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A cGMP-dependent protein kinase, encoded by the *Drosophila foraging* gene, regulates neurotransmission through changes in synaptic structure and function

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ABSTRACT

A cGMP-dependent protein kinase (PKG) encoded by the Drosophila foraging (for) gene regulates both synaptic structure (nerve terminal growth) and function (neurotransmission) through independent mechanisms at the Drosophila larval neuromuscular junction (nmj). Glial for is known to restrict nerve terminal growth, whereas presynaptic for inhibits synaptic vesicle (SV) exocytosis during low frequency stimulation. Presynaptic for also facilitates SV endocytosis during high frequency stimulation. for's effects on neurotransmission can occur independent of any changes in nerve terminal growth. However, it remains unclear if for's effects on neurotransmission affect nerve terminal growth. Furthermore, it's possible that for's effects on synaptic structure contribute to changes in neurotransmission. In the present study, we examined these questions using RNA interference to selectively knockdown for in presynaptic neurons or glia at the Drosophila larval nmj. Consistent with our previous findings, presynaptic knockdown of for impaired SV endocytosis, whereas knockdown of glial for had no effect on SV endocytosis. Surprisingly, we found that knockdown of either presynaptic or glial for increased neurotransmitter release in response to low frequency stimulation. Knockdown of presynaptic for did not affect nerve terminal growth, demonstrating that for's effects on neurotransmission does not alter nerve terminal growth. In contrast, knockdown of glial for enhanced nerve terminal growth. This enhanced nerve terminal growth was likely the cause of the enhanced neurotransmitter release seen following knockdown of glial for. Overall, we show that for can affect neurotransmitter release by regulating both synaptic structure and function.

Introduction

Synaptic strength is influenced by changes in both synaptic structure (nerve terminal growth) and function (neurotransmission). The Drosophila larval neuromuscular junction (nmj) is a well-established model system for studying synaptic structure and function (reviewed in Harris & Littleton, 2015). Effects on these processes have been shown to be interrelated in some cases (Budnik, Zhong, & Wu, 1990; Davis & Goodman, 1998; Sigrist, Thiel, Reiff, & Schuster, 2002; Stewart, Schuster, Goodman, & Atwood, 1996; Zhong, Budnik, & Wu, 1992; Zhong & Wu, 2004), but not others (Dason et al., 2009; Romero-Pozuelo, Dason, Atwood, & Ferrús, 2007; Romero-Pozuelo et al., 2014). While most studies using this model have focused on presynaptic and postsynaptic regulation of synaptic structure and function, the importance of glia in these processes is becoming increasingly apparent (Brink, Gilbert, Xie, Petley-Ragan, & Auld, 2012; Dason, Allen, Vasquez, & Sokolowski, 2019; Fuentes-Medel et al., 2009; Keller et al., 2011; Kerr et al., 2014).

The Drosophila foraging (for) gene encodes a cGMPdependent protein kinase (PKG) (Osborne et al., 1997) that ARTICLE HISTORY

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KEYWORDS

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affects numerous behaviours (see Anreiter & Sokolowski, 2019 for a detailed review). for's influence on these behaviours likely arise from for's effects at synapses. We recently found that for is expressed in both presynaptic neurons and glia at the Drosophila larval nmj and affects both synaptic structure and function (Dason et al., 2019). Specifically, we found that for⁰ null mutants have increased nerve terminal growth (Dason et al., 2019). Tissue specific rescues demonstrated that this increased nerve terminal growth was due to the absence of for in glia (Dason et al., 2019). Furthermore, for⁰ null mutants have increased neurotransmission in response to low frequency stimulation due to increased presynaptic Ca²⁺ entry and decreased neurotransmission in response to high frequency stimulation due to reduced synaptic vesicle (SV) recycling (Dason et al., 2019). Tissue specific rescues showed that both effects were due to presynaptic for (Dason et al., 2019).

Fluorescein-assisted light inactivation of *for* revealed that *for*'s effects on neurotransmission can occur independent of its effects on nerve terminal growth (Dason *et al.*, 2019). However, it remains possible that *for*'s effects on nerve terminal growth can also affect neurotransmission and *for*'s effects on neurotransmission may contribute to changes in

nerve terminal growth. To address these possibilities, we used RNA interference (RNAi) to selectively knock down all *for* transcripts in neurons or glia at the *Drosophila* larval nmj and examined whether *for*'s effects on synaptic structure and function were interrelated.

Material and methods

Fly stocks

Fly stocks were grown on molasses based fly medium in uncrowded conditions at 25 °C (Anreiter, Vasquez, Allen, & Sokolowski, 2016). Mid third instar larvae were used for all experiments. All transgenic lines were backcrossed into a wild-type *sitter* background (Allen, Anreiter, Neville, & Sokolowski, 2017). The GAL4/UAS system was used for neuron or glial specific expression of transgenes (Brand & Perrimon, 1993). A UAS-forRNAi line targeting all for transcripts and UAS-Dcr were used to knock down for in selected tissues (Dason *et al.*, 2020). *n-syb-GAL4* (Verstreken *et al.*, 2009) was used to drive expression of UAS-forRNAi, UAS-Dcr in neurons and Repo-GAL4 (Sepp, Schulte, & Auld, 2001) was used to drive expression of UAS-forRNAi, UAS-Dcr in glia.

Immunohistochemistry

Immunohistochemistry was performed as previously described (Cantarutti, Burgess, Brill, & Dason, 2018). Larvae were dissected, fixed and then incubated overnight at 4° C with primary antibodies diluted in blocking solution. FITC-conjugated anti-horseradish peroxidase (HRP) antibody (1:800 dilution; Jackson ImmunoResearch) was used to visualize neurons and the mouse monoclonal bruchpilot (brp) antibody (1:100 dilution; Iowa Hybridoma Bank; Wagh *et al.*, 2006) was used to visualize active zones. Larval fillet preparations were mounted in Permafluor (Immunon, Pittsburgh, PA) on a glass slide with a cover slip. Preparations were viewed under a TCS SP5 confocal laser-scanning microscope (Leica, Heidelberg, Germany) with a $63 \times$ oil-immersion objective (1.4 NA).

Electrophysiology

Intracellular recordings were performed in HL6 saline (Macleod, Hegstrom-Wojtowicz, Charlton, & Atwood, 2002) supplemented with 0.5 mM CaCl₂ as previously described (Dason, Smith, Marin, & Charlton, 2014). Sharp glass electrode filled with 3 M KCl (\sim 40M Ω) were used to measure spontaneously occurring miniature excitatory junction potentials (mEJPs) and stimulus-evoked excitatory junction potentials (EJPs) from the ventral longitudinal muscle fiber 6 (abdominal segment 3) of dissected larvae. A suction electrode was used to stimulate cut segmental nerves at 0.05 Hz. Electrical signals were recorded using the MacLab/4S data acquisition system (ADInstruments).

FM1-43 imaging

FM1-43 experiments were performed as previously described (Dason, Smith, Marin, & Charlton, 2010). The following high K⁺ saline: 25 mM NaCl, 90 mM KCl, 10 mM NaHCO₃, 5 mM HEPES, 30 mM sucrose, 5 mM trehalose, 10 mM MgCl₂, 2 mM CaCl₂, pH 7.2 (Verstreken, Ohyama, & Bellen, 2008) was to induce high K⁺ depolarization for FM1-43 loading and unloading. Presynaptic boutons were loaded with FM1-43 (Invitrogen) by high K⁺ depolarization for 2 min and subsequently washed for 5 min in Ca^{2+} free HL6. 75 µM Advasep-7 was included for the first 1 min of the wash to reduce background fluorescence from extracellular FM1-43 (Kay et al., 1999). A Leica TCS SP5 confocal laserscanning microscope with a 63X water dipping objective (0.9 NA) was used to take an image of FM1-43 uptake and then high K^+ depolarization for 2 min was used to induce exocytosis. A second image was then taken to document FM1-43 unloading. The released fraction was calculated using the following formula: (fluorescence of load - fluorescence of unload)/fluorescence of load.

Statistical analysis

SigmaPlot (version 11.0; Systat Software) was used for statistical analysis. One-way ANOVA tests (with a Holm–Sidak *post hoc* test) were used for comparing datasets. Error bars in all figures represent \pm standard error of the mean (sem).

Results

Glial for negatively regulates nerve terminal growth and evoked neurotransmission

We previously found that for expression in glia rescued the increased nerve terminal growth seen in the for^0 null mutant (Dason et al., 2019). We hypothesized that knockdown of glial for would phenocopy the increased nerve terminal growth seen in the for^0 null mutant. We used RNAi to knockdown for in glia at the Drosophila larval nmj and then stained nmjs with HRP (a neuronal membrane marker) and brp (an active zone marker). We counted the number of synaptic boutons on segment 3 of muscle fibers 6 and 7 in experimental and control genotypes. Two types of axons innervate these muscle fibers and result in 1b and 1s boutons, which differ in their morphological and physiological properties (Atwood, Govind, & Wu, 1993; Kurdyak et al., 1994). As predicted, we found that glial knockdown of for increased the number of 1b and 1s boutons compared to controls (Figure 1(A,B)). The number of active zones per bouton was not significantly different between genotypes (Figure 1(C,D)). We next examined the role of glial for in neurotransmitter release by recording the compound EJP generated by tonic-like type 1 b and phasic-like type 1 s boutons from segment 3 of muscle fiber 6 in third-instar larvae in response to low frequency stimulation (0.05 Hz). Surprisingly, we found glial knockdown of *for* significantly increased the amplitude of EJPs compared to controls (Figure 1(E,F)). There were no significant differences in the



Figure 1. Glial *for* negatively regulates neurotransmitter release and nerve terminal growth. A, Fixed larval nmjs stained with FITC-conjugated anti-HRP antibody. B, Glial specific knockdown of *for* (+;+;*UAS-for RNAi, UAS-dcr/Repo-GAL4*) significantly increased the number of 1 b (F(2,26) = 7.162, p < .05; n = 7-12) and 1 s boutons (F(2,26) = 5.166, p < .05; n = 7-12) in comparison to controls (+;+;*UAS-for RNAi, UAS-dcr/+* and +;+;*Repo-GAL4*/+). C, Representative images of 1 b and 1 s boutons stained with FITC-conjugated anti-HRP antibody and anti-BRP. D, The number of active zones per 1 b (F(2,34) = 0.725, p > .05; n = 10-14) or 1 s boutons (F(2,35) = 0.159, p > .05; n = 11-14) was not significantly different between experimental and control genotypes. E, Representative traces of EJPs and mEJPs. Preparations were maintained in HL6 (0.5 mM Ca²⁺) saline. F, The amplitude of evoked EJPs were significantly enhanced when glial *for* was knocked down in comparison to control genotypes (F(2,28) = 9.427, p < .05; n = 7-11) or frequency (F(2,24) = 0.713, p > .05; n = 7-11) between genotypes. Error bars represent ± sem.

amplitude (Figure 1(E,G)) or frequency (Figure 1(E,H)) of mEJPs between genotypes. Thus, glial *for* negatively regulates both nerve terminal growth and evoked neurotransmission.

this increased neurotransmission does not affect nerve terminal growth.

Presynaptic for negatively regulates neurotransmission

Presynaptic for negatively regulates evoked neurotransmission in response to low frequency stimulation (Dason et al., 2019). To determine if this increased neurotransmission alters nerve terminal growth, we used RNAi to knock down for in presynaptic neurons at the Drosophila larval nmj and assessed whether presynaptic for regulated nerve terminal growth. Presynaptic knockdown of for had no effect on the number of 1b and 1s boutons or the number of active zones per bouton compared to controls (Figure 2(A-D)). We next examined the role of presynaptic for in neurotransmitter release by recording EJPs in response to low frequency stimulation (0.05 Hz). As expected, we found that presynaptic knockdown of for significantly increased the amplitude of EJPs compared to controls (Figure 2(E,F)). There were no significant differences in the amplitude (Figure 2(E,G)) or frequency (Figure 2(E,H)) of spontaneously occurring mEJPs between genotypes. Thus, presynaptic for negatively regulates evoked neurotransmission and

Presynaptic for is required for SV endocytosis

We previously found that expression of presynaptic for, but not glial for, could rescue the impaired SV endocytosis seen in the for⁰ null mutant (Dason et al., 2019). To determine if reduced levels of presynaptic or glial for phenocopies the SV endocytosis impairment of the for⁰ null, we used the lipophilic dye FM1-43 (Betz & Bewick, 1992) to monitor SV cycling. We assessed SV cycling in control and experimental genotypes by measuring FM1-43 uptake by stimulating preparations with high K⁺ saline for 2 min in the presence of FM1-43 (Figure 3(A)). FM1-43 uptake was significantly reduced when presynaptic for was knocked down in comparison to controls (Figure 3(A,B)). No effect was observed when glial for was knocked down (Figure 3(B)). We next measured FM1-43 unloading by applying high K⁺ saline for 2 min to determine if the impairment seen when presynaptic for was knocked down was due to a defect in SV exocytosis. We found that a similar fraction was released by all genotypes (Figure 3(A,C)) demonstrating that an impairment in SV exocytosis was not the cause of the reduced FM1-43



Figure 2. Presynaptic *for* negatively regulates neurotransmitter release but has no effect on nerve terminal growth. A, Fixed larval nmjs stained with FITC-conjugated anti-HRP antibody. B, Neuronal specific knockdown of *for* (+;+;*UAS-for RNAi, UAS-dcr/n-syb-GAL4*) had no effect on the number of 1 b (F(2,21) = 0.166, p > .05; n = 7-9) and 1 s boutons (F(2,21) = 0.488, p > .05; n = 7-9) in comparison to controls (+;+;*UAS-for RNAi, UAS-dcr/+* and +;+;*n-syb-GAL4*/+). C, Representative images of 1 b and 1 s boutons stained with FITC-conjugated anti-HRP antibody and anti-BRP. D, The number of active zones per 1 b (F(2,20) = 0.313, p > .05; n = 10-13) or 1 s bouton (F(2,27) = 0.954, p > .05; n = 9-12) was not significantly different between experimental and control genotypes. E, Representative traces of EJPs and mEJPs. Preparations were maintained in HL6 (0.5 mM Ca²⁺) saline. F, The amplitude of evoked EJPs were significantly enhanced when presynaptic *for* was knocked down in comparison to control genotypes (F(2,25) = 7.117, p < .05; n = 7-11). G,H, There were no significant differences in mEJP amplitude (F(2,22) = 3.015, p > .05; n = 6-11) or frequency (F(2,22) = 0.577, p > .05; n = 6-11) between genotypes. Error bars represent ± sem.

uptake seen in larvae with presynaptic *for* knocked down. Thus, presynaptic *for* is required for SV endocytosis.

Discussion

Our study characterized the effects of presynaptic or glial knockdown of *for* on synaptic structure and function. In agreement with our previous findings using a for^0 null mutant, we found presynaptic *for* negatively regulates neuro-transmitter release in response to low frequency stimulation and is required for SV endocytosis, whereas glial *for* functions to restrain nerve terminal growth. Expanding on our previous work, we found that the enhanced neurotransmission resulting from presynaptic knockdown of *for* did not alter nerve terminal growth. Furthermore, we found that glial *for* can affect neurotransmission through its effects on nerve terminal growth.

for and axonal growth

Data from our study (Figure 1) and several others demonstrates that *for* functions to restrain axonal growth (Dason *et al.*, 2019; Peng *et al.*, 2016; Renger, Yao, Sokolowski, & Wu, 1999; Song *et al.*, 2019). Higher levels of *for* in allelic for variants have fewer ectopic nerve entry points in muscles at the Drosophila larval nmj (Renger et al., 1999). Drosophila for⁰ null mutant larvae have increased nerve terminal growth and this increased nerve terminal growth is rescued by expression of for in glia (Dason et al., 2019). Consistent with these findings, we show that selectively knocking down for in glia enhanced nerve terminal growth (Figure 1(A,B)). for's effects on nerve terminal growth are not a consequence of increased neurotransmission, as presynaptic knockdown of for increases neurotransmission without affecting nerve terminal growth (Figure 2). Genetic experiments in Drosophila suggest that for regulates nerve terminal growth by modulating TGF- β /BMP retrograde signaling (Dason *et al.*, 2019). Specifically, for may affect the secretion of the glial TGF- β ligand, May, which regulates nerve terminal growth through muscle derived Glass bottom boat (Gbb), the Drosophila TGF- β /BMP homologue (Fuentes-Medel *et al.*, 2012). Consistent with these Drosophila studies, a role for glial PKG in synaptogenesis was demonstrated in Xenopus laevis tadpole (Sild, Van Horn, Schohl, Jia, & Ruthazer, 2016). for is also required for restraining axonal growth in Drosophila embryos. for²⁰⁻²⁹ null mutants display axonal overgrowth, incorrect axon pathfinding and incorrect target recognition (Peng et al., 2016). for physically interacts with the transcription factor longitudinal lacking (lola) (Peng et al., 2016),

which is required for repelling longitudinal axons away from the midline (Crowner, Madden, Goeke, & Giniger, 2002). *for* regulates axon guidance by antagonizing the effects of Lola in *Drosophila* embryos (Peng *et al.*, 2016).

Drosophila for was also recently shown to restrict axon regeneration. Specifically, Drosophila for inhibits axon regeneration of larval sensory neurons (Song *et al.*, 2019). During axon regeneration, a mechanosensitive ion channel called Piezo is activated and induces local Ca^{2+} transients at the growth cone, leading to activation of nitric oxide synthase and *for*, which restrict axon regeneration demonstrating that *for* functions as a brake for axon regeneration (Song *et al.*, 2019). These findings are consistent with the reported roles of *for* negatively regulating axonal growth (Dason *et al.*, 2019; Peng *et al.*, 2016; this study). Collectively, these studies suggest that *for* negatively regulates axonal growth and that *for* could be a potential target for increasing axon regeneration and modulating nervous system repair.

Presynaptic for and neurotransmission

We found that knockdown of for in presynaptic neurons increased evoked neurotransmission in response to low frequency stimulation (Figure 2(E,F)). These findings are in agreement with our previous work that demonstrated that for⁰ null mutants have increased evoked neurotransmission and presynaptic Ca^{2+} entry in response to low frequency stimulation and that these effects could be rescued by expressing for in presynaptic neurons (Dason et al., 2019). Consistent with a role of for in negatively regulating evoked neurotransmission, an earlier study found that overexpression of for reduced evoked neurotransmission (Renger et al., 1999). Cultured Drosophila neurons of allelic variants of for with reduced PKG activity were previously shown to have decreased voltage-dependent K⁺ currents (Renger et al., 1999). A similar reduction of K^+ currents in for⁰ null mutants or larvae with for knock downed in presynaptic neurons could increase action potential duration, leading to increased presynaptic Ca²⁺ entry and increased evoked neurotransmission.

Knockdown of *for* in presynaptic neurons did not affect spontaneous neurotransmission (Figure 2(E,G,H)). Similarly, we previously found that spontaneous neurotransmission was not altered in *for*^{ρ} null mutants (Dason *et al.*, 2019). Our data suggests that *for* has a selective effect on evoked neurotransmission. These findings are consistent with growing evidence that evoked and spontaneous neurotransmission have distinct molecular mechanisms (Kavalali, 2015; Melom, Akbergenova, Gavornik, & Littleton, 2013; Pang *et al.*, 2011; Ramirez, Khvotchev, Trauterman, & Kavalali, 2012).

We previously used fluorescein-assisted light inactivation to acutely inactivate *for* and found that *for's* effects on neurotransmission can occur independent of any changes in nerve terminal growth (Dason *et al.*, 2019). In agreement with this, knockdown of presynaptic *for* had no effect on nerve terminal growth (Figure 1(A,B)), but did increase

(A) UAS-for RNAi UAS-for RNAi/ n-syb-GAL4



Figure 3. Presynaptic *for* regulates SV endocytosis. A, Representative images of presynaptic boutons loaded with FM1–43 during high K⁺ stimulation for 2 min. High K⁺ saline was then reapplied for 2 min and fluorescence was measured again (unload). B, Neuronal specific knockdown of *for* (+;+;*UAS-for RNAi, UAS-dcr/n-syb-GAL4*) significantly reduced the amount of FM1-43 uptake compared to controls (+;+;*UAS-for RNAi, UAS-dcr/+* and +;+;*n-syb-GAL4/+*) (F(2,17) = 10.170, *p* < .05; *n* = 4–10), demonstrating impaired SV recycling. Glial specific knockdown (+;+;*UAS-for RNAi, UAS-dcr/Repo-GAL4*) had no effect in comparison to controls (+;+;*UAS-for RNAi, UAS-dcr/+* and +;+;*Repo-GAL4/+*) (F(2,15) = 0.0197, *p* > .05; *n* = 4–8). C, A similar fraction of FM1-43 was released in controls and RNAi lines, demonstrating that recycled SVs could undergo exocytosis (presynaptic knockdown: (F(2,17) = 3.301, *p* > .05; *n* = 4–10 and glial knockdown: (F(2,15) = 1.702, *p* > .05; *n* = 4–8). Fluorescence (F) was reported with background F subtracted. Scale bars represent 4 µm. Error bars represent ± sem.

evoked neurotransmission (Figure 1(E,F)). Thus, presynaptic *for* directly regulates evoked neurotransmission.

Presynaptic for and SV endocytosis

We found that knockdown of presynaptic for impaired SV endocytosis (Figure 3). We previously used fluoresceinassisted light inactivation of FOR and a temperature-sensitive dynamin mutant, *shibire^{ts1}*, to demonstrate that FOR is necessary for endocytosis of SVs that have undergone exocytosis using a functional FOR protein (Dason et al., 2019). These data show that FOR's effects on SV endocytosis are not simply a consequence of altered SV exocytosis. These findings are consistent with several studies that propose that PKG plays a key role in balancing SV exocytosis and endocytosis (Collado-Alsina, Ramírez-Franco, Sánchez-Prieto, & Torres, 2014; Dason et al., 2019; Eguchi, Nakanishi, Takagi, Taoufiq, & Takahashi, 2012; Petrov, Giniatullin, Sitdikova, & Zefirov, 2008; Taoufiq, Eguchi, & Takahashi, 2013). During periods of sustained synaptic transmission, SV recycling is upregulated through a retrograde pathway that involves the release of nitric oxide (NO) from the postsynaptic cell and a subsequent increase in presynaptic cGMP and PIP₂ in cultured hippocampal neurons and synapses of the rat Calyx of Held (Eguchi et al., 2012; Micheva, Buchanan, Holz, & Smith, 2003). Similarly, application of NO donors at the Drosophila nmj induces cGMP immunoreactivity in presynaptic boutons (Wildemann & Bicker, 1999) and both presynaptic cGMP (Shakiryanova & Levitan, 2008) and PIP₂ levels (Dason et al., 2014) were found to be elevated during high frequency stimulation. PKG is activated by cGMP and thought to upregulate PIP₂ levels through a Rho-kinase (Eguchi et al., 2012; Taoufiq et al., 2013). PIP₂ recruits AP-2 and clathrin to sites of endocytosis (Micheva et al., 2003). Thus, a reduction in PIP₂ is likely the cause of the impaired SV endocytosis observed following the inhibition of PKG in mammalian studies or the absence of for in Drosophila studies.

Glial for's effects on synaptic structure and function

Growing evidence demonstrates a role for glia in regulating both synaptic structure and function. Specifically, glia are important mediators of processes such as neurotransmission (Panatier et al., 2011), synapse formation (Mauch et al., 2001) and synaptic plasticity (Henneberger, Papouin, Oliet, & Rusakov, 2010). We previously found that expressing for in glia in a for⁰ null mutant background rescued the enhanced nerve terminal growth seen in the for^0 null mutant but did not rescue the enhanced neurotransmission seen in the for⁰ null mutant (Dason et al., 2019). This clearly demonstrates that glial for is not having a direct effect on neurotransmission. In the present study, we found that knocking down glial for enhanced both nerve terminal growth and neurotransmission (Figure 1). This strongly suggests that while presynaptic for has direct effects on neurotransmission, glial for regulates neurotransmission through its effects on nerve terminal growth.

The for gene and behaviours

The for gene is required for survival (Anreiter et al., 2021) and known to affect many behaviours, such as foraging (Allen et al., 2017; Anreiter, Kramer, & Sokolowski, 2017; Anreiter & Sokolowski, 2018), learning and memory (Kaun, Hendel, Gerber, & Sokolowski, 2007; Mery, Belay, So, Sokolowski, & Kawecki, 2007), stress responses (Caplan, Milton, & Dawson-Scully, 2013; Dawson-Scully, Armstrong, Kent, Robertson, & Sokolowski, 2007; Krill & Dawson-Scully, 2016), sleep (Donlea et al., 2012) and nociception (Dason et al., 2020). Neuronal and glial for likely contribute in regulating many of these responses and behaviours. for is expressed in both neurons and glia in the larval CNS (Allen, Anreiter, Vesterberg, Douglas, & Sokolowski, 2018; Dason et al., 2020). Interestingly, for appears to be expressed primarily in glia in the adult CNS (Allen & Sokolowski, 2021). However, some expression of for in neurons in the adult CNS has also been reported (Eddison, Belay, Sokolowski, & Heberlein, 2012). In recent years, the importance of glia in flies and mammals have been shown in behaviours such as learning and memory (Adamsky et al., 2018; Frankland & Josselyn, 2020; Kol et al., 2020), sleep (Artiushin & Sehgal, 2020) and pain (Salter & Beggs, 2014). Future behavioural studies on FOR/PKG should examine the potential contributions of both neuronal and glial FOR/PKG. Our data demonstrates that for has pleiotropic effects at synapses, with presynaptic for directly regulating synaptic function, while glial for regulates synaptic function through effects on nerve terminal growth at the larval nmj.

Disclosure statement

No potential conflict of interest was reported by the author(s).

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