

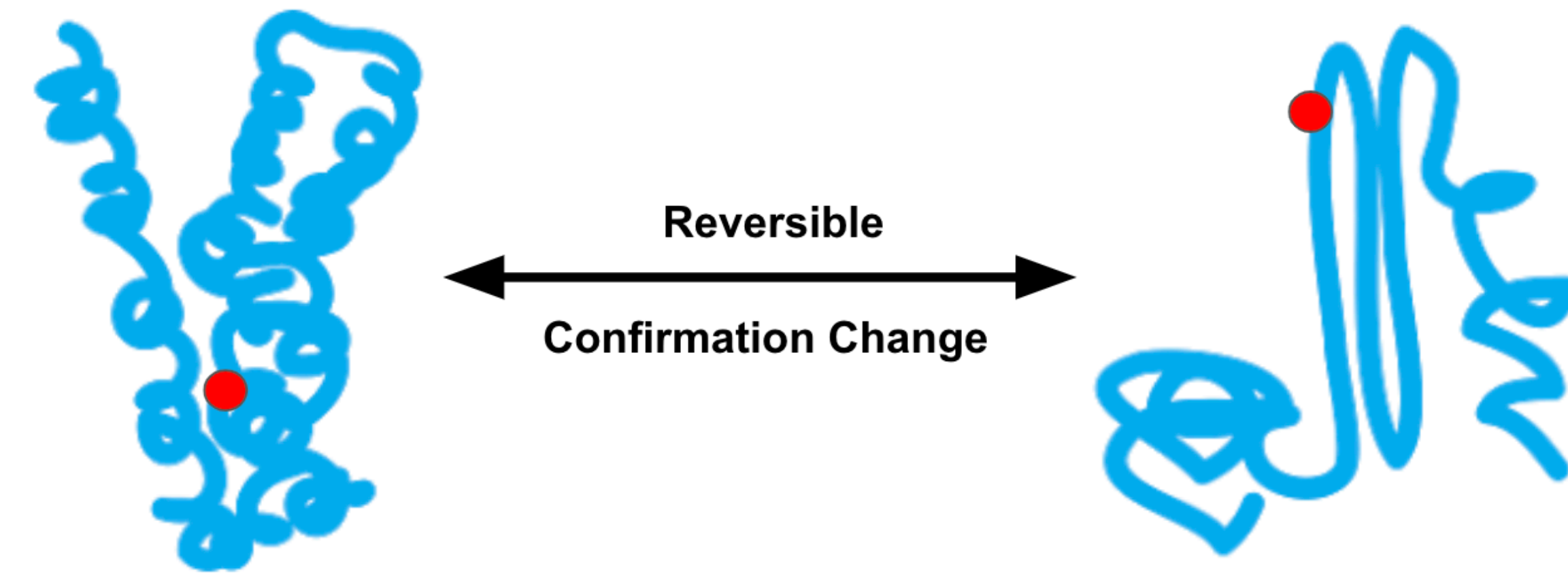
Propagating Putative Prion States in Budding Yeast

Jacob Evarts¹, David Garcia^{1, 2}

¹University of Oregon, ²Department of Biology

Prion proteins

- Prions are misfolded proteins that can cause other proteins of the same type to also misfold



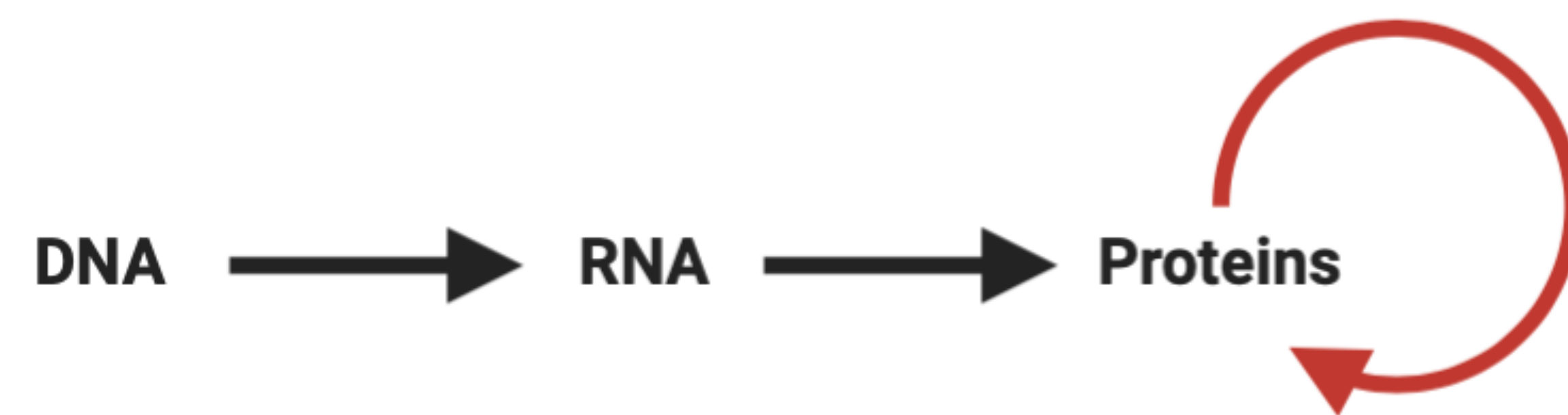
Naive or Normal Protein

Prion Protein

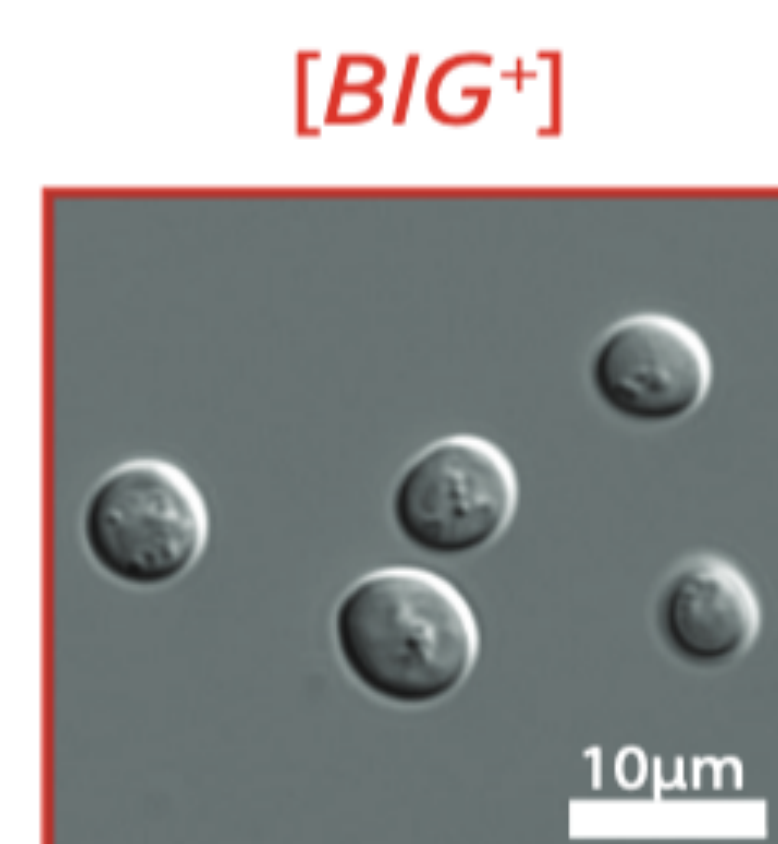
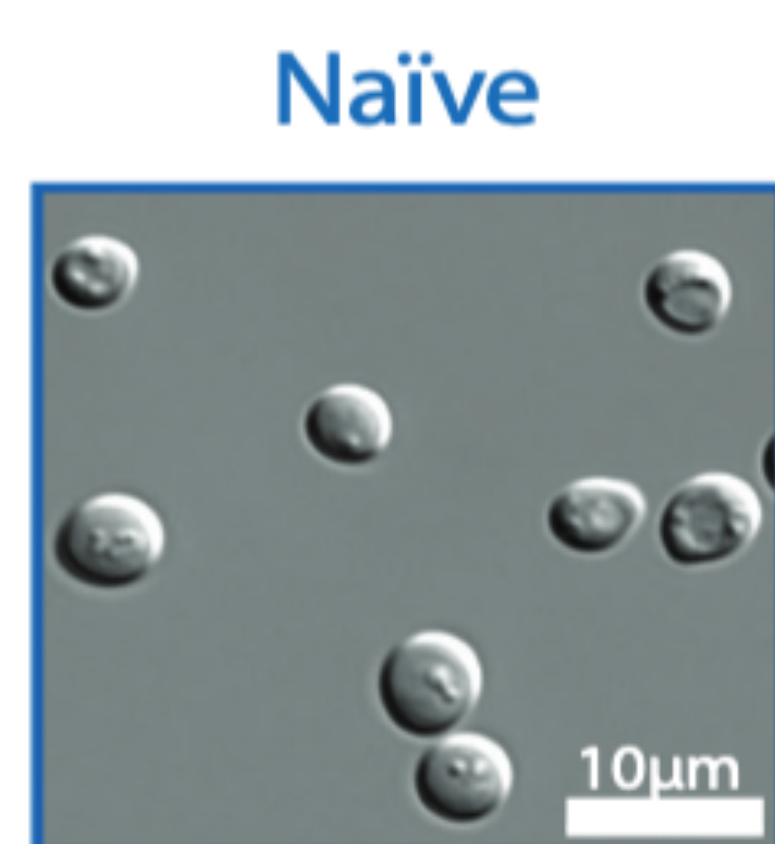
- Previously believed to be only disease causing agents, such as Bovine Spongiform Encephalopathy (Mad Cow Disease)

Challenging the convention

- Recent evidence suggests that prions provide an additional class of epigenetic mechanisms that work at a rapid pace

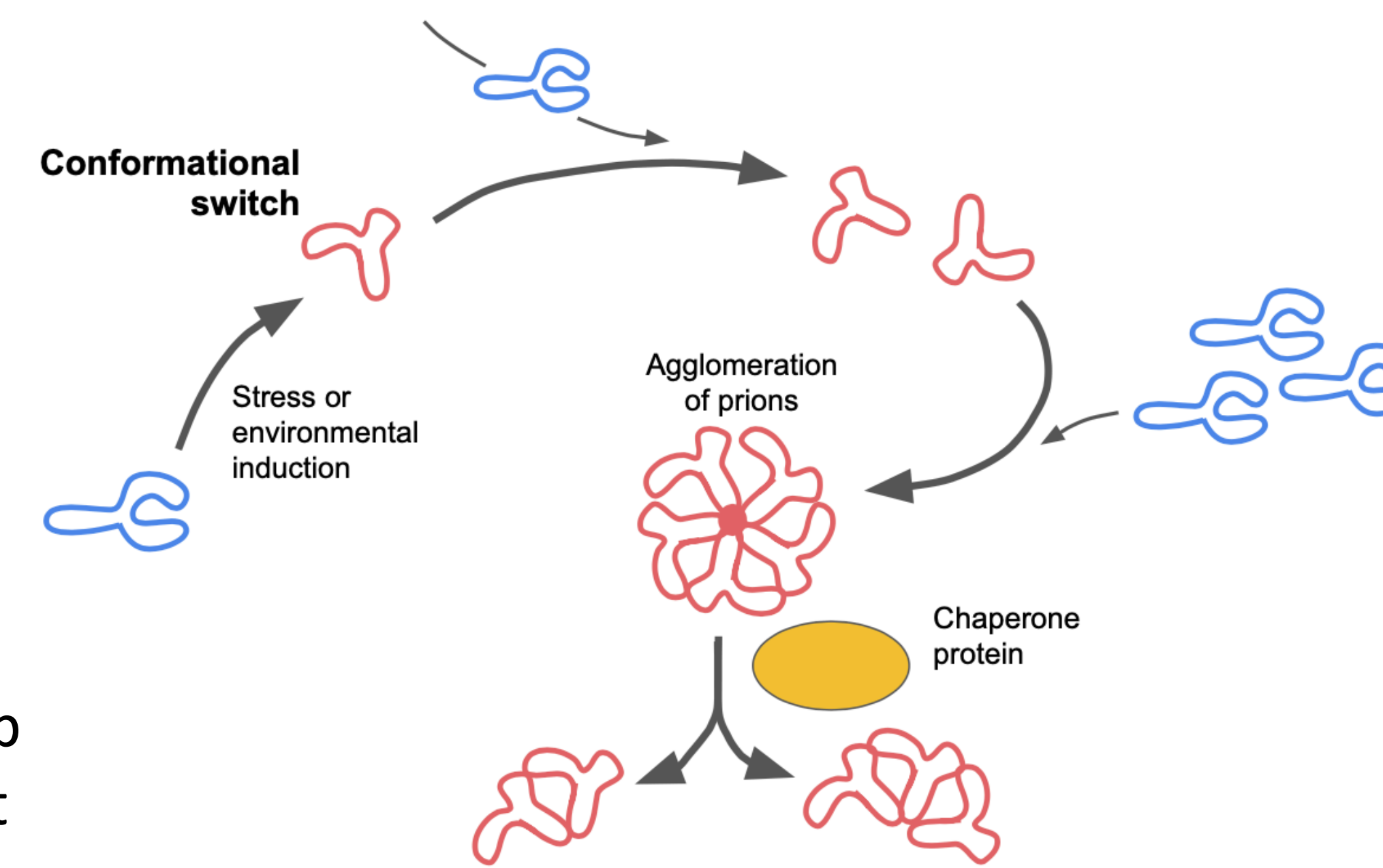


- The Garcia Lab and others recently discovered prions states, such as [BIG⁺] that allow cells to grow faster and larger



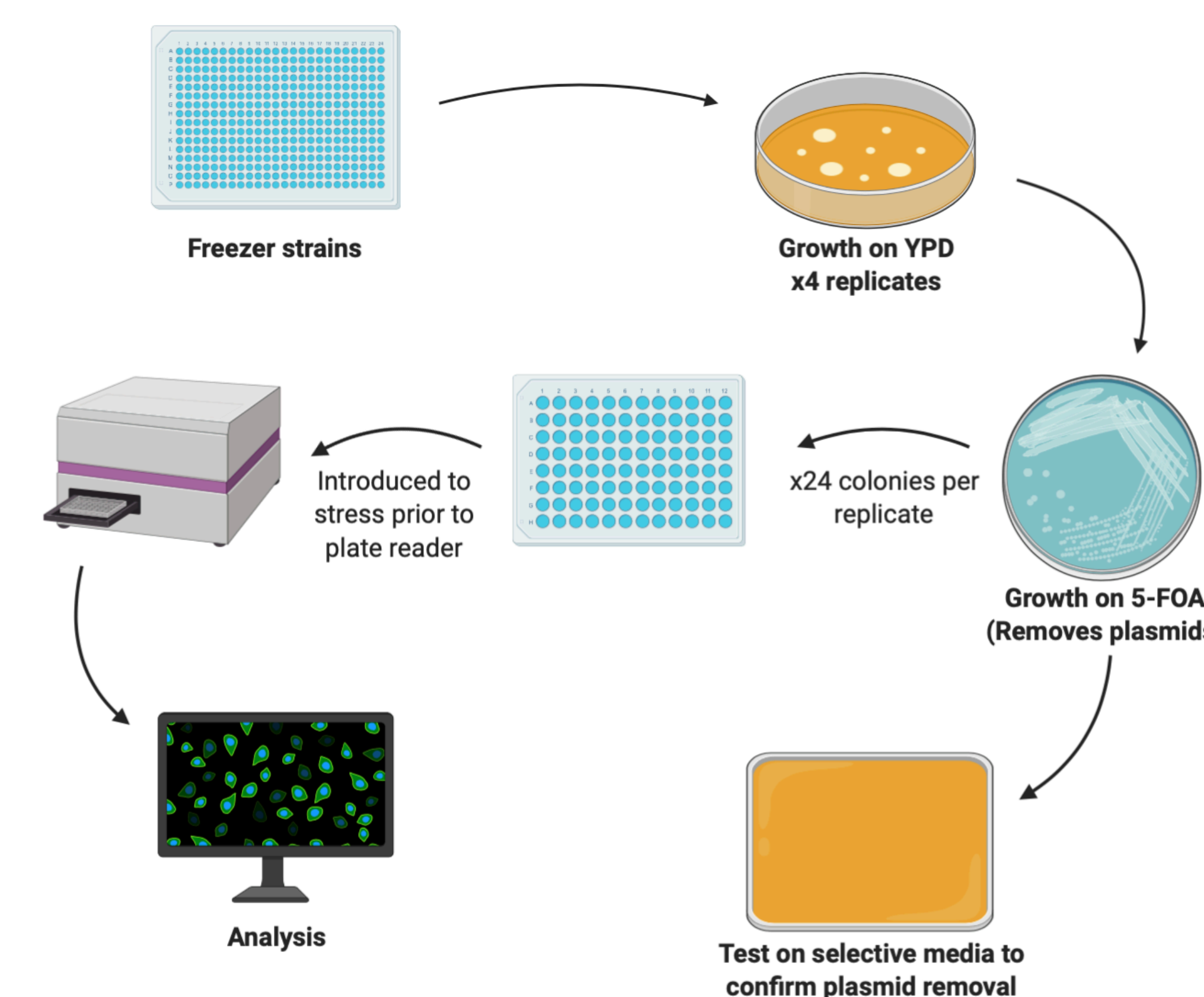
Molecular propagation of prion proteins

- Prions formations occur much more frequently than genetic mutation
- Stress from the environment and protein overexpression can induce prion formation
- Chaperone proteins help 'seed' prions to the next generation of cells



Screen for new prion-like proteins

- The goal of my project is to identify new RNA modifying enzyme (RME) prions



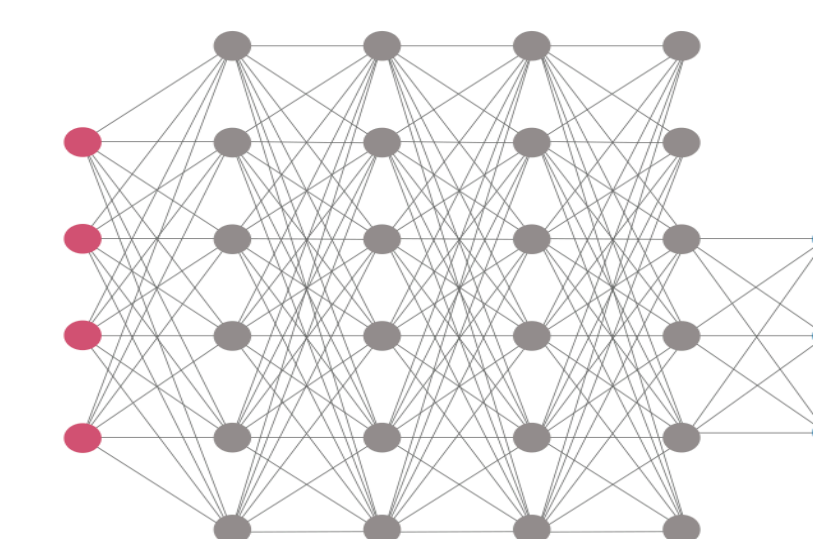
~23% of disease associated genes have close ortholog in yeast

~50% of mutations in RMEs are linked to diseases in humans

>5mil data points collected measuring growth rate and carrying capacity

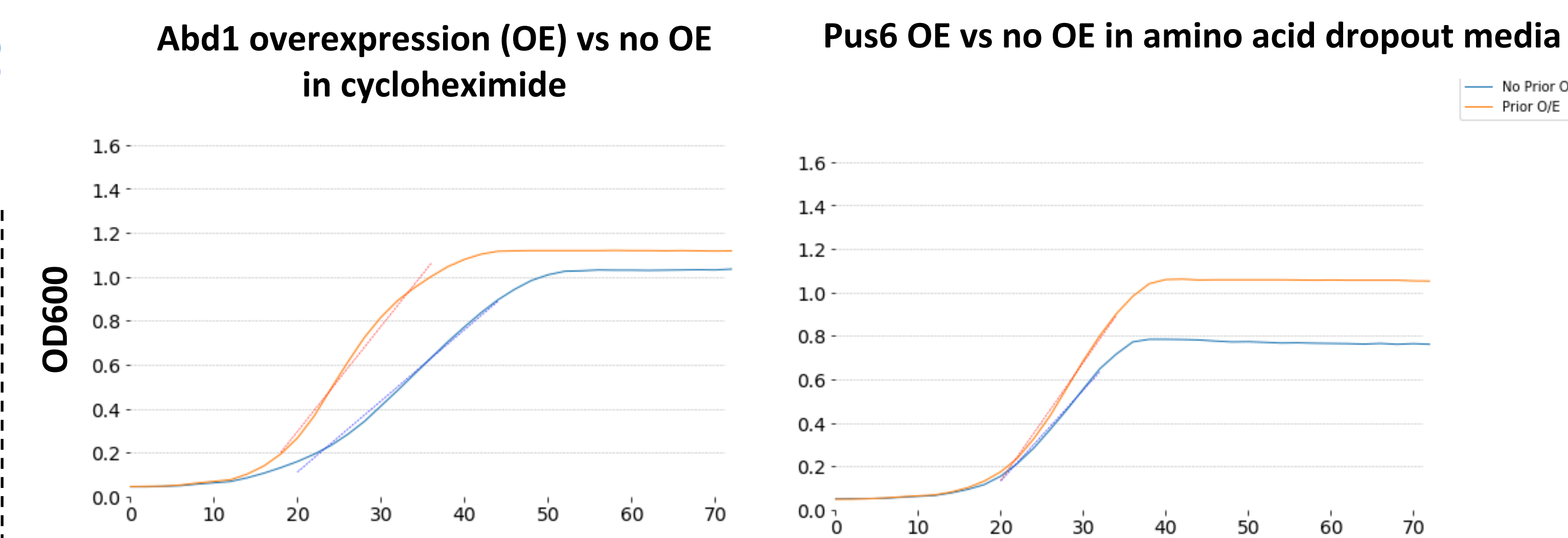
- Cell strains had previous overexpression of RMEs and were exposed to various stress conditions (heat, chemicals, nutrient deficiency, etc.)

- Data analytics and machine-learning toolsets are used to sort and classify the extreme amounts of raw data produced



Results

- Prion states appear to be a common form of inheritance in yeast
- Nine different strains showed significant resistance to stress after having previous overexpression of an RME



Trm5 overexpression vs. no overexpression in cycloheximide



Better growth in replicants 3 & 4

Future Experiments

- Use fluorescence microscopy to see if these proteins are agglomerating
- Test human homologs in similar assays in yeast
- Try other machine-learning models on predicting prion phenotypes