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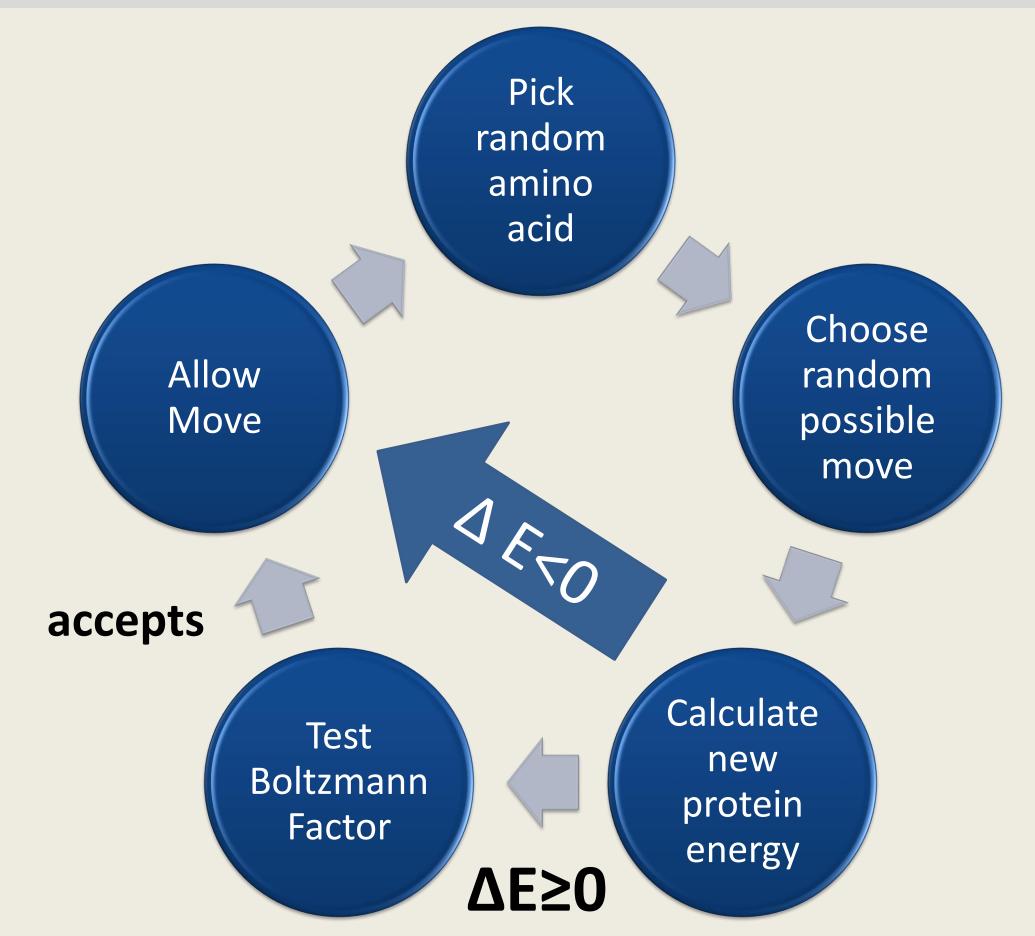
Protein Folding & Self-Organized Criticality

Arun Bajracharya; Faculty Advisor: Joelle Murray Linfield College, McMinnville, OR

INTRODUCTION

Proteins are known to fold into tertiary structures that determine their functionality in living organisms. However, the complex dynamics of protein folding and the way they consistently fold into the same structures is unknown. Self-organized criticality (SOC) has provided a framework for understanding complex systems in various scientific disciplines through scale invariance and the associated "fractal" power law behavior. In this research, we use a simple hydrophobic-polar lattice-bound computational model to investigate self-organized criticality as a possible mechanism for generating complexity in protein folding.

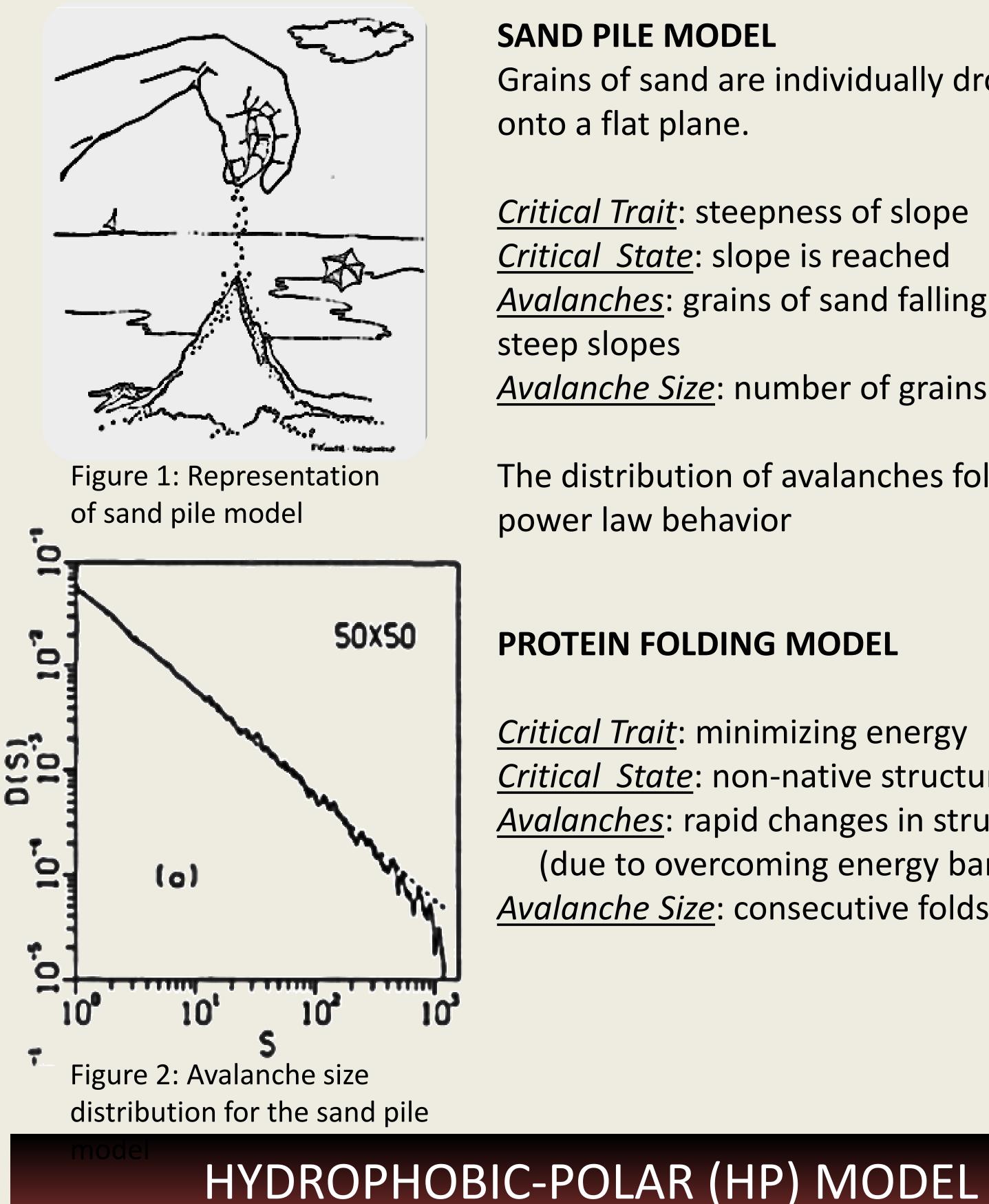
ALGORITHM



SELF-ORGANIZED CRITICALITY (SOC)

SOC system is a dynamical system at a critical state that is characterized by power law behavior. It follows that these systems are both time and scale invariant.

The Sand Pile Model is the first and most well known example of a selforganizing system.



SAND PILE MODEL Grains of sand are individually dropped onto a flat plane.

<u>Critical Trait</u>: steepness of slope <u>Critical State</u>: slope is reached Avalanches: grains of sand falling from steep slopes Avalanche Size: number of grains falling

Figure 4: Flow diagram representing the computation sequence of the folding process

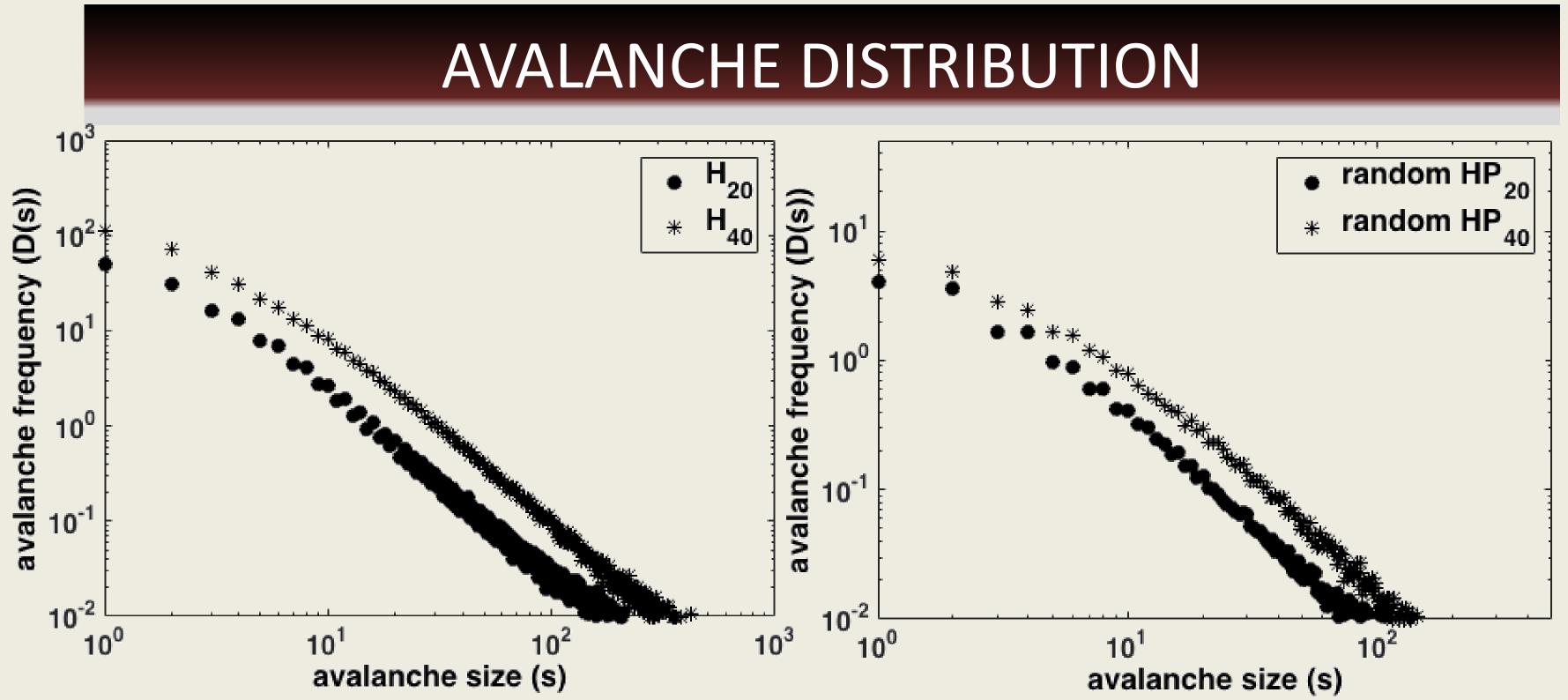


Figure 5: Avalanche distributions for 20 and 40 amino acid chain. Since SOC systems follow a power law, an avalanche distribution on a log plot should show a straight line

The distribution of avalanches follow a power law behavior

PROTEIN FOLDING MODEL

<u>Critical Trait</u>: minimizing energy

Critical State: non-native structure

Avalanche Size: consecutive folds

Avalanches: rapid changes in structure

(due to overcoming energy barriers)

• 20% * 50% (D(s)) + 80% ♦ 0% 10⁰ 10⁻¹ 10^{3} 10^{0} 10¹ avalanche size (s)

Figure 6: Stopping avalanches to inspect SOC resilience

DISCUSSION

Avalanche stopping alters SOC

characteristics

FUTURE RESEARCH

 Avalanche distributions in figure 5 Does the model exhibit other SOC \bullet

Simplified model for understanding how a protein

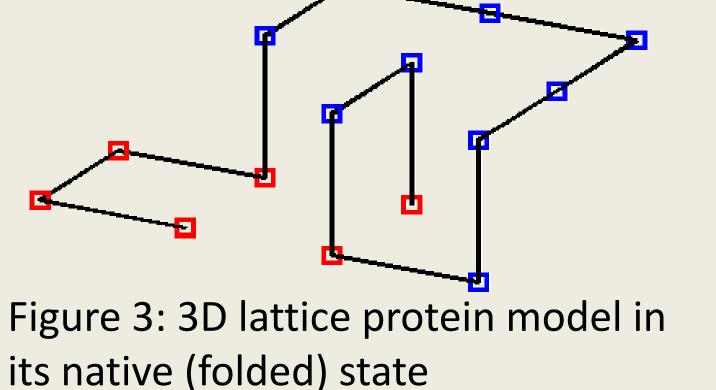


AVALANCHE STOPPING

- Individual folds are randomly prevented throughout the folding process
 - What impact does random stopping have on the avalanche distribution?

folds in space.

•3D lattice: 1 bond length, no stretching •HP model: 2 types of amino acidshydrophobic (H) and polar (P) •One amino acid moves at a time



exhibits power law, although this feature alone doesn't support the presence of SOC

features? For longer protein chains?

Do more sophisticated models exhibit SOC?

Is tertiary structure connected to SOC?

¹P. Bak, C. Tang, and K. Wiesenfeld, Phys. Rev. Lett. **59**, 381 (1987) K. A. Dill, Polym. Prepr. Am. Chem. Soc. Div. Polym. Chem. 36,) H. Frauenfelder, Proc. Natl. Acad. Sci. U. S. A. **99 Suppl 1**, 2479 (2635 (1995002)