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

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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 D2O content. In (B), we use a representation similar to the one used by Stuhmann,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, $1/R_{CM}$. Details about the calculation of $R_{g,CM}^2$ and $1/R_{CM}$ 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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DOI: 10.1039/c4sm01705f
www.rsc.org/softmatter

806 | Soft Matter, 2015, 11, 806

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