

Multiple sensory G proteins in the olfactory, gustatory and nociceptive neurons modulate longevity in *Caenorhabditis elegans*

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Abstract

The life span of the nematode *Caenorhabditis elegans* is under control of sensory signals detected by the amphid neurons. In these neurons, *C. elegans* expresses at least 13 $G\alpha$ subunits and a $G\gamma$ subunit, which are involved in the transduction and modulation of sensory signals. Here, we show that loss-of-function mutations in the $G\alpha$ subunits *odr-3*, *gpa-1* and *gpa-9*, in the $G\gamma$ subunit *gpc-1* and the introduction of extra copies of the $G\alpha$ subunit *gpa-11* extend the life span of *C. elegans*. Loss-of-function of *odr-3* and extra copies of *gpa-11* act synergistically and can together extend life span more than two-fold, indicating that sensory signals play an important role in regulating life span. We show that *gpa-1*, *gpa-11*, *odr-3* and *gpc-1* all signal via the *daf-16* FOXO family transcription factor. In addition, *odr-3* and *gpa-11* might suppress life span extension partially independent of the insulin/IGF-1 like receptor homologue *daf-2*. Our results suggest that the previously unanticipated nociceptive ASH and/or ADL neurons regulate longevity. We expect that the implication of specific G proteins will eventually contribute to the identification of the sensory cues that determine the rate of aging in *C. elegans*.

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Introduction

Genetic analysis of the life span of the nematode *Caenorhabditis elegans* has greatly aided in understanding the mechanisms that regulate aging. An intriguing finding is that *C. elegans* life span is under control of sensory perception (Apfeld and Kenyon, 1999; Alcedo and Kenyon, 2004). *C. elegans* perceives its environment through 12 bilateral pairs of ciliated amphid neurons. Mutations that disrupt the cilia of these neurons, resulting in the inability to respond to sensory cues, increase the life span of *C. elegans* (Apfeld and Kenyon, 1999). Furthermore, laser ablation of some of the neurons that detect environmental cues, such as the AWA, AWC, ASI and ASG neurons, extends life span (Alcedo and Kenyon, 2004). However, the sensory cues and precise sensory signaling mechanisms that regulate longevity are not yet known.

As in mammals, many sensory cues in *C. elegans* are probably detected by G protein coupled receptors, which, in response, activate multiple G protein mediated signaling cascades (Troemel et al., 1995; Jansen et al., 1999). The genome of *C. elegans* encodes 21 $G\alpha$ subunits, 2 $G\beta$ subunits and 2 $G\gamma$ subunits (Jansen et al., 1999; Cuppen et al., 2003). Of these, 13 $G\alpha$ subunits and 1 $G\gamma$ subunit are specifically expressed in the amphid sensory neurons (Table 1; Zwaal et al., 1997; Roayaie et al., 1998; Jansen et al., 1999, 2002). Thus far, for eight $G\alpha$ subunits and the $G\gamma$ subunit a role in sensory perception has been demonstrated (Zwaal et al., 1997; Roayaie et al., 1998; Jansen et al., 1999, 2002; Chao et al., 2004; Hilliard et al., 2004; Lans et al., 2004; Hukema et al., 2006). For example, the $G\alpha$ subunit ODR-3 mediates the detection of olfactory and nociceptive stimuli (Roayaie et al., 1998; Hilliard et al., 2004). Furthermore, the $G\alpha$ subunits ODR-3 and GPA-1 and the $G\gamma$ subunit GPC-1 play a role in gustatory plasticity, a behavioral response to NaCl (Jansen et al., 2002; Hukema et al., 2006). The $G\alpha$ subunit GPA-11 has been shown to regulate octanol sensitivity in response to a food/serotonin signal (Chao et al., 2004).

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Table 1
Mean life spans of G protein mutant and transgenic strains, at 25 °C^a

Strain ^b	Mean±SEM (days)	75th perc. (days) ^c	n ^d	p value ^e	Amphid neuron expression ^f	Sensory function ^f
<i>Amsterdam</i> ^g						
Wild type	13.4±0.2	15	203			
<i>gpa-1</i>	14.4±0.3	16	78	0.007	ADL, ASH, ASI, ASJ	Salt perception
<i>gpa-1XS</i>	12.3±0.4	14	44	0.004		
<i>gpa-2</i>	9.5±0.6	12	40	<0.0001	AWC	Olfaction, dauer pheromone
<i>gpa-2XS</i>	15.1±0.4	17	73	<0.0001		
<i>gpa-3</i>	11.4±0.5	14	40	0.001	AWA, AWC, ADF, ADL, ASE, ASG, ASH, ASI, ASJ, ASK	Nociception, olfaction, dauer pheromone
<i>gpa-3XS</i>	13.1±0.3	15	39	0.144		
<i>gpa-3QL</i>	11.6±1.2	14	28	0.432		
<i>gpa-4</i>	14.1±0.4	16	43	0.053	ASI	
<i>gpa-4XS</i>	12.7±0.6	15	44	0.981		
<i>gpa-5</i>	14.2±0.4	16	43	0.036	AWA, ADL	Olfaction
<i>gpa-5XS</i>	12.1±0.5	14	42	0.052		
<i>gpa-6</i>	12.0±0.5	13	41	0.036	AWA, AWB, ASH, ADL	
<i>gpa-6XS</i>	13.0±0.5	15	46	0.912		
<i>gpa-7</i>	13.4±0.5	15	36	0.976		
<i>gpa-7XS</i>	14.1±0.4	16	22	0.465		
<i>gpa-8</i>	10.7±0.5	13	35	<0.001	None	
<i>gpa-8XS</i>	11.5±0.5	13	42	0.003		
<i>gpa-9</i>	17.5±0.4	19	42	<0.0001	ASJ	
<i>gpa-9XS</i>	11.0±0.6	14	41	0.001		
<i>gpa-10</i>	11.0±0.3	12	50	<0.0001	ADF, ASI, ASJ	
<i>gpa-11</i>	12.8±0.4	18	46	0.278	ADL, ASH	Serotonin/food
<i>gpa-11XS</i>	19.5±0.7	26	65	<0.0001		
<i>gpa-12</i>	13.2±0.5	16	37	0.906	Undetermined	
<i>gpa-13</i>	12.9±0.4	16	38	0.261	ADF, ASH, AWC	Olfaction
<i>gpa-13XS</i>	12.4±0.7	16	46	0.727		
<i>gpa-14</i>	13.7±0.4	16	44	0.668	ASI, ASJ, ASH, ASK	
<i>gpa-14XS</i>	12.9±0.3	13	44	0.118		
<i>gpa-15</i>	13.3±0.4	16	46	0.907	ADL, ASH, ASK	
<i>gpa-15XS</i>	11.1±0.6	13	21	<0.0001		
<i>odr-3</i>	18.0±0.4	19	80	<0.0001	AWA, AWB, AWC, ADF, ASH	Olfaction, salt, nociception
<i>gpc-1</i>	14.7±0.4	17	45	0.002	AWB, ADL, ASH, ASI, ASJ, AFD	Salt perception
<i>gpc-1XS</i>	15.2±0.5	17	33	0.0001		
<i>gpa-11; odr-3</i>	14.3±0.5	17	35	*0.037	**<0.0001	
<i>Rotterdam</i> ^b						
wild type	17.5±0.3	20	89			
<i>gpa-9</i>	19.6±0.3	22	87	<0.0001		
<i>gpa-9XS</i>	19.4±0.4	21	71	0.0001		
<i>gpa-11</i>	17.0±0.6	19	37	0.843		
<i>gpa-11XS</i>	21.1±0.5	24	90	<0.0001		
<i>gpa-11XS(pkIs540)</i>	22.8±0.6	24	93	<0.0001		
<i>odr-3</i>	19.7±0.5	24	84	<0.0001		
<i>gpa-11; odr-3</i>	17.2±0.4	19	69	*0.823	**<0.0001	
<i>odr-3; gpa-11XS</i>	34.7±1.6	42	48	**<0.0001	***<0.0001	
<i>odr-3; gpa-11XS(pkIs540)</i>	25.4±1.1	32	42	**<0.0001	°0.132	

^a Cumulative results of life span analysis at 25 °C.

^b All strains carry loss-of-function mutations that represent putative null alleles. This has been verified using antibodies in the case of *odr-3*, *gpa-11*, *gpa-3*, *gpa-5*, *gpa-6*, *gpa-13* and *gpc-1* (Lans et al., 2004; S.R. and G.J., personal communication). Strains that have significantly extended life span are indicated in bold. Two independent integrated arrays of *gpa-11XS* were tested, to confirm that additional copies of the *gpa-11* gene caused life span extension. None of the strains, except *gpa-3QL*, shows defective dye filling (Jansen et al., 1999, 2002). XS denotes extra copies of the gene as a transgene, QL a constitutive activating mutation.

^c The 75th perc. (percentile) is the age when the fraction of animals alive reaches 0.25.

^d n denotes number of animals tested.

^e p values are compared to wild type, except * compared to *gpa-11*, ** compared to *odr-3*, *** compared to *gpa-11XS* and ° compared to *gpa-11XS(pkIs540)*.

^f Expression patterns and functions were published previously (Zwaal et al., 1997; Roayaie et al., 1998; Jansen et al., 1999, 2002; Yau et al., 2003; Chao et al., 2004; Hilliard et al., 2004; Lans et al., 2004; Hukema et al., 2006).

^g During the course of our experiments, our lab moved from the city of Amsterdam to Rotterdam. Animals survived significantly longer in Rotterdam, possibly because of different environmental conditions. Results obtained at the same location, however, never showed large variations, emphasizing their significance. Importantly, the outcome of experiments performed at both locations was similar: animals that showed extended life span in Amsterdam showed extended life span in Rotterdam. All assays in Amsterdam were performed once, except assays with *gpa-1*, *gpa-2XS*, *gpa-4XS*, which were performed twice and with wild type, which were performed six times. All assays in Rotterdam were performed twice except assays with *gpa-11*, *odr-3*; *gpa-11XS* and *odr-3*; *gpa-11XS(pkIs540)*, which were performed once.

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