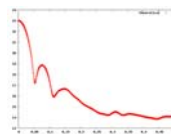




Fast SAXS Profile Computation with Debye Formula



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PDB files

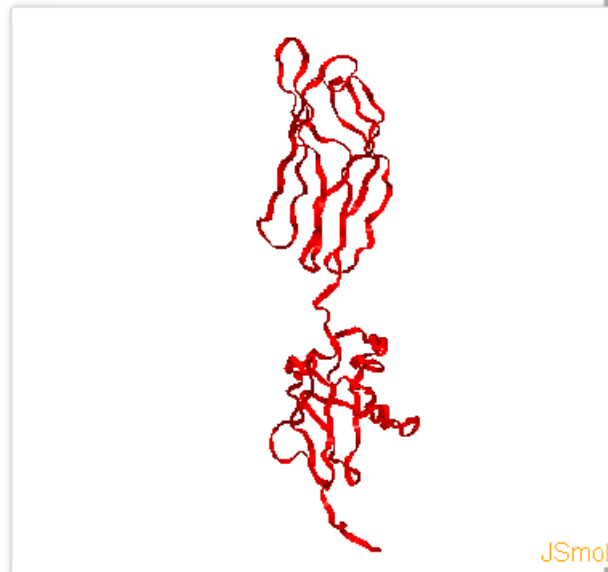
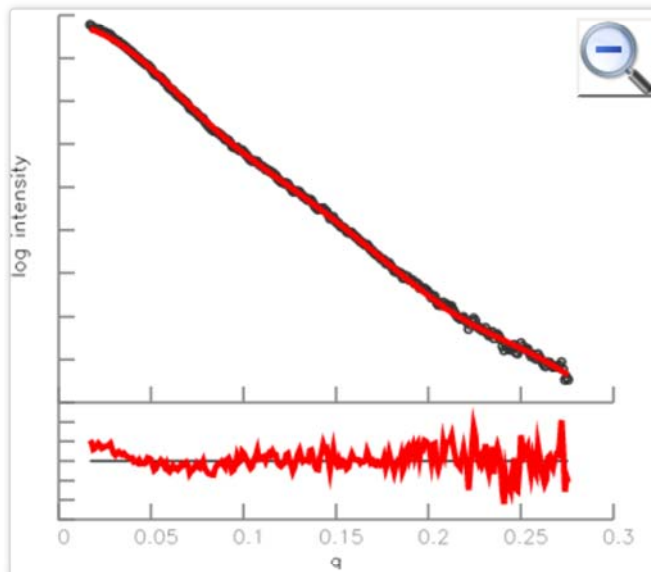
[26994.B99990039.pdb](#)

Profile file

[SAXS_26994_merged_q27.dat](#)

-

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



PDB file	show/hide	χ	c_1	c_2	R_g	# atoms	fit file
26994.B99990039	<input checked="" type="checkbox"/>	1.79	1.03	-0.81	25.73	1682	26994.B99990039 SAXS_26994_merged_q27.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: