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Susa, Arijana; Bose, Ranjita; Grande, Antonio; van der Zwaag, Sybrand; Garcia Espallargas, Santiago

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Effect of the dianhydride/branched diamine ratio on the architecture and room temperature healing behaviour of polyetherimides

A. Susa, R. K. Bose, A. M. Grande, S. van der Zwaag, S. J. Garcia*

Novel Aerospace Materials group, Faculty of Aerospace Engineering,

Delft University of Technology, Kluyverweg 1, 2629 HS, Delft, The Netherlands

*e-mail: s.j.garciaespallargas@tudelft.nl

List of figures and tables:

Figure S1.	Full IR spectra of the four PEIs as a function of the offset from the theoretical stoichiometric ratio.	S-2
Figure S2.	¹ H NMR spectra of the monomers: ODPa in DMSO and DD1 in CDCl ₃ and their assignment to the molecular structure.	S-3
Figure S3.	¹ H NMR spectra of the four polymers in CDCl ₃ and their assignment to the polymer structure.	S-4, S-5
Figure S4:	WAXS and temperature controlled SAXS diffractograms.	S-6
Figure S5:	TGA curves and DSC curves from the second heating cycle showing glass transitions.	S-5
Table S-I.	Results of the swelling tests.	S-7
Figure S6:	a) van Gurp-Palmen plot; b) horizontal shift factors (a_T) from the TTS mastercurves, $T_{ref}=25^\circ\text{C}$; c) loss tangent ($\tan\delta$) as a function of temperature.	S-8
Table S-II:	Tensile properties of the virgin polymers.	S-9
Figure S7.	Stress-strain curves showing the D-0.9 healing behaviour at the higher healing T.	S-9
Table S-III.	Effect of branching on the generic polymer properties.	S-10
Figure S8.	a) TTS mastercurve of ND-1.0 and b) van Gurp-Palmen plots of the referent non-branched PEI ND-1.0 in comparison to its branched counterpart D-1.0	S-10
Table S-IV.	Characteristic parameters obtained from rheology for the non-branched PEI (ND-1.0) as compared to its branched counterpart (D-1.0).	S-10
Figure S9.	Temperature dependant rheological behaviour of the neat DD1.	S-11

ATR/FTIR spectra

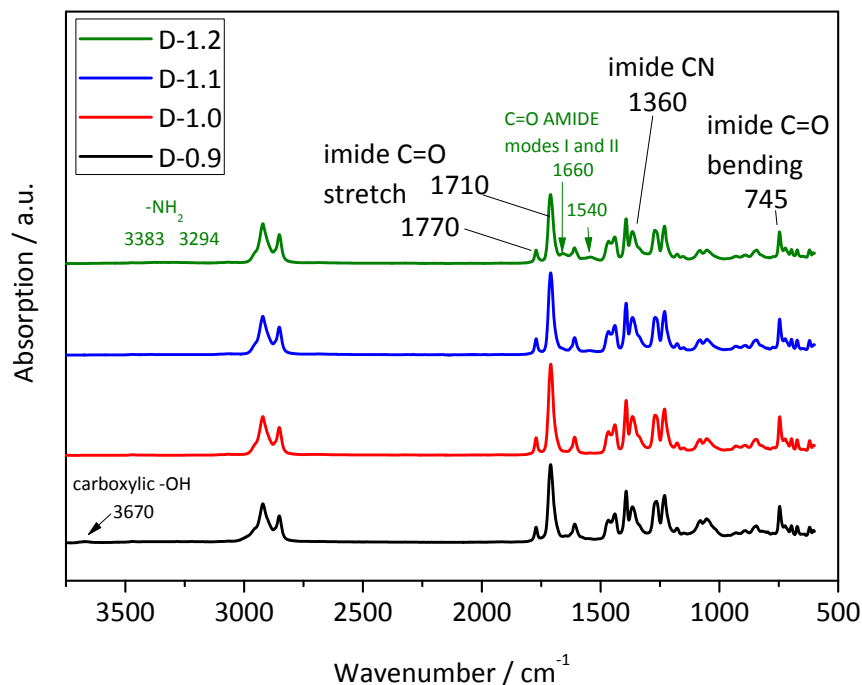


Figure S1. Full IR spectra of the four PEIs as a function of the offset from the theoretical stoichiometric ratio.

Yield

Percent yield of the polymer was calculated according to equation:

$$\text{Percent yield (\%)} = 100 \left(\frac{\text{Actual mass of the product}}{\text{Predicted mass of the product}} \right)$$

where predicted mass of the product was calculated according to the stoichiometric balance, assuming that 1 mol of ODPA and 1 mol of DD1 give 1 mol of PEI and 2 mol of water (4.25 wt% of water):

$$\text{Predicted mass of the product/g} = m(\text{ODPA}) + m(\text{DD1}) - m(\text{H}_2\text{O})$$

¹H NMR

Solution state ¹H NMR spectra were collected using the Agilent-400 MR DD2 at 25°C at 400 MHz. The solutions of polymers and DD1 were prepared in CDCl₃, and ODPA was measured in deuterated DMSO. Spectra were referenced to the solvent residual peak for DMSO and to TMS for CDCl₃, respectively. Spectra were not normalized. PAA's were not tested due to the insolubility in the available NMR solvents. Furthermore, D-1.2 was insoluble as well, but due to the partial crosslinked nature.

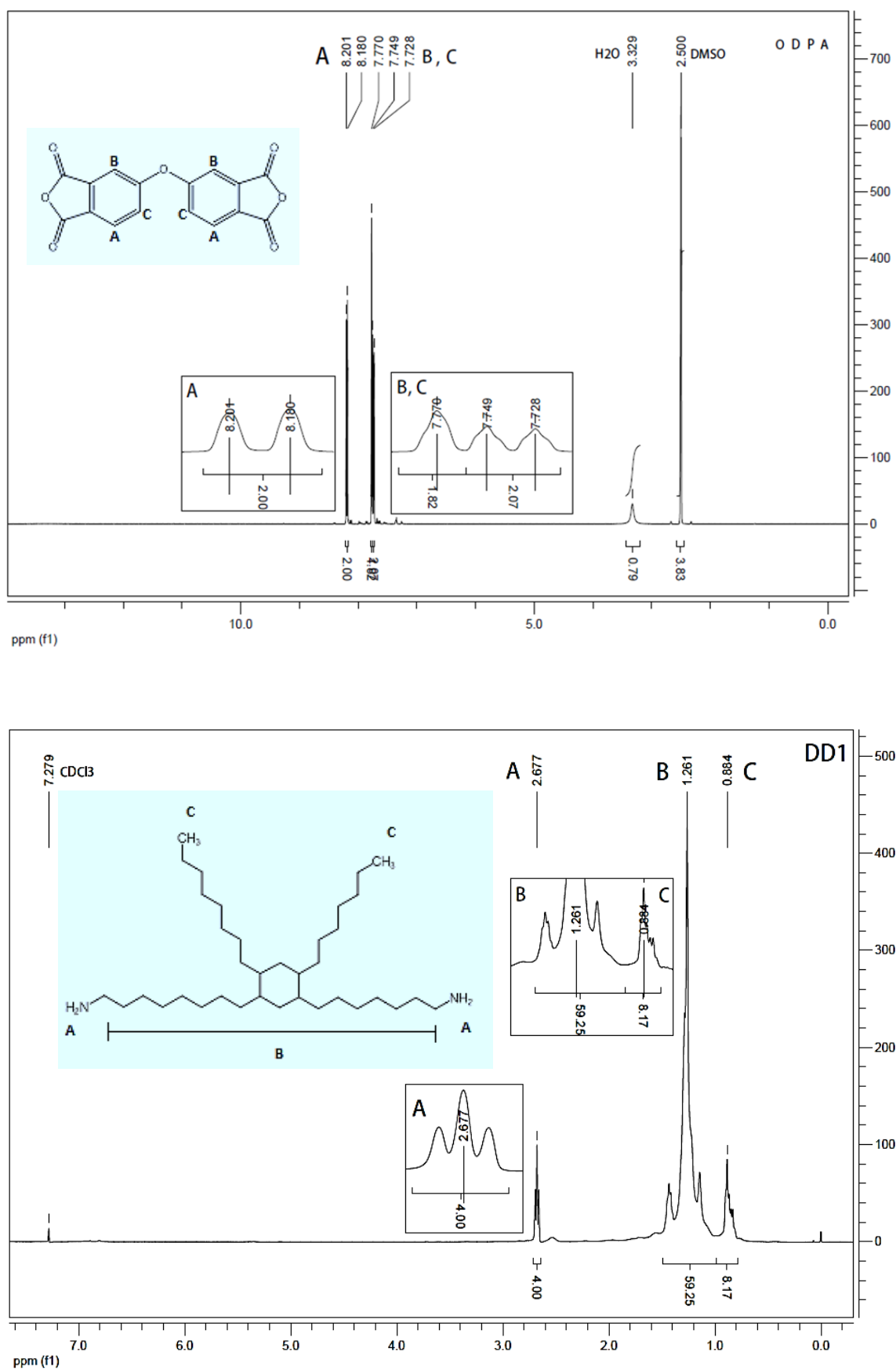
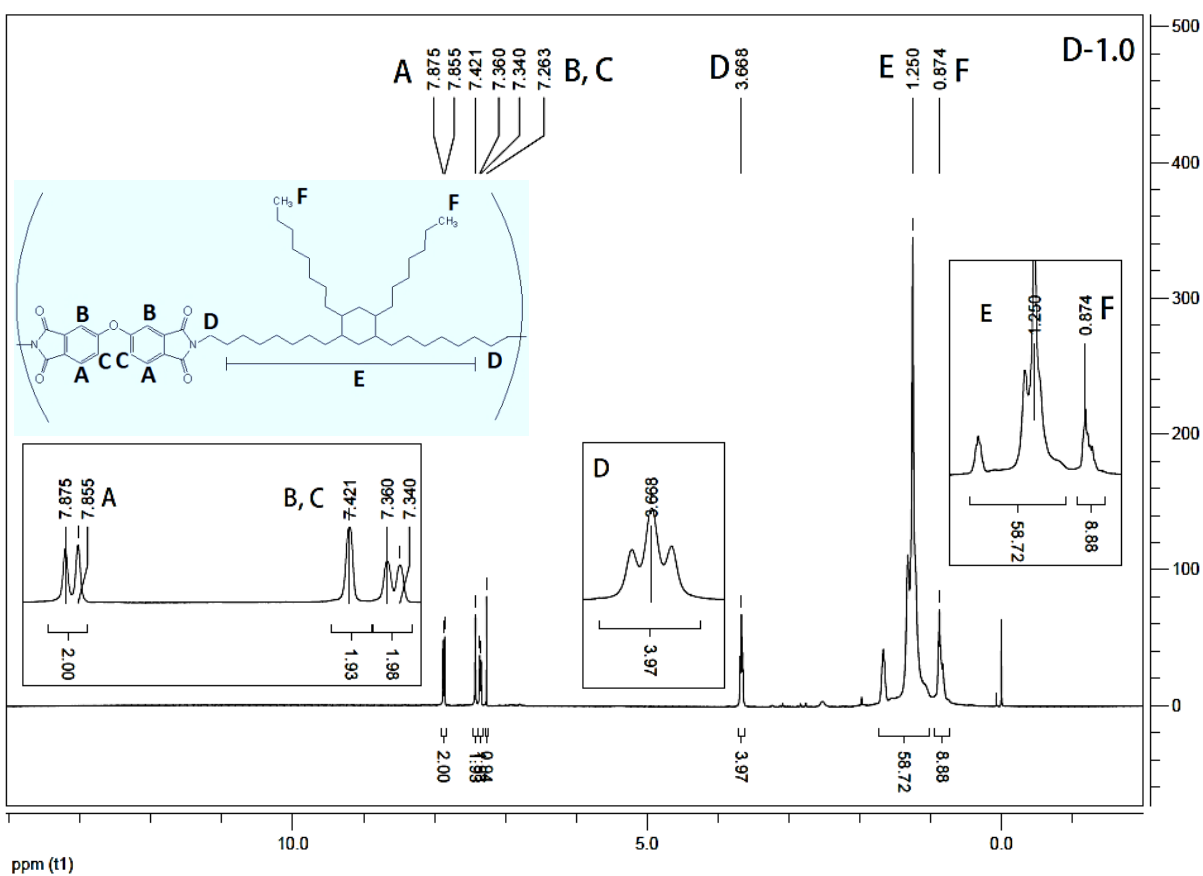
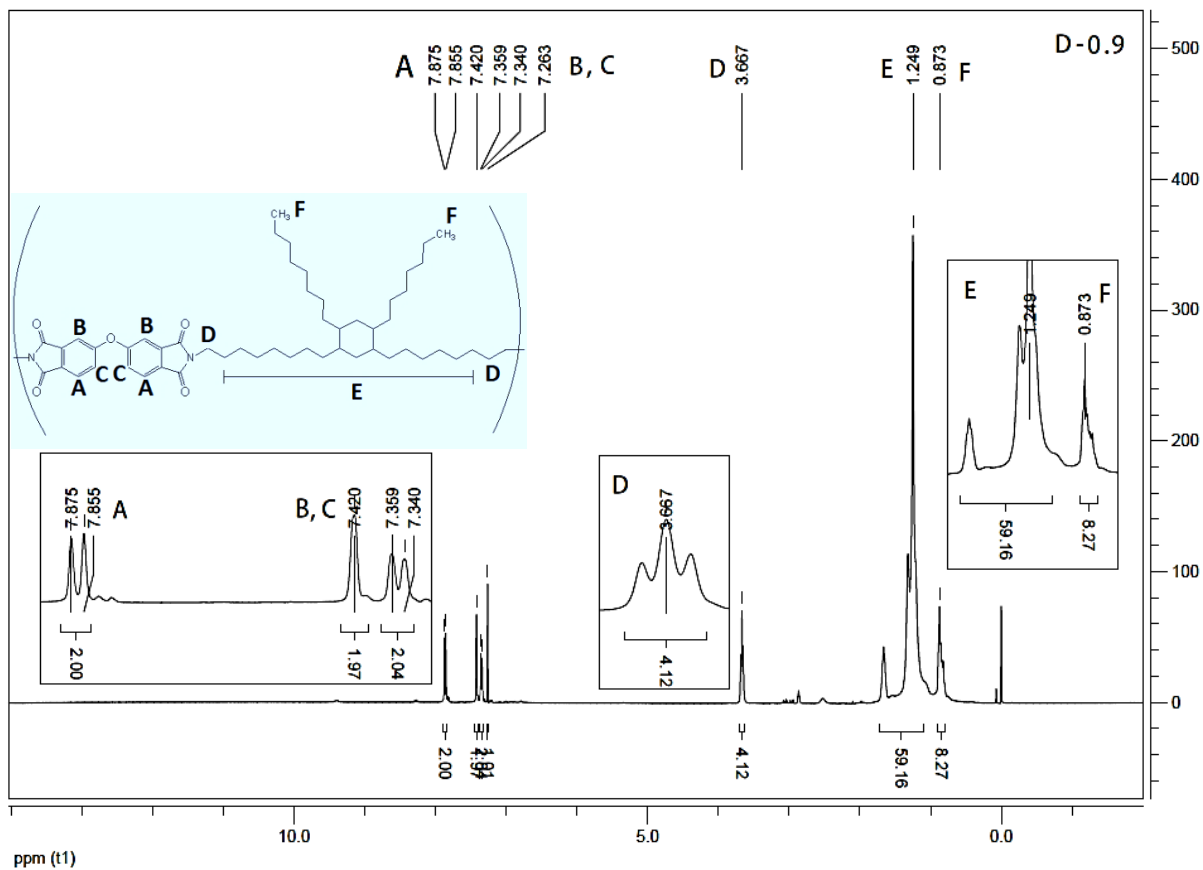


Figure S2. ¹H NMR spectra of the monomers: ODPA in DMSO and DD1 in CDCl₃ and their assignment to the molecular structure.



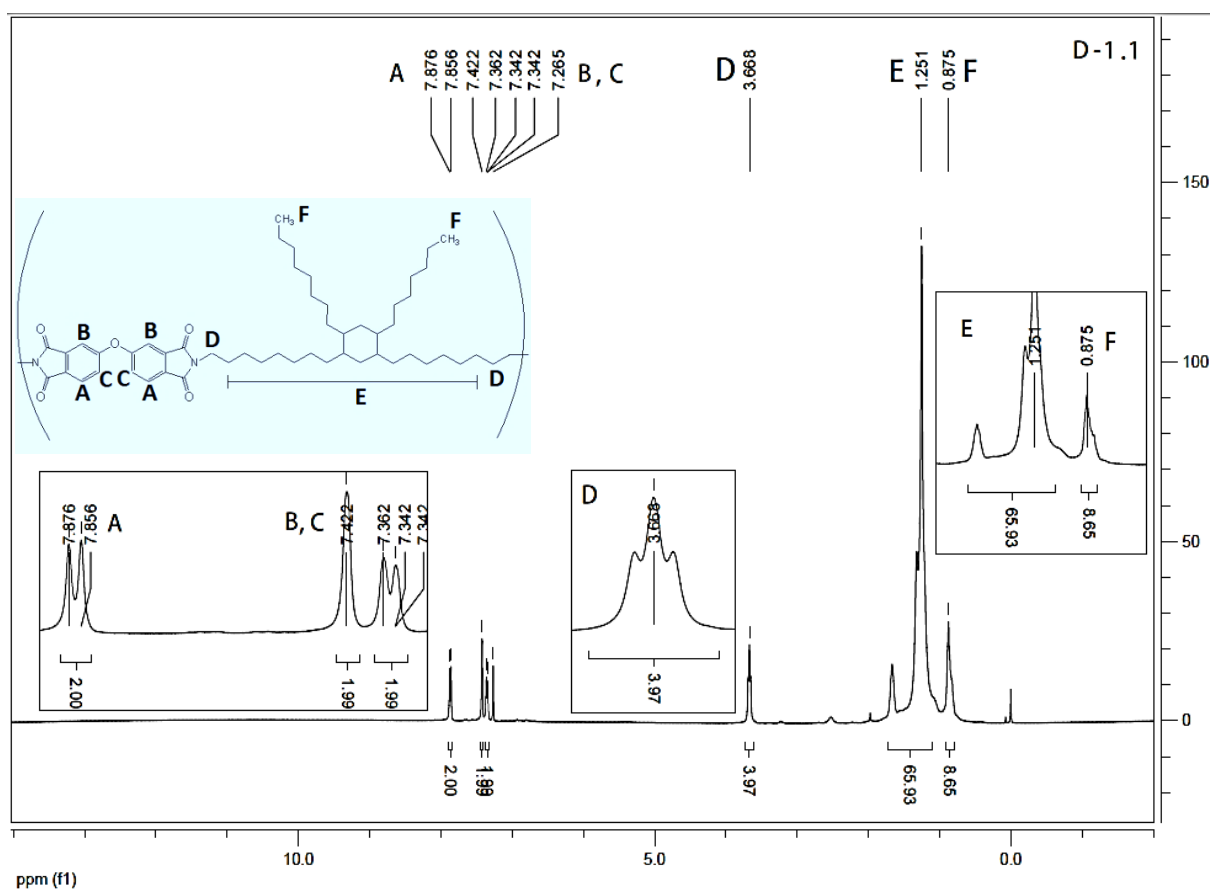


Figure S3. ^1H NMR spectra of the three polymers in CDCl_3 and their assignments to the polymer structure.

X-ray diffraction measurements

Wide Angle X-ray diffraction (WAXS) data were collected on a Bruker D8 Advance diffractometer with Co K α radiation at the room temperature. Temperature controlled Small Angle X-ray Scattering (SAXS) was conducted using an AXS D8 Discover instrument from Bruker AG. SAXS scans were collected at five different temperatures: preliminary scan at room temperature followed by heating to 60°C. In a cooling ramp at the 0.5°C/min rate, the scans were further collected at 50, 25 and 5°C.

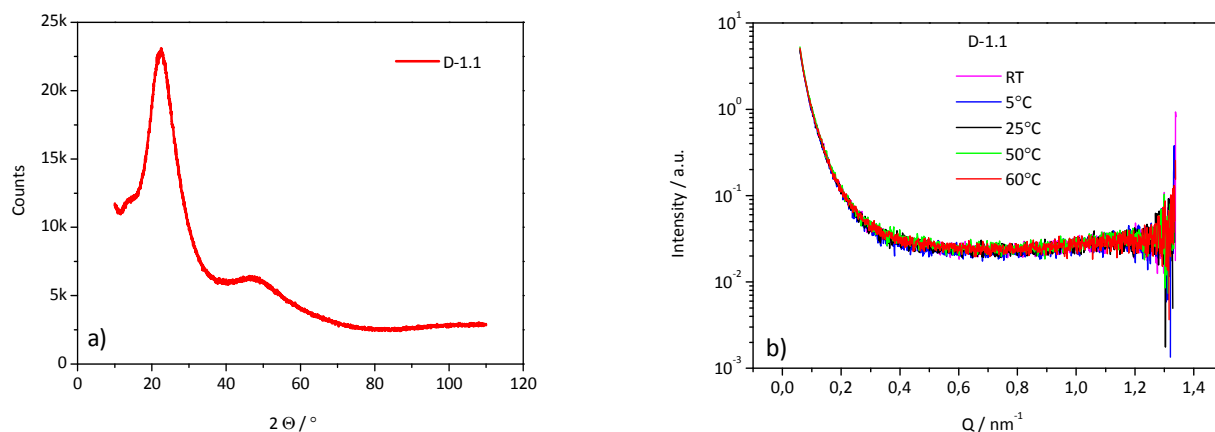


Figure S4. a) WAXS and b) temperature controlled SAXS diffractogram of the D-1.1 polymer with representative patterns of all polymers in this work.

Thermal analysis

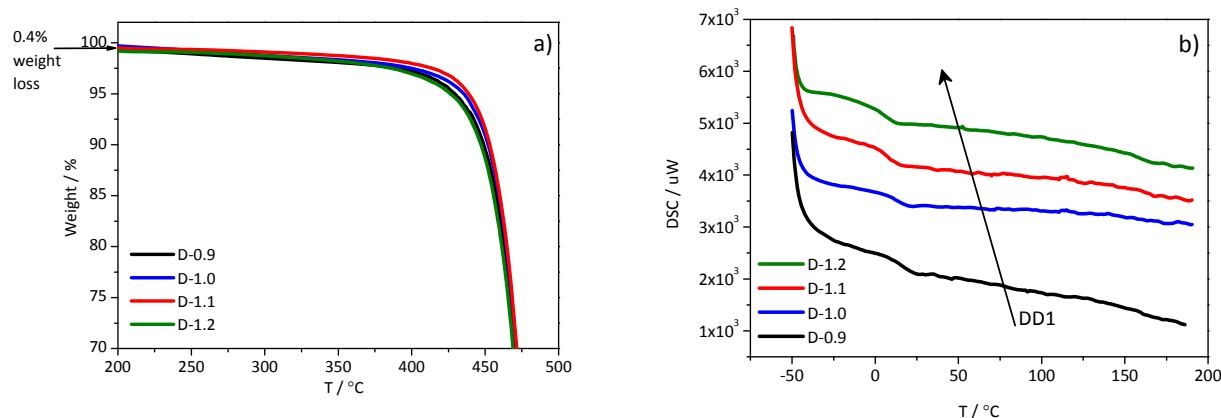


Figure S5. a) TGA curves showing weight loss of SH-PEIs in the high temperature range; b) DSC curves from the second heating cycle showing glass transitions.

Swelling tests

The crosslinking density was determined by solvent-swelling measurements for 72h and calculated according to the Flory–Rehner equation (eq. S1).

$$\ln(1 - \Phi_r) + \Phi_r + \chi\Phi_r^2 = -\frac{\rho_r}{M_c} V_s \left(\Phi_r^{1/3} - \frac{2\Phi_r}{f} \right) \quad \text{eq-S1}$$

where:

Φ_r = volume fraction of swollen polymer

χ = polymer-solvent interaction parameter. In this work a value of 0.3 (cis-polybutadiene raw elastomer-toluene¹) was taken as reference due to the lack of existing values for this new class of polymers.

ρ_r = polymer density

M_c = molecular weight of polymer between two crosslinks

V_s = molar volume of solvent (106.8 mL/mol for toluene¹)

f = functional cross-links (3)

The volume fraction (Φ_r) of swollen polymer was calculated according to:

$$\Phi_r = \frac{w_i/\rho_r}{w_i/\rho_r + (w_s - w_d)/\rho_s} \quad \text{eq-S2}$$

where:

w_i = initial weight of the sample

w_s = swollen weight of the sample after 72h of immersion

w_d = weight of the sample dried at 60°C in vacuum for 72h

ρ_s = solvent density (0.87 g/cm³ for toluene)

The crosslink density (ν) is then calculated as:

$$\nu = \frac{1}{2M_c} \quad \text{eq-S3}$$

Table S-I. Results of the swelling tests.

Polymer	Q, Swelling ratio after 72h (%)	Soluble fraction (%)	M_c (g/mol)	ν , Crosslinking density (mol/cm ³)
D-0.9	/	100	/	/
D-1.0	/	100	/	/
D-1.1	2680 ± 186	60 ± 8	2609 ± 62	1.92 ± 0.05
D-1.2	441 ± 8	13 ± 0,3	929 ± 14	5.39 ± 0.08

¹ Hansen, C.M., *Hansen solubility parameters: a user's handbook*, CRC Press LLC: USA, 2000.

Rheological parameters

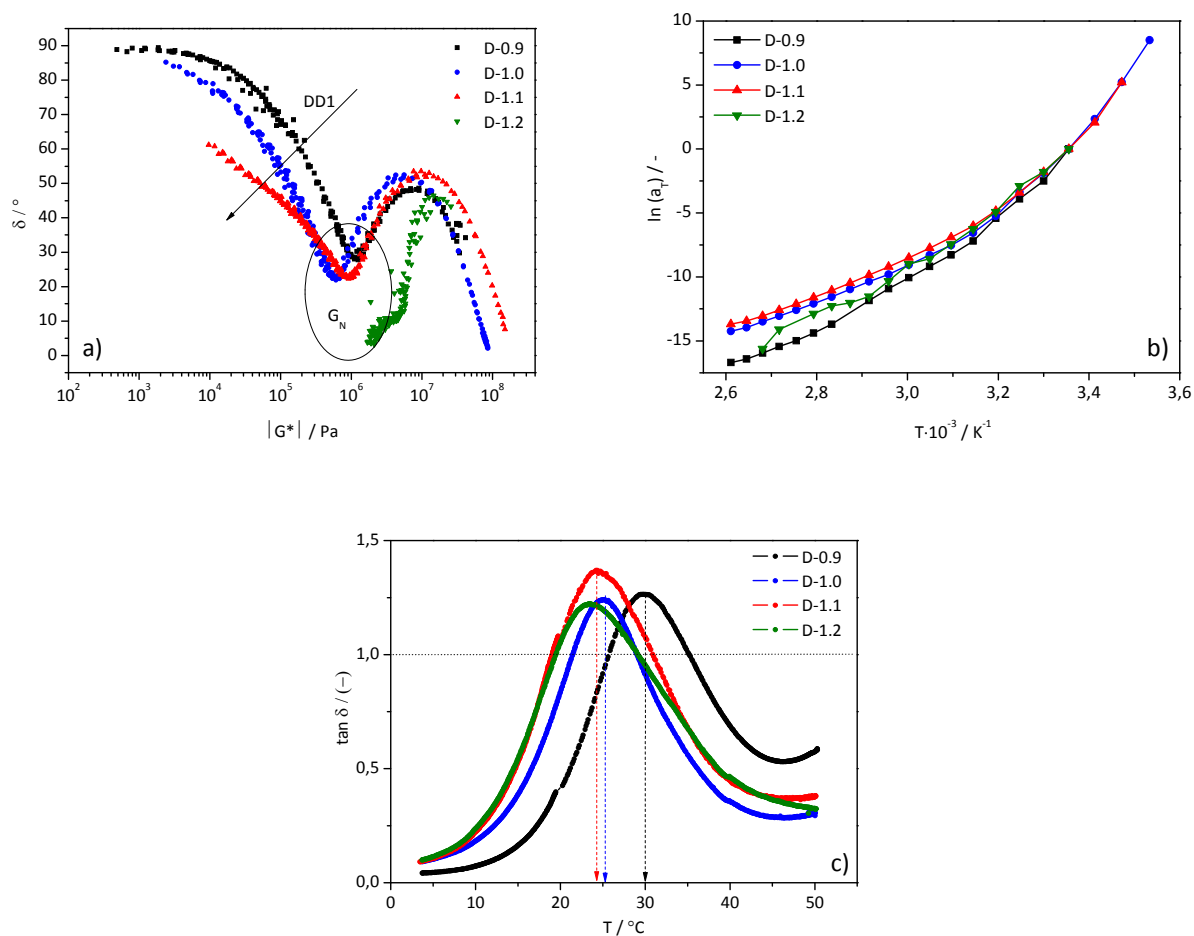


Figure S6: **a)** van Gurp-Palmen plot; **b)** horizontal shift factors (a_T) from the TTS mastercurves, $T_{\text{ref}}=25^\circ\text{C}$; **c)** loss tangent ($\tan\delta$) as a function of temperature (T) from the temperature sweep experiments. Dotted lines are guidance for a reader's eye showing the elastic/viscous dominance region determined by $\tan\delta=1$. Arrows point the values of T ($\tan\delta_{\text{max}}$) for the non-crosslinked polymers that heal at 25 $^\circ\text{C}$ (red and blue) and does not heal at 25 $^\circ\text{C}$ (black).

Tensile test

Table S-II: Tensile properties of the virgin polymers.

Polymer	σ_b (MPa)	ϵ_b (%)	E (MPa)
D-0.9	5.9±0.2	330±20	110
D-1.0	5.7±0.1	440±20	54
D-1.1	4.4±0.2	560±10	33
D-1.2	2.0±0.4	360±50	6.5

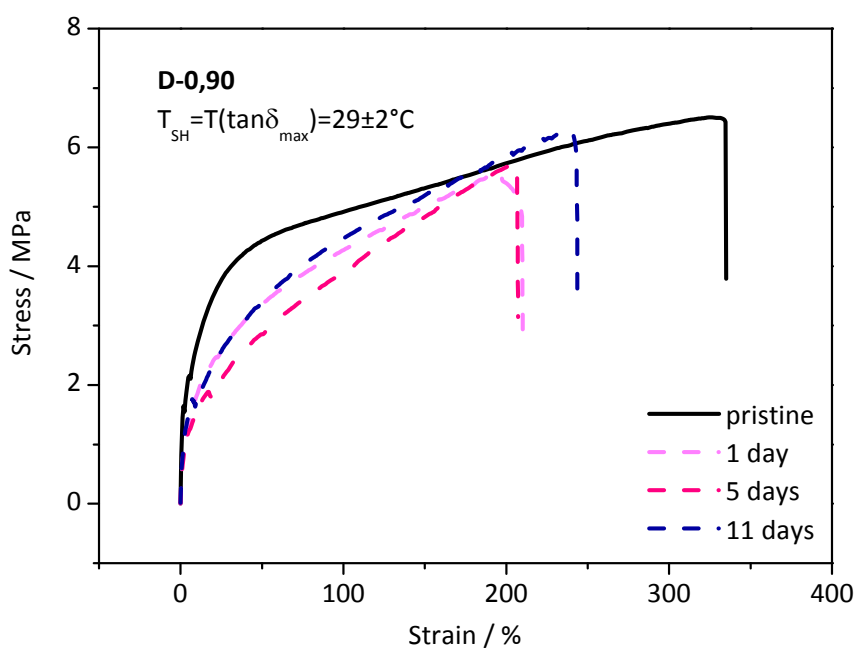
Healing tests in tension at elevated T 

Figure S7. Stress-strain curves showing the **D-0.9** healing behaviour at the higher healing T , that corresponds to T of the maximum of $\tan\delta$ determined from the rheological T -sweep test, as function of the healing time. Full lines represent pristine (—) and dashed lines represent healed (---) samples after given healing time.

Reference: non-branched PEI

Table S-III. Effect of branching on the generic polymer properties. Where ND-1.0 contains a linear non-branched C12 aliphatic diamine and D-1.0 a branched C18 aliphatic diamine (DD1) in (theoretical) stoichiometric ratio.

polymer	Mw (g/mol)	Mn (g/mol)	PDI	DSC-T _g (°C)	TGA-T (2% weight loss) (°C)	Density (g/cm ³)
ND-1.0	*	*	*	69	435	1.20
D-1.0	32k	16k	2,0	13	380	1.05

* GPC data not available since ND-1.0 is not soluble in the GPC solvents available.

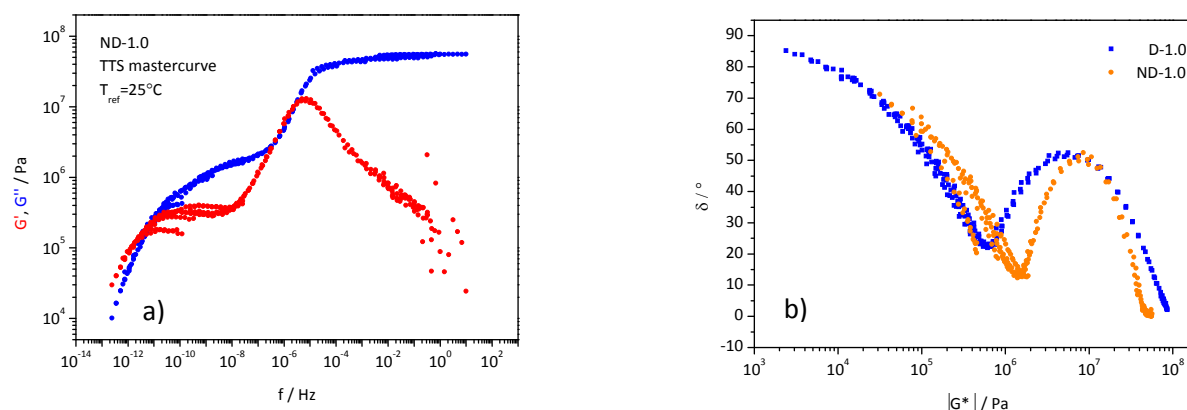


Figure S8. a) TTS mastercurve of ND-1.0 and b) van Gorp-Palmen plots of the referent non-branched PEI (orange) ND-1.0 in comparison to its branched counterpart D-1.0 (blue). T_{ref}=25°C

Table S-IV. Characteristic parameters obtained from rheology for the non-branched PEI (ND-1.0) as compared to its branched counterpart (D-1.0).

Polymer	f _s (Hz)	f _d (Hz)	f _g (Hz)	τ _s (s)	τ _d (s)	τ _g (s)	G' slope at f < f _s	G'' slope at f < f _s	G _N * (Pa)	M _e ** (g/mol)
ND-1.0	7.5·10 ⁻¹²	3.8·10 ⁻⁷	2.8·10 ⁻⁶	13.3·10 ¹¹	2.6·10 ⁶	3.6·10 ⁵	1.06	0.49	1.39·10 ⁶	2140
D-1.0	2.0·10 ⁻⁵	0.035	1.5	5.0·10 ⁴	28.0	0.7	1.25	0.83	6.09·10 ⁵	4270

* G_N calculated from the Van Gorp-Palmen plot, δ(|G*|)^{2,3}

** M_e were calculated using experimentally determined densities (Table S-III) according to M_e = ρRT / G_N (Doi and Edwards) equation.

² Ahmadi, M.; Hawke, L. G. D.; Goldansaz, H.; van Ruymbeke, E. *Macromolecules* **2015**, *48*, 7300.

³ Trinkle, S.; Friedrich, C. *Rheol. Acta* **2001**, *40*, 322.

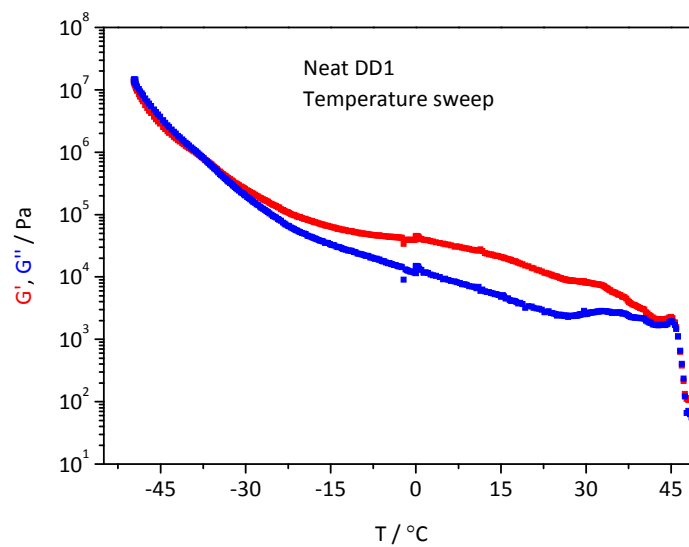
Neat branched dimer diamine (DD1); T-sweep

Figure S9. Temperature dependant rheological behaviour of the neat DD1.