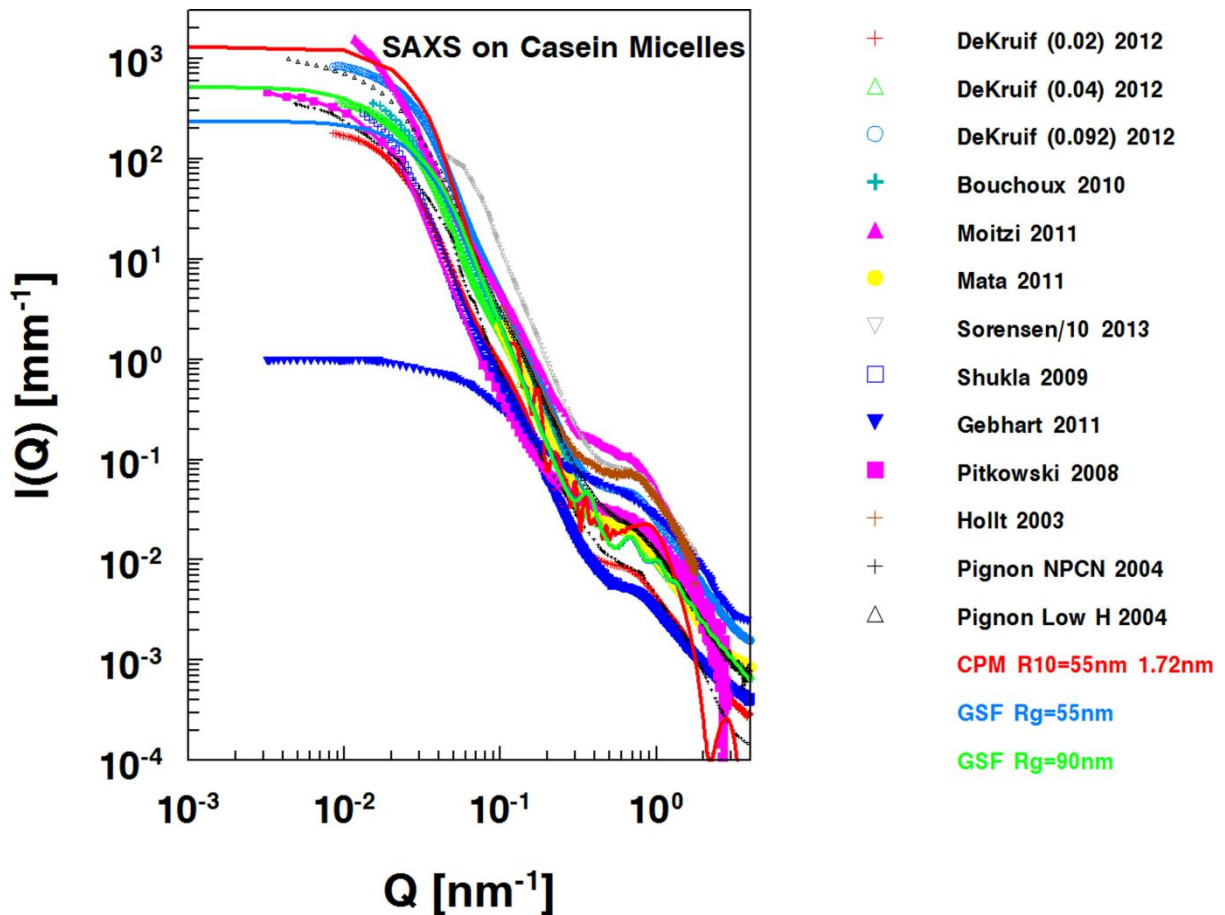


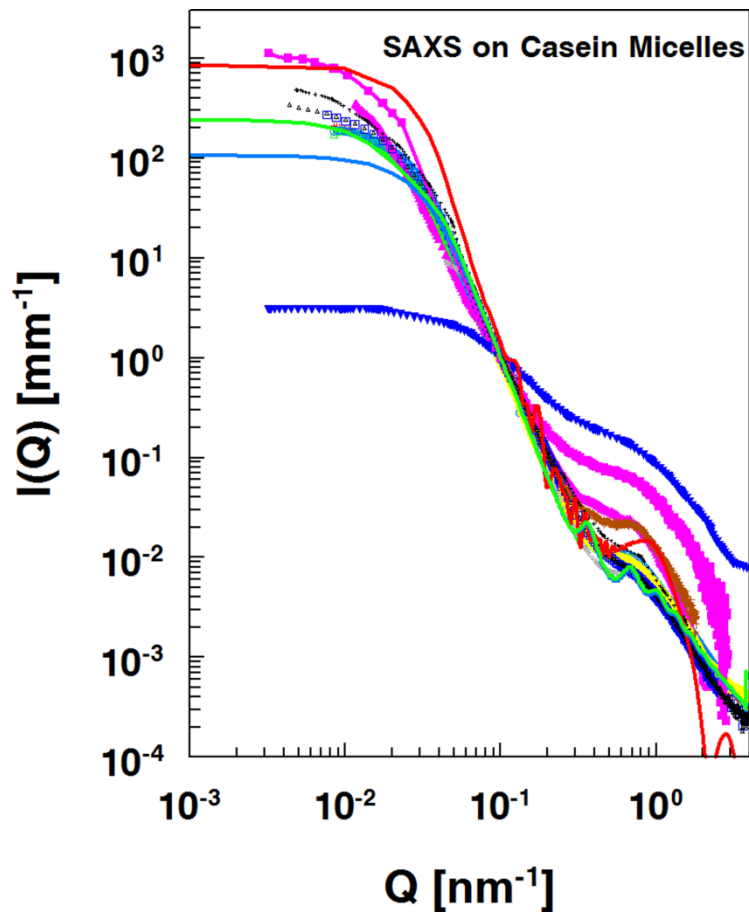
# The structure of casein micelles: a review of small angle scattering data.

## Supplementary figures

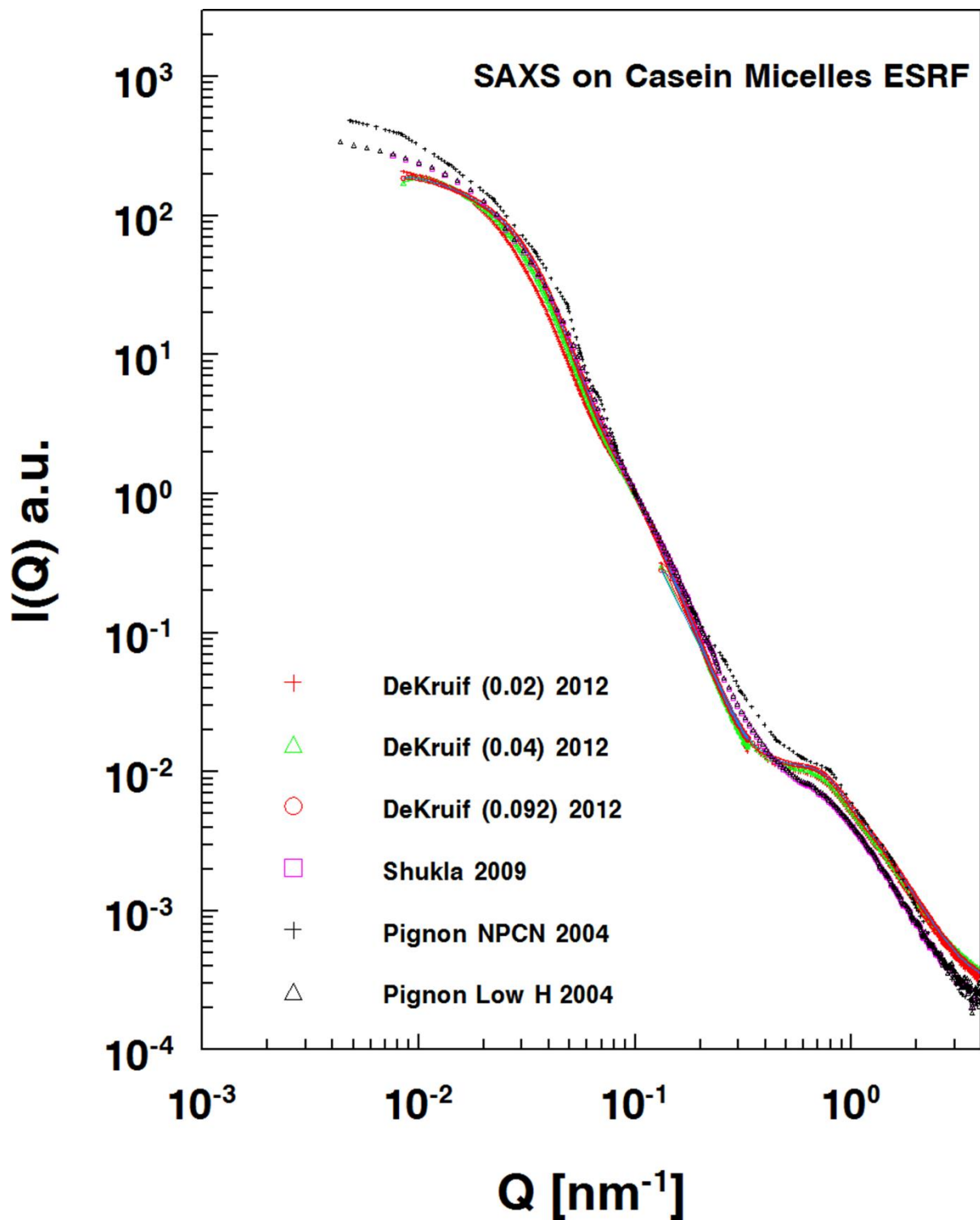
C.G. (Kees) De Kruif

**Figure 1.** SAXS data on casein micelles. (Top) Unscaled data and (below) all data scaled to  $I(Q = 0.1 \text{ nm}^{-1})=1$ . The wiggles in the calculated spectra are remnants of the numerical integration of a polydisperse sphere. In practice they will be smeared out.





**Figure 2.** Comparison of SAXS data on casein micelles obtained at the ESRF, beamline ID02, from three different investigations ( De Kruif *et al.*, 2012; Shukla *et al.*, 2009; Pignon *et al.*, 2004). The data of Shukla and Pignon at low heat superimpose perfectly at all Q.



**Figure 3.** SANS spectra at various contrasts, as indicated in the lower left-hand corner of each panel. The letters CK = Cornelis De Kruif (De Kruif *et al.*, 2012), CH = Carl Holt (Holt *et al.*, 2003), PS = Philip Stothart (1989), SH = Steen Hansen (Hansen *et al.*, 1996), HT = Hans Tromp (2013), TH = Thom Huppertz (Huppertz & De Kruif, 2008) and AJ = Andrew Jackson (Jackson & McGillivray, 2011).

