Neuron-Specific Regulation of Associative Learning and Memory by MAGI-1 in C. elegans

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Abstract

Background: Identifying the molecular mechanisms and neural circuits that control learning and memory are major challenges in neuroscience. Mammalian MAGI/S-SCAM is a multi-PDZ domain synaptic scaffolding protein that interacts with a number of postsynaptic signaling proteins and is thereby thought to regulate synaptic plasticity [1,2,3].

Principal Findings: While investigating the behavioral defects of C. elegans nematodes carrying a mutation in the single MAGI ortholog magi-1, we have identified specific neurons that require MAGI-1 function for different aspects of associative learning and memory. Various sensory stimuli and a food deprivation signal are associated in RI interneurons during learning, while additional expression of MAGI-1 in glutamatergic AVA, AVD and possibly AVE interneurons is required for efficient memory consolidation, i.e. the ability to retain the conditioned changes in behavior over time. During associative learning, MAGI-1 in RIA neurons controls in a cell non-autonomous fashion the dynamic remodeling of AVA, AVD and AVE synapses containing the ionotropic glutamate receptor (iGluR) GLR-1 [4]. During memory consolidation, however, MAGI-1 controls GLR-1 clustering in AVA and AVD interneurons cell-autonomously and depends on the ability to interact with the β-catenin HMP-2.

Significance: Together, these results indicate that different aspects of associative learning and memory in C. elegans are likely carried out by distinct subsets of interneurons. The synaptic scaffolding protein MAGI-1 plays a critical role in these processes in part by regulating the clustering of iGluRs at synapses.

Introduction

Dynamic changes involving the formation of new synapses, morphological changes of dendrites, increased numbers of dendritic spines and the redistribution of synaptic proteins create the remarkable plasticity of the nervous system [5,6]. Activation of neurotransmitter receptors on the postsynaptic side of the synapse also triggers biochemical pathways that lead to changes in neuronal function. NMDA and AMPA subtypes of ionotropic glutamate receptors (iGluR) play crucial role in vertebrate synaptic plasticity, and are linked by membrane-associated guanylate kinases (MAGUKs) into large signaling complexes. However, the in vivo role of MAGUKs in LTP formation and learning is largely unknown. Mammalian MAGI/S-SCAM is a postsynaptic multi-PDZ domain-containing MAGUK with a unique inverted domain organization [2]. MAGI was reported to interact with a multitude of neuronal proteins including NMDA receptors, transmembrane AMPA receptor regulating proteins (TARPs), a number of other neuronal proteins, and β-catenin [1,7,8,9]. Therefore MAGI proteins are thought to play an important role in synaptic plasticity and memory formation.

The relatively simple nervous system of C. elegans composed of 302 neurons allows non-associative (adaptation and habituation) as well as associative learning between a variety of volatile or soluble chemoattractants, or cultivation temperature, and food. In order to investigate associative learning in C. elegans, several learning assays have been established, for example, salt chemotaxis learning [10], olfactory associative learning [11,12], and temperature learning [13,14]. Previous studies have also shown that regulators of learning and memory are conserved between mammals and C. elegans [11,12,15,16]. AMPA type of ionotropic glutamate receptors not only play a critical role in vertebrate synaptic plasticity, but glutamate neurotransmission has also been shown to be involved in habituation of the tap withdrawal response as well as in olfactory associative learning in C. elegans [12,16]. Therefore, the analysis of genes found in C. elegans can provide important insights into the mechanisms of learning and memory in vertebrates including humans.

Vertebrate genomes encode three MAGI isoforms, while a single MAGI-1 ortholog exists in C. elegans. Therefore, C. elegans is an ideal model to study the function of the MAGI-1 protein in associative learning and neural plasticity. In this study, we


Editor: Georges Chapouthier, L’Université Pierre et Marie Curie, France
Received March 16, 2009; Accepted May 25, 2009; Published June 24, 2009
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Funding: This research was supported by grants from the Swiss National Science Foundation to A.H and the Kanton of Zurich and from the National Institutes of Health, USA (CA095281) to S.vdH. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The authors have declared that no competing interests exist.

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generated a deletion mutant of magi-1 and investigated the learning and memory defect in the mutant worms. We found that loss of magi-1 function, regardless of the sensory input, impairs associative learning in C. elegans. Expression of the wild-type MAGI-1 protein in a single pair of interneurons can rescue the learning defects. In addition, magi-1 mutant worms show defects in memory consolidation, which requires the wild-type MAGI-1 protein in a distinct set of interneurons. We also found that MAGI-1 controls the distribution of ionotropic glutamate receptor, GLR-1 during learning and memory consolidation. We propose a molecular mechanism for MAGI-1 during neural plasticity, which modulatory function may be conserved in mammals.

Results

Loss of MAGI-1 function impairs associative learning independent of sensory input

In order to study MAGI-1 function in a simple animal model, we generated a knockout allele (zh66) of the single C. elegans MAGI ortholog magi-1 (Fig. S1b). The magi-1 locus encodes two mRNAs that are transcribed from two alternative promoters (Fig. S1a, b). The magi-1(zh66) deletion allele removes most of the region common to both isoforms. Total protein extracts from zh66 animals lacked both isoforms as determined by Western blot analysis (Fig. S1c). Thus, the zh66 deletion likely represents a magi-1 loss-of-function (lf) allele. magi-1(lf) mutants appear healthy, fertile and display no obvious morphological or locomotory defects (data not shown).

To investigate a potential role of MAGI-1 in C. elegans olfactory associative learning, we first tested the chemotaxis of magi-1(lf) animals towards different compounds. The chemotaxis of magi-1(lf) mutants to six different volatile attractants, one soluble attractant, and a repellent was comparable to the response of the wild-type N2 strain (Fig. S2a). Furthermore, both wild-type and magi-1(lf) mutants responded similarly to food starvation, indicating that magi-1(lf) mutants have no obvious sensory or motor defects (Fig. S2b). To test the role of MAGI-1 in associative learning, we used an established context-dependent starvation conditioning protocol [14,17]. In the olfactory learning assay, unconditioned wild-type and magi-1(lf) animals both exhibited strong chemotaxis towards diacetyl (DA) (Fig. 1a). However, after a one-hour starvation period in the presence of DA, wild-type animals displayed a strongly reduced attraction to DA, whereas starvation-conditioned magi-1(lf) mutants exhibited only a partial reduction in chemotaxis towards DA (Fig. 1a). We obtained similar results when using isovaleral alcohol as the attractant (data not shown). The magi-1(tm446) deletion that removes only the long isoform, on the other hand had no significant effect on olfactory learning (Fig. 1a). This result suggests that the two MAGI-1 isoforms act in a redundant manner during associative learning.

To exclude the possibility that the conditioning defect in magi-1(lf) mutants was due to changes in adaptation rather than association, we conditioned magi-1(lf) mutants and wild-type controls with DA in the presence of abundant food (Fig. 1b). In this experimental context, magi-1(lf) mutants were not defective in DA adaptation when compared to wild-type (Fig. 1b).

As a next step, we wondered if the observed associative learning defect was specific to volatile chemo-attractants or reflected a more general learning deficit. Indeed, magi-1(lf) mutants also showed strong defects in gustatory (salt chemotaxis) and thermotaxis conditioning assays. Starvation of wild-type animals in the presence of NaCl induced a strong avoidance of NaCl in subsequent chemotaxis assays (Fig. 1c). magi-1(lf) mutants, on the other hand, showed a greatly reduced avoidance of NaCl after starvation conditioning (Fig. 1c). In addition, magi-1(lf) animals failed to associate low (15°C) or high (25°C) temperatures with food starvation (Fig. 1d, e). We conclude that the loss of MAGI-1 function does not affect adaptation but impairs associative learning independently of the type of sensory input, as we found that MAGI-1 is required for the association of a food starvation signal with diverse environmental stimuli such as smell, taste or temperature.

MAGI-1 acts in RIA interneurons during associative learning

To study the MAGI-1 expression pattern, we generated a construct encoding the long isoform of magi-1 fused to GFP (Fig. S1b). This magi-1::gfp minigene construct rescued the behavioral defects of magi-1(lf) mutants (Fig. 1a, c). MAGI-1::GFP was expressed in several interneurons, including AVA, AVD, AVE, RIM, and RIA (Fig. 2a, b). These neurons are all known to express ionotropic glutamate receptors (6GluRs) [4]. Therefore, we performed tissue-specific rescue experiments to determine the cellular focus of action. We tested empty pPD95.67 vector, or expressed the magi-1::gfp coding region under the control of a 2.7 kb fragment of the glr-1 promoter, which is active in RIM, AVA, AVD and AVE, or the glr-3 or glr-6 promoters, which are exclusively active in RIA neurons (Fig. 3c). In the olfactory learning paradigm, Pgbr-3 orPgbr-6-driven magi-1::gfp rescued the learning defects of magi-1(lf) mutants, while no rescue was observed with Pgbr-1-driven magi-1::gfp (Fig. 3a,d). Expression of magi-1 in RIA neurons also rescued the learning defect in the salt chemotaxis learning paradigm (Fig. 3b). Therefore, expression of magi-1 in RIA was sufficient to restore associative learning to magi-1(lf) mutants in an immediate recall test. To further confirm that RIA plays a central role in associative learning, we ablated the RIA neuron using glr-3 promoter-driven activated ICE caspase (Fig. 4a, b). In order to visualize the efficient ablation of RIA we co-expressed glr-3 driven GFP together with ICE caspase. Expression of the activated caspase led to apoptotic cell death of RIA neurons in early larvae in 59.6% of the transgenic animals (Fig. 4a, b). Transgenic worms where ICE activity was insufficient to kill RIA were used as internal controls. Genetic ablation of RIA neurons caused associative learning defects both in the volatile attractant and salt chemotaxis learning paradigms (Fig. 4c, d). Together with previous findings [13], these results confirm the central role of RIA neurons in a neural circuit mediating associative learning.

Efficient memory consolidation requires MAGI-1 function in AVA and AVD interneurons

We next tested the memory consolidation, the capability of the animals to retain a conditioned behavior over time, by starvation conditioning the animals in the presence of DA and letting the animals recover for 30 minutes in the absence of DA before testing their chemotaxis [18]. In wild-type animals, the negative association of DA with starvation persisted during the 30-minute recovery period (Fig. 3c). Interestingly, although expression of magi-1 in RIA neurons using the glr-6 or glr-3 promoters rescued conditioning in magi-1 mutants (Fig. 3d), it was not sufficient to restore memory consolidation over the 30 minutes recovery phase (Fig. 3c). Since MAGI-1 is not exclusively expressed in RIA, we speculated that memory consolidation might require MAGI-1 expression in additional neurons such as AVA, AVD, AVE, or RIM. We therefore co-expressed MAGI-1 under the control of different neuron-specific promoters (Fig. 3c) together with the RIA-specific glr-6 promoter construct and tested the double transgenic animals for memory consolidation. Co-expression of
magi-1::GFP under the glr-6 promoter with either of the glr-1, unc-42 or nmr-1 promoters rescued the memory consolidation defects of magi-1(lf) mutants (Fig. 3d). Since no MAGI-1::GFP expression was detected in AVE and the unc-42 promoter is not active in AVE and RIM, furthermore, AVE is mainly presynaptic in the ventral nerve chord, magi-1 acts most likely predominantly in AVA and AVD during memory consolidation. However, our results cannot fully exclude that participation of additional neurons during memory consolidation are also required. AVA, AVD, and AVE interneurons express the GLR-1 glutamate receptor subunit [4], which has been shown to be essential for associative learning [16] as well as for the nose-touch response [19]. Similar to glr-1(lf) mutants, magi-1(lf) mutants showed a defect in their nose-touch response that could be rescued by expressing magi-1::gfp under the control of the glr-1 promoter (Fig. S3), suggesting that MAGI-1 is required together with GLR-1 in AVA and AVD interneurons for a proper nose touch response. AVA, AVD and AVE neurons project their processes along the ventral nerve chord (VNC). GLR-1 localizes in AVA, AVD, AVE and other VNC synapses in punctae corresponding to receptor clusters in post-synaptic regions [20]. Furthermore, MAGI-1::YFP was localized in the VNC adjacent to the presynaptic marker synaptobrevin (SNB-1), most likely in the postsynaptic region as well as in distinct dots flanking the synaptic cleft (Fig. 2d–f). Thus, efficient memory consolidation requires MAGI-1 function in glutamatergic AVA and AVD and possibly in AVE interneurons.

Figure 1. MAGI-1 is essential for associative learning. a, Context-dependent starvation conditioning of wild-type or mutant worms was assayed towards 0.1% DA volatile chemo-attractant without (naive) or with (conditioned) pre-incubation with DA. Chemotaxis index was calculated as CI = (worms at DA - worms at solvent)/total number of worms. b, Chemotaxis towards DA was tested in the presence or absence of food. c, Associative learning of magi-1(lf) and wild-type animals by NaCl starvation conditioning. d and e, magi-1(lf) animals were tested in a thermotaxis association paradigm by starvation conditioning the animals at 15°C or 25°C. All experiments were done in triplicate and repeated at least three times. Error bars indicate average±S.E.M. Significance between datasets was tested with two-tailed student’s t-test (** p<0.01, *** p<0.001). doi:10.1371/journal.pone.0006019.g001

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MAGI-1 regulates the remodeling of GLR-1 containing synapses during associative learning

Since MAGI-1 function is required in AVA and AVD neurons for memory consolidation, we asked whether GLR-1 containing AVA and AVD synapses in the VNC posterior to the vulva are remodeled during associative learning using a functional glr-1::gfp reporter [20]. In wild-type animals, synaptic GLR-1::GFP cluster size in the posterior VNC remained unchanged after starvation or exposure to DA in the presence of abundant food alone (Fig. 5a, c) (naïve: 1.705±0.158 μm, n = 79; starved: 1.73±0.115 μm, n = 83, p = 0.9062, two-tailed t-test compared to untreated dataset; DA with food: 1.76±0.145 μm, n = 173, p = 0.7836, two-tailed t-test compared to untreated dataset). However, GLR-1::GFP cluster size was reduced upon starvation conditioning with DA (1.22±0.0688 μm, n = 181, p = 0.000562, two-tailed t-test compared to untreated dataset; DA and starvation: 1.26±0.1 μm, n = 180, p = 0.000562, two-tailed t-test compared to untreated dataset) and stayed small during the 30 minute recovery phase (Fig. 5c) (1.24±0.0875 μm, n = 118, p = 0.000889, two-tailed t-test compared to untreated dataset). Thus, persistent synaptic remodeling in the VNC is induced in wild-type animals by olfactory associative conditioning but not by food starvation or olfactory adaptation alone.

We next tested if MAGI-1 regulates GLR-1 distribution in the VNC of wild-type and magi-1(0) mutants. In magi-1(0) mutants the size of the GLR-1::GFP clusters was already enlarged in naïve animals and did not decrease after starvation conditioning (Fig. 5d). We then analyzed the size of GLR-1::GFP clusters in conditioned magi-1(0) worms carrying the Pglr-1::magi-1, the Pglr-6::magi-1 or both transgenes together. Pglr-1::magi-1 alone reduced the size of GLR-1::GFP clusters to the size observed in naïve wild-type animals, but cluster size did not decrease further upon starvation conditioning (Fig. 5d). Expression of MAGI-1 in RIA using the glr-6 promoter not only restored associative learning (Fig. 5a, d) but also the reduction of GLR-1::GFP cluster size upon starvation conditioning (untreated: 2.14±0.1 μm n = 202; conditioned: 1.473±0.0596 μm, n = 202, p = 1.54*10^{-8}, two-tailed t-test compared to untreated dataset). However, GLR-1::GFP cluster size in naïve Pglr-6::magi-1 animals was still increased similar to non-transgenic magi-1(0) mutants (Fig. 5d) (magi-1(0): 2.328±0.19 μm n = 85; magi-1(0), Ex/Pglr-1::magi-1::gfp: 2.143±0.1 μm, n = 202, p = 0.349, two-tailed t-test) Furthermore, GLR-1::GFP clusters in Pglr-6::magi-1 animals reverted to nearly the size observed in unconditioned animals after 30 minutes of recovery, suggesting that expression of MAGI-1 in RIA alone was not sufficient to mediate a persisting size reduction of GLR-1 clusters at AVA and AVD synapses in the VNC (1.94±0.084 μm, n = 147, p = 5.18*10^{-6}, two-tailed t-test compared to conditioned dataset). Finally, all aspects of the conditioned changes in GLR-1 clustering were rescued when magi-1 was simultaneously expressed in RIA, AVA and AVD using the glr-1 and glr-6 promoters (Fig. 5d) (1.3±0.0682 μm, n = 170, p = 5.83*10^{-5}, two-tailed t-test compared to untreated dataset). Thus, MAGI-1 plays distinct functions in RIA, AVA and AVD neurons during learning and memory. Expression of MAGI-1 in RIA induces during associative learning the remodeling of GLR-1 containing synapses in AVA and AVD in a cell non-autonomous fashion. At the same time, MAGI-1 acts cell-autonomously in AVA and AVD to consolidate the conditioned changes in GLR-1 clustering during memory formation.

Efficient memory consolidation requires the HMP-2 β-catenin interaction domain of MAGI-1

Even though MAGI-1 regulates the dynamic changes in GLR-1 clustering at synapses, we did not detect a direct interaction between MAGI-1 and GLR-1, neither in vitro nor in vivo (data not shown). Since the vertebrate MAGI protein interacts with β-catenin [3], we tested if C. elegans MAGI-1 might bind to the β-catenin homolog HMP-2, the only one of the three C. elegans β-catenins that is predicted to contain a C-terminal PDZ binding motif (A.H. and A.S. personal observation). GST pull-down experiments revealed binding of HMP-2 to MAGI-1 (Fig. 6a). This interaction required the PDZ binding motif at the extreme C-terminus of HMP-2 as well as the fifth PDZ domain of MAGI-1 (Fig. 6a). To investigate the physiological significance of the HMP-2/MAGI-1 interaction, we generated a truncated MAGI-1 rescue construct lacking PDZ domain 5 [a deletion of amino acids 954 to 1092 in MAGI-1]. The MAGI-1ΔPDZ5::GFP protein localized similar to full-length MAGI-1::GFP (data not shown) and was able to rescue the associative learning defect (Fig. 6b), but neither the memory consolidation nor the nose touch-response defects of magi-1(0) mutants (Fig. 6c and data not shown). In contrast, a deletion mutant of MAGI-1 lacking PDZ1-3 was able to interact with...
HMP-2 β-catenin (Fig. 6a), but was not sufficient to rescue the associative learning defect of magi-1(lf) mutants (Fig. 6b). We conclude that MAGI-1 plays distinct molecular functions in RIA, AVA and AVD neurons, as memory consolidation and nose touch-response but not associative learning require the HMP-2 β-catenin-binding domain in MAGI-1.

**Discussion**

In summary, we demonstrate that RIA, AVA, AVD and likely AVE interneurons are responsible for different aspects of associative learning in *C. elegans* (Fig. 6d). While previous data suggested that RIA mediates the association of temperature with food presentation, we show that AVA and AVD are also involved in associative learning. This finding highlights the complexity of the neuronal network involved in associative learning in *C. elegans*.
feeding status [15], our results show that RIA plays a central role also in olfactory and gustatory associative learning. Thus, RIA may serve to integrate and associate multiple environmental inputs with food starvation. Upon associative conditioning, RIA probably relays a signal to other neurons including the command interneurons AVA, AVD, and likely AVE. Since, there is no direct synaptic connection between RIA and AVA, AVD or AVE interneurons, likely other neurons are also required to mediate signals from RIA to AVA, AVD and AVE. Our results suggest that the multi PDZ-domain protein MAGI-1 is required in RIA for the integration of those environmental inputs during associative learning. On the other hand, memory consolidation, the retention of the conditioned behavior over time, requires the GLR-1 expressing interneurons AVA, AVD and likely AVE, where MAGI-1 is necessary to induce persisting changes in the synaptic GLR-1 cluster size. During associative conditioning, the size but not the number of GLR-1 clusters decrease, which might reflect an increased density of glutamate receptors at post-synaptic membranes. Recently, the long isoform of the C.elegans MAGI-1 was demonstrated to influence mechanosensory habituation and GLR-1 receptor degradation through ubiquitination [21]. During habituation the number of GLR-1 positive synapses decrease in the MAGI-1 mutant worms carrying a deletion in the long isoform, suggesting a decrease in glutamate signaling. On the other hand overall number of synapses remains unaffected. In contrast to habituation, associative learning increase synaptic sensitivity and cause a dynamic remodeling of synapses. In our work we demonstrate that complete loss of both isoforms of MAGI-1 impairs mechanosensation through the command interneurons, demonstrated by the lack of response to nose touch in magi-1(0) mutant worms. Furthermore, loss of MAGI-1 has additional effect on neuronal plasticity besides the demonstrated role in non-associative learning [21]. In contrast to the changes observed in the number of GLR-1 punctae during repeated long-term mechano-stimulation, we found that associative learning regulates the average size but not the number of GLR-1 positive synapses. Computational modeling suggested that the glutamate concentration decreases rapidly with increasing distance from the site of release [22,23]. Hence, slight local shifts in receptor density can lead to large changes in the postsynaptic response. The so-called flexible matrix model describes rapid and continuous changes of the synaptic architecture driven by the actin cytoskeleton [24]. Upon associative learning, a signal from the RIA interneuron induces a transient GLR-1 cluster remodeling in AVA, AVD interneurons that could sensitize the postsynaptic densities. This process is, however, independent of MAGI-1 function in AVA and AVD. On the other hand, the persisting changes in GLR-1 cluster size that correlate with memory...
The changes in synapse structure and composition. Further analysis of the role of a MAGI-1/β-catenin complex might give insight into a mechanism of memory formation conserved between *C. elegans* and humans.

Materials and Methods

General methods and strains used

Standard methods were used for maintaining and manipulating *C. elegans* [25]. Experiments were conducted at 20°C otherwise indicated. The *C. elegans* Bristol strain, variety N2, was used as the wild-type reference strain in all experiments. Alleles and transgenes

- **Figure S1.** MAGI-1 regulates GLR-1 localization during associative learning and memory formation. Confocal images of the anterior half of the ventral nerve cord of a 4 days old adult **a**, Naive wild-type and **b**, magi-1(lf) worms carrying the glr-1::gfp transgene. **c**, Average size of GLR-1::GFP clusters in the posterior VNC upon starvation in the absence (starved, n = 174) or presence of 0.1% DA (conditioned, n = 181), conditioning followed by 30 minutes recovery in the absence of DA (conditioned+30 min recovery, n = 119), or conditioning in the presence of abundant food (conditioned+food, n = 84). **d**, Average size of GLR-1::GFP clusters in the posterior VNC upon starvation without (control) or with 0.1% DA (conditioned) or conditioning followed by 30 minutes recovery in the absence of DA (conditioned+recovery) in wild-type and magi-1(lf) animals carrying the indicated transgenes (wild type: control n = 80, conditioned n = 181; magi-1(lf): control n = 86, conditioned n = 55; magi-1(lf); Ex[glr-1::magi-1]: control n = 182, conditioned n = 165; magi-1(lf); Ex[glr-6::magi-1]: control n = 202, conditioned n = 288, conditioned+recovery n = 186; magi-1(lf); Ex[glr-1::magi-1, glr-6::magi-1]: control n = 233, conditioned n = 271, conditioned+recovery n = 253). Three independent experiments were quantified, and three independent transgenic lines were tested for each construct. Error bars indicate average±S.E.M. Significance between datasets as indicated was tested with two-tailed student’s t-test (n.s. p > 0.05, ** p < 0.01, *** p < 0.001). doi:10.1371/journal.pone.0006019.g005

consolidation require the presence of MAGI-1 in AVA and AVD. Interestingly, AVA and AVD are backward command interneurons, therefore an increase in synaptic sensitivity in these neurons could be the direct cause of the avoidance behavior upon associative learning. We also show that the same domain in MAGI-1 that is necessary for the interaction with the β-catenin HMP-2 is also required to retain the conditioned behavior over time, but dispensable for associative learning per se. Hence, MAGI-1 could serve as a scaffold and indirectly control glutamate receptor signaling in AVA, AVD and AVE neurons through interaction with the cadherin/catenin complex, for example by the consolidating rearrangement of the actin cytoskeleton and thereby...
used were magi-1 (tm446), nl25 [20], akEx211 [gph-3::gfp; glr-3::ICE; lin-13], lin-13[n765ts], magi-1(zh66), magi-1(lf); nl25, magi-1(zh66); ubEx9 [magi-1::gfp, sur-5::mDsRed], magi-1(zh66); ubEx2 [pglr-1::magi-1::gfp, sur-5::mDsRed], magi-1(zh66); ubEx3 [pglr-6::magi-1::gfp, sur-5::mDsRed], magi-1(zh66); ubEx12 [pglr-3::magi-1::gfp, sur-5::mDsRed], magi-1(zh66); ubEx10 [pglr-6::magi-1::gfp, pglr-1::magi-1::flag, sur-5::mDsRed], magi-1(zh66); ubEx13 [pglr-6::magi-1::gfp, punc-42::magi-1::gfp, sur-5::mDsRed], magi-1(zh66); ubEx4 [pglr-6::magi-1::gfp, sur-5::mDsRed], magi-1(zh66); ubEx14 [pglr-6::magi-1::gfp, pmw-1::magi-1::gfp, sur-5::mDsRed], magi-1(zh66); ubEx5 [pglr-6::magi-1::gfp, sur-5::mDsRed], magi-1(zh66); ubEx6 [pglr-6::magi-1::gfp, ubc-11::pglr-1::flag, sur-5::mDsRed], magi-1(zh66); ubEx17 [pglr-1::flag, sur-5::mDsRed], magi-1(zh66); ubEx19 [pglr-1::APD5::gfp, sur-5::mDsRed], magi-1(zh66). All experiments were done in triplicate and repeated at least in two independent experiments. At least two independent transgenic lines were tested for each construct. Error bars indicate average ± S.E.M. Datasets were compared to magi-1(lf) mutants using two-tailed student’s t-test (** p<0.001).

Isolation of magi-1(zh66)
To isolate a deletion mutation in magi-1, a library consisting of ~5×10⁴ mutagenized F1 animals was screened as described [27].
with some modifications. Wild-type worms were mutagenized with a combination of 30 μg/ml psoralen (TMP) and 2000 μJ/cm² UV, allowed to recover for 16 hours and five F₀ animals were plated on each of 900 NGM plates. Cultures were grown and one third of the animals were harvested for DNA isolation. To identify the plate containing the magi-1(ze66) deletion, DNA extracts from 12 cultures was pooled and each pool was tested by nested PCR with outer primers OAS-364 [5`TTCCCCGCGTTGAATTCGG-GACCCCAAGG TTG]³, OAS-422 [5`CAGGATGATGTCCCTCATCCTATAC]³, and inner primers OAS-423 [5`GGTCCCG-TCCGACTJAGTTCCGGAGTC]³, OAS-424 [5`CCATCGGAA-GAACCTGTTCCAG CTG]³. The remaining animals from the culture positive for the ze66 deletion were subjected to three rounds of sib-selection until a single homozygous magi-1(ze66) animal had been identified. Before further analysis, the magi-1(ze66) mutation was backcrossed five times against N2 animals.

Olfactory conditioning was assessed as described [17] with some modifications. Starvation conditioning was performed without food in the presence of 2 μl undiluted chemo-attractor spotted on a piece of filter paper and attached to the lid of the plate for 1 hour on 10 cm CTX plates (5 mM KH₂PO₄/K₂HPO₄ pH 6.0, 1 mM CaCl₂, 1 mM MgSO₄, 2% agar). Naive and conditioned worms were given a choice between a spot of 0.1% (vol/vol) DA or ethanol and sodium-azide. After one hour the animals were counted and a chemotaxis index was calculated as described [28].

Chemotaxis to water-soluble compounds was assessed as described with some modifications [29]. Pairs of opposite quadrants of four-quadrant Petri plates (Falcon X plate, Becton Dickinson Labware) were filled with buffered agar (2% agar, 5 mM KH₂PO₄/K₂HPO₄ pH 6.0, 1 mM CaCl₂ and 1 mM MgSO₄), either containing 25 mM NaCl or no salt. Adjacent quadrants were connected with a thin layer of molten agar. A population of well-fed, young adults was washed three times with CTX buffer (5 mM KH₂PO₄/K₂HPO₄ pH 6.0, 1 mM CaCl₂ and 1 mM MgSO₄) and 100–200 worms were placed at the intersection of the four quadrants. The distribution of the worms over the four quadrants was determined after 10 minutes. For NaCl conditioning, animals were exposed to 25 mM NaCl in CTX buffer for 4 hours.

Thermotaxis assay

A steep thermal gradient on a thin agar plate was established as described previously [13]. After build up of the gradient, plates were separated into “20°C”, “25°C” and “15°C” regions where “20°C” corresponded to the 19°C to 21°C zone, “25°C” to a warmer and “15°C” to a cooler region. For conditioning, animals were grown at the indicated temperature for 2 days and starved for 4 hrs before testing. The numbers of worms in each of the three temperature zones were determined after 10 minutes, and the thermotaxis was calculated as described [15].

**Microscopy**

GFp (or GFP variants)-tagged proteins were detected with a Zeiss Axiovert 200M LSM 5 Pascal confocal microscope. Animals were immobilized in sodium-azide and GLR-1::GFP was recorded posterior to the vulva, along the z-axis. Quantification of GLR-1::GFP cluster size was performed on the projected z-sections using Openlab 5.0 software package (Improvision).

**Molecular biology**

Promoter-gene fusions were generated by PCR fusion as described [31]. All fragments were amplified using proof reading polymerase (LA Taq, Takara) from C. elegans genomic DNA. Primer sequences are available on request. GST-fusion proteins were expressed in E. coli and purified on glutathione sepharose using standard protocols. After incubation with total worm protein extracts prepared by lysing worms in lysis buffer (50 mM Tris-HCl pH 7.4, 1% Triton-X 100, 10% glycerol, 150 mM NaCl, 1 mM NaF, 1 mM Na-orthovanadate, and protease inhibitor cocktail), bound MAGI-1::GFP proteins were detected on Western blots using anti-GFP antibodies (Roche).

**Supporting Information**

**Figure S1**

Found at: doi:10.1371/journal.pone.0006019.s001 (0.30 MB DOC)

**Figure S2**

Found at: doi:10.1371/journal.pone.0006019.s002 (0.13 MB DOC)

**Figure S3**

Found at: doi:10.1371/journal.pone.0006019.s003 (0.20 MB DOC)

**Acknowledgments**

We wish to thank members of our groups for critical discussion and comments on the manuscript. We are also grateful to the C. elegans genetics centre and Shohei Mitani for providing strains, to Andrew Fire for GFP vectors and Christopher Rongo for MAGI-1 antibodies.

**Author Contributions**

Conceived and designed the experiments: AS. Performed the experiments: AS AH. Analyzed the data: AS AH. Contributed reagents/materials/analysis tools: FH AVM SvdH. Wrote the paper: AS FH AVM SvdH AH.

**References**

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