

Regulation of gametic gene expression across development via **RNAi-to-RNAi cascade in C. elegans** Trilotma Sen, Cara McCormick, and Alicia K. Rogers The Department of Biology, The University of Texas at Arlington, Texas

25°C

Figure 1: Temperature sensitive sterility and

lower brood size in *mut-16* mutants

400

300

200 -

Abstract

Coordination of gene regulatory networks is necessary for proper execution of cellular programs throughout development. RNA interference (RNAi) is an essential regulatory mechanism in all metazoans. Proper RNAi-mediated gene regulation requires coordination of several RNAi branches to ensure homeostasis. For example, in *Caenorhabditis elegans*, the Argonautes, ALG-3 and ALG-4, are expressed specifically during spermatogenesis (L4 stage) and bind small interfering RNAs (siRNAs) complementary to sperm-enriched genes. We find that *alg-3* and *alg-4* are regulated by siRNAs. Our work shows that gene switches are operated via these siRNAs to regulate the Argonautes' expression in a temporal manner. This RNAi-to-RNAi regulatory cascade is essential for coordinating ALG-3/4 pathway function, particularly during heat stress, to provide thermotolerant sperm-based fertility. This work provides insight into one regulatory motif used to maintain RNAi homeostasis, across developmental stages, despite environmental stressors. As RNAi pathways are evolutionarily conserved, other species likely use similar regulatory architectures to



Fluorescence microscopy was done for L4s and young







Results

Figure 2: Sperm genes are downregulated in the germline of heat-stressed L4 *mut-16* mutants







Future Directions & Impact

• Our goal is identify how the ALG-3/4 pathway is regulated and how do these pathways control sperm development?

- To dissect the regulatory architecture of regulation of *alg-3* and *alg-4*, we will use guide RNA based feeding assay to overexpress *alg-3* and *alg-4* for our future experiments. We will also use CRISPR-directed mutagenesis to uncouple the *Mutator* complex's influence of ALG-3/4 pathway.
- Together, these studies will provide a map of molecular interactions in RNAi pathway functions deepening our understanding of genetic regulation by small RNAs.
- This research would add to our incomplete understanding of underlying factors contributing to infertiliy and heritable dis-
- Such studies in model metazoans could have far-reaching translational implications for human reproductive health.

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