

College of Saint Benedict and Saint John's University

DigitalCommons@CSB/SJU

Celebrating Scholarship and Creativity Day

Undergraduate Research

4-26-2018

Analysis of potential archaeal NER endonuclease homologs using *Saccharomyces cerevisiae*

Toni R. Gohman

College of Saint Benedict/Saint John's University, trgohman@csbsju.edu

Follow this and additional works at: https://digitalcommons.csbsju.edu/ur_cscday



Part of the [Biology Commons](#)

Recommended Citation

Gohman, Toni R., "Analysis of potential archaeal NER endonuclease homologs using *Saccharomyces cerevisiae*" (2018). *Celebrating Scholarship and Creativity Day*. 8.

https://digitalcommons.csbsju.edu/ur_cscday/8

This Presentation is brought to you for free and open access by DigitalCommons@CSB/SJU. It has been accepted for inclusion in Celebrating Scholarship and Creativity Day by an authorized administrator of DigitalCommons@CSB/SJU. For more information, please contact digitalcommons@csbsju.edu.

Analysis of potential archaeal NER endonuclease homologs using *Saccharomyces cerevisiae*

Toni Gohman, advisor Dr. Michael Reagan

Introduction

- Environmental agents can create distorting lesions in DNA and disrupt cell function¹
- Nucleotide Excision Repair (NER) involves complex cooperation of proteins to remove DNA lesions²
- NER process and proteins involved are well understood for eukaryotes, but not for archaea³
- rad1 (known as XPF in eukarya) is 5' endonuclease in NER and is found in all eukaryotes²
- Based on amino acid sequencing and biochemical function, archaeal proteins Bax1 and Hef1 could perform rad1 role^{4,5}

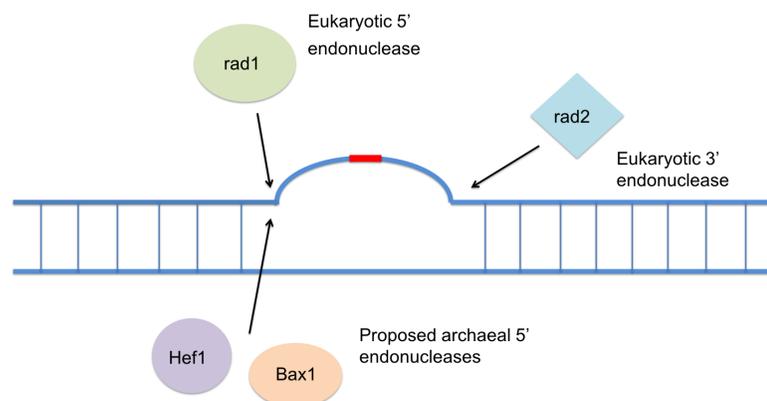


Figure 1. Eukaryotic nuclease function and proposed archaeal homologs in NER. rad1 functions as a 5' endonuclease in eukaryotic NER; Bax1 and Hef1 are proposed to perform the same function based on structure and nuclease activity.

Methods

- rad1 deletion (Δ rad1) *S. cerevisiae* strain was obtained and archaeal genes were inserted using pYES2/NT vector⁶
- Growth in Uracil-deficient media ensures that plasmid will be maintained in cells
- *S. cerevisiae* strains were grown with dextrose for 72 hours at 30°C
- Transferred to galactose to induce expression of archaeal gene and grown at 30°C
- After 24 hours, cells were plated on agar and exposed to Ultraviolet (UV) light
- Cells were grown for 72-120 hours at 30°C and survival rates were calculated

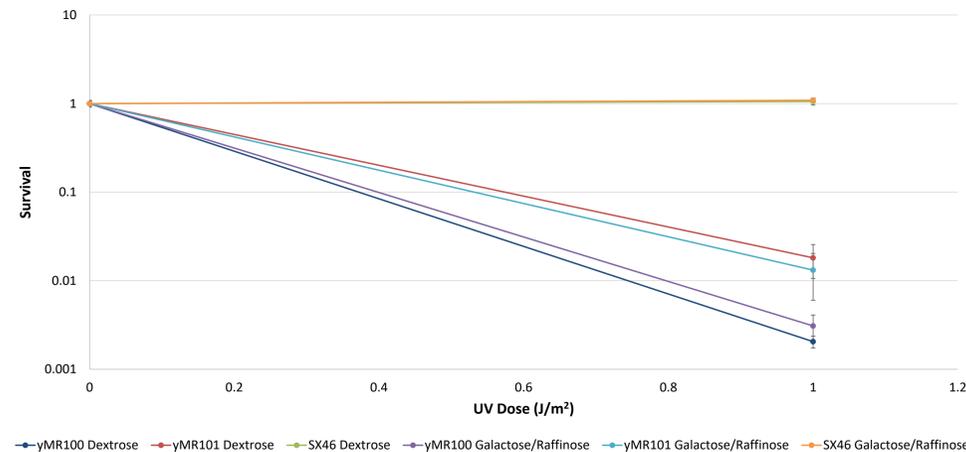


Figure 2. Survival of Δ rad1 *S. cerevisiae* + *M. voltae* Hef1 (yMR100) when utilizing different carbohydrate sources. yMR100, wild type (WT), and Δ rad1+pYES2/NT vector (yMR101) were plated with dextrose or galactose+raffinose and exposed to UV. Error bars represent standard error of the mean.

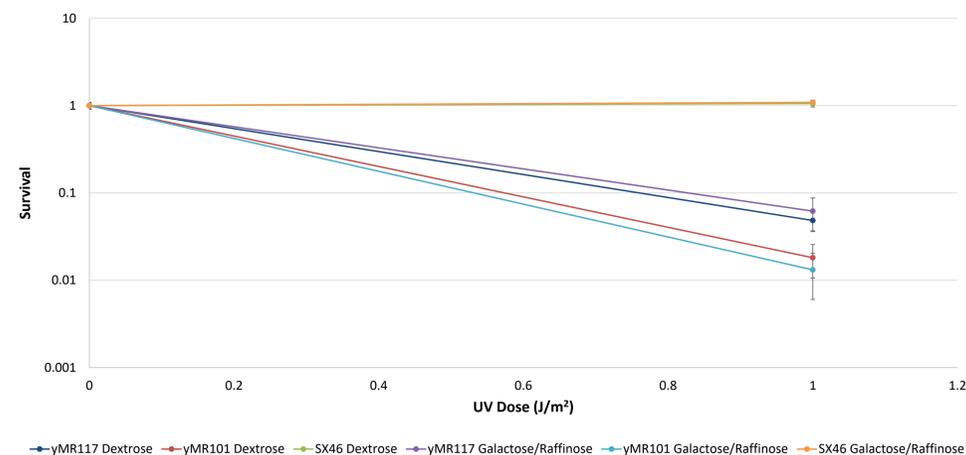


Figure 3. Survival of Δ rad1 *S. cerevisiae* + *M. acetivorans* Hef1 (yMR117) when utilizing different carbohydrate sources. yMR117, WT, and Δ rad1+pYES2/NT vector (yMR101) were plated with dextrose or galactose+raffinose and exposed to UV. Error bars represent standard error of the mean.

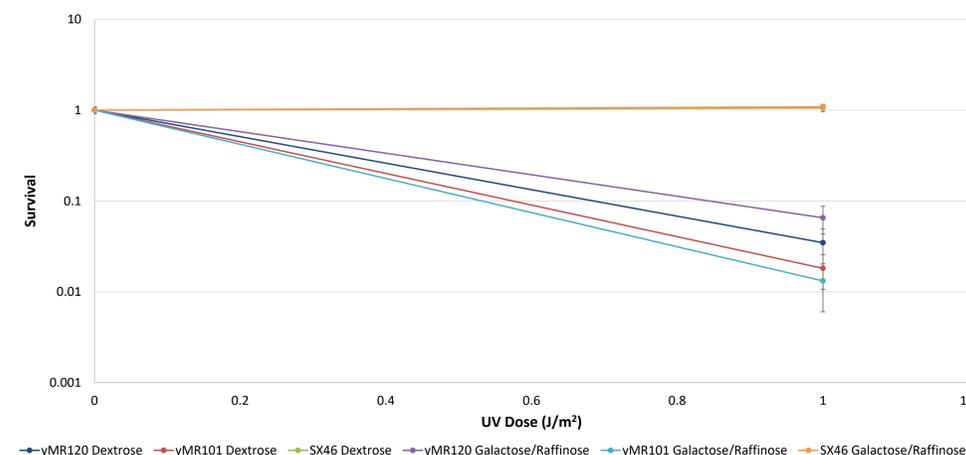


Figure 4. Survival of Δ rad1 *S. cerevisiae* + *M. acetivorans* Bax1 (yMR120) when utilizing different carbohydrate sources. yMR120, WT, and Δ rad1+pYES2/NT vector (yMR101) were plated with dextrose or galactose+raffinose and exposed to UV. Error bars represent standard error of the mean.

Results

- Δ rad1 + pYES2/NT vector shows high UV sensitivity consistent with nonfunctional NER
- Δ rad1 + *M. voltae* Hef1 did not have a higher rate of survival than Δ rad1 + pYES2/NT vector with either carbohydrate source (Fig. 2)
- Δ rad1 + *M. acetivorans* Hef1 had significantly higher survival rates than Δ rad1 + pYES2/NT vector with both carbohydrate sources (Fig. 3)
- Δ rad1 + *M. acetivorans* Bax1 had significantly higher rates of survival than Δ rad1 + pYES2/NT vector on galactose/raffinose agar (Fig.4)
- Δ rad1 + *M. acetivorans* Bax1 plated on galactose/raffinose agar had higher rates of survival than when plated on dextrose agar (Fig. 4)

Discussion

- Increased survival of Δ rad1 + *M. acetivorans* Bax1 and Hef1 strains indicates that Bax1 and Hef1 have potential to perform rad1 function in archaea
- Increased survival of Δ rad1 + *M. acetivorans* Bax1 on galactose+raffinose agar suggests that plasmid maintenance and expression is critical for repair in initial hours after UV exposure
- *M. acetivorans* Hef1 was synthesized to correct for the codon bias in *S. cerevisiae* and can account for higher survival than uncorrected *M. voltae* Hef1

References

- Costa, R. M.A., Chiganças, V., da Silva Galhardo, R., Carvalho, H., & Menck, C. F.M. (2003). The eukaryotic nucleotide excision repair pathway. *Biochimie*, 85, 1083-1099.
- Rouillon, C., & White, M. F. (2011). The evolution and mechanisms of nucleotide excision repair proteins. *Research in Microbiology*, 162, 19-26.
- Kelman, Z., & White, M. F. (2005). Archaeal DNA replication and repair. *Current Opinion in Microbiology*, 8, 669-676.
- Roth, H. M., Tessmer, I., Van Houten, B., & Kisker, C. (2009). Bax1 is a novel endonuclease: Implication for Archaeal nucleotide excision repair. *The Journal of Biological Chemistry*, 284(47), 32272-32278.
- Roberts, J. A. & White, M. F. (2005). An Archaeal Endonuclease Displays Key Properties of Both Eukaryal XPF ERCC1 and Mus81. *The Journal of Biological Chemistry*, 280(7), 5924-5928.
- Invitrogen. (2012). pYES2/NT A, B, and C, pYC2NT A, B, C: Yeast expression vectors with N-terminal tags: User Guide. Carlsbad, CA: Life Technologies.

Acknowledgements:

I would like to thank Dr. Michael Reagan for his mentorship on this project, as well as for the use of his lab space and materials.