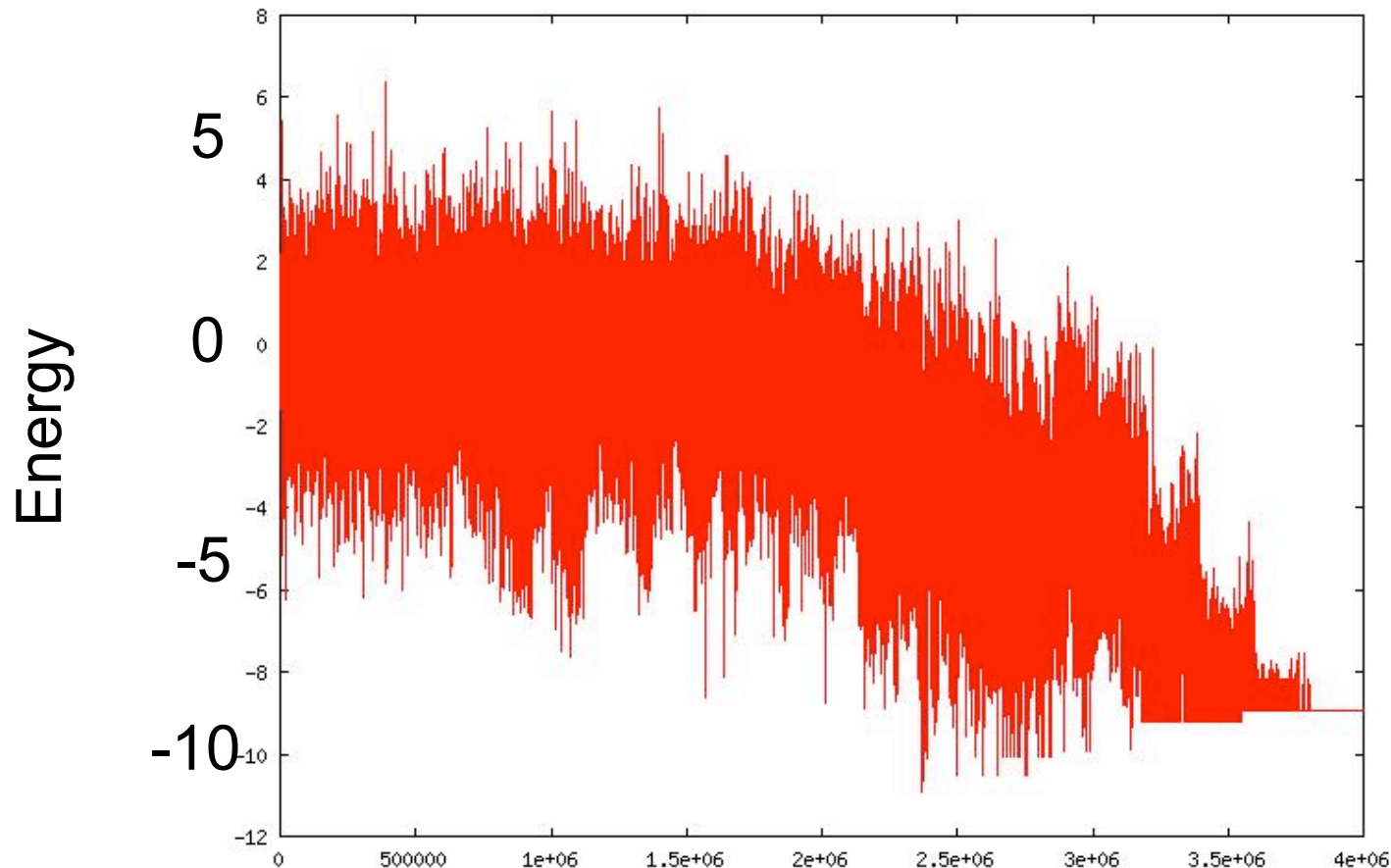
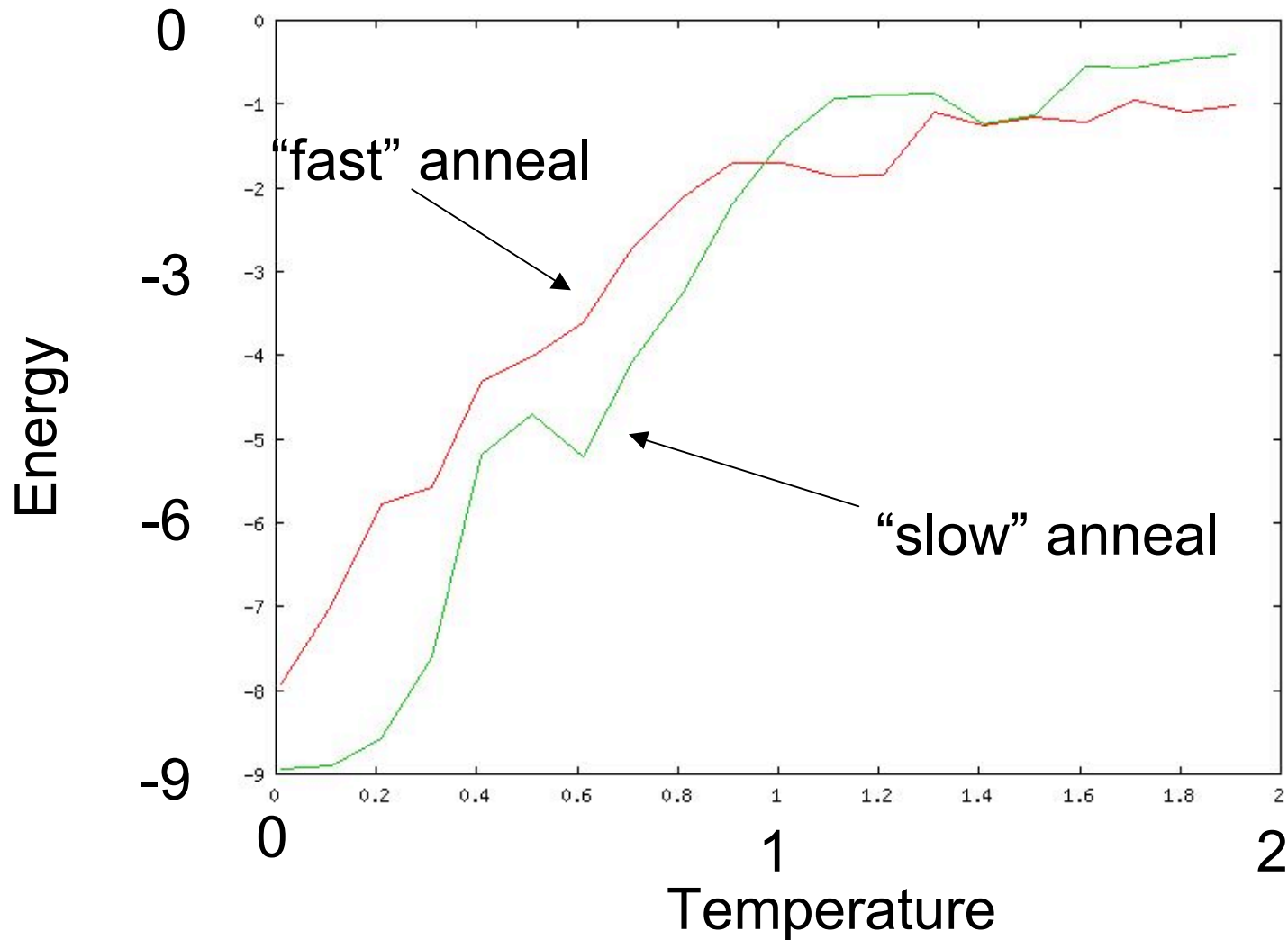


Protein folding results



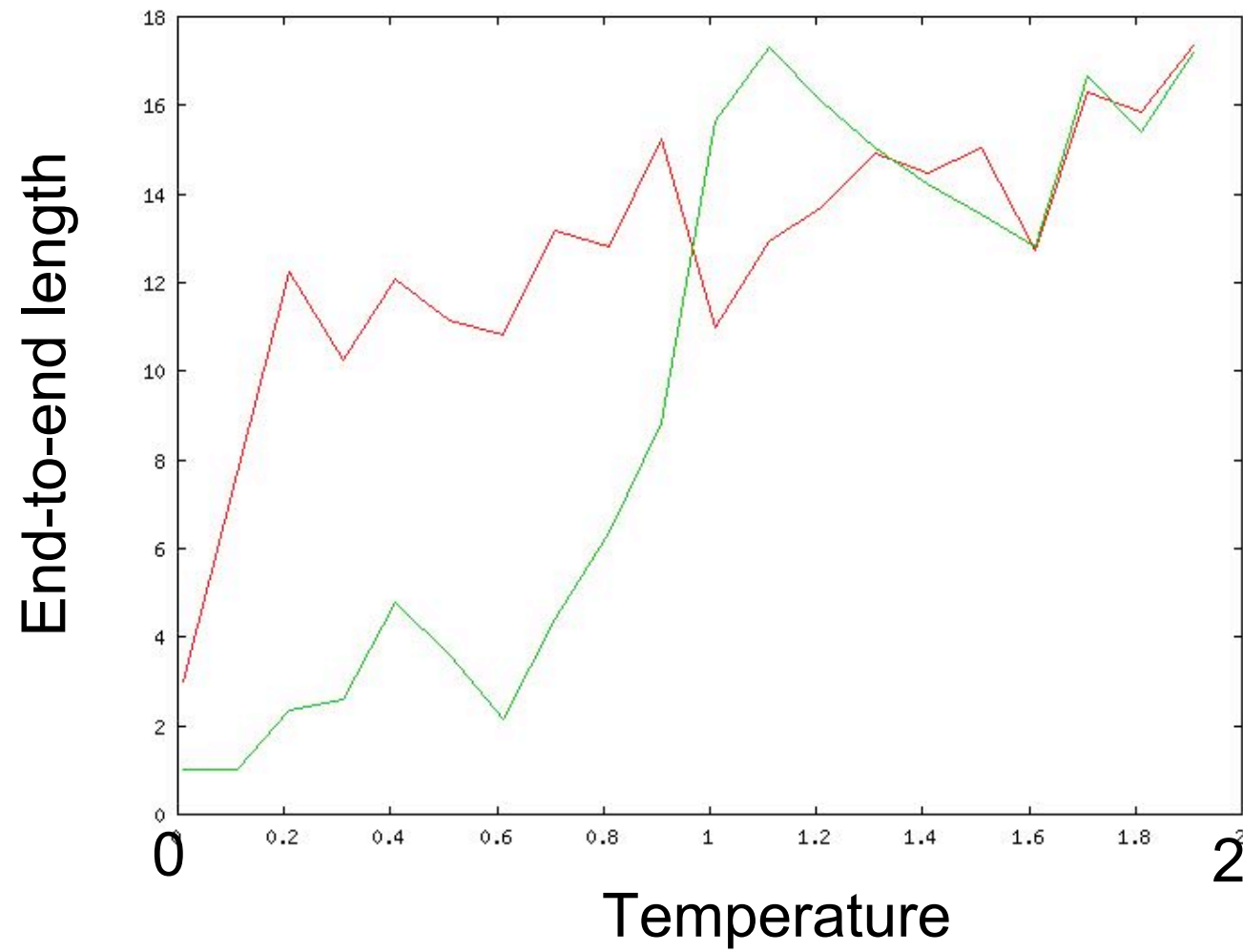
- “Slow” anneal, 4 million steps! Fast anneal 1/2 as many
- Still obviously could do better!
- Temperature decreased from $T=2$ to $T=0$

Average energy



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

Average length



Final structure from “slow” anneal

