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EFF-1 promotes muscle fusion, paralysis and retargets

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infection by AFF-1-coated viruses in *C. elegans*

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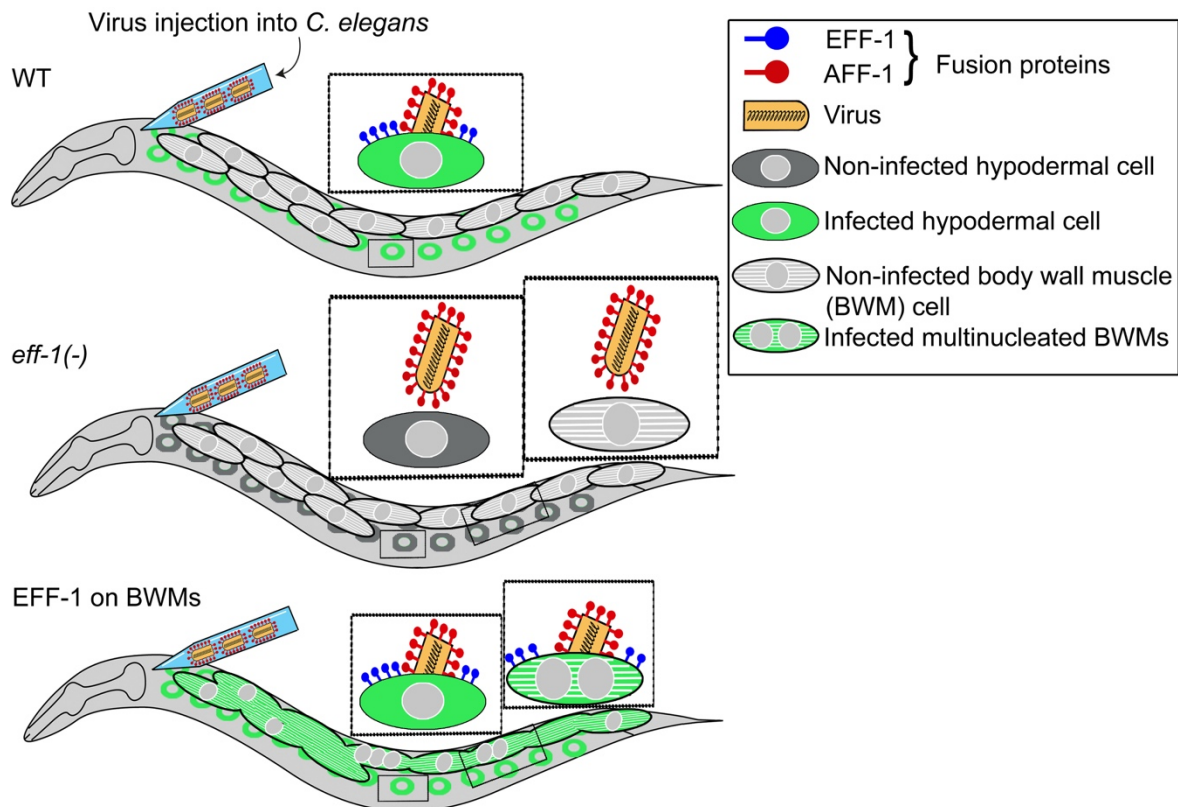
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14 Running title: Abnormally fused muscles infected by fusexin-coated virus

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16 **Graphical abstract:**



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19 **A hallmark of muscle development is that myoblasts fuse to form myofibers. However,**
20 **smooth muscles and cardiomyocytes do not generally fuse. In *C. elegans*, the body wall**
21 **muscles (BWMs), the physiological equivalents of skeletal muscles, are mononuclear.**
22 **Here, to determine what would be the consequences of fusing BWMs, we express the**
23 **cell-cell fusogen EFF-1 in these cells. We find that EFF-1 induces paralysis and dumpy**
24 **phenotypes. To determine whether EFF-1-induced muscle fusion results in these**
25 **pathologies we injected viruses pseudotyped with AFF-1, a paralog of EFF-1, into the**
26 **pseudocoelom of *C. elegans*. When these engineered viruses encounter cells**
27 **expressing EFF-1 or AFF-1 they are able to infect them as revealed by GFP expression**
28 **from the viral genome. We find that AFF-1 viruses can fuse to EFF-1-expressing**
29 **muscles revealing multinucleated fibers that cause paralysis and abnormal muscle**
30 **morphogenesis. Thus, aberrant fusion of otherwise non-syncytial muscle cells may**
31 **lead to pathological conditions.**

32

33 **Significance statement**

34 Most cells are individual units that do not mix their cytoplasm. However, some cells fuse to
35 become multinucleated in placenta, bones and muscles. In most animals, muscles are formed
36 by myofibers that originate by cell-cell fusion. In contrast, in *C. elegans* the body wall muscles
37 are mononucleated cells that mediate worm-like movement. EFF-1 and AFF-1 fusogens
38 mediate physiological cell fusion in *C. elegans*. By ectopically expressing EFF-1 in body wall
39 muscles we induce their fusion resulting in behavioral and morphological deleterious effects,
40 revealing possible causes of congenital myopathies in humans. Using AFF-1-coated
41 pseudoviruses we infect EFF-1-expressing muscle cells retargeting viral infection into these
42 cells. We suggest that virus retargeting can be utilized to study myogenesis, neuronal
43 regeneration, gamete fusion and screens for new fusogens in different organisms. In addition,
44 our virus retargeting system can be used in gene-therapy, viral-based oncolysis and to study
45 viral-host interactions.

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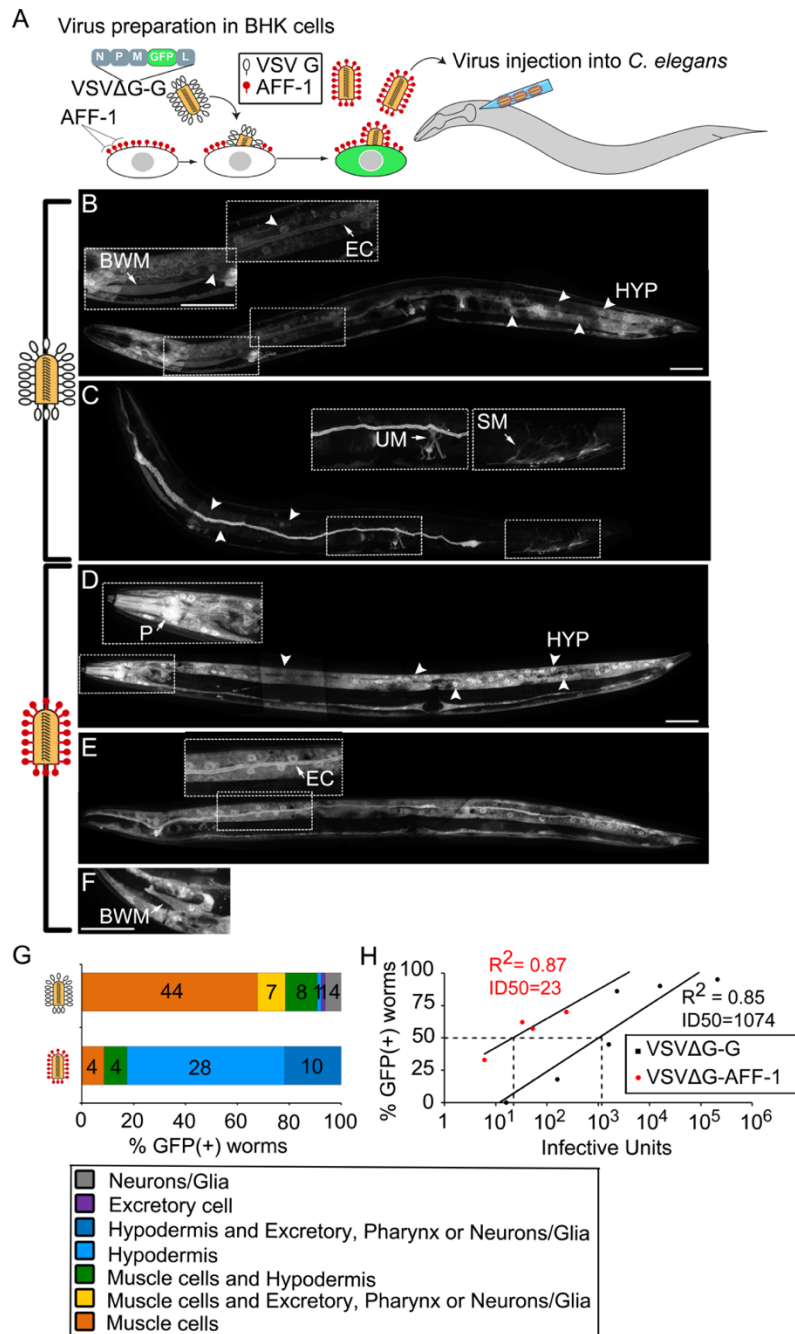
48 **Introduction**

49 Vesicular Stomatitis Virus (VSV) is an enveloped negative strand RNA rhabdovirus. VSV
50 utilizes its surface glycoprotein G (VSV-G) to infect vertebrates and invertebrates and lyses
51 many cell lines tested to date [1–3]. VSV is widely used for pseudotyping other viruses and
52 has a high transduction efficiency [1–4]. These properties turned VSVΔG into a promising
53 vector for gene therapy, tissue regeneration, viral-based oncolysis, and VSV-based vaccines,
54 some of which successfully completed phase III clinical trials [1,5]. Moreover, VSV-based
55 vectors are useful for studying mechanisms of transcription and replication of RNA viruses,
56 cellular trafficking, antiviral responses and fusion proteins (fusogens) [6–9]. However, the
57 major bottleneck for both applied and fundamental research purposes is achieving an efficient
58 and specific targeting of VSV-based vectors into desired cells/tissues of a live, multicellular
59 organism [3]. Like other viruses, VSV lacks specificity for desired target cells as it suggestively
60 enters cells through highly ubiquitous receptors such as the LDL receptor [10]. Retargeting
61 VSV into particular tissues of interest therefore requires blocking the virus' natural interactions
62 providing new, cell-specific interactions. Indeed, mutation or substitution of VSV-G with
63 glycoproteins from other viruses or chimeric glycoproteins coupled with antibodies can
64 retarget VSV to specific cells [3,4,11–13].

65 VSV-G is a class III viral fusogen whereas the *Caenorhabditis elegans* EFF-1 and AFF-
66 1, which fuse somatic cells during development, are structural homologs of class II viral

67 proteins and gamete HAP2(GCS1) fusogens from the fusexin family [14–22]. EFF-1 and AFF-
68 1 also participate in maintaining neuronal architecture and neuronal reconnection following
69 injury [23–27]. Studied viral glycoproteins, including VSV-G, use a unilateral fusion
70 mechanism that depends on the expression of receptors only on the target cells [28,29]. In
71 contrast, EFF-1 and AFF-1 are bilateral fusogens- their presence is required on the
72 membranes of both apposing cells to mediate fusion [6,16,18,30,31]. These two fusogens can
73 act in either a homotypic or a heterotypic manner [6,16,18,21] and mediate heterotypic cell
74 fusion of Sf9 insect and Baby Hamster Kidney (BHK) cells [6,16]. Finally, VSV viruses
75 containing a *GFP* substituting the *VSV-G* coding sequence (*VSVΔG*) [32,33] that are coated
76 with AFF-1 (*VSVΔG-AFF-1*) specifically infect AFF-1 or EFF-1 expressing BHK cells [6]. Thus,
77 in contrast to the pseudotyped virus coated with the native, unilateral G glycoprotein (*VSVΔG-*
78 *G*), infection by *VSVΔG-AFF-1* requires fusogen expression on the host cell membrane. To
79 date, however, *VSVΔG* coated with AFF-1, EFF-1 or any other non-viral fusogen have not
80 been tested for infection in a living organism. Recently *VSVΔG-G* was demonstrated to infect
81 living *C. elegans* [7,34]. Given that: (i) *VSVΔG-AFF-1* requires a fusogen on the target cell for
82 infection, (ii) a detailed cellular atlas of AFF-1 and EFF-1 expression and function in *C. elegans*
83 is known and (iii) *C. elegans eff-1* and *aff-1* mutants are available, we test whether *VSVΔG-*
84 *AFF-1* can be retargeted to specific cells in living *C. elegans*. We found that *AFF-1*-coated
85 viruses infect *C. elegans* and specifically target cells that express functional EFF-1 or AFF-1.
86 Furthermore, *AFF-1*-coated pseudoviruses can be redirected to mononucleated body wall
87 muscles (BWMs) that ectopically express EFF-1. Thus, the new delivery system enabled us
88 to observe that EFF-1-induced BWMs merger and formation of non-functional syncytial
89 muscle fibers, demonstrating the consequences of aberrant muscle fusion in an animal that
90 normally uses mononucleated muscles for gait. Based on our results, we propose that in *C.*
91 *elegans* a layer of longitudinal mononucleated muscles electrically coupled by gap junctions
92 efficiently mediate wavelike movement and ectopic fusion disrupts the morphology and
93 physiology of the muscular system. We also suggest additional applications for *VSVΔG-AFF-*
94 *1* including finding new fusogens and fusogen-expressing tissues, studying fusogen-fusogen
95 interactions and fusogen-interacting proteins *in vivo*, studying neuronal regeneration
96 processes and using specific cell-delivery approaches for other viruses such as coronaviruses
97 and retroviruses in *C. elegans* and in other organisms.

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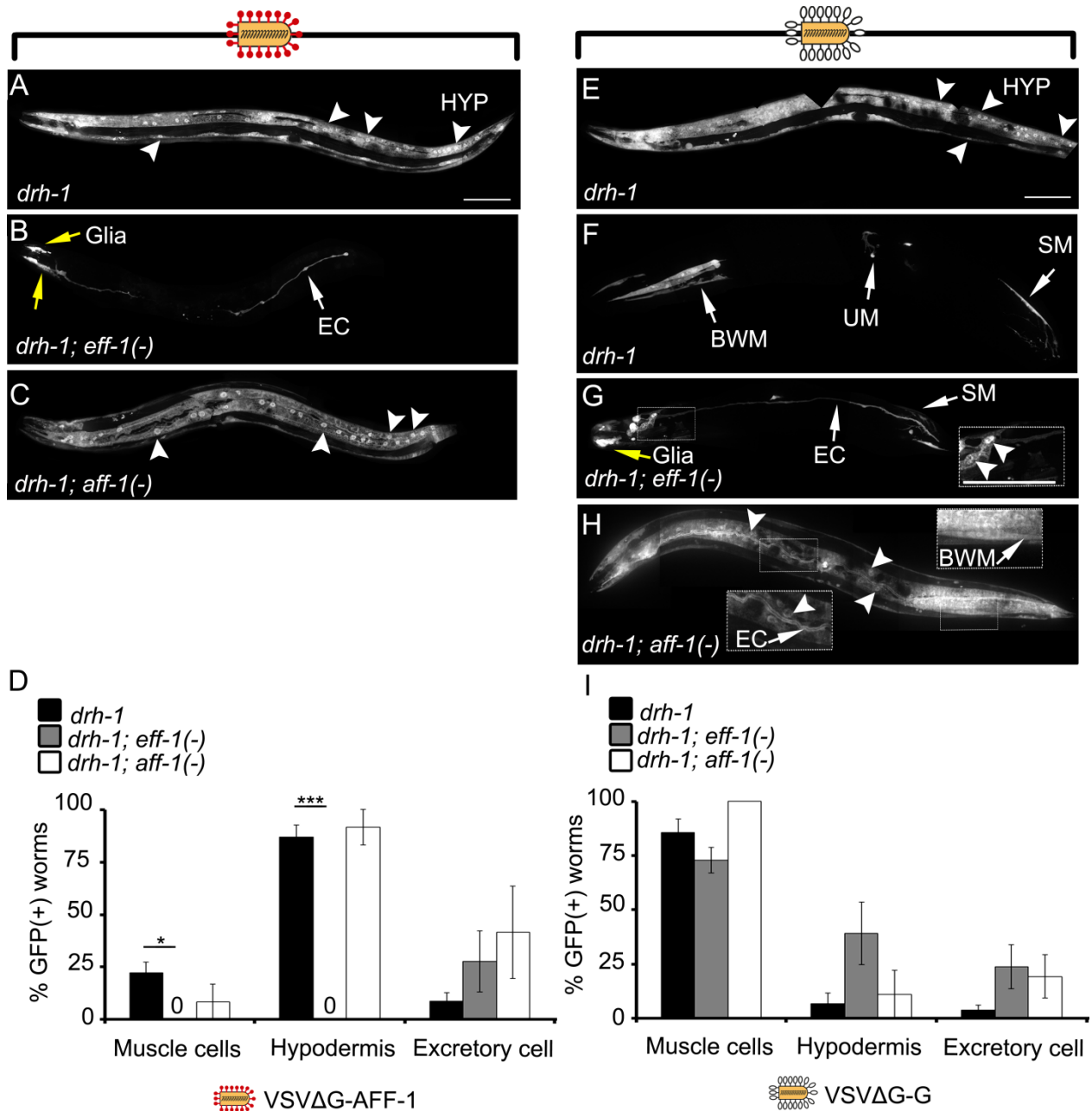
Figure 1. VSVΔG-AFF-1 and VSVΔG-G have different tropisms

102 **(A)** BHK cells expressing AFF-1 (red pins) were infected with VSVΔG-G. The viral genome
 103 encodes GFP replacing VSV-G (white pins). GFP(+) (green), infected cells. VSVΔG-AFF-1
 104 pseudoviruses were harvested from the supernatant, titrated and microinjected into *C.*
 105 *elegans*. **(B-F)** Confocal Z-stack projections of *drh-1* worms injected with VSVΔG-G (2300-
 106 16000 IU) or VSVΔG-AFF-1 (33-240 IU) and imaged 48-72 h later (VSVΔG-G, n=65; VSVΔG-
 107 AFF-1, n=46 from 5 and 6 independent experiments respectively). Images and their insets are
 108 dot boxed. Arrowheads, hypodermal (HYP) nuclei. Arrows, infected cell. BWM-Body Wall
 109 Muscle, UM-Uterine Muscle, SM-Stomatointestinal Muscle, EC-Excretory Cell, P-Pharynx.
 110 Scale bars, 50 μm. **(G)** Distribution of worms with specified infected, GFP(+) cell types. **(H)**
 111 Fraction of GFP(+) worms injected with indicated IU of VSVΔG-G or VSVΔG-AFF-1. Each dot
 112 represents an independent experiment (n=10-28 worms per experiment). R² and the IU doses
 113 producing 50% GFP(+) worms (ID₅₀) are indicated. See also Figure S1.

114 **Results**

115 **VSV Δ G-AFF-1 and VSV Δ G-G infect mostly hypodermal and muscle cells respectively**

116 Wildtype, recombinant or pseudotyped VSV has been shown to infect rodents, fish, farm
117 animals, and primates [1,35–37]. Recently, VSV Δ G-G was shown to infect *C. elegans* [7].
118 Previously, pseudotyped VSV expressing the *C. elegans* somatic fusogen AFF-1 (VSV Δ G-
119 AFF-1) was shown to infect AFF-1 or EFF-1 expressing BHK cells [6], but was not tested in a
120 living organism. To test whether pseudotyped VSV Δ G-AFF-1 [6,33,38] can infect *C. elegans*,
121 we prepared a recombinant VSV strain encoding the fluorescent reporter GFP, coated by AFF-
122 1 fusion protein (VSV Δ G-AFF-1). Briefly, BHK cells expressing AFF-1 were infected with
123 VSV Δ G-G helper virus (Fig. 1A). Newly generated virions were coated by plasma membrane-
124 bound AFF-1, thereby producing VSV Δ G-AFF-1 pseudotyped viruses. Then, we titered the
125 viruses determining the number of viral Infective Units (IU)/ml in BHK-AFF-1 cells. Finally,
126 VSV Δ G-AFF-1 viruses were injected into worm's pseudocoelom; a body cavity filled with fluid
127 that surrounds the internal organs. We expected that virus injected into the pseudocoelom will
128 encounter different cells inside the worm (Figure 1A). Worms were injected with VSV Δ G-AFF-
129 1 that was pre-incubated with α VSV-G antibody (see materials and methods and Figure S4),
130 VSV Δ G-G (as a positive control) or DMEM medium (as negative control). We used *drh-1(-)*
131 (Dicer Related Helicase -1) mutant worms, as they are more susceptible to VSV infection and
132 do not have any observable phenotypes [7,34]. Taken that VSV-G works unilaterally, while
133 AFF-1 and EFF-1 fusogens have to be present in both fusing membranes [6,16,18,30], we
134 hypothesized that VSV Δ G-G and VSV Δ G-AFF-1 will produce different infection patterns.
135 Hence, we characterized and compared the GFP positive (GFP(+)) cells infected by VSV Δ G-
136 G or VSV Δ G-AFF-1. VSV Δ G-G preferentially infected muscle cells, including BWM [7], uterine
137 muscles (UM) and stomatointestinal muscles (SM, Figures 1B and 1C). VSV Δ G-G also
138 infected the epidermal cells (hypodermis, HYP), excretory canal cell (EC, Figure 1B), glia and
139 neurons in the head (Figures S1A-S1C). In contrast, VSV Δ G-AFF-1 infected mostly HYP cells
140 (Figure 1D), and also excretory cell (Figure 1E) pharyngeal muscles (P) (Figure 1D), BWM
141 (Figure 1F) and head neurons and glia (Figures S1D-S1F). VSV Δ G-G infects muscles that do
142 not express any known fusogen (67% of infected, GFP(+) worms had only BWM infection),
143 whereas VSV Δ G-AFF-1 mainly infects the EFF-1-expressing hypodermis (61% of GFP(+)
144 worms had only hyp infection) (Figure 1G). Moreover, the calculated ID50 (infection dose
145 producing 50% GFP(+) worms) for VSV Δ G-G and VSV Δ G-AFF-1 were approximately 1000
146 and 20 IU respectively (Figure 1H). Therefore, VSV Δ G-AFF-1, a virus pseudotyped with a
147 bilateral fusogen, can efficiently infect a multicellular organism such as *C. elegans*. Moreover,
148 VSV Δ G-AFF-1 and VSV Δ G-G are specialized for different cellular targets and have different
149 infectivity levels.



150

151

152 **Figure 2. VSVΔG-AFF-1 requires EFF-1 on target cells for infection**

153 (A-C) Confocal Z-stack projections of *C. elegans* infected with VSVΔG-AFF-1 (33-240 IU) and
 154 imaged 48 h later. *drh-1* (n=46), *drh-1;eff-1(-)* (n=15) and *drh-1;aff-1(-)* (n=10).

155 (D) Fraction of GFP(+) worms from (A-C).

156 (E-H) Confocal Z-stack projections of *C. elegans* infected with VSVΔG-G (2300-4700 IU) 48
 157 h post-injection. *drh-1* (n=56), *drh-1;eff-1(-)* (n=39) or *drh-1;aff-1(-)* (n=15).

158 (I) Fraction of GFP(+) worms from (E-H).

159 Arrowheads, HYP nuclei. Arrows, infected cell. See Figure 1 for cell types. Scale bars, 100
 160 μm. Error bars represent mean ± SEM. *p<0.05, ***p<0.001 (Student's t- test).

161 See also Figure S2.

162

163 **VSV Δ G-AFF-1 requires EFF-1 in host cells for infection**

164 We showed that cells infected by VSV Δ G-AFF-1 included hypodermis, excretory cell, neurons
165 in the head, glia and pharyngeal cells. It is known that in adults pharyngeal cells, head
166 neurons, glia and hypodermal cells express EFF-1 [14], while pharyngeal cells, excretory duct
167 cell and glia express AFF-1 [18,21] and lastly, excretory cell, head neurons, hypodermal cells
168 and pharyngeal cells express EFF-2, a recent duplication of EFF-1 with unknown function
169 (BG, Oren-Suissa and BP unpublished data). Importantly, BWM cells that do not express any
170 known fusogen in *C. elegans* were infected by VSV Δ G-AFF-1 in only 9% of the worms
171 compared to 67% of the worms infected with VSV Δ G-G (Figure 1G). These findings suggest
172 that infection of living *C. elegans* by VSV Δ G-AFF-1 follows a bilateral action mechanism, and
173 that VSV Δ G-AFF-1 is capable of interacting with several members of the fusexin family on the
174 target cell. Therefore, we hypothesized that EFF-1-expressing hypodermal cells could not be
175 infected by VSV Δ G-AFF-1 in the absence of EFF-1, while cells expressing AFF-1 or EFF-2
176 would still be targeted. To test this hypothesis, we injected either VSV Δ G-G or VSV Δ G-AFF-
177 1 into the pseudocoelom of wt, null *eff-1(ok1021)* mutants (*(eff-1(-);[14]*), null *aff-1(tm2214)*
178 mutants (*(aff-1(-);[18]*) or putative null *eff-2(hy51)* (*eff-2(-)*; see materials and methods) worms,
179 and quantified the percentage of worms with GFP(+) cells. We detected VSV Δ G-AFF-1
180 infected hypodermal cells in *eff-1(+)* (Figure 2A) but not in *eff-1(-)* worms (Figure 2B), while
181 GFP(+) excretory and glia cell were detected even in the absence of EFF-1 (Figure 2B). *aff-*
182 *1(-)* worms had GFP(+) hypodermal cells as in wildtype (Figure 2C). Overall, the fraction of
183 worms showing VSV Δ G-AFF-1-infected hypodermal cells was similarly high in wildtype and in
184 *aff-1(-)* (87% and 92% for wt and *aff-1(-)*, respectively) but was absent in *eff-1(-)* mutants
185 (Figure 2D). In addition, muscles were inefficiently infected in wildtype and *aff-1(-)* mutants,
186 and not infected in *eff-1(-)* animals (Figure 2D). The fraction of worms with GFP(+) excretory
187 cells was not significantly different in all backgrounds (Figure 2D). Finally, the fraction of
188 worms with GFP(+) glia/neuron cells was significantly higher in *eff-1(-)*, compared to wt and
189 *aff-1(-)* animals (Figures S1D-S1F). Moreover, *eff-2(-)* and wildtype animals were similarly
190 infected in their hypodermal, muscle and excretory cells (Figure S2). These results show that
191 VSV Δ G-AFF-1 requires EFF-1 expressed on hypodermal cells, for their infection. Moreover,
192 we show evidence for AFF-1-EFF-1 bilateral heterotypic interactions *in vivo*.

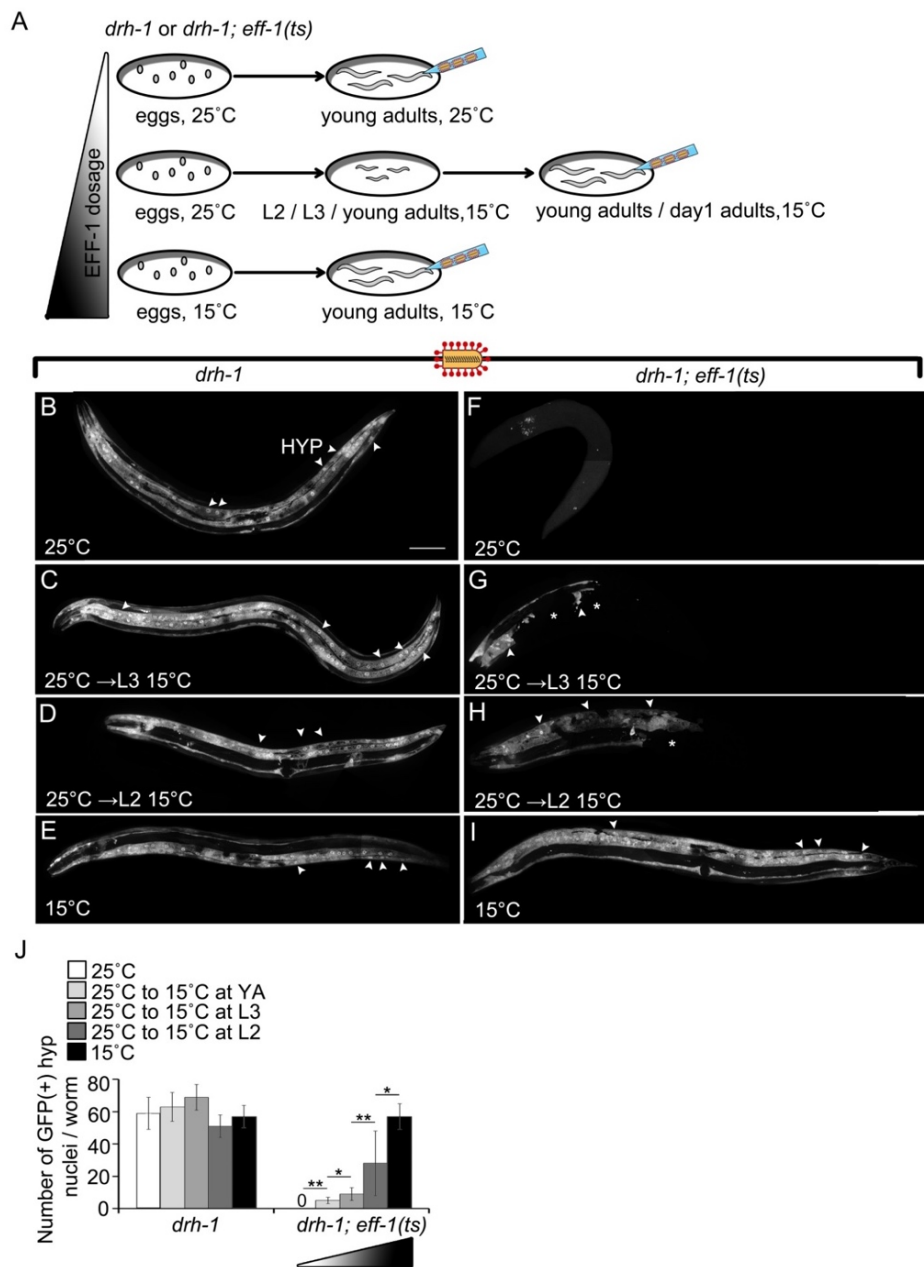
193 EFF-1 mediates formation of hyp7 syncytium by fusing 139 hypodermal cells during
194 embryonic and larval development and *eff-1(-)* worms have mononucleated hypodermal cells
195 instead of this syncytium [14]. To exclude the possibility that hypodermal cells of *eff-1(-)* worms
196 have a reduced susceptibility to viral infection, we injected wt, *aff-1(-)* and *eff-1(-)* worms with
197 VSV Δ G-G. We found that wild-type worms injected with VSV Δ G-G, presented a full body-
198 length covered by infected hypodermal cells along with muscles and excretory cells (Figures

199 2E-2F), while *eff-1(-)* worms had only few GFP(+) hyp nuclei localized closer to the injection
200 region- the head, as well as muscles and excretory cells (Figure 2G). *aff-1(-)* worms had
201 hypodermal infection similar to wt animals (Figure 2H). Yet, the fraction of worms injected with
202 VSV Δ G-G that showed GFP(+) muscle and hypodermal cells was not significantly different in
203 wt, *eff-1(-)* and *aff-1(-)* backgrounds (Figure 2I). Finally, we observed a small fraction of worms
204 with GFP(+) glia/neurons in the head with no significant difference between all tested
205 backgrounds (Figures S1A-S1C). Thus, hypodermal cells are not compromised for VSV Δ G-G
206 infection even in the absence of EFF-1. In addition, VSV Δ G-G with its unilateral viral fusogen
207 infects different cells including muscles, hypodermis, excretory cell, and glia/neurons in an
208 EFF-1- and AFF-1-independent manner. This is in contrast to VSV Δ G-AFF-1 that uses a
209 bilateral mechanism and infects cells that normally express EFF-1 or AFF-1 in adult animals.
210

211 **VSV Δ G-AFF-1-mediated infection depends on *eff-1* activity in target cells**

212 We have shown VSV Δ G-AFF-1 fails to infect hypodermal cells in *eff-1(-)* animals (Figure 2).
213 To study whether conditional induction of *eff-1* can trigger infection by VSV Δ G-AFF-1 we
214 varied the dosage of *eff-1* using temperature shifts in a conditional temperature sensitive (*ts*)
215 mutant [14,25]. In *eff-1(hy21ts)* animals grown at the permissive 15°C, most epidermal cells
216 fuse to form multinucleated hypodermis and vulva [14]. However, when *eff-1(hy21)* animals
217 are grown at the restrictive 25°C, epidermal cells fail to fuse, producing phenotypes similar to
218 the null *eff-1(ok1021)* worms [16]. Therefore, *eff-1(ts)* worms can serve as a temperature
219 inducible system for modification of EFF-1 dosage *in vivo* [14,19,25]. To obtain varying
220 expression of EFF-1, we maintained *eff-1(ts)* animals at 15°C, 25°C or performed 25°C to
221 15°C downshifts at different developmental stages and then injected VSV Δ G-AFF-1 into
222 young adult worms (Figure 3A). Wildtype worms in all experimental conditions had GFP(+) hyp
223 nuclei spanning the worm's body from head to tail (Figures 3B-3E), while *eff-1(ts)* animals
224 maintained at the restrictive temperature and then shifted down from 25°C to 15°C, had a
225 gradual increase in infected GFP(+) hyp cells (Figures 3F-3I). In wt worms maintained in all
226 conditions, the number of GFP(+) hypodermal nuclei was about 69, consistent with the number
227 of hyp7 nuclei found on one side of the animals body [39,40]. In contrast, for *eff-1(ts)*, the
228 longer time the worms developed at the permissive temperature (15°C), the more GFP(+) hyp
229 nuclei were scored (Figure 3J). Thus, hypodermal infection by VSV Δ G-AFF-1 increases with
230 conditional induction of *eff-1*.

231



232

233

234 **Figure 3. Infection by VSVΔG-AFF-1 increases with induction of EFF-1 function**

235 **(A)** Wt or temperature sensitive *eff-1(ts)* animals were maintained at permissive (15°C) or
 236 restrictive (25°C) temperatures for different times during their development. Each row
 237 represents a different experimental condition. Blue needles indicate virus injection. L2, Larval
 238 stage 2; L3, Larval stage 3.

239 **(B-I)** Confocal Z-stack projections of worms infected with VSVΔG-AFF-1. *drh-1* or *drh-1*;*eff-1(ts)*
 240 animals were injected with VSVΔG-AFF-1 (7-125 IU) as adults, and imaged 48 h later.
 241 Arrowheads, HYP nuclei. Asterisks, patches of GFP(-) surrounded by GFP(+) hypodermal
 242 cells. Scale bar, 100 μm.

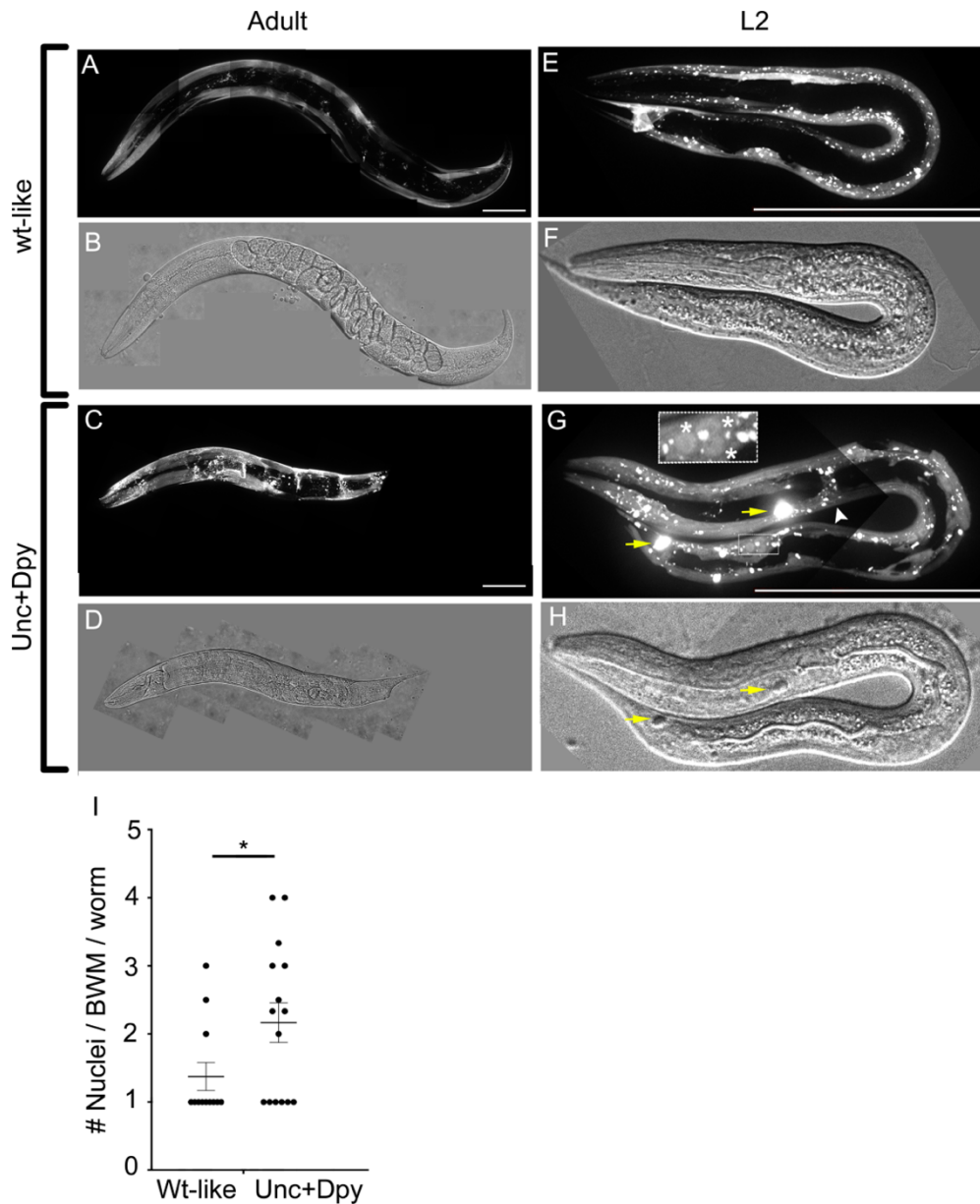
243 **(J)** Quantitation of experiments. Error bars average ± SEM. Student t-test. **p*<0.05, ***p*<0.001.
 244 *n*=3-10 worms per condition. Triangle, dosage of *eff-1*.

245

246 **Ectopic EFF-1 expression in BWMs produces dumpy and uncoordinated worms**

247 We demonstrated that VSV Δ G-AFF-1 specifically infects EFF-1 expressing cells and rarely
248 infects muscle cells. In addition, EFF-1-dependent infection could be induced by EFF-1(ts)
249 conditional expression in target cells (Figure 3). Hence, we propose that ectopic EFF-1
250 expression in host cells, such as body wall muscles (BWMs), could retarget VSV Δ G-AFF-1
251 into these cells. *C. elegans* has 95 mononucleated rhomboid BWMs, arranged as staggered
252 pairs and bundled in four quadrants that run along the worm's body [41–43]. While most
253 vertebrates and invertebrates have syncytial striated muscles composed of long
254 multinucleated myofibers, in *C. elegans* and other nematodes the BWMs have not been
255 described to fuse and do not form multinucleated myofibers
256 (<https://www.wormatlas.org/hermaphrodite/muscleintro/MusIntroframeset.html>). Therefore,
257 we hypothesized that EFF-1 expression in BWMs can induce their ectopic fusion, alter their
258 structure, produce muscle-related phenotypes and increase VSV Δ G-AFF-1 infection of
259 BWMs. First, we monitored animals with an extra-chromosomal array containing genomic
260 EFF-1 under the muscle-specific *myosin-3* promoter (*myo-3p::EFF-1*), together with *myo-*
261 *3p::mCherry* which labels cytoplasm and nuclei in BWMs, enteric, gonadal and vulval muscles.
262 Most adult animals, including *myo-3p::mCherry* (+) and (-), were wild-type-like (Table S1,
263 Figures 4A-4B and Movie S1), while about 10% of animals were both *myo-3p::mCherry*(+)
264 and had an uncoordinated and dumpy (Unc+Dpy) phenotype (Table S1, Figures 4C-4D and
265 Movie S1). Most *myo-3p::mCherry*(+), Unc+Dpy animals arrested during larval development
266 (Table S1) with additional phenotypes including bridged BWMs behind the terminal bulb (0/12
267 wt-like worms and 9/15 Unc+Dpy worms), mCherry-labeled aggregates that could be
268 observed with DIC (0/12 wt-like worms and 9/15 Unc+Dpy worms) (Figures 4E-4H), *myo-*
269 *3p::mCherry*(+) BWMs with clustered nuclei (3/12 wt-like worms and 9/15 Unc+Dpy worms)
270 and elevated number of nuclei/ BWM cell (1.4 ± 0.2 and 2.2 ± 0.3 nuclei/BWM cell in wt-like vs
271 Unc+Dpy larvae respectively) (Figures 4E-4I). While ectopic *hsp::EFF-1* expression causes
272 embryonic lethality [14,19,25], we found no difference between the fractions of *myo-*
273 *3p::mCherry* (+) and (-) unhatched eggs (Table S1). Thus, EFF-1 expression in BWMs
274 accounts for the Unc+Dpy phenotypes, larval arrest and clustered BWM nuclei, which
275 supports the hypothesis that ectopic EFF-1 induces BWMs multinucleation by cell-cell fusion
276 causing behavioral and morphological phenotypes.

277



278

279 **Figure 4. EFF-1 expression in BWMs produces Unc+Dpy worms with multinucleated**
 280 **cells**

281 (A-H) Images of fluorescent Z-stack projections and respective DIC of animals with
 282 extrachromosomal *myo-3p::EFF-1*, *myo-3p::mCherry*. (G) White arrowhead, bridge formed
 283 between 2 BWMs from opposing quadrants. Yellow arrows, *myo-3p::mCherry* accumulations
 284 also in DIC (H). Asterisks, clustered nuclei within one BWM. Scale bars, 100 μ m.

285 (I) Number of nuclei per *myo-3p::mCherry* (+) BWM cell in L2s. wt-like (n=12); Unc+Dpy
 286 (n=15). Each dot represents the average number of nuclei/BWM cell/worm. Total average \pm
 287 SEM for each phenotype. Two tailed Student's t-test $p < 0.05^*$.

288 See also Table S1.

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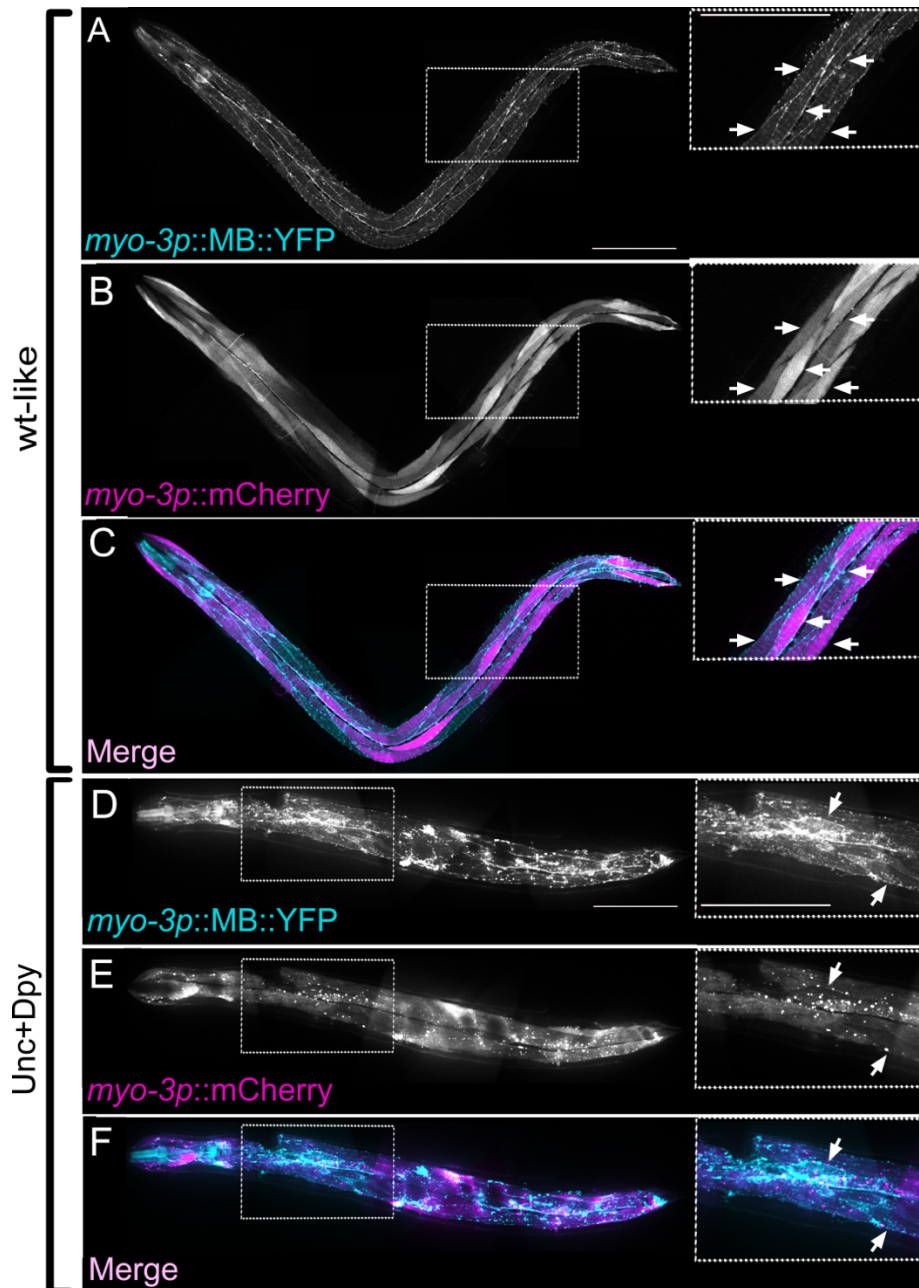
290 **EFF-1 expression in BWMs induces their fusion and retargets VSV Δ G-AFF-1 to muscles**

291 To determine whether EFF-1 expression in BWMs can induce their fusion, we imaged BWMs
 292 expressing a membrane-bound YFP (MB::YFP), *myo-3p::EFF-1* and *myo-3p::mCherry*, (see
 293 materials and methods). BWMs of wt-like worms have a normal spindle-shaped morphology

294 with cellular membranes between adjacent BWMs showing different levels of cytoplasmic
295 *myo-3p::mCherry* (as *myo-3p::mCherry* is an extrachromosomal array) (Figures 5A-5C). In
296 contrast, *Unc+Dpy* worms had disordered muscle fiber structure, the membranes surrounding
297 the BWMs were indistinguishable and the BWMs showed an evenly distributed cytoplasmic
298 *myo-3p::mCherry* indicating fusion and content mixing between these cells during
299 development (Figures 5D-5F). Thus, EFF-1 expression in BWMs induces disappearance of
300 cellular membranes and cytoplasmic merger, suggesting that EFF-1-mediated cell-cell fusion
301 alters BWM structure, likely resulting in *Unc+Dpy* and larval arrest phenotypes.

302 We showed that VSV Δ G-AFF-1 infects EFF-1-expressing hypodermal cells in >80%
303 of the infected worms but infects muscle cells that do not express EFF-1 in about 20% of the
304 infected animals (Figure 2D). Moreover, EFF-1 expression in BWMs induced their fusion, and
305 generated *Unc+Dpy* worms (Table S1, Movie S1 and Figures 4A-4D and 5A-5F). Taken that
306 EFF-1 and AFF-1 are bilateral fusogens and VSVG is a unilateral fusogen, we hypothesized
307 that ectopic expression of EFF-1 in BWMs would increase their infection by VSV Δ G-AFF-1
308 but not by VSV Δ G-G. Therefore, we injected either VSV Δ G-AFF-1 or VSV Δ G-G into the
309 pseudocoelom of worms ectopically expressing EFF-1 and mCherry in BWMs and imaged
310 them. For VSV Δ G-AFF-1 but not for VSV Δ G-G, the fraction of infected BWMs was significantly
311 higher in *Unc+Dpy* worms (expressing the *myo-3p::mCherry* marker) compared to wt-like
312 (expressing the *myo-3p::mCherry* marker) and wt animals (Figure 6). Thus, EFF-1 expression
313 in BWMs retargets VSV Δ G-AFF-1 into muscles but does not affect the natural VSV Δ G-G
314 tropism for BWMs.

315 We showed that VSV Δ G-AFF-1 infects neuron and glia cells in the head (Figures S1D-
316 S1F). In order to assess the infection of other fusogen-expressing neurons, we chose the
317 arborized neuron pair PVD, previously shown to express EFF-1, which spans the length of the
318 worm [25], VSV Δ G-AFF-1-infected PVD cells were not observed. Nevertheless, we found 3
319 wt animals (out of >100 infected worms), where PVD cells were infected with VSV Δ G-G
320 following injections with high dosage of pseudotyped viruses (Movie S2). Moreover, we tested
321 whether ectopic AFF-1 expression in PVD or ectopic EFF-1 expression in 12 mechanosensory
322 and chemosensory neurons in *eff-1(ts)* background at the restrictive 25°C, could induce
323 VSV Δ G-AFF-1 infection in these cells. None of these treatments produced VSV Δ G-AFF-1
324 infection in the neurons ectopically expressing EFF-1 or AFF-1 (Figure S3 and Table S2).
325 There are several possibilities to explain why VSV Δ G-AFF-1 did not infect these neurons; (i)
326 relatively low titer of VSV Δ G-AFF-1 preps, (ii) a physical barrier such as hypodermis, sheath
327 cells or glia cells that surround neurons or (iii) diffusion of the GFP signal in long neuronal
328 processes.



329

330

331 **Figure 5. EFF-1 expression in BWMs induces their fusion**

332 (A-C) Confocal images of wt-like adult worms with membrane bound (MB) *myo-3p::MB::YFP*

333 (cyan) and extrachromosomal array containing *myo-3p::EFF-1*, *myo-3p::mCherry* (magenta).

334 (D-F) Confocal images of Unc+Dpy [*myo-3p::MB::YFP* (cyan); *myo-3p::EFF-1*, *myo-*

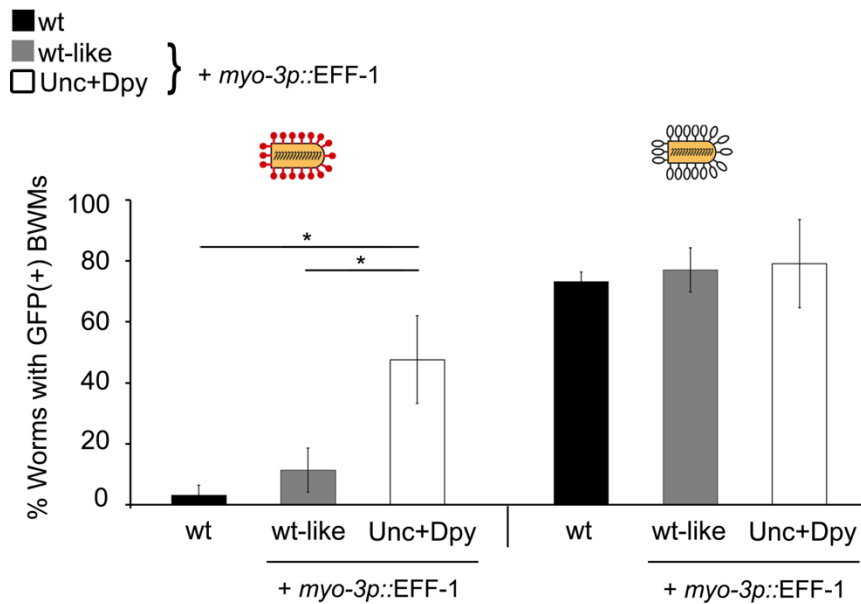
335 *3p::mCherry*]. Arrows, unfused BWMs with MB (cyan). Note only two unfused BWMs, all the

336 others appear fused with no MB separating them. Insets correspond to white-dotted area.

337 Scale bars 100 μ m.

338 See also Movie S1.

339



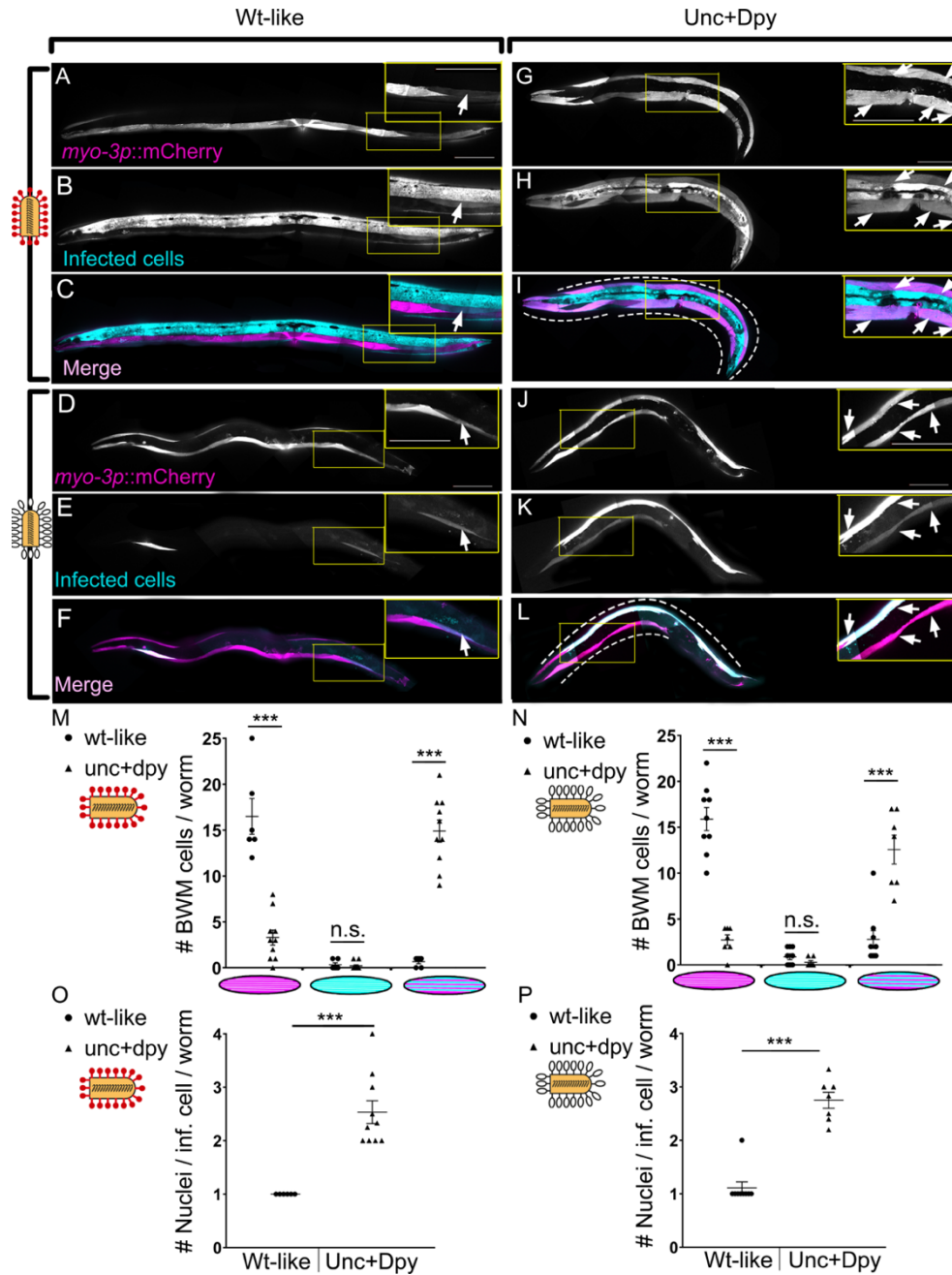
340

341 **Figure 6. Retargeting of VSVΔG-AFF-1 to body wall muscle cells**

342 Wild-type worms and animals with extrachromosomal array containing *myo-3p::EFF-1* and
343 *myo-3p::mCherry* were injected with VSVΔG-AFF-1 (35-63 IU, red pins; n=39 wt, n=50 wt-like
344 and n=27 Unc+Dpy worms) or VSVΔG-G (3×10^5 IU, white pins; n=30 wt-like and 14 Unc+Dpy)
345 respectively. Wt worms injected with VSVΔG-G (2300-4700 IU, n=56) were taken from figure
346 2I. Animals were analysed by SDC microscopy. Data represents average percentage of
347 worms with GFP(+) BWMs \pm SEM. Student's t-test: *p<0.05.
348

349 **EFF-1 expression in fused BWMs enables VSVΔG-AFF-1 and VSVΔG-G spreading**

350 Based on the aberrant muscle fusion phenotype seen in Unc+Dpy worms following EFF-1
351 expression in BWMs, we hypothesized that viral spreading through such novel BWM syncytia
352 would be enhanced. To test this hypothesis, we injected VSVΔG-AFF-1 or VSVΔG-G into wt-
353 like or Unc+Dpy worms and quantified the number of BWM cells that were either (i) mCherry(+)
354 only (expressing EFF-1 and *myo-3p::mCherry*(+)), (ii) GFP(+) only (infected but not
355 expressing EFF-1) or (iii) mCherry(+) and GFP(+) expression in the same cells (infected and
356 expressing EFF-1). We found that for both VSVΔG-AFF-1 and VSVΔG-G, wt-like worms had
357 individual GFP(+) BWM cells, that were mostly mCherry (-) (Figures 7A-7F, arrows), while
358 Unc+Dpy animals had continuous GFP(+) BWMs that overlapped with mCherry along their
359 body length (Figures 7G-7L, dashed lines and arrows). Compared to wt-like animals, the
360 Unc+Dpy worms had about five-times fewer non-infected mCherry(+) only BWMs, a similar
361 number of infected-GFP(+) only cells and about fifteen-fold more infected BWM cells that also
362 express mCherry (Figures 7M-7N). Lastly, Unc+Dpy worms had ~2.5 nuclei per VSVΔG-AFF-
363 1- or VSVΔG-G- infected BWMs, compared to wt-like animals with 1 nucleus per BWM cell
364 (Figures 7O-7P). Therefore, EFF-1 expression in BWMs transforms mononucleated BWMs
365 into syncytial muscle fibres enabling viral spreading within the fused cells and increasing the
366 number of both VSVΔG-AFF-1- and VSVΔG-G-infected BWM cells.



367

368 **Figure 7. EFF-1 expression in BWM enables VSV Δ G-AFF-1 and VSV Δ G-G spreading**
 369 **along fused muscles**

370 (A-L) Z-stack projections of wt-like (A-F) and Unc+Dpy animals (G-L) expressing *myo-*
 371 *3p::EFF-1* and *myo-3p::mCherry* infected with VSV Δ G-AFF-1 (35-63 IU red pins) or VSV Δ G-
 372 G (3×10^5 IU; white pins). Insets and their corresponding images (yellow frames). Arrows,
 373 individual infected (cyan) BWMs. Dashed lines outline grouped BWMs that express *myo-*
 374 *3p::mCherry* and *myo-3p::EFF-1* (magenta) and infected with virus (cyan) showing spreading
 375 of GFP. Scale bars, 100 μ m. (M-N) Number of BWM cells/worm expressing EFF-1 (magenta
 376 cell), infected (cyan cell) or expressing EFF-1 and infected (magenta and cyan). wt-like
 377 (circles) and Unc+Dpy (triangles). Each point represents a single worm. (O-P) Quantitation of
 378 multinucleation of infected BWMs. Each dot represents an average number of nuclei/ GFP(+)
 379 BWM, calculated from 1-6 multinucleated BWMs of a single worm. (M and O) wt-like n=6 and
 380 Unc+Dpy n=10 animals. (N and P) wt-like n=9 and Unc+Dpy n=7 animals.
 381 Black horizontal lines, average \pm SEM. Student's t-test, *** $p < 0.0001$; n.s., not-significant.

382 **Discussion**

383 **Utilizing VSV Δ G-AFF-1 to study muscle fusion and muscle fusogens**

384 Myogenesis of striated muscle involves the formation of multinucleated myofibers by
385 myoblast-myoblast fusion. Myoblast fusion is also essential for muscle tissue patterning,
386 maintenance and damage repair [62,63]. In contrast, *C. elegans* BWMs do not fuse and exist
387 as mononucleated cells, contracting in synchrony that mediate gait [64]. We used VSV Δ G-
388 AFF-1 to label EFF-1-expressing BWMs and found that EFF-1-mediated BWMs fusion results
389 in loss of coordination, although we did not determine whether the observed locomotory
390 defects result from aberrant joining of separately-innervated muscles or from a loss of
391 contractility. These results demonstrate that a locomotory circuit may be optimally adjusted for
392 both separate (as in *C. elegans*) and syncytial (as in vertebrate) striated muscles. Indeed,
393 failure of myoblast fusion in vertebrates may have equally severe phenotypes [47,48]. Despite
394 the importance of muscle fusion, the studies of muscle fusion proteins are just emerging [55–
395 58]. We suggest that VSV Δ G-AFF-1 can be used in screens to identify additional muscle
396 fusogens in invertebrates (e.g. *Drosophila*). Alternatively, viruses coated with the bilateral
397 Myomaker, can be a tool to screen for muscle-expressed Myomaker-interacting proteins in
398 vertebrates.

399

400 **VSV Δ G-AFF-1 infects, and can be retargeted to muscles expressing EFF-1 or AFF-1**

401 Previously described viral-vector-based systems used viral unilateral fusogens to target
402 desired cells. For instance, fusogens that are either native, modified or from different viral
403 origin, change the specificity of the viruses targets and facilitate the delivery process [1–3]. In
404 addition, a potent targeting of cancer cell lines was achieved with adenovirus (non-enveloped
405 virus) armed with an adaptor composed from 2 designed ankyrin repeat proteins (DARPs),
406 one binding to the viral fiber knob and the second binding to cancer-specific markers [44]. Our
407 work presents a new approach to retarget viral vectors into muscle cells, based on viruses
408 coated with a nematode bilateral fusogen, namely AFF-1 and expression of bilateral fusogens
409 such as EFF-1/AFF-1 in desired host cells. VSV Δ G-AFF-1 has a high degree of specificity
410 towards cells that express EFF-1/AFF-1 and can be redirected by expressing them on a
411 certain cell (e.g. BWMs). Moreover, VSV Δ G-AFF-1-mediated infection can be manipulated
412 quantitatively by changing the amount of a partner fusogen expressed on the target cells. VSV
413 and VSV-pseudotypes successfully infect invertebrate and vertebrate cell-lines and model
414 organisms [1,2,35–37]. Although EFF-1 and AFF-1 are not present, and have no known
415 homologs in vertebrates, VSV Δ G-AFF-1 infects mammalian BHK cells expressing AFF-1/EFF-
416 1 [6]. It will be interesting to test if VSV Δ G-AFF-1 can specifically target cells ectopically
417 expressing AFF-1/EFF-1 in a vertebrate model. Similarly, pseudotyped VSV Δ G-SARS-CoV-

418 2-S-glycoprotein [4] could be targeted to humanized worms or mice expressing hACE2 to
419 study entry mechanisms, pathogenesis and potential treatments for viral infections, including
420 COVID19 in model organisms.

421

422 **Additional applications for VSV Δ G-AFF-1**

423 In the recent decade, extracellular vesicles (EVs) emerged as intercellular communication
424 carriers, containing variable cargoes (e.g lipids, DNA, RNA, toxins and proteins) and involved
425 in different biological processes including cell-cell interactions, cancer, tissue and neuronal
426 regeneration. EVs can serve as biomarkers and are potent vehicles for gene and drug delivery
427 [26,45,46]. However, work with EVs-based vectors has to tackle the issues of specific
428 targeting, loading of desired cargo, presence of undesired/non-specific content and
429 quantification of produced EVs. Our system consisting of enveloped pseudotyped virus coated
430 with bilateral fusogens and host muscle cells expressing bilateral fusogens, demonstrates cell-
431 specific targeting. Moreover, VSV Δ G-AFF-1 has the benefits of viral-based vectors, namely
432 loading a specific and tailored cargo including fluorescent markers such as GFP-coding RNA
433 sequence, simple vector quantification and cargo amplification within the host cell. Therefore,
434 VSV Δ G-AFF-1 may serve as a paradigm for EVs-mediated transport. It can be utilized to
435 explore how EVs deliver their cargoes and to study whether and how EVs are involved in
436 different biological processes.

437 Exoplasmic membrane fusion is a fundamental biological process involved in
438 myogenesis, fertilization, vulva development, bone formation and resorption, hypodermis
439 morphogenesis, placentation, tubulogenesis, neuronal regeneration and viral infection
440 [28,31,47,48]. Despite the importance of these processes, very few cell-cell fusogens have
441 been identified and characterized so far. These include: (i) Syncytins involved in placenta
442 formation [49–51]; (ii) Fusexins including EFF-1 and AFF-1 from *C. elegans* [14,17–22,31] and
443 HAP2/GCS1 which mediates gamete fusion in protists and flowering plants [38,52–54] and
444 (iii) Myomaker (TMEM8c) and myomerger (myomixer/minion) proteins that fuse muscle cells
445 in vertebrates [55–58]. We demonstrate that VSV Δ G-AFF-1 injected into *C. elegans*
446 encounters and infects different EFF-1/AFF-1-expressing cells. Ectopic expression of EFF-1
447 in BWMs induces these mononucleated muscles to form syncytial myofibers, however, these
448 multinucleated muscles are pathological and the worms become deformed and paralyzed.
449 Moreover, bilateral heterotypic fusion has been described between EFF-1 and AFF-1 and also
450 between EFF-1 and sperm HAP2/GCS1 from *Arabidopsis* expressed on two populations of
451 BHKs [38]. Therefore, VSV Δ G-AFF-1/EFF-1 may be exploited in screens to identify new
452 bilateral fusogens or fusogen-interacting proteins in different organisms and to understand

453 how and why some myoblasts fuse (e.g. skeletal muscles) while others function optimally as
454 individual mononucleated cells (e.g. smooth muscles and BWMs).
455

456 **Materials and methods**

457 **Nematode strains**

458 Unless otherwise stated, all nematodes were maintained at 20°C, according to standard
459 protocols [65,66]. *drh-1(tm1329)* served as the wild type background. Mutations and strains
460 that were used in this study are listed in Table S3.

461

462 **DNA constructs**

463 The *myo-3p::EFF-1* plasmid was constructed by cloning the *myo-3* promoter region from *myo-*
464 *3p::mCherry* plasmid with Sal I (New England BioLabs Cat#R3138) and Nhe I (ThermoFisher
465 Cat# FD0974) and inserting it into the *hsp16-2::EFF-1* plasmid cut with the same enzymes to
466 replace the original heat shock promoter. To produce *eff-2 (hy51)* mutant worms with CRISPR,
467 the following DNA constructs were generated:

- 468 1. pBG115 plasmid encoding single guide targeting *eff-2* to insert *mNeonGreen* was
469 generated by cloning *eff-2* targeting sequence into the CAS9 plasmid pDD162 with BG84
470 and BG85 primers.
- 471 2. BG123 conversion oligonucleotide encoding wt *pha-1* fragment was PAGE purified.
- 472 3. PCR amplicon encoding for the mNeonGreen and *unc-54* 3'UTR flanked by homology
473 arms to the *eff-2* gene, generated by amplification of mNeonGreen from plasmid X, with
474 BG129 and BG130 primers containing a 50 bp homology arms to *eff-2*.

475 Sequences of primers and oligonucleotides mentioned above are found in Table S3.

476

477 **Transgenic animals**

478 For standard extrachromosomal transgenes, germline transformation was performed using
479 standard protocols [67]. Transgenic lines were kept as extrachromosomal arrays and
480 maintained by following the expression from either *myo-3p::mCherry*, *myo-2p::GFP*, or *mec-*
481 *4p::dsRed*, *odr-1p::dsRed* plasmids that were co-injected as transformation markers. *myo-*
482 *3p::mCherry* encodes mCherry expression in body wall muscle and vulva muscle cells. *myo-*
483 *2p::GFP* encodes GFP expression specifically expressed in pharyngeal muscles. *pmec-*
484 *4::dsRed* encodes dsRed expression in six touch receptor neurons. *odr-1p::dsRed* encodes
485 dsRed expression in two odor sensory neurons. Plasmids mentioned above and strains
486 containing these arrays are listed in Table S3. Transgenic lines with extrachromosomal arrays
487 were generated as follows:

- 488 • BP2126: *drh-1(tm1329);eff-1(hy21)* injected with 20ng/μl *pmyo-2::GFP* as
489 transformation marker and 0.1ng/μl *pdes-2::AFF-1*.
- 490 • BP2131-3: *drh-1(tm1329);eff-1(hy21)* injected 10ng/μl *pmec-4::dsRed* and 10ng/μl
491 *podr-1::dsRed* as transformation markers and to label sensory neurons, 1ng/μl *pmec-*
492 *4::EFF-1* and 1ng/μl *podr-1::EFF-1*.
- 493 • BP2137: *drh-1(tm1329)* injected 10ng/μl *myo-3p::mCherry* as transformation marker
494 and to label Body Wall Muscle cells, and 1ng/μl *myo-3p::EFF-1*.
- 495 *eff-2(hy51)* allele was generated by CRISPR/Cas9 [65,66] insertion of mNeonGreen into the
496 first exon of *eff-2* gene. *pha-1* was used as a conversion marker. *pha-1(e2123)* temperature
497 sensitive worms were injected with 50 ng/μl pBG115 targeting *eff-2* and containing CAS9, 50
498 ng/μl pJW1285 sgRNA plasmid against *pha-1* that contains the CAS9, 20 ng/μl BG123
499 conversion oligonucleotide encoding wt *pha-1* fragment and 20 ng/μl amplicon encoding for
500 the mNeonGreen and *unc-54* 3'UTR flanked by homology arms to the *eff-2* gene. Worms that
501 survived development at the restrictive temperature (25°C) were screened and sequenced for
502 mutations in the *eff-2* gene. *hy51* is the result of an imprecise partial inverted insertion of the
503 PCR fragment into the *eff-2* locus. One bp was deleted at position +31 and 488bp of the *unc-*
504 *54* 3'UTR and mNeonGreen gene were inserted into *eff-2* coding region. The product of the
505 inverted insertion is a non-functional fluorescent protein that causes a frame shift in *eff-2* and
506 terminates its translation at Amino Acid #18. The *eff-2* targeting sequence was added to
507 primers (see Table S3) and inserted to the linearized vector using restriction-free cloning
508 technique.

509

510 **Live imaging of worms**

511 For imaging of viral infection and BWM fusion in *C. elegans*, worms were analyzed by
512 Nomarski optics and fluorescence microscopy using Nikon eclipse Ti inverted microscope with
513 Yokogawa CSU-X1 spinning disk confocal (SDC) as described previously [26,27]. Briefly,
514 animals were anesthetized in 0.01-0.05% tetramisole in M9 solution for 20-30 min and then
515 picked with an eyelash attached to a toothpick and transferred to a 5 μl droplet of M9 solution
516 placed on 3% agar slide. Images were acquired with Metamorph software, when using the
517 spinning disk confocal. Z-stacks were taken with Plan Fluor 40x NA=1.3 or Apochromat 60x
518 NA=1.4 objectives. Excitation of GFP was achieved with 488 nm wavelength laser (2-8%
519 intensity, 100 ms exposure time). mCherry was excited with 561 nm wavelength (15-20%
520 intensity, 100 ms exposure time). ~0.5 μm z-steps were recorded with iXon3 EMCCD camera
521 (Andor). Multidimensional data were reconstructed as maximum intensity projections using Fiji
522 software (NIH ImageJ). For live imaging of worms with fused BWM phenotypes, plates with

523 worms were placed on the stage of Zeiss stereo Discovery V8 stereo microscope. Images and
524 movies were captured at x8 magnification with additional magnification from PlanApo S 2.3X
525 objective and a Hamamatsu ORCA-ER camera controlled by micromanager software
526 (<https://micro-manager.org>). Figures were prepared using Fiji, Adobe Photoshop CS5 and
527 GraphPad Prism 8.

528

529 **DNA transformation and viral infection by microinjection**

530 Microinjections were performed as described [7,34] with some modifications. Shortly, late L4
531 or young adult worms were placed into droplet of halocarbon oil on 3% agarose pads. Pulled
532 capillary needles were secured onto a Nikon DIAPHOT 300 microscope equipped with a
533 micromanipulator and regulated pressure source (Narishige). For DNA transformation,
534 needles were loaded with 0.8 μ l with DNA solution containing TE buffer, the target construct
535 DNA, a co-injection marker DNA and the required amount of with pKSI-1 (empty vector) DNA
536 to reach a total concentration of 100 ng/ μ l DNA. For experiments with viral infection, needles
537 were loaded 0.8 μ l DMEM (mock-infections) or pseudo viruses in DMEM+5%FBS. Infection
538 doses of VSV are based on VSV titration on BHK cells and used 10nL volume as a single
539 microinjection dose [7]. Agar pads with worms were placed on stage of Nikon DIAPHOT 300
540 microscope. Worms were observed under x40 objective. For DNA transformation, animals
541 were injected into the gonad and immediately placed into droplet of M9 buffer placed on NGM
542 plates seeded with OP50-1 *E. coli*. For experiments with viral infection worms were injected
543 into pseudocoelom -behind the terminal bulb of the pharynx and were immediately placed into
544 droplet of M9 buffer placed on NGM plates with 50 μ g/mL FUdR (Sigma), seeded with OP50-
545 1 *E. coli*. Unless otherwise stated, animals were maintained at 25°C until scoring of infection.

546

547 **Scoring viral infection**

548 Scoring viral infection using fluorescence assay was performed as described [7,34] with some
549 modifications. Briefly, 48-72 hours post injection worms were processed for live imaging as
550 described above. Animals that were unresponsive to prodding by a platinum wire worm pick
551 were considered dead and were removed from the experiment. Animals that crawled off the
552 plate or were lost during the experiment were censored. Worms with ≥ 1 GFP (+) cells were
553 considered as infected worms, while not injected or DMEM injected worms served as a
554 negative controls. For different experiments, either the number of GFP(+) cells/animals or the
555 type of infected tissues were observed and quantified.

556 **Temperature shift experiments**

557 Temperature sensitive *eff-1(hy21ts)* mutant worms were synchronized by hypochlorite
558 treatment of adult worms. The obtained eggs were left for overnight L1 hatching on NGM
559 plates without food at 20° C. L1 animals were then transferred to NGM plates with OP50
560 bacteria at 15° C or 25° C incubators until reaching the desired developmental stage. The
561 plates were either downshifted from 25° C to 15° C, or left at 25° C or at 15° C throughout the
562 experiment. The developmental stages were determined by analyzing gonadal size and
563 structure using Nomarski optics. Finally, worms were injected with VSV Δ G-AFF-1, maintained
564 and imaged as described above.

565

566 **Counting overlapping cells**

567 To test if EFF-1 expressing BWMs were infected by VSV Δ G-AFF-1, we utilized worms with an
568 extrachromosomal array expressing EFF-1 under the *myo-3* promoter and a plasmid
569 containing *myo-3p::mCherry*. We found that most mCherry (+) worms had a wt-like phenotype,
570 but a small fraction of these worms were Unc+Dpy. Importantly, all sibling mCherry(-) worms
571 were wt-like (Table S1). Hence, for the experiment we utilized a fluorescence
572 stereomicroscope and chose late L4/ young adult, wt-like and Unc+Dpy animals that had
573 continuous *myo-3p::mCherry*(+) BWMs (which could express EFF-1 and fuse with each other).
574 Next, these worms were injected with VSV Δ G-AFF-1 or VSV Δ G-G. 48-72h later, Z-stack
575 images of the worms were obtained with SDC microscope (as described in live imaging
576 section). Worms with ≥ 1 infected BWM were selected. For each worm, we counted the number
577 of BWMs that were mCherry(+) only (express EFF-1), GFP(+) only (infected by virus) and
578 overlapping- with both mCherry(+) and GFP(+) (express EFF-1 and infected).

579

580 **Counting number of nuclei per BWM**

581 To test if ectopic EFF-1 expression in BWMs produces multinucleated BWMs, wt-like and
582 Unc+Dpy animals that had continuous *myo-3p::mCherry*(+) BWMs (which could express EFF-
583 1 and fuse with each other) were imaged by SDC microscope as described above. In one set
584 of experiments we imaged L2 worms and counted number of nuclei per *myo-3p::mCherry* cell.
585 In a second set of experiments, we imaged adult worms with BWMs infected by either VSV Δ G-
586 AFF-1 or VSV Δ G-G and counted number of nuclei per infected (GFP(+)) BWM cell. For both
587 sets of experiments, in worms with at least one multinucleated BWM we monitored all
588 multinucleated BWMs and calculated the average number of nuclei/BWM for each worm.
589 Worms in which multinucleated cells were not observed, were considered as having 1
590 nucleus/BWM.

591

592 **Counting phenotypes in worms expressing EFF-1 in BWMs**

593 To find whether ectopic expression of EFF-1 in BWM cells produces any special phenotypes,
594 we used BP2137 *drh-1(tm1329)IV*; *hyEx375[myo-3p::EFF-1, myo-3p::mCherry, KS*
595 *bluescript]*. We first isolated 4 wt-like *myo-3p::mCherry* L4 hermaphrodite worms, one worm
596 per NGM plate+Op50. These worms were led to lay eggs and were transferred to a fresh
597 plate 1-2 times per day during 5 days. Progeny including eggs, larvae and adults were divided
598 into two groups, namely *myo-3p::mCherry(+)* and *myo-3p::mCherry(-)*. We counted the total
599 number of progeny with a certain phenotype (e.g. wt-like worms, adult Unc+Dpy worms, larval
600 arrested Unc+Dpy worms and unhatched eggs) for each of the 4 mothers. Finally, we
601 calculated the average number of worms with certain phenotype \pm SEM and the fraction \pm SE
602 of worms with a certain phenotype in each group.

603

604 **Cell culture and preparation of pseudoviruses**

605 Baby Hamster Kidney cells (BHK), BHK-21(ATCC) were cultured in Dulbecco's Modified
606 Eagles Medium (DMEM) and recombinant viruses were prepared as described [6] with some
607 modifications. Briefly, BHK cells were grown to 70% confluence on 10 cm plates and then
608 transfected using Fugene HD + OptiMEM at ratio 1:4 (Fugene:DNA), with plasmids encoding
609 pOA20 (pCAGGS::*aff-1::FLAG*, 2 μ g/ml final concentration) [6] or pOA28 (pCAGGS::*VSV-G*
610 *Indiana*, 1 μ g/ml final concentration)[33]. Following 24 h incubation at 37°C in 5% CO₂, cells
611 were infected with VSVG-complemented VSV Δ G recombinant virus (VSV Δ G-G) at a
612 multiplicity of infection (MOI) of 5, for 1 hour at 37°C in a 5% CO₂ incubator in serum free
613 DMEM. Virus infected cells were washed 3-6 times with PBS (+ Ca⁺⁺ & Mg⁺⁺) to remove
614 unabsorbed VSV Δ G-G virus. Following a 24 h incubation period at 37°C, the supernatant
615 containing the VSV Δ G-G, or VSV Δ G-AFF-1 pseudoviruses were harvested without scraping
616 the cells and centrifuged at 600 g for 10 min at 4°C to clear cell debris. VSV Δ G-AFF-1 Virions
617 were filtered through 0.22 μ m filter unit and then double concentrated. First, by pelleting at
618 100,000 g through a 20% sucrose cushion, and resuspension in 10%FBS DMEM, second, by
619 pelleting at 100,000 g through a 10% sucrose cushion, and final resuspension in 45 μ l DMEM.
620 Finally, VSV Δ G-AFF-1 was incubated with anti-VSV-G antibody mAb diluted 1:1000 to inhibit
621 infection due to residual presence of VSV-G. The effective blocking of VSV-G was confirmed
622 by titrating pseudoviruses in BHK cells and by injection of VSV Δ G-G incubated with anti-VSV-
623 G into worms (Figure S4).

624

625 **Titering VSV pseudotyped viruses**

626 Titering VSV pseudotyped viruses was performed as described [6] with some modifications.
627 Briefly, 5x10³ BHK cells were plated into each well of a 96 well tissue culture plate (NUNC,
628 cat# 167008). To determine the titer of VSV Δ G-AFF-1, BHK cells were initially transfected with

629 2 µg/ml pOA20 (pCAGGS::AFF-1::FLAG). Cells transfected with empty vector served as
630 control. Eight serial x2 dilutions of the virus were performed and added to cells. After 18-24
631 hours of incubation, GFP expressing cells were counted in at least three dilutions using x20
632 objective of Zeiss Axiovert 200M fluorescence microscope. Inoculation was performed in the
633 presence of anti-VSV-G antibody mAb diluted 1:1000 to inhibit infection due to residual
634 presence of VSV-G.

635

636 **Statistical tests**

637 The specific tests used are described in the figure captions and the results section. The graphs
638 show mean ± SEM unless noted otherwise. For each experiment at least two biological
639 replicates were performed and the number of animals per experiment is stated in the figure
640 legends.

641

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853

854 **Abbreviations**

855	AFF-1	Anchor cell Fusion Failure
856	BHK	Baby Hamster Kidney cells
857	BWM	Body Wall Muscle
858	Dpy	Dumpy phenotype
859	EC	Excretory Canal cell
860	EFF-1	Epithelial Fusion Failure 1
861	EFF-2	Epithelial Fusion Failure 2
862	FUSEXINS	FUSion proteins essential for sexual reproduction and Exoplasmic merger of
863		plasma membranes
864	HYP	Hypodermis (epidermis)
865	SM	Somatointestinal Muscle
866	ts	temperature sensitive
867	UM	Uterine Muscle
868	Unc	Uncoordinated phenotype
869	VSV	Vesicular Stomatitis Virus
870	VSV-G	VSV-glycoprotein G
871	VSVΔG	pseudoviruses in which the glycoprotein G gene was deleted
872	VSVΔG-AFF-1	pseudovirus coated with AFF-1
873	VSVΔG-G	pseudovirus coated with G glycoprotein

874

875 **Acknowledgements**

876 We thank Don Gammon for providing the *drh-1(-)* worms, Massimo Hilliard for providing *mec-*
877 *4p::mCherry*, *mec-4p::EFF-1*, *odr-1p::dsRed* and *odr-1p::EFF-1* plasmids and the
878 Caenorhabditis Genetics Center for nematode strains. We thank Andy Fire for suggesting
879 studying syncytial BWMs and their potential phenotypes. We also thank Sharon Inberg, Yael
880 Iosilevskii, Rosina Giordano-Santini, Dan Cassel and Sivan Korenblit for helpful discussions
881 and for critically reading the manuscript.

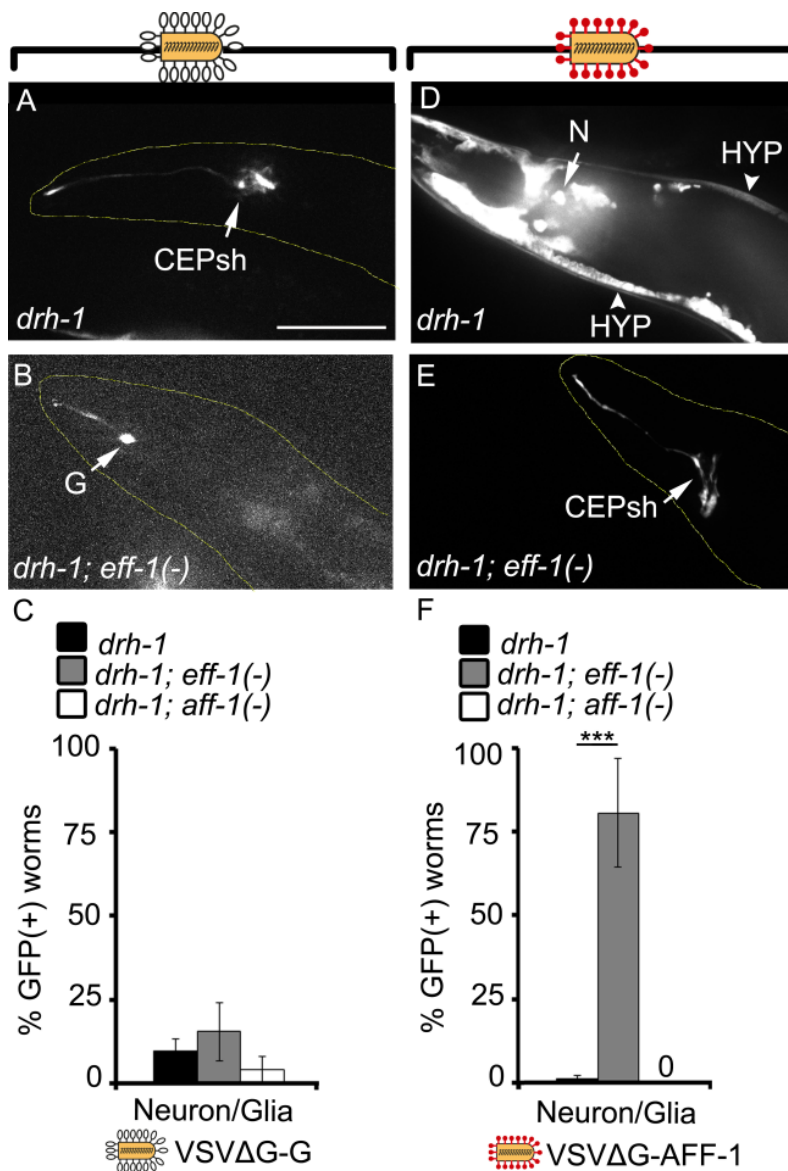
882

883 **Author contributions**

884 A.M. and B.P. conceived the project. A.M. performed all experiments unless otherwise
885 specified. X.L. generated *myo-3p::EFF-1* Plasmid and BP2171-3 nematode strains and
886 performed imaging of worms in Figure 5. E.M. improved viral preparation protocol. B.G.
887 generated EFF-2 CRISPR plasmid and O.K. produced *eff-2(hy51)* worms. A.M. and B.P.
888 analyzed data and wrote the manuscript, with input from all authors.

889

890 Online supplemental material



891

892

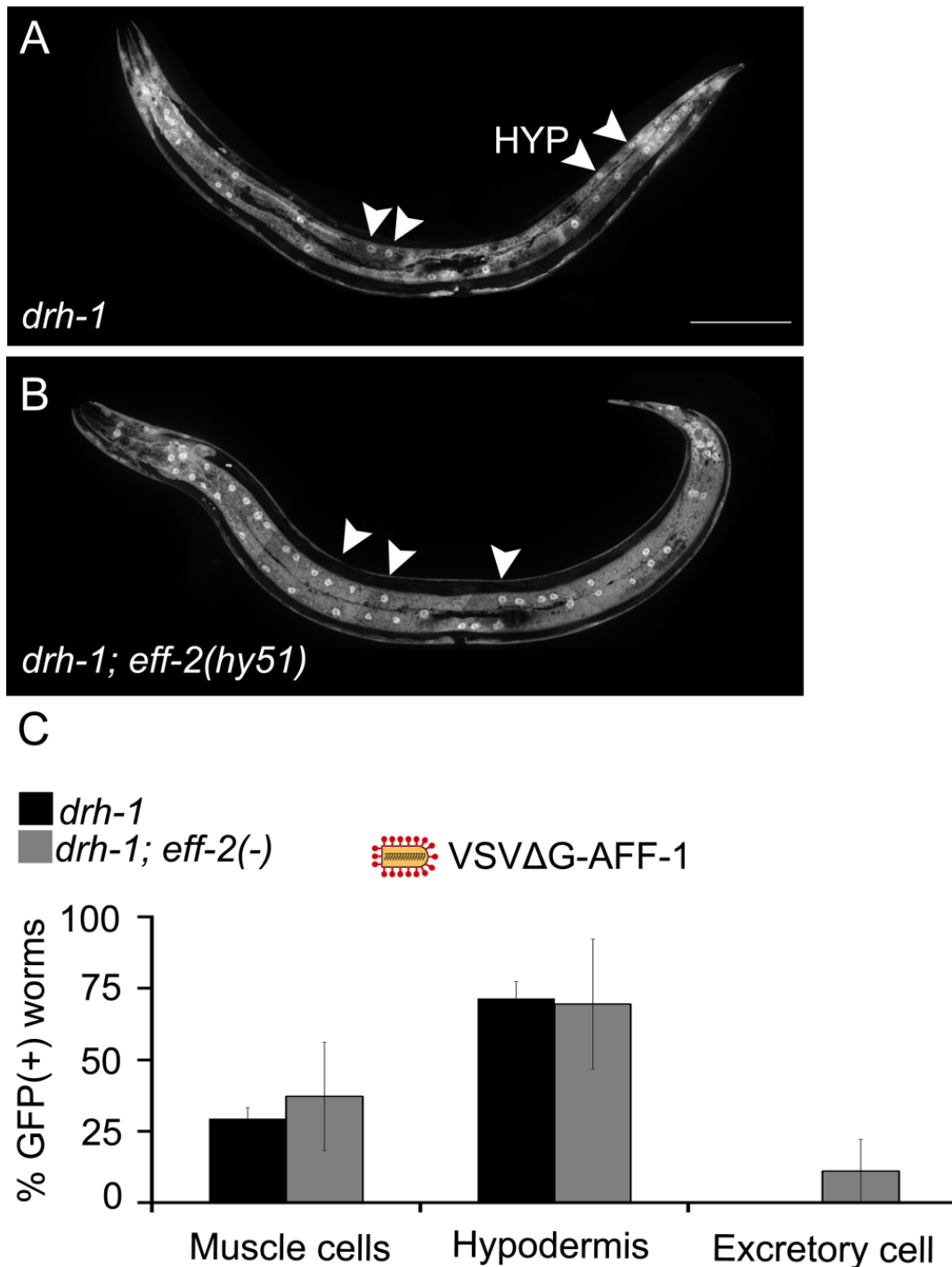
893 **Figure S1: VSVΔG-AFF-1 and VSVΔG-G infect glia and neuron cells**

894 **(A-B)** SDC microscope Z-stack projections of *drh-1* and *drh-1; eff-1(-)* worms injected in the
 895 pseudocoelom with 2300-4700 IU VSVΔG-G (white pins) and their heads imaged 48 h later.

896 **(C)** Fraction of GFP(+) worms infected with VSVΔG-G in neuron/glia cells in the specified
 897 background. Animals were injected with VSVΔG-G and analyzed as in (A-B).

898 **(D-E)** SDC microscope Z-stack projections of *C. elegans* cells infected with 33-240 IU VSVΔG-
 899 AFF-1 (red pins). *drh-1* or *drh-1; eff-1(-)* worms were injected in the pseudocoelom and
 900 imaged 48 h later.

901 **(F)** Fraction of GFP(+) worms infected with VSVΔG-AFF-1 in neuron/glia cells. Animals were
 902 analyzed as in (D-E). For (A-B) and (D-E) arrowheads point to hypodermal nuclei (HYP) and
 903 arrows point to indicated infected cell. CEPsh- Cephalic Sheath (glia), G- glia, N- neuron.
 904 Scale bar, 50 μm. In (A, B and E) yellow dashed lines outline worm's anterior body (head). In
 905 (C) and (F), bars represent average ± SEM. ***P<0.001 (Student's T- test). For VSVΔG-AFF-
 906 1: n=46,15 and 10 for *drh-1*,*drh-1;eff-1(-)* and *drh-1;aff-1(-)* respectively, and for VSVΔG-G:
 907 n= 56, 39 and 15 for *drh-1*, *drh-1;eff-1(-)* and *drh-1;aff-1(-)* respectively).



908

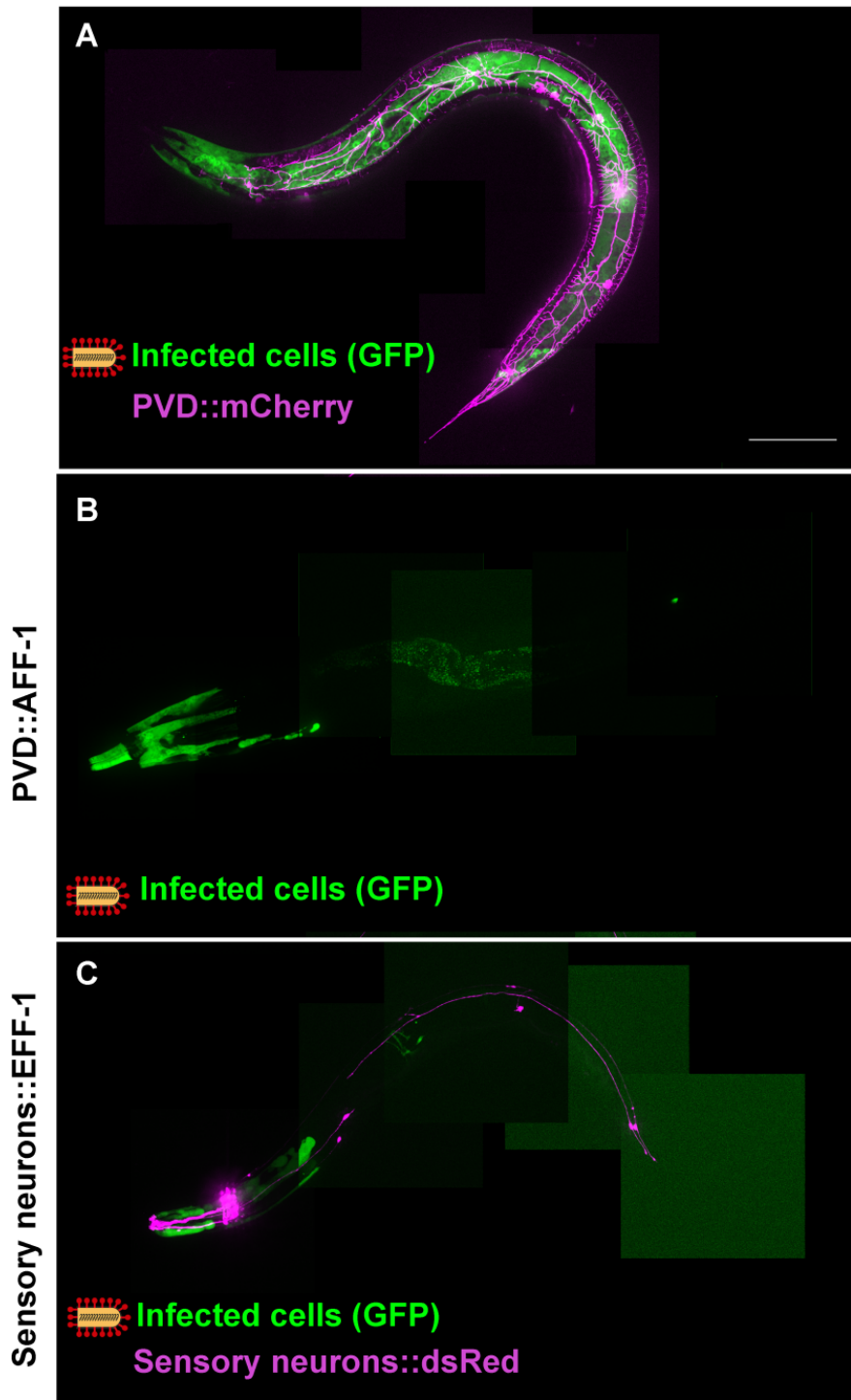
909

910 **Figure S2: VSVΔG-AFF-1 infects *eff-2(-)* worms**

911

912 **(A-B)** SDC microscope Z-stack projections of *drh-1* or *drh-1;eff-2(-)* worms injected with
913 VSVΔG-AFF-1 (35-67 IU) and imaged 48 h later.

914 **(C)** Fraction of GFP(+) worms infected with VSVΔG-AFF-1. Animals were injected with
915 VSVΔG-AFF-1 and analyzed as in (A-B). Arrowheads point to hypodermal (HYP) nuclei. Scale
916 bar, 100 μm. In (C) bars represent average ± SEM. n=23 and 20 for *drh-1* and *drh-1;eff-2(-)*
917 respectively. For all types of infected cells, there is no significant difference between % GFP(+)
918 worms from *drh-1* and *drh-1;eff-2 (-)* backgrounds with two-tailed Student's t-test (p<0.05).
919



920

921 **Figure S3. VSV Δ G-AFF-1 does not infect PVD and other sensory neurons ectopically**
922 **expressing AFF-1/EFF-1**

923 (A-C) SDC microscope Z-stack projections of animals infected with 82-103 IU VSV Δ G-AFF-1
924 (red pins). (For genotypes and quantitation see Table S2). Scale bar, 100 μ m.

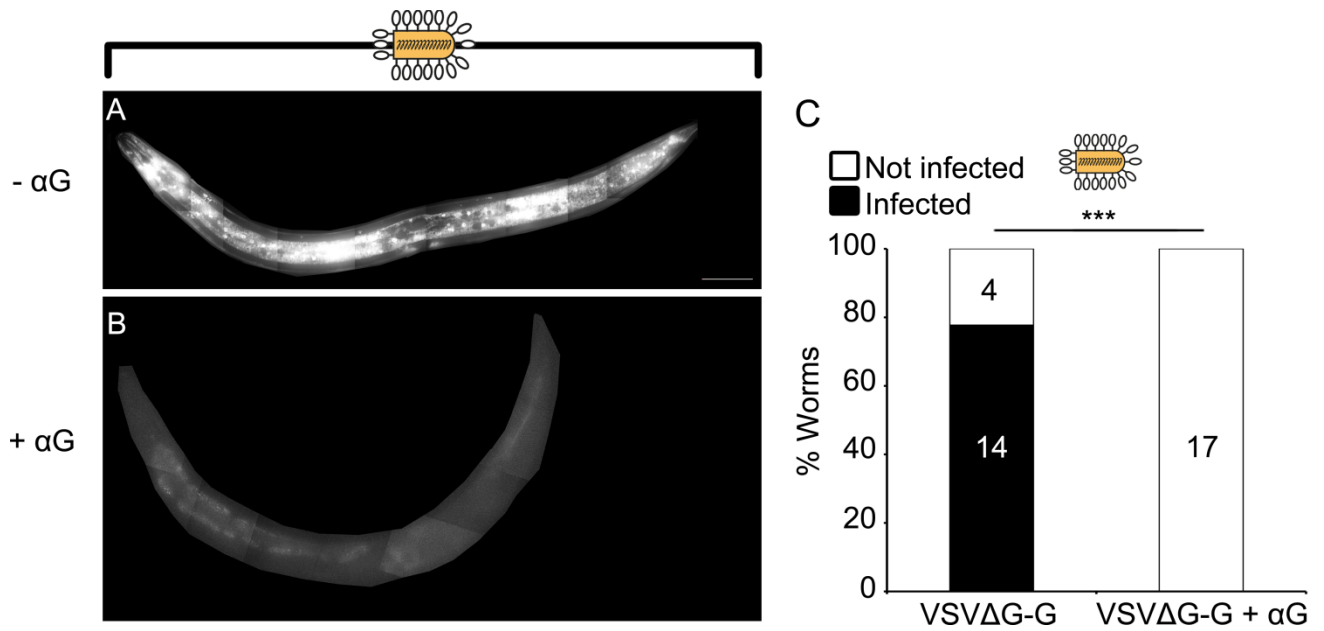
925 (A) Young adult expressing mCherry in PVD.

926 (B) *eff-1(ts)* adult expressing AFF-1 in PVD.

927 (C) *eff-1(ts)* adult expressing EFF-1 and dsRed in 12 sensory neurons.

928 See also Table S2 and Movie S2.

929



930

931

932 **Figure S4. Anti-VSV-G antibody blocks VSVΔG-G infection in living worms**

933 **(A-B)** SDC microscope Z-stack projections of *drh-1* worms injected with 4700 IU VSVΔG-G
934 that was either not preincubated (-αG), or preincubated with αVSV-G antibody (+αG) and
935 imaged 48 h later. Scale bar, 50 μm.

936 **(C)** Fraction of worms that are infected or not infected with VSVΔG-G or VSVΔG-G + αG.
937 Animals were treated as in (A-B). n=18 and 17 animals for VSVΔG-G and VSVΔG-G+αG
938 respectively. Fisher exact test. p***<0.001.
939

940 **Supplementary tables**

941 **Table S1. Phenotypes of worms with EFF-1 expressed in BWMs**

942

Genotype	<i>drh-1(-); hyEx375[myo-3p::mCherry, myo-3p::EFF-1]</i>		
Phenotype/ group	<i>myo-3p::mCherry(+)</i>	<i>myo-3p::mCherry(-)</i>	P-value
wt-like animals	110 ± 12 (81±3%)	60 ± 10 (87±6%)	0.39258
Adult Unc+Dpy	3 ± 1 (2±1%)	0 (0%)	0.05316
Larval-arrested Unc+Dpy	15 ± 2 (11±1%)	0 (0%)	0.00029***
Unhatched eggs	7 ± 2 (5±1%)	8 ± 3 (13±6%)	0.23501

943

944 Phenotype count of total progeny from 4 hermaphrodites with extrachromosomal array
945 containing *myo-3p::EFF-1* and *myo-3p::mCherry*. Data presented as average of the progenies
946 from 4 mothers ±SEM and the percentage out of total relevant group (e.g. *myo-3p::mCherry(+)*
947 or *myo-3p::mCherry(-)*). P-value calculated for certain phenotype fraction (%) in *myo-*
948 *3p::mCherry(+)* vs *myo-3p::mCherry(-)* groups. Two tailed Student's t-test. p<0.01***.

949

950 **Table S2. AFF-1/EFF-1 expression in sensory neurons does not induce their infection**
 951 **with VSV Δ G-AFF-1**
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Strain	Genotype	Description	total # of worms	# VSV Δ G-AFF-1-infected worms	# worms with GFP(+) cells overlapping with EFF-1/AFF-1 expressing cells
BP2126	<i>drh-1(tm1329); eff-1(hy21); hyEx373 [pmyo-2::GFP, pdes-2::AFF-1(pME4), KS]</i>	AFF-1 expressed in PVD (Oren-Suissa et al 2017) , in <i>eff-1(ts)</i> background	18	6	0
BP2131	<i>drh-1(tm1329); eff-1(hy21); hyEx374 [pmec-4::EFF-1, pmec-4::dsRed, podr-1::EFF-1, podr-1::dsRed, KS] line1</i>	EFF-1 expressed in 12 sensory neurons (kind gift from M. Hilliard lab), in <i>eff-1(ts)</i> background	28	15	0
BP2132	<i>drh-1(tm1329); eff-1(hy21); hyEx374 [pmec-4::EFF-1, pmec-4::dsRed, podr-1::EFF-1, podr-1::dsRed, KS] line2</i>	EFF-1 expressed in 12 sensory neurons, in <i>eff-1(ts)</i> background	22	7	0
BP2133	<i>drh-1(tm1329); eff-1(hy21); hyEx374 [pmec-4::EFF-1, pmec-4::dsRed, podr-1::EFF-1, podr-1::dsRed, KS] line3</i>	EFF-1 expressed in 12 sensory neurons, in <i>eff-1(ts)</i> background	35	13	0

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955 **Table S3. Reagents, strains, oligos, plasmids and software**

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Antibodies		
Anti-VSV-G [8G5F11] Antibody	Kerafast	Cat#8G5F-11
Bacterial and Virus Strains and tissue culture cells		
Recombinant Vesicular Stomatitis Virus VSVΔG	[68] Caenorhabditis Genetics Center	VSVΔG
<i>E. coli</i> : OP50-1		OP50-1
BHK-21(ATCC)	[69]	BHK-21(ATCC)
Chemicals, Peptides, and Recombinant Proteins		
FUdR	Sigma	Cat#F0503-1G
Tetramisole	Sigma	Cat#L9756-5G
Dulbecco's Modified Eagles Medium (DMEM) High glucose, Gibco™	Thermo Scientific	Cat#41965039
Fugene HD	Promega	Cat#E2311
OptiMEM	Gibco	Cat#31985070
Dulbecco's Phosphate Buffered Saline (DPBS) With Calcium and Magnesium	Biological Industries	Cat#02-020-1A
Certified Fetal Bovine Serum (FBS), Heat Inactivated	Biological Industries	Cat#04-121-1A
C. elegans strains		
<i>drh-1(tm1329) IV</i>	[7]	<i>drh-1(tm1329)</i>
<i>eff-1(ok1021)II</i>	[14]	BP347
<i>drh-1(tm1329) IV; eff-1(ok1021)II</i>	This paper	BP2122
Strain <i>eff-1(hy21)II</i>	[14, 19]	BP75
<i>drh-1(tm1329) IV; eff-1(hy21)II</i>	This paper	BP2123
<i>drh-1(tm1329) IV; aff-1(tm2214)II</i>	This paper	BP2124
<i>dziS53[F49H124p::mCherry] Unknown chromosome</i>	[70]	EB2110
<i>drh-1(tm1329) IV; dziS53[F49H124p::mCherry]</i>	This paper	BP2121
<i>trls10 [myo-3p::MB::YFP + myo-2p::YFP + ceh-23::HcRed + unc-25::DsRed + unc-129nsp::CFP]</i>	Caenorhabditis Genetics Center	RP1
<i>eff-2 (hy51) II; mcls46[dlg-1::RFP;unc-119(+)]III</i>	This paper	BP2104
<i>drh-1(tm1329) IV ; eff-2(hy51) II</i>	This paper	BP2125
<i>drh-1(tm1329)IV; hyEx375 [myo-3p::EFF-1, myo-3p::mCherry, KS bluescript]</i>	This paper	BP2137
<i>trls10 [myo-3p::MB::YFP + myo-2p::YFP + ceh-23::HcRed + unc-25::DsRed + unc-129nsp::CFP];hyEx375 [myo-3p::EFF-1, myo-3p::mCherry, KS bluescript]</i>	This paper	BP2171-3
<i>drh-1(tm1329) IV; eff-1(hy21) II; hyEx373 [myo-2p::GFP, des-2p::AFF-1, KS bluescript]</i>	This paper	BP2126
<i>drh-1(tm1329) IV; eff-1(hy21) II; hyEx374 [mec-4p::EFF-1, pmec-4::dsRed, podr-1::EFF-1, odr-1p::dsRed, KS bluescript]</i>	This paper	BP2131-3
Oligonucleotides		
Primer: Cloning <i>myo-3p</i> from <i>myo3p::mCherry</i> Plasmid with <i>Sall</i> restriction enzyme: ACGCGTCGAC AGTGATTATAGTCTGTTTTTC	This paper	LXH60
Primer: Cloning <i>myo-3p</i> from <i>myo3p::mCherry</i> Plasmid with <i>NheI</i> restriction enzyme: CTAGCTAGCCATTCTAGATGGATCTAGTG	This paper	LXH61
Oligonucleotide: Conversion oligonucleotide encoding wt <i>pha-1</i> repair fragment for CRISPR: <i>caaaatacgaatcgaagactcaaaaagagtatgctgtatgattacagatgttcaagttattcataaatcatgatag</i>	This paper	BG123
Primer: Addition of homology arms to <i>eff-2</i> around <i>mNeonGreen</i> for CRISPR: <i>ttcctgaagttcatcaaatcgaagctggcGcgtatttcagaagcATGGTTCCAAGGAGAAAGAGG</i>	This paper	BG129
Primer: Addition of homology arms to <i>eff-2</i> around <i>mNeonGreen</i> for CRISPR: <i>agattctaagtgtcagatagcttgatgatCtgCgagagaataactaGACTAGTAGGAAACAGTTATGTTTGG</i>	This paper	BG130
Primer: Production of sgRNA plasmid targeting <i>eff-2</i> at locus AGAAGCAGAATAAACTGGGGAGG for CRISPR: AGAAGCAGAATAAACTGGGGGTTTTAGAGCTAGAAATAGC	This paper	BG84
Primer: Production of sgRNA plasmid targeting <i>eff-2</i> at locus AGAAGCAGAATAAACTGGGGAGG for CRISPR: CCCAGTTTATTCTGCTTCTAAACATTTAGATTGCAATTC	This paper	BG85

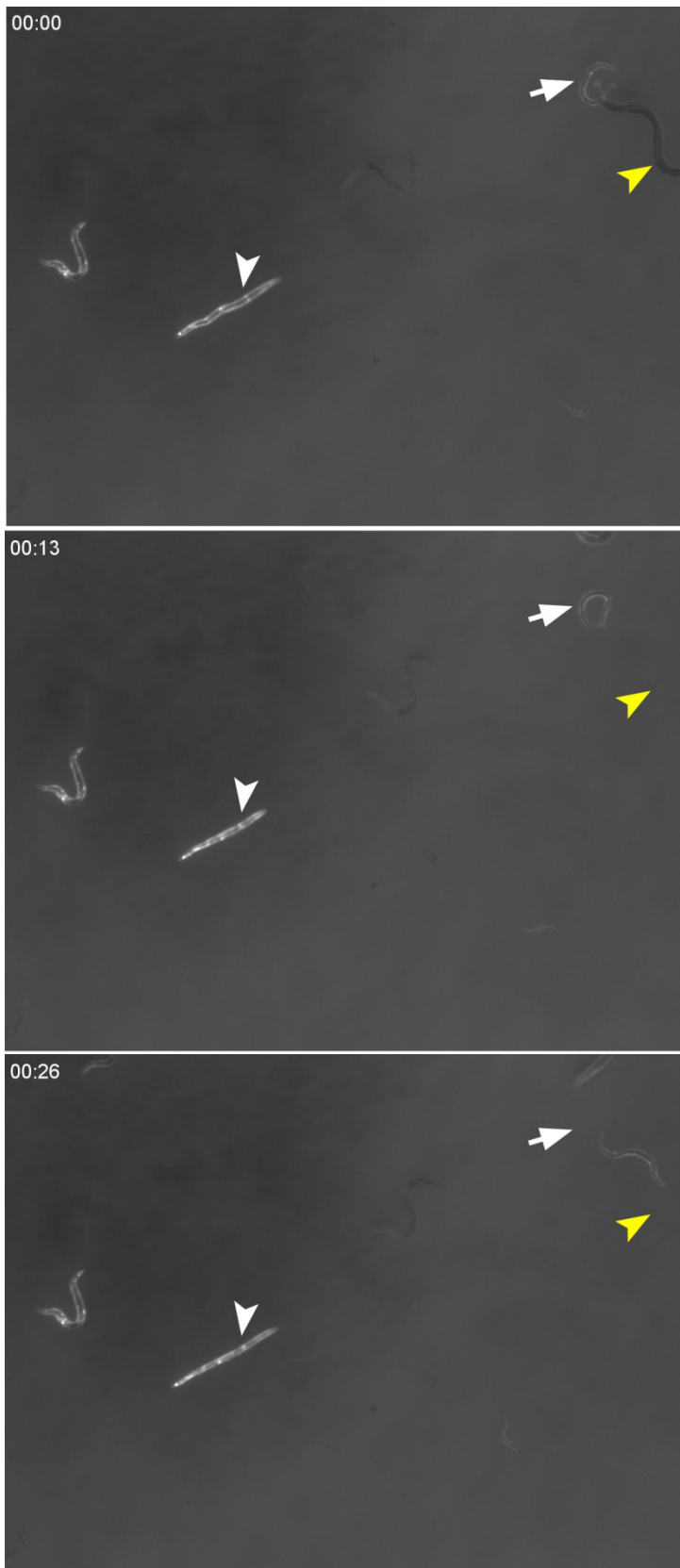
Recombinant DNA

Plasmid pCAGGS-Gind: <i>pCAGGS::VSV-G Indiana</i>	[68]	pCAGGS-Gind
Plasmid pOA20: <i>pCAGGS::aff-1::FLAG</i>	[6]	pOA20
Plasmid pKSI-1: pBluescript empty vector	[71]	Addgene plasmid #51725
Plasmid pXH20: <i>myo-3p::EFF-1</i>	This paper	pXH20
Plasmid pME1: <i>hsp16.2::EFF-1</i>	[19]	pME1
Plasmid pCFJ104: <i>myo-3p::mCherry</i>	[72]	Addgene plasmid #19328
Plasmid pBG115: <i>CeU6p::eff-2_sgRNA for CRISPR</i>	This paper	pBG115
Plasmid pDD162: <i>eft-3p::Cas9 + Empty sgRNA for CRISPR</i>	[73]	Addgene plasmid #47549
Plasmid pJW1285: <i>eft-3p::Cas-9-SV40 NLSNLS::tbb-2 3'UTR + CeU6p::pha-1_sgRNA for CRISPR</i>	[74]	Addgene plasmid #61252
Plasmid pNG1: <i>dlg-1p::Lifeact::mNeonGreen</i>	Kind gift of A. Hajnal	pNG1
PCR amplicon: <i>mNeonGreen flanked by homology arms to the eff-2 gene for CRISPR</i>	This paper	N/A
Plasmid : <i>odr-1p::dsRed</i>	[24]	odr-1p::dsRed
Plasmid: <i>odr-1p::EFF-1</i>	[24]	odr-1p::EFF-1
Plasmid: <i>mec-4p::EFF-1</i>	[24]	mec-4p::EFF-1
Plasmid: <i>mec-4p::mCherry</i>	[24]	pmec-4::mCherry

Software

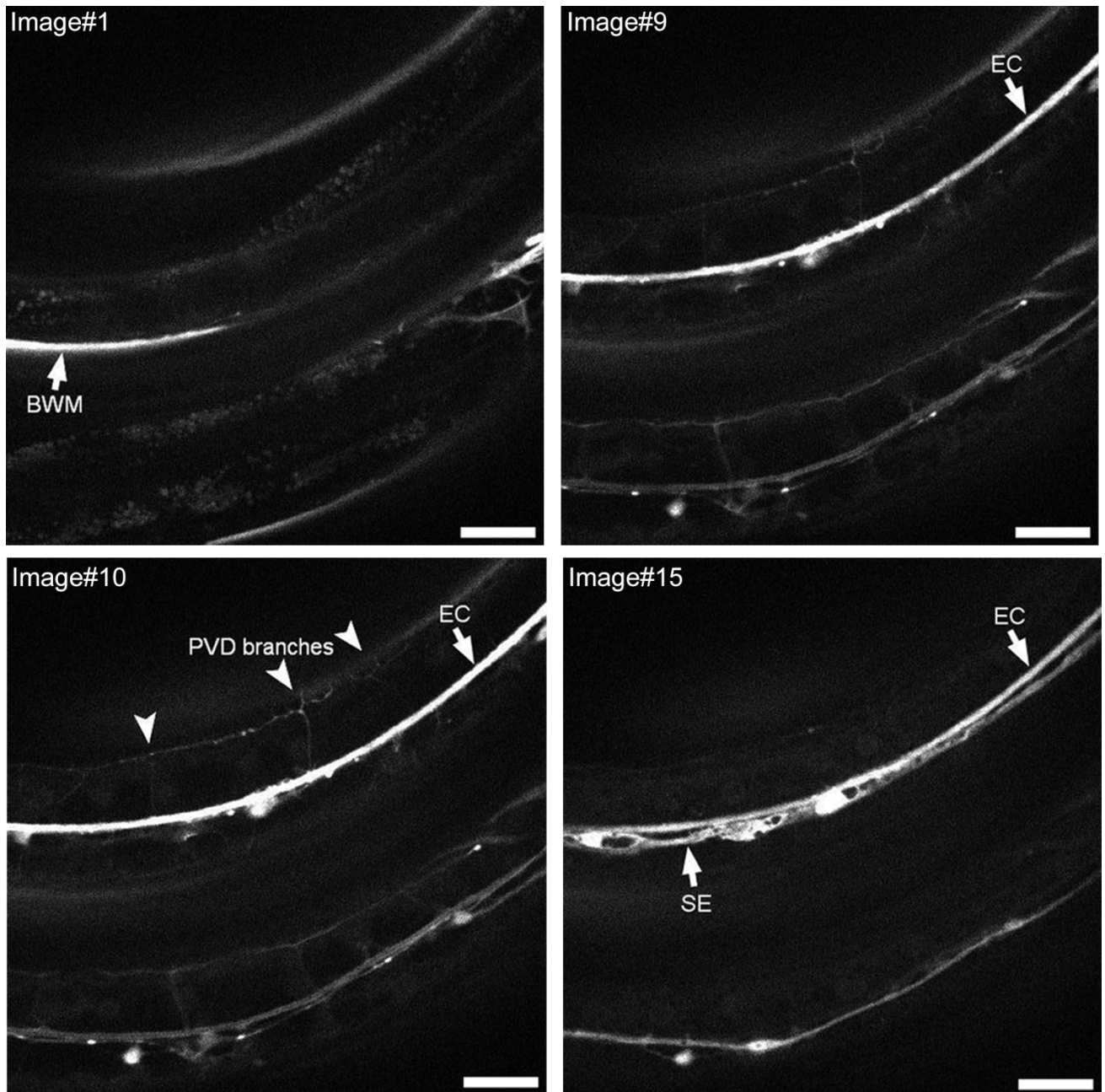
Fiji	NIH image	
GraphPad Prism 8	GraphPad Software, Inc.	
Adobe Photoshop CS5 and CS6	Adobe	
MetaMorph 7.8.1.0	Molecular Devices	https://www.moleculardevices
Micro-Manager	[75]	https://micro-manager.org

957 **Data for movies**



Movie S1. EFF-1 ectopic expression in BWMs results in Uncoordinated and Dumpy (Unc+Dpy) phenotypes

Mixed population of worms with extrachromosomal *pmyo-3::mCherry* and *pmyo-3::EFF-1*. White arrowhead point to mCherry(+) worm that is Unc+Dpy. White arrow point to mCherry(+) worm that is wt-like. Yellow arrowhead points to a wt-like mCherry(-) worm, which left the frame within seconds. Elapsed time (seconds) indicated in top left corner.



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979 **Movie S2. VSV Δ G-G infects PVD neuron**

980 SDC microscope Z-stack of young adult worm injected with 10^6 IU VSV Δ G-G and imaged 48h
981 later. Arrowheads, infected PVD's candelabra/menorahs arborized branches. Arrows, infected
982 cell. BWM-Body Wall Muscle, EC-Excretory Cell, SE-Seam cell syncytium. Scale bars, 25 μ m.
983 Note that there is a second worm (bottom, not indicated in images), also showing infected
984 PVD branches, EC, SE and a muscle cell.