

Supplementary Information

Molecular dynamics simulation and non-isothermal crystallization kinetics of polyamide 4 and different bio-based polyamide blends

Yajing Zhang,^{ab} Mingda Wang,^{ab} Liquan Wang,^d Tao Chen^{bd}, Weisheng Feng,^d Tianyi Wang^d and Liming Zhao^{*abc}

^a State Key Laboratory of Bioreactor Engineering, East China University of Science and Technology, Shanghai 200237, China. E-mail: zhaoliming@ecust.edu.cn.

^b Key Laboratory of Biobased Material Engineering, China National Light Industry, East China University of Science and Technology, Shanghai 200237, China.

^c Shanghai Collaborative Innovation Center for Biomanufacturing Technology (SCICBT), Shanghai 200237, China.

^d Shanghai Key Laboratory of Advanced Polymeric Materials, School of Materials Science and Engineering, East China University of Science and Technology, Shanghai 200237, China.

Methods

The simulations were conducted using LJ units, and all quantities are unitless. The units of mass, length, time, and energy are defined by m , r_c , τ , and $k_B T$, respectively. The time unit τ can be formulated by^{S1}

$$\tau = \sqrt{m r_c^2 / k_B T} \quad (\text{S1})$$

where T and k_B are the absolute temperature and Boltzmann constant, respectively. $k_B T$ and m are equal to 1.0 in our simulations. This simulation only considers the nonbonding potential energy (U_{nb}), bond potential energy (U_b), bond-angle-bending potential energy (U_A), and hydrogen-bond interaction potential energy (U_{hb}). The parameters of the beads, nonbonding potential energy, bonds, and angles are shown in Table S1-S4.

Table S1 Parameters for beads.

Bead type	Description	m	ε	σ
A	N atom in the amide group	1.0	1.0	1.0
B	Carbonyl group in the amide group	1.0	1.0	1.0
C	Methylene group in the molecular chain of PA4	1.0	1.0	1.0
D	Methylene group in the molecular chain of PAX	1.0	1.0	1.0
H	H atoms attached to the N atoms	1.0	1.0	1.0

Table S2 Parameters for nonbonding potential energy.

Bead 1	Bead 2	Δ	ε	σ	r_{nb}
A	A	0	1.0	1.0	$4.5 r_c$
A	B	0	1.0	1.0	$4.5 r_c$
A	C	0	1.0	1.0	$4.5 r_c$
A	D	0	1.0	1.0	$4.5 r_c$
A	H	-0.8	1.0	1.0	$4.5 r_c$
B	B	0	1.0	1.0	$4.5 r_c$
B	C	0	1.0	1.0	$4.5 r_c$
B	D	0	1.0	1.0	$4.5 r_c$
B	H	-0.8	1.0	1.0	$4.5 r_c$
C	C	0	1.0	1.0	$4.5 r_c$
C	D	0	1.0	1.0	$4.5 r_c$
C	H	-0.8	1.0	1.0	$4.5 r_c$
D	D	0	1.0	1.0	$4.5 r_c$
D	H	-0.8	1.0	1.0	$4.5 r_c$
H	H	-1.6	1.0	1.0	$4.5 r_c$

Table S3 Parameters for bonds.

Bead 1	Bead 2	K_b	R_b
A	B	$30\varepsilon/\sigma^2$	1.5σ
A	H	$30\varepsilon/\sigma^2$	1.5σ
B	C	$30\varepsilon/\sigma^2$	1.5σ
B	D	$30\varepsilon/\sigma^2$	1.5σ

Table S4 Parameters for angles.

Bead 1	Bead 2	Bead 3	θ_0	K_A
A	B	C	180	10.0
A	B	D	180	10.0
B	C	C	180	10.0
B	D	D	180	10.0
C	A	B	180	10.0
D	A	B	180	10.0