

# daf-2 modulates regeneration of mechanosensory neurons II

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## **Description:**

The *daf-2* gene encodes an insulin-like growth factor/IGF-1 receptor that regulates *C. elegans* embryonic and larval development. It has previously been shown that DAF-2 inhibits neurite regeneration of the GABAergic motor neurons and PVD sensory neurons in an age-dependent fashion [1, 2]. Following injury, the posterior lateral microtubule (PLM) neurons are capable of regenerating through axonal fusion, a highly efficient regrowth mechanism in which separated fragments fuse back together [3-6]. We previously established that a critical event for axonal fusion to occur is the exposure of injury-induced phosphatidylserine (PS) 'save-me' signals [5]. The level of PS exposed increases with advancing age [3]. To determine if *daf-2* is involved in this age-dependent modulation of PS exposure, we visualised and quantified the level of PS exposed after PLM axotomy using a secreted, tagged version of Annexin V [5, 7]. Mutation of *daf-2* had no effect on PS exposure 1 h post-axotomy, with no significant differences observed on either the distal or proximal axon segments (Table 1).

**Table 1.** Quantification of the relative level of PS exposed 1 h post-axotomy.

Genotype	PS exposed on distal axon (relative to pre-axotomy)	n	PS exposed on proximal axon (relative to pre-axotomy)	n
wild-type	$1.53 \pm 0.105$	28	$1.44 \pm 0.0855$	28
daf-2(e1370)	$1.51 \pm 0.167$	26	$1.57 \pm 0.166$	26

#### Reagents

One-day-old adult hermaphrodites were used for all experiments, and were grown under standard conditions at 20°C. The BXN301 [daf-2(e1370); smIs95(Phsp-16.2::sAnxV::mRFP); zdIs5(Pmec-4::GFP)] strain was used along with the CU4204 [smIs95(Phsp-16.2::sAnxV::mRFP); zdIs5(Pmec-4::GFP)] control strain. The daf-2(e1370) allele has been considered temperature sensitive for the dauer phenotype, but not for the long-lived phenotype. At 20°C, daf-2(e1370) animals display a greater than 2-fold increase in lifespan compared to the wild-type [8]. Laser axotomy, microscopy and quantification of data was performed as previously described [3].

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12/01/2017 - Open Access

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#### **Funding:**

This work was supported by National Health and Medical Research Council (NHMRC) Project Grant 1101974.

### Acknowledgements

We thank Ding Xue for sharing strains.

## Reviewed by Rachid El Bejjani

**Received** 11/10/2017, **Accepted** 11/26/2017. **Available** starting <u>WormBase</u> release WS264, **Published Online** 12/01/2017.

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Citation: Abay, ZC; Yu-Ying Wong, M; Neumann B. (2017): daf-2 modulates regeneration of mechanosensory neurons II. Micropublication: biology. Dataset. https://doi.org/10.17912/W2SM1T