# Estimating Evolutionary Parameters for Protein Low Complexity Regions using an Approximate Bayesian Computation

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## Overview

Background information
Research Questions/Explorations
Experimental Approach
Results
6 1 1 1 5 11 1
Conclusion and Future Work

## What are LCR's?

#### Saccharomyces cerevisiae SRP40 Protein LCRs

- >CAA82171.1(298-316) complexity=2.18 (15/1.90/2.20) tpassnestpsasssssan

## Shannon's Entropy - MAYBE

$$H = -L \sum p_i log_2(p_i)$$

# LCR's Present in Unique Ways

## Homorepeats

Consecutive iterations of a single residue



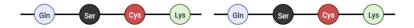
## Direpeats

Consecutive iterations of two ordered, different residues



## Tandem Repeats

Sequence of residues which are repeated a number of times



# LCR's are Hypermutable

# Proposed Mechanisms of LCR Evolution

## Why Care about LCRs and Their Evolution?

# What will this Study Explore?

# What Approach will be Taken?

# Why use an ABC-MCMC

## MCMC for ABC

- $\bullet \ \, \text{Propose a move from } \theta \text{ to } \theta' \text{ according to a transition kernel } q(\theta,\theta').$
- ② Generate simulated dataset D' using  $\theta'$  and calculate S'.
- **3** If  $\rho(S', S) \le \epsilon$  continue to 4, otherwise remain at  $\theta$  and go to 1.
- Calculate

$$\alpha(\theta, \theta') = min(1, \frac{\pi(\theta')q(\theta', \theta)}{\pi(\theta)q(\theta, \theta')})$$

- **5** Accept  $\theta'$  with probability  $\alpha$ , otherwise stay at  $\theta$ .
- Return to 1.

## ABC-MCMC Algorithm - MY VERSION

- 1
- (3)
- 4
- 5

# Simulation Step

## Results

# Conclusion/Future Work

## Blocks in Beamer

#### Standard Block

This is a standard block.

## Alert Message

This block presents alert message.

An example of typesetting tool

Example: MS Word, LATEX