On Assignment 6, Part 1

- Q. What's pos.d?
- A. Your pdf.cu will open pos.d, and compute pair distribution function for the atomic positions in it; after compilation nvcc -o pdf pdf.cu

the executable pdf & input data pos.d must be placed in the same directory as the Slurm script pdf.sl

Big picture: Make the doubly-nested big for loops into a kernel

On Assignment 6, Part 1

- Q. What's the email on Friday?
- A. Addendum to Friday's lecture: I forgot to add the last piece of change to avoid race condition (slide 9 in http://cacs.usc.edu/education/cs596/CUDA-PDF.pdf)

End of the doubly-nested atom loops

- Q. Should I worry about the warnings: variables dr, rij, i, j were declared but never referenced?
- A. No. I didn't bother to remove their declaration, when moving the main computation of histogram() to kernel. They will do no harm, or remove their declarations at the beginning of histogram().

On Assignment 6, Part 1

- Q. What to plot?
- A. Pair distribution function g(r) (right column) vs. atomic-pair distance r (left column) that will be output into pdf.d by your pdf.cu.

Use the force! Reduced variance estimators for densities, radial distribution functions, and local mobilities in molecular simulations •

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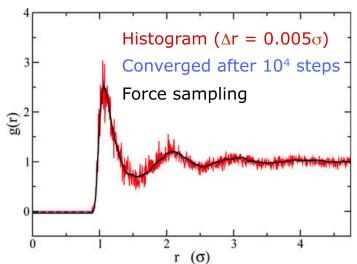




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http://cacs.usc.edu/education/cs596/Rotenberg-UseTheForce-JCP20.pdf