

1 ***INX-18 and INX-19 play distinct roles in electrical synapses that modulate***  
2 ***aversive behavior in Caenorhabditis elegans***

3 **Short Title: Electrical synapses in behavior modulation**

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25 **Abstract**

26 In order to respond to changing environments and fluctuations in internal states, animals adjust  
27 their behavior through diverse neuromodulatory mechanisms. In this study we show that electrical  
28 synapses between the ASH primary quinine-detecting sensory neurons and the neighboring ASK  
29 neurons are required for modulating the aversive response to the bitter tastant quinine in *C.*  
30 *elegans*. Mutant worms that lack the electrical synapse proteins INX-18 and INX-19 become  
31 hypersensitive to dilute quinine. Cell-specific rescue experiments indicate that *inx-18* operates in  
32 ASK while *inx-19* is required in both ASK and ASH for proper quinine sensitivity. Imaging analyses  
33 find that INX-19 in ASK and ASH localizes to the same regions in the nerve ring, suggesting that  
34 both sides of ASK-ASH electrical synapses contain INX-19. While *inx-18* and *inx-19* mutant animals  
35 have a similar behavioral phenotype, several lines of evidence suggest the proteins encoded by  
36 these genes play different roles in modulating the aversive quinine response. First, INX-18 and INX-  
37 19 localize to different regions of the nerve ring, indicating that they are not present in the same  
38 synapses. Second, removing *inx-18* disrupts the distribution of INX-19, while removing *inx-19* does  
39 not alter INX-18 localization. Finally, by using a fluorescent cGMP reporter, we find that INX-18 and  
40 INX-19 have distinct roles in establishing cGMP levels in ASK and ASH. Together, these results  
41 demonstrate that electrical synapses containing INX-18 and INX-19 facilitate modulation of ASH  
42 nociceptive signaling. Our findings support the idea that a network of electrical synapses mediates  
43 cGMP exchange between neurons, enabling modulation of sensory responses and behavior.

44

45 **Author Summary**

46 Animals are constantly adjusting their behavior to respond to changes in the environment or to  
47 their internal state. This behavior modulation is achieved by altering the activity of neurons and  
48 circuits through a variety of neuroplasticity mechanisms. Chemical synapses are known to impact  
49 neuroplasticity in several different ways, but the diversity of mechanisms by which electrical  
50 synapses contribute is still being investigated. Electrical synapses are specialized sites of  
51 connection between neurons where ions and small signaling molecules can pass directly from one  
52 cell to the next. By passing small molecules through electrical synapses, neurons may be able to  
53 modify the activity of their neighbors. In this study we identify two genes that contribute to  
54 electrical synapses between two sensory neurons in *C. elegans*. We show that these electrical  
55 synapses are crucial for proper modulation of sensory responses, as without them animals are  
56 overly responsive to an aversive stimulus. In addition to pinpointing their sites of action, we  
57 present evidence that they may be contributing to neuromodulation by facilitating passage of the  
58 small molecule cGMP between neurons. Our work provides evidence for a role of electrical  
59 synapses in regulating animal behavior.

60 **Introduction**

61 A defining feature of animal behavior is its plasticity. Animals adapt their behavior in order to  
62 respond to environmental challenges and physiological changes. Such behavioral plasticity is  
63 essential for animal survival and is achieved by changing the activity of neurons and circuits in a  
64 variety of ways. One way is through neuromodulation, whereby diffusible signals such as  
65 neuropeptides, dopamine, and serotonin are used to tune brain activity in broad regions[1–3]. By  
66 contrast, neuronal activity can be altered locally by changing the strength of individual synapses[4,  
67 5]. In order to understand dynamic brain function, it is crucial to uncover mechanisms that drive  
68 neuroplasticity at various levels.

69        Electrical synapses (also known as gap junctions) are composed of membrane channels that  
70        join the cytoplasm of two cells[6]. They are found throughout vertebrate and invertebrate nervous  
71        systems[6–9] where they pass both electrical and chemical signals between connected cells[10].  
72        Electrical synapses have been primarily studied for their ability to synchronize electrical activity  
73        between pairs or groups of neurons[11–13], but can also pass small molecules such as calcium[14,  
74        15], cAMP[16–19], cGMP[17, 20], IP<sub>3</sub>[15, 21], and even small miRNA[22, 23]. Interestingly, while  
75        electrical synapses share similar function and protein topology in vertebrates and  
76        invertebrates[24], genes encoding electrical synapse components are evolutionarily unrelated[6,  
77        10]. As a result, electrical synapses in vertebrates are composed of connexins, while those in  
78        invertebrates are composed of innexins (INXs). The separate evolution of electrical synapses  
79        suggests the functional necessity of these channels, although their role in neural plasticity and brain  
80        function is not fully understood.

81        Recently, it was discovered that innexin networks play a crucial role in cGMP-dependent  
82        sensory modulation in *Caenorhabditis elegans*[25]. Krzyzanowski and colleagues found that cGMP  
83        functions within the sensory neuron ASH to dampen nociceptive sensitivity but is produced in  
84        neighboring neurons[26]. They further showed that cGMP-mediated dampening of ASH nociceptive  
85        sensitivity requires an innexin-based network[25]. These findings uncover a new strategy of  
86        network regulation that may contribute to the modulation of neural activity. ASH is the primary  
87        nociceptive neuron pair in *C. elegans* and responds with increased calcium levels to diverse  
88        aversive stimuli including hyperosmolarity, nose touch, heavy metals such as copper, volatile  
89        repellents such as octanol and alkaloids such as quinine[27–33]. ASH controls movement away  
90        from noxious stimuli through synapses on the forward and backward command interneurons.[34,  
91        35] Nociception in ASH is extensively modulated, and reactivity to aversive stimuli such as quinine  
92        is regulated by the presence of food and the satiety state of the worm[25, 36–40]. Notably, ASH

93 forms electrical synapses with multiple other sensory neurons and a few interneurons[41, 42],  
94 suggesting electrical synapses may be crucial in modulating its activity.

95 We investigated the impact of electrical synapses between ASH and its neighbor ASK on  
96 behavioral sensitivity to the bitter tastant quinine. ASK forms multiple electrical synapses with  
97 ASH[42] and expresses several innexins[8, 43, 44], making it a candidate for directly modifying ASH  
98 activity. Results of this study show that the electrical synapse proteins INX-18 and INX-19 function  
99 within ASK and ASH to allow for modulation of the quinine avoidance response. Through imaging,  
100 we found that INX-18 and INX-19 localize to known sites of electrical synapses. Our data further  
101 suggest that INX-19 plays a principle role in diffusion of cGMP from ASK to ASH. Our study  
102 identifies a direct connection between two sensory neurons that modulates neuronal activity and  
103 thus regulates behavior in *C. elegans*.

104 **Results**

105 ***Innixin-18 and innixin-19 are required for modulation of the quinine response***

106 A recent study suggests that a network of electrical synapses is involved in modulation of the  
107 quinine response[25], however the exact composition of those electrical synapses has not been  
108 determined. ASH is a multimodal nociceptive neuron that responds to quinine and forms direct  
109 electrical synaptic connections with the sensory neuron ASK[41, 42], which is also involved in  
110 quinine sensation[32]. To explore whether the electrical synapses between ASK and ASH play a role  
111 in modulating quinine sensitivity, we investigated the innexins INX-18 and INX-19 that are  
112 expressed in these two sensory neurons[8, 43, 44]. While INX-4 is also expressed in ASH, we did  
113 not include it in our analyses as it has already been explored in a previous study[25].

114 To determine whether INX-18 and/or INX-19 play a role in modulating the behavioral response  
115 to quinine, we assayed *inx-18(ok2454)*, *inx-19(ky634)* and *inx-19(tm1896)* mutant animals (figure

116 1A-B) for quinine sensitivity. We placed drops of quinine solution in front of freely crawling worms  
117 and recorded their responses as “responding” if they reverse or “non-responding” if they continue  
118 forward[32, 45]. We found that these mutant animals were hypersensitive to 1 mM quinine in the  
119 quinine drop test (figure 1C). As a negative control, we examined the response of mutant animals to  
120 M13 buffer. Both *inx-18(ok2454)* and *inx-19(tm1896)* animals responded to M13 buffer at similar  
121 levels to wild-type (N2) animals, *inx-19(ky634)* animals, however, were slightly more responsive  
122 than wild-type animals (figure S1A). This may be because this strain has mildly increased  
123 spontaneous reversal rates (see below). As a positive control, we tested the response of mutant  
124 animals to a high concentration of quinine (10 mM) that is strongly aversive to wild-type  
125 animals. We found that all strains respond similarly to presentation of 10 mM quinine (figure  
126 S1B). Together, these data show that *inx-18(ok2454)*, *inx-19(ky634)* and *inx-19(tm1896)* mutant  
127 animals have increased quinine avoidance, suggesting that ASH activity is increased in the absence  
128 of these electrical synapse components.

129 ***The inx-19(tm1896) allele alters quinine responses without affecting locomotion***

130 Two different *inx-19* alleles (*tm1896* and *ky634*) have been identified and implicated in sensory  
131 neuron function[43]. While mutant animals with either allele show increased response to 1 mM  
132 quinine (figure 1C), these two alleles have different impacts on locomotion. First, *inx-19(ky634)*  
133 mutant animals exhibited more reversals in response to M13 (figure S1A). Second, during  
134 locomotion, *inx-19(ky634)* animals spontaneously reversed more frequently in the absence of  
135 stimuli (figure S2A). Third, the average crawling velocity of *inx-19(ky634)* mutant animals was  
136 lower than that of wild-type animals (figure S2B). These data suggest that *inx-19(ky634)* animals  
137 have altered movement in addition to changes in quinine response. At a molecular level, *inx-*  
138 *19(ky634)* is a G→A single nucleotide polymorphism causing an E70K substitution within the first  
139 extracellular loop of INX-19, while *inx-19(tm1896)* is a 546 basepair deletion that removes the  
140 majority of the first intracellular loop and a portion of the second transmembrane domain of INX-

141 19 (figure 1A). Because the function of innexins requires their transmembrane domains, *tm1896* is  
142 likely to be a strong loss-of-function or null allele. By contrast, a substitution within the  
143 extracellular docking domain may have a more complicated effect on protein function. For this  
144 reason, *inx-19(tm1896)* animals were utilized for the remainder of the experiments.

145 ***Inx-19 is required in both ASK and ASH for modulation of the quinine response***

146 *Inx-19* is expressed in multiple tissues such as neurons and muscles. Even within the nervous  
147 system, *inx-19* is expressed in ASH as well as a number of other neurons, including ASK, which has  
148 been implicated in quinine sensation and its regulation[32, 43, 44]. To determine the site of action  
149 of INX-19, we performed a series of rescue experiments with *inx-19* cDNA fused to fluorophores in  
150 the *inx-19(tm1896)* background. We found that, under the control of the native *inx-19*  
151 promoter[43], expression of *inx-19* cDNA fully rescued quinine hypersensitivity in response to 1  
152 mM quinine (figure 2A). This demonstrates that *inx-19* cDNA is functional and the *inx-19* mutation  
153 is responsible for the quinine hypersensitivity phenotype. Interestingly, these worms also showed  
154 reduced response to 10 mM quinine, suggesting that INX-19 overexpression could cause over-  
155 correction of the quinine sensitivity defects (figure S3A).

156 We then expressed GFP or mCherry-tagged *inx-19* cDNA under the control of cell-selective  
157 promoters to determine in which neurons INX-19 acts to regulate quinine sensitivity. We found that  
158 expression of *inx-19* cDNA in either ASK or ASH (using *Psra-9*[46] and *Posm-10*[47, 48],  
159 respectively) did not significantly restore the quinine response to 1 mM quinine in *inx-19(tm1896)*  
160 animals. In contrast, simultaneous expression of *inx-19* in both ASK and ASH brought 1 mM quinine  
161 response rates back to wild-type levels (figure 2A). As controls, we tested the response of these  
162 animals to M13 buffer and 10 mM quinine and found no change in sensitivity (figure S3A, B). These  
163 data indicate that INX-19 is required in both ASK and ASH for appropriate modulation of quinine  
164 sensitivity.

165 ***Inx-18 is required in ASK for modulation of the quinine response***

166 *Inx-18* is expressed in a subset of neurons including ASK[8, 44]. However, unlike *inx-19*, *inx-18*  
167 is not expressed in ASH, indicating that its site of action resides outside of ASH. To determine  
168 whether the altered quinine response rate of *inx-18* mutant animals is due to the lack of INX-18  
169 function, we performed rescue experiments using *inx-18*. *Inx-18* does not have an obvious  
170 promoter, as several genes lie directly upstream of its genomic position. However, the second  
171 intron has been successfully used to drive its expression[49]. To test whether the *inx-18(ok2454)*  
172 mutation is responsible for the quinine hypersensitivity phenotype, we cloned *inx-18* gDNA, which  
173 included the intronic regions. Expression of *inx-18* gDNA was sufficient to restore responses to 1  
174 mM quinine in *inx-18(ok2454)* mutant animals to wild-type levels, indicating that loss of *inx-18* is  
175 the reason for quinine hypersensitivity (figure 2B). Next, we found that the site of action of *inx-18* is  
176 in ASK, as expression of *inx-18* cDNA fused to GFP using the *Psra-9* promoter rescued the quinine  
177 hypersensitivity phenotype (figure 2B). As controls, we tested the response of these animals to M13  
178 buffer and 10 mM quinine and found no change in sensitivity (figure S3C, D) These results show  
179 that *inx-18* and *inx-19* have distinct, but partially overlapping, sites of action. Combined, our data  
180 indicate that INX-19 must be present in both ASK and ASH, while INX-18 in ASK alone is sufficient  
181 to modulate the quinine response.

182 ***ASK INX-19 and ASH INX-19 localize to the same regions in neighboring axons.***

183 The *C. elegans* wiring diagram suggests that the ASK and ASH neurons form electrical synapses  
184 with one another in the nerve ring[41, 42], which raises the possibility that INX-18 and INX-19 are  
185 components of these electrical synapses. As our behavioral results show that *inx-19* functions in  
186 both ASK and ASH, we examined the subcellular localization of INX-19 in these two neurons using  
187 fluorescence microscopy. We drove expression of GFP-tagged INX-19 in ASK and mCherry-tagged  
188 INX-19 in ASH. These fluorophore-tagged INX-19 constructs are functional as they can restore  
189 quinine responses in *inx-19(tm1896)* mutant animals (figure 2A). If INX-19 is a component of

190 electrical synapses between ASK and ASH, we reasoned that INX-19 expressed in ASK would  
191 localize to the same regions of the nerve ring as INX-19 expressed in ASH. Our imaging data show  
192 that INX-19 forms punctate structures along the axons in the nerve ring when expressed in both  
193 cells. As expected, most ASK INX-19 and ASH INX-19 is localized to overlapping puncta, despite the  
194 fact that these innexin proteins are in two distinct neurons (figure 3A-D). Quantification of these  
195 images show that INX-19 expressed in ASK and ASH produces puncta that colocalize 67% of the  
196 time (figure 3H). These data indicate that INX-19 is present on both sides of the ASK-ASH electrical  
197 synapses.

198 ***INX-18 rarely colocalizes with INX-19***

199 Our behavioral results indicate that INX-18 functions within ASK to modulate the behavioral  
200 response to quinine. To investigate where INX-18 resides in ASK, and whether it is functioning in  
201 the same synapses as INX-19, we expressed GFP-tagged INX-18 and asked whether it colocalizes  
202 with INX-19 (figure 3E-G). We found that, like INX-19, GFP-tagged INX-18 forms puncta along the  
203 axons (figure 3F). However, INX-18 showed low levels of colocalization with mCherry-tagged INX-  
204 19 expressed in ASH (~4% colocalization, figure 3H), demonstrating that the vast majority of INX-  
205 18 is not in the same synapses as INX-19 in adult animals.

206 ***INX-19 localization in ASK requires both *inx-18* and *inx-19****

207 To determine the relationship between INX-18 and INX-19 localization, we investigated  
208 whether the expression patterns of INX-18 and INX-19 are influenced by one another. We  
209 expressed fluorescently-tagged *inx-18* and *inx-19* cDNA in ASK and ASH individually and examined  
210 their expression patterns in mutant backgrounds. We found that the number of INX-19 puncta in  
211 the ASK axon was significantly reduced in *inx-18* mutant animals (figure 4A). In addition,  
212 localization of INX-19 within ASK requires INX-19 in other neurons, as the number of ASK INX-19  
213 puncta was diminished in *inx-19(tm1896)* mutant animals (figure 4A). In no cases were the puncta

214 fully eliminated, indicating that only some electrical synapses are affected in each case. We did not  
215 observe significant differences in the number of INX-19 puncta in ASH in *inx-18(ok2454)* or *inx-*  
216 *19(tm1896)* animals, although the downward trend (figure 4B) suggests that INX-19 localization in  
217 ASH may need both *inx-18* and *inx-19*. In contrast, INX-18 localization does not appear to require  
218 INX-19, as the number of INX-18 puncta in the nerve ring remained unchanged in *inx-19(tm1896)*  
219 mutant animals (figure 4C). This indicates that the localization of INX-18 is independent of INX-19.  
220 Taken together, these data suggest that *inx-18* plays a role in INX-19 electrical synapse assembly  
221 and/or maintenance. Perhaps INX-18 is transiently present in the ASK-ASH synapses during  
222 development, but by adulthood INX-18 has been removed from these synapses. Indeed, a number of  
223 studies have shown that innexin expression can be developmentally controlled[8, 43, 44].

224 ***Inx-18 and inx-19 have largely overlapping functions***

225 To investigate the functional relationship between *inx-18* and *inx-19*, we assessed the  
226 behavioral responses of *inx-18; inx-19* double mutant animals. If these two genes act in parallel to  
227 regulate quinine sensitivity, the phenotype of the double mutant should be stronger than that of the  
228 single mutants. If, however, *inx-18* and *inx-19* are acting together in the same pathway, we would  
229 expect animals with mutations in both genes to have a phenotype of similar strength to the single  
230 mutant animals. The *inx-19(tm1896); inx-18(ok254)* double mutants responded at somewhat higher  
231 rates than both the *inx-18(ok2454)* and *inx-19(tm1896)* single mutants (figure 4D), but this  
232 difference was statistically insignificant. This suggests that the two genes function largely in the  
233 same pathway to modulate the quinine response. Together with the visualization data, these  
234 findings suggest that while INX-18 is localized to different electrical synapses than INX-19, its  
235 primary function is to set up or maintain INX-19 localization.

236 ***Three different possibilities for the function of the ASK-ASH electrical synapses in***  
237 ***quinine regulation***

238 In order to determine how *inx-18* and *inx-19* affect ASH activity, we considered three potential  
239 mechanisms: First, *inx-18* and *inx-19* mutations may alter the cell fate of ASK or ASH, leading to  
240 changes in the quinine sensing circuit. Second, the ASK-ASH electrical synapses could function to  
241 shunt calcium, depressing ASH activity by allowing calcium ions to flow out to ASK. In this case, we  
242 expect that removal of ASK-ASH electrical synapses would result in increased  $\text{Ca}^{2+}$  signals in ASH  
243 and decreased  $\text{Ca}^{2+}$  levels in ASK. Finally, the ASK-ASH electrical synapses could pass cGMP from  
244 ASK to ASH, thus down-regulating the quinine response in ASH. Indeed, it was previously  
245 demonstrated that expressing the guanylyl cyclase GCY-27 in ASK rescued the quinine  
246 hypersensitivity in *gcy-27(ok3653)* mutant animals[26], suggesting an important role of cGMP in  
247 ASK in modulating quinine responses. We tested these three possibilities by examining cell fate  
248 markers, the calcium indicator GCaMP6s, and the fluorescent cGMP reporter FlincG3 in ASK and  
249 ASH.

250 ***ASK and ASH cell fate and morphology are unchanged in *inx-19* and *inx-18* mutant  
251 animals***

252 Electrical synapse channels are known to regulate cell fate decisions during development[50,  
253 51], in particular, *inx-19* has been shown to regulate neural differentiation in *C. elegans*[43]. Thus, it  
254 is possible that *inx-19* or *inx-18* also impacts ASK and/or ASH cell fate or morphology. To test this  
255 possibility, we expressed mCherry in ASK (using the *sra-9* promoter) and mTagBFP2 in ASH (using  
256 the *osm-10* promoter, which also expresses weakly in ASI). We found that the cell fate of ASK and  
257 ASH remained the same in the *inx-18(ok2454)* and *inx-19(tm1896)* mutant animals, as the number  
258 of neurons that expressed these fluorescent markers and their positions were unaltered (figure 5).  
259 Furthermore, we showed that the morphology of ASK and ASH were identical between wild-type  
260 and the mutant animals. Specifically, both ASK and ASH have cell bodies near the terminal bulb of  
261 the pharynx, while dendrites extend to the nose tip and axons project into the nerve ring.  
262 Additionally, the cell bodies, dendrites, and axons remained clearly visible In wild-type, *inx-*

263 *19(tm1896)* and *inx-18(ok2454)* mutant animals (figure 5B). Together, these data indicate that  
264 there is no gross morphological or cell fate changes to either ASK or ASH upon removal of INX-18  
265 and INX-19.

266 ***ASK calcium responses remain unchanged upon removal of ASK-ASH electrical***  
267 ***synapses***

268 We examined the possibility that the ASK-ASH electrical synapses function to shunt calcium,  
269 thus decreasing behavioral responses to quinine. Previous studies have shown that the ASH  
270 neurons respond strongly to quinine with an increase in intracellular calcium[27]. While ASK is  
271 known to be a minor player in the quinine response[32], the calcium response of ASK neurons to  
272 quinine is unknown. In ASK, attractive stimuli typically result in a decrease in calcium levels, while  
273 the aversive stimulus SDS results in a calcium increase[52]. Thus, it is possible that the aversive  
274 stimulus quinine also directly triggers a calcium increase in ASK. Alternatively, ASK may receive  
275 calcium ions from the primary quinine-sensing neuron ASH via the ASK-ASH electrical synapses. If  
276 the ASK-ASH electrical synapses pass calcium from ASH to ASK, this shunting effect would decrease  
277 ASH calcium levels in response to quinine as some of the calcium ions in ASH would flow to ASK in  
278 wild-type worms. In contrast, in animals lacking the ASK-ASH electrical synapses, we would expect  
279 increased calcium levels in ASH as the flow to ASK would be blocked. If ASK receives calcium from  
280 ASH, we would expect any quinine-induced calcium signal in ASK to decrease in mutant animals  
281 lacking the ASK-ASH electrical synapses.

282 We expressed GCaMP6s in ASK and ASH to visualize calcium dynamics in those cells in response  
283 to quinine presentation. Because both ASK and ASH are involved in blue-light avoidance  
284 behavior[53], the GCaMP6s experiments were carried out in a *lite-1(ce314)* background to eliminate  
285 blue-light induced changes of GCaMP6s fluorescence in ASK and ASH. Our results showed that  
286 GCaMP6 fluorescence in ASK and ASH increased after switching from buffer to quinine, indicating

287 increased  $\text{Ca}^{2+}$  levels in response to quinine (figure 6A-B, blue traces). However,  $\text{Ca}^{2+}$  signals in ASH  
288 were much more robust than those in ASK, consistent with the role of ASH as the primary quinine-  
289 sensing neuron[32].

290 To examine the impact of electrical synapses on  $\text{Ca}^{2+}$  dynamics, we monitored ASK and ASH  
291 GCaMP6s fluorescence in mutant *inx-18(ok2454)* and *inx-19(tm1896)* animals. We found that the  
292 increase in ASK GCaMP6s fluorescence remained the same between wild-type and mutant worms  
293 (figure 6B, 6D, 6F), suggesting that the ASK-ASH electrical synapses are not a main conduit for the  
294 ASK  $\text{Ca}^{2+}$  signal. When we imaged GCaMP6s fluorescence in ASH, we found the increase in ASH  
295 GCaMP6s fluorescence were enhanced in *inx-18 (ok2454)* and *inx-19(tm1896)* animals (figure 6A,  
296 6C, 6E). These results are consistent with the behavioral quinine hypersensitivity observed in these  
297 mutant worms. Together, these data show that ASK  $\text{Ca}^{2+}$  signals do not rely on the ASK-ASH  
298 electrical synapses, indicating that  $\text{Ca}^{2+}$  shunting to ASK is not the primary mechanism of quinine  
299 response regulation.

300 ***cGMP levels in ASK and ASH are influenced by ASK-ASH electrical synapses***

301 cGMP is required within ASH for down regulation of the quinine response[26]. However, ASH is  
302 not known to express any guanylyl cyclases, which produce cGMP. Recently, two studies suggested  
303 that guanylyl cyclase expression in other neurons plays a key role in modulating the quinine  
304 response[25, 26]. These findings prompted us to examine whether ASH acquires cGMP through the  
305 ASK-ASH electrical synapses. Indeed, ASK expresses the guanylyl cyclases ODR-1 and GCY-27[54],  
306 both of which are known to modify the quinine response[25, 26]. If ASK supplies ASH with cGMP  
307 through the ASK-ASH electrical synapses, we would expect to observe diminished levels of cGMP in  
308 ASH with a compensatory increase within ASK in *inx-18(ok2454)* and *inx-19(tm1896)* mutant  
309 animals.

310 To visualize levels of cGMP within ASK and ASH, we utilized the *C. elegans* codon-optimized  
311 version of FlincG3, which contains the cGMP binding domains of protein kinase G1 $\alpha$  fused to  
312 cpEGFP[55, 56]. Binding of cGMP increases FlincG3 fluorescence. We co-expressed FlincG3 and the  
313 red fluorescent protein mScarlet under control of the same promoters in ASK and ASH in the *lite-*  
314 *1(ce314)* background (figure 7A). After crossing the transgenes into *inx-18(ok2454)* and *inx-*  
315 *19(tm1896)*, we imaged FlincG3 fluorescence in ASK and ASH. FlincG3 fluorescence was compared  
316 to mScarlet fluorescence to account for variations in expression levels. We found that ASH FlincG3  
317 fluorescence was decreased in both *inx-18(ok2454)* and *inx-19(tm1896)* mutant animals (figure 7B),  
318 suggesting a reduction of the basal cGMP levels in ASH. These data are consistent with the  
319 behavioral hyper-responsiveness of *inx-18* and *inx-19* mutant worms to dilute quinine, as decreased  
320 cGMP levels could lead to increased ASH calcium levels in response to quinine[25, 26]. In ASK,  
321 FlincG3 fluorescence was increased in *inx-19(tm1896)* mutant animals but was unchanged in *inx-*  
322 *18(ok2454)* animals (figure 7C), suggesting that INX-19-based electrical synapses are primarily  
323 responsible for supplying ASH with cGMP from ASK. Together, our data suggest that INX-18 and  
324 INX-19 are major components of the ASK-ASH electrical synapses that modulate behavioral  
325 sensitivity to quinine, and that they do so by affecting transport of cGMP into ASH.

326 **Discussion**

327 We showed that electrical synapses between the *C. elegans* sensory neurons ASK and ASH play  
328 an active role in modifying nociceptive behavior via the passage of cGMP between cells. We found  
329 that the innexins INX-18 and INX-19 are required within ASK and ASH for proper modulation of the  
330 quinine response, as mutant animals lacking these innexins are hyperresponsive to quinine. These  
331 innexins form electrical synapses between ASK and ASH, in which INX-19 is a major component,  
332 though INX-18 is important for correct localization of INX-19 synapses in ASK. Our study supports  
333 a model in which ASK-ASH electrical synapses facilitate the passage of cGMP from ASK to ASH.  
334 Within ASH, cGMP downregulates calcium signals in response to quinine stimulation, likely by

335 binding to and activating the cGMP-dependent protein kinase EGL-4[26], ultimately leading to a  
336 reduction neural activity and thus aversive behavior (figure 8).

337 Electrical synapses can be made of different combinations of innexin subunits. Homotypic  
338 channels contain hemichannels that are composed of the same innexins, while heterotypic channels  
339 are made up of hemichannels that are composed of different innexins. The channel composition  
340 determines permeability, as heterotypic channels are thought to produce rectified electrical  
341 synapses: those that preferentially pass ions and small molecules in one direction rather than  
342 equally in both[57–59]. Our data suggest that INX-19 is a major component of the ASK-ASH  
343 electrical synapses. One possibility is that INX-19 forms homotypic channels. However, some INX-  
344 19 synapses do contain INX-18, suggesting that at least some are heterotypic. Though the number  
345 of electrical synapses containing both INX-18 and INX-19 is quite small, it is possible that levels of  
346 INX-18 within such synapses are generally low, making their visualization difficult. INX-18 could  
347 also make electrical synapses with other innexins in ASH. Nonetheless, our results suggest that the  
348 main function of INX-18 is carried out through its regulation of INX-19, as the *inx-18* and *inx-19*  
349 mutants do not show additive responses to quinine.

350 The structural makeup of the ASK-ASH electrical synapses has functional implications for ASH  
351 modulation. The composition of electrical synapses is key in determining their permeability, and  
352 heterotypic composition is a major cause of rectification[57, 59–61]. If the ASK-ASH electrical  
353 synapses are heterotypic (*i.e.*, consist of both INX-18 and INX-19 hemichannels) and rectified, this  
354 could explain why ASK cGMP levels, but not calcium levels, are affected by *inx-18* and *inx-19*  
355 mutations. Rectified channels bias the direction of movement of ions and molecules, making it more  
356 likely for signals to travel in one direction. If small molecule signals could easily pass from ASK to  
357 ASH but not in the reverse direction, cGMP may be more likely to travel from ASK to ASH than  $\text{Ca}^{2+}$   
358 would be from ASH to ASK. This mechanism could explain why our data suggest movement of cGMP

359 but not  $\text{Ca}^{2+}$ . Additionally, the permeability of electrical synapses is dependent on the subunits that  
360 make up the channels[17, 62]. While the permeability of most innixin-based channels is unknown,  
361 it is possible that the ASK-ASH electrical synapses are more permeable to cGMP than  $\text{Ca}^{2+}$ ,  
362 particularly given the timescales upon which each operate. Electrical synapses have long been  
363 considered low-pass filters, preferentially passing signals that change over longer time periods as  
364 opposed to quick oscillations[63, 64]. Regardless of the molecular reason, the selectivity of  
365 electrical synapses to either particular molecules or directions means that they can be sophisticated  
366 players within neural circuits. Changes in innixin composition during development or in mature  
367 circuits could dramatically impact how the neurons are regulated through the electrical synaptic  
368 network.

369 Electrical synapses are not static structures; they are regulated developmentally as well as in  
370 mature circuits[44, 63, 65–67]. Our data suggest that innexins can impact the localization of other  
371 innexins even if they are not a permanent part of the same synapses. INX-18 plays a crucial role in  
372 the localization of INX-19. Thus, its main impact on modulating the quinine response may be in  
373 supporting the function of INX-19. While INX-18 is required for proper localization of INX-19, an  
374 *inx-18(ok2454)* mutation does not eliminate INX-19 synapses completely. This may explain why the *inx-*  
375 *18(ok2454)* mutation does not have an impact on cGMP levels in ASK, as some signaling could still  
376 occur through the remaining INX-19-based electrical synapses even in the absence of INX-18.

377 ASH activity is modulated by cGMP, and yet ASH is not known to express any guanylyl cyclases,  
378 which produce cGMP[54, 68, 69]. This suggests that other neurons may regulate its activity. Such  
379 modulation occurs in the context of a larger sensory neuron network that simultaneously assesses  
380 many different sensory inputs, any of which could be affecting baseline levels of cGMP within  
381 sensory neurons. Thus, by being sensitive to changes in cGMP levels, ASH is able to receive  
382 modulatory information from many neurons simultaneously. ASH receives cGMP from its

383 immediate neighbor ASK as well as other neurons[25], suggesting that cGMP levels within ASH (and  
384 thus nociceptive sensitivity) are under the control of a number of external signals. If this is the case,  
385 cGMP could be a general signal of the state of the worm, integrating multiple signals to indicate  
386 whether it is in a favorable or unfavorable circumstance[70–74]. Our data support the notion that  
387 electrical synapses regulate function in a sensory neuron network by modulating the passage of  
388 small molecules into neurons such as ASH. In this way, multiple sensory inputs such as availability  
389 of food or sexual partners, presence of pathogens or other environmental conditions could alter  
390 various different behaviors at once.

391 **Figures**

392 **Figure 1: Mutations in *inx-19* and *inx-18* result in hypersensitivity to quinine.**

393 **A,B)** Diagram of *inx-19* and *inx-18* alleles used. Innexin genes code for proteins that consist of 4  
394 transmembrane helices with intracellular N and C tails. *Inx-19(ky634)* is a SNP resulting in an E>K  
395 substitution within the first extracellular loop, while *inx-19(tm1896)* is an in-frame deletion of  
396 546bp that removes most of the intracellular loop and a portion of the third transmembrane  
397 domain. *Inx-18(ok2454)* is a ~1800bp deletion that removes the second-fourth transmembrane  
398 domains and a portion of the C-terminus. **C)** Quinine Drop Test with 1 mM quinine. *Inx-19(ky634)*,  
399 *inx-19(tm1896)*, and *inx-18(ok2454)* mutant animals are hypersensitive to 1 mM quinine,  
400 responding a greater percentage of the time. N2 (wild-type)=18%, n=510; *inx-19(ky634)*=65%,  
401 n=120, p<0.0001; *inx-19(tm1896)*=44%, n=390, p<0.0001; *inx-18(ok2454)*=44%, n=350, p<0.0001.  
402 All groups were compared with a Chi-square test (p<0.0001,  $\alpha=0.05$ ), and post-hoc Fisher's Exact  
403 tests with Bonferroni's correction ( $\alpha=0.0167$ ) were computed to compare each group to the  
404 control.

405 **Figure 2: Expression of *inx-19* and *inx-18* in ASK and ASH restores wild-type quinine  
406 sensitivity.**

407 **A)** Expression of *inx-19* isoform A cDNA under the native promoter in *inx-19(tm1896)* animals  
408 rescued quinine sensitivity to N2 (wild-type) levels. Expression in ASK (*Psra-9*, which expresses  
409 solely in ASK[46]) or ASH (*Posm-10*, which also expresses in the tail neurons PHA and PHB as well  
410 as weakly in ASI[47, 48]) alone did not significantly rescue the behavior, while simultaneous  
411 expression did. N2=15%, n=220; *inx-19(tm1896)*=46%, n=210; *inx-19;Pinx-19::inx-19cDNA*=18%,  
412 n=100, p=0.62 vs N2, p<0.0001 vs *inx-19*; *inx-19;Psra-9::inx-19cDNA*=32%, n=100, p=0.0009 vs N2,  
413 p=0.02 vs *inx-19*; *inx-19;Posm-10::inx-19cDNA*=37%, n=110, p<0.0001 vs N2, p=0.13 vs *inx-19*; *inx-*  
414 *19;Psra-9::inx-19cDNA*; *Posm-10::inx-19cDNA* =22%, n=110, p=0.16 vs N2, p<0.0001 vs *inx-19*. All  
415 groups were compared with a Chi-square test (p<0.0001,  $\alpha$ =0.05), and post-hoc Fisher's Exact tests  
416 with Bonferroni's correction ( $\alpha$ =0.006) were computed to compare each group to N2 and *inx-*  
417 *19(tm1896)*. All rescues were performed with C-terminal mCherry- or GFP-tagged INX-19 and  
418 expression was verified visually before behavioral experiments. **B)** Expression of *inx-18* gDNA in  
419 *inx-18(ok2454)* animals rescued the quinine hypersensitivity phenotype, as did expression of *inx-18*  
420 cDNA in ASK (*Psra-9*). N2=13%, n=120; *inx-18(ok2454)*=48%, n=120; *inx-18;inx-18gDNA*=12%,  
421 n=100, p=0.84 vs N2, p<0.0001 vs *inx-18*; *inx-18;Psra-9::inx-18cDNA*=14%, n=120, p>0.99 vs N2,  
422 p<0.0001 vs *inx-18*. All groups were compared with a Chi-square test (p<0.0001,  $\alpha$ =0.05), and post-  
423 hoc Fisher's Exact tests with Bonferroni's correction ( $\alpha$ =0.013) were computed to compare each  
424 group to N2 and *inx-18(ok2454)*. All rescues except for gDNA were performed with C-terminal GFP-  
425 tagged INX-18 and expression was verified visually before behavioral experiments.

426 **Figure 3: INX-19 and INX-18 colocalize in the nerve ring when expressed in ASK and  
427 ASH**

428 **A)** Diagram of the *C. elegans* head in a dorsal view. Dashed box indicates the location of imaging  
429 of ASK and ASH axons in the nerve ring. **B-D)** INX-19 expressed in both ASK (where it is tagged with

430 GFP) **(B)** and ASH (where it is tagged with mCherry) **(C)** forms multiple puncta that colocalize along  
431 the ASK-ASH axons. Points of colocalization are indicated with white arrowheads. ASK and ASH are  
432 additionally expressing cytosolic mTagBFP2, seen in the axons that traverse the image, highlighted  
433 in **D. E-G**) INX-19 tagged with mCherry expressed in ASH **(E)** colocalizes in the nerve ring with GFP-  
434 tagged INX-18 expressed in ASK **(F)**. A white arrowhead indicates a point of colocalization.  
435 Cytosolic BFP fills the ASK-ASH axons, highlighted in **G. H**) Quantification of colocalization. In  
436 worms expressing INX-19 in ASK and ASH, 67% of nerve ring puncta colocalize (n=144 puncta in 14  
437 animals). In worms expressing INX-18 in ASK and INX-19 in ASH, ~4% of nerve ring puncta  
438 colocalize (n=81 puncta in 10 animals). Each dot represents an individual worm, and error bars are  
439  $\pm$ SEM.

440 **Figure 4: inx-18 and inx-19 play distinct roles in ASK-ASH electrical synapse localization  
441 and function**

442 **A)** *inx-19* cDNA was expressed using *Psra-9* and fluorescent puncta in the nerve ring were  
443 counted in N2 (wild-type), *inx-18(ok2454)* and *inx-19(tm1896)* backgrounds. Each dot represents  
444 an individual worm and error bars are  $\pm$ SEM. Ordinary one-way ANOVA between three groups  
445 showed significant differences ( $F[2,12]=5.763$ ,  $p=0.02$ ,  $\alpha=0.05$ ). Dunnett's multiple comparison test  
446 showed that INX-19 ASK puncta were decreased in *inx-18(ok2454)* ( $n=5$ ,  $p=0.01$ ) and in *inx-*  
447 *19(tm1896)* ( $n=5$ ,  $p=0.05$ ) in comparison to N2 ( $n=5$ ). **B)** *inx-19* cDNA was expressed using *Psrd-10*  
448 and puncta in the nerve ring were counted in N2, *inx-18(ok2454)* and *inx-19(tm1896)* backgrounds.  
449 Each dot represents an individual worm and error bars are  $\pm$ SEM. Ordinary one-way ANOVA  
450 between three groups showed no significant differences ( $F[2,14]=0.814$ ,  $p=0.46$ ,  $\alpha=0.05$ ). **C)** *inx-18*  
451 cDNA was expressed using *Psra-9* and puncta in the nerve ring were counted in N2, *inx-18(ok2454)*  
452 and *inx-19(tm1896)* backgrounds. Each dot represents an individual worm and error bars are  
453  $\pm$ SEM. Ordinary one-way ANOVA between three groups showed no significant differences  
454 ( $F[2,13]=1.637$ ,  $p=0.23$ ,  $\alpha=0.05$ ). **D)** *Inx-18(ok2454);inx-19(tm1896)* double mutant animals were

455 assayed for sensitivity to 1 mM quinine using the quinine drop test. Double mutants responded at  
456 higher rates than either *inx-18* or *inx-19* single mutants. N2=18%, n=510; *inx-19(tm1896)*=44%,  
457 n=390; *inx-18(ok2454)*=44%, n=350; *inx-19;inx-18*=53%, n=180, p=0.05 vs *inx-19*, p=0.05 vs *inx-18*.  
458 All groups were compared with a Chi-square test (p<0.0001,  $\alpha=0.05$ ), and post-hoc Fisher's Exact  
459 tests with Bonferroni's correction ( $\alpha=0.025$ ) were computed to compare the double mutant to  
460 single mutant animals.

461 **Figure 5: ASK and ASH architecture is unaltered in *inx-18* and *inx-19* mutant animals**

462 **A)** Diagram of neural architecture of ASK, ASH, and ASI in the *C. elegans* head. The dendrites  
463 reach out to the nose while the axons extend from the cell body into the nerve ring around the  
464 isthmus of the pharynx. **B-D)** Representative confocal images of the worm head with *Psra-*  
465 *9::mCherry* (ASK) and *Posm-10::bfp* (ASH and weakly in ASI) show cell bodies, dendrites extending  
466 to the nose, and axons projecting into the nerve ring. Images on the left include maximum intensity  
467 projections of the mCherry and BFP images superimposed upon a brightfield image to show  
468 location of cells; images on the right are maximum intensity projections of the mCherry and BFP  
469 channels without the brightfield image to show details of the cell architecture. Comparison between  
470 N2 (wild-type), *inx-19(tm1896)*, and *inx-18(ok2454)* (15-20 animals per genotype were imaged)  
471 show no major differences in cell architecture.

472 **Figure 6: ASK  $\text{Ca}^{2+}$  responses to quinine presentation are unaltered in *inx-18* and *inx-19*  
473 mutant animals while ASH  $\text{Ca}^{2+}$  responses are heightened in both**

474 **A)** GCaMP6s fluorescence intensity in ASH in response to 10 mM quinine. Cells were imaged for  
475 30s with presentation of quinine at 10s. The *lite-1(ce314)* mutation was included to eliminate blue-  
476 light induced calcium responses in ASK and ASH. All genotypes showed an increase in ASH  
477 GCaMP6s fluorescence in response to quinine presentation, though for *lite-1;inx-19(tm1896)* and  
478 *lite-1;inx-18(ok2454)* animals the response is larger and faster than that of *lite-1(ce314)*. Averaged

479 GCaMP6s traces are shown and error bars are  $\pm$ SEM. n=48 animals for all genotypes tested. **B)**  
480 GCaMP6s fluorescence intensity in ASK in response to 10 mM quinine. ASK showed small increases  
481 of GCaMP6s signals and there were no significant differences between genotypes. Averaged GCaMP  
482 traces are shown and error bars are  $\pm$ SEM. n=24, n=21 and n=22 animals imaged for *lite-1(ce314)*,  
483 *lite-1;inx-19* and *lite-1;inx-18*, respectively. **C, D)** Heatmaps showing individual traces from all  
484 worms analyzed. Data points in the heatmaps represent GCaMP6s signals normalized to the  
485 averaged fluorescence intensity of the first 3 seconds of imaging. **E)** Quantification of ASH  
486 fluorescence change at four seconds after quinine stimulation. One-way ANOVA between three  
487 groups showed significant differences ( $F[2,141]=3.89$ ,  $p=0.02$ ,  $\alpha=0.05$ ), and Dunnett's multiple  
488 comparison test showed that mean ASH GCaMP6s fluorescence change in *lite-1(ce314)* animals  
489 (n=48) differed from both *lite-1;inx-19* (n=48,  $p=0.02$ ) and *lite-1;inx-18* (n=48,  $p=0.05$ ) animals. **F)**  
490 Quantification of ASK fluorescence change four seconds after quinine stimulation. One-way ANOVA  
491 between three groups showed no significant differences in ASK GCaMP6s fluorescence  
492 ( $F[2,64]=0.202$ ,  $p=0.817$ ,  $\alpha=0.05$ ) between *lite-1(ce314)* (n=24), *lite-1;inx-19* (n=21) and *lite-1;inx-*  
493 *18* animals (n=22).

494 **Figure 7: Mutations in inx-18 and inx-19 disrupt endogenous cGMP levels in ASK and**  
495 **ASH**

496 **A)** Diagram of FlincG3 .The cGMP binding domains of PKG 1 $\alpha$  (blue) are followed by circularly  
497 permuted EGFP (green) and a short PKG 1 $\alpha$  tail (blue). WingG2 increases in brightness in response  
498 to cGMP. **B)** Example of FlincG3 and mScarlet expression within ASH. Ellipses were drawn around  
499 the cell body to measure fluorescence intensity. **C)** cGMP levels within the ASH cell body. The ratio  
500 between mean fluorescence intensity of FlincG3 and mScarlet signals was determined for each  
501 genotype. Decreases in ASH FlincG3 fluorescence were found in *inx-18(ok2454)* and *inx-19(tm1896)*  
502 mutant animals when compared to wild-type worms. Each data point was obtained from a single  
503 cell; error bars are  $\pm$ SEM. One-way ANOVA between three groups showed significant differences

504 (F[2,68]=3.643, p=0.03,  $\alpha$ =0.05), and Dunnett's multiple comparison test showed that mean  
505 fluorescence intensity in *lite-1(ce314)* (n=24) cells differed from both *lite-1;inx-18* cells (n=24,  
506 p=0.05) and *lite-1;inx-19* cells (n=23, p=0.04). **D)** cGMP levels within the ASK cell body. ASK FlincG3  
507 fluorescence was not altered in *inx-18(ok2454)* mutant animals, and increased in *inx-19(tm1896)*  
508 mutant animals when compared to wild-type animals. Each data point was obtained from a single  
509 cell; error bars are  $\pm$ SEM. One-way ANOVA between three groups showed significant differences  
510 (F[2,72]=8.115, p=0.0007,  $\alpha$ =0.05), and Dunnett's multiple comparison test showed that mean  
511 fluorescence intensity in *lite-1(ce314)* cells (n=26) did not differ from *lite-1;inx-18* cells (n=25,  
512 p=0.87) but was increased in *lite-1;inx-19* cells (n=24, p=0.0008).

513 **Figure 8: Model of ASK-ASH electrical synapse facilitation of ASH modulation**

514 Our study supports a model in which ASK-ASH electrical synapses facilitate the passage of cGMP  
515 from ASK to ASH. Within ASH, cGMP downregulates calcium signals in response to quinine  
516 stimulation, leading to a reduction in aversive behavior. INX-19 (orange) is shown on both sides of  
517 the ASK-ASH electrical synapses while INX-18 (purple) is shown joining with an unknown innexin  
518 and contributing to INX-19-based synapse localization.

519 **Figure S1: *inx-18* and *inx-19* mutant animals respond normally to control solutions**

520 **A)** *inx-19(tm1896)* and *inx-18(ok2454)* mutant animals respond at N2 (wild-type) levels when  
521 presented with M13 buffer, while *inx-19(ky634)* animals respond slightly more than wild-type  
522 animals. N2=13%, n=330; *inx-19(ky634)*=23%, n=120, p=0.012 ; *inx-19(tm1896)*=19%, n=210,  
523 p=0.07; *inx-18(ok2454)*=16%, n=160, p=0.33. All groups were compared with a Chi-square test  
524 (p=0.05,  $\alpha$ =0.05), and post-hoc Fisher's Exact tests with Bonferroni's correction ( $\alpha$ =0.017) were  
525 computed to compare each group to the control. **B)** *inx-19(ky634)*, *inx-19(tm1896)*, and *inx-*  
526 *18(ok2454)* mutant animals respond at wild-type levels when presented with 10 mM quinine.  
527 N2=93%, n=330; *inx-19(ky634)*=97%, n=120, p=0.18; *inx-19(tm1896)*=97%, n=210, p=0.03; *inx-*

528  $18(ok2454)=98\%$ ,  $n=120$ ,  $p=0.02$ . All groups were compared with a Chi-square test ( $p=0.02$ ,  
529  $\alpha=0.05$ ), and post-hoc Fisher's Exact tests with Bonferroni's correction ( $\alpha=0.017$ ) were computed  
530 to compare each group to the control.

531 **Figure S2: *inx-19(ky634)* mutant animals have movement defects**

532 **A)** *Inx-19(ky634)* mutant animals reverse more frequently than N2 (wild-type) animals.  
533 Number of reversals were counted from a one-minute video. One-way ANOVA between three  
534 groups showed significant differences ( $F[2,99]=6.943$ ,  $p=0.0015$ ,  $\alpha=0.05$ ), and Dunnett's multiple  
535 comparison test showed that N2 ( $n=34$ ) differed from *inx-19(ky634)* ( $n=33$ ,  $p=0.0006$ ) but not *inx-*  
536 *19(tm1896)* ( $n=35$ ,  $p=0.097$ ). **B)** *inx-19(ky634)* mutant animals have lower average movement  
537 velocity than N2 animals. One-way ANOVA between three groups showed significant differences  
538 ( $F[2,99]=6.089$ ,  $p=0.003$ ,  $\alpha=0.05$ ), and Dunnett's multiple comparison test showed that N2 ( $n=34$ )  
539 differed from *inx-19(ky634)* ( $n=33$ ,  $p=0.021$ ) but not *inx-19(tm1896)* ( $n=35$ ,  $p=0.677$ ). Each data  
540 point represents a single worm and error bars are  $\pm$ SEM.

541 **Figure S3: Responses of worms carrying rescue transgenes to negative and positive  
542 control solutions**

543 **A)** *Inx-19(tm1896)* animals carrying rescue transgenes behaved like N2 (wild-type) animals  
544 when presented with M13 buffer. N2=14%,  $n=220$ ; *inx-19(tm1896)*=19%,  $n=210$ ; *inx-19;Pinx-*  
545 *19::inx-19cDNA*=10%,  $n=100$ ; *inx-19;Psra-9::inx-19cDNA*=10%,  $n=100$ ; *inx-19;Posm-10::inx-*  
546 *19cDNA*=11%,  $n=110$ ; *inx-19;Psra-9::inx-19cDNA; Posm-10::inx-19cDNA*=10%,  $n=110$ . All groups  
547 were compared with a Chi-square test ( $p=0.12$ ,  $\alpha=0.05$ ) **B)** *Inx-18(ok2454)* animals carrying rescue  
548 transgenes behaved like N2 animals when presented with M13 buffer. N2=12%,  $n=120$ ; *inx-*  
549 *18(ok2454)*=7%,  $n=120$ ; *inx-18;inx-18gDNA*=4%,  $n=100$ ; *inx-18;Psra-9::inx-18cDNA*=9%,  $n=120$ . All  
550 groups were compared with a Chi-square test ( $p=0.16$ ,  $\alpha=0.05$ ). **C)** *Inx-19(tm1896)* animal carrying  
551 neuron-specific transgenes behaved like N2 animals when presented with 10 mM quinine, but

552 expression of *inx-19* cDNA using the native promoter reduced the responses to 10 mM quinine  
553 below wild-type levels. N2=96%, n=220; *inx-19(tm1896)*=97%, n=210; *inx-19;Pinx-19::inx-*  
554 *19cDNA*=85%, n=100, p=0.002 vs N2, p=0.0004 vs *inx-19*; *inx-19;Psra-9::inx-19cDNA*=91%, n=100,  
555 p=0.10 vs N2, p=0.04 vs *inx-19*; *inx-19;Posm-10::inx-19cDNA*=97%, n=110, p=0.76 vs N2, p>0.99 vs  
556 *inx-19*; *inx-19;Psra-9::inx-19cDNA*; *Posm-10::inx-19cDNA* =96%, n=110, p>0.99 vs N2, p=0.74 vs *inx-*  
557 *19*. All groups were compared with a Chi-square test (p<0.0002,  $\alpha=0.05$ ), and post-hoc Fisher's  
558 Exact tests with Bonferroni's correction ( $\alpha=0.006$ ) were computed to compare each group to N2  
559 and *inx-19(tm1896)*. **D**) When expressing *inx-18* cDNA under the native promoter or in ASK, *inx-*  
560 *18(ok2454)* animals behaved like wild-type when presented with 10 mM quinine. N2=97%, n=120;  
561 *inx-18(ok2454)*=95%, n=120; *inx-18;inx-18gDNA*=91%, n=100; *inx-18;Psra-9::inx-18cDNA*=91%,  
562 n=120. All groups were compared with a Chi-square test (p=0.21,  $\alpha=0.05$ )

563 **Materials and Methods**

564 **C. elegans Culture**

565 Strains were maintained at room temperature (20-21°C) on NGM agar plates seeded with OP50  
566 *E. coli* bacteria. The N2 strain (Bristol, England) was used as wild type. The following mutant strains  
567 were used in this study: CX6161 *inx-19 (ky634)* I, FX01896 *inx-19 (tm1896)* I, RB1896 *inx-18*  
568 (*ok2454*) IV, BJH2183 *inx-18 (ok2454)* IV; *inx-19(tm1896)* I, BJH2259 *lite-1 (ce314)* X, BJH2304 *lite-*  
569 *1(ce314);inx-19(tm1896)*, and BJH2303 *lite-1(ce314);inx-18(ok2454)*.

570 **Transgenes**

571 Transgenic strains for rescue experiments were generated by microinjection[75] of various  
572 innexin-containing plasmids (30-40 ng/ $\mu$ l) together with co-injection markers. The co-injection  
573 markers were *Punc-122::gfp* (BJP-I15, 20-40 ng/ $\mu$ l) and *Punc-122::mcherry* (BJP-I14, 30-40 ng/ $\mu$ l).  
574 Cytoplasmic fluorophores (mCherry, mTagBFP2, and mScarlet) were injected at 30-40ng/ $\mu$ l. For

575 GCaMP imaging experiments, plasmids (BJP-L136, *Psrbc-66::GCaMP6s::SL2::mCherry::let-858utr* or  
576 BJP-L137, *Posm-10::GCaMP6s::SL2::mCherry::let-858utr*) were injected at 70 ng/μl into the light-  
577 insensitive *lite-1(ce314)* worms. To quantify cGMP levels, FlincG3 plasmids (pFG270, *Psrbc-*  
578 *66::FlincG3::unc-54utr* or pFG250, *Psrd-10::FlincG3::unc-54utr*) were injected at 20 ng/μl into *lite-*  
579 *1(ce314)* worms.

580 ***Behavioral Assays***

581 Well-fed day 1 adults were used for all analyses. To ensure uniformity of worm age and feeding  
582 status, L4 animals were picked onto fresh plates the afternoon before behavior tests. Behavior  
583 assays were performed on at least 5 separate days in parallel with controls.

584 ***Quinine Drop Test***

585 The quinine drop test was performed as described previously[31, 32, 45]. Quinine HCl (Sigma-  
586 Aldrich Q1125) was dissolved in M13 Buffer pH 7.4 (30 mM Tris-HCl pH 7.0, 100 mM NaCl, 10 mM  
587 KCl) to 10 mM. Aliquots were frozen at -20°C. Aliquots were defrosted on the day of the experiment  
588 and allowed to reach room temperature before use. Solutions were loaded into glass needles via  
589 mouth pipetting through long silicone tubing. Needles were formed from 1.5 mm filamented glass  
590 capillaries (World Precision Instruments, Inc.) with a Sutter micropipette puller and the tips  
591 opened by breaking with fine forceps. 10cm NGM plates were brought to room temperature on the  
592 bench overnight and then left open at room temperature to dry for 2.5-4 hours before being used  
593 (plates are appropriately dry when worms leave tracks on the agar that do not immediately  
594 disappear). For each assay, 15 worms were placed on a plate and allowed to acclimate for 30 min.  
595 Small drops (approximately 1 body length in diameter) of M13, 1 mM quinine, or 10 mM quinine  
596 were then delivered via glass needle approximately 1 body length in front of worms. When worms  
597 encountered the drop, they were scored as avoiding the drop if they initiated a reversal within 4 s  
598 and reversed at least half a body length backwards. To avoid desensitization, worms were not

599 tested with a new solution within 2min of initial drop presentation. The experimenter was blind to  
600 the strain when scoring reversals.

601 ***Movement Assays***

602 5 worms at a time were placed on 10 cm NGM plates and allowed to acclimate for 1 minute.  
603 Video capturing was then carried out using an imaging set up from MBF Bioscience. Freely crawling  
604 worms were monitored for 60 seconds at 5 frames per second. Moving velocity at each frame was  
605 calculated by the WormLab 4.1 from MBF Bioscience after confirming correct assignment of head  
606 location throughout the video. Reversals were denoted with negative values. Comparison of  
607 number of reversals/min and mean velocity was calculated using an ordinary one-way ANOVA  
608 using Dunnett's correction for multiple comparisons between all groups. The alpha value was set at  
609 0.05.

610 ***Confocal Microscopy for Imaging Synapse and Cell Architecture***

611 Young adults were paralyzed using 30 mg/ml 2,3-butanedione monoxime (BDM) dissolved in  
612 M9. Worms were imaged using an Olympus FV1000 confocal system with a 60x oil objective (NA  
613 1.4). Z-stacks of fluorescent images (0.40  $\mu$ m step-size for synapses, or 1.20  $\mu$ m step-size for cell  
614 architecture) were taken at the region of interest. Maximum intensity projections of images were  
615 obtained using Fiji. For colocalization analysis, the number of puncta within the nerve ring in each  
616 channel was counted, and scored as colocalizing (containing signal from both channels) or non-  
617 colocalizing (containing signal from a single channel). Percentage colocalization was calculated by  
618 determining the ratio between the number of colocalizing puncta and the total number of puncta in  
619 each maximum intensity projection.

620 ***Calcium Imaging***

621 GCaMP6s[76] was used for all calcium imaging. *Lite-1(ce314)* worms were injected with either  
622 *Psra9::GCaMP6s::SL2::mCherry::let-858utr* (ASK) or *Posm-10::GCaMP6s::SL2::mCherry::let-858utr*  
623 (ASH) along with the co-injection marker *Punc-122::mCherry*. Transgenic lines were crossed with  
624 mutant animals to generate *lite-1(ce314);inx-19(tm1896)* and *lite-1(ce314);inx-18(ok2454)*, which  
625 carry the identical extrachromosomal arrays for imaging. Worms were imaged using a microfluidic  
626 olfactory chip[77]. M13 buffer was used to load worms into the chip, and their nose tips were  
627 washed with M13 buffer for 30 seconds before each recording. At the start of the recording, animals  
628 were exposed to M13 buffer for 10 s before 10 mM Quinine dissolved in M13 was washed in to the  
629 chip. The images were captured at 5 frames per second with an exposure time of 100ms on a Leica  
630 DMI3000B inverted microscope with a 63x Oil objective and a QImaging OptiMOS camera. The  
631 region of interest was defined as a square-shaped area surrounding the desired cell body.  
632 Background-subtracted fluorescence intensity values were collected from every sample's ROI and  
633 stored into MATLAB formatted files. Change in fluorescence intensity ( $\Delta F/F\%$ ) was calculated by  
634 dividing each value by the average intensity of the first 3 seconds of imaging.

635 **cGMP Imaging**

636 FlincG3[55, 56] was used for cGMP imaging. *Lite-1(ce314)* worms were injected with either  
637 *Psrbc-66::FlincG3::unc-54utr* and *Psrbc-66::mScarlet::unc-54utr* (ASK) or *Psrd-10::FlincG3::unc-54utr*  
638 and *Psrd-10::mScarlet::unc-54utr* (ASH) along with the co-injection marker *Punc-122::mCherry*.  
639 Transgenic lines were crossed with mutant animals to generate *lite-1(ce314);inx-19(tm1896)* and  
640 *lite-1(ce314);inx-18(ok2454)*, which carry the identical extrachromosomal arrays for imaging. L4  
641 worms were picked onto fresh OP50-seeded NGM plates 6 hours before imaging to ensure  
642 synchronization of age and feeding status. Young adults were paralyzed with 30 mg/ml BDM  
643 dissolved in M9. Immobilized worms were imaged using an Olympus FV1000 confocal microscope  
644 with a 60x Water objective. Kalman filtering was used to reduce noise. Z-stacks (1.28  $\mu\text{m}$  step-size)

645 were taken through the cell body. Maximum intensity projections were obtained using Fiji[78]. Two  
646 elliptical ROIs were drawn in the mScarlet channel: one surrounding the cell body and one  
647 capturing background fluorescence from a region near the cell body that did not contain an axon or  
648 dendrite. Mean pixel intensity in both the FlincG3 and mScarlet channels was calculated using Fiji  
649 and background intensity was subtracted from cell body intensity. The ratio between FlincG3 and  
650 mScarlet mean intensity was calculated to control for expression variation.

651 **DNA constructs**

Name	Construct	Construction Notes
BJP-L109	<i>Pinx-19::inx-19a::gfp::unc-54utr</i>	<i>Pinx-19</i> (5556bp) is from Dr. Cornelia Bargmann and primers were: GATAAGCGCGGATGCTCCT and TGACAGTGCTCTCAGAGGGA. <i>Inx-19a</i> cDNA is from Dr. Cornelia Bargmann and primers were: ATGTGGCGGACACCAGCATC and AAGAAACGATTTCGTCTGTCCAGGA.
BJP-I15	<i>Punc-122::gfp::unc-54utr</i>	
BJP-L99	<i>Psra-9::inx-19a::mCherry::gdp-2utr</i>	<i>Psra-9</i> is 3012bp and primers were: GCATGCTATATTCCACCAAA and GAAATCTTGAAACTGAAAAATACA
BJP-L112	<i>Psra-9::inx-19a::gfp::unc-54utr</i>	
BJP-L125	<i>Psra-6::inx-19a::mCherry::gdp-2utr</i>	<i>Psra-6</i> is 2018bp and primers were TTCCAGTGCTCTGAAAATCTTG and GGCAAAATCTGAAATAATAAATATT
BJP-L114	<i>Posm-10::inx-19a::gfp::unc-54utr</i>	<i>Posm-10</i> (900bp) is from Dr. Josh Kaplan and primers were: CTTGACACCGACTGGCAC and GCGTCGACACCTTGTAAAGAT
BJP-L120	<i>Psrd-10::inx-19a::gfp::unc-54utr</i>	<i>Psrd-10</i> (1841bp) is from Dr. Denise Ferkey and primers were: AGCCACGGCTAGCTACAG and GTTGAATTGGTCTGTGAGCT
	<i>inx-18 gDNA PCR</i>	<i>Inx18</i> gDNA (7646bp) used the primers: ACAGTCGAGTCGTCGTCG and TAATTTGAAACAAAATCGGAAAGAA
BJP-L46	<i>Psra-9::inx-18::gfp::unc-54utr</i>	<i>Inx-18</i> cDNA (1308bp) is from Dr. Zhao-Wen Wang and primers were: ATGGTCGGTGGATTCCG and AACATAATGTGTCCGTGTCGGA
BJP-L115	<i>Psrbc-66::mTagBFP2::unc-54utr</i>	<i>Psrbc-66</i> is 2055bp and used the primers:

		CAACGATGAAATATTGATCGTACAAA and TTCTGAGACACCTGACTTCTGTC
BJP-L116	<i>Posm-10::mTagBFP2::unc-54utr</i>	
BJP-L143	<i>Psrbc-66::mScarlet::unc-54utr</i>	
BJP-L142	<i>Psrd-10::mScarlet::unc-54utr</i>	
BJP-L139	<i>Psra-9::mCherry::unc-54utr</i>	
BJP-L136	<i>Psrbc-66::GCaMP6s::SL2::mCherry::let-858utr</i>	
BJP-L137	<i>Posm-10::GCaMP6s::SL2::mCherry::let-858utr</i>	
pFG270	<i>Psrbc-66::FlincG3::unc-54utr</i>	Received from Dr. Denise Ferkey
pFG250	<i>Psrd-10::FlincG3::unc-54utr</i>	Received from Dr. Denise Ferkey

652 **Statistical Analyses**

653 Statistical analyses for all experiments except calcium imaging were carried out as described in  
654 the legends for each figure using GraphPad Prism Statistical analysis of the calcium imaging  
655 experiments was carried out using a custom written MATLAB program and GraphPad Prism.

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661 Kaplan, and Dr. Zhao-Wen Wang for plasmids.

662 **Authors' Contributions**

663 L.V., I.R., and J.B. conceived of experiments. L.V. designed and performed experiments and  
664 analyzed data. B.U. performed and analyzed GCaMP6s experiments. S.W., D.M.F., and N.D.L.  
665 provided unpublished reagents and assisted with experimental design. L.V. wrote the manuscript  
666 with I.R. and J.B.

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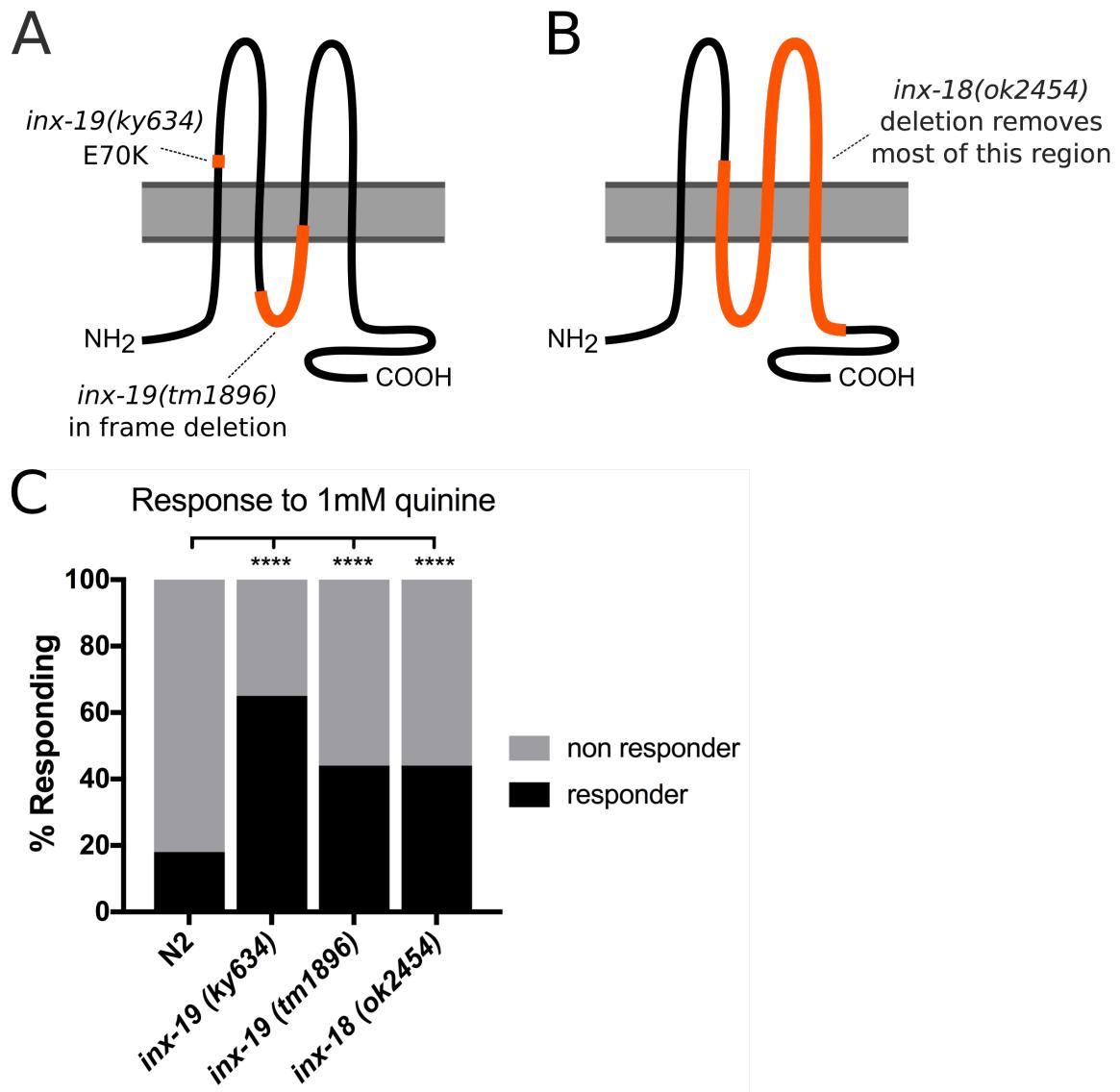
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833

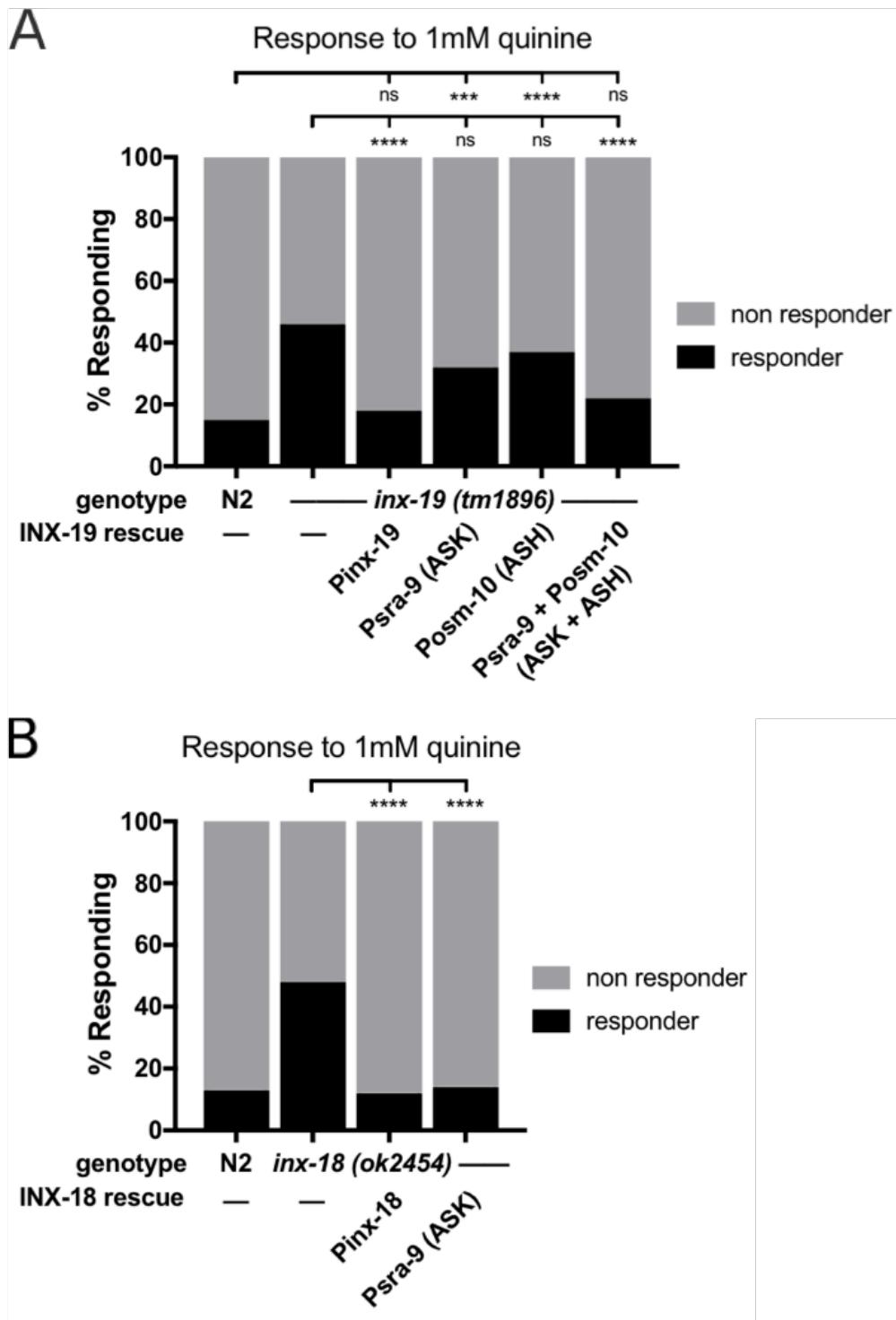
834

835 **Fig 1**



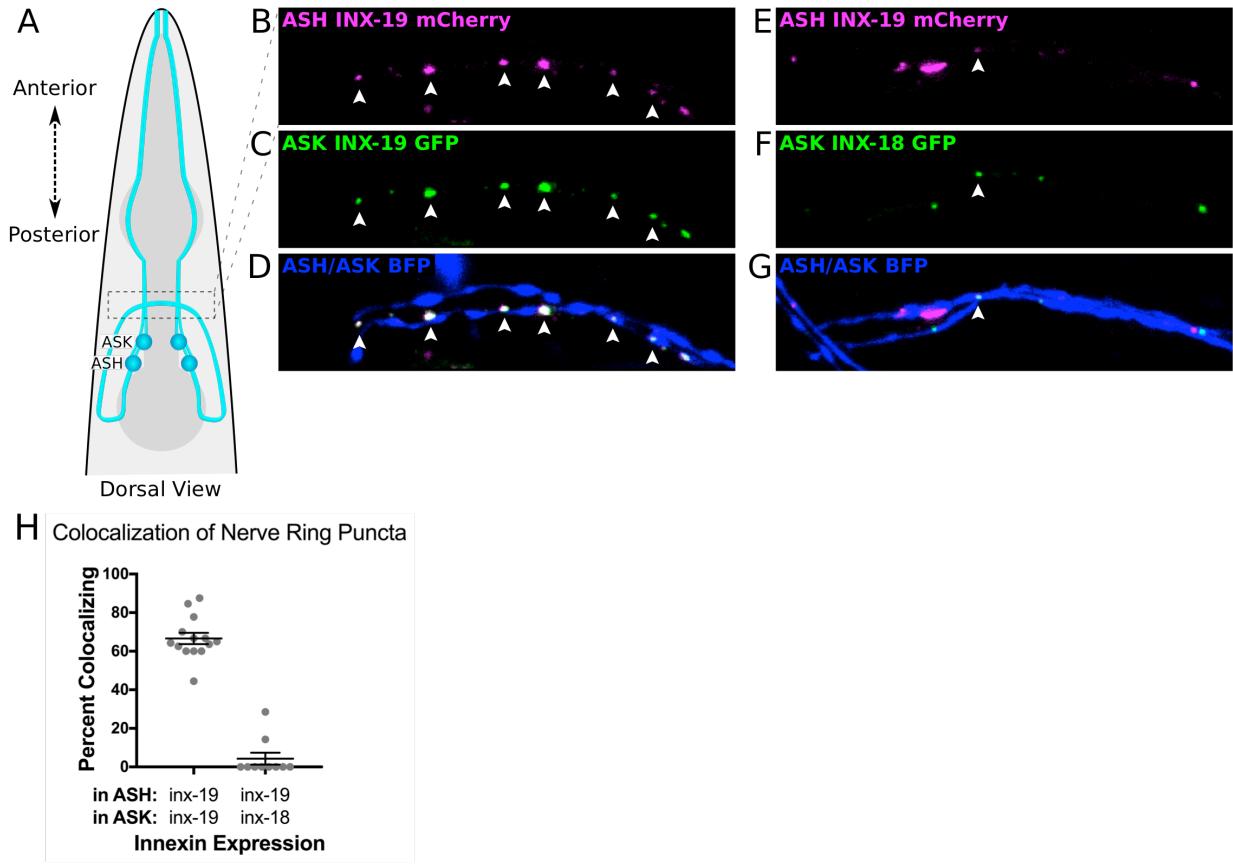
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837 **Fig 2**



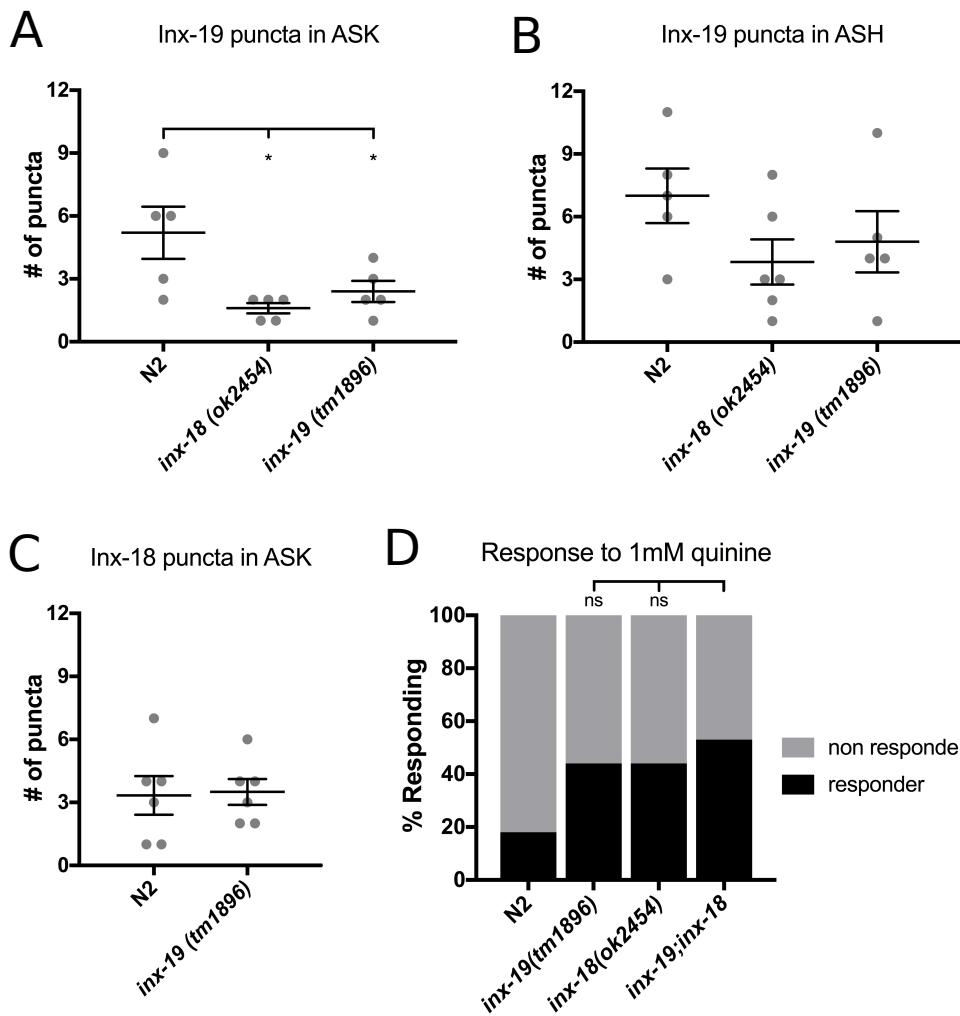
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839 **Fig 3**



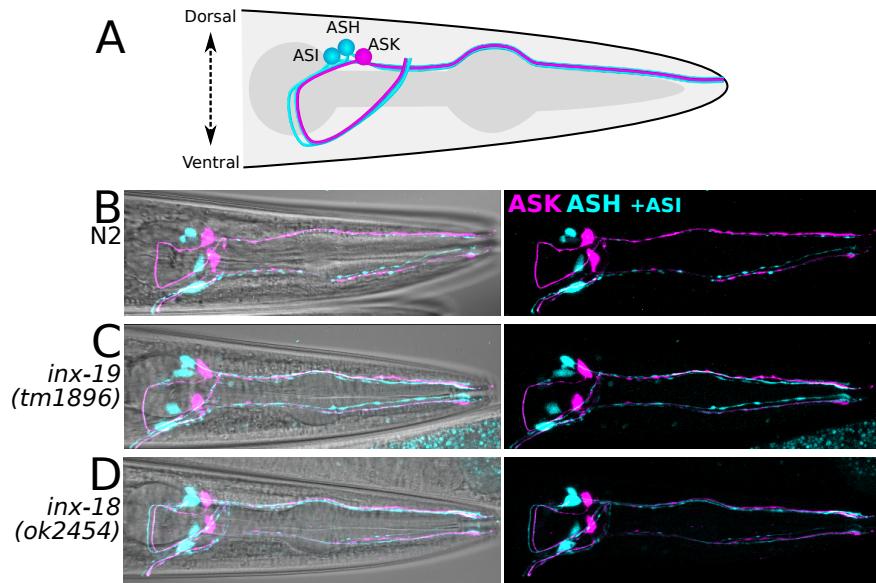
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841 **Fig 4**



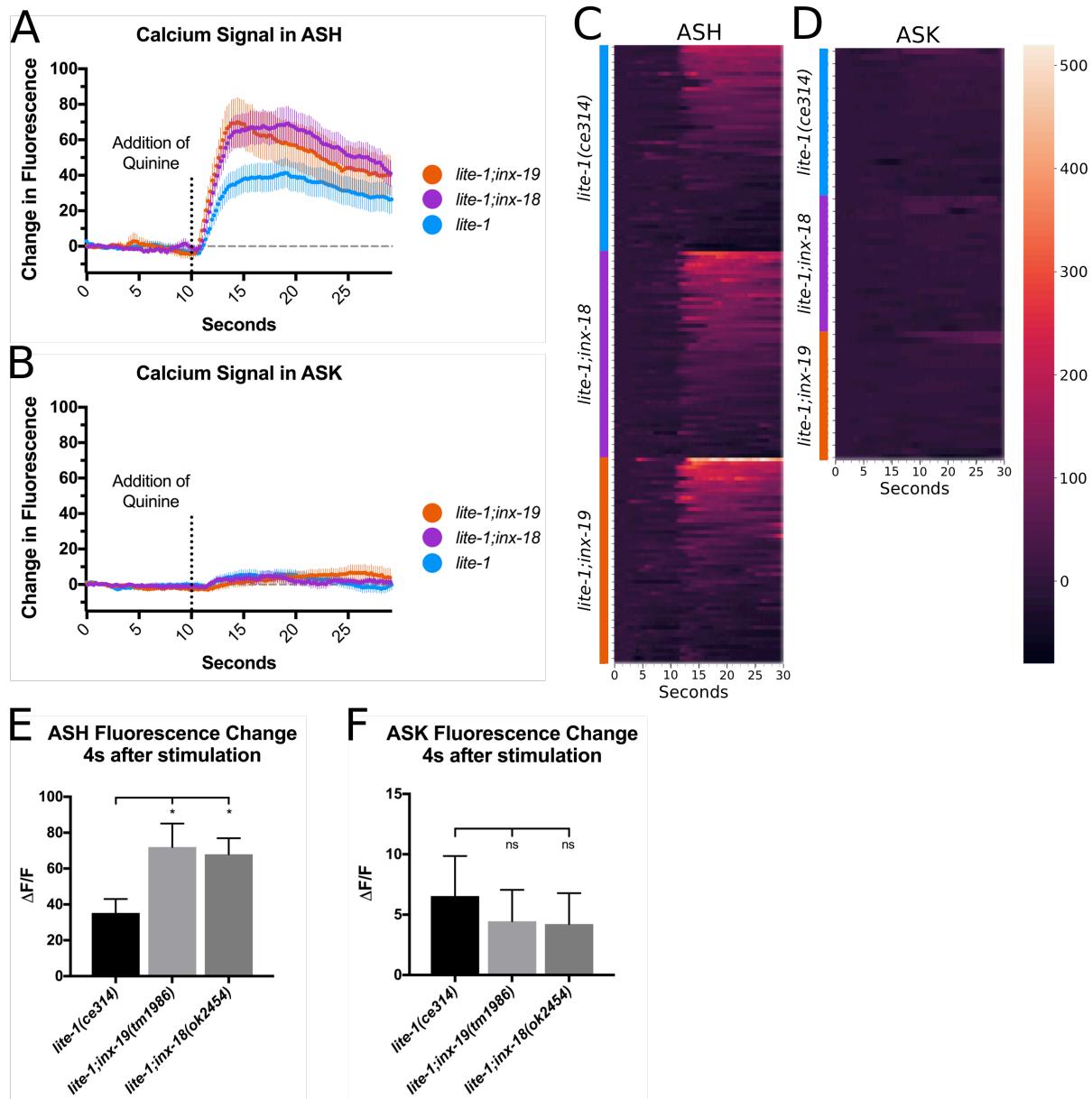
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843 **Fig 5**



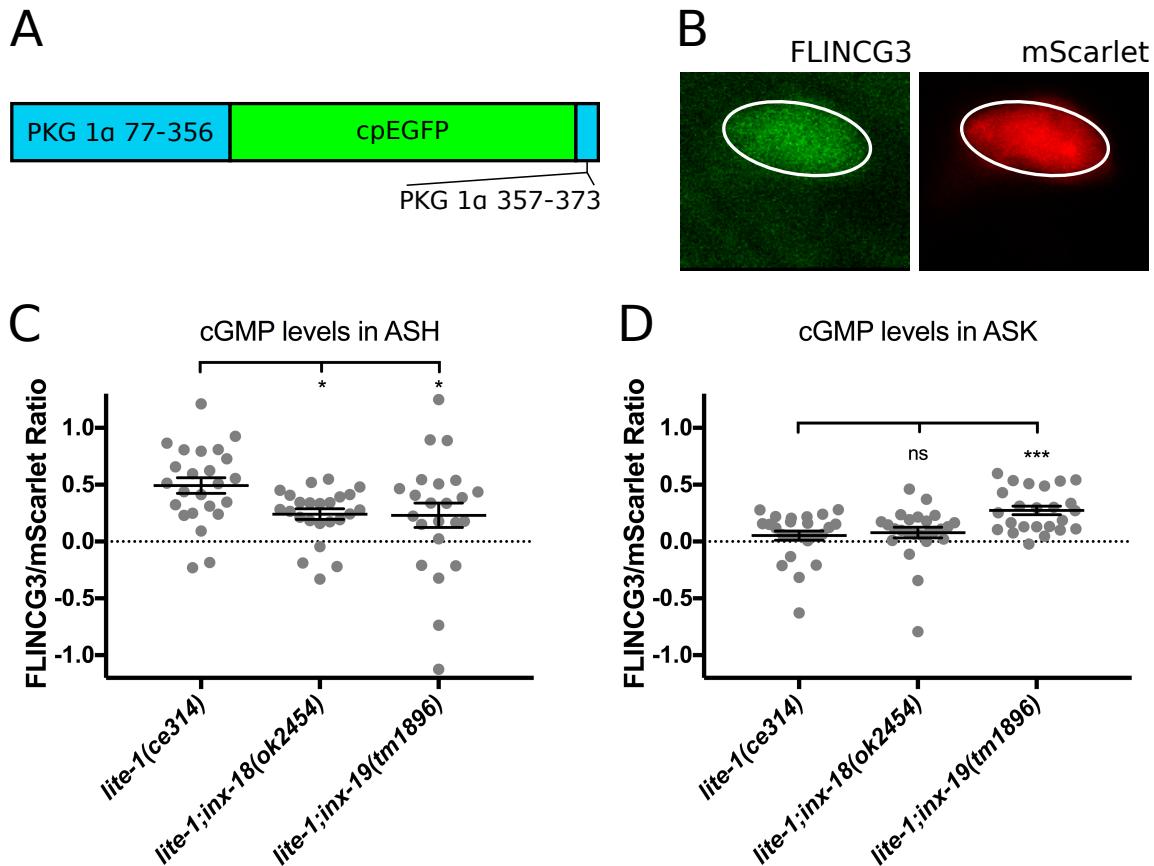
844

845 **Fig 6**



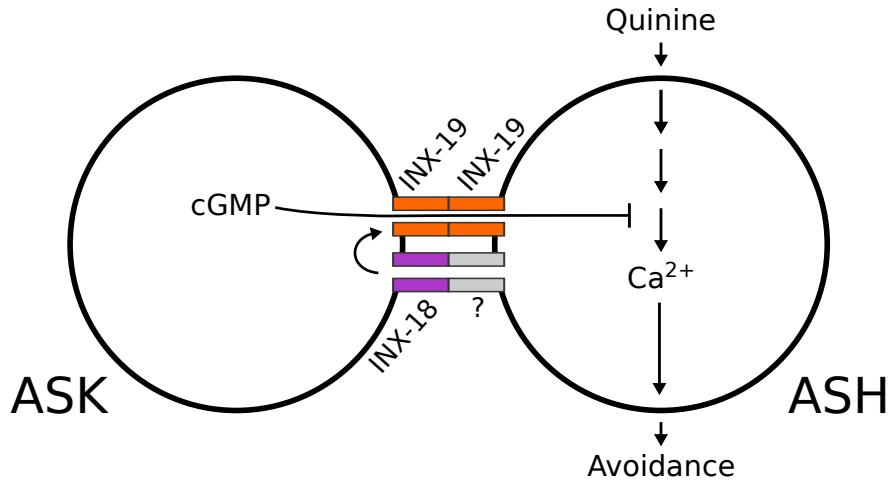
846

847 **Fig 7**



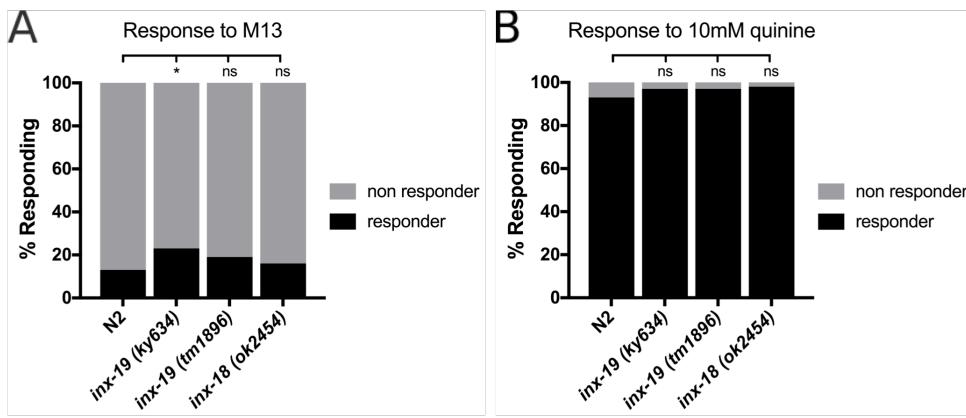
848

849 **Fig 8**



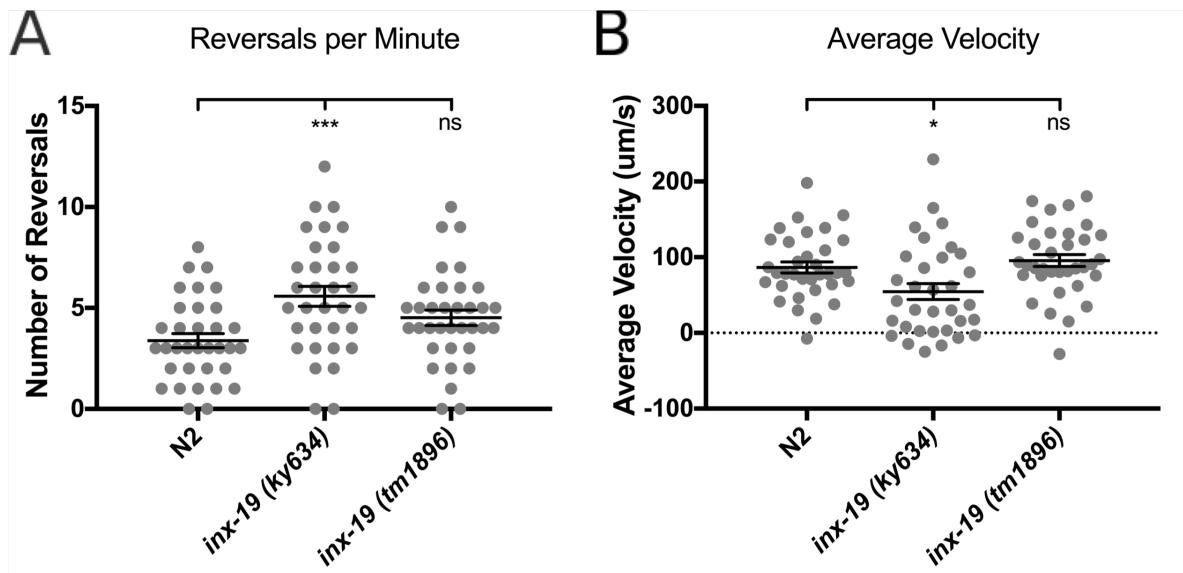
850

851 **Fig S1**



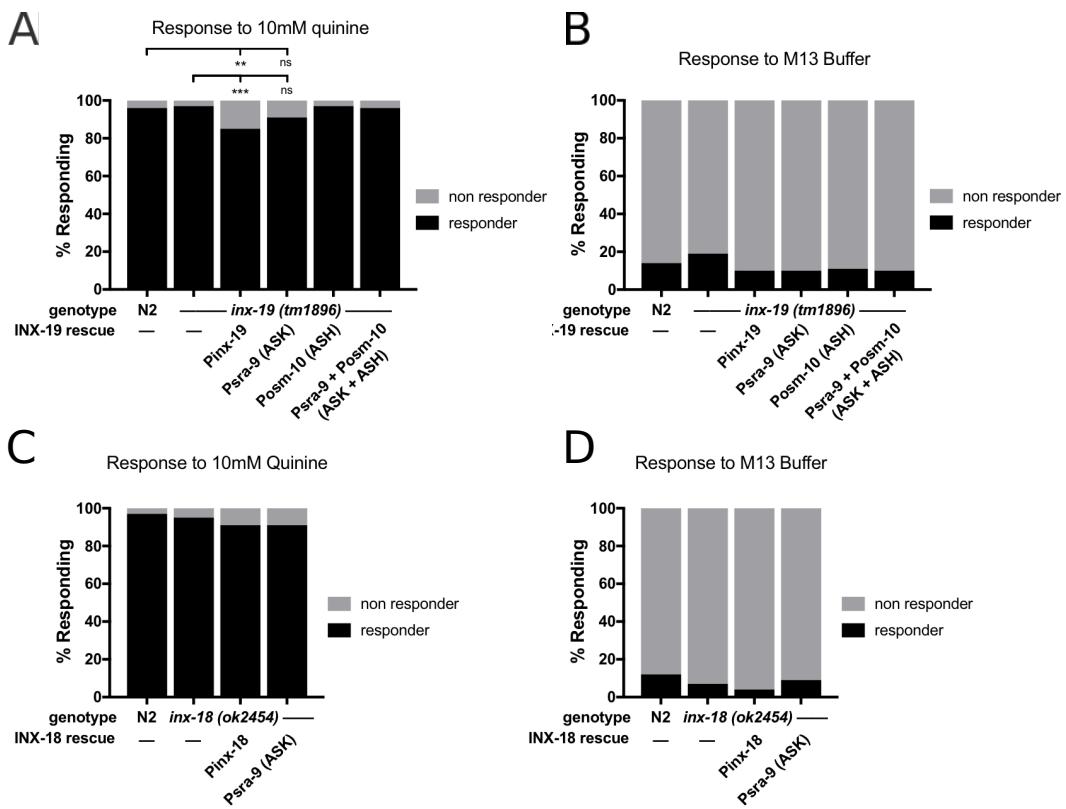
852

853 **Fig S2**



854

855 **Fig S3**



856