Cellular/Molecular

Extrasynaptic Muscarinic Acetylcholine Receptors on Neuronal Cell Bodies Regulate Presynaptic Function in Caenorhabditis elegans

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Acetylcholine (ACh) is a potent neuromodulator in the brain, and its effects on cognition and memory formation are largely performed through muscarinic acetylcholine receptors (mAChRs). mAChRs are often preferentially distributed on specialized membrane regions in neurons, but the significance of mAChR localization in modulating neuronal function is not known. Here we show that the Caenorhabditis elegans homolog of the M1/M3/M5 family of mAChRs, gar-3, is expressed in cholinergic motor neurons, and GAR-3-GFP fusion proteins localize to cell bodies where they are enriched at extrasynaptic regions that are in contact with the basal lamina. The GAR-3 N-terminal extracellular domain is necessary and sufficient for this asymmetric distribution, and mutation of a predicted N-linked glycosylation site within the N-terminus disrupts GAR-3-GFP localization. In transgenic animals expressing GAR-3 variants that are no longer asymmetrically localized, synaptic transmission at neuromuscular junctions is impaired and there is a reduction in the abundance of the presynaptic protein sphingosine kinase at release sites. Finally, GAR-3 can be activated by endogenously produced ACh released from neurons that do not directly contact cholinergic motor neurons. Together, our results suggest that humoral activation of asymmetrically localized mAChRs by ACh is an evolutionarily conserved mechanism by which ACh modulates neuronal function.

Introduction

Muscarinic acetylcholine (ACh) receptors are G-protein coupled receptors (GPCRs) that mediate many of the modulatory functions of ACh in the CNS. Muscarinic signaling affects multiple behaviors—including alertness, cognition, memory, and motor control—by altering neuronal excitability, synaptic protein composition, and the strength of communication between neurons and other tissues (Picciotto et al., 2012). The five subtypes of mammalian muscarinic ACh receptors (mAChRs), M1-M5, differ in ligand affinity, subcellular localization, and signaling capacity. Generally, M1, M3, and M5 couple to the $G\alpha q/11$ heterotrimeric G-protein, and their activation leads to an increase in neuronal activity; M2 and M4, on the other hand, couple to $G\alpha i/o$ and negatively regulate neurotransmitter release from presynaptic terminals (Eglen, 2006; Wess et al., 2007).

Muscarinic receptor subtypes are differentially distributed on specialized membrane regions within neurons (Levey et al., 1991; Rouse et al., 1998; Oldenburg and Ding, 2011). Immunohisto-

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chemical studies of M1 receptors, for example, show preferential localization on cell bodies and dendrites (Yamasaki et al., 2010), whereas M2 receptor abundance is generally greater on axon terminals (Rouse et al., 1997). At these specialized membrane regions, mAChRs can be found at either synaptic or extrasynaptic sites. For example, ultrastructural studies show that M2/M4 receptors can localize to extrasynaptic sites of presynaptic terminals, where they can function as autoreceptors or heteroreceptors (Rouse et al., 2000; Picciotto et al., 2012). On the other hand, M1 receptors can localize to extrasynaptic regions on dendritic spines and shafts (Mrzljak et al., 1993; Yamasaki et al., 2010), where they primarily function as heteroreceptors. The highly specialized mAChR receptor distribution in the brain suggests that receptor localization may be a critical component underlying cholinergic modulation of neuronal physiology.

Extrasynaptic receptors can respond to ACh released from distant sources, such as neurotransmitter that escapes the synapse (spillover) or diffuses through the extracellular space from distant sites (Agnati et al., 2010; Vizi et al., 2010). This signaling, termed "volume transmission," is supported by observations that central cholinergic nerve endings are distributed at lower densities than AChR binding sites, and cholinergic terminals are often not associated with postsynaptic membrane specializations (Mrzljak et al., 1993; Descarries and Mechawar, 2000). In volume transmission, one neuron secreting ACh can act on extrasynaptic mAChRs of many targets to modulate aspects of cholinergic signaling, such as those associated with arousal and attention (Sarter et al., 2009; Vizi et al., 2010; Fuxe et al., 2012). However, the molecular mechanisms underlying mAChR localization and the

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functional importance of mAChR localization in neurons are still unclear.

Here, we show that a *Caenorhabditis elegans* mAChR M1/M3/M5 family member, GAR-3, is asymmetrically enriched on extrasynaptic membrane surfaces of motor neuron cell bodies, and this asymmetric localization is necessary for its effects on neuromuscular function. We find that GAR-3 is in contact with the pseudocoelom, a cavity analogous to the circulatory system, and endogenous ACh released from a distance can activate GAR-3 on motor neurons. We propose a model whereby ACh released into the pseudocoelom activates GAR-3 to modulate cholinergic neurotransmission.

Materials and Methods

C. elegans strains. All behavioral experiments were performed on young adult hermaphrodites, unless otherwise noted. Some strains were provided by the Caenorhabditis Genetic Center, which is funded by NIH Office of Research Infrastructure Programs (P40 OD010440). The wild-type reference strain was N2 Bristol. The following strains were used: gar-3(gk305), nre-1(hd20) lin15b(hd126), unc-2(lj1), unc-13(s69), unc-17(e245), unc-80(e1272), unc-104(e1265), nuIs197[Pmyo-2::GFP, Punc-129::sphk-1-GFP], nuIs321[Punc-17::mCherry], vjIs50[Pttx-3:mRFP, Punc-129::GAR-3-GFP], vjIs14[Punc-129::mCherry], vjSi06[Punc-129::gar-3-gfp, cb-unc-119(+)]. All strains used were outcrossed >4 times and integrated transgenes were outcrossed 10 times.

Molecular biology. The gar-3 promoter fragment was amplified by PCR from mixed stage genomic DNA, and GAR-3 chimeras were made using gar-3b and gar-1 cDNA. All primer sequences are available upon request. The following plasmids were made by standard molecular biology techniques (plasmid name[promoter-gene-marker]): pDS221[Punc-129::sphk-1-gfp], pDS322[Punc-129::gar-3b-gfp], pDS333[Punc-17:: unc-17], pDS324[Punc-17::gar-3b], pDS336[Punc-129::gar-1-gfp], pDS339[Punc-129::gar-3(NT3)-gfp], pDS340[Punc-129::gar-3(2NQ)-gfp], pDS341[Punc-129::gar-1-3-gfp], pDS342[Punc-129::gar-3-1-gfp], pDS343 [Punc-129::gar-3(1i3)-gfp], pDS345[Punc-129::gar-3(N33Q)-gfp], pDS346[Punc-129::gar-3(N28Q)-gfp], pDS347[Punc-17::gar-3b-gfp], pDS348[Punc-129::gar-3(NT1)-gfp], pDS349[Punc-129::gar-3(NT2)gfp], pDS354[Psra-6::mCherry], pDS364[Pgar-3(8.5kB)::gfp], pDS368 [*Pgar-3*(8.5kB)::gar-3b-gfp], pDS369[*Pdfk-2b::mCherry*], pDS371[*Punc-17:*: gar-1], pDS372[Punc-17::gar-1-3b], pDS373[Pmyo-3::gar-3-gfp], pDS374 [*Pmyo-3::gar-1-3-gfp*], pDS399[*Punc-47::mCherry*], pDS400[*Pglr-1::* mCherry], pDS401[$Punc-17(\Delta)$::mCherry]. pDS402[$Punc-17(\Delta)$::unc-17]. To make a MosSCI plasmid (pDS404), Punc-129::gar-3b-gfp was cloned into pCFJ151.

Transgenic lines. Transgenic strains were generated by injecting either N2, gar-3(gk305), or nuIs197 animals with expression constructs (10–25 ng/µl) and a coinjection marker [KP#708 (Pttx-3::rfp, 40 ng/µl) or KP#1106 (Pmyo-2::nls-gfp, 10 ng/µl)]. Microinjection was performed using standard techniques as previously described (Mello et al., 1991). At least three lines for each transgene were examined and a single, representative transgene was used for the experiments. The following transgenic lines were made (name-[plasmid]): vjEx579[PDS333], vjEx601[pDS364], vjEx602[pDS368], vjEx603[pDS369], vjEx635[pDS324], vjEx732[pDS322], vjEx733[pDS340], vjEx734[pDS341], vjEx735[pDS342], vjEx736[pDS343], vjEx737[pDS345], vjEx738[pDS346], vjEx741[pDS354], vjEx742[pDS374], vjEx743[pDS371], vjEx744[pDS372], vjEx745[pDS347], vjEx746[pDS399], vjEx747[pDS400], vjEx748[pDS373], vjEx749[pDS348], vjEx750[pDS349], vjEx751[pDS339], vjEx753[pDS336], vjEx754[pDS401], vjEx755[pDS402].

Behavioral analysis. For analysis of neurotransmission, animals were tested for their sensitivity to aldicarb (Bayer CropScience), an inhibitor of acetylcholinesterase. Animals paralyze after acute exposure to 1.0 mm aldicarb, and paralysis of adult worms was scored every 10 min, starting at 40 min. For each experiment, animals were exposed to NGM plates supplemented with aldicarb, using ~20 worms per plate in triplicate. The number of worms paralyzed on each plate was counted to extract the percentage paralyzed at each time-point per genotype. The percentages were averaged at each time point per genotype and plotted graphically. For each experiment, the genotype was blind to the scorer, and the anal-

ysis was repeated at least two times. For experiments using arecoline, worms were pretreated with plates containing either control M9 or arecoline (15 mm in M9) for 2 h. This pretreatment duration was determined to be the minimum concentration that produced maximal aldicarb hypersensitivity, to minimize nonspecific effects of the drug.

RNAi. RNAi feeding was performed as previously described (Kamath et al., 2001). Briefly, RNAi feeding clones were grown for 12 h at 37°C in LB culture supplemented with 50 μ g/ml carbenicillin. RNAi clones were then seeded onto agar plates containing IPTG (1 mm) and carbenicillin (50 μ g/ml). Eggs were isolated from adult state worms [vjIs50[Punc-129::gar-3b-gfp];nre-1(hd20) lin-15b(hd126)] using a sodium hypochlorite solution. Eggs were washed with M9, and \sim 100 eggs were added to each seeded RNAi plate. After 3-4 d, animals were removed from bacteria and assessed for GAR-3-GFP localization. To ensure effective RNAi, we screened animals fed the gar-3 RNAi clone, as a positive control, which resulted in a 64% reduction in fluorescence compared with feeding with an empty RNAi vector (L4440). For each RNAi treatment, we determined the number of motor neurons in which GAR-3-GFP adopted a uniform fluorescence distribution. Approximately 11.3% (6/53) of cells examined in animals not treated with RNAi expressed nonlocalized GAR-3-GFP. Statistical differences were determined by χ^2 (ψ^2) test, and significance was set at p < 0.05. Results are reported in Table 1.

Microscopy and analysis. Fluorescence microscopy experiments were performed as previously described (Ch'ng et al., 2008). Briefly, adult worms (unless otherwise stated) were paralyzed using 2,3-Butanedione monoxime (30 μ g/ μ l, Sigma-Aldrich) and mounted on 2% agarose pads for imaging. For all fluorescence microscopy experiments, images were captured with a Nikon eclipse 90i microscope equipped with a Nikon PlanApo 100× objective (NA = 1.4) and a Photometrics Coolsnap ES camera. MetaMorph 7.0 software (Universal Imaging; Molecular Devices) was used to analyze images.

For analyses of GAR-3-GFP on cell bodies, images were captured from the ventral motor neurons throughout the anterior-posterior axis of the animals. For quantification of asymmetry, pixel intensities were recorded manually (using MetaMorph), and the background fluorescence was subtracted. For quantification of total GAR-3-GFP membrane fluorescence, pixel values were recorded along the surface of the neuronal cell body manually (using MetaMorph), averaged, and the background fluorescence was subtracted for the final fluorescence value. For treatment of animals with drugs, animals were dissected as described previously (Richmond, 2006) to expose the extracellular matrix (ECM). Either control solution (M9), collagenase type IV (Sigma-Aldrich), or trypsin (Cellgro) were added to the worm before analysis of GAR-3-GFP. For colocalizing GAR-3-GFP with wheat germ agglutinin (WGA; Alexa Fluor 594; Invitrogen), the WGA conjugate was injected into the pseudocoelomic region, allowed 15 min for recovery, and analyzed. For analysis following RNAi treatment, DA/DB cells throughout the A/P axis of the animal were scored as either asymmetrical or uniform.

For analysis of SPHK-1-GFP at synapses, serial image stacks were captured from dorsal axons of DA/DB neurons near the posterior gonadal bend of adult worms, and the maximum intensity projection was used for analysis of the dorsal cords. Line scans of the maximum intensity projection image were also recorded using MetaMorph. The fluorescence intensity values were then quantified using Puncta 6.0 software written with Igor Pro (Wavemetrics), as previously described (Dittman and Kaplan, 2006; Ch'ng et al., 2008). Quantification spanned ~500 synapses for all experiments, and specific n values are labels in graphs. For all experiments, fluorescence values were normalized to the values of 0.5 µm FluoSphere beads (Invitrogen) captured during each imaging session. This was performed to provide a standard for comparing absolute fluorescence levels between animals from different sessions. We found that the conditions used for imaging showed that SPHK-1-GFP were predominantly localized to synaptic puncta, as determined previously (Chan et al., 2012). There was little or no difference between their interpunctal fluorescence (in axons) and the auto-fluorescence observed at *C*. elegans axons. Therefore, we excluded the interpunctal fluorescence in our analyses of SPHK-1-GFP.

Table 1. RNAi analysis of GAR-3-GFP asymmetry

C. elegans gene	Cosmid	Protein ortholog	% Cells uniform	Total cells examined	ψ^2
Cell polarity					
lgl-1	F56F10.4	Lethal giant larvae	27.3	22	2.944
lin-2	F17E5.1a	(MAGUK) family	6.7	30	0.476
mom-2	F38E1.7	Wnt family	40.0	30	9.277*
mom-5	T23D8.1	Frizzled receptor	7.1	28	0.359
mig-5	T05C12.6	Disheveled	8.3	24	0.158
par-1	H39E23.1a	Serine-threonine kinase	10.0	50	0.047
par-2	F58B6.3a	E3 ubiquitin ligase	9.6	73	0.100
par-5	M117.2	14-3-3	12.8	39	0.048
Unknown	C25A8.5	FER kinase	5.6	18	0.503
vang-1	B0410.2	Strabismus	7.1	28	0.359
Glycosylation					
aman-3	F48C1.1	Processing glycosydase	13.8	29	0.107
dad-1	F57B10.10	Oligosaccharyltransferase complex	11.4	35	0.000
gly-13	B0416.6	N-acetylglycosaminyl-transferase	23.3	30	2.093
gly-14	M01F1.1	N-acetylglycosaminyl-transferase	18.9	37	1.018
gly-20	C55B7.2	β 1,6 GnT V	15.8	57	0.466
ostd-1	M01A10.3	Oligosaccharyltransferase complex	26.7	60	4.227*
Unknown	T22D1.4	OST1/ribophorin	11.4	35	0.000
Unknown	F41D3.6	Galectin	22.9	48	2.419
Unknown	C47E12.3	lpha 1,2 mannosidase l	15.0	20	0.182
Unknown	F13H10.4	lpha1,2 glycosidase l	11.8	17	0.003
Unknown	F26H9.8	Glycoprotein GlcT	17.4	23	0.517
Unknown	ZC506.1	Processing glycosydase	5.4	37	0.941
Unknown	T03G11.4	Processing glycosydase	14.0	43	0.150
Unknown	ZK686.3	Oligosaccharyltransferase complex	6.5	31	0.538
Unknown	T12A2.2	Oligosaccharyltransferase complex	12.8	39	0.048
Unknown	T19B12.6	Heparin sulphate	18.2	33	0.797

GAR-3-GFP was analyzed following RNAi knockdown of select genes involved in cell polarity and glycosylation. The effect of the RNAi treatment was calculated by scoring the number of non-asymmetric cells (equal fluorescence throughout the motor neuron cell body) over the total number of cells examined. We analyzed the neurons throughout the A/P axis, as depicted in Figure 2. A χ^2 test was performed based on the number of uniform cells observed in WT animals (6/53, 11.3%).

Statistical analysis. A Student's t test was used to determine significance when comparing the fluorescence of SPHK-1-GFP or GAR-3-GFP in different conditions (significance was set at p < 0.05 and determined by Student's t tests). A χ^2 test was performed to determine the significance of GAR-3-GFP asymmetry in the RNAi experiments. SEM (SEM) was performed for all comparisons and displayed graphically. Quantification of synaptic marker abundance at puncta (punctal fluorescence) and synaptic marker density in axons (density) were performed as previously described (Ch'ng et al., 2008) and are useful measures of the structural and functional properties of presynaptic terminals.

Results

gar-3 functions in cholinergic motor neurons to regulate aldicarb responsiveness

GAR-3 signaling regulates locomotion and the recruitment of SPHK-1/sphingosine kinase to presynaptic terminals of cholinergic neurons at neuromuscular junctions (NMJs; Dittman and Kaplan, 2008; Chan et al., 2012). To determine in which cells GAR-3 acts to regulate NMJ function, we investigated the expression pattern of a transcriptional gar-3 reporter (P gar-3::gfp) in which an 8.5 kb promoter fragment immediately upstream of the gar-3 start site drives the expression of gfp. This promoter fragment is longer than promoters previously used to examine gar-3 localization (Fig. 1A; Steger and Avery, 2004; Liu et al., 2007). In transgenic animals expressing P gar-3::gfp, fluorescence was detected in body wall and pharynx muscles, as well as in motor neurons of the ventral nerve cord (Fig. 1A,B and data not shown). GFP fluorescence was detected in all cholinergic motor neurons of the ventral nerve cord (labeled with mCherry) but not in GABAergic motor neurons (Fig. 1B).

To determine whether GAR-3 functions in cholinergic motor neurons or in body wall muscles to regulate NMJ function, we examined the responses of gar-3 mutants to drugs that promote neurotransmission at NMJs. Aldicarb is an acetylcholine esterase inhibitor that causes the accumulation of ACh at cholinergic synapses, which leads to muscle hypercontraction and paralysis (Miller et al., 1996; Mahoney et al., 2006). Conditions or mutations that delay aldicarb-induced paralysis (resistance to aldicarb) generally reduce ACh release from motor neurons or reduce the excitability of body wall muscles. Conversely, conditions or mutations that accelerate the rate of aldicarbinduced paralysis (hypersensitivity to aldicarb) generally have increased neurotransmitter release or enhanced muscle excitability (Miller et al., 1996; Mahoney et al., 2006). gar-3(gk305) mutants, which have a 452 nucleotide deletion predicted to remove the C-terminal 153 aa of the receptor, were resistant to aldicarb compared with wild-type controls (Fig. 1C; Chan et al., 2012). Expression of full-length gar-3b cDNA in cholinergic motor neurons (using the unc-17 promoter) caused gar-3 mutants to become hypersensitive to aldicarb, whereas expression of gar-3b cDNA in muscles (using the myo-3 promoter) did not cause aldicarb hypersensitivity (Fig. 1*C*,*D*).

The nicotinic agonist levamisole activates receptors that cause body wall muscle contraction, and levamisole treatment results in paralysis due to muscle hypercontraction (Lewis et al., 1980). The rate of paralysis of gar-3 mutants treated with levamisole was similar to that of wild-type controls, and expression of gar-3 cDNA in muscles did not alter responses of gar-3 mutants to levamisole (Fig. 1E). Together, these data suggest that gar-3 mu-

^{*}Significant genes (p < 0.05).

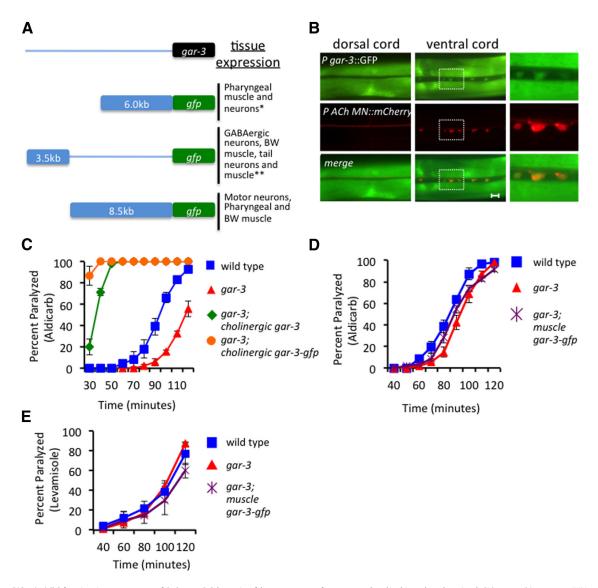


Figure 1. GAR-3/mAChR functions in motor neurons of *C. elegans. A*, Schematics of the *gar-3* reporter fragments analyzed in this study and previously (*Steger and Avery, 2004; ***Liu et al., 2007), with the tissues with reported *gar-3* expression indicated. *B*, Representative images showing the expression of the *gar-3* reporter *vjEx601*[*Pgar-3(8.5 kb)::GFP*] in cholinergic motor neurons and body wall muscles. *C*, *D*, Rates of paralysis of wild-type, *gar-3* mutants, and transgenic *gar-3* mutants upon exposure to the acetylcholine esterase inhibitor aldicarb (1.0 mm). Cholinergic denotes *vjEx635*[*Punc-17::gar-3b]* or *vjEx745*[*Punc-17::gar-3b-gfp*]. Muscle denotes *vjEx748* [*Pmyo-3::gar-3b-gfp*]. *E*, Paralysis of wild-type and *gar-3* mutants upon exposure of the cholinergic agonist levamisole (200 μm). Scale bar, 50 μm. *C*, *D*, Error bars indicate ± SEM.

tants have defects in ACh release from motor neurons. This is consistent with our previous findings that *gar-3* functions in cholinergic neurons to recruit SPHK-1/sphingosine kinase to presynaptic terminals (Chan et al., 2012). However, our data do not rule out other levamisole-independent aspects of GAR-3 function in body wall muscles.

GAR-3 adopts asymmetric localization on motor neuron cell bodies

We next examined the subcellular localization of a C-terminally GFP-tagged GAR-3 fusion protein (GAR-3-GFP). Expression of GAR-3-GFP in all cholinergic neurons (using the *unc-17* promoter) of *gar-3* mutants caused similar hypersensitivity to aldicarb as untagged GAR-3 (Fig. 1*C*), suggesting that this fusion protein is fully functional. GAR-3-GFP adopted a striking pattern of localization in motor neurons and muscles when its expression was driven by the 8.5 kb *gar-3* promoter: Fluorescence was observed in the ventral nerve cord and in motor neuron

cell bodies in discrete patches that run parallel to the ventral nerve cord (Fig. 2*A*).

To determine the localization pattern of GAR-3-GFP more precisely, we expressed GAR-3-GFP either in a subset of cholinergic motor neurons (the DA/DB class, using the unc-129 promoter) or in muscles (using the myo-3 promoter). In motor neurons, GAR-3-GFP primarily localized to narrow regions that extended longitudinally along the dorsal side of each motor neuron cell body (Fig. 2A). This pattern is suggestive of cell surface localization of GAR-3-GFP because the fluorescent regions were found on the outer edge of cell bodies marked by soluble mCherry (Fig. 2B). Similar localization patterns were observed in DA/DB motor neurons of animals expressing single-copy GAR-3-gfp (by MosSCI insertion of Punc-129::GAR-3-gfp; Fig. 2A). In body wall muscles, GAR-3-GFP fluorescence was also asymmetrically distributed at sites of contact between muscle cells, on muscle arms, and in postsynaptic regions where muscle arms contact the dorsal and ventral nerve cords (Fig. 2E).

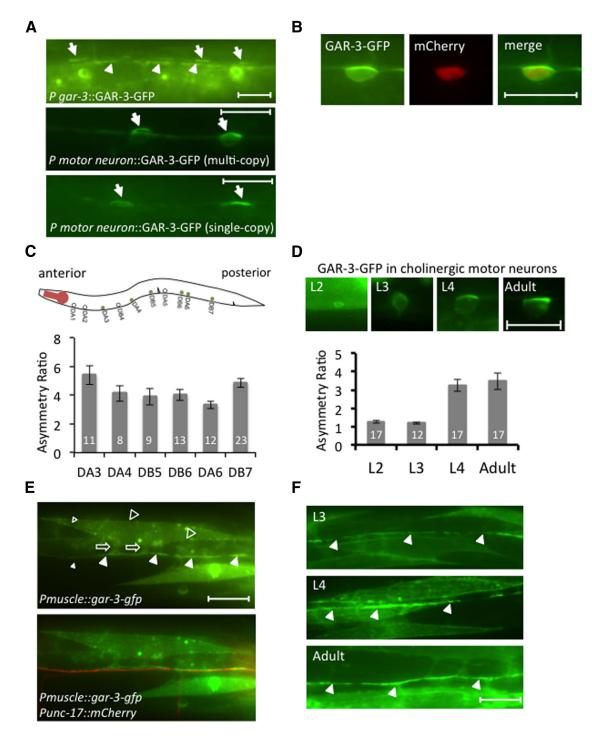


Figure 2. GAR-3/mAChR is asymmetrically localized on membranes of motor neurons and body wall muscles. *A*, Images showing the localization of GAR-3-GFP expressed under its endogenous promoter (*vjEx602*[*P gar-3(8.5)::gar-3b-gfp*)], as a multicopy integrant under a motor neuron promoter (*vjEx602*[*P unc-129::gar-3b-gfp*]), or as a single copy integrant under a motor neuron promoter (*vjIs50*[*P unc-129::gar-3b-gfp*]). *B*, Representative images showing the localization of GAR-3-GFP (*vjIs50*) and mCherry (*vjIs14*[*P unc-129::mCherry*]) in DA/DB cell bodies. *C*, Top, Diagram and quantification of the DA/DB neurons examined along the A/P axis of the worm. Cells examined in this study are filled-in circles. *D*, Representative images and quantification of GAR-3-GFP localization in motor neurons (*vjIs50*) from L2 to adult stages. *E*, Representative images showing the colocalization of GAR-3-GFP in body wall muscles (*vjEx748*[*P myo-3::gar-3b-gfp*]) with mCherry expressed in cholinergic neurons (*nuIs321*[*P unc-17::mCherry*]). *F*, Representative images showing GAR-3-GFP in muscles (*vjEx748*) of L3 to adult stages. *A*, *E*, *F*, Filled arrows indicate motor neuron cell bodies, filled arrowheads indicate postsynaptic membranes of in the dorsal nerve cord, open arrowheads indicate muscle cell membranes, and open arrows indicate muscle arms. *C*, *D*, Numbers in bars represent the number of animals examined and error bars indicate ± SEM. Scale bars, 10 μm.

To quantify the asymmetric distribution of GAR-3-GFP in neuron cell bodies, we compared fluorescence on the dorsal and ventral sides of cell bodies to establish an asymmetry ratio (D/V). The asymmetry ratio in DA/DB class neurons expressing multicopy GAR-3-gfp was 4.9 \pm 0.3 (expressed as mean \pm SEM hence-

forth, n=53). In neurons expressing single-copy *GAR-3-gfp* in DA/DB class, the ratio was 6.2 ± 0.9 (n=17). GAR-3-GFP was asymmetrically distributed in each of the DA/DB class neurons examined along the anterior–posterior length of the ventral nerve cord (Fig. 2*C*). GAR-3-GFP fluorescence was not detected in ax-

ons in single-copy transgenic animals (0 of 15 axons) and was rarely detected in axons of multicopy animals (3 of 23 axons). GAR-3-GFP was also not detected in dendrites of single-copy transgenic animals, and was occasionally detected in proximal dendrites of multicopy animals (data not shown). When axonal or dendritic GAR-3-GFP fluorescence was detected, it adopted a diffuse pattern of localization (Fig. 2*A*, *B*, and data not shown). Based on these results, we conclude that GAR-3 is highly enriched in cholinergic motor neuron cell bodies, where it asymmetrically distributes to the dorsal surfaces, but is likely to be very low or absent from neuronal processes.

GAR-3-GFP localization was not asymmetric early in development (asymmetry ratio 1.19 \pm 0.6 in the L3 stage), but it became asymmetric starting at the L4 stage (3.25 \pm 0.34) and maintained asymmetry in adults (Fig. 2*D*). GAR-3-GFP on muscle membranes was visible in young animals (L3 stage) and appeared enhanced in older animals (Fig. 2*F*). These observations suggest that the asymmetric localization of GAR-3 in motor neuron cell bodies and muscles may be under developmental control.

The N-terminal extracellular domain of GAR-3 is necessary and sufficient for asymmetric localization

To investigate mechanisms underlying GAR-3 asymmetry, we first examined the localization of chimeras in which domains of GAR-3 were replaced with homologous domains from another mAChR family member, GAR-1. Unlike GAR-3, GAR-1-GFP fusion proteins adopted a uniform (symmetric) localization pattern in motor neuron cell bodies (asymmetry ratio of 1.20 \pm 0.04; Fig. 3 *A*, *B*). The third intracellular domain of GPCRs, the i3 loop, specifies G-protein coupling and has been shown to be important for GPCR trafficking (Nathanson, 2008). GAR-3 chimeras containing the intracellular loop of GAR-1 [GAR-3(1-i3)] were similarly asymmetric as GAR-3-GFP (4.92 \pm 0.95), suggesting that the i3 loop is not necessary for GAR-3 asymmetry. Chimeras in which the N-terminal domain of GAR-3 was replaced with the GAR-1 N-terminus (GAR-1-3 chimeras) adopted a uniform distribution (asymmetry ratio of 1.11 \pm 0.05), indistinguishable from that of GAR-1 (Fig. 3A, B). On the other hand, chimeras in which the N-terminal region of GAR-1 was replaced with that of GAR-3 (GAR-3-1 chimeras) adopted a highly asymmetric distribution (4.49 \pm 0.70). Together these results show that the N-terminus of GAR-3 is both necessary and sufficient for the asymmetric localization of GAR-3-GFP in neuronal cell bodies.

The reduced asymmetrical distribution of the GAR-1-3 chimera was not a secondary effect of a reduction in expression or stability of the chimera: First, in three independently generated transgenic lines, total fluorescence on cell-body surfaces was similar in GAR-3-GFP and GAR-1-3-GFP-expressing animals (Fig. 3C). Second, GAR-1-3-GFP fluorescence appeared to be higher in axons (16 of 21 axons examined) compared with GAR-3-GFP. Thus, the uniform distribution of the GAR-1-3-GFP chimera in cell bodies cannot be explained by a reduction in its expression. The GAR-1-3 chimera also adopted a uniform distribution in body wall muscles, with no enrichment in muscle arms or in postsynaptic membranes of the ventral nerve cord, suggesting that the N-terminus of GAR-3 is required for asymmetric localization in both neurons and muscles (Fig. 3D).

To further map the sequence determinants for GAR-3 localization, we analyzed the cell body distribution of GAR-3-GFP variants with different N-terminal deletions. GAR-3-GFP variants lacking amino acids 13–23 [GAR-3(Δ NT1)] or amino acids 24–39 [GAR-3(Δ NT2)] adopted a uniform pattern of localization

tion (asymmetry ratios 1.10 ± 0.03 and 1.13 ± 0.04 , respectively), whereas deletion of amino acids 54-66 [GAR-3(Δ NT3)] remained asymmetrically distributed (2.94 \pm 0.39; Fig. 3*A*,*B*). Thus, residues between positions 13 and 39 are necessary for the asymmetrical distribution of GAR-3 in neuron cell bodies.

Glycosylation contributes to asymmetric GAR-3 localization

Two predicted *N*-glycosylation sites, N28 and N33, are present within the N-terminal extracellular region required for GAR-3 asymmetry (consensus: N,X,S/T). Because glycosylation regulates the post-translational processing of mammalian muscarinic receptors and influences receptor trafficking (Liles and Nathanson, 1986), we examined the effects of altering these sites on GAR-3 localization. GAR-3-GFP variants carrying the N33Q substitution were significantly less asymmetric than wild-type GAR-3-GFP (2.59 \pm 0.26). Variants containing the N28Q substitution, however, remained asymmetric (3.41 \pm 0.38; Fig. 3*A*,*B*). GAR-3-GFP variants with mutations in both asparagines (N28Q and N33Q) adopted similar reductions in asymmetry as GAR-3-GFP(N33Q; 2.57 \pm 0.31). Thus, glycosylation at N33 contributes to GAR-3 asymmetry, but additional N-terminal determinants must also specify GAR-3 localization.

To further explore the contribution of glycosylation on GAR-3 localization, we knocked down enzymes involved in protein glycosylation by RNA interference (RNAi; Table 1). We examined 16 genes predicted to be involved in various aspects of biosynthesis, transfer, and modification of glycosyl groups in the ER. We found that knock down of one gene, ostd-1, the δ subunit of an oligosaccharyltransferase (OST) complex protein, decreased GAR-3-GFP asymmetry (Table 1). OST proteins participate in protein N-glycosylation by catalyzing the transfer of branched oligosaccharides to Asn residues on nascent polypeptides in the ER (Chavan and Lennarz, 2006).

We next examined other cellular pathways that might regulate GAR-3 asymmetric localization. GPCR receptor activation often leads to a reduction in their cell-surface expression through receptor internalization or redistribution (Ritter and Hall, 2009; Reiner and Nathanson, 2012). Acutely increasing ACh levels with aldicarb, which causes the accumulation of ACh at cholinergic synapses (Mahoney et al., 2006), did not significantly change the GAR-3-GFP asymmetry ratio compared with nontreated controls (Fig. 4A). Similarly, neither genetically decreasing ACh biosynthesis (either acutely in temperature-sensitive mutants for *cha-1*/choline acetyltransferase or chronically in *unc-17*/VAChT mutants) nor ACh release (in *unc-2*/VGCC) significantly altered the asymmetry ratios (Fig. 4A,B). Given these results, it seems likely that GAR-3 asymmetry is not regulated by activity-dependent mechanisms.

The planar cell polarity (PCP) and Wnt signaling pathways regulate cell polarity by establishing asymmetry on specific regions of cell membranes (Hardin and King, 2008; Nance and Zallen, 2011). To explore whether these pathways regulate GAR-3 distribution, we examined the effects of RNAi-mediated knockdown of PCP or Wnt pathway genes on GAR-3-GFP asymmetry (Table 1). Although, RNAi of most PCP and Wnt pathway genes tested did not generate a significant change in GAR-3-GFP distribution (Table 1), we observed that RNAi of *mom-2*, one of five secreted Wnt ligands (Green et al., 2008), resulted in a significant reduction of GAR-3-GFP asymmetry (Table 1). Together these results suggest that extracellular determinants, including glycosylation, and possibly specific Wnt signaling pathways, appear to play a role in the establishment or maintenance of asymmetric GAR-3 in neuronal cell bodies.

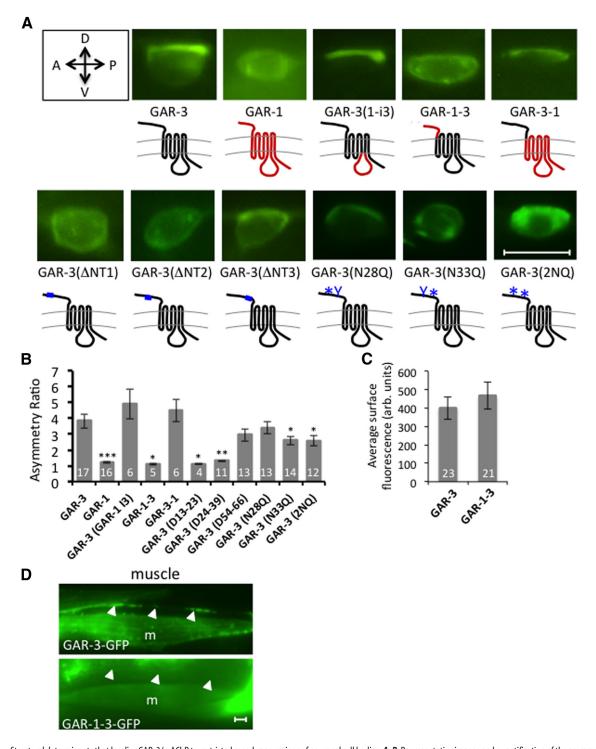


Figure 3. Structural determinants that localize GAR-3/mAChR to restricted membrane regions of neuronal cell bodies. *A, B,* Representative images and quantification of the asymmetry ratio of GAR-3-GFP and GAR-3-GFP variants in DA/DB motor neuron cell bodies. *C,* Average fluorescence on outer edge of DA/DB cell bodies of GAR-3-GFP and GAR-1-3 chimeras (three independent lines each). *D,* Representative images of GAR-3-GFP (*vjEx748*) and GAR-1-3-GFP (*vjEx742*) in body wall muscles. "m" denotes muscle and arrowhead denotes the ventral nerve cord. *A,* Asterisks denote mutated glycosylation sites. *B, C,* Numbers in bars represent the number of animals examined and error bars are ± SEM. *B,* **p* < 0.005, ****p* < 0.0005; Student's *t* tests. Scale bar, 5 μm.

Localization of GAR-3 to the basal lamina of the pseudocoelom

We reasoned that interactions between the N-terminus of GAR-3 and adjacent tissues might be involved in GAR-3 localization. 3-D reconstructions from serial electron microscopy show that the cell bodies of all ventral cord motor neurons are in direct contact with skin (ventral hypodermal ridge), neuronal processes, and the basal lamina of the pseudocoelom (David Hall,

Albert Einstein College of Medicine, New York, personal communication; White et al., 1976). Reconstructions of the DA3 and DA7 motor neurons shows that the skin contacts a large area of the cell bodies along their ventral and medial surfaces (Fig. 5*A*; White et al., 1976). Several neuronal processes running longitudinally along the ventral nerve cord, including those of DVB, AVF, and PVQ, contact the DA7 cell body along its lateral surface. Finally, the basal lamina is a small region of the ECM that sepa-

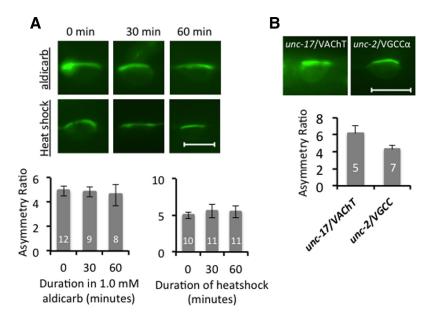


Figure 4. GAR-3 asymmetry is not affected by changes in synaptic ACh levels. **A**, Representative images (top) and quantification (bottom) showing the localization of GAR-3-GFP in DA/DB motor neurons (vjls50) following 0, 30, and 60 min of aldicarb treatment or heat shock. **B**, Representative images (top) and quantification (bottom) showing the localization of GAR-3-GFP in DA/DB motor neurons (vjls50) in the indicated mutants. Scale bars, 5 μ m. Numbers in bars denote the number of animals examined, and error bars are \pm SEM.

rates the pseudocoelomic cavity from the dorsal surface of motor neuron cell bodies.

Because GAR-3-GFP localized to narrow, dorsal regions of DA/DB cell bodies, it seemed most likely that GAR-3 contacts either one of the dorsally positioned neuronal processes or the basal lamina of the pseudocoelom. We found that GAR-3-GFP fluorescence did not colocalize with the processes of mCherrylabeled DVB, AVF, AVB, or PVQ neurons (Fig. 5A). However, GAR-3-GFP consistently colocalized with rhodamine-conjugated wheat germ agglutinin (WGA) that had been injected into the pseudocoelom. WGA binds to sugar moieties in the ECM, and WGA-labeled membranes appeared as a fluorescent band that lined the pseudocoelom (Fig. 5A). Together, these results suggest that GAR-3 is associated with the basal lamina. Consistent with this, when animals were dissected to expose the DA/DB cell bodies, acute (1 min) treatment with trypsin or collagenase caused GAR-3-GFP to become more uniformly distributed on the surface of the cell body (Fig. 5B). In the case of trypsin, the asymmetry ratio was significantly reduced from 6.48 \pm 1.05 in controls to 2.05 \pm 0.44 in trypsin treated animals (p < 0.05). Together these results suggest that GAR-3 is not in contact with neuronal processes in the ventral cord, but rather contacts the basal lamina separating the pseudocoelomic cavity and motor neuron cell body.

Asymmetric localization of GAR-3 is necessary for neuromuscular function

To address whether the asymmetric localization of GAR-3 in motor neuron cell bodies is important for synaptic function, we examined the effects of mislocalized GAR-3 variants on aldicarb responses and SPHK-1/sphingosine kinase recruitment. The GAR-1-3 chimera is composed of the N-terminal extracellular domain of GAR-1 fused to GAR-3 at the first TM domain (Fig. 3A). Structure–function studies of mammalian mAChRs have found that the N-terminal extracellular domain does not impact agonist binding, receptor dimerization, or G-protein coupling

(Romero-Fernandez et al., 2011; Leach et al., 2012). Expression of the GAR-1-3 chimera in motor neurons failed to rescue the aldicarb resistance of *gar-3* mutants. In contrast, GAR-3 expression in *gar-3* mutants caused aldicarb hypersensitivity (Fig. 6A).

Arecoline is a muscarinic agonist that causes gar-3-dependent aldicarb hypersensitivity (Fig. 6A; Lackner et al., 1999; Chan et al., 2012). Arecoline sensitivity was restored to gar-3 mutants by cholinergic expression of wild-type GAR-3 (Fig. 6A). We found that animals expressing GAR-1-3 were as responsive to arecoline treatment as GAR-3-expressing animals since the shifts in aldicarb hypersensitivity following arecoline treatment were similar (Fig. 6A). We conclude that the GAR-1-3 chimeras retain normal agonist responses but they fail to restore aldicarb responsiveness to gar-3 mutants because they are not asymmetrically localized.

We next examined the ability of the mislocalized GAR-1-3 chimera to recruit SPHK-1/sphingosine kinase to presynaptic terminals. Previous studies have shown that *gar-3* signaling in cholinergic motor

neurons increases the synaptic abundance of SPHK-1-GFP fusion proteins. SPHK-1-GFP adopts a punctate pattern of fluorescence in axons, and each puncta represents SPHK-1 at a release site (Chan et al., 2012). Cholinergic GAR-3 expression in *gar-3* mutants increased SPHK-1-GFP punctal fluorescence by 50% compared with controls (Fig. 6*B*), without altering the number of SPHK-1-GFP puncta (2.57 \pm 0.08 puncta per 10 μ m in control vs 2.46 \pm 0.07 in *gar-3* expressed, p > 0.05). In contrast, cholinergic expression of the mislocalized GAR-1-3 chimera in *gar-3* mutants did not significantly increase SPHK-1-GFP abundance (Fig. 6*B*). Thus, asymmetrically localized GAR-3 is necessary for both the behavioral and synaptic functions mediated by neuronal GAR-3 signaling.

Axonal calcium influx is necessary for GAR-3-mediated SPHK-1 recruitment

If the activation of GAR-3 on cholinergic neuron cell bodies leads to changes in presynaptic properties, we predict that disrupting signal propagation along axons would block the effects of GAR-3 activation on presynaptic function. In C. elegans, the voltageinsensitive cation leak channel, NALCN (encoded by nca-1 and nca-2), is required for calcium propagation along axons from cell bodies to nerve terminals (Yeh et al., 2008). unc-80 encodes a conserved NALCN-interacting protein required for the function of the NALCN channel. *unc-80* is expressed in ventral cord motor neurons, and unc-80 mutants have defects in calcium propagation along motor axons (Yeh et al., 2008). In mutants lacking unc-80, steady-state SPHK-1-GFP punctal fluorescence was slightly reduced. Activating GAR-3 by acute arecoline treatment results in gar-3-dependent increases in SPHK-1-GFP punctal fluorescence (Fig. 7A; Chan et al., 2012). The arecoline-mediated increase in SPHK-1-GFP punctal fluorescence was abolished in unc-80 mutants (Fig. 7A). These results show that recruitment of SPHK-1 to synapses by muscarinic signaling requires calcium propagation along the axon.

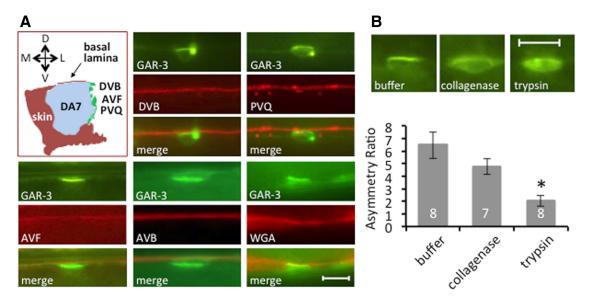


Figure 5. GAR-3 localizes near the extracellular matrix of the basal lamina. *A*, Diagram (top left) of the DA7 motor neuron cell body and cells that it contacts (constructed from White et al., 1976). Red is the hypodermis (skin), green are neuronal processes, and red line is basal lamina. M, Medial; L, lateral; D, dorsal; V, ventral. Representative images of GAR-3-GFP in DA/DB motor neurons (*vjls50*) colocalized with mCherry expressed in DVB (*vjEx746[P unc-47::mCherry]*), PVQ (*vjEx741[P sra-6::mCherry]*), AVF (*vjEx603[P dfk-2::mCherry]*), or AVB (*vjEx474[P glr-1::mCherry]*) and WGA injected into the pseudocoelomic cavity. *B*, Representative images and quantification of asymmetry of GAR-3-GFP in DA/DB motor neurons in dissected animals following treatment of control (M9), collagenase (1 mm), and trypsin (0.05%) for 1 min. **p* < 0.05, Student's *t* tests. Numbers in bars denote the number of animals examined. Scale bar, 5 μm.

Humoral ACh can activate GAR-3

The localization of GAR-3 to the basal lamina, which lacks cholinergic synapses, raises the possibility that GAR-3 may be activated by nonsynaptic ACh. To explore this idea, we determined whether ACh released from a source distant from motor neuron cell bodies could activate GAR-3. To do this, we drove ACh release from a subset of cholinergic neurons that do not directly contact the ventral nerve cord using a truncated *unc-17* promoter fragment [*Punc-17*(Δ)]. *Punc-17*(Δ) expresses specifically in head neurons (Rand, 1989; Tsalik and Hobert, 2003). We first confirmed that $Punc-17(\Delta)$ expression does not overlap with gar-3 expression in the ventral nerve cord by examining fluorescence patterns of animals coexpressing Punc-17(Δ)-mCherry reporters and GAR-3-gfp. We found that mCherry fluorescence was abundant in neurons of the head and in processes in the lateral cords, but was not observed in the ventral nerve cord (Fig. 7*B*).

To drive release of endogenously produced ACh in these neurons, we overexpressed unc-17, the C. elegans vesicular acetylcholine transporter (VAChT). VAChTs are synaptic vesicleassociated transmembrane proteins and overexpression of VAChT in hippocampal neurons increases ACh release by increasing the quantal content of vesicles (Song et al., 1997; Nagy and Aubert, 2012). We found that expressing unc-17/VAChT under the *Punc-17*(Δ) promoter caused aldicarb hypersensitivity (Fig. 7C), implying that the release of ACh from head/lateral neurons by unc-17/VAChT can increase synaptic transmission at NMJs. Similarly, unc-17/VAChT expression under the Punc- $17(\Delta)$ promoter significantly increased SPHK-1-GFP abundance at synapses (Fig. 7D). Both the increases in aldicarb sensitivity and SPHK-1 abundance caused by unc-17/VAChT expression were completely blocked by gar-3 mutations (Fig. 7C,D). Together, these results indicate that activation of GAR-3 by endogenous ACh released from distant neurons can positively regulate transmitter release form presynaptic motor terminals.

Discussion

In this study, we show that GAR-3, a *C. elegans* homolog of the M1/M3/M5 muscarinic receptor family, is preferentially localized on motor neuron cell body membranes that contact the basal lamina of the pseudocoelom. This localization is dependent on the GAR-3 N-terminal extracellular domain and is critical for *gar-3*-mediated regulation of presynaptic function. These results indicate that specialized membrane localization of mAChRs in neurons may be an evolutionarily conserved feature of neuromodulation by muscarinic signaling.

The function of asymmetric GAR-3 in motor neuron cell bodies

Examination of the subcellular localization of GAR-3-GFP in motor neurons showed several striking features. First, GAR-3 was preferentially concentrated on cell bodies and not detectable on dendrites or axons using multicopy overexpression and single-copy insertion of GAR-3-GFP. Second, GAR-3 is likely to be on the surface of the cell, because GAR-3-GFP-expressing animals responded robustly to exogenously applied arecoline, and digestion of the ECM caused rapid redistribution of GAR-3-GFP. Third, GAR-3-GFP adopted an asymmetric localization pattern along the dorsal region of neuronal cell bodies, and mislocalized GAR-3 variants failed to restore proper aldicarb responsiveness and presynaptic SPHK-1 recruitment to gar-3 mutants. Finally, in unc-80 mutants, which have disrupted calcium signal propagation from cell bodies to axons (Yeh et al., 2008), GAR-3 activation failed to regulate SPHK-1 presynaptic abundance.

One model to explain these results is illustrated in Figure 7*E*. ACh in the pseudocoelom acts on GAR-3 on motor neuron cell bodies, which can propagate a signal to axon terminals. This propagation is likely to be dependent on calcium currents mediated by UNC-80/NCA-1/NCA-2. Subsequently, activation of presynaptic voltage gated calcium channels promotes neurotransmitter release in part by regulating the levels of SPHK-1 at release sites. SPHK-1 is an enzyme that converts sphingosine to

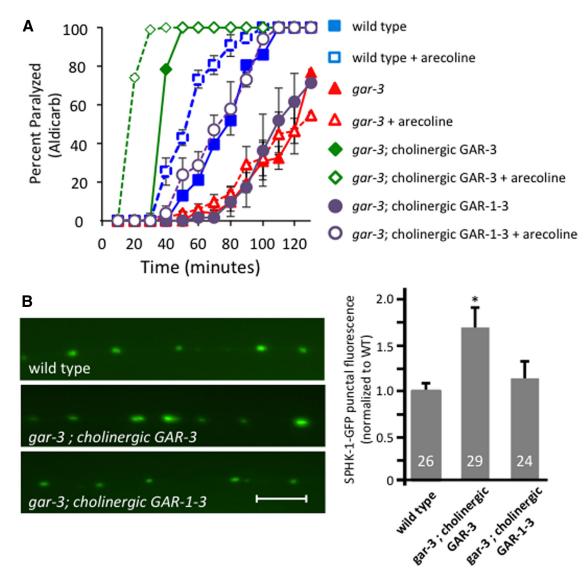


Figure 6. Behavioral and synaptic defects caused by mislocalized GAR-3/mAChR variants. *A*, Rates of worm paralysis of wild-type, *gar-3*, and *gar-3* animals expressing the cholinergic GAR-3 (*vjEx635[P unc-17::gar-3b]*) or cholinergic GAR-1-3 (*vjEx744[P unc-17::GAR-1-3b]*) transgenes upon exposure to the acetylcholine esterase inhibitor aldicarb (1.0 mm) following a 2 h pretreatment with control M9 (solid lines) or arecoline (15 mm in M9; dotted lines). *B*, Representative images and quantification of SPHK-1-GFP punctal fluorescence at synapses of DA/DB motor neurons of *gar-3* mutants expressing the indicated transgenes. Numbers in bars denote the number of animals examined, error bars are ± SEM. Scale bars, 10 μ m. *p < 0.05, Student's t tests.

sphingosine-1-phosphate, and local generation of sphingosine-1-phosphate at presynaptic terminals is proposed to promote neurotransmitter release (Brailoiu et al., 2002; Chan and Sieburth, 2012; Chan et al., 2012) Genetic and biochemical studies have identified several signaling components that are likely to act downstream of GAR-3 to regulate synaptic SPHK-1 including the heterotrimeric G-protein EGL-30/G α q, the EGL-30 effector UNC-73/Trio, the calcium/calmodulin binding protein CALM-1/CIB, and the voltage gated calcium channel UNC-2 (Chan and Sieburth, 2012; Chan et al., 2012). It remains to be determined where within the motor neurons these components act to transduce the signals from GAR-3 to the synapse. It is also possible that low levels of GAR-3 are found at synapses, below fluorescence detection, and contribute to changes in neurotransmission. Our results also do not rule out the possibility that NCA-1/NCA-2 may act in parallel to GAR-3 signaling to regulate presynaptic release. Interestingly, the NALCN channel is activated by the M3 mAChR in pancreatic β cells, and activation of NALCN appears to be independent of heterotrimeric G-proteins but dependent on the nonreceptor tyrosine kinase Src in β cells and in pyramidal cells of the hippocampus (Lu et al., 2009; Swayne et al., 2009).

Mechanisms of asymmetry

We found that the extracellular N-terminus of GAR-3 was necessary and sufficient for the fivefold enrichment of GAR-3-GFP in dorsal neuronal cell bodies and was necessary for membrane localization in muscles. We identified a putative N-linked glycosylation site (N33) within the N-terminal that was partially required for GAR-3-GFP localization. Although the N-terminus of GAR-3 does not share significant overall homology with other *C. elegans* or mammalian mAChR family members, human M3 and M4 mAChRs have been shown to be glycosylated *in vivo* (Nathanson, 2008; Romero-Fernandez et al., 2011). In COS cells, M3 mAChR is glycosylated on four Asn residues in its N-terminus, and mutation of these residues

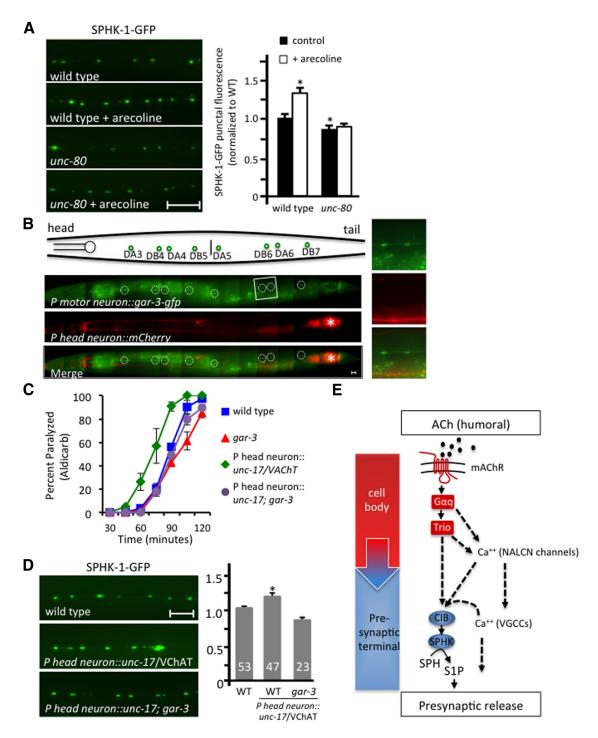


Figure 7. Endogenous ACh released form head neurons causes aldicarb hypersensitivity and recruits SPHK-1 to synapses. *A,* Representative images and quantification of SPHK-1-GFP punctal fluorescence in DA/DB motor neuron axons in the indicated strains treated with either control (M9) or arecoline (15 mm in M9) 2 h before analysis. *B,* Top, Schematic of DA/DB neuron positions as viewed from the ventral side of the animal. Bottom, Images of animals expressing *P motorneuron::GAR-3-gfp (vjls50)* or *P head neuron::mCherry (vjEx754)*. Right, Close-up images of the boxed area. Asterisks denote nonspecific gut fluorescence. *C,* Aldicarb responsiveness of the indicated strains. Head neuron *unc-17*/VAChT is the vjEx755 transgene. *D,* Representative images and quantification of SPHK-1-GFP puncta in the indicated genotypes. *E,* Model of the GAR-3 signaling pathway in motor neurons, in which GAR-3 is proposed to be activated by humoral ACh to modulate presynaptic release. Dotted lines denote indirect or unconfirmed interactions. Red is indicative of cell body localization and blue of synaptic localization within cells in the model. CIB, Calcium and integrin binding protein; SPHK, sphingosine kinase; SPH, sphingosine; S1P, sphingosine-1-phosphate; VGCCs, voltage gated calcium channels. Scale bars, 10 μm. Error bars indicate ± SEM. *p < 0.05, Student's *t* tests.

disrupts transport of the receptor to the plasma membrane (Romero-Fernandez et al., 2011).

We identified OSTD-1/OST- δ as a candidate for positively regulating GAR-3 asymmetry. OST- δ is a component of the OST complex that catalyzes the initial step in the biosynthesis of

N-linked glycoproteins in the lumen of the rough endoplasmic reticulum (Chavan and Lennarz, 2006). OSTD-1/OST- δ is expressed ubiquitously in *C. elegans*, including in motor neurons, and knockdown of this gene leads to locomotion defects (www. wormbase.org; Simmer et al., 2003). More studies will help de-

termine whether OSTD-1 directly participates in the localization of GAR-3 and whether other subunits of this complex also contribute to GAR-3 localization.

Our finding that the third cytoplasmic loop (the i3 loop) of GAR-3 was not necessary for the asymmetric localization of GAR-3 contrasts with the mechanisms by which M2 and M3 mAChRs are sorted in polarized epithelial cells. In Madin–Darby canine kidney cells, M2 and M3 mAChRs are delivered to the apical and basolateral membranes, respectively, using short sorting signals located in their i3 loops (Nadler et al., 2001; Iverson et al., 2005). It is thought that the i3 loop interacts with cytosolic factors that direct the polarized trafficking of vesicles as they go through the secretory pathway (Nathanson, 2008; Reiner and Nathanson, 2012). Thus, we speculate that GAR-3 localization is not regulated at the level of vesicle trafficking but rather by an interaction between the N-terminal extracellular domain and the ECM once GAR-3 is delivered to the cell surface.

Based on our colocalization studies, we believe that the N-terminus of GAR-3 interacts with a component of the basal lamina. NID-1/Nidogen and CLE-1/type XVIII collagen are components of the basal lamina surrounding cell bodies of motor neurons, and both proteins are proposed to regulate synaptic development (Kramer, 1994; Ackley et al., 2003). However, these proteins are enriched in lateral regions of the basal lamina that are not in contact with motor neuron cell bodies (Kramer, 1994; Ackley et al., 2003), and mutations in the genes encoding these proteins did not alter GAR-3-GFP localization (D. Sieburth, unpublished observations), making it unlikely that they are involved in GAR-3 localization. A member of the adhesion GPCR family, GPR56, binds collagen III at its N-terminal extracellular domain, and this interaction is critical for GPR56 activation and proper cell surface expression (Jin et al., 2007; Singer et al., 2013). Interestingly GAR-3 residues 38-48 show 82% similarity with residues in the GPR56 collagen-binding site. However, the GPR56 collagen-binding site did not confer localization to GAR-3 lacking its N-terminus (D. Sieburth, unpublished observations). Identifying components of the basal lamina that regulate GAR-3 localization will help elucidate mechanisms underlying the establishment and maintenance of GAR-3 in cell bodies.

Nonsynaptic ACh activates extrasynaptic GAR-3

Cholinergic transmission in the brain can occur by wired transmission, mediated by ACh released into the synaptic cleft, or by volume transmission by ambient ACh (Sarter et al., 2009; Agnati et al., 2010; Trueta and De-Miguel, 2012). Cholinergic signaling through volume transmission is supported by observations that there are fewer cholinergic fibers than cells expressing AChRs. Also, cholinergic neurons make few synaptic specializations with target cells and M3 receptors in cortical and hippocampal neurons are largely extrasynaptic (Descarries and Mechawar, 2000). Our results argue that GAR-3 is activated by extrasynaptic ACh released from distant cells, supporting a role for volume transmission in mediating motor function. First, GAR-3 localized to extrasynaptic regions of motor neuron cell bodies that are devoid of cholinergic synaptic contacts, as shown by electron microscopy (White et al., 1976). Second, colocalization studies showed that GAR-3 is exposed to the basal lamina, an ECM that separates motor neuron cell bodies from the pseudocoelom. Third, driving ACh release from cholinergic neurons in the head and lateral ganglia (using a truncated unc-17 promoter) was sufficient for GAR-3-dependent effects on NMJ protein composition and function. Because motor neurons receive direct synaptic input from command interneurons, but not cholinergic neurons of the head or lateral cords, the ACh released from these cells is likely to activate GAR-3 via an indirect mechanism.

What is the cellular source of ACh that activates GAR-3 on motor neurons? The pseudocoelom is a body cavity of the worm that supplies gases, nutrients, and other molecules to tissues throughout the animal, and indirect evidence supports the idea that ACh may be present at low levels in the pseudocoelom. First, a putative ACh gated ion channel, *cup-4*, functions in scavenger cells that are in direct contact with the pseudocoelom (Patton et al., 2005). Second, the choline transporter, *cho-1*, which mediates the re-uptake of the breakdown product of ACh, may be expressed in noncholinergic cell types including the intestine, which is exposed to the pseudocoelom (Mullen et al., 2007; Duerr et al., 2008). Finally, numerous sensory and interneurons are in contact with the pseudocoelom (Cheung et al., 2004; Kao et al., 2007), raising the possibility that certain cholinergic neurons in the head and tail may also release ACh into the pseudocoelom. gar-3 mutants have weak aldicarb resistance phenotypes and no detectable defect in SPHK-1 recruitment (Chan et al., 2012), suggesting that under normal conditions, ACh levels may be low. Interestingly, the binding affinities of agonists to GAR-3 and to M1 mAChRs (Park et al., 2006), which mediate volume transmission in the murine hippocampus (Yamasaki et al., 2010), are comparable. Perhaps the highly concentrated localization of GAR-3 at the basal lamina maximizes response to low ACh levels in the pseudocoelom. Interestingly, humoral transmission by other neuromodulators is proposed to control motor neuron activity in C. elegans: dopamine receptors controlling locomotion are not postsynaptic to any dopaminergic neuron terminal (Chase et al., 2004), and serotonin receptors mediating learned feeding responses to familiar foods are found far from the primary serotonin release sites (Song et al., 2013). Humoral transmission may provide a means to mediate modulatory aspects of neuronal networks on slower timescales (Sarter et al., 2009; Agnati et al., 2010; Fuxe et al., 2010). Indeed, our data support a modulatory function of GAR-3 in the regulation of neurotransmission because the functional defects of gar-3 mutants are more obvious under activity-dependent states (e.g., arecoline treatment, unc-17/VChAT overexpression). Identifying the cellular source of ACh that activates GAR-3 signaling in motor neurons will help to clarify the role that muscarinic signaling plays in regulating behavior.

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