

S1: Cremer-Pople analysis for the twenty 10 μ s explicit solvent molecular dynamics simulations.

For each simulation, the puckering parameters Q and θ are plotted against simulation time, θ and φ are plotted against each other and $\cos\langle\theta\rangle$ is plotted against simulation time (reports convergence of conformation).

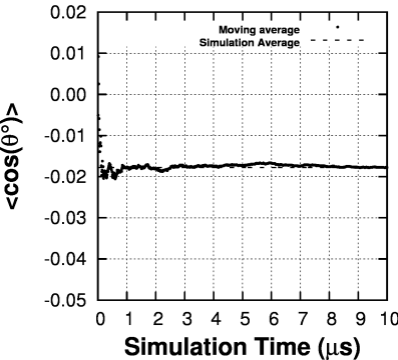
Key:

pages 2-41: α -anomers (1-OH), denoted alpha-D-X

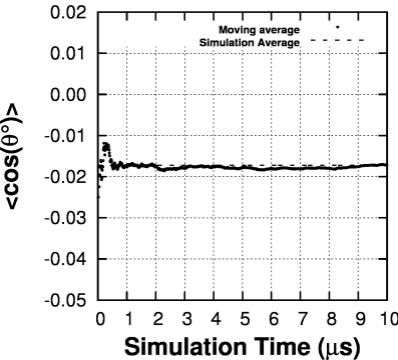
pages 42-81: 1-O-methyl glycosides

-1C4 postfix:¹C₄-initiated simulations, otherwise ⁴C₁-

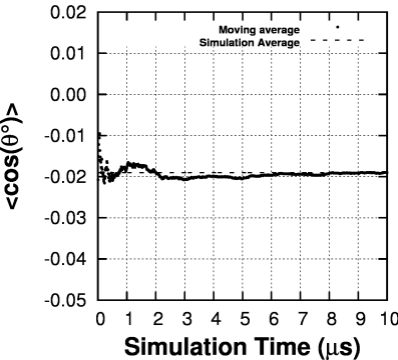
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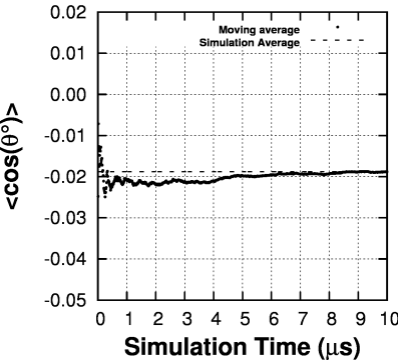
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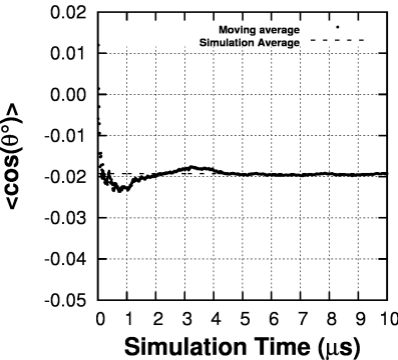
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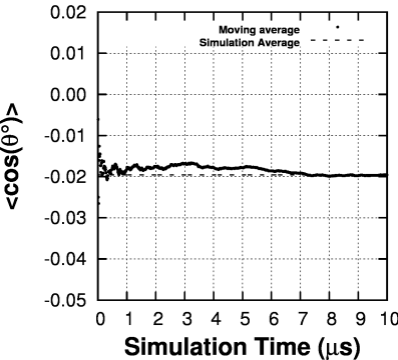
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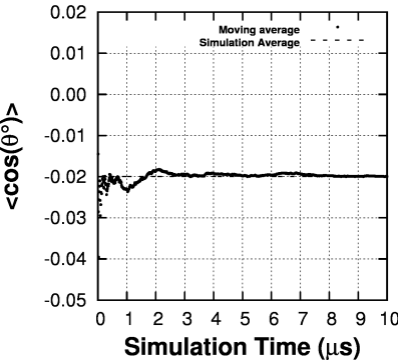
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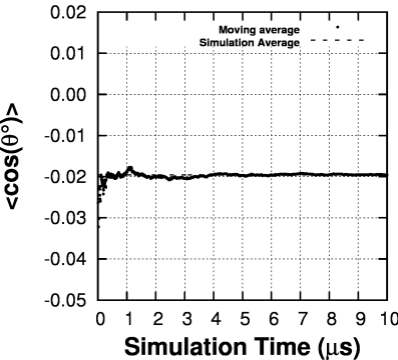
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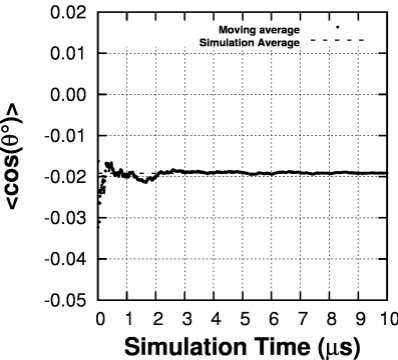
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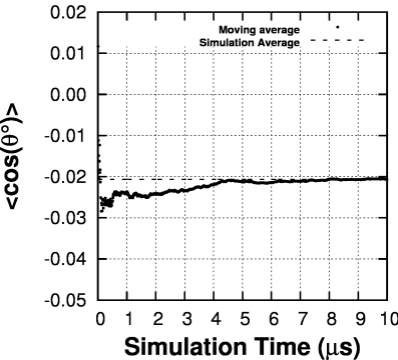
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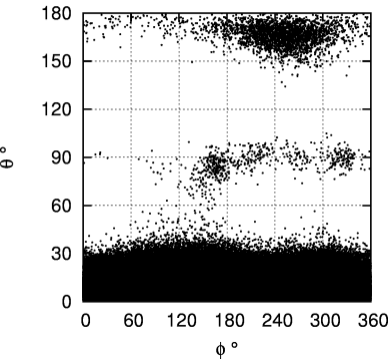
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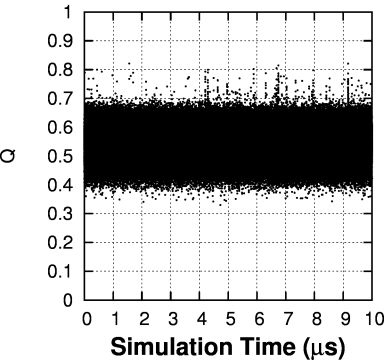
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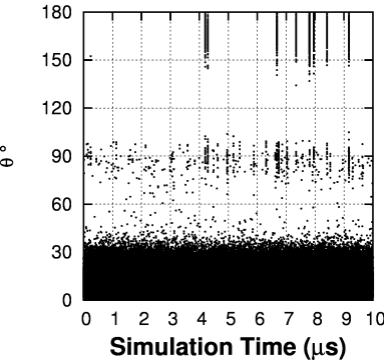
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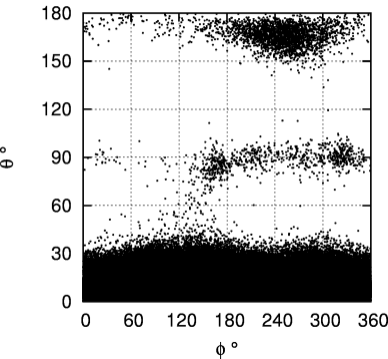
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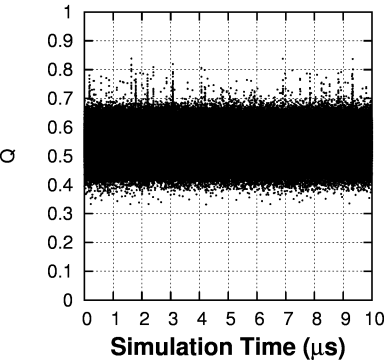
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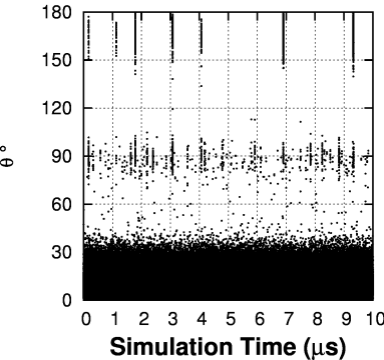
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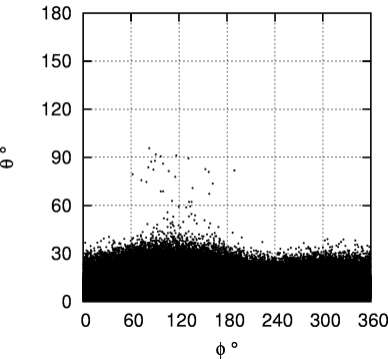
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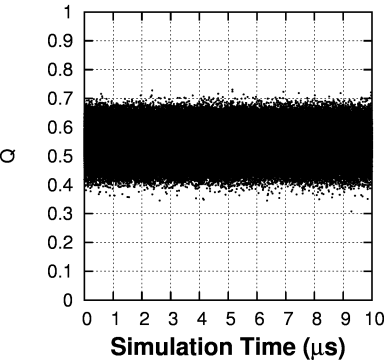
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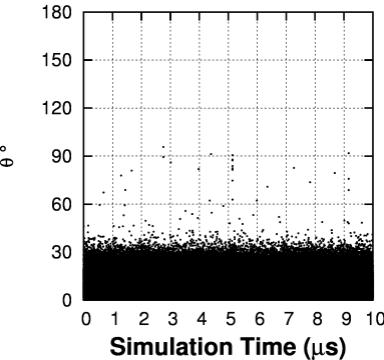
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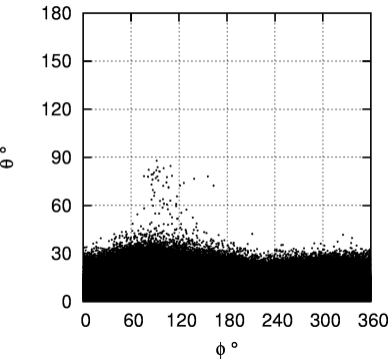
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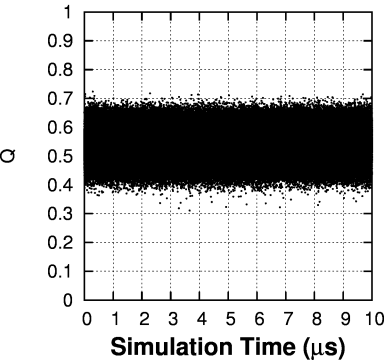
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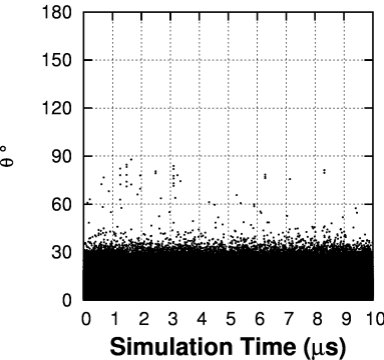
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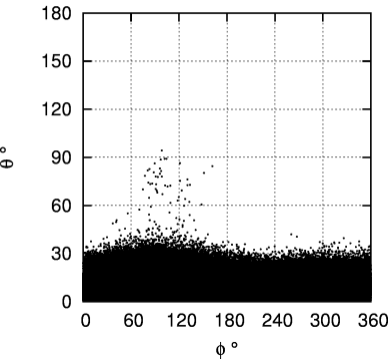
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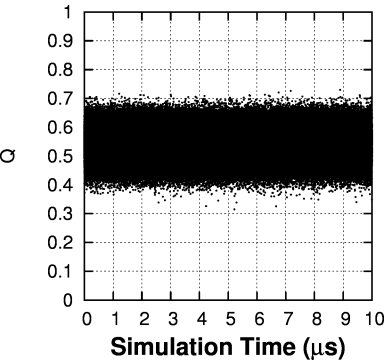
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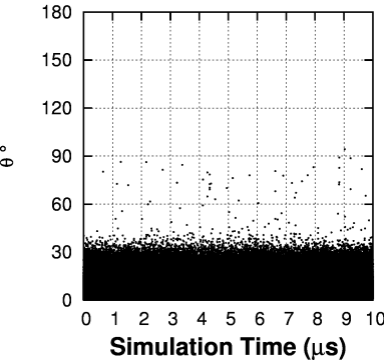
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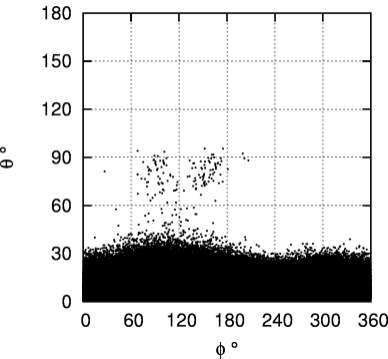
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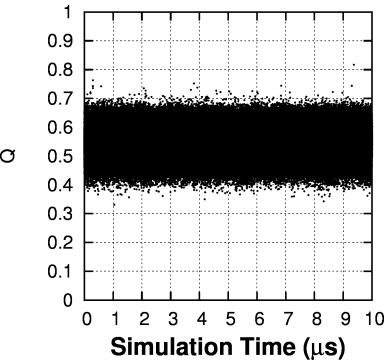
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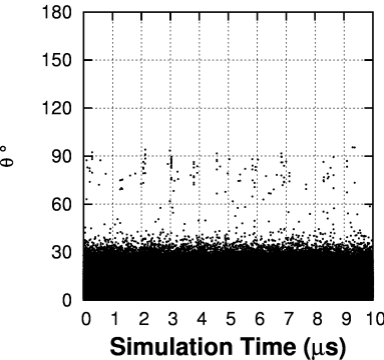
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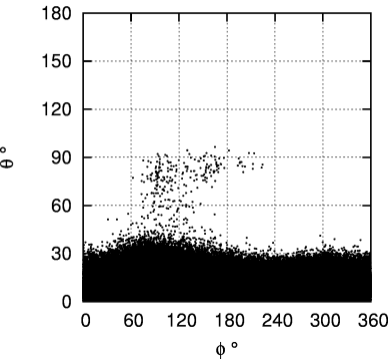
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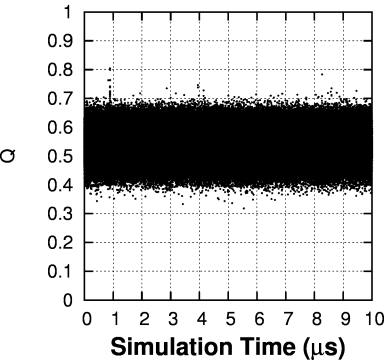
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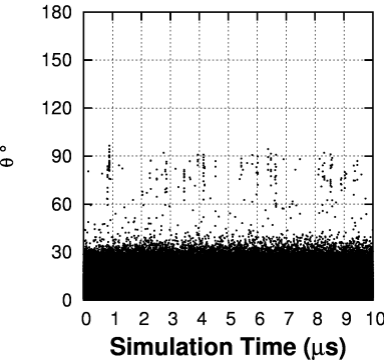
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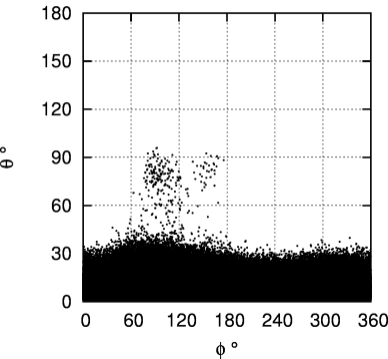
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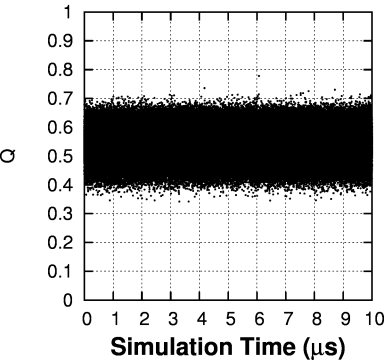
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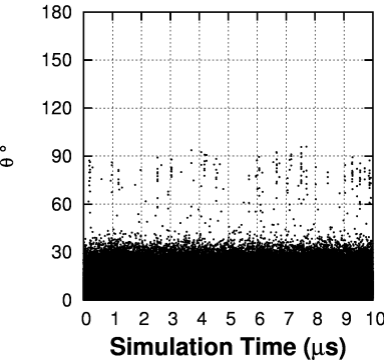
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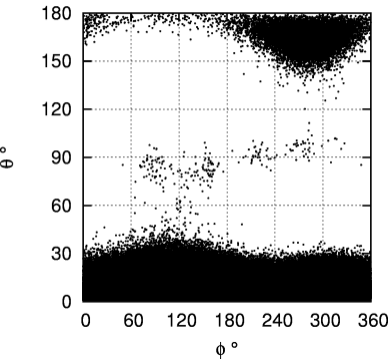
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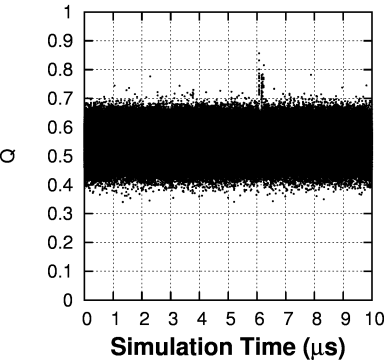
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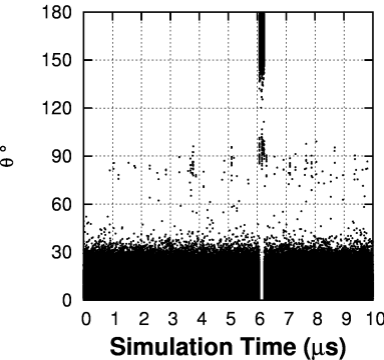
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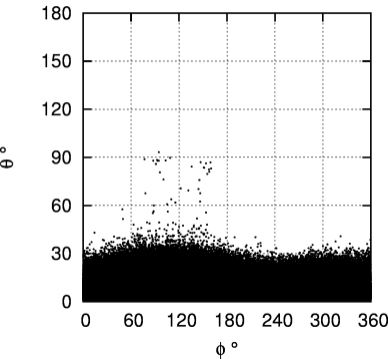
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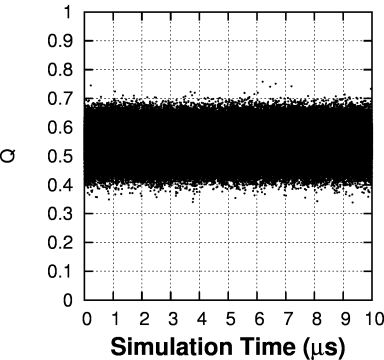
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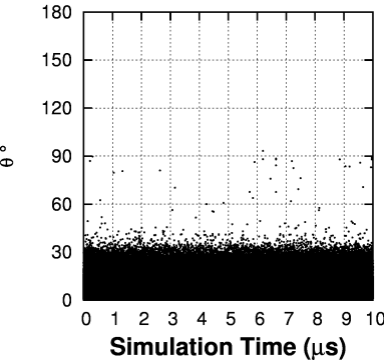
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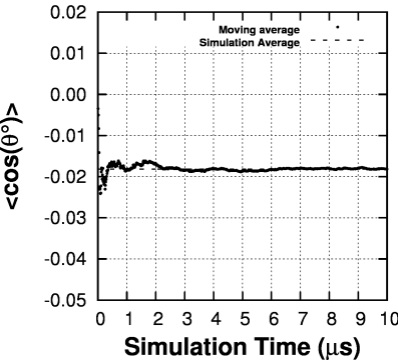
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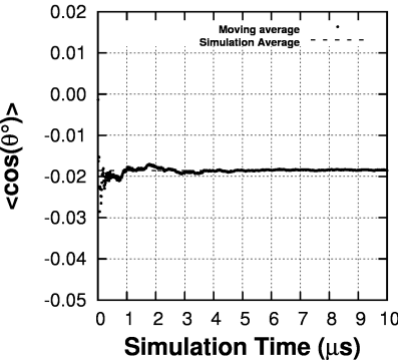
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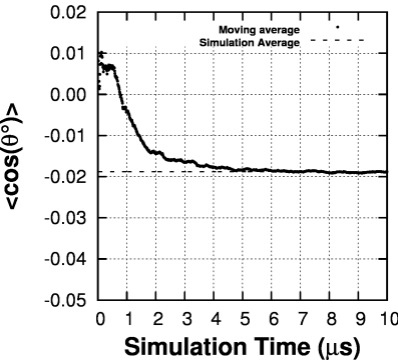
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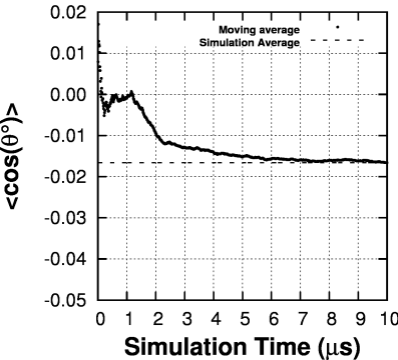
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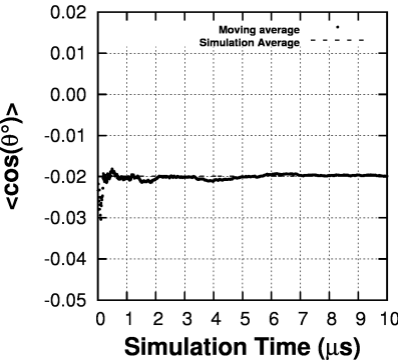
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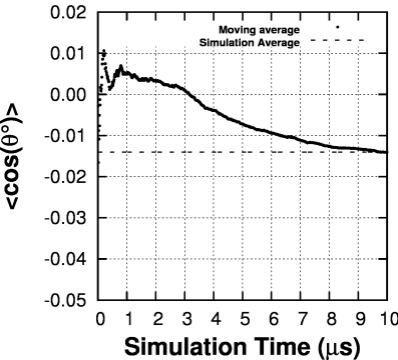
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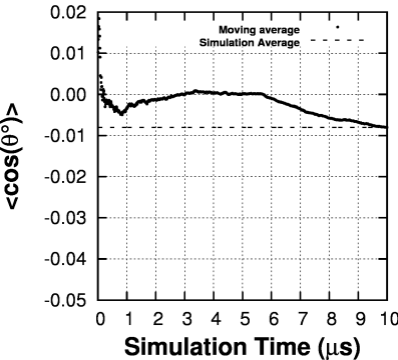
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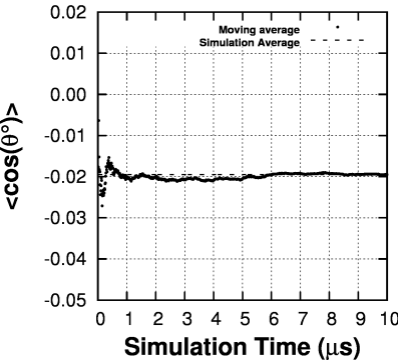
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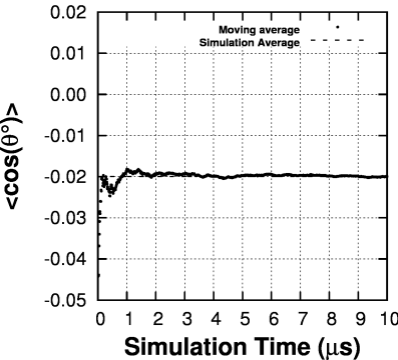
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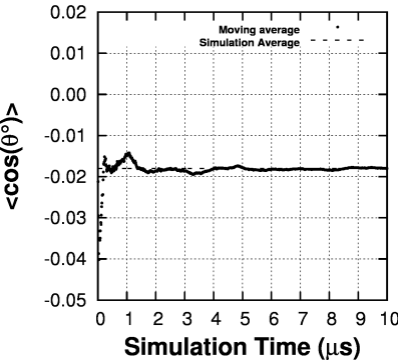
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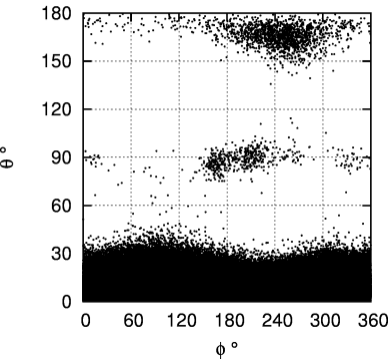
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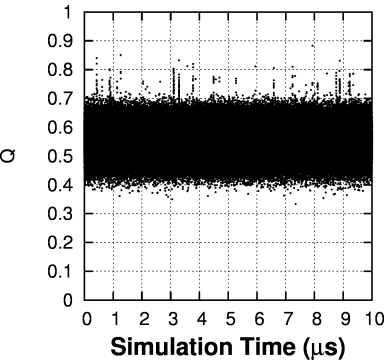
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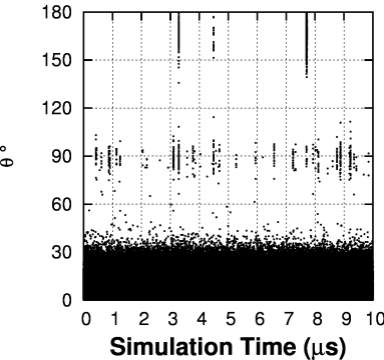
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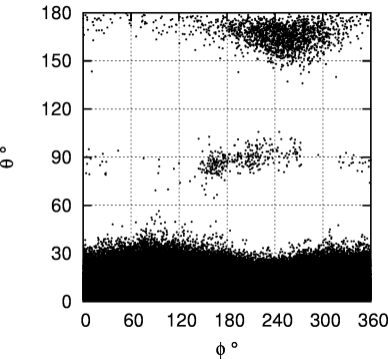
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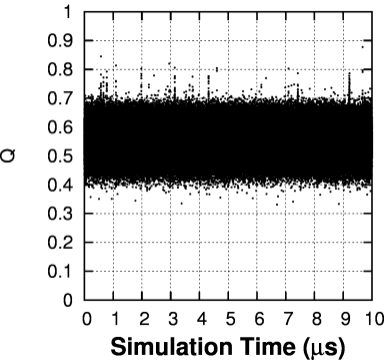
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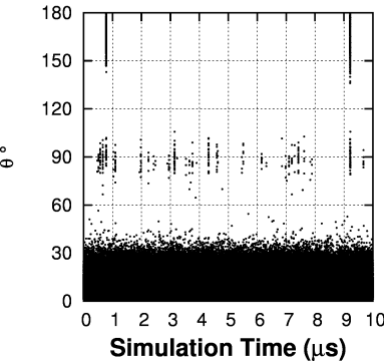
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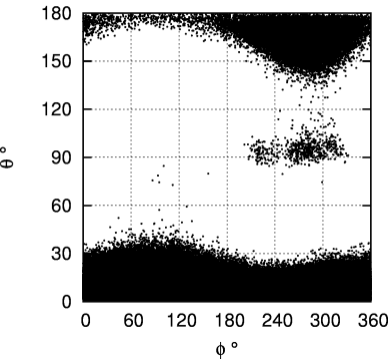
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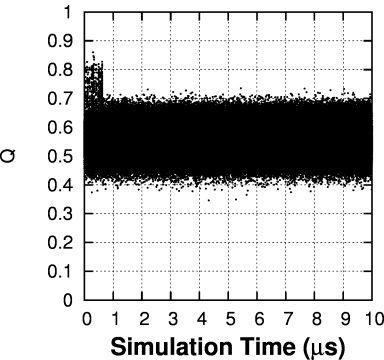
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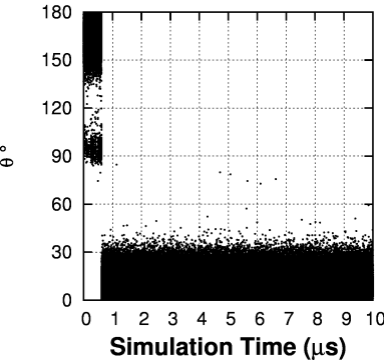
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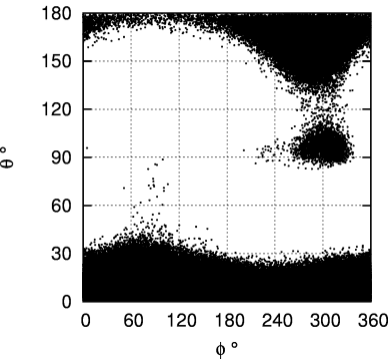
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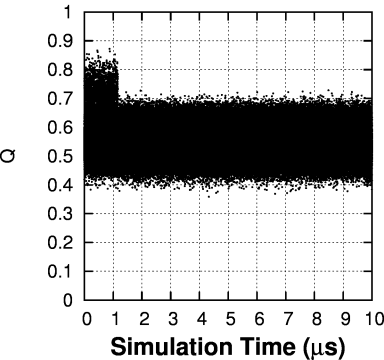
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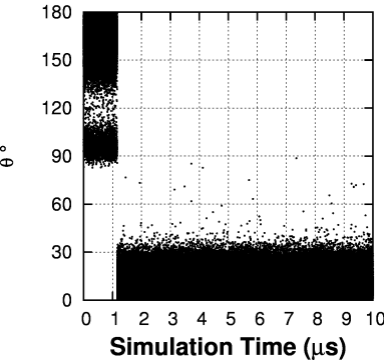
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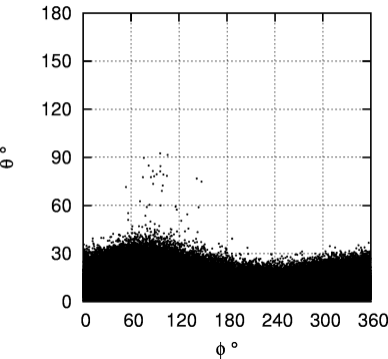
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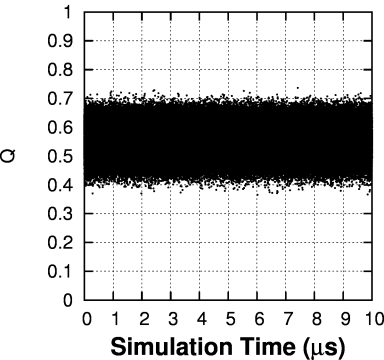
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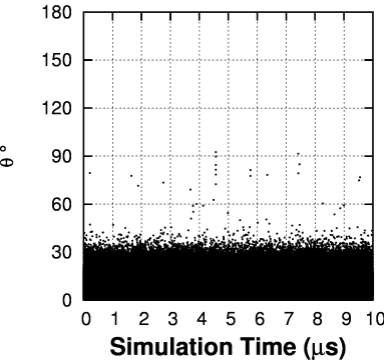
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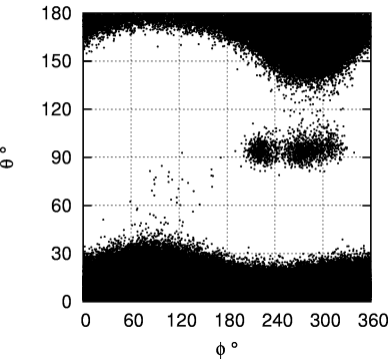
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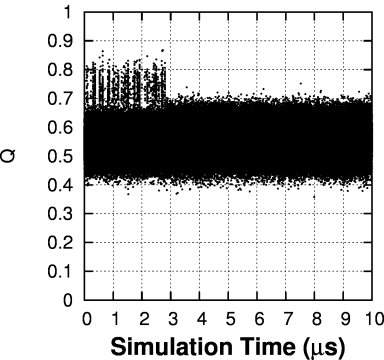
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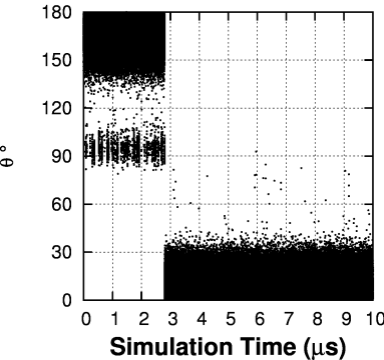
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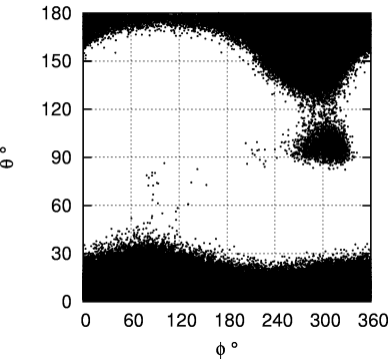
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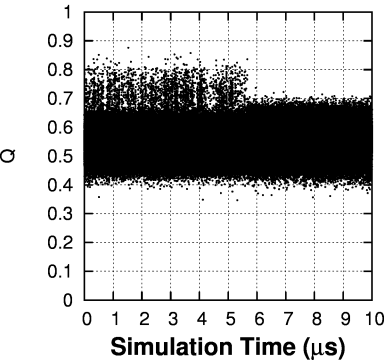
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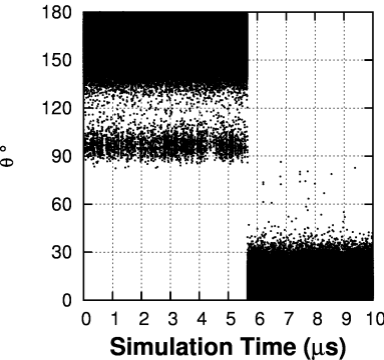
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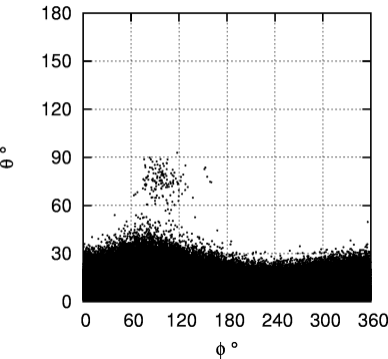
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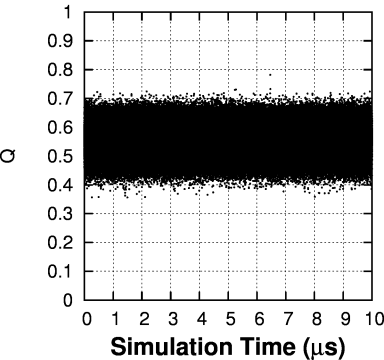
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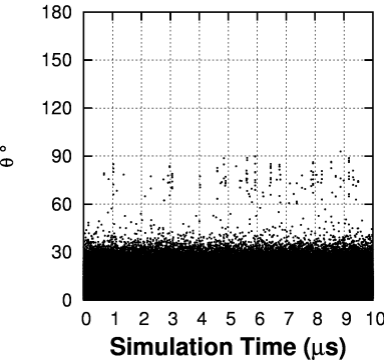
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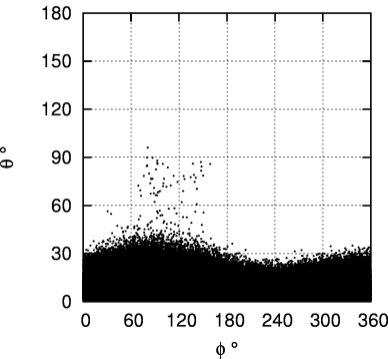
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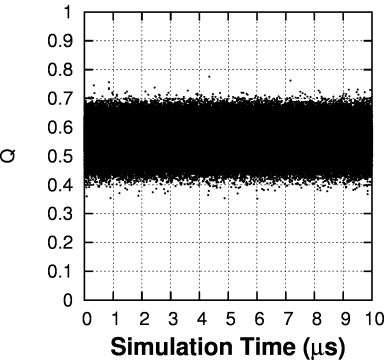
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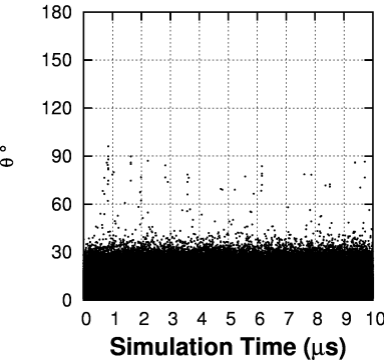
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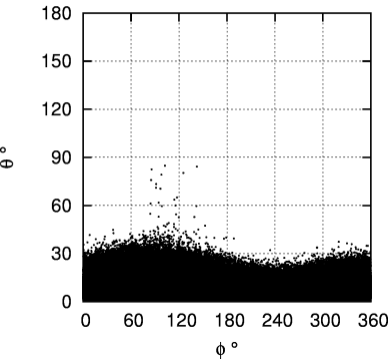
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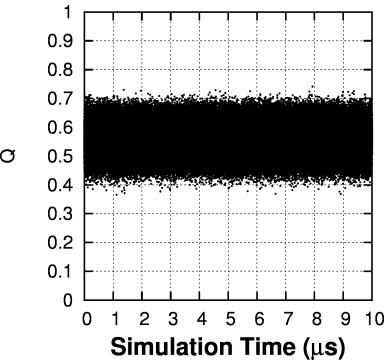
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glcNS



glcNS



glcNS

