

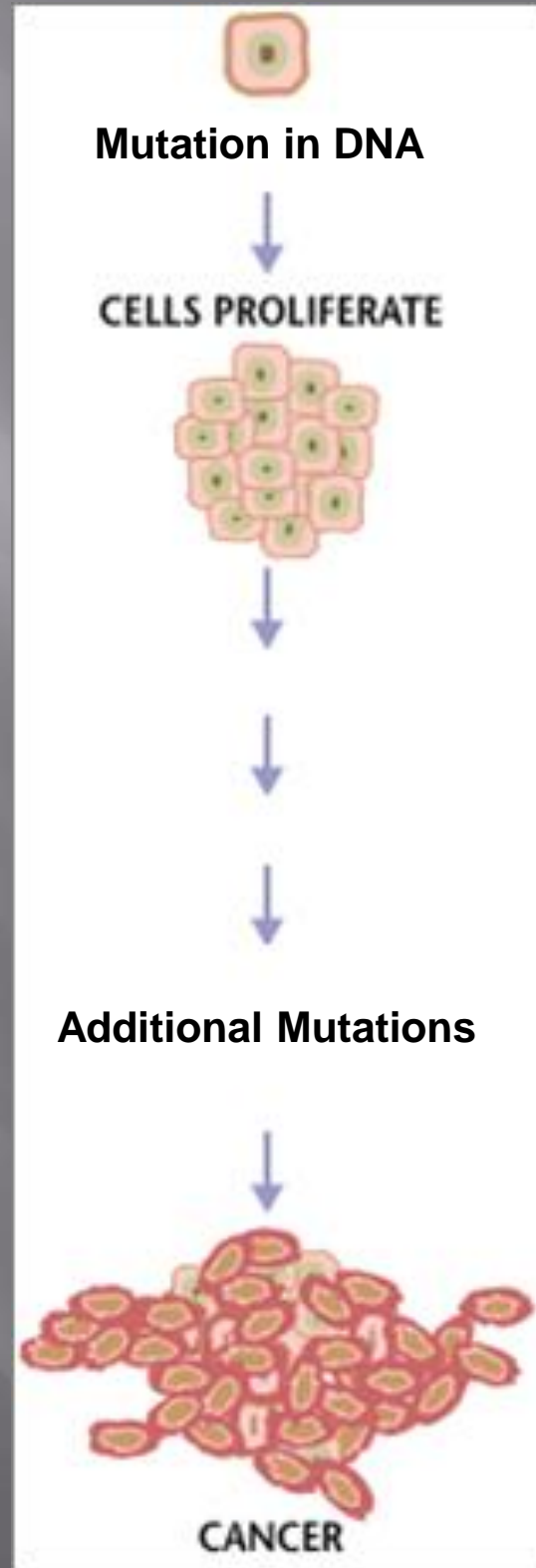
Modeling the role of cell fusion in cancer development

by Andrew Kim and Dash Elhauge

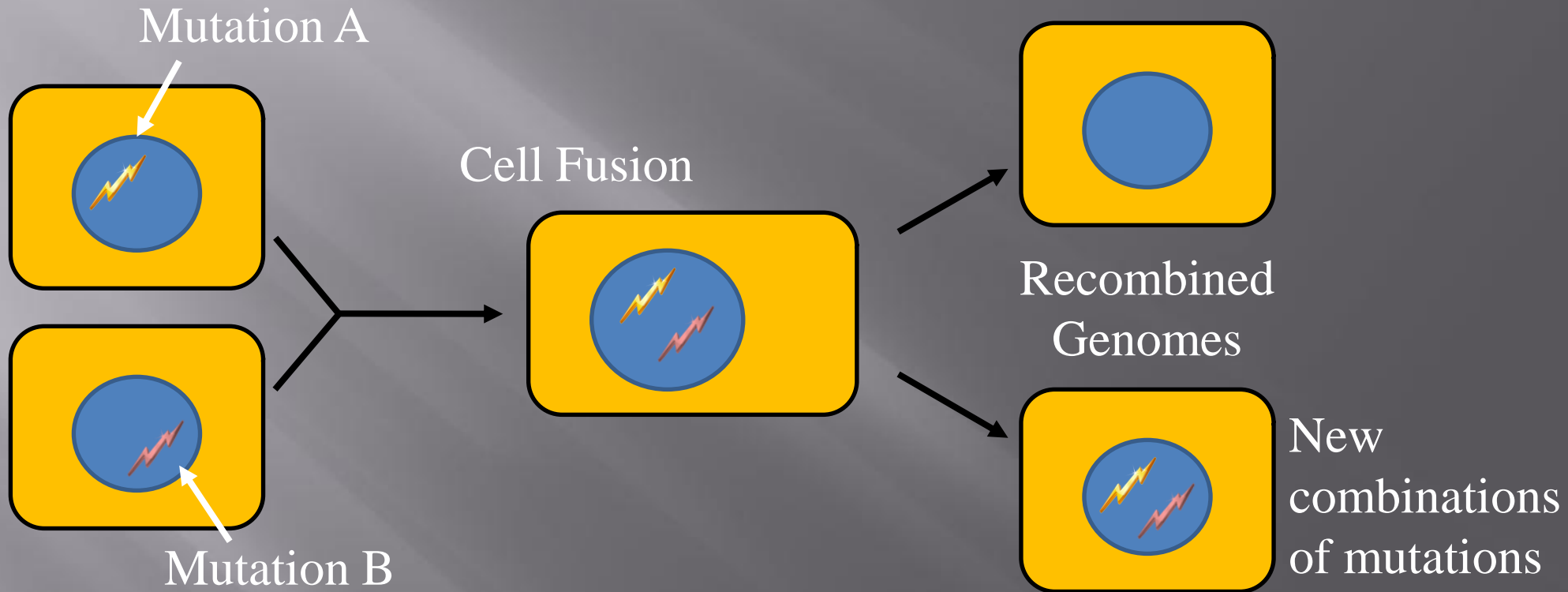
MIT PRIMES

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Introduction



Why does cell fusion occur in cancer?



Hypothesis: Cell fusion allows for recombination of cancer causing genes from different cell lineages

Our Generic Cancer Model

The Goal: *To get cancer to evolve from a population*

Homogeneous
Population (size ~1000)



Apply Mutations (frequency ~ 10^{-6})



Apply Cell fusion (rate ~ 1%)



Apply Evolutionary
Advantage to Most
Cancerous Cells



Next Generation

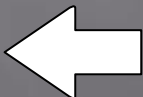


NO

Is cancer
found?



YES



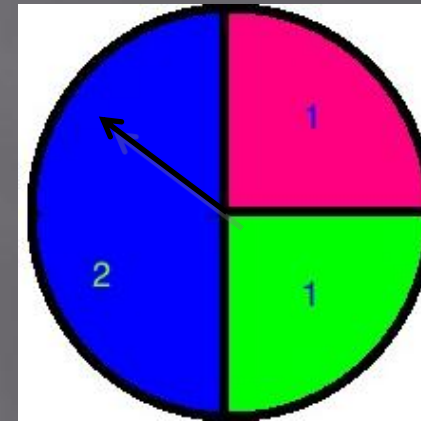
FINISH



What defines advantage?

-Each generation, select cells for the next generation randomly with probability proportional to cell's fitness, keeping population size constant

Sum Fitness: 4



Spin 3 times: Blue, Blue, Green

What defines fitness?

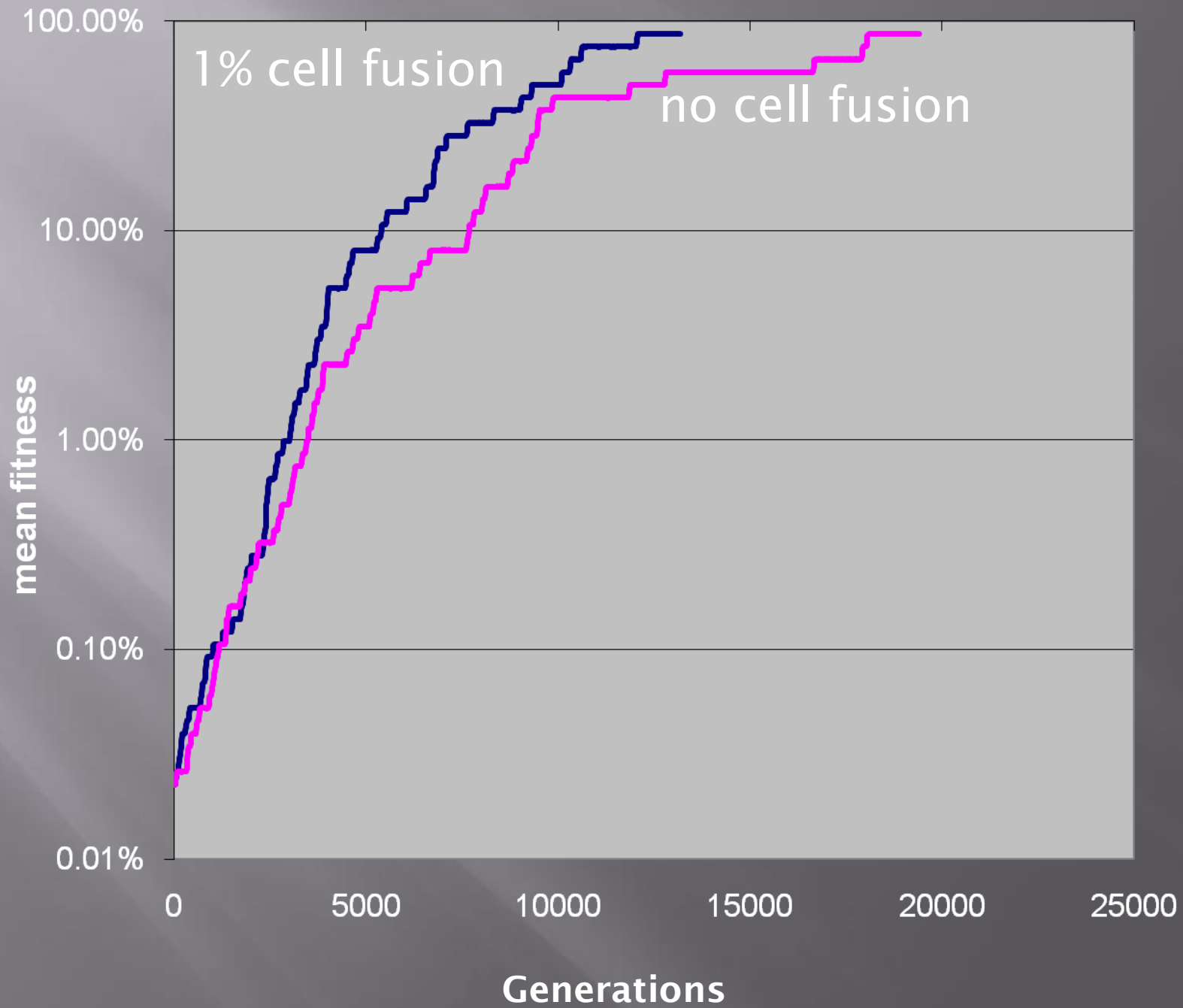
-multiply by a constant s every step towards cancer, i.e.:

$$\text{fitness} \propto e^{-\text{Const} * (\text{distance to cancer})}$$

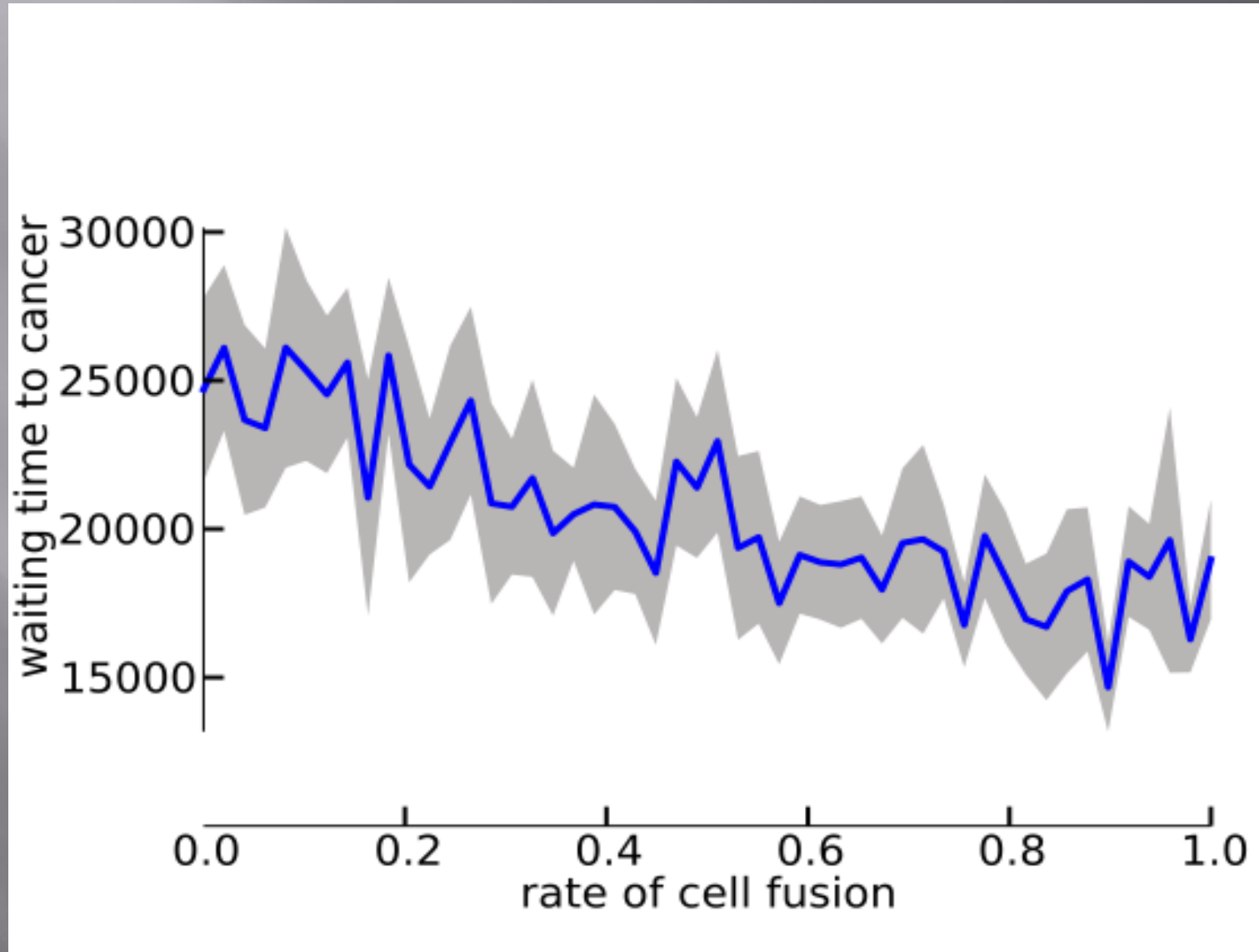
Distance is
number of
discrepancies:

Cancer: 0 0 1 1 0 1 0
 ↑ ↑ ↑
Cell: 1 0 0 1 0 1 1
 = 3

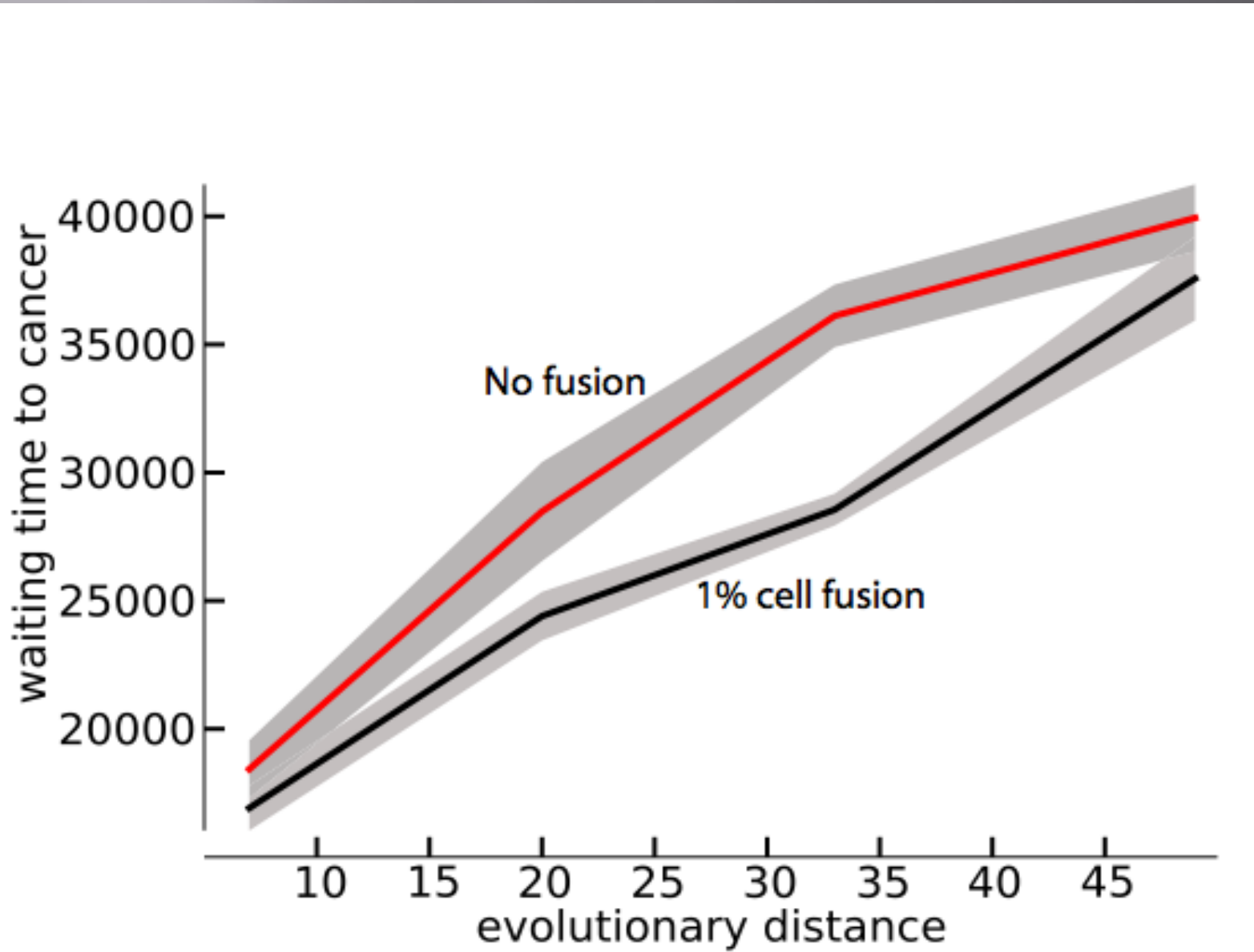
Trajectory of Mean Fitness



Waiting time to cancer decreases as rate of cell fusion increases



Evolutionary distance between starting population and cancer state affects waiting time



Interpreting and Future Studies

- Genetic recombination is universally important in evolution
- Introduction of more environmental, spacial factors
- Limiting the number of assumptions
- Try different cell fusion methods
- Redefining fitness to be more realistic and model complex traits



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