to gravimetric weight loss determination. Degradation experiments using higher IsPEtase concentrations up to 50 µg enzyme and lower LC-cutinase amount down to 5 µg enzyme were also prepared as control samples which led to significantly lower weight losses of the PET films (data not shown) than using the enzyme amounts mentioned above in main text.

**Data availability**

The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

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Author contributions
R.W. and C.S. designed the experiments; C.S., D.G., and P.B. performed the solid-state NMR analysis and the corresponding evaluation; T.S. and D.B. carried out the cloning, expression and purification of recombinant polyester hydrolases; T.S. and R.W. performed and evaluated the enzymatic PET hydrolysis; R.W., J.M., U.B., and W.Z. supervised the study; R.W. and C.S. wrote the manuscript; U.B., W.Z., D.B., and J.M. edited the manuscript.

Competing interests
The authors declare no competing interests.

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