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:

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

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$_2$O content. In (B), we use a representation similar to the one used by Stuhmann, and that consists of plotting the squared radius of gyration of the casein micelle population $R_g^{CM2}$ (i.e., without the contribution of fat droplets) as a function of the reciprocal of the contrast of the micelles, $1/R_g^{CM}$. Details about the calculation of $R_g^{CM2}$ and $1/R_g^{CM}$ from the experimental data are given in ESI part F.

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