

Degradation-Driven Failure Minimization in Genetic Circuits Through Model Checking

Landon Taylor, Zhen Zhang, and Lukas Buecherl

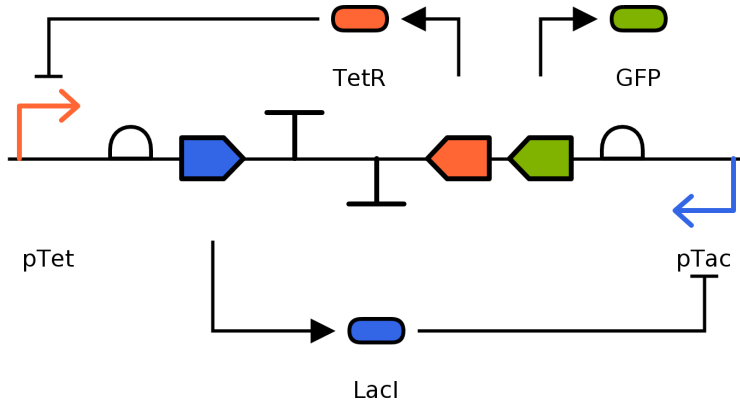
Utah State University

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College of Engineering
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Genetic Toggle Switch



Genetic Toggle Switch

Desired Behavior

The toggle switch should stay in the specified (on/off) state.
Unintended state reversal is a failure (undesired state).

Our Goal

Research Question

Can we modify degradation rates of LacI, TetR, and YFP to reduce failure probability?

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Why degradation?

We can tune performance by utilizing degradation tags.

CTMC Models

- Two models represent the circuit initialized in the “on” (YFP high) and “off” (YFP low) states
- Desired behaviors are mutually exclusive
 - Calls for using two models
 - Creates an optimization problem
- Models are represented as *continuous-time Markov chains* (CTMCs)

CTMC Models

In a CTMC model:

- Each assignment of variable values is a *state*
- Each reaction/update is a *transition*
- Transitions occur at a specified *rate*
- Transition rates correspond to a *probability* of choosing a certain transition

CTMC Models

- CTMC transient analysis gives an *exact* probability
- We use the PRISM probabilistic model checker to do transient analysis
- We specify behavior in *Continuous Stochastic Logic* (CSL):
YFP Low: $P_{=?}[\text{true } U^{[0,2^{100}]} \text{YFP} \geq 30]$
YFP High $P_{=?}[\text{true } U^{[0,2^{100}]} \text{YFP} < 10]$

Rate Variation

- Varied degradation rates (k_d) of YFP, LacI, and TetR across 9 values:
[0.00375, 0.005, 0.00625, 0.0075, 0.00875, 0.01, 0.01125, 0.0125, 0.015]
- Values are spaced around iBioSim's default k_d of 0.0075
- We explore 3^9 combinations:
(0.00375, 0.00375, 0.00375)
(0.00375, 0.00375, 0.005)
(0.00375, 0.005, 0.00375)
(0.005, 0.00375, 0.00375)
(0.00375, 0.00375, 0.00625)
...

Methodology

For each combination of degradation rates for YFP, LacI, and TetR:

- Python script updates the model's degradation rates
- PRISM CTMC transient analysis finds the probability of an undesired state switch within about one cell cycle

Methodology

After all models are tested:

- Script processes and ranks resulting failure probabilities, considering:
 - Each rate combination affects circuit behavior differently
 - Improving the YFP high model may harm the YFP low model
 - A metric is needed to weigh the probability pairs
- Script recommends optimal combinations of degradation rates

Metrics

Metrics penalize both high probabilities and imbalanced probabilities. They include:

- Simple sum $(p_{\text{high}} + p_{\text{low}})$,
- Average plus difference $((p_{\text{high}} + p_{\text{low}})/2 + \text{abs}(p_{\text{high}} - p_{\text{low}}))$,
- Sum of squares $(p_{\text{high}}^2 + p_{\text{low}}^2)$ and cubes $(p_{\text{high}}^3 + p_{\text{low}}^3)$,
- and several others

The variety of metrics allows flexibility: engineers can choose their priorities

For example:

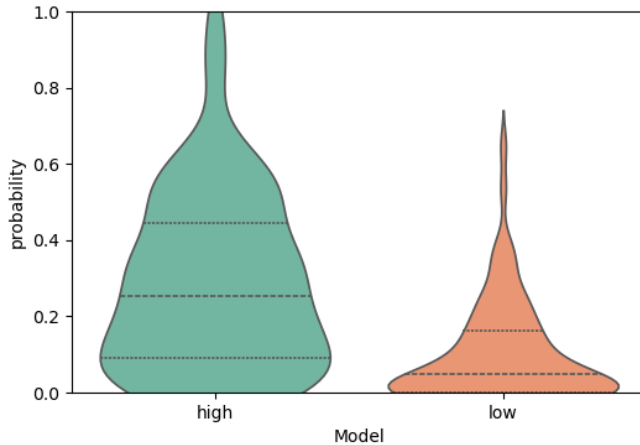
- $p_{\text{high}} = 0.3$ and $p_{\text{low}} = 0.3$ is balanced
- $p_{\text{high}} = 0.6$ and $p_{\text{low}} = 0.0$ has the same average, but is not balanced

Probability Distribution

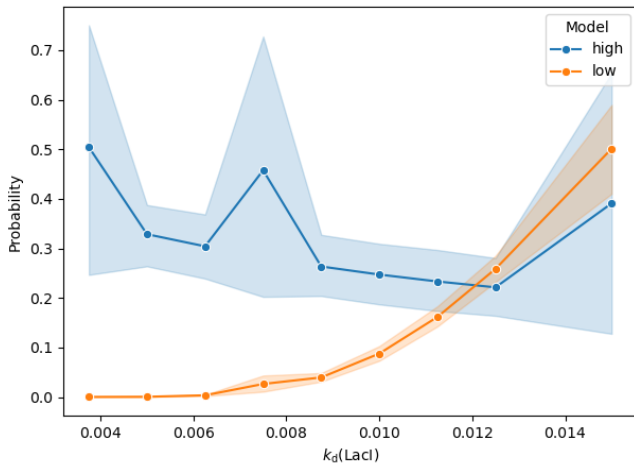
Preliminary analysis shows that:

- Each degradation rate has a unique impact on failure probability
- The YFP high model sees more variation than the YFP low model
- The YFP low model sees a generally-low failure probability

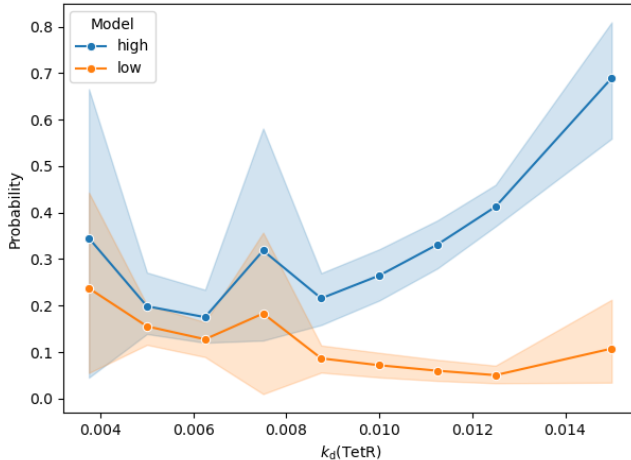
Probability Distribution



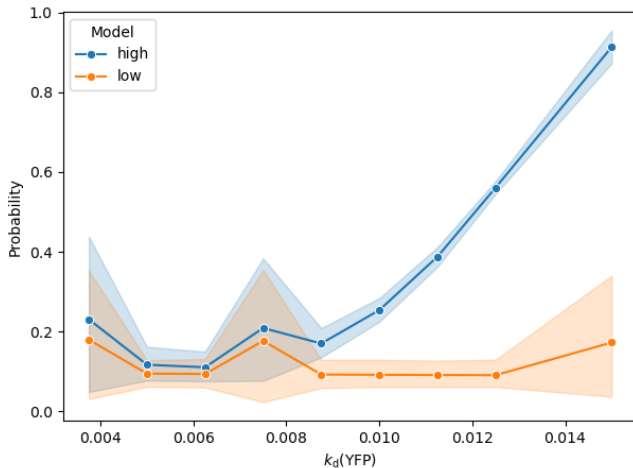
Probability vs k_d



Probability vs TetR



Probability vs YFP



Observations

- Optimizing robustness in one model reduces stability in the other
- This model calls for a balanced profile across YFP, LacI, and TetR
- Each degradation rate has a unique impact
- Modifying degradation rates can greatly impact probability of failure

Best Combinations

For 20 metrics over all parameter combinations, winning combinations for YFP, LacI, TetR are:

- (0.00375, 0.00375, 0.00375) – 5 metrics
- (0.00375, 0.015, 0.00375) – 5 metrics
- (0.015, 0.00375, 0.015) – 5 metrics
- (0.00625, 0.00625, 0.005) – 2 metrics
- (0.015, 0.00375, 0.00375) – 2 metrics
- (0.015, 0.015, 0.00375) – 1 metric

Conclusion

We choose $(YFP, LacI, TetR) = (0.00375, 0.00375, 0.00375)$ because:

- It was selected by five metrics
- Manual validation found it to be best overall
- It provides a reasonable balance between probability of failure for both the high and low models

Limitations:

- We rely on a combinatorial analysis of degradation rates
- This approach is minimally scalable

Future Work:

- Validate these results in the lab
- Use parameter synthesis techniques and heuristics to scale to more complex genetic circuits

Thank You

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