

Role of the p38 MAP kinase pathway in *C. elegans* surface antigen switching

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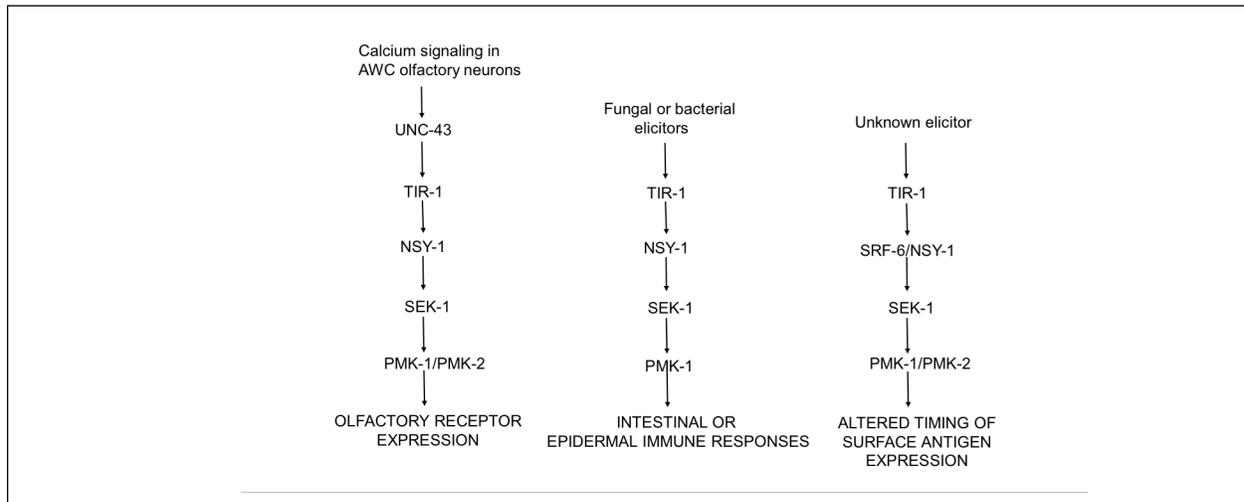


Figure 1 Involvement of the p38 MAP kinase pathway in three different processes in *C. elegans*. The p38 pathway *per se* consists of a MAPKKK (NSY-1), a MAPKK (SEK-1), and a MAPK (PMK-1). On the left, the p38 pathway is being used to determine cell fate in the AWC chemosensory neuron pair (Troemel, Sagasti, and Bargmann 1999). In the center, the p38 pathway is being used to regulate immune responses to fungal and bacterial pathogens. These include immune responses that occur in the epidermis or the intestine (reviewed in Partridge, Gravato-Nobre, and Hodgkin 2010). On the right, the p38 pathway is being used to regulate expression of an L1-specific surface epitope recognized by a monoclonal antibody (Foley et al 2019). In addition to the MAP kinase cascade, *C. elegans* processes mediated by the p38 pathway utilize upstream signaling proteins such as the adapter protein TIR-1, as well as G proteins, phospholipase C, and protein kinase C (not shown, reviewed in Partridge, Gravato-Nobre and Hodgkin 2010).

Description

In Van Sciver et al., 2019 and Honzel et al., 2019, we showed that the gene previously described as *srf-6* is actually *nsy-1*, which encodes NSY-1, the MAPKKK in the *C. elegans* p38 pathway. In earlier work, we had shown that *srf-6* mutations affect timing of expression of a surface epitope (Hemmer et al., 1991). Wild-type worms express the epitope only at the L1 stage, but *srf-6* mutants express it additionally at stages L2-L4 (called CLD for Constitutive Larval Display). In addition, we reported that wild-type worms can display the L1-specific epitope on stages L2-L4 when grown on a modified medium containing the concentrated extract of liquid nematode culture medium (called ILD for Inducible Larval Display, Grenache et al., 1996). Thus the expression of the L1-specific epitope appears to be controlled by an inducible switch that is under control of the *srf-6* gene, which, as we now know, is a component of the p38 MAP kinase pathway in *C. elegans*. The *srf-6* mutant phenotype, according to this model, corresponds to a switch that is constitutively “on”. *srf-6(yj13)* has a CLD phenotype similar to that of a large *nsy-1* deletion, suggesting that SRF-6 may function to inhibit expression of the L1-specific epitope after the L1 stage.

The modulation of this switch by an extract of liquid nematode culture medium (Grenache et al., 1996) suggested to us that it might be triggered by environmental signals detected by the nematodes’ chemical senses. Genes such as *daf-4* and *daf-7* encode components of a TGF beta pathway that control formation of the *C. elegans* dauer larva in response to dauer pheromone, which is secreted by worms and detected by worm chemosensation (reviewed in Patterson and Padgett 2000). *Daf-4* and *daf-7* mutants also show the CLD phenotype (Grenache et al., 1996). This led us to test *srf-6* mutants for chemosensory defects directly (Olsen et al., 2007). We determined that *srf-6(yj13)* mutants are defective in chemotaxis to both water-soluble and volatile attractants. Conversely, we also tested

07/04/2019 – Open Access

chemosensory mutants for ILD and found that genes required for integrity of the chemosensory ciliated nerve endings are also required for ILD (Olsen et al., 2007). However, genes required for olfaction were not required for ILD. We note that *nsy-1* is expressed in other neurons in addition to AWC (Sagasti et al., 2001), and is required in the ADF amphid neurons for pathogen-induced induction of serotonin biosynthesis (Shivers et al., 2009). The tissue of expression and time of action of *nsy-1/srf-6* in relation to ILD remain to be determined.

A clue as to how *srf-6* might modulate surface antigen expression is found in the fact that neither *srf-3(yj10)* nor *srf-3(yj10); srf-6(yj43)* double mutants show immunofluorescence of an L1-specific epitope at any developmental stage (Hemmer et al., 1991). The *srf-3* gene encodes a nucleotide sugar transporter, and the pathogenic bacteria *Yersinia* and *Microbacterium nematophilum* are unable to infect *srf-3* mutants (Hoflich et al., 2004). Furthermore, *srf-3* mutants are deficient in glycoconjugates (Cipollo et al., 2004). Thus *srf-6* might control the expression of specific glycosylation enzymes via sensing of environmental chemical conditions.

It is well established that the outer surface of parasitic nematodes is covered in a glycoprotein surface coat. Similarities can be found between the L1-specific epitope of *C. elegans* and the stage-specific expression of parasitic nematode “excretory-secretory antigens”. These are also found in association with the surface of the parasite. A good example is the *Toxocara canis* infective larva (L3), which has a glycoprotein coat composed primarily of an abundant mucin-like protein, TES120 (Page and Maizels 1992; Page, Rudin, and Maizels 1992). TES120 can also be found in the culture medium as a secretory product of this developmental stage (Page and Maizels 1992). The parasite has the ability to shed its TES120-containing surface coat in response to antibody binding (Smith et al 1981). We have found that mAb M37 staining is only visible on the surface of the *C. elegans* L1 when worms are incubated with the antibody at 0-4° C. When the sample is allowed to warm on the microscope stage, worms shed large fluorescent flakes or patches and eventually appear completely unstained (Politz and Philipp 1992). Thus a *C. elegans* surface epitope shows similar stage-specificity and ability to be released, as do surface coat molecules of parasitic nematodes. This raises the possibility that stage-specificity of the surface antigens of parasitic nematodes might also be controlled by a MAP kinase pathway.

Reagents

C. elegans strains used in this work were listed in Van Sciver et al, 2019, Honzel et al, 2019, and Foley et al, 2019. Strains will be sent to the CGC.

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Acknowledgments

I acknowledge the many contributions of researchers in the Politz lab, published and unpublished, that contributed to the story described here.

Funding

I acknowledge the Office of the Dean of Arts and Sciences, Worcester Polytechnic Institute, for partial support for this project.

Author Contributions: The author conceptualized, visualized, and wrote this article.

Reviewed by Maria Gravato-Nobre

Received 06/18/2019. **Accepted** 06/27/2019. **Published Online** 07/04/2019.

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Citation:

Politz, SM (2019). Role of the p38 MAP kinase pathway in *C. elegans* surface antigen switching. microPublication Biology. 10.17912/micropub.biology.000130