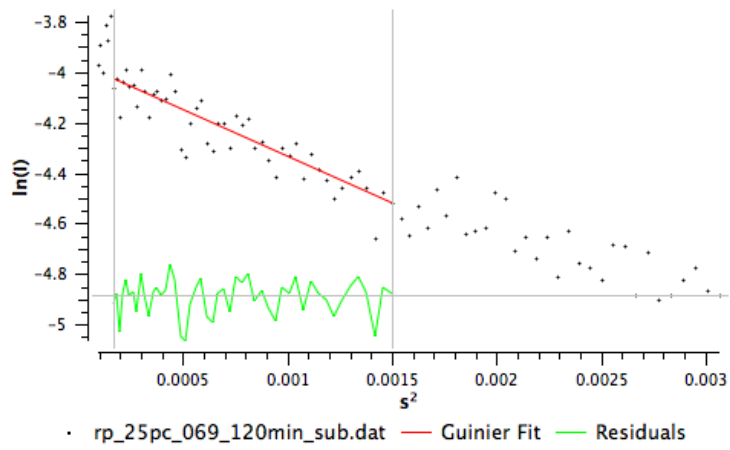
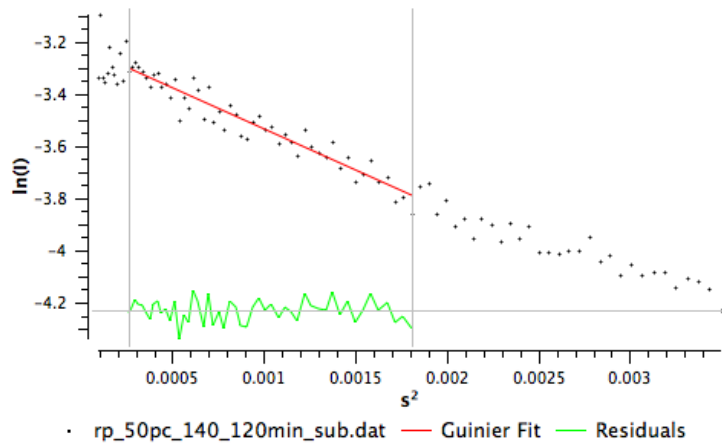


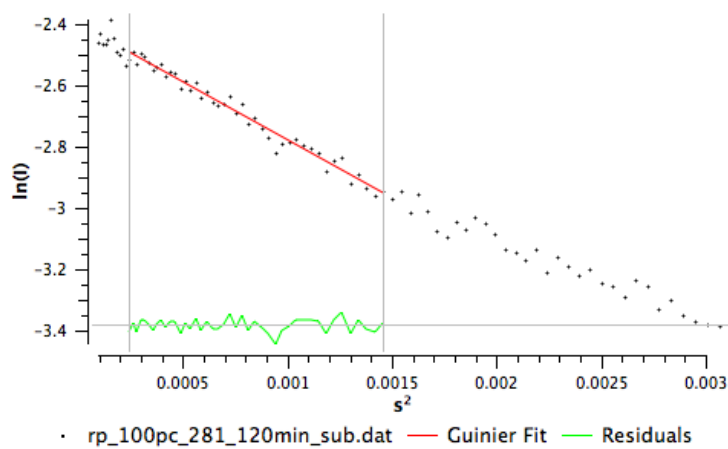
Guinier Plot



2.25mg/mL

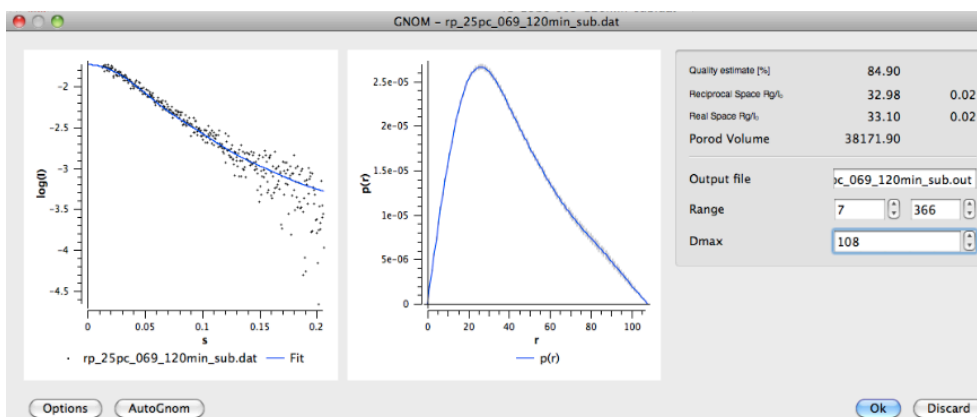


4.5mg/mL

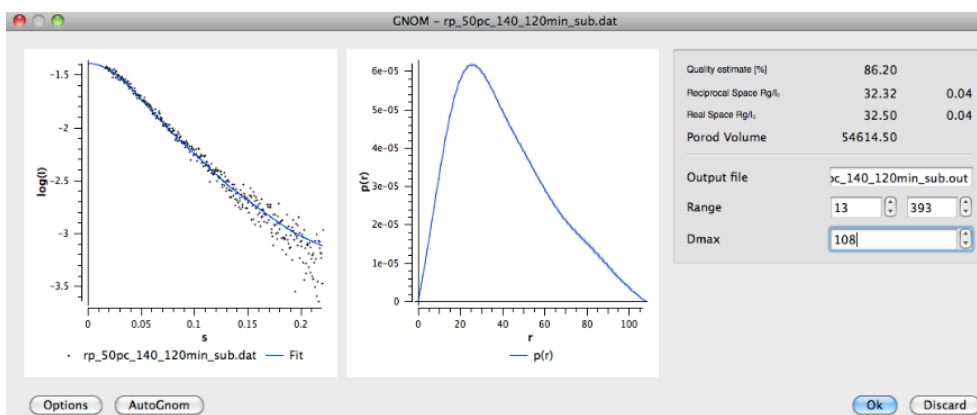


9mg/mL

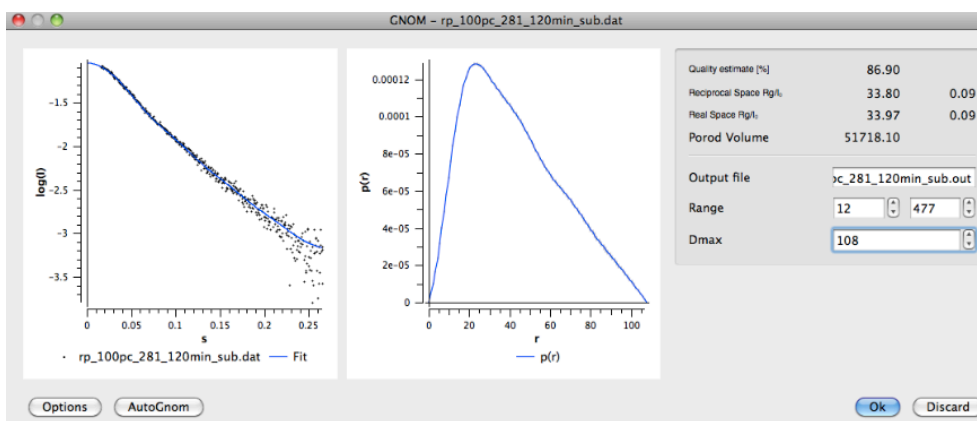
Pair-distance distribution



2.25mg/mL



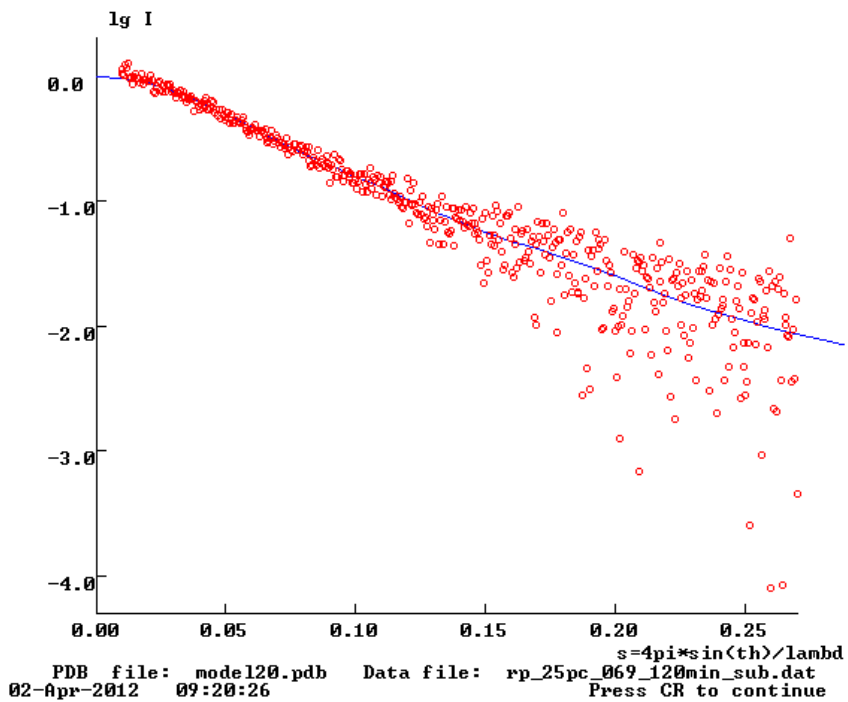
4.5mg/mL



9mg/mL

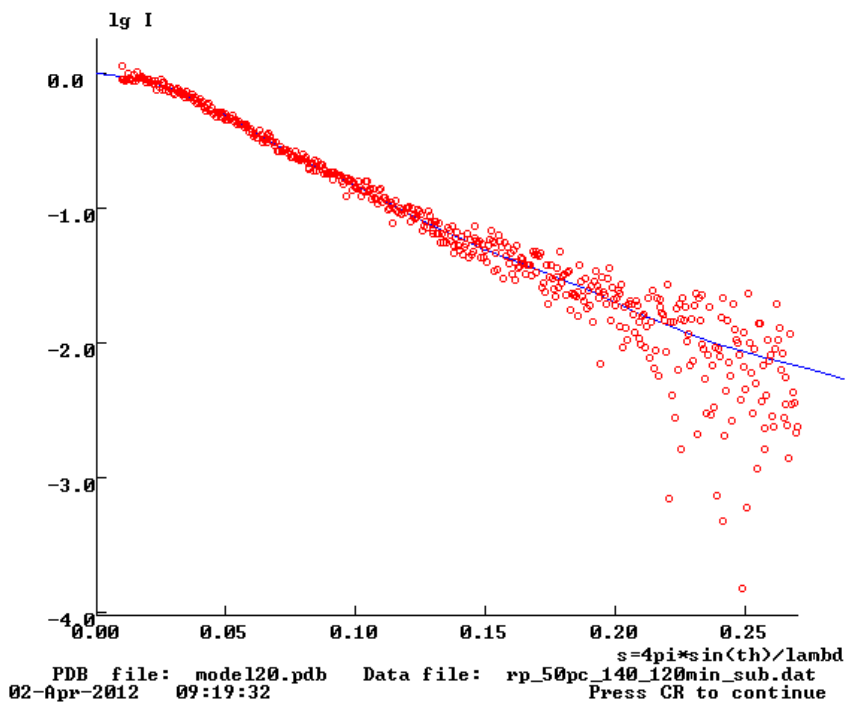
Crysol. Crystal structure: Dimer+DNA

Dro: 0.020 Ra: 1.800 RGE: 31.29 RGT: 29.07 Vol: 29732. Chi: 0.987



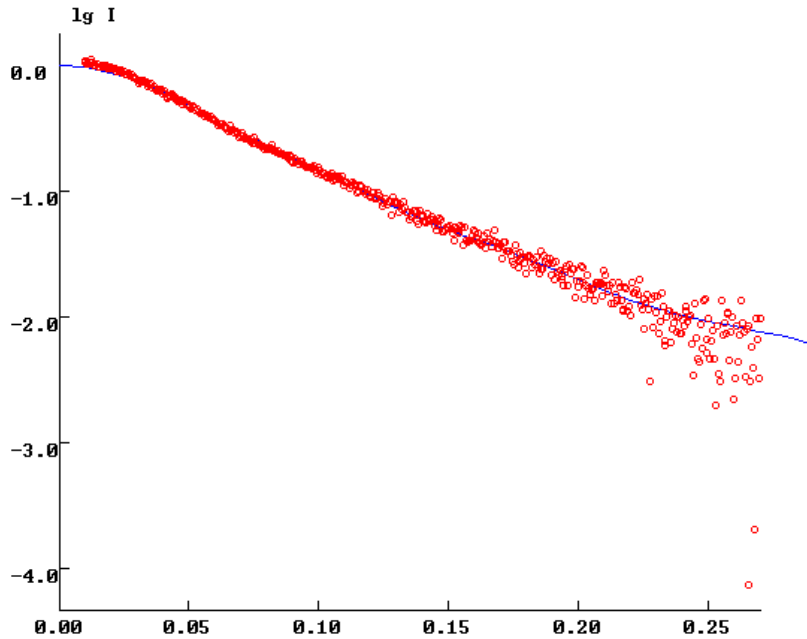
2.25mg/mL

Dro: 0.068 Ra: 1.400 RGE: 30.23 RGT: 29.50 Vol: 29732. Chi: 1.022



4.5mg/mL

Dro: 0.075 Ra: 1.400 RGE: 32.01 RGT: 29.31 Vol: 32081. Chi: 1.420

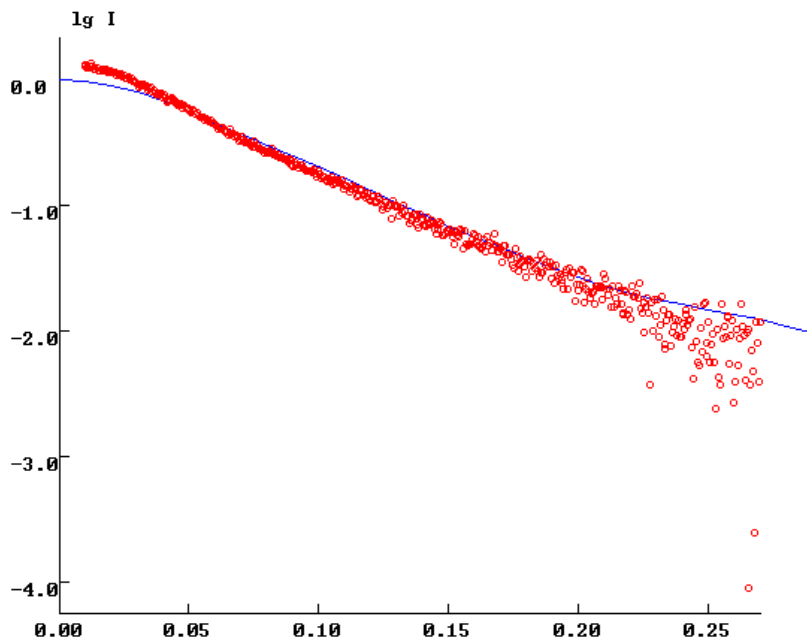


PDB file: MafA_refmt.pdb Data file: rp_100pc_281_120min_sub.dat
23-Apr-2012 13:58:01 s=4pi*sin<th>/lamdb
Press CR to continue

9mg/mL

Crysol. Model: Monomer+DNA (I deleted one protein chain in the .pdb file)

Dro: 0.075 Ra: 1.400 RGE: 31.23 RGT: 26.32 Vol: 20891. Chi: 3.201



PDB file: MafA_mono.pdb Data file: rp_100pc_281_120min_sub.dat
23-Apr-2012 15:58:59 s=4pi*sin<th>/lamdb
Press CR to continue

9mg/mL