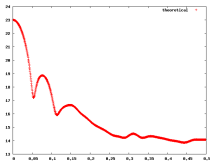




Fast SAXS Profile Computation with Debye Formula



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PDB files

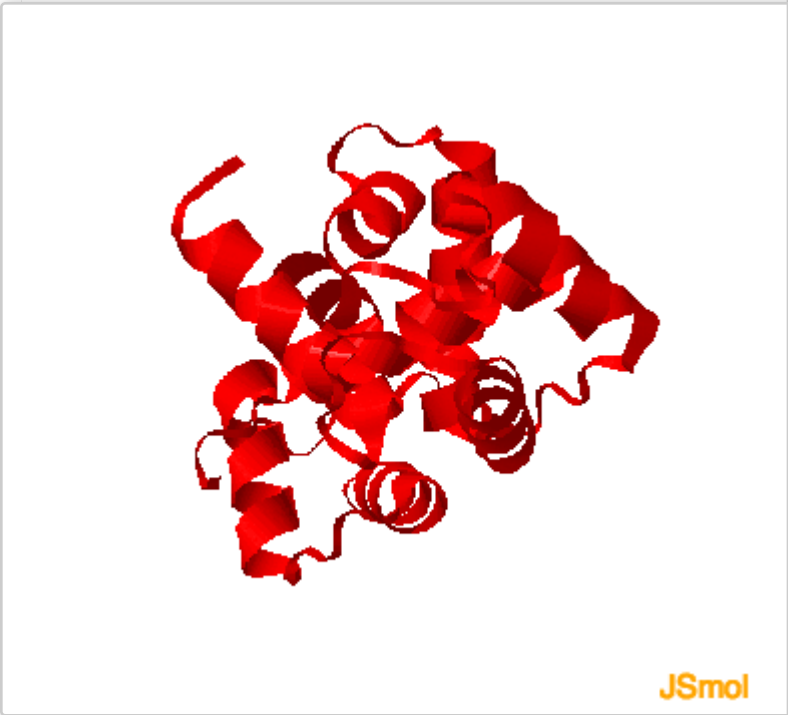
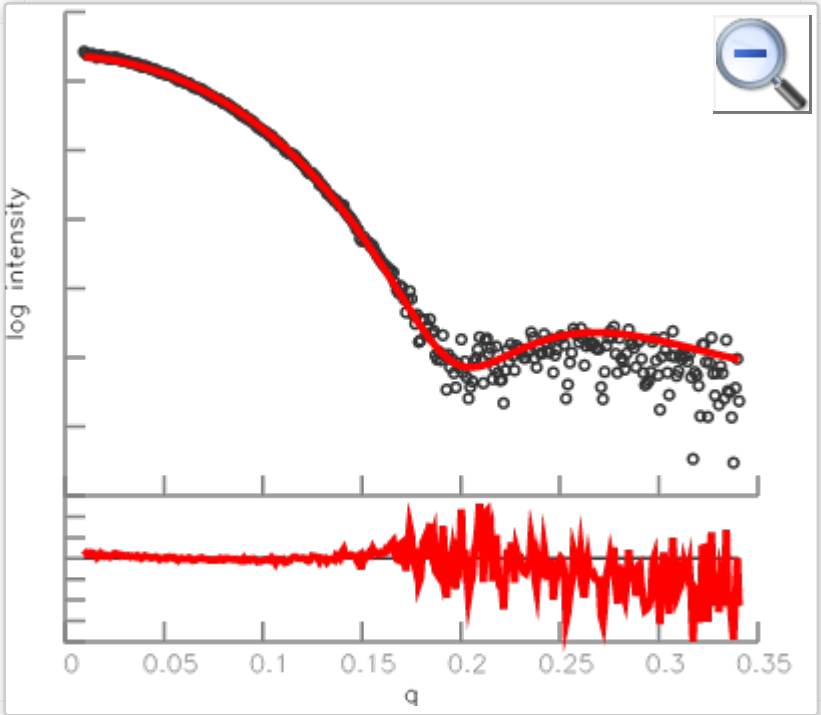
[4ds7_tr.pdb](#)

Profile file

[4ds7_tr_01C_S072_0_01.sub](#)

-

Can't see interactive display? Use [old interface](#)



JSmol

PDB file	show/hide	χ	c_1	c_2	R_g	# atoms	fit file
4ds7_tr	<input checked="" type="checkbox"/>	1.90	1.03	4.00	15.26	1269	4ds7_tr_4ds7_tr_01C_S072_0_01.dat

If you use FoXS, please cite:

D. Schneidman-Duhovny, M. Hammel, JA. Tainer, and A. Sali. Accurate SAXS profile computation and its assessment by contrast variation experiments. Biophysical Journal 2013.

D. Schneidman-Duhovny, M. Hammel, and A. Sali. FoXS: A Web server for Rapid Computation and Fitting of SAXS Profiles. NAR 2010.38 Suppl:W540-4

Contact: