## Synthesis of biodegradable PGA-PBC-PGA triblock copolymers and closed-loop recycling via a thermal depolymerization strategy

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## Supplementary Information 1-Further explanation on DMA

As the PGA content decreases, the storage modulus of the triblock copolymer gradually decreases. This indicates a gradual decrease in the stiffness of the copolymers. This is consistent with the high PGA modulus and low PBC modulus. In addition, within the low temperature range (<0 °C), as the PGA content decreases, The T<sub>g</sub> of the PBC segments gradually moves towards the high-temperature direction in the loss modulus and (Tan $\delta$ ). However, the T<sub>g</sub> of PGA (40.7 °C by DSC) and the T<sub>m</sub> of PBC (52.3 °C by DSC) are too close. Therefore, within the range of 40 °C -80 °C, there is a transition from glass state to high elastic state in the PGA segments, as well as a transition from high elastic state to viscous flow state in the PBC segments. The T<sub>g</sub> of PGA in DMA is affected by the melting of PBC segments, and the results are not accurate. It is difficult to completely distinguish between the two, which affects the accuracy of the T<sub>g</sub> of PGA. But within -50 °C -0 °C, there is only a phase transition of PBC from glass state to high elastic state, so the T<sub>g</sub> of PBC is accurate. We can observe the T<sub>g</sub> of the PBC chain segments in PGA-PBC-PGA90 in DMA, which is not observed in DSC. It also can be seen that the changes in T<sub>g</sub> are consistent with those measured by DSC from the DMA curves.

Time (min)	15	20	25	30	45	60	90	120	150	180
Conversion (%)	87.9	90.6	93.4	95.7	98.4	98.6	98.7	98.7	98.8	98.8

Table S1	Conversion	rate of GL a	t different times.
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## Table S2 Diffusion coefficients of different characteristic peaks of PGA-PBC-PGA.

Sample	Diffusion coefficient of	Diffusion coefficient of	Diffusion coefficient of peak d		
Sample	peak b	peak c			
PGA-PBC-PGA40	5.91×10 <sup>-12</sup>	5.50×10 <sup>-12</sup>	5.50×10 <sup>-12</sup>		
PGA-PBC-PGA50	4.93×10 <sup>-12</sup>	4.23×10 <sup>-12</sup>	4.23×10 <sup>-12</sup>		
PGA-PBC-PGA60	4.42×10 <sup>-12</sup>	3.82×10 <sup>-12</sup>	3.82×10 <sup>-12</sup>		
PGA-PBC-PGA70	1.65×10 <sup>-12</sup>	1.19×10 <sup>-12</sup>	1.19×10 <sup>-12</sup>		

Table S3 Degradation residual weights of PGA-PBC-PGA in the presence and absence of enzymes.

Samples	Residue weight (%)								
Conditions	PGA	PGA90	PGA80	PGA70	PGA60	PGA50	PGA40	PBC	
Enzymatic	3.71	36.56	58.16	65.36	74.80	83.30	86.62	95.84	
Without enzymatic	13.97	55.98	66.19	73.90	77.32	89.88	91.34	99.04	

## Table S4 The structural composition of PGA-PBC-PGA copolymers before degradation, after enzymatic

degradation, and after non-enzymatic degradation

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Samples Conditions		PGA90	PGA80	PGA70	PGA60	PGA50	PGA40
Mass percentage of GA units in PGA- PBC-PGA (%)	Before degradation	90.0	80.0	70.0	60.0	50.0	40.0
	Enzymatic catalyzed degradation	79.6	70.2	57.0	51.9	42.7	34.2
	Non-enzymatic catalyzed degradation	78.4	69.9	56.7	49.9	42.4	33.3



Fig.S1 <sup>1</sup>H NMR spectra of PBC-OH.











Fig.S7 Fitting curves of diffusion coefficients for different characteristic peaks of PGA-PBC-PGA.



Fig.S8 DMA curves of PGA-PBC-PGA.



**Fig.S9** <sup>1</sup>H NMR spectra of PGA-PBC-PGA60 before degradation, after enzymatic degradation, and after non enzymatic degradation.



Fig.S10 Pictures of degradation of PGA-PBC-PGA at different times.



