Protein folding results

- “Slow” anneal, 4 million steps! Fast anneal 1/2 as many
- Still obviously could do better!
- Temperature decreased from $T=2$ to $T=0$
Average energy

- Some equilibration at each temperature before averaging
- Slower annealing results in lower energy
Average length

End-to-end length vs. Temperature
Final structure from “slow” anneal