

# Things left:

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- Write essay
- Accept DNA sequence from the user at program start (By default use the Insulin, note that other DNA sequences might fail as the pre-mRNA splicing algorithm is very heuristic and prone for failure in exotic inputs)
- Fail while iterating in the simulation when getting to the end of the mRNA chain when an error happened while translating the chain (Due to the nonsense-mediated-decay)
- Fix simulation with spacings and try to make it Shimon-esque

For now that is all.