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Correction: Toward realistic computer modeling of paraffin-based composite materials: critical assessment of atomic-scale models of paraffins

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 Correction for 'Toward realistic computer modeling of paraffin-based composite materials: critical assessment of atomic-scale models of paraffins' by Artyom D. Glova *et al.*, *RSC Adv.*, 2019, 9, 38834–38847, DOI: 10.1039/C9RA07325F.

The authors regret that the 1–4 Lennard-Jones interactions were not accounted for in the molecular dynamics simulations of *n*-eicosane samples with the use of the united-atom GROMOS force field, as required by the original parametrization of this force field.¹ This led to the abnormal behavior of the corresponding systems, namely: the *n*-eicosane samples did not crystallize within the temperature range of 200–450 K. After the 1–4 interactions were turned on, the GROMOS force field allowed us to observe the crystallization of *n*-eicosane with the transition temperature of 270 ± 1 K, see Fig. 1.

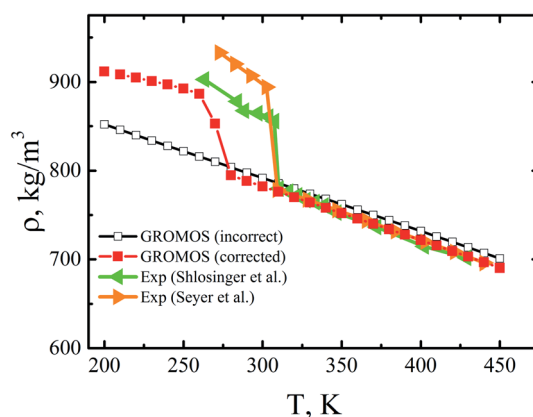


Fig. 1 Mass density of *n*-eicosane samples as a function of temperature for the GROMOS force field, *cf.* Fig. 2(b) of the original publication. The open black squares show the original incorrect data, and the closed red squares correspond to the corrected results. For the sake of comparison, the experimental data^{2,3} is also presented.

Most structural and dynamic characteristics also changed after the 1–4 interactions were properly accounted for. Below we present the updated versions of Tables 1 and 2 of the original publication.

Overall, the GROMOS force field does not show abnormal behavior for the *n*-eicosane sample anymore and performs rather similarly to other united-atom force fields such as *e.g.* PYS. The main conclusions of the original publication remain unchanged since the use of all-atom general-purpose force fields was shown to provide a more realistic description for *n*-eicosane as compared to their united-atom counterparts.



Table 1 The crystallization temperature (T_c), the coefficient of volumetric thermal expansion (CTE) measured in the temperature interval 400–450 K, the radius of gyration (R_g), and the end-to-end distance H of n -eicosane chains for different force fields. The corrected data for the GROMOS force field is highlighted in bold

Force field	T_c , K	CTE, 10^{-4} K^{-1}	R_g , nm ($T = 250 \text{ K}$)	R_g , nm ($T = 450 \text{ K}$)	H , nm ($T = 250 \text{ K}$)	H , nm ($T = 450 \text{ K}$)
GAFF	330 ± 1	17.4 ± 0.1	0.75 ± 0.01	0.60 ± 0.01	2.43 ± 0.01	1.70 ± 0.01
GAFF2	365 ± 2	14.3 ± 0.1	0.75 ± 0.01	0.62 ± 0.01	2.43 ± 0.01	1.82 ± 0.01
OPLS-AA	365 ± 1	14.7 ± 0.1	0.74 ± 0.01	0.62 ± 0.01	2.42 ± 0.01	1.80 ± 0.01
L-OPLS-AA	265 ± 3	12.3 ± 0.1	0.72 ± 0.01	0.57 ± 0.01	2.29 ± 0.01	1.63 ± 0.01
CHARMM36	320 ± 1	12.4 ± 0.1	0.74 ± 0.01	0.59 ± 0.01	2.39 ± 0.01	1.69 ± 0.01
GROMOS	270 ± 1	9.1 ± 0.1	0.72 ± 0.01	0.58 ± 0.01	2.34 ± 0.01	1.66 ± 0.01
NERD	270 ± 1	11.6 ± 0.1	0.74 ± 0.01	0.58 ± 0.01	2.41 ± 0.01	1.64 ± 0.01
OPLS-UA	310 ± 1	7.6 ± 0.1	0.73 ± 0.01	0.57 ± 0.01	2.38 ± 0.01	1.63 ± 0.01
PYS	270 ± 10	9.0 ± 0.1	0.72 ± 0.01	0.55 ± 0.01	2.34 ± 0.01	1.57 ± 0.01
TraPPE	280 ± 1	9.6 ± 0.1	0.74 ± 0.01	0.58 ± 0.01	2.41 ± 0.01	1.65 ± 0.01
Experiment	310 (ref. 4)	8.8–8.9 (ref. 2 and 3)	—	—	2.43 (ref. 5)	—

Table 2 The shear viscosity, the diffusion coefficient and the mass density for n -eicosane samples in the liquid state simulated with different force fields at $T = 450 \text{ K}$. The corrected data for the GROMOS force field is highlighted in bold

Force field	η , mPa s	D , $10^{-5} \text{ cm}^2 \text{ s}^{-1}$	ρ , kg m^{-3}
GAFF	0.42 ± 0.04	2.9 ± 0.2	592.3 ± 0.1
GAFF2	0.59 ± 0.01	2.1 ± 0.1	633.5 ± 0.1
OPLS-AA	0.83 ± 0.07	1.6 ± 0.1	668.2 ± 0.1
L-OPLS-AA	0.64 ± 0.08	2.2 ± 0.2	656.4 ± 0.1
CHARMM36	0.60 ± 0.04	2.1 ± 0.2	658.1 ± 0.1
GROMOS	0.52 ± 0.01	2.7 ± 0.1	690.3 ± 0.1
NERD	0.32 ± 0.01	3.6 ± 0.1	661.0 ± 0.1
OPLS-UA	0.79 ± 0.02	1.8 ± 0.1	753.1 ± 0.1
PYS	0.53 ± 0.01	2.6 ± 0.1	698.8 ± 0.1
TraPPE	0.45 ± 0.04	2.9 ± 0.1	693.7 ± 0.1
Experiment	0.594 ($T = 453 \text{ K}$) ⁶	2.2 ($T = 443 \text{ K}$) ⁷	696.0 ($T = 440 \text{ K}$) ³

The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.

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