

Supplementary Information

Structure-property relationships of aromatic polyamides and polythioamides: comparative consideration with those of analogous polyesters, polythioesters and polydithioesters

Masayuki Nagasawa, Tatsuya Ishii, Daisuke Abe and Yuji Sasanuma*

Department of Applied Chemistry and Biotechnology, Graduate School and Faculty of Engineering, Chiba University, 1-33 Yayoi-cho, Inage-ku, Chiba 263-8522, Japan

Appendix A Statistical Weight Matrices U_j 's (j , bond number) of PA2T and PTA2T

For the bond numbers, see Fig. 2. Here, the symbol \otimes stands for direct product.

$$U_1 = C_1 \otimes R_3 \quad (\text{A1})$$

$$U_2 = C_3 \otimes R_1 \quad (\text{A2})$$

$$U_3 = C_1 \otimes R_\gamma \quad (\text{A3})$$

$$U_4 = C_6 \otimes R_1 \quad (\text{A4})$$

$$U_5 = C_1 \otimes R_3 \quad (\text{A5})$$

$$U_6 = I_3 \otimes R_3 \quad (\text{A6})$$

$$U_7 = U_f = \begin{bmatrix} \sigma_{ttt} & \sigma_{tttg+} & \sigma_{ttg-} & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & \sigma_{tg+t} & \sigma_{tg+g+} & \sigma_{tg+g-} & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & \sigma_{tg-t} & \sigma_{tg-g+} & \sigma_{tg-g-} \\ \sigma_{g+tt} & \sigma_{g+tg+} & \sigma_{g+tg-} & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & \sigma_{g+g+t} & \sigma_{g+g+g+} & \sigma_{g+g+g-} & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & \sigma_{g+g-t} & \sigma_{g+g-g+} & \sigma_{g+g-g-} \\ \sigma_{g-tt} & \sigma_{g-tg+} & \sigma_{g-tg-} & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & \sigma_{g-g+t} & \sigma_{g-g+g+} & \sigma_{g-g+g-} & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & \sigma_{g-g-t} & \sigma_{g-g-g+} & \sigma_{g-g-g-} \end{bmatrix} \quad (\text{A7})$$

$$U_a = C_3 \otimes I_3 \otimes R_1 \quad (\text{A8})$$

$$U_b = C_3 \otimes S_1 \otimes R_\gamma \quad (\text{A9})$$

$$U_c = C_1 \otimes I_6 \otimes R_1 \quad (\text{A10})$$

$$U_d = C_6 \otimes S_1 \otimes R_3 \quad (\text{A11})$$

$$U_e = C_1 \otimes I_3 \otimes R_3 \quad (\text{A12})$$

and

$$U_n = C_3 \otimes I_3 \otimes R_1 \quad (\text{A13})$$

where

$$C_1 = \begin{bmatrix} 1 \\ 0 \\ 0 \end{bmatrix} \quad (\text{A14})$$

$$C_3 = \begin{bmatrix} 1 \\ 1 \\ 1 \end{bmatrix} \quad (\text{A15})$$

$$C_6 = \begin{bmatrix} 1 \\ 1 \\ 1 \\ 1 \\ 1 \\ 1 \end{bmatrix} \quad (\text{A16})$$

$$R_1 = \begin{bmatrix} 1 & 0 & 0 \end{bmatrix} \quad (\text{A17})$$

$$R_3 = \begin{bmatrix} 1 & 1 & 1 \end{bmatrix} \quad (\text{A18})$$

$$R_\gamma = \begin{bmatrix} \gamma_{t-t} & \gamma_{(t-c)^+} & \gamma_{(t-c)^-} & \gamma_{(c-t)^+} & \gamma_{(c-t)^-} & \gamma_{c-c} \end{bmatrix} \quad (\text{A19})$$

$$S_1 = \begin{bmatrix} 1 & 0 & 0 \\ 0 & 0 & 0 \\ 0 & 0 & 0 \end{bmatrix} \quad (\text{A20})$$

and I_3 and I_6 are the identity matrices of sizes 3 and 6 respectively, with $\gamma_\xi = \exp(-\Delta G_\xi/RT)$ (ξ : orientation around bond 3 or b) and $\sigma_k = \exp(-\Delta G_k/RT)$ (k : conformations in bonds 5, 6, and 7 or bonds d, e, and f).

Appendix B Statistical Weight Matrices U_j 's (j , bond number) of PA3T ans PTA3T

The U_1 to U_6 matrices are the same as those of PA2T ans PTA2T.

$$U_7 = I_3 \otimes I_3 \otimes R_3 \quad (\text{A21})$$

$$U_a = C_3 \otimes I_3 \otimes I_3 \otimes R_1 \quad (\text{A22})$$

$$U_b = C_3 \otimes I_3 \otimes S_1 \otimes R_\gamma \quad (\text{A23})$$

$$U_c = C_1 \otimes S_1 \otimes I_6 \otimes R_1 \quad (\text{A24})$$

$$U_d = C_1 \otimes I_6 \otimes S_1 \otimes R_3 \quad (\text{A25})$$

$$U_e = C_6 \otimes S_1 \otimes I_3 \otimes R_3 \quad (\text{A26})$$

$$U_f = C_1 \otimes I_3 \otimes I_3 \otimes R_3 \quad (\text{A27})$$

and

$$U_n = C_3 \otimes I_3 \otimes I_3 \otimes R_1 \quad (\text{A28})$$

The U_8 and U_g matrices are defined on the next page.

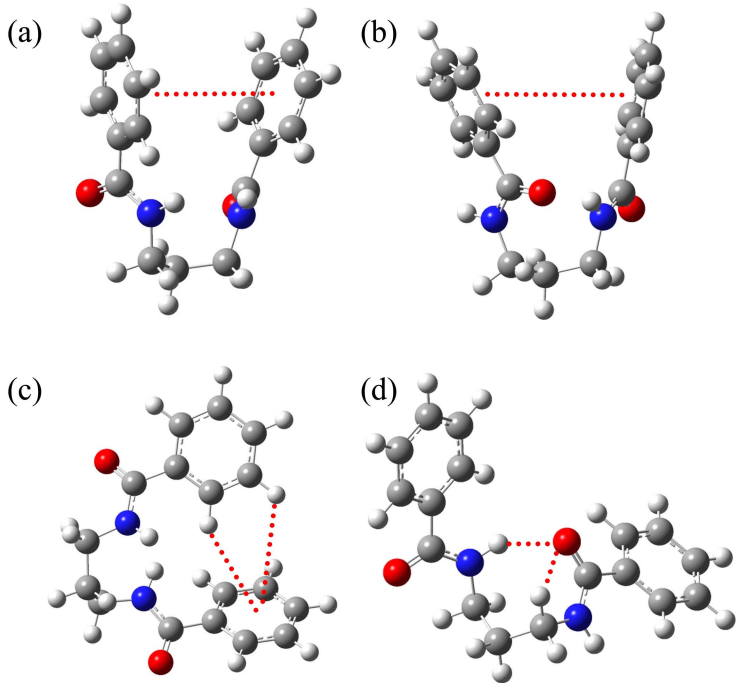


Fig. S1 Intramolecular interactions of 3DBA: π/π of (a) g+g+g-g- and (b) g+g+g-g+; C-H... π of (c) tg+g-g-; NH...O=C of g+g+g+g-.

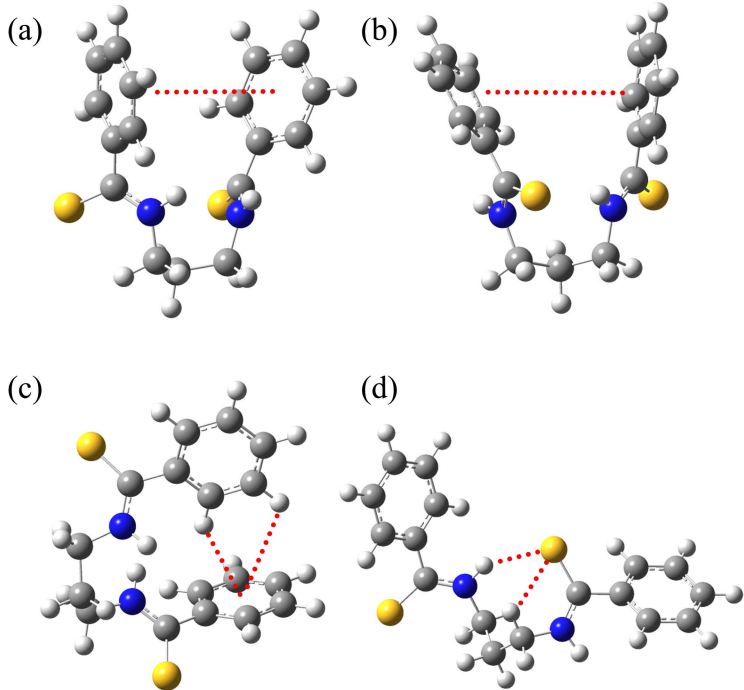


Fig. S2 Intramolecular interactions of 3DBTA: π/π of (a) g+g+g-g- and (b) g+g+g-g+; C-H... π of (c) tg+g-g-; NH...S=C of g+g+g-g-.

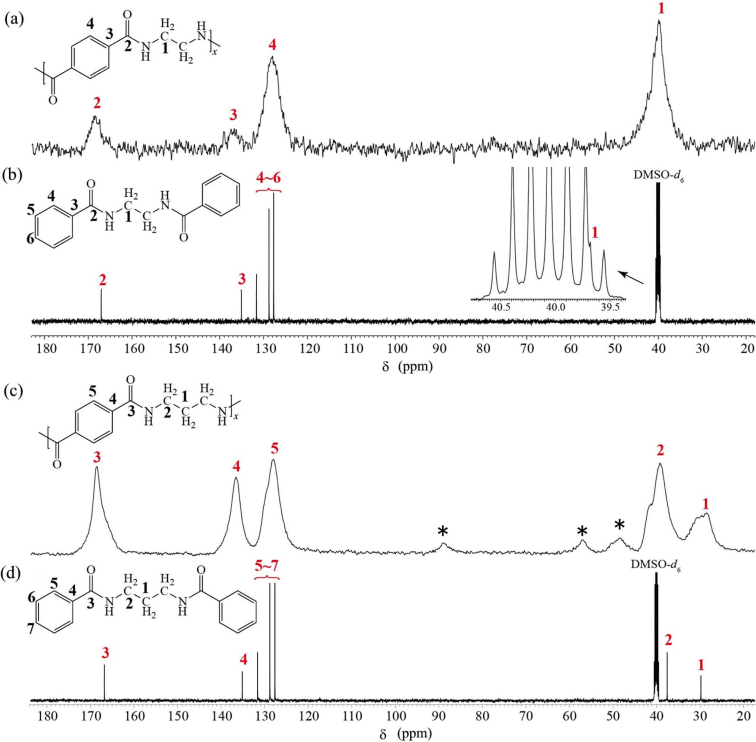


Fig. S3 Solid-state ^{13}C NMR spectra observed from (a) PA2T and (c) PA3T, compared with solution spectra from (b) 2DBA and (d) 3DBA dissolved in $\text{DMSO}-d_6$. The peaks were assigned as indicated, and the asterisk represents the spinning sideband.

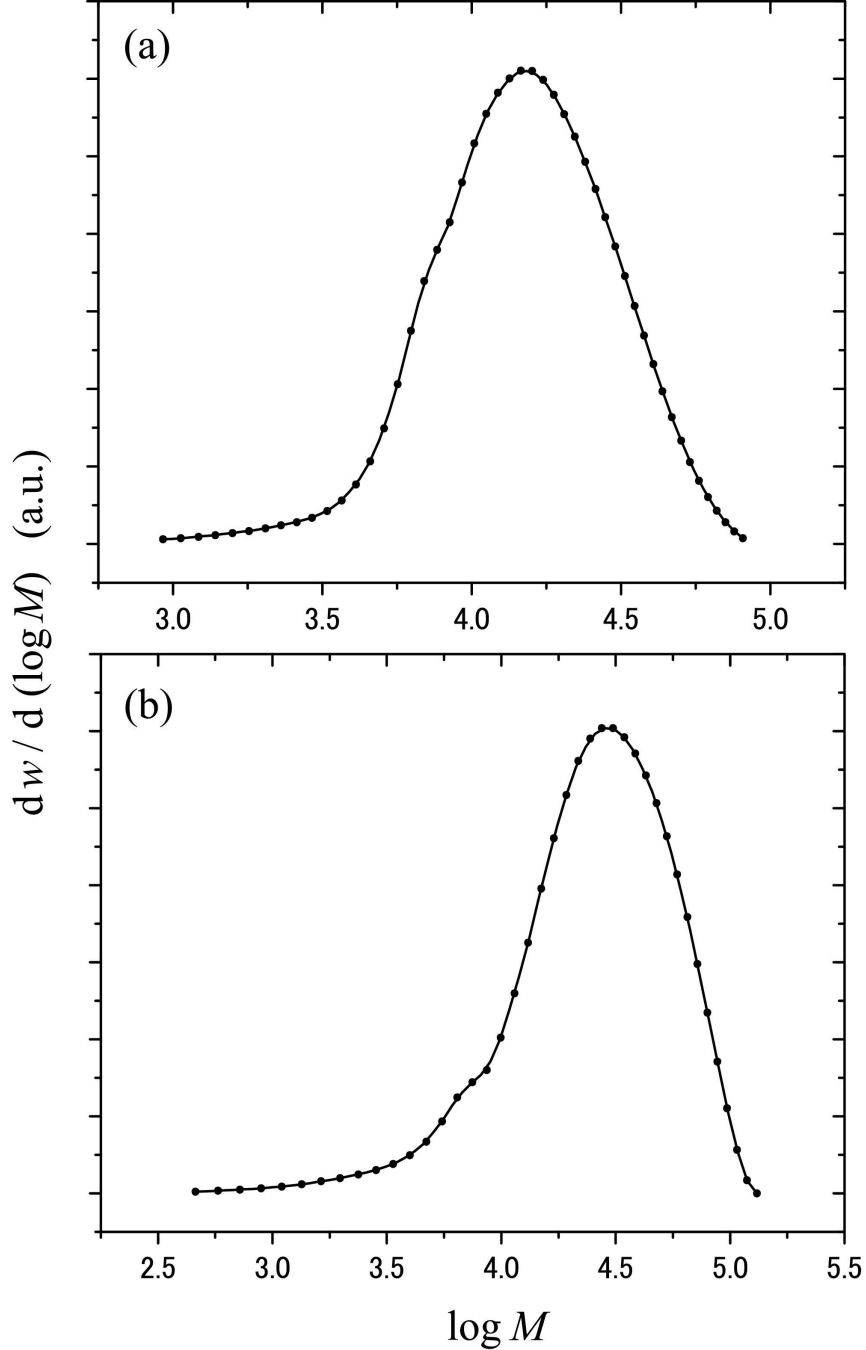


Fig. S4 Molecular weight distributions of (a) PTA2T and (b) PTA3T.

Table S1 Geometrical parameters of PA2T, used in the refined RIS calculations

conformation						
bond ^a			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond a (c)						
t	t	t-t	1.352	121.5	116.8	1.5
g ⁺			1.362	122.3	117.8	-1.9
g ⁻			1.362	122.3	117.8	1.9
t	t	(t-c) ⁺	1.352	121.5	116.7	0.3
g ⁺			1.361	122.4	117.9	-0.7
g ⁻			1.361	122.4	117.9	-0.7
t	t	(t-c) ⁻	1.352	121.5	116.7	-0.3
g ⁺			1.361	122.4	117.9	0.7
g ⁻			1.361	122.4	117.9	0.7
t	t	(c-t) ⁺				
g ⁺			1.361	122.4	119.1	0.2
g ⁻			1.361	122.4	119.1	0.2
t	t	(c-t) ⁻				
g ⁺			1.361	122.4	119.1	-0.2
g ⁻			1.361	122.4	119.1	-0.2
t	t	c-c	1.352	121.5	116.7	1.0
g ⁺			1.362	122.3	119.1	-2.3
g ⁻			1.362	122.3	119.1	2.3
bond b						
t	t-t	t	5.811	116.8	116.8	0.0
	(t-c) ⁺		5.811	116.7	116.7	53.6
	(t-c) ⁻		5.811	116.7	116.7	-53.6
	(c-t) ⁺		5.809	116.8	116.8	123.0
	(c-t) ⁻		5.809	116.8	116.8	-123.0
	c-c		5.809	116.7	116.7	180.0
bond d (f)						
t	t	t				
	g ⁺		1.453	123.2	112.3	87.7
	g ⁻		1.453	123.2	112.3	-87.7
	t	g ⁺	1.457	122.1	111.1	3.3
	g ⁺		1.456	122.9	113.7	94.1
	g ⁻		1.462	123.2	114.6	-93.8
	t	g ⁻	1.457	122.1	111.1	-3.3
	g ⁺		1.462	123.2	114.6	93.8
	g ⁻		1.456	122.9	113.7	-94.1

conformation						
bond ^a			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond e						
t	t	t				
	g ⁺					
	g ⁻					
g ⁺	t					
	g ⁺		1.525		111.1	-112.7
	g ⁻					
g ⁻	t					
	g ⁺		1.525		111.1	112.7
	g ⁻					
t	t	g ⁺				
	g ⁺					
	g ⁻		1.525		114.6	-112.7
g ⁺	t		1.538		112.3	2.5
	g ⁺		1.534		113.7	116.8
	g ⁻		1.532		113.6	-119.0
g ⁻	t		1.538		112.1	0.5
	g ⁺		1.533		113.7	116.7
	g ⁻		1.533		113.0	-116.7
t	t	g ⁻				
	g ⁺		1.525		114.6	112.7
	g ⁻					
g ⁺	t		1.538		112.1	0.5
	g ⁺		1.533		113.0	116.7
	g ⁻		1.533		113.7	-116.7
g ⁻	t		1.538		112.3	-2.5
	g ⁺		1.532		113.6	119.0
	g ⁻		1.534		113.7	-116.8

^a j denotes the current bond. The blank line represents that the conformation is inexistent. ^bLength of bond j in Å. ^cAngle formed between bonds $j-1$ and j in deg. ^dAngle formed between bonds j and $j+1$ in deg. ^eDihedral angle of bond j in deg.

Table S2 Geometrical parameters of PTA2T, used in the refined RIS calculations

conformation						
bond ^a			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond a (c)						
t	t	t-t	1.336	124.3	115.4	1.8
g ⁺			1.342	126.2	114.8	3.5
g ⁻			1.342	126.2	114.8	-3.5
t	t	(t-c) ⁺	1.336	124.6	115.5	2.8
g ⁺			1.341	126.4	114.4	3.0
g ⁻			1.341	126.4	114.4	3.0
	t	(t-c) ⁻	1.336	124.6	115.5	-2.8
g ⁺			1.341	126.4	114.4	-3.0
g ⁻			1.341	126.4	114.4	-3.0
t	t	(c-t) ⁺	1.335	124.6	115.6	-2.4
g ⁺			1.341	126.3	114.8	-2.9
g ⁻			1.341	126.3	114.8	-2.9
t	t	(c-t) ⁻	1.335	124.6	115.6	2.4
g ⁺			1.341	126.3	114.8	2.9
g ⁻			1.341	126.3	114.8	2.9
t	t	c-c	1.337	124.3	115.8	1.7
g ⁺			1.342	126.2	114.9	3.7
g ⁻			1.342	126.2	114.9	-3.7
bond b						
t	t-t	t	5.793	115.4	115.4	0.0
	(t-c) ⁺		5.794	115.5	115.5	79.6
	(t-c) ⁻		5.794	115.5	115.5	-79.6
	(c-t) ⁺		5.792	115.6	115.6	98.0
	(c-t) ⁻		5.792	115.6	115.6	-98.0
	c-c		5.792	115.8	115.8	180.0
bond d (f)						
t	t	t	1.456	125.3	109.7	-9.3
	g ⁺		1.454	126.2	111.5	89.9
	g ⁻		1.454	126.2	111.5	-89.9
	t	g ⁺	1.458	124.6	111.8	0.2
	g ⁺		1.457	126.2	113.6	93.1
	g ⁻		1.461	126.8	114.2	-81.8
	t	g ⁻	1.458	124.6	111.8	-0.2
	g ⁺		1.461	126.8	114.2	81.8
	g ⁻		1.457	126.2	113.6	-93.1

conformation						
bond ^a			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond e						
t	t	t	1.526		109.6	0.4
	g ⁺		1.520		111.7	111.5
	g ⁻		1.520		111.7	-111.5
g ⁺	t		1.531		109.7	-1.5
	g ⁺		1.530		112.1	113.8
	g ⁻		1.525		111.8	-112.1
g ⁻	t		1.531		109.7	1.5
	g ⁺		1.525		111.8	112.1
	g ⁻		1.530		112.1	-113.8
t	t	g ⁺	1.531		111.6	-1.5
	g ⁺		1.530		113.5	113.8
	g ⁻		1.525		114.2	-112.1
g ⁺	t		1.539		111.6	9.8
	g ⁺		1.533		113.6	116.8
	g ⁻					
g ⁻	t		1.538		111.5	-0.7
	g ⁺					
	g ⁻					
t	t	g ⁻	1.531		111.6	1.5
	g ⁺		1.525		114.2	112.1
	g ⁻		1.530		113.5	-113.8
g ⁺	t		1.538		111.5	0.7
	g ⁺					
	g ⁻					
g ⁻	t		1.539		111.6	-9.8
	g ⁺					
	g ⁻		1.533		113.6	-116.8

^a j denotes the current bond. The blank line represents that the conformation is inexistent. ^bLength of bond j in Å. ^cAngle formed between bonds $j-1$ and j in deg. ^dAngle formed between bonds j and $j+1$ in deg. ^eDihedral angle of bond j in deg.

Table S3 Geometrical parameters of PA3T, used in the refined RIS calculations

conformation						
bond ^a			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond a (c)						
t	t	t-t				
g ⁺			1.360	123.4	117.6	-0.6
g ⁻			1.360	123.4	117.6	0.6
t	t	(t-c) ⁺				
g ⁺			1.360	122.3	117.9	-1.8
g ⁻			1.360	122.3	117.9	-1.8
t	t	(t-c) ⁻				
g ⁺			1.360	122.3	117.9	1.8
g ⁻			1.360	122.3	117.9	1.8
t	t	(c-t) ⁺				
g ⁺			1.359	122.7	119.3	-1.6
g ⁻			1.359	122.7	119.3	-1.6
t	t	(c-t) ⁻				
g ⁺			1.359	122.7	119.3	1.6
g ⁻			1.359	122.7	119.3	1.6
t	t	c-c				
g ⁺			1.359	122.8	119.1	-1.6
g ⁻			1.359	122.8	119.1	1.6
bond b						
t	t-t	t	5.825	118.1	118.1	0.0
	(t-c) ⁺		5.824	118.0	118.0	46.9
	(t-c) ⁻		5.824	118.0	118.0	-46.9
	(c-t) ⁺		5.823	119.4	119.4	129.2
	(c-t) ⁻		5.823	119.4	119.4	-129.2
	c-c		5.822	119.2	119.2	180.0
bond d (g)						
t	t	t	1.458	122.6	110.9	-19.0
	g ⁺		1.457	123.3	113.1	88.8
	g ⁻		1.457	123.3	113.1	-88.8
t	t	g ⁺				
g ⁺			1.456	123.8	113.6	76.9
g ⁻			1.461	123.5	113.3	-34.8
t	t	g ⁻				
g ⁺			1.461	123.5	113.3	34.8
g ⁻			1.456	123.8	113.6	-76.9

conformation						
bond ^a			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond e (f)						
t	t	t	1.526	110.9	111.1	-1.0
	g ⁺		1.530	112.6	113.1	113.1
	g ⁻		1.530	112.6	113.1	-113.1
g ⁺	t		1.532	113.1	111.4	1.1
	g ⁺		1.532	114.0	113.2	114.1
	g ⁻		1.531	113.8	113.3	-113.7
g ⁻	t		1.532	113.1	111.4	-1.1
	g ⁺		1.531	113.8	113.3	113.7
	g ⁻		1.532	114.0	113.2	-114.1
t	t	g ⁺	1.527	111.1	113.3	-2.0
	g ⁺		1.529	112.3	114.8	114.2
	g ⁻		1.528	113.2	115.7	-109.4
g ⁺	t		1.532	112.7	113.2	0.5
	g ⁺		1.533	113.6	114.9	124.4
	g ⁻		1.528	113.3	115.7	-113.3
g ⁻	t		1.532	113.0	113.0	-2.6
	g ⁺		1.530	112.5	114.8	114.5
	g ⁻		1.534	115.1	115.7	-105.4
t	t	g ⁻	1.527	111.1	113.3	2.0
	g ⁺		1.528	113.2	115.7	109.4
	g ⁻		1.529	112.3	114.8	-114.2
g ⁺	t		1.532	113.0	113.0	2.6
	g ⁺		1.534	115.1	115.7	105.4
	g ⁻		1.530	112.5	114.8	-114.5
g ⁻	t		1.532	112.7	113.2	-0.5
	g ⁺		1.528	113.3	115.7	113.3
	g ⁻		1.533	113.6	114.9	-124.4

^a j denotes the current bond. The blank line represents that the conformation is inexistent. ^b Length of bond j in Å. ^c Angle formed between bonds $j-1$ and j in deg. ^d Angle formed between bonds j and $j+1$ in deg. ^e Dihedral angle of bond j in deg.

Table S4 Geometrical parameters of PTA3T, used in the refined RIS calculations

conformation						
bond ^a			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond a (c)						
t	t	t-t	1.338	125.3	115.1	2.4
g ⁺			1.339	126.6	114.6	2.1
g ⁻			1.339	126.6	114.6	-2.1
t	t	(t-c) ⁺	1.338	125.3	115.2	2.5
g ⁺			1.339	126.6	114.5	3.9
g ⁻			1.339	126.6	114.5	3.9
t	t	(t-c) ⁻	1.338	125.3	115.2	-2.5
g ⁺			1.339	126.6	114.5	-3.9
g ⁻			1.339	126.6	114.5	-3.9
t	t	(c-t) ⁺	1.338	125.3	115.3	-2.0
g ⁺			1.339	126.6	114.6	-1.8
g ⁻			1.339	126.6	114.6	-1.8
t	t	(c-t) ⁻	1.338	125.3	115.3	2.0
g ⁺			1.339	126.6	114.6	1.8
g ⁻			1.339	126.6	114.6	1.8
t	t	c-c	1.338	125.4	115.1	2.2
g ⁺			1.339	126.6	114.5	1.7
g ⁻			1.339	126.6	114.5	-1.7
bond b						
t	t-t	t	5.809	115.1	115.1	0.0
	(t-c) ⁺		5.809	115.2	115.2	75.8
	(t-c) ⁻		5.809	115.2	115.2	-75.8
	(c-t) ⁺		5.808	115.3	115.3	102.5
	(c-t) ⁻		5.808	115.3	115.3	-102.5
	c-c		5.808	115.1	115.1	180.0
bond d (g)						
t	t	t	1.458	125.4	110.5	6.2
	g ⁺		1.458	126.4	112.3	88.9
	g ⁻		1.458	126.4	112.3	-88.9
	t	g ⁺	1.461	125.6	111.6	-9.9
	g ⁺		1.461	126.1	114.0	112.7
	g ⁻		1.460	127.4	112.5	-59.0
	t	g ⁻	1.461	125.6	111.6	9.9
	g ⁺		1.460	127.4	112.5	59.0
	g ⁻		1.461	126.1	114.0	-112.7

conformation						
bond ^a			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond e (f)						
t	t	t	1.525	110.5	110.7	-0.6
	g ⁺		1.525	111.6	113.1	109.8
	g ⁻		1.525	111.6	113.1	-109.8
g ⁺	t		1.532	112.6	110.8	-0.3
	g ⁺		1.532	114.0	113.1	112.7
	g ⁻		1.530	112.5	113.0	-113.0
g ⁻	t		1.532	112.6	110.8	0.3
	g ⁺		1.530	112.5	113.0	113.0
	g ⁻		1.532	114.0	113.1	-112.7
t	t	g ⁺	1.528	110.4	113.1	0.4
	g ⁺		1.528	111.4	115.1	110.7
	g ⁻		1.535	113.4	116.1	-91.1
g ⁺	t		1.535	112.3	113.0	-8.2
	g ⁺		1.534	113.8	115.3	115.2
	g ⁻		1.535	113.4	116.1	-91.1
g ⁻	t		1.533	112.4	113.2	-1.0
	g ⁺					
	g ⁻		1.533	115.3	116.0	-110.4
t	t	g ⁻	1.528	110.4	113.1	-0.4
	g ⁺		1.535	113.4	116.1	91.1
	g ⁻		1.528	111.4	115.1	-110.7
g ⁺	t		1.533	112.4	113.2	1.0
	g ⁺		1.533	115.3	116.0	110.4
	g ⁻					
g ⁻	t		1.535	112.3	113.0	8.2
	g ⁺		1.535	113.4	116.1	91.1
	g ⁻		1.534	113.8	115.3	-115.2

^a j denotes the current bond. The blank line represents that the conformation is inexistent. ^b Length of bond j in Å. ^c Angle formed between bonds $j-1$ and j in deg. ^d Angle formed between bonds j and $j+1$ in deg. ^e Dihedral angle of bond j in deg.

Table S5 Geometrical parameters of P3TS₂, used in the refined RIS calculations

conformation			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond a (c)						
t	t	trans	1.807	98.4	117.9	0.0
g ⁺			1.809	99.0	117.7	0.2
g ⁻			1.809	99.0	117.7	-0.2
t	t	cis	1.808	98.5	118.8	0.0
g ⁺			1.804	99.6	118.6	-0.1
g ⁻			1.804	99.6	118.6	0.1
bond b						
t	trans	t	5.798	117.9	117.9	0.0
	cis		5.787	118.8	118.8	180.0
bond d (g)						
t	t	t	1.838	98.4	109.7	0.0
	g ⁺		1.835	98.9	113.5	97.4
	g ⁻		1.835	98.9	113.5	-97.4
	t	g ⁺	1.841	98.6	111.4	-12.6
	g ⁺		1.838	98.7	114.7	96.9
	g ⁻		1.838	100.0	115.1	-81.1
	t	g ⁻	1.841	98.6	111.4	12.6
	g ⁺		1.838	100.0	115.1	81.1
	g ⁻		1.838	98.7	114.7	-96.9

conformation			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond e (f)						
t	t	t	1.532	109.7	110.3	0.0
	g ⁺		1.534	111.4	113.3	111.8
	g ⁻		1.534	111.4	113.3	-111.8
σg ⁺	t		1.532	113.5	110.5	-1.6
	g ⁺		1.533	114.7	113.1	113.4
	g ⁻		1.535	115.1	113.6	-114.2
σg ⁻	t		1.532	113.5	110.5	1.6
	g ⁺		1.535	115.1	113.6	114.2
	g ⁻		1.533	114.7	113.1	-113.4
t	t	σg ⁺	1.530	109.6	113.1	-0.1
	g ⁺		1.532	111.0	116.0	114.2
	g ⁻		1.533	115.9	116.5	-92.7
σg ⁺	t		1.528	113.6	113.4	6.2
	g ⁺		1.533	114.0	115.2	107.4
	g ⁻		1.534	117.1	116.8	-105.2
σg ⁻	t		1.530	113.3	113.2	-0.3
	g ⁺		1.533	114.0	115.2	107.4
	g ⁻		1.536	116.1	116.8	-96.4
t	t	σg ⁻	1.530	109.6	113.1	0.1
	g ⁺		1.533	115.9	116.5	92.7
	g ⁻		1.532	111.0	116.0	-114.2
σg ⁺	t		1.530	113.3	113.2	0.3
	g ⁺		1.536	116.1	116.8	96.4
	g ⁻		1.533	114.0	115.2	-107.4
σg ⁻	t		1.528	113.6	113.4	-6.2
	g ⁺		1.534	117.1	116.8	105.2
	g ⁻		1.533	114.0	115.2	-107.4

^a j denotes the current bond. The blank line represents that the conformation is inexistent. ^bLength of bond j in Å. ^cAngle formed between bonds $j-1$ and j in deg. ^dAngle formed between bonds j and $j+1$ in deg. ^eDihedral angle of bond j in deg.

Table S6 Geometrical parameters of P3TS₄, used in the refined RIS calculations

conformation						
bond ^a			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond a (c)						
t	t	t-t	1.758	103.9	113.0	3.1
g ⁺			1.770	104.7	112.7	2.1
g ⁻			1.770	104.7	112.7	-2.1
t	t	(t-c) ⁺	1.759	103.9	113.2	4.0
g ⁺			1.772	104.2	112.1	5.9
g ⁻			1.772	104.2	112.1	5.9
t	t	(t-c) ⁻	1.759	103.9	113.2	-4.0
g ⁺			1.772	104.2	112.1	-5.9
g ⁻			1.772	104.2	112.1	-5.9
t	t	(c-t) ⁺	1.758	103.9	113.4	-3.1
g ⁺			1.771	104.7	113.0	-5.5
g ⁻			1.771	104.7	113.0	-5.5
t	t	(c-t) ⁻	1.758	103.9	113.4	3.1
g ⁺			1.771	104.7	113.0	5.5
g ⁻			1.771	104.7	113.0	5.5
t	t	c-c	1.759	103.8	113.3	-3.1
g ⁺			1.771	104.7	112.9	2.0
g ⁻			1.771	104.7	112.9	-2.0
bond b						
t	t-t	t	5.789	113.0	113.0	0.0
	(t-c) ⁺		5.789	113.2	113.2	70.8
	(t-c) ⁻		5.789	113.2	113.2	-70.8
	(c-t) ⁺		5.787	113.4	113.4	106.3
	(c-t) ⁻		5.787	113.4	113.4	-106.3
	c-c		5.788	113.3	113.3	180.0
bond d (g)						
t	t	t	1.831	103.6	109.1	3.7
	g ⁺		1.830	104.4	113.5	96.9
	g ⁻		1.830	104.4	113.5	-96.9
	t	g ⁺	1.834	103.5	110.7	1.3
	g ⁺		1.833	104.2	114.8	96.0
	g ⁻		1.835	105.4	113.7	-68.8
	t	g ⁻	1.834	103.5	110.7	-1.3
	g ⁺		1.835	105.4	113.7	68.8
	g ⁻		1.833	104.2	114.8	-96.0

conformation						
bond ^a			l_j^b	θ_{j-1}^c	θ_j^d	ϕ_j^e
$j-1$	j	$j+1$				
bond e (f)						
t	t	t	1.534	109.1	110.1	-0.8
	g ⁺		1.535	110.7	113.2	112.6
	g ⁻		1.535	110.7	113.2	-112.6
g ⁺	t		1.533	113.5	110.2	1.5
	g ⁺		1.535	114.8	112.6	113.2
	g ⁻		1.536	113.7	113.2	-112.5
g ⁻	t		1.533	113.5	110.2	-1.5
	g ⁺		1.536	113.7	113.2	112.5
	g ⁻		1.535	114.8	112.6	-113.2
t	t	g ⁺	1.534	109.1	113.2	4.3
	g ⁺		1.535	110.4	116.2	114.0
	g ⁻		1.536	112.5	117.1	-101.3
g ⁺	t		1.533	113.6	113.2	5.9
	g ⁺		1.534	114.6	115.9	117.8
	g ⁻		1.535	115.6	117.1	-104.7
g ⁻	t		1.533	113.2	113.2	2.8
	g ⁺		1.534	113.0	115.1	107.4
	g ⁻		1.536	115.9	116.3	-98.2
t	t	g ⁻	1.534	109.1	113.2	-4.3
	g ⁺		1.536	112.5	117.1	101.3
	g ⁻		1.535	110.4	116.2	-114.0
g ⁺	t		1.533	113.2	113.2	-2.8
	g ⁺		1.536	115.9	116.3	98.2
	g ⁻		1.534	113.0	115.1	-107.4
g ⁻	t		1.533	113.6	113.2	-5.9
	g ⁺		1.535	115.6	117.1	104.7
	g ⁻		1.534	114.6	115.9	-117.8

^a j denotes the current bond. The blank line represents that the conformation is inexistent. ^b Length of bond j in Å. ^c Angle formed between bonds $j-1$ and j in deg. ^d Angle formed between bonds j and $j+1$ in deg. ^e Dihedral angle of bond j in deg.

Table S7 Average geometrical parameters of PA2T, PA3T, PTA2T, and PTA3T

	bond	bond length (Å)	bond angle (°)	dihedral angle (°)			
				ϕ_t or ϕ_{t-t}	ϕ_g^\pm or $\phi_{(t-c)^\pm}$	$\phi_{(c-t)^\pm}$	ϕ_{c-c}
PA2T	a	1.360	117.5	0.1			
	b	5.810	117.5	0.0	± 53.6	± 123.0	180.0
	c	1.360	122.5	0.1			
	d	1.458	112.9	0.0	± 92.2		
	e	1.529	112.9	0.2	± 113.9		
	f	1.458	122.5	0.0	± 92.2		
PA3T	a	1.359	118.7	0.0			
	b	5.823	118.7	0.0	± 46.9	± 129.2	180.0
	c	1.359	123.0	0.0			
	d	1.458	112.8	-19.0	± 72.1		
	e	1.531	112.8	-0.2	± 116.2		
	f	1.531	112.8	-0.2	± 116.2		
	g	1.458	123.0	-19.0	± 72.1		
PTA2T	a	1.340	115.3	0.1			
	b	5.793	115.3	0.0	± 79.6	± 98.0	180.0
	c	1.340	125.8	0.1			
	d	1.457	112.4	-2.2	± 88.4		
	e	1.529	112.4	0.0	± 113.3		
	f	1.457	125.8	-2.2	± 88.4		
PTA3T	a	1.338	115.1	0.3			
	b	5.808	115.1	0.0	± 75.8	± 102.5	180.0
	c	1.338	125.9	0.3			
	d	1.459	111.7	4.0	± 88.4		
	e	1.529	112.4	-0.2	± 111.3		
	f	1.529	111.7	-0.2	± 111.3		
	g	1.459	125.9	4.0	± 88.4		

Table S8 Average geometrical parameters of P2TS₂, P3TS₂, P2TS₄, and P3TS₄

	bond	bond length (Å)	bond angle (°)	dihedral angle (°)			
				ϕ_t or ϕ_{t-t}	ϕ_{g^\pm} or $\phi_{(t-c)^\pm}$	$\phi_{(c-t)^\pm}$	ϕ_{c-c}
P2TS ₂	a	1.802	117.9	0.0			
	b	5.787	117.9	0.0			180.0
	c	1.802	99.4	0.0			
	d	1.832	113.1	-0.2	±97.3		
	e	1.523	113.1	0.0	±115.0		
	f	1.832	99.4	-0.2	±97.3		
P3TS ₂	a	1.807	118.2	0.0			
	b	5.793	118.2	0.0			180.0
	c	1.807	99.0	0.0			
	d	1.837	113.6	0.0	±95.2		
	e	1.532	113.3	0.0	±110.8		
	f	1.532	113.6	0.0	±110.8		
	g	1.837	99.0	0.0	±95.2		
P2TS ₄	a	1.762	113.0	0.1			
	b	5.787	113.0	0.0	±71.5	±106.3	180.0
	c	1.762	104.5	0.1			
	d	1.826	113.3	1.2	±85.1		
	e	1.525	113.3	0.0	±112.8		
	f	1.826	104.5	1.2	±85.1		
P3TS ₄	a	1.768	113.0	0.0			
	b	5.788	113.0	0.0	±70.8	±106.3	180.0
	c	1.768	104.3	0.0			
	d	1.832	113.3	0.8	±91.3		
	e	1.534	113.7	0.0	±110.4		
	f	1.534	113.3	0.0	±110.4		
	g	1.832	104.3	0.8	±91.3		