TEXAS WOMAN'S $\mathbf{U} \mathbf{N} \mathbf{I} \mathbf{V} \mathbf{E} \mathbf{R} \mathbf{S} \mathbf{I} \mathbf{T} \mathbf{Y}^{\mathsf{T}}$

Protective role of the *C. elegans* DBL-1/TGF-β signaling pathway in innate immune defenses against a panel of Gram-negative and Gram-positive bacteria Bhoomi J. Madhu*, Laura K. Hanson, and Tina L. Gumienny Texas Woman's University, Department of Biology, Denton, TX

Background

C. elegans exhibits different types of responses to immune challenges; the first response is avoidance of pathogens, which also shows the ability of animals to differentiate food sources, because these roundworms eat bacteria¹. The second mechanism is protection through physical (exoskeleton) and mechanical means (pharyngeal grinding of bacteria)². The third mechanism is induction of genes that include antimicrobial genes². One of the signaling pathways involved in the innate immune response is the TGF-B pathway. DBL-1 is a ligand of a TGF- β signaling pathway in C. elegans known to regulate innate immunity genes^{3,4}. Previous studies show DBL-1 regulates expression of innate immune response genes in response to Gram-negative Serratia marcescens Db11⁵. Is DBL-1 signaling required for an effective response to a range of bacterial pathogens or to a specific type of pathogen?



Figure 1. Canonical DBL-1/TGF- β signaling pathway in *C. elegans*. Adapted from Gumienny and Savage-Dunn (2005)³.

Hypothesis

We propose that DBL-1 is required to mount an effective innate immune response to a variety of pathogenic organisms. The panel of pathogenic bacteria to test our hypothesis includes:

Gram-positive bacteria	Gram-negative bacteria
Bacillus megaterium	Enterobacter cloacae
Enterococcus faecalis	Klebsiella oxytoca
Staphylococcus epidermidis	Serratia marcescens

Results





Wild-type N2 animals and *dbl-1* mutant animals were picked onto EZ plates spotted with test or control bacteria. Number of worms on the bacterial spot and off the bacterial spot at every time interval was recorded and aversion ratio was calculated (A= number of animals off the lawn/total number of animals). Assays were performed in triplicates and statistical analyses were performed by repeated measures ANOVA.

2. Does DBL-1 affect the feeding response of nematodes to pathogenic bacteria?

A: Reduced feeding by loss of DBL-1 and in response to Gram-positive bacteria is additive



Figure 3. Pharyngeal pumping rate of the three *dbl-1* genotypes of *C. elegans* to an array of Gram-negative and -positive bacteria *E. coli* OP50 is the control. Error bars represent standard deviation.*, p< 0.05 compared to the wild type on the same bacterial strain. #, p< 0.001 compared to the wild type on E. coli.

Wild-type N2 animals and *dbl-1* mutants were picked onto EZ plates spotted with test or control bacteria. The number of pharyngeal contractions were counted for 20 seconds. Two counts were made for each animal and averaged. Assays were performed in triplicates and statistical analyses were performed by two-way ANOVA.



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Inducible antibacterial defense system in *C. elegans*. *Current Biology*. 12:1209–1214.

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4. Is a DBL-1 pathway reporter affected by

A: Select pathogens further increase reporter

S. epidermidis 102%

113%

199%

* #



Figure 6. DBL-1 pathway activity reporter in wild-type and *dbl-1(-)* backgrounds fed on Gram-positive and -negative bacteria Animals expressing the green fluorescent protein reporter (texls127 [spp-9p::gfp]) in wild-type and dbl-1 mutant backgrounds were picked onto EZ plates spotted with test or control bacteria. tex/s127 is derived from wkEx52 ⁶.Statistical analyses were performed by two-way ANOVA. Error bars represent standard deviation. *, p<0.05 compared to the wild type on the same bacteria.

#, p<0.05 compared to the same worm strain on the E. coli OP50. Percent

Conclusions

Gram +: Avoidance response is independent of DBL-1 Gram -: Animals lacking DBL-1 display high avoidance

Gram +: Loss of DBL-1 makes nematodes more susceptible to

Gram -: Loss of DBL-1 makes nematodes more susceptible

Gram +: Loss of DBL-1 and infection is additive Gram -: Loss of DBL-1 and K. oxytoca infection is additive

DBL-1 is required for induction of common and specific antimicrobial genes in response to Gram-positive and -negative bacteria

Exposure to some Gram-positive and -negative bacteria further

Future directions

Characterize role of spp-9 in the context of immunity. Determine role of DBL-1-mediated effects on surface coat (1155B) and cuticle (447B), which help protect animals

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