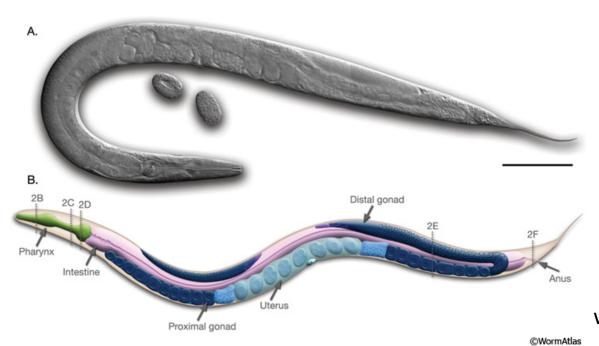


Creation of *odd-skipped* fluorescent reporter strains in *C. elegans* via CRISPR/Cas9 genome editing By Lillian McCormick, Levi Duhaime, and Amy C. Groth, PhD. Methods: Co-CRISPR Introduction **Future Directions** Approaches for increasing efficiency: WT Injection mix includes repair templates Using single-stranded repair templates for *dpy-10* and for desired modification Using ribonucleoprotein complexes (RNPs) = Cas9 protein + WT (MOD) guide RNA **Nested CRISPR** strategy: • Higher success rate than original CRISPR/Cas9 strategy⁶ dpy-10(rol)MOD dpy-10(rol)MOD WJ • 1st insertion (single-stranded) contains beginning and end dpy-10(WT)WT *dpy*-10(*mut*) ' WT WT of the MOD and homology from DNA • Uses user-designed sgRNA • 2nd insertion (double-stranded) uses whole MOD as repair wormatlas.com sequence Homozygotes identified Beginning and end of MOD act as homology by PCR and isolated Uses known/tested sgRNA for GFP — DNA ODD-1 Promoter dpy-10(WT)dpy-10(rol)MOD MOD **Cas9** target sequence dpy-10(WT) 'MOD dpy-10(WT/rol)' MOD 3 HOM HOM dpy-10(WT) MOD MOD dpy-10(rol)Cas9 target sequence dpy-10(WT) 'WT dpy-10(WT/rol)'WT dpy-10(rol). <u>WT</u> dpy-10(WT) WT dpy-10(WT/rol) 'WT dpy-10(WT) 'WT GFP (Figure adapted from Dickinson and Goldstein, 2016) ODD-1 GFP Promoter • Allows screening by phenotype: screen for non-dumpy **Hybrid CRISPR** strategy: rollers in F1 and for wild type in F2⁵ Melting/annealing of repair sequences makes "free ends" Desired modification (MOD) is either *odd-2::mCherry* or 50% of repair mix is partially single-stranded DNA = better odd-1::GFP repair/higher insertion success⁷ • >150 worms were injected and screened using this method Repair Mix 120 bp homology Results Melt/ Anneal Results from Co-CRISPR injections: F1 PCR results: Arrow shows positive result MOD should be in Acknowledgments/References chromosome of positive F1 NA individuals Acknowledgments Offspring had no insertions Offspring singled from positive F1:

- *. elegans* microscopic nematodes; model organism
- Easy visualization due to transparency
- Easy to grow/maintain
- Entire genome has been sequenced¹



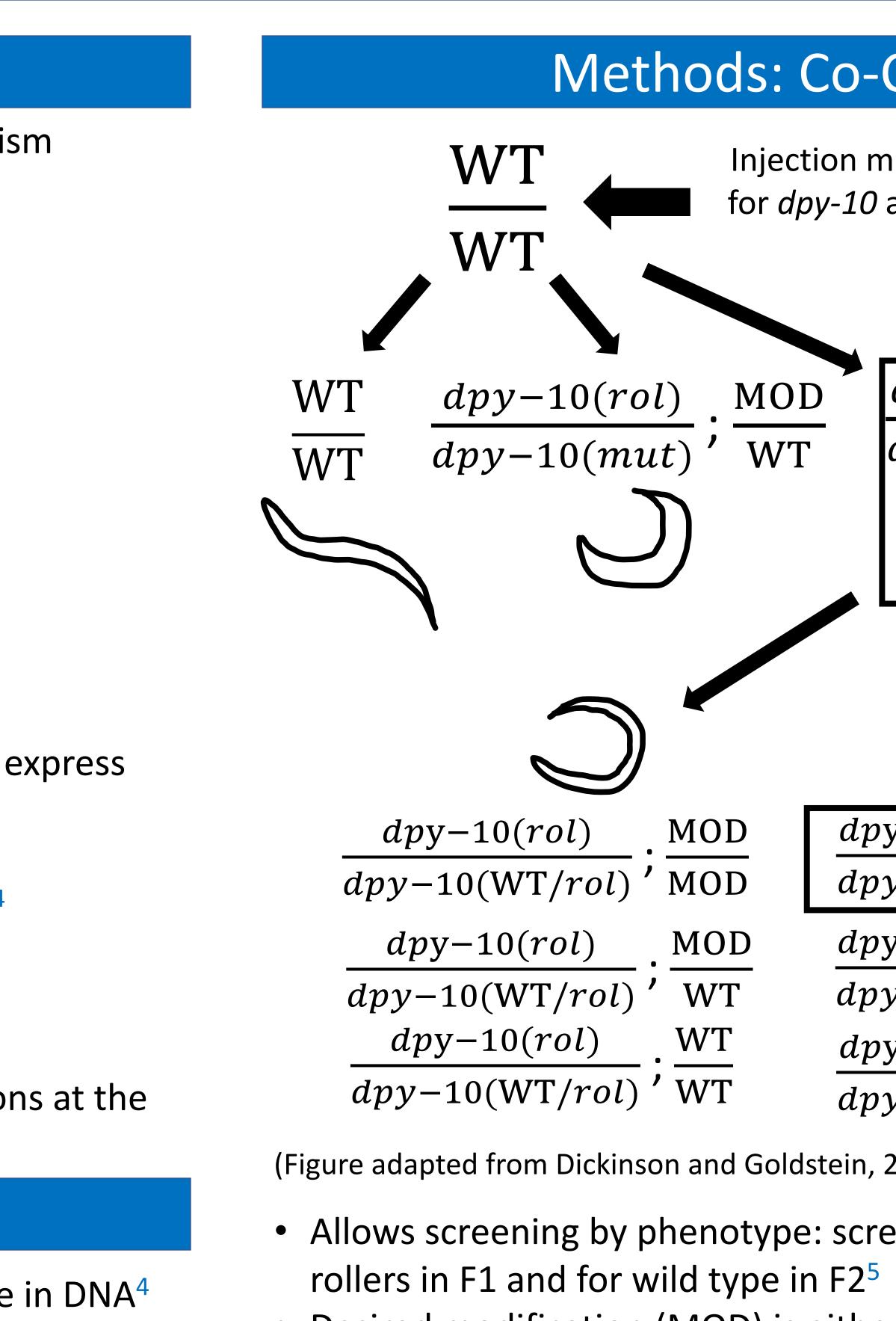
- **Odd-skipped** gene family –
- Odd-skipped related genes in mammals
- Odd-1 and odd-2 expressed in gut of C. elegans²
- Odd gene array-based³ reporter strains exist but express inconsistently.
- **CRISPR/Cas9** system used for **genetic engineering**⁴
- CRISPR clustered regularly interspaced short palindromic repeats

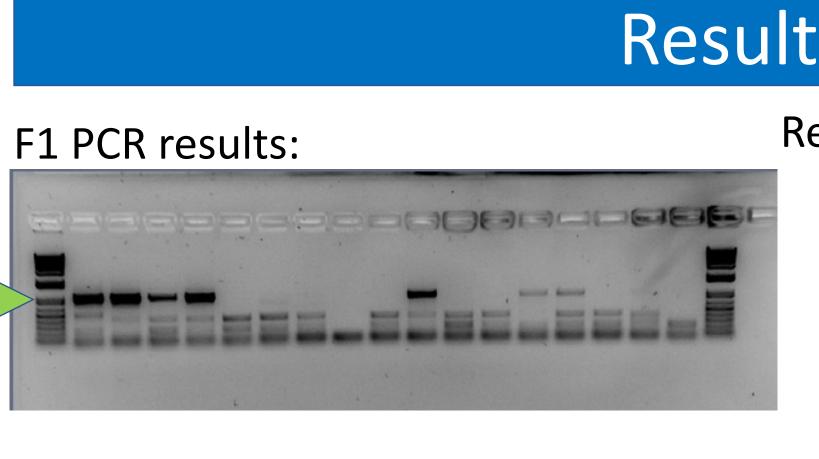
GOAL: Make reporter strains with fluorescence insertions at the beginning of the *odd-1* and *odd-2* genes

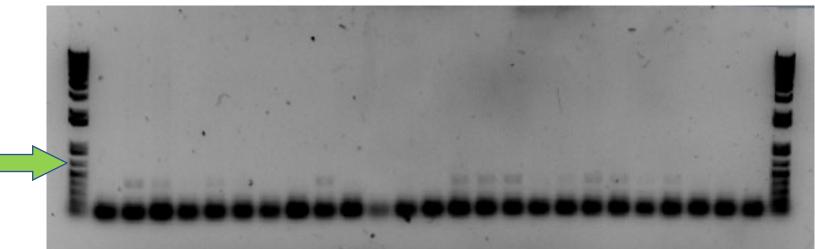
Methods: CRISPR/Cas-9

- Modified guide-RNA directs Cas9 to target sequence in DNA⁴ Both delivered via plasmid in the injection mix
- Cas9 (an endonuclease) makes double-strand breaks in DNA
- The repair sequence includes:
- Fluorescent protein (and epitope tag, if desired)
- Homology regions on both sides matching sequence around the double-strand break (HOM)
- Either short homology (40 bp) or long homology (500 bp)

	Promoter			ODD-1					DN
Cas9 target sequence									
	HOM	3X	Flag	GFP		HOM			Rep Sec
Pro	omoter <mark>3</mark> X		GF	GFP		ODD		_	– Res
									inc







epair equence

esult: DNA insertion

Not successful: MOD not incorporated into chromosome in F1s, so offspring were negative for MOD

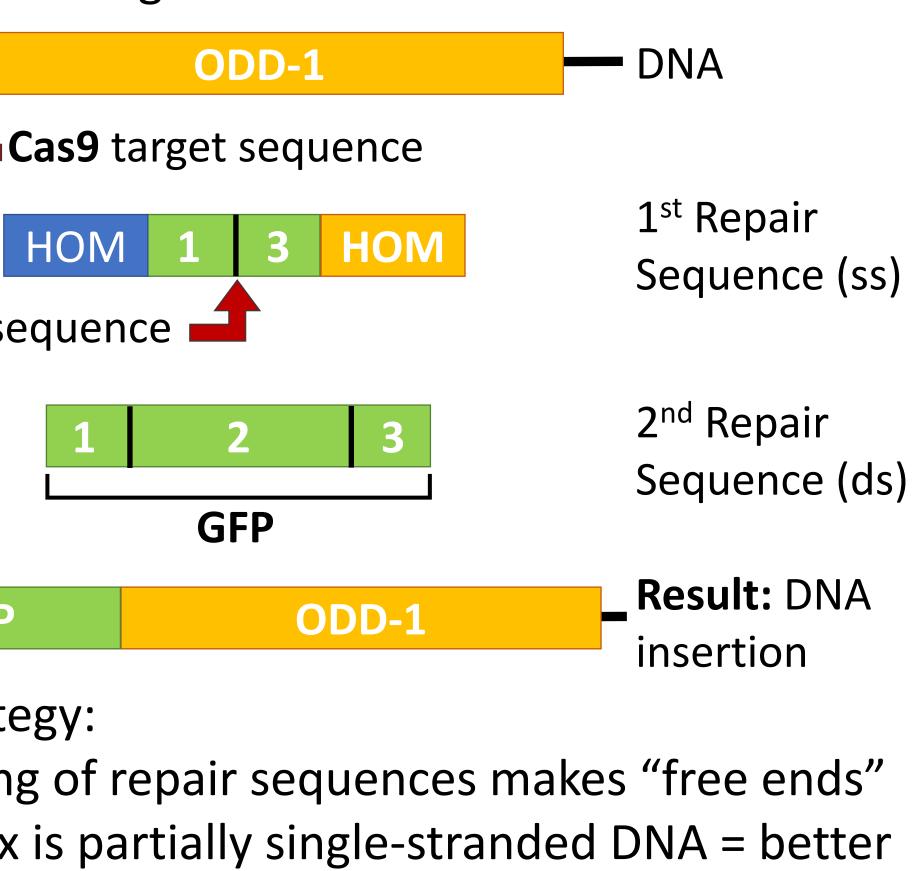
The Biology Department at ECSU and the AAUP Faculty Research Grant. Some strains were provided by the CGC, which is funded by NIH Office of Research Infrastructure Programs (P40 OD010440).

References

- 1. Wormbook.org

- 5. Dickinson, D.J. and Goldstein, B. 2016. Wormbook.





2. Buckley, M.S., Chau, J., Hoppe, P.E., and Coulter, D.E. 2004. Dev. Gen. Evo. 214:10-18. 3. Mello, C. C., Kramer, J. M., et al. 1991. EMBO Jour. 10:3959-3970. 4. Farboud, B., Severson, A. F., & Meyer, B. J. 2019. *Genetics.* **211**:431-457. 6. Vicencio, J., Martinez-Fernandez, C., Serrat, X., & Ceron, J. 2019. Genetics. 211:1143-1154. 7. Dokshin, G. A., Ghanta, K. S., Piscopo, K. M., & Mello, C. C. 2018. *Genetics.* **210**:781-787.