

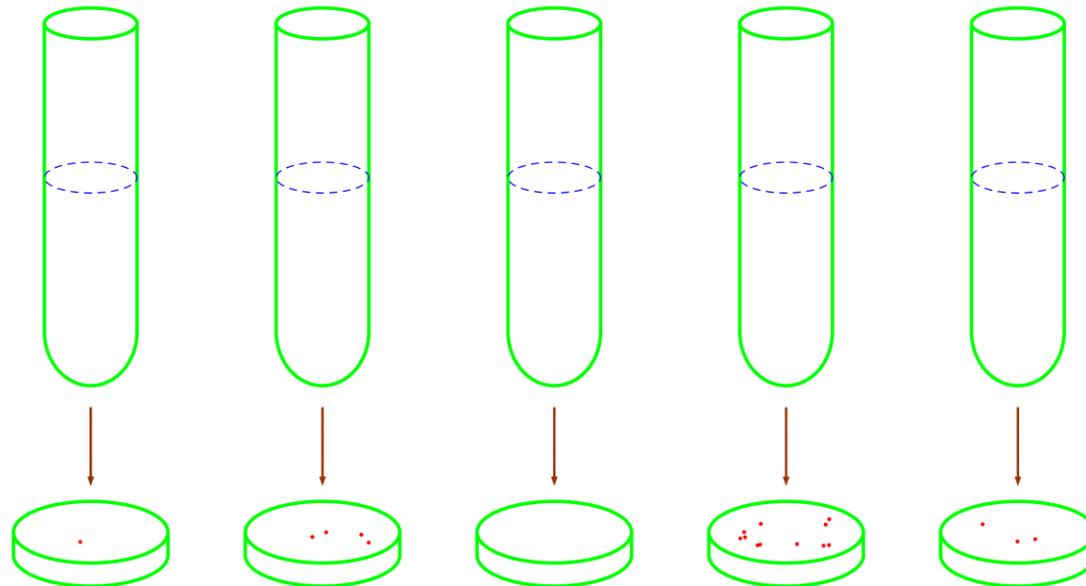
Simulating Chromosome Segregation

– Qi Zheng



What motivated this study?

- The fluctuation experiment of Luria and Delbrück
- Growing wild type cells in tubes and let mutation occur
- Selecting mutants (e.g., drug resistant mutants) on plates
- Computing a mutation rate from the number of mutants





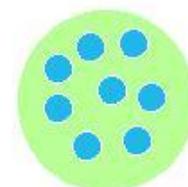
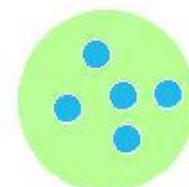
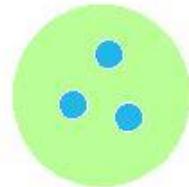
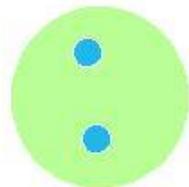
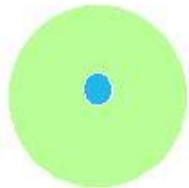
Simulating Chromosome Segregation

Qi Zheng

Department of Epidemiology & Biostatistics, Texas A&M University

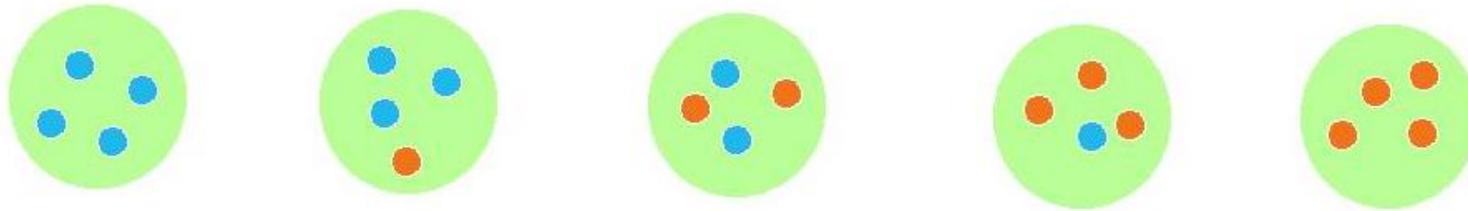
Why care about polyploidy?

- A cell may have more than one chromosome
- A cell has a ploidy value of k if it has k chromosomes
- The classic Luria-Delbrück protocol assumes $k=1$
- Polyploidy complicates the estimation of mutation rates
- So we need to take into account the ploidy value when calculating mutation rates
- The following cells have a ploidy value of 1, 2, 3, 5 and 8



What is the key issue?

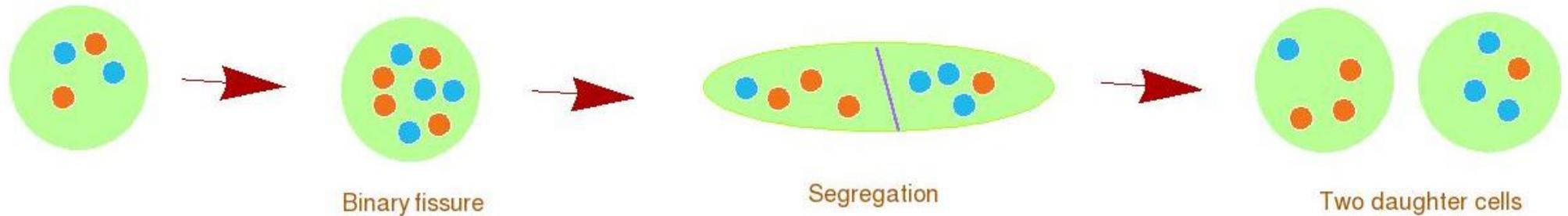
- A cell having k mutated chromosomes is called a type k cell
- With ploidy value = 4, there are 5 types of cells: type 0, type 1, ..., type 4.



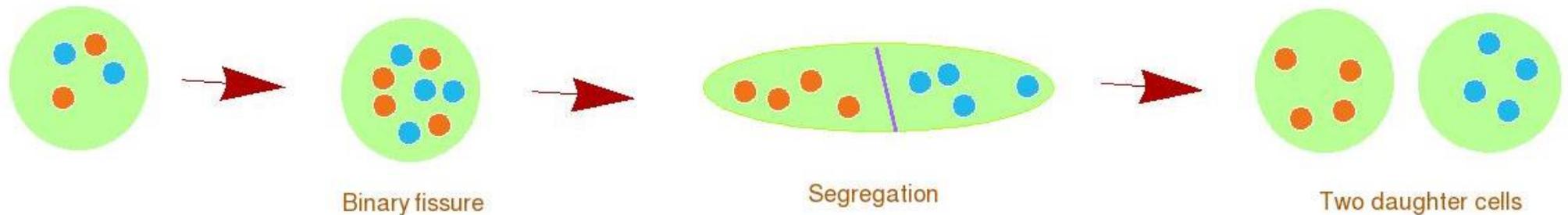
- If a cell population starts with a type 1 cell, it may eventually have all possible types of cells
- With a small probability μ , a wild type chromosome may generate a mutated daughter chromosome
- How does a cell population evolve starting with a type 1 cell?



What is random chromosome segregation?



- In this example cells have a ploidy value of 4
- 8 daughter chromosomes are randomly assigned to 2 daughter cells
- A type 2 cell may lead to a type 3 cell and a type 1 cell
- But there are other possibilities, e.g.:





Simulating Chromosome Segregation

Qi Zheng

Department of Epidemiology & Biostatistics, Texas A&M University

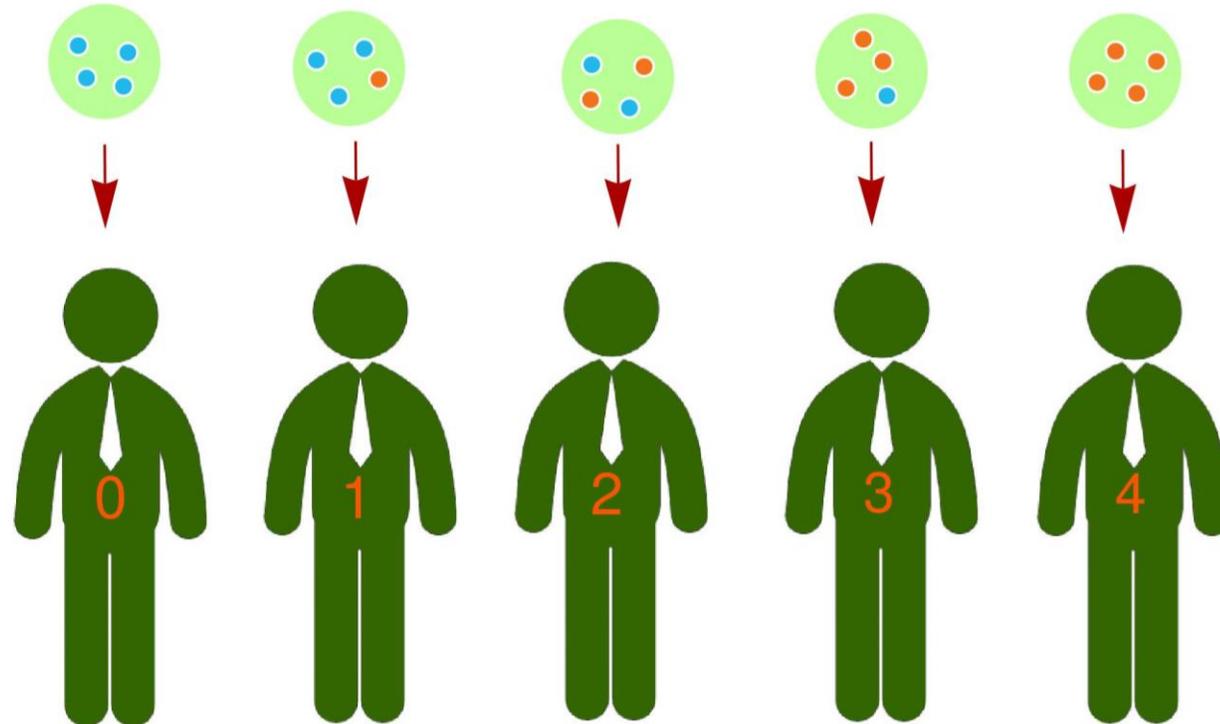
The fraction of homozygous cells

- If a cell contains only mutated chromosomes, call it a homozygous cell for convenience
- The fraction of homozygous cells is a function of g , the elapsed number of cell generations, and the mutation probability μ
- Current belief: this fraction approaches 1.0 as g increases, at least for cases where the ploidy value is a power of 2
- This claim was derived intuitively
- This claim has not yet been subjected to theoretical verification or rigorous simulation testing



Tackling the key issue by simulation

- Agent based simulation: let each cell be an agent having the number of mutated chromosomes as its attribute
- The model was encoded with **NetLogo**





Simulating Chromosome Segregation

Qi Zheng

Department of Epidemiology & Biostatistics, Texas A&M University

What resources were needed?

- For $g=25$ generations, more than 33 million agents will be generated for each cycle
- With so many agents, running the simulation requires a node having 1 TB of random memory
- For reliable results, at least 500 cycles are needed
- For $g=27$, it requires a node having 2 TB of memory
- It then generates 134 million agents
- One cycle of simulation consumes about 1 hour of CPU time





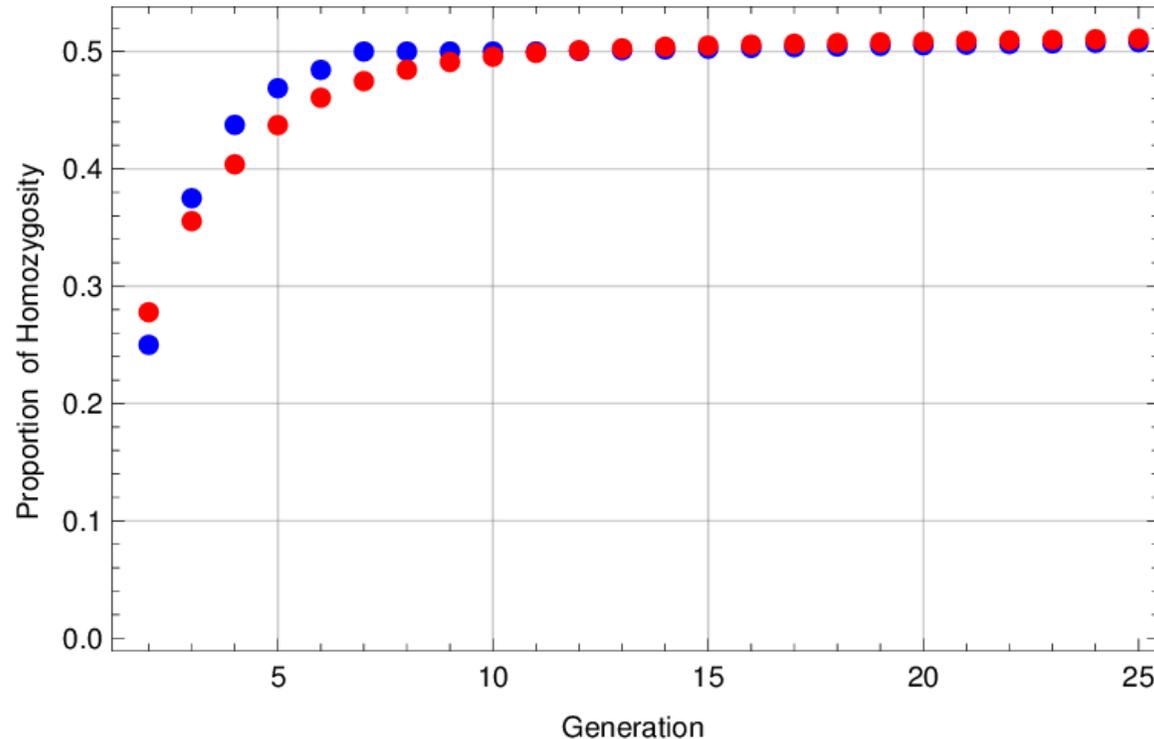
Simulating Chromosome Segregation

Qi Zheng

Department of Epidemiology & Biostatistics, Texas A&M University

What were discovered?

- It challenges current belief: the proportion of homozygous cells does not approach 1.0
- When ploidy value = 2, $\mu=0.001$, the asymptotic proportion is about 0.5
- Blue = median proportion, red = mean proportion





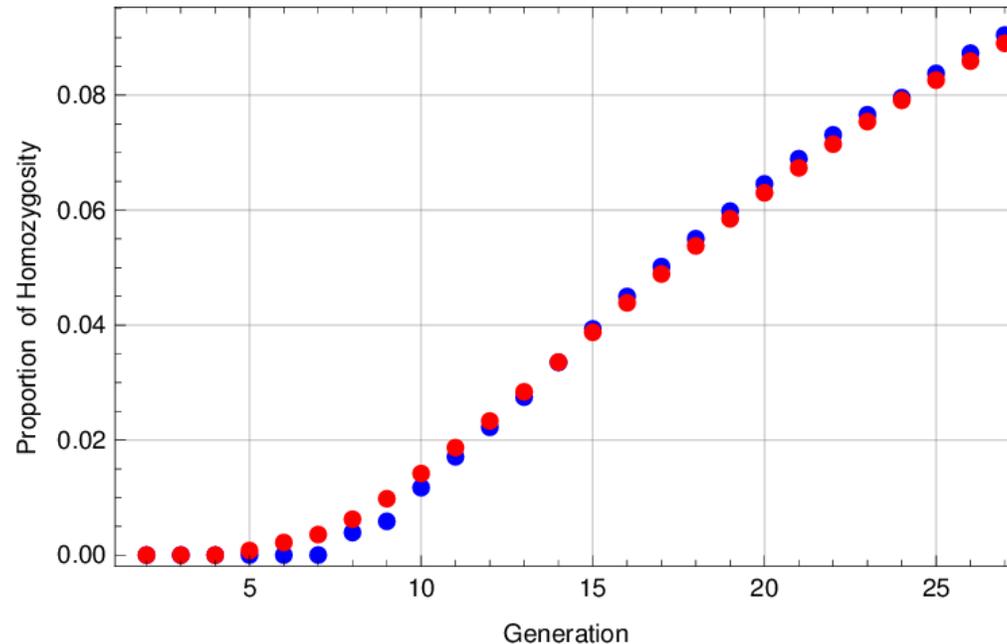
Simulating Chromosome Segregation

Qi Zheng

Department of Epidemiology & Biostatistics, Texas A&M University

Conjectures and challenges

- Simulations have generated interesting conjectures, e.g.:
- There may be unknown simple relations between the ploidy value and the fraction of homozygous cells
- Challenges: with ploidy value = 8, $g=27$ may not be sufficient



The author thanks the Texas A&M High Performance Research Computing for technical support. The simulations were performed on an IBM NeXtScale cluster. Contact: qzheng@sph.tamhsc.edu

