

CORRECTION

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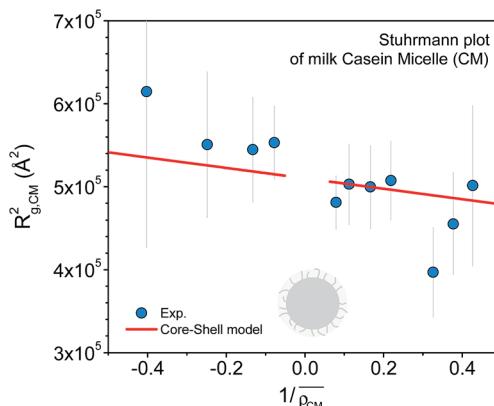
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Correction: Structural heterogeneity of milk casein micelles: a SANS contrast variation study

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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.

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

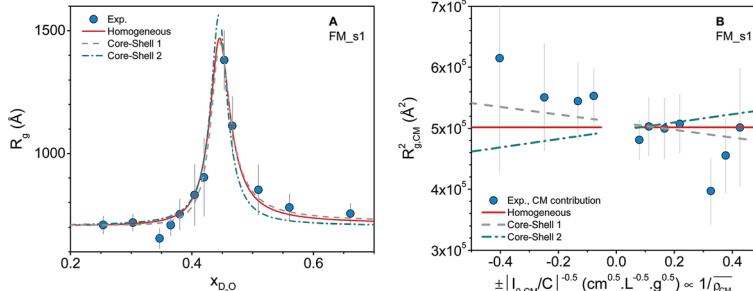


Fig. 7 Using non-homogeneous structural models for modeling the variation of R_g with contrast: an example with casein micelles from fresh milk (FM_s1). The description of the core–shell models is in the text. (A) gives the variation of the apparent radius of gyration R_g (i.e., including the contribution 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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