

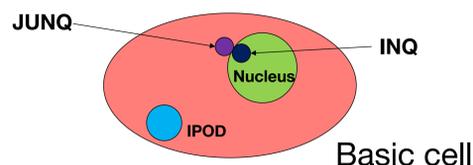
What's With That JUNQ?: Protein Sequestration and Aggregation in Yeast

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Introduction

What are we doing and why?

- Protein misfolding and the presence of sequestration sites has been linked to many neurodegenerative disorders (Alzheimer's, Huntington's, Parkinson's)
- What's pathogenic? The sites themselves or does the presence of the sites already indicate pathogenicity?
- Understanding how two specific misfolded protein sequestration sites interact may allow us to relate this to neurodegenerative diseases in general



Inspirations and Past Conclusions

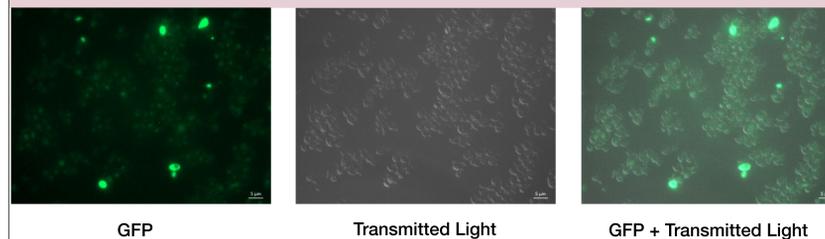
- Eukaryotic cells sort their misfolded proteins into distinct compartments – soluble proteins to the JUNQ and insoluble to the IPOD (Kaganovich et al, 2008)¹
- JUNQ inclusion bodies are asymmetrically inherited, so daughter cells do not get misfolded proteins (Ogrodnik et al, 2014)²
- The JUNQ and INQ home to each other but are not physically connected (Sontag, unpublished)

Terms to Know

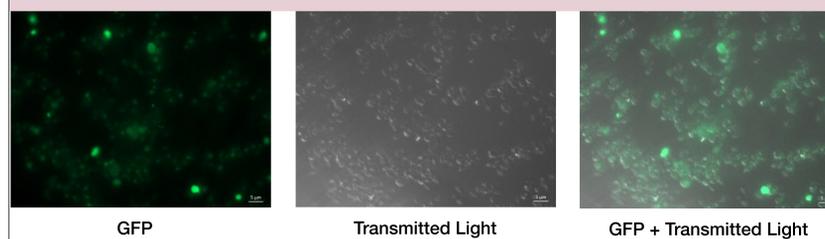
- *juxtannuclear quality control compartment (JUNQ)*: body of aggregated misfolded proteins localized just outside the nucleus
- *intranuclear quality control compartment (INQ)*: body of aggregated misfolded proteins located inside the nucleus
- *nuclear localization signal (NLS)*: localization tag that forces a protein to hone to the nucleus
- *nuclear export signal (NES)*: localization tag that forces a protein out of the nucleus
- *luciferase TS (luci)*: protein engineered to misfold above 37°C
- *Von Hippel Lindau (VHL)*: protein that will misfold in yeast

Results

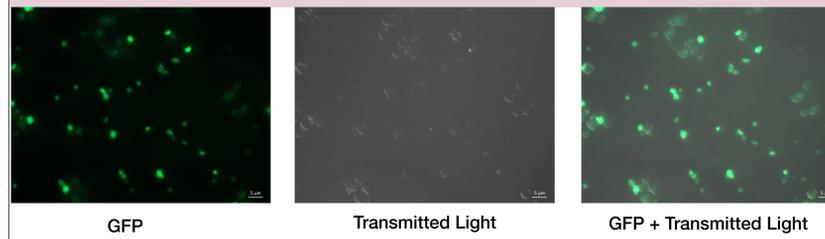
W303 + NLS-GFP-Luci



W303 + NLS-GFP-VHL



Y7092 + NLS-GFP-Luci



Conclusions

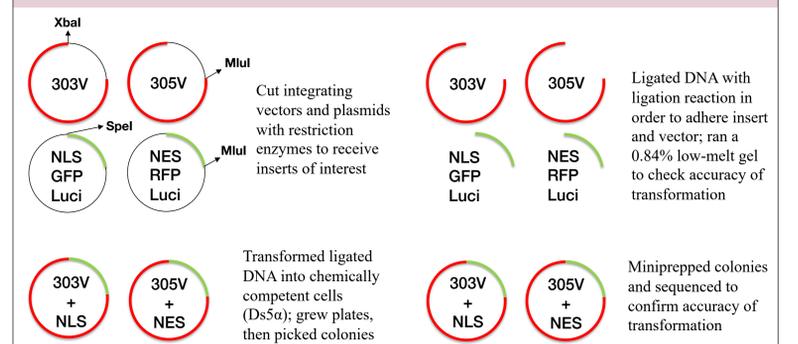
- Successful creation of integrating vectors pAG303-NLS-GFP-Luci, pAG303-NLS-GFP-VHL, pAG305-NES-RFP-Luci, pAG305-NES-RFP-VHL
- Successful integration of NLS-GFP-Luci into yeast strains W303 and Y7092 (evidenced by the imaging, where GFP marks the cell nuclei) and NLS-GFP-VHL into W303

Next Steps

- Integrating the NES-RFP-Luci and NES-RFP-VHL into the NLS-integrated W303 and Y7092, and doing a library screen of deletion collection with the Gitler lab to study JUNQ/INQ
- Creating a singular integrating vector with NLS and NES linked by a self-cleaving T2A peptide
- Reattempting the Y7092 + NLS-GFP-VHL integration

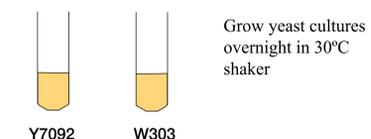
Methods

NLS-GFP-Luci and NES-RFP-Luci + Vectors

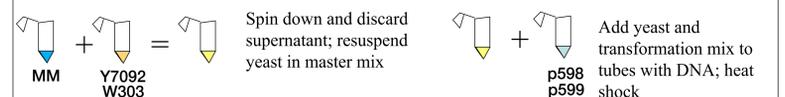
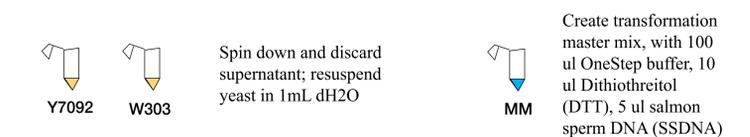
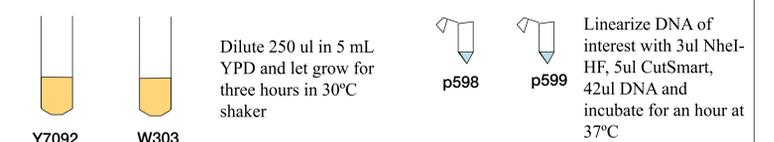


Integrating into Y7092 and W303

Day 0



Day 1



Resuspend mix in 1.5 mL YPD; recover for 5 hours at 30°C, then plate on selective plates

References

- 1: Kaganovich, Daniel et al. "Misfolded proteins partition between two distinct quality control compartments." *Nature* vol. 454, 7208 (2008): 1088-95. doi:10.1038/nature07195
- 2: Ogrodnik M, et al. "Dynamic JUNQ inclusion bodies are asymmetrically inherited in mammalian cell lines through the asymmetric partitioning of vimentin". *Proceedings of the National Academy of Sciences of the United States of America*. 111 (22) (2014): 8049-54. doi:10.1073/pnas.1324035111.

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