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Mushroom body defect is required in parallel to Netrin for midline axon guidance in Drosophila

Marie-Sophie Cate1, Sangeetha Gajendra¹, Samantha Alsbury¹,*, Thomas Raabe2, Guy Tear¹,‡,* and Kevin J. Mitchell³,⁴,*

ABSTRACT

The outgrowth of many neurons within the central nervous system is initially directed towards or away from the cells lying at the midline. Recent genetic evidence suggests that a simple model of differential sensitivity to the conserved Netrin attractants and Slit repellents is insufficient to explain the guidance of all axons at the midline. In the Drosophila embryonic ventral nerve cord, many axons still cross the midline in the absence of the Netrin genes (NetA and NetB) or their receptor frazzled. Here we show that mutation of mushroom body defect (mud) dramatically enhances the phenotype of Netrin or frazzled mutants, resulting in many more axons failing to cross the midline, although mutations in mud alone have little effect. This suggests that mud, which encodes a microtubule-binding coiled-coil protein homologous to NuMA and LIN-5, is an essential component of a Netrin-independent pathway that acts in parallel to promote midline crossing. We demonstrate that this novel role of Mud in axon guidance is independent of its previously described role in neural precursor development. These studies identify a parallel pathway controlling midline guidance in Drosophila and highlight a novel role for Mud potentially acting downstream of Frazzled to aid axon guidance.

KEY WORDS: Drosophila, Axon guidance, Midline, Mud, NuMA, LIN-5, Netrin

INTRODUCTION

In the central nervous system (CNS) of vertebrates and invertebrates most neurons extend across the midline to form commissures, while some remain extend on their own side (Tear, 1999; Kaprielian et al., 2001; Garbe and Bashaw, 2004; Evans and Bashaw, 2010). This decision depends in part on the responsiveness of the growth cone to Netrin attractants and Slit repellents, both secreted from cells at the midline, although additional mechanisms also exist to direct axons across the midline (Andrews et al., 2008; Dickson and Zou, 2010; Evans and Bashaw, 2010; Spitzweck et al., 2010; Organisti et al., 2015).

The Netrins act as chemoattractants to bring axons to the midline in flies, worms and vertebrates. In all these organisms, Netrin loss-of-function causes defects in the projection of axons towards the midline (Hedgecock et al., 1990; Harris et al., 1996; Mitchell et al., 1996; Serafini et al., 1996), and the same is true of mutations in the Netrin receptors unc-40, DCC and frazzled (Hedgecock et al., 1990; Kolodziej et al., 1996; Fazeli et al., 1997). However, their activity does not fully account for the guidance of all commissural axons across the midline, suggesting the existence of additional mechanisms (Brankatschk and Dickson, 2006).

In Drosophila, a number of components of Netrin-independent mechanisms that attract axons to the midline have been identified. Removal of either Decam, fni (stan – FlyBase) or robo2 significantly enhances the failure of midline crossing caused by the absence of frazzled alone (Andrews et al., 2008; Spitzweck et al., 2010; Organisti et al., 2015). However, the loss of any of these genes individually does not lead to a significant midline guidance defect. Thus, the role of these additional pathways in directing commissural axons across the midline is only revealed in the absence of Netrin signalling.

Here we show that Mushroom body defect (Mud) also has a role in a Netrin-independent signalling pathway directing commissural axons to the midline in Drosophila. Mud has previously been identified to function within neuroblasts and sensory organ precursors to couple the orientation of the mitotic spindle to both intrinsic and extrinsic cues (Bowman et al., 2006; Izumi et al., 2006; Siller et al., 2006; Siller and Doe, 2009; Segalen et al., 2010). We show that its role in axon outgrowth is independent of its activity within neuroblasts. Mud is expressed within postmitotic neurons, where it may act downstream of Frazzled to influence intrinsic neuronal polarity necessary for axonal outgrowth.

RESULTS AND DISCUSSION

Netrin deficiencies reveal the presence of an additional activity mediating axon guidance across the midline

Drosophila has two Netrin genes, NetA and NetB, which are adjacent on the X chromosome (Fig. 1A). The original studies investigating the role of the Drosophila Netrins reported a difference in phenotype between a small deficiency Df(1)NP5 that removed both Netrin genes and a slightly larger deficiency, Df(1)KA9, that extends further than NP5 (Fig. 1A) (Harris et al., 1996; Mitchell et al., 1996). Embryos hemizygous for the smaller deficiency Df(1)NP5 have thinner or occasionally absent axon commissures in the ventral nerve cord, with the posterior commissure being more strongly affected, and occasional breaks in the longitudinal connectives (Fig. 1B, Table 1). This phenotype is similar to that seen in embryos where only NetA and NetB have been removed (Fig. 1B, Table 1) (Brankatschk and Dickson, 2006), although Df(1)NP5 is slightly more severe (Andrews et al., 2008). By contrast, embryos hemizygous for the slightly larger deficiency
Df(1)KA9 exhibit a more severe phenotype, with a near complete loss of midline crossing in some commissures (Fig. 1B, Table 1). The larger deficiency affects the guidance of anterior and posterior commissural axons at the midline.

Restoration of either Netrin gene at the midline is sufficient to completely rescue the Df(1)KA9 phenotype, while rescuing the Df(1)NP5 to near wild type (Harris et al., 1996; Mitchell et al., 1996). These findings indicate the existence of a gene activity, also deleted in Df(1)KA9, that enhances midline crossing defects caused by the absence of the Netrins, but which has a mild phenotype when removed alone.

Markers for specific subsets of commissural neurons confirm the increased severity of commissural defects in Df(1)KA9 compared with Df(1)NP5 embryos (Table 1, Fig. 2). The Eg-GAL4 driver identifies the EG cluster of 10-12 cells that extend axons in the anterior commissure and the EW cluster of four cells that project in the posterior commissure (Higashijima et al., 1996; Dittrich et al., 1997; Garbe and Bashaw, 2007).

Midline crossing by Eagle-positive EG and EW neurons is significantly more disrupted in Df(1)KA9 than in Df(1)NP5, due both to stalling of axons prior to midline crossing and to misguidance, where axons extend across the midline along an aberrant trajectory (Table 1, Fig. 2). In Df(1)NP5 embryos the EW axons fail to cross the midline in 37% of segments, while 20% of the EG axons do not cross. In Df(1)KA9 embryos the number of segments where EW axons fail to cross the midline is increased to 78% (Fig. 2). The outgrowth of the SP1 neuron, one of the earliest axons to cross the midline in the anterior commissure, was examined using anti-Connectin (Meadows et al., 1994). Behaviour of the SP1 neurons mirrors that of the EG axons, with significantly more failing to cross the midline in Df(1)KA9 (57%) compared with Df(1)NP5 (21%).

**Mud is the enhancer of Netrins**

We used overlapping synthetic deficiencies in the region (Livingstone, 1985) to map the enhancer activity (Fig. 1A).
Table 1. Quantification of commissural phenotypes

<table>
<thead>
<tr>
<th>Genotype</th>
<th>Anterior commissure</th>
<th>Posterior commissure</th>
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<tbody>
<tr>
<td></td>
<td>Normal (%)</td>
<td>Thin (%)</td>
</tr>
<tr>
<td>BP102 axon staining</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Df(1)NP5</td>
<td>42</td>
<td>55</td>
</tr>
<tr>
<td>Df(1)KA9</td>
<td>10</td>
<td>56</td>
</tr>
<tr>
<td>mud1</td>
<td>90</td>
<td>10</td>
</tr>
<tr>
<td>fra957</td>
<td>43</td>
<td>29</td>
</tr>
<tr>
<td>pinsP6</td>
<td>89</td>
<td>11</td>
</tr>
<tr>
<td>fra957PinsP62</td>
<td>31</td>
<td>62</td>
</tr>
<tr>
<td>mud1frafra957</td>
<td>18</td>
<td>47</td>
</tr>
<tr>
<td>Df(1)KA9+mud rescue</td>
<td>44</td>
<td>55</td>
</tr>
<tr>
<td>mud4+mud rescue</td>
<td>100</td>
<td>0</td>
</tr>
</tbody>
</table>

Embryos stained with BP102 were quantified by characterising the thickness of the anterior and posterior commissures within each segment as normal, thin or absent. Thin commissures are those where the commissure is reduced in width to 70% or less of their normal size. Formation of the commissural pathways as revealed by Eagle (Eg)-GAL4 driving expression of Venus-YFP was quantified by characterising the thickness of the anterior (EG) and posterior (EW) commissures within each segment as normal, thin or absent. Thin commissures are those where the commissure is reduced in width to 70% or less of normal size. The posterior commissure phenotype of Df(1)KA9 is significantly different to that of Df(1)NP5 (chi-squared test, P<0.0001). Embryos stained with anti-Connectin were quantified by counting the number of hemisegments where the identified SP1 neuron is able to extend across the midline. The SP1 neuron phenotypes are significantly different (one-way ANOVA with Bonferroni correction, P<0.0001).

To see whether mud encodes the additional midline guidance activity removed in Df(1)KA9 we examined embryos double mutant for mud and frazzled (fra). fra embryos have a similar commissural axon guidance phenotype to small Netrin deficiencies (Fig. 1B) (Kolodziej et al., 1996). When mud alleles, or the small deficiency Df(1)B24-B54 that deletes the mud region, are combined with fra alleles the double-mutant embryos fail to form the majority of commissures – a phenotype indistinguishable from that of Df(1)KA9.

Confirmation that mud is necessary for the formation of the commissures that form in Df(1)NP5 embryos was demonstrated by reintroducing mud as a transgene into Df(1)KA9 embryos, using a BAC construct that contains the mud genomic region. This resulted in a rescue of the BP102 phenotype in Df(1)KA9 embryos to one resembling that of the smaller Df(1)NP5 deficiency (Fig. 1B). Thus, mud encodes the enhancer activity that accounts for the more severe phenotype observed in the larger Df(1)KA9 deficiency and is necessary to enable axons to cross the midline in this background. This places Mud as a component within an additional signalling pathway that directs axons to the midline, the role of which becomes apparent in the absence of Netrin signalling.

***mud mutation has direct effects on axon extension and guidance***

Mud has previously been shown to be required during the asymmetric division of embryonic neuroblasts, where it couples mitotic spindle orientation to cortical polarity at metaphase (Bowman et al., 2006; Izumi et al., 2006; Siller et al., 2006). Initial defects in this process in mud mutants are largely recovered by a realignment of the spindle during telophase, and only minor consequences have been reported on subsequent neuronal number and fate. The pattern of expression of Even-skipped is largely unchanged in mud mutants, Df(1)KA9 or Df(1)NP5 (Izumi et al., 2006) (data not shown). Similarly, the neurons identified by anti-Futsch (22C10) and anti-Fasciclin 2 (1D4) form as normal (data not shown), indicating little or no change in cell fate. To test further whether the roles of Mud during neuroblast division and axon outgrowth are separable, the ventral nerve cord phenotypes of fra;pins double mutants were examined. Pins functions with Mud in asymmetric neuroblast division (Siller et al., 2006) and if loss of mud during neuroblast division leads to defects in subsequent axon outgrowth, then a pins mutant should have a similar effect on axon guidance and would enhance fra phenotypes. However, the ventral nerve cord phenotype of fra;pins double mutants is no more severe than that of fra mutants alone (Fig. 3A). The effects on axon guidance due to the absence of Mud are thus not attributable to a disruption of cell fate.

We also examined the consequence of manipulating the levels of Mud activity within neuroblasts and neurons using the UAS-GAL4 system (Brand and Perrimon, 1993). We made use of a P-element insertion (EY20197) that inserts a UAS immediately upstream of mud. Increasing Mud expression in neuroblasts using the Sca-GAL4 driver did not result in axon outgrowth defects (data not shown). However, increasing Mud expression in Eagle-positive neurons
caused a reduction in the number of axons extending across the midline through the anterior commissure (Fig. 3B) suggesting that Mud acts in a dose-dependent manner in axons. The decrease in axon number is not due to a loss of the cells. This disruption is consistent with a direct role of Mud in axonal projection or guidance in postmitotic neurons.

**Mud is expressed in postmitotic neurons**

Mud contains multiple coiled-coil domains and a microtubule-binding domain and shares similarity to the vertebrate protein NuMA (Numa1) and LIN-5 of *C. elegans* (Guan et al., 2000; Bowman et al., 2006; Izumi et al., 2006; Siller et al., 2006). The *Drosophila* gene encodes seven isoforms, and a probe that detects all isoforms shows that mud transcripts are expressed throughout embryonic development. Zygotic expression is restricted to the ventral nerve cord from stage 11 and mud remains expressed in the ventral nerve cord until the end of embryogenesis.
Mud protein expression has previously been described as localised to both the apical cortex and the centrosome of neuroblasts (Izumi et al., 2006). We find that Mud is also expressed within neurons, where it is localised within a punctate pattern within the soma. Mud is expressed in most, if not all, neurons and is also present within midline cells and members of the longitudinal glia (Fig. 3C). NuMA has similarly been reported to be expressed in a particulate distribution within the somatodendritic compartment of postmitotic sympathetic and hippocampal neurons, a distribution that requires intact microtubules (Ferhat et al., 1998). Mud, in common with its LIN-5 and NuMA homologues, is able to bind microtubules, has a conserved role in regulating mitotic spindle formation and is able to link intrinsic or extrinsic cues to orient spindle formation (Du et al., 2001; Segalen et al., 2010). Mud also has an ability to recruit dynein/dynactin (Siller and Doe, 2009) and functions in the planar cell polarity pathway (Segalen et al., 2010; Johnston et al., 2013), raising the possibility that Mud might function in neurons to link polarity information with the dynnein/ dynactin complex to orient microtubule structures within neurons to encourage directed outgrowth.

**Commissure formation utilises a variety of partially redundant pathways**

Multiple signalling pathways, in addition to Netrins, cooperate to direct axon outgrowth towards and across the midline. Mutations in the genes encoding the transmembrane proteins Dscam1, Robo2 and Fmi or the intracellular proteins Abi and Trio and now Mud can all dramatically enhance the reduction of axonal midline crossing in Netrin or fra mutants (Forsthoefel et al., 2005; Andrews et al., 2008; Spitzweck et al., 2010; Evans et al., 2015; Organisti et al., 2015).

Because Mud has previously been implicated in a signalling pathway downstream of Frizzled (Fz) in planar cell polarity (Segalen et al., 2010), we tested for genetic interactions between Fz and Netrins and between Fz and mud. We find, as recently independently reported (Organisti et al., 2015), that NetAB/Fz double mutants display a severe lack of commissures, similar to mud/fra or Df(1)K949 mutants, whereas mud/fz mutants are not appreciably more severe than either mud or fz mutants alone (Fig. 3D). These data are consistent with a model whereby Fz and Mud both operate in a common, parallel pathway to the Netrins, possibly in concert with Fmi (Organisti et al., 2015). Double mutants of mud with Robo2 or with Dscam1 did not show a significant increase in midline crossing defects (data not shown), which formally suggests that Mud might also act in common pathways with these proteins, but this straightforward interpretation is complicated by the multiple functions demonstrated for Robo2 and Dscam1 in midline guidance (Andrews et al., 2008; Evans et al., 2015).

Further investigations are necessary to gain a better understanding of the mechanisms through which these multiple pathways are integrated within growth cones to enable the precise navigation of commissural axons at the midline.

**MATERIALS AND METHODS**

**Genetics**

The following Drosophila stocks were used: (1) mud<sup>1</sup>/Fm7cGal, (2) mud<sup>1</sup>/Fm7cGal, (3) fra<sup>277</sup>/CyO;glacZ, (4) Df-NP5/Fm7cGal, (5) Df-KA9/Fm7cGal, (6) NetAB/Fm7bactin (courtesy of B. Dickson, IMP, Vienna), (7) fra<sup>1</sup>/TM6bAbdAcZ, (8) mud<sup>1</sup>/Fm7c.fra<sup>957</sup>/CyO;wgBgal, (9) Pins<sup>966</sup>/TM6bAbdAcZ, (10) fra<sup>957</sup>/CyO;wgBgal;Pins<sup>P261</sup>/TM6bAbdAcZ, (11) mud<sup>1</sup>/Fm7cGAL;Pins<sup>P966</sup>/TM6bAbdAcZ, (12) P[EPgy2/EY20197, (13) egGAL4::UASCD8GFP, (14) elavGAL4 on II, (15) egGAL4, (17) mud<sup>1</sup>/Fm7c::GFP and (18) fra<sup>957</sup>/CyO;D6fEYFP. X-Y translocation stocks with breakpoints in the 12E-13A region, originally constructed by Stewart and Merriam (1973), were used to generate deficiencies for defined regions of the X chromosome as described by Ashburner (1989). Unless otherwise stated, stocks were obtained from the Bloomington Stock Center.

**Molecular biology**

A genomic rescue construct for mud was created in P[acman] (Venken et al., 2006). 17.5 kb was retrieved from BAC CH322-147E14 (BACPAC Resources Center) (Venken et al., 2009) covering chromosome arm X from 14138384 to 14157868. The rescue construct includes the promoter region of mud, located 147 bp downstream of CG32599 to 1546 bp upstream of mud, the mud gene and region downstream to 363 bp upstream of the closest downstream gene, CG1461. 500 bp homology arms homologous to the left and right ends of the transgene were subcloned into P[acman] to create a targeting construct. MW005 cells [courtesy of Colin Dolphin, King’s College, London; described in Westenberg et al. (2010)] were made competent for recombinering by inducing the Red recombinase essentially as described (Dolphin and Hope, 2006). After verification of the correct integration of genomic DNA into P[acman] by sequencing, transgenic flies containing this P[acman]-mud construct inserted into the VK6 attP site at 19E7 on the X chromosome were obtained by BestGene. A Venus-YFP tag was inserted at the N-terminus of Mud using Gibson assembly and the construct inserted at attP40 by BestGene.

**Immunohistochemistry**

Embryos were collected, fixed and stained as previously described by Patel (1994). The following primary antibodies were used: monoclonal antibody BP102 (Developmental Studies Hybridoma Bank; 1:20), mouse anti-β-Gal (Promega, Z3781; 1:300), mouse anti-Connectin [courtesy of Robert White, University of Cambridge, UK (Meadows et al., 1994); 1:20] and rabbit anti-GFP (ThermoFisher Scientific, A6455; 1:300). Secondary antibodies were purchased from Molecular Probes. Stacks of images were obtained using a Zeiss LSM 510 confocal microscope and processed using Velocity 5.2 imaging software (Improvement).

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**Competing interests**

The authors declare no competing or financial interests.

**Author contributions**

M.-S.C. and K.J.M. contributed equally to the experimental work. S.G. and S.A. also contributed to the experimental work. T.R. contributed reagents. K.J.M. and G.T. conceived the experiments, obtained funding and prepared the manuscript.

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**References**


