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Correction: Structural heterogeneity of milk casein micelles: a SANS contrast variation study

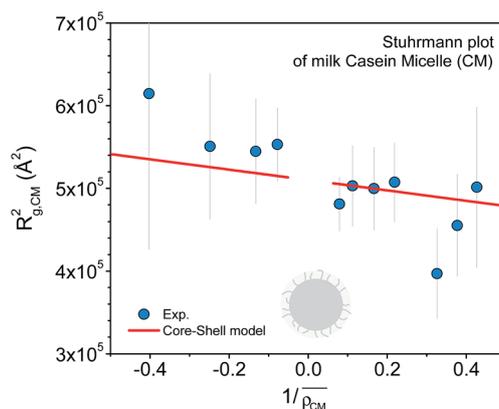
Antoine Bouchoux,^{‡*ab} Jorge Ventureira,^{ab} Geneviève Gésan-Guiziou,^{ab} Fabienne Garnier-Lambrouin,^{ab} Peng Qu,^{ab} Coralie Pasquier,^{ab} Stéphane Pézenec,^{ab} Ralf Schweins^c and Bernard Cabane^d

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Correction for 'Structural heterogeneity of milk casein micelles: a SANS contrast variation study' by Antoine Bouchoux *et al.*, *Soft Matter*, 2015, DOI: 10.1039/c4sm01705f.

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The original manuscript contained an error in the labelling of the y-axis in Fig. 7, and in the graphical abstract. Please see the corrected figures below:



Graphical Abstract

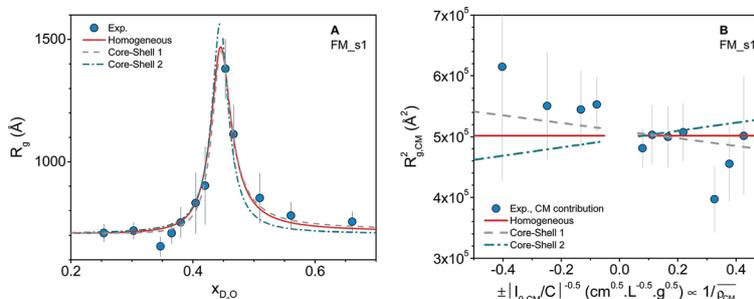


Fig. 7 Using non-homogeneous structural models for modeling the variation of R_g with contrast: an example with casein micelles from fresh milk (FM_{s1}). The description of the core-shell models is in the text. (A) gives the variation of the apparent radius of gyration R_g (*i.e.*, including the contribution of fat droplets) as a function of D_2O content. In (B), we use a representation similar to the one used by Stuhrmann,¹⁰ and that consists of plotting the squared radius of gyration of the casein micelle population $R_{g,CM}^2$ (*i.e.*, without the contribution of fat droplets) as a function of the reciprocal of the contrast of the micelles, $1/\rho_{CM}$. Details about the calculation of $R_{g,CM}^2$ and $1/\rho_{CM}$ from the experimental data are given in ESI part F.†

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

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