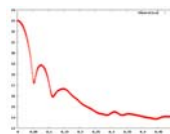




## Fast SAXS Profile Computation with Debye Formula



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PDB files

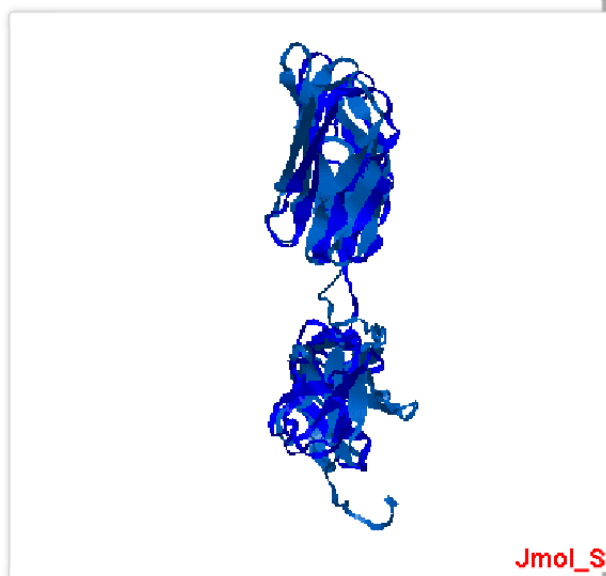
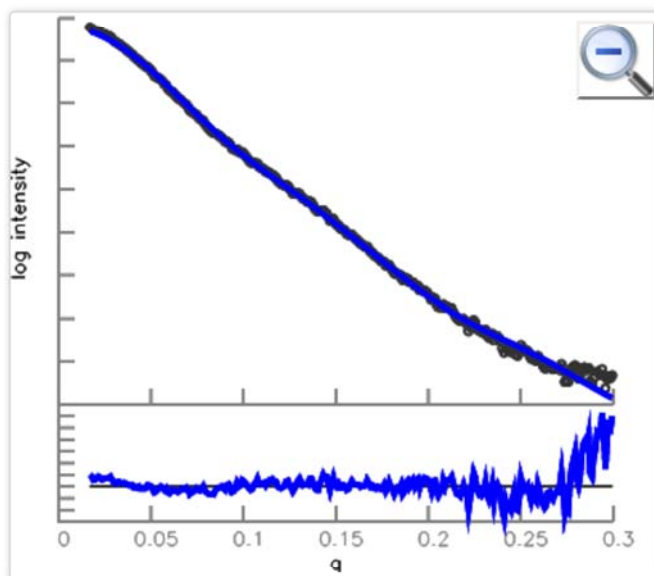
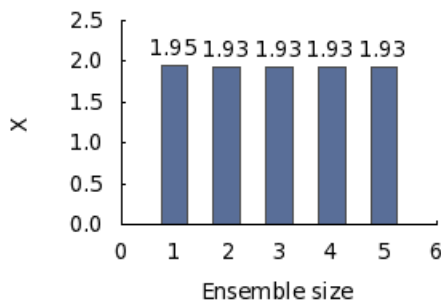
26994.zip

Profile file

[SAXS\\_26994\\_merged\\_q3.dat](#)

-

### MES (Minimal Ensemble Search) Fit



| Ensemble size | <input type="checkbox"/> show all/<br><input checked="" type="checkbox"/> hide all | χ    | models and weights  |      | fit file   |
|---------------|--|------|---------------------|------|--|
| 1             | <input type="checkbox"/>   | 1.95 | 26994.B99990039.pdb | 1.00 | <a href="#">26994.B99990039_SAXS_26994_merged_q3.dat</a> |
| 2             | <input checked="" type="checkbox"/>  | 1.93 | 26994.B99990088.pdb | 0.14 | <a href="#">avg_2</a>                                    |
|               |  |      | 26994.B99990039.pdb | 0.86 |  |
| 3             | <input type="checkbox"/>   | 1.93 | 26994.B99990039.pdb | 0.86 | <a href="#">avg_3</a>                                    |
|               |  |      | 26994.B99990088.pdb | 0.14 |  |
|               |  |      | 26994.B99990012.pdb | 0.00 |  |
| 4             | <input type="checkbox"/>   | 1.93 | 26994.B99990027.pdb | 0.00 | <a href="#">avg_4</a>                                    |
|               |  |      | 26994.B99990039.pdb | 0.86 |  |
|               |  |      | 26994.B99990088.pdb | 0.14 |  |
|               |  |      | 26994.B99990066.pdb | 0.00 |  |
| 5             | <input type="checkbox"/>   | 1.93 | 26994.B99990066.pdb | 0.00 | <a href="#">avg_5</a>                                    |
|               |  |      | 26994.B99990039.pdb | 0.86 |  |

|                     |      |
|---------------------|------|
| 26994.B99990088.pdb | 0.14 |
| 26994.B99990011.pdb | 0.00 |
| 26994.B99990074.pdb | 0.00 |

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If you use [MES](#), please cite:

Pelikan M, Hura GL, Hammel M. Structure and Flexibility within proteins as identified through small angle X-ray scattering. *General Physiology and Biophysics*. 2009;28:174 - 89.