ADDITIONS AND CORRECTIONS

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William L. Jorgensen* and Julian Tirado-Rives*: Monte Carlo vs Molecular Dynamics for Conformational Sampling

The units for the computer timings should have been minutes rather than seconds. No qualitative changes are needed.

Page 14510. The calculations required 95.4 minutes of CPU time for 10.0 ps of MD or for 1.63×10^6 configurations of MC.

Page 14512. From the behavior of the torsional energy profiles and conformer populations over the full runs, the roughly 10×10^6 configurations of MC needed for conformational convergence required 584 minutes, while the corresponding roughly 100 ps of MD required 954 minutes or a factor of 1.63 more time. From the periods for the *trans* populations to decay from 100% to 80%, the MC20 calculation required 97 minutes, while the MD run needed 370 minutes or a factor of 3.81 more time.