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## Lateral neuron-glia interactions steer the response of axons to the Robo code

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### Abstract

*Glia are required for axon pathfinding along longitudinal trajectories, but it is unknown how this relates to the molecular paradigm of axon guidance across the midline. Most interneuron axons in bilateral organisms cross the midline only once. Preventing them from recrossing the midline requires the expression of Robo receptors on the axons. These sense the repulsive signal Slit, which is produced by the midline. The lateral positioning of longitudinal axons depends on the response to Slit by the combination of Robo receptors expressed by the axons, on selective fasciculation, and on longitudinal (lateral) glia. Here, we analyse how longitudinal glia influence reading of the 'Robo code' by axons. We show that whereas loss of *robo1* alone only affects the most medial axons, loss of both glial cells missing (*gcm*) and *robo1* causes a severe midline collapse of longitudinal axons, similar to that caused by the loss of multiple Robo receptors. Furthermore, whereas ectopic expression of *robo2* is sufficient to displace the medial MP2 axons along a more lateral trajectory, this does not occur in *gcm-robo1* double-mutant embryos, where axons either do not extend at all or they misroute exiting the CNS. Hence, lateral neuron-glia interactions steer the response of axons to the Robo code.*

### Keywords

Glia; axon guidance; *robo*; *gcm*; neuron-glia interactions

## INTRODUCTION

The 'decision' of growth cones to cross the midline and never to recross it is a successful paradigm for the molecular dissection of growth-cone guidance (Bagri *et al.*, 2002; Culverwell and Karlstrom, 2002; Dickson, 2002; Guthrie, 2001; Tear, 1999a; Tear, 1999b). Similar molecular mechanisms operate in other contexts in which either axons or cells navigate and migrate towards a target (Englund *et al.*, 2002; Gilthorpe *et al.*, 2002; Kinrade *et al.*, 2001; Kramer *et al.*, 2001; Wong *et al.*, 2002). Coordination of left-right movement in all bilateral organisms requires that interneuron axons linking the motor systems with the brain cross the midline once only. The midline is the source of a repulsive diffusible signal that is encoded by *slit*. The Slit gradient is read by axons as they upregulate the expression of a combination of Robo receptors on crossing the midline (Ba-Charvet *et al.*, 1999; Batty *et al.*, 1999; Brose *et al.*, 1999; Kidd *et al.*, 1999; Kidd *et al.*, 1998a; Kidd *et al.*, 1998b; Rajagopalan *et al.*, 2000; Simpson *et al.*, 2000a; Simpson *et al.*, 2000b). The expression by a neuron of a particular combination of *robo1*, *robo2* and *robo3* receptors determines the lateral position of its axon in response to different thresholds of the Slit gradient (Rajagopalan *et al.*, 2000; Simpson *et al.*, 2000a; Simpson *et al.*, 2000b). Thus, axons that

contain Robo1, Robo2 and Robo3 will project further away from the midline than those with Robo1 and Robo3, and, in turn, these will be further from the midline than axons with only Robo1.

Robo receptors are also thought to be involved in fasciculation (Rajagopalan *et al.*, 2000; Simpson *et al.*, 2000a; Simpson *et al.*, 2000b). As an interneuron axon extends longitudinally, it 'chooses' which trajectory to follow, a process known as 'selective fasciculation' (Raper *et al.*, 1984). Accordingly, an axon might follow a more lateral or medial trajectory, depending on selective fasciculation, which is dictated by the combination of Robo receptors expressed and other cell-adhesion molecules that are present in the axons, such as FasII (Rajagopalan *et al.*, 2000; Simpson *et al.*, 2000a; Simpson *et al.*, 2000b). The intracellular domains of the different Robo receptors differ and Robo receptors can form homodimers (Hivert *et al.*, 2002). Conceivably, Robos might also interact with other cell-surface molecules.

The ability of Robo receptors to determine axonal positioning has been demonstrated by ectopic expression of Robo2 and Robo3 in axons that project medially in normal embryos, which causes the lateral displacement of these axons (Rajagopalan *et al.*, 2000; Simpson *et al.*, 2000a). Normally, Robo2 is expressed over the lateral FasII fascicle, in the axons that project furthest away from the midline (Rajagopalan *et al.*, 2000; Simpson *et al.*, 2000a). The dMP2 and vMP2 axons at the end of embryogenesis normally project along the first, medial fascicle closest to the midline (Hidalgo and Brand, 1997). Ectopic expression of Robo2 in the MP2 axons displaces their trajectory laterally, to the central fascicle (Simpson *et al.*, 2000a). However, ectopic expression of Robo2 does not displace MP2 axons sufficiently far from the midline to reach the domain of axons that express Robo2 endogenously. Although this might reflect either insufficient Robo2 levels, because increasing Robo2 concentration shifts the lateral displacement of other axons (Rajagopalan *et al.*, 2000), or the need for these axons to also express *robo3* in order to reach the most lateral domain, it might also indicate that other factors restrict the lateral positioning of axons.

Glia are necessary for axon guidance and axonal fasciculation (Bastiani *et al.*, 1986; Hidalgo and Booth, 2000; Poeck *et al.*, 2001; Rangarajan *et al.*, 1999; Sepp and Auld, 2003; Sepp *et al.*, 2001; Silver *et al.*, 1982). Longitudinal glia (LG) overlie the longitudinal axons of the ventral nerve cord of the *Drosophila* embryo, whereas midline glia enwrap the commissural portion of the axons as they cross the midline. LG occupy positions where growth cones 'make steering decisions' during growth cone guidance (called choice points); they trigger fasciculation and defasciculation events; and they maintain neuronal survival and the integrity of the neuropile (Booth *et al.*, 2000; Hidalgo and Booth, 2000). The major axonal fascicles along which the different *robo*-expressing axons extend are formed by interactions between the LG and the pioneer axons (Hidalgo and Booth, 2000). Multiple inputs, such as cell-survival interactions and cell contact between axons and glia confine axonal trajectories regardless of the presence of the Robo1 receptor on axons (Kinrade *et al.*, 2001). In fact, when lateral neuron-glia interactions are disrupted, axons misroute across the midline despite the fact that they express *robo1* and that the midline still produces Slit. This shows that interactions between neurons and glia have a crucial role in lateral positioning of axons that is independent of the Slit gradient. Here, we ask whether LG can influence the reading of the Robo code by axons.

Slit and Robo1 are involved in glial migration before axon guidance (Kinrade *et al.*, 2001). LG migrate in two distinct phases (Hidalgo and Booth, 2000; Kinrade *et al.*, 2001): in the first, neuron-independent, phase, the LG migrate medially from their site of origin at the edge of the neuroectoderm towards the midline, until they contact the cell bodies of the

pioneer neurons (medial migration); in the second phase (during axon guidance), the LG migrate longitudinally, along the antero-posterior axis in close contact with extending pioneer growth cones (longitudinal migration). Longitudinal migration correlates with the movements of axons (Hidalgo and Booth, 2000) and if normal axonal patterns are disrupted LG will migrate to retain axonal contact (Kinrade *et al.*, 2001). Medial migration takes place within domains of *robo1*-expressing cells. The glia stop migrating at a fixed distance from the midline as a consequence of Slit signalling, which coincides with the moment in which the LG contact the pioneer neuron cell bodies. In fact, LG severely collapse over the midline in *slit* mutants, and overexpression of *slit* in neurons can prevent the medial migration of LG (Kinrade *et al.*, 2001). The expression of *robo1* then declines in the LG, which do not express *robo1* during longitudinal migration. Thus, during axon guidance, LG migration does not require Robo1. However, in many cases, Slits are required for cell migration (Englund *et al.*, 2002; Gilthorpe *et al.*, 2002; Kramer *et al.*, 2001; Wong *et al.*, 2002). Thus, it is possible that Robo2 and Robo3 influence glial migration during axon guidance.

Taking into account the Robo code and that neuron-glia interactions are required for axon guidance, fasciculation and lateral axonal positioning, there are two conceivable mechanisms organising longitudinal axonal patterns: (1) both axons and glia respond to the Slit gradient, LG produce Robo receptors and, thus, Slit is the overall organiser of the CNS neuropile. In this case, LG influence axon guidance but they might not influence the reading of the Robo code by axons. (2) Only axons respond to Slit and glial positioning follows other cues. In this later case, neuron-glia interactions constitute a separate input to which axons respond and LG might influence the output of the Robo code.

## OBJECTIVE

Here, we ask whether LG influence the reading of the Robo code by axons. To approach this question we first asked whether LG might themselves respond directly to the Robo code. Hence, we tested whether the LG express *robo2* and/or *robo3* during axon guidance. Since the glia do not express *robo2* and *robo3* but they are involved in axon guidance, we next asked whether the absence of functional glial cells might alter the lateral displacements normally observed in conditions of loss and gain of function of Robo receptors. For this we looked at axonal patterns in *gcm-robo1* double-mutant embryos and at the effects of expressing *robo2* in medial neurons in these embryos.

## METHODS

### Flies

The following fly stocks were used: (1) *gcm*<sup>ΔP1</sup>/CyOlacZ; (2) *robo*<sup>Z14</sup>/CyOlacZ; (3) *robo2*<sup>8</sup>/Cyo wglacZ; (4) *robo3*<sup>1</sup>/CyolacZ; (5) F263 (LG-lacZ reporter) (Jacobs *et al.*, 1989); (6) 15J2 (MP2 GAL4); (7) UASrobo2; (8) UASGAPGFP. The following stocks were generated by conventional genetics: (1) *gcm*<sup>ΔP1</sup> *robo*<sup>Z14</sup>/CyOlacZ; (2) *gcm*<sup>DP1</sup>/CyO; UASGAPGFP; (3) *gcm*<sup>ΔP1</sup>/CyO; 15J2; (4) *gcm*<sup>ΔP1</sup> *robo*<sup>Z14</sup>/CyOlacZ; 15J2; (5) *gcm*<sup>ΔP1</sup> *robo*<sup>Z14</sup>/CyOlacZ; UASGAPGFP; (6) *gcm*<sup>ΔP1</sup> *robo*<sup>Z14</sup>/CyOlacZ; UASrobo2-myc; (7) *robo*<sup>Z14</sup>/CyOlacZ; 15J2; (8) *robo*<sup>Z14</sup>/CyOlacZ; UASGAPGFP; (9) *robo*<sup>Z14</sup>/CyOlacZ; UASrobo2. Mutant embryos were identified by staining the reporter-lacZ marked balancers with anti-Bgal antibodies.

### Immunohistochemistry

Antibody stainings were performed following a standard protocol (Patel, 1994). Antibodies were used in the following dilutions for fluorescence detection (for HRP detection, primary antibody concentrations were halved): rabbit anti-Bgal (Cappel), 1:2500; rabbit anti-Repo, 1:100; mouse anti-Repo, 1:10; mouse FasII, 1:2; mouse anti-Myc (9E10, Iowa Hybridoma Bank), 1:10; rabbit anti-GFP (AbCam), 1:250; mouse anti-Robo2 (HRP, Iowa), 1:500;

mouse anti-Robo3 (HRP), 1:10; mouse 22c10, 1:5; mouse anti-Slit (HRP), 1:10. As secondary antibodies we used biotinylated antibodies at 1:300 (Jackson and Vector Labs) followed by streptavidin Alexa 488 and Alexa 546 at 1:250 or the Elite kit (Vector Labs), or directly conjugated to Alexa 488 and Alexa 546 at 1:250 (Molecular Probes). Anti-Myc was detected using the Tyramide Signal Amplification kit (PerkinElmer).

### Microscopy and imaging

All fluorescence microscopy used a BioRad Radiance 2000 laser scanning confocal microscope. 3D imaging used Volocity (Improvision).

## RESULTS

### LG do not require the Robo code for longitudinal migration

To investigate whether LG respond to the Robo code, we looked at glial migration in the absence of signalling by Robo1, Robo2 and Robo3. We find that glial migration follows the positions of axons in all genetic backgrounds (Fig. 1). In *robo1* mutants, the medial axonal fascicle misroutes across the midline, whereas the intermediate and lateral fascicles extend laterally: some LG migrate over the midline whereas some retain their lateral positions (Fig. 1B, 100% penetrance,  $n=5$  embryos). In *robo3* mutants, the medial and intermediate axonal fascicles fuse, some LG line up over the fused fascicles, whereas some remain laterally together with remaining lateral axons (Fig. 1C, 100% penetrance,  $n=9$  embryos). In *robo2* mutants, the axons are defasciculated and the LG are disorganised, but the phenotype is rather weak (Fig. 1D, 26% penetrance,  $n=96$  hemisegments) and the glia do not migrate abnormally closer to the midline. Because LG migration is affected in both *robo1* and *robo3* mutants, LG migration might depend on either the Robo code or on contact with axons.

We already know that LG do not express *robo1* at this stage (Kinrade *et al.*, 2001), which indicates that longitudinal migration of LG requires contact with axons. To verify whether *robo2* and *robo3* are expressed in the LG during axon guidance, we looked at their distribution with anti-Robo2 and anti-Robo3 antibodies and visualised the LG with either the nuclear glial marker anti-Repo or anti- $\beta$ gal in embryos bearing the LG-lacZ reporter (Fig. 2) (Jacobs *et al.*, 1989). Robo2 is expressed at low levels from the end of stage 12. At this stage, the Robo2 signal that appears to surround LG cells is erratic (Fig. 2A,B) and more ventral than that of Repo in the LG (Fig. 2A,G). From stage 13/14 on, the Robo2 signal is more ventral (Fig. 2C) than the LG (Fig. 2D), and a lateral view shows that Robo2 signal is physically separated from the LG nuclei, which lie dorsally (Fig. 2H). At stage 16 and 17, the Robo2 signal appears to partially surround the LG from a frontal view (Fig. 2I), but from a transverse view it is clear that Repo<sup>+</sup> LG lie dorsally over the Robo2 axons (Fig. 2J). These temporal profiles show that Robo2 is not expressed in the LG during axon guidance.

Robo3 is expressed first at stage 15, and expression is maintained in the same pattern to the end of embryogenesis. By *in situ* hybridisation, we find that *robo3* transcripts are not located in the dorsal plane of the neuropile (Fig. 2M) but in a ventral plane (Fig. 2N). Thus, Robo3 transcripts are not in the plane along which LG are located, implying that *robo3* is not expressed in the LG. Fluorescent imaging of Robo3 together with the glial markers reveals apparent colocalization in the LG when the whole stack is projected onto one single image. However, this is deceiving. From a frontal view, the domains of Robo3 include the LG, and they surround nuclei that express Repo (Fig. 2O) but, again, the axonal staining of Robo3 is ventral and physically separated from the overlying Repo signal in LG (Fig. 2K). When LG are visualised with the lacZ reporter and anti- $\beta$ gal antibodies, the cytoplasmic projections of the LG are seen to enwrap, but not mix with, the axonal bundles that express Robo3 (Fig. 2P), which can also be seen in transverse sections (Fig. 2L). Thus, Robo3 is not expressed in the LG.

To further test that glia do not express *robo2* and *robo3*, we checked whether they are expressed in embryos that lack functional glia in *glial cells missing (gcm)* mutants. We found that at stage 13 *robo2* is expressed in clusters in the same antero-posterior location as the LG (Fig. 3A). In *gcm* mutants (Fig. 3D), Robo2 signal is still present in all hemisegments, confirming that at stage 12 *robo2* expression is neuronal. Nevertheless, there is a slight decrease in Robo2 signal in these embryos, which could be caused by loss of neurons through apoptosis in the absence of glia (Booth *et al.*, 2000). Later, at stage 16, *robo2* is still expressed in clusters in *gcm* embryos (Fig. 3B,E). However it is absent from the axons, which is consistent with the fact that axons cannot extend in the absence of glia, and with an increase in neuronal death (Booth *et al.*, 2000; Hidalgo and Booth, 2000). *Robo3* expression starts later, at stage 15 and the pattern does not change for the remainder of embryogenesis (Fig. 3C). Robo3 is still present in clusters in *gcm*-mutant embryos (Fig. 3F). However, as with *robo2*, signal is lost along the axonal tracts. Interestingly, we do not see overexpression of either *robo2* or *robo3* in *gcm*-mutant embryos, in which glia are presumed to adopt a neuronal fate. These data confirm that expression of *robo2* and *robo3* is restricted to neurons.

Because *robo2* and *robo3* are not expressed in the LG, longitudinal migration of LG does not rely on the integration of the Robo code cell-autonomously in the glia. Glia are always displaced following axons in the *robo1*, *robo2* and *robo3*-mutant embryos, which indicates that glial migration depends on interactions with axons and on other molecular mechanisms.

### Loss of glia induces midline collapse in loss-of-function *robo1* mutants

We wondered whether, reciprocally, glial-cell function might explain the different effects of *robo1* mutations on axonal trajectories. In *robo1* mutants, only the medial FasII fascicle misroutes over the midline; the outermost fascicles extend normally, parallel to the midline (Kidd *et al.*, 1998a) (Fig. 4B). Double-mutant *robo2-robo1* embryos have a dramatic midline collapse of the outer fascicles as well, supporting the notion that multiple Robo receptors keep axons away from the midline (Simpson *et al.*, 2000a; Simpson *et al.*, 2000b). We wondered if LG might also help the lateral fascicles retain their positions in *robo1* mutants.

We looked at the FasII fascicles in *robo1* mutants that also lack normal glial function, that is, *gcm-robo1* double-mutant embryos (Fig. 4). To our surprise, in *gcm-robo1* double-mutant embryos all fascicles can collapse along the midline (Fig. 4D), with full collapse observed in 12.5% of cases ( $n=144$  segments) and collapse from one side of the embryo in 22.6% ( $n=288$  hemisegments). This phenotype is never observed in either mutant alone. In *gcm-robo1* double mutants, 31% of axons also misroute towards the muscle ( $n=288$  hemisegments) (Fig. 4D, asterisk), a phenotype observed with less frequency in *gcm* mutants. Thus, in a *gcm-robo1* double-mutant background, the absence of glia causes a disruption to the lateral fascicles that is comparable to the removal of multiple Robo receptors.

Axons might project along the midline in *gcm-robo1* double mutants if the either midline glia are absent, despite the fact that *gcm* is not expressed in the midline glia, or if no repulsive Slit signal is produced. Thus, we verified whether Slit is produced in the double mutants. We find that, in *gcm-robo1* double-mutant embryos, Slit signal is produced (Fig. 4E,F) and the midline glia are present but disorganised. These results show that midline collapse in *gcm-robo1* double mutants is not caused by lack of repulsion from the midline.

Axons might collapse over the midline because of extensive loss of Robo2 and Robo3 neurons, caused by neuronal apoptosis induced by lack of glia (Booth *et al.*, 2000). To test whether Robo3 neurons, which are amongst the most abundant, are still present in *gcm-robo1* double-mutants embryos, we stained embryos with anti-Robo3 antibodies (Fig. 4F).

We find abundant distribution of Robo3 axons. Interestingly, most of these axons now cross the midline aberrantly (Fig. 4F). This means that midline collapse in *gcm-robo1* double-mutant embryos is not only caused by extensive neuronal apoptosis.

Longitudinal axons project along the midline in *gcm-robo1* mutant embryos even though they express Robo3. This is consistent with previous findings that show that interfering with lateral neuron-glia interactions causes midline crossing despite *robo1* expression and repulsion from the midline (Kinrade *et al.*, 2001). Thus, in the absence of glia, axons misroute to find alternative trajectories, regardless of midline-derived repulsion and the expression of Robo receptors by the axons. Because Robo3 labels only follower axons, LG are necessary to keep follower axons away from the midline independent of the Robo code.

### Different requirements for *robo1* and glia among pioneer axons

To determine whether these dramatic axonal defects are specific to follower neurons or due to primary defects in axon guidance that can be traced back to abnormal function of pioneer neurons, we looked at the extension of pioneer axons in *gcm-robo1* double mutants. In wild-type the descending dMP2/MP1 pioneer fascicle meets the ascending vMP2/pCC pioneer fascicle to form a characteristic chain-like structure (Fig. 5A) (Hidalgo and Brand, 1997). In *gcm* mutants the vMP2/pCC fascicle extends normally, but the dMP2/MP1 fascicle extends outwards straight towards the muscle (88.2%,  $n=144$  hemisegments) (Fig. 5B) (see also Hidalgo and Booth, 2000). In *robo1* mutants the converse is found and only the pCC/vMP2 fascicle is affected, collapsing over the midline (100%,  $n=191$ ) (Fig. 5C). In *gcm-robo1* double-mutant embryos both fascicles are affected, as the vMP2/pCC fascicle collapses completely over the midline (100%,  $n=174$ ) and the dMP2/MP1 fascicle either does not extend or misroutes towards the muscle (89%,  $n=174$ ) (Fig. 4D). Slightly later, at stage 14, these phenotypes become even clearer. Wild-type embryos have two distinct fascicles, the outer dMP2/MP1 and the inner vMP2/pCC fascicles (Fig. 5F). In *gcm* mutants only the vMP2/pCC fascicle is present (68.5%,  $n=92$ ) (Fig. 5G), whereas in *robo1* mutants only the dMP2/MP1 fascicle extends normally (27.7%,  $n=94$ ) (Fig. 5H). In *gcm-robo1* double-mutants, there is a synergistic effect on both fascicles (Fig. 5I): the vMP2/pCC fascicle is completely collapsed over the midline (96%,  $n=156$ ), and the dMP2/MP1 fascicle is missing (97%,  $n=156$ ), meaning that it might have either misrouted out of the CNS or collapsed over the midline. These results demonstrate that loss of glia in a *robo1*-mutant background alters the formation of axonal trajectories from the beginning of pioneer-growth-cone extension. Thus, absence of functional glia in a *robo1* loss-of-function background synergistically affects axon guidance of both pioneer and follower neurons.

Our results also show that *robo1* is not required in the early stages of dMP2/MP1 fascicle extension. In fact, we find that whereas in *robo* mutants the vMP2/pCC fascicle is trapped in the midline, the dMP2/MP1 fascicle begins to extend normally with its characteristic curved trajectory (Fig. 5C). Slightly later in wild-type embryos, the dMP2/MP1 fascicle can be seen as it separates from the vMP2/pCC fascicle (Fig. 5E). In *robo1* mutants, the dMP2/MP1 fascicle extends its normal trajectory, and curves medially slightly more than in wild-type only at the most posterior end (Fig. 5J), possibly because of the lack of the ascending vMP2/pCC fascicle that the dMP2/MP1 axons normally encounter. Later still, at stage 14, the vMP2/pCC fascicle is always collapsed over the midline in *robo1* mutants, but the dMP2/MP1 fascicle can form normally (Fig. 5H). This indicates that the pioneer neurons dMP2 and MP1 do not require *robo1* during growth-cone guidance.

The finding that *robo1* is not required in the pioneer dMP2/MP1 fascicle contrasts with previous reports suggesting that *robo1* is required by all pioneer axons, despite the fact that *robo1* has not been shown to be expressed in the MP1 and dMP2 neurons (Kidd *et al.*, 1998a; Rajagopalan *et al.*, 2000; Simpson *et al.*, 2000b).

## Lateral displacement of axons upon ectopic expression of *robo2* requires glia

The ectopic expression of *robo2* in neurons that normally project over the medial fascicle is sufficient to displace their trajectory laterally to the central fascicle (Fig. 6E and Fig. 7B; Rajagopalan *et al.*, 2000; Simpson *et al.*, 2000a). This is one of the key experiments that demonstrates the nature of the Robo code. We wondered whether the response of axons to gain-of-function in the Robo code might require normal glial environment and function.

To test whether Robo2 can displace axons laterally along a longitudinal trajectory in embryos that lack glia and in which fascicles misroute over the midline, we targeted the expression of Robo2 in *gcm-robo1* double-mutant embryos (Fig. 6). In controls, ectopic expression of *robo2* can displace axons laterally and longitudinally in *robo1*-mutant embryos (Fig. 6G and Fig. 7C), indicating that *robo1* is not required for the displacement induced by *robo2* expression. In *gcm* mutants, the three FasII fascicles might be defasciculated but are distinctly present, and MP2 axons visualised with GFP project normally over the medial fascicle (78.5%,  $n=28$  hemisegments) (Fig. 6B). Ectopic expression of *robo2* in *gcm*-mutant embryos might cause: (1) lateral displacement in embryos that have a mild phenotype (the expressivity of the *gcm* phenotype is variable); (2) in embryos with a more severe phenotype, the axons that misexpress *robo2* project over a single, fused, FasII fascicle (35%,  $n=20$  hemisegments) (Fig. 6F); and (3) axons misroute to the muscle exiting the CNS (10%,  $n=20$  hemisegments) (Fig. 6F). Thus, axons that express *robo2* ectopically are not necessarily displaced over a lateral longitudinal trajectory in the absence of functional glia.

The effect of glial loss is exacerbated in severe *gcm-robo1* double-mutant embryos in which the longitudinal fascicles either collapse along the midline or misroute severely towards the periphery and exit the CNS (Fig. 6D and Fig. 7D). When the MP2 axons in these double mutants are visualised with GFP, they are seen to project all along the midline (63.3%,  $n=30$  segments) (Fig. 6D) or misroute out of the CNS (13.3%,  $n=60$  hemisegments). However, when MP2 axons in these embryos misexpress *robo2*, they do not extend longitudinally. Instead, they either do not grow (36.6%,  $n=166$  hemisegments) and remain stunted at the base of the commissures, or they leave the CNS and misroute towards the muscle (30.7%,  $n=166$  hemisegments) (Fig. 6H). Therefore, in the absence of glia as well as *robo1*, ectopic Robo2 cannot define a more lateral position for axons along a longitudinal pathway. Instead, the fact that longitudinal axons either do not grow or exit the CNS, means that axons respond to ectopic *robo2* by being repelled along other pre-existing major fascicles and that in the absence of glia there are not sufficient fasciculation cues for axons to extend longitudinally.

## CONCLUSIONS

- Longitudinal migration of LG does not require the Robo code cell-autonomously. Thus, a Slit gradient is not sufficient to define neuropile structure.
- Longitudinal axons collapse along the midline in *gcm-robo1* double mutants, which mimics loss-of-function of multiple Robo receptors.
- In *gcm-robo1* double-mutant embryos, axons that ectopically express *robo2* cannot extend longitudinally and they either do not grow or exit the CNS.

These findings mean that axons require LG to read the Robo code.

## DISCUSSION

### Glia are required for the reading of the Robo code by axons

We demonstrate here that interfering with neuron-glia interactions alters how axons respond to the Robo code. Thus, normal neuron-glia interactions are necessary for the correct functioning of the axonal Robo code.

We have shown that LG do not migrate normally during axon guidance in embryos mutant in the different *robo* genes. Instead, glial positions follow axonal pattern changes in these mutants. We have shown that neither *robo2* nor *robo3* are expressed in the LG during their longitudinal migration. Previously, we have shown that *robo1* is expressed in LG during their medial migration before axon guidance, but not later during longitudinal migration (Kinrade *et al.*, 2001). Furthermore, both *robo2* and *robo3* are still expressed in the absence of glia in *gcm*-mutant embryos, meaning that loss of glia does not cause loss of the *robo* code. Thus, these findings indicate that the lateral position of LG relative to the midline does not depend directly on Slit and the Robo code, but that it depends on interactions with axons and on other signalling mechanisms. This means that the Slit gradient is not sufficient to define all aspects of neuropile structure. Because glia are necessary for axon guidance, but longitudinal migration of glia is not controlled by Slit, axons receive multiple, Slit-dependent and Slit-independent inputs that define their lateral trajectories.

We have shown that lateral neuron-glia interactions influence how axons read the Robo code. We have shown that removing functional glia with *gcm* mutations in *robo1* loss-of-function mutants causes a severe, midline-collapse phenotype, comparable to the removal of multiple Robo receptors. This means that glia are necessary for the reading of the Robo code by axons. It also means that, in addition to Robo-code signalling on axons, LG are necessary to keep axons away from the midline. We have also shown that in *gcm-robo* embryos with severe, midline-collapse phenotypes, the ectopic expression of *robo2* does not redirect axons to a more lateral, but still longitudinal pathway. Instead, MP2 axons either do not grow at all or they leave the CNS and project towards the muscle. Either way, they do not project longitudinally. Thus, glia are also required for axonal displacement caused by gain-of-function of Robo receptors. This demonstrates that neuron-glia interactions along the longitudinal pathways are necessary for the function of the axonal Robo code, and to define and maintain the longitudinal trajectories of axons. Both follower and pioneer neurons require inputs from both the Robo code and glia to define their trajectories, thus, these two inputs are necessary for the establishment of longitudinal trajectories from the beginning and throughout axon guidance.

### Different requirements for the Robo code among pioneer neurons

Different pioneer axons have different requirements for glia and the Robo code. Loss of glia in *gcm* mutants prevents the dMP2/MP1 fascicle from descending longitudinally. Instead, loss of *robo* causes a very dramatic collapse on the midline of the pCC/vMP2 fascicle, whereas the dMP2 growth cone extends normally. The fact that *robo1* mutations do not affect early dMP2/MP1 pioneer growth-cone guidance might help explain why the intermediate (MP1) FasII fascicle is not affected in *robo* mutants. In fact, the intermediate fasII fascicle originates from neurons that never required *robo1*, and where *robo1* has not been shown to be expressed.

Later, when dMP2 fasciculates with pCC along the medial fascicle, it does require *robo* function. The fact that ectopic expression of *robo2* in dMP2 and vMP2 neurons causes displacement only to the intermediate fascicle and not more laterally, could be because the medial and intermediate fascicles become fasciculated together earlier (Hidalgo and Brand,



1997) and the ectopic expression of Robo2 prevents their defasciculation (Simpson *et al.*, 2000a). This is consistent with a primary role of Robo2 in fasciculation.

The differential requirement for *robo1* in the pioneer neurons also helps to define the different shape of their trajectories: the pCC axon extends very straight and parallel to the midline, and it clearly requires *robo1* whereas the dMP2 axons take a turn outwards before turning back inwards (Hidalgo and Brand, 1997): such a complex trajectory might not be achieved with a constant activation of Robo signalling, because this would ensure a constant distance from the midline, hence a parallel trajectory.

### Possible mechanisms involving glia and fasciculation

Our results have shown that the Robo code is insufficient to determine the longitudinal trajectories of axons, and that axons also sense a second input based on local interactions with glia. In the context of neuron-glia interactions, the Slit gradient could still be the driving force organising axonal patterns if the Robo code directly determines glial migration as well as axonal positioning. However, we have shown that LG do not express Robo receptors and that their migration depends on local and other cues. Thus, our data support the notion that axons receive two instructions that define their longitudinal trajectories: one in the form of signalling from Robo receptors, and one from interactions with LG. Furthermore, being an interactive system, glia can influence the outcome of the axonal Robo code.

The molecular mechanism by which glia steer the response of axons to midline signalling is unknown. We have shown that axons can cross the midline in the absence of glia and neuron-glia interactions despite the expression of multiple Robo receptors (Kinrade *et al.*, 2001; and this work) and that axons of interneurons that do not normally cross the midline either do not extend at all or misroute to the muscle when forced to overexpress *robo2* in *gcm-robo* double mutants. Perhaps Robo receptors are involved directly in fasciculation/adhesion between axons and glia. Although Robo receptors have been shown to function as homophilic binding molecules (Hivert *et al.*, 2002), they might also interact with other proteins on the glial cell membrane, causing different responses in the axon in the absence and presence of glia. Another possibility is that glia change growth-cone behaviour by influencing the small GTPases (Kuhn *et al.*, 1999) that function downstream of Robo (Patel and Van Vactor, 2002) so that, in the absence of glia, normal signalling by Robo is abrogated. Perhaps Robo receptors interact with tyrosine kinases (Wong *et al.*, 2003) in the growth cone that respond to interactions with glia and, in the absence of glia, the input to the Robo code is lost. Last, glia are likely to influence the molecular environment of extending axons. For example, loss of glia might change the relative concentration of heparan sulfate proteoglycans and other extracellular components that influence signalling by Robo (Johnson *et al.*, 2004; Steigemann *et al.*, 2004) and, presumably, other pathways involved in neuron-glia interactions. Together, it appears that the roles played by the Robos are dependent on context. If the cellular environment changes then the Robos cannot display the normal repulsive signalling function as a read-out of distance from the midline.

Whatever the mechanism, we conclude that axons require fasciculation cues present in both glia and axons to trace their trajectories. Thus, loss of fasciculation cues in *gcm-robo* double mutants prevents the longitudinal extension of *robo2*-expressing axons independently of midline repulsion. Given that axons do extend longitudinally in either *gcm* or *robo* single mutants, the loss of both glia and axonal fasciculation cues in *gcm-robo1* double mutants exacerbates the disorientation of the axons. In *gcm-robo1* double mutants, interneuron longitudinal axons can extend over the motoraxons that exit the CNS, which indicates that there they encounter fasciculation cues that they can adhere to. This is consistent with Robo2 playing a key role in fasciculation. It has been observed in other contexts that the

roles of the different Robo receptors vary greatly (Godenschewege *et al.*, 2002; Zlatic *et al.*, 2003) and that both Robo receptors and Slits have relevant roles in fasciculation and axonal growth (Hivert *et al.*, 2002; Hutson and Chien, 2002; Shu *et al.*, 2003). Our results support the notion that Robo2 and, perhaps, Robo3 are involved in fasciculation, and that fasciculation might also involve direct interactions with membrane proteins on the glial surfaces.

Given that glia are necessary for axon guidance and extension of axons in the correct direction (Hidalgo and Booth, 2000), it is paradoxical that the reciprocal interactions between axons and glia during extension and migration acquire directionality. One mechanism that confers some directionality is the non-autonomous maintenance of cell survival, which places an instructive positional pressure on cells. In fact, the survival of some LG depends on neurons. The descending fascicle of the dMP2 pioneer neuron extends the longest distance during axon guidance, and it does so completely surrounded and anticipated by glial cells (Hidalgo and Booth, 2000). Neuregulin Vein produced by the dMP2 neurons maintains the survival of a subset of these glia (Hidalgo *et al.*, 2001). Follower-neuron survival also depends on glial cells, but pioneer-neuron survival does not (Booth *et al.*, 2000). Because the pioneer neurons are the first neurons to navigate and trace out a scaffolding of the major axonal fascicles, their independence from glia for survival provides directionality during axon guidance (Booth *et al.*, 2000).

It is conceivable that in *gcm-robo1* double mutants the collapse of axons over the midline might correlate with an increase in neuronal apoptosis because there is extensive apoptosis in *gcm*-mutant embryos. We have shown previously that interfering with cell survival disrupts the lateral positions of axons. However, the induction of apoptosis cannot be the main cause of midline collapse in the absence of *robo1* and *gcm*. First, we observe axon-guidance defects in the pioneer axons, which do not undergo apoptosis in *gcm-robo1* double mutants. Second, complete lack of cell death in H99 embryos that also lack glia (*gcm-H99* double mutants) also induces severe midline crossing by axons. This means that neuronal apoptosis might be a correlation but not a cause of the *gcm-robo1* double-mutant phenotype. Instead, these findings support the notion that neuron-glia interactions maintain the survival of both cell types and restrict their positions.

### Glia during vertebrate axon guidance

In vertebrates, neuron-glia interactions might play an analogous role in the confinement of axonal trajectories and influencing the final outcome of other guidance cues. For example, the longitudinal axons of the spinal cord in vertebrates are also associated with a type of glial cells, oligodendrocytes, whose cell bodies are found all along the length of the spinal cord. As with ablation of glia in *Drosophila*, elimination of oligodendrocytes causes disruptions to axonal patterns, such as defasciculation and misrouting, which indicate that myelinating cells create a 'channel' along which axons extend (Schwab and Schnell, 1991). Similarly, motor axons also use mechanical cues, in the form of non-neuronal 'tubes', to find their trajectories during regeneration, implying that axons follow cues of diverse nature during pathfinding (Nguyen *et al.*, 2002). Regeneration experiments have also shown that re-growth of broken axonal tracts requires that oligodendrocytes are aligned in the direction of axonal growth, and it has been shown that transplanting glial cells to the sites of axonal injury can restore axonal trajectories and neuronal function (Ramon-Cueto *et al.*, 2000). These and other findings indicate that glia are also required for the formation of axonal tracts in vertebrates and that axons navigate by sensing multiple inputs from their environment. Slits and Robos are conserved in vertebrates. Understanding the complexity of cellular and molecular events that regulate Slit-Robo mechanisms and neuron-glia interactions during axon guidance is necessary for the therapeutic implementation of CNS repair.

To conclude, we have shown that in the absence of glia and, most dramatically, in a *gcm-robo1* double-mutant background, axons lose their ability to extend longitudinally in conditions of either loss- or gain-of-function of the Robo code. Our results show that neuron-glia interactions are necessary for the correct functioning of the axonal Robo code.

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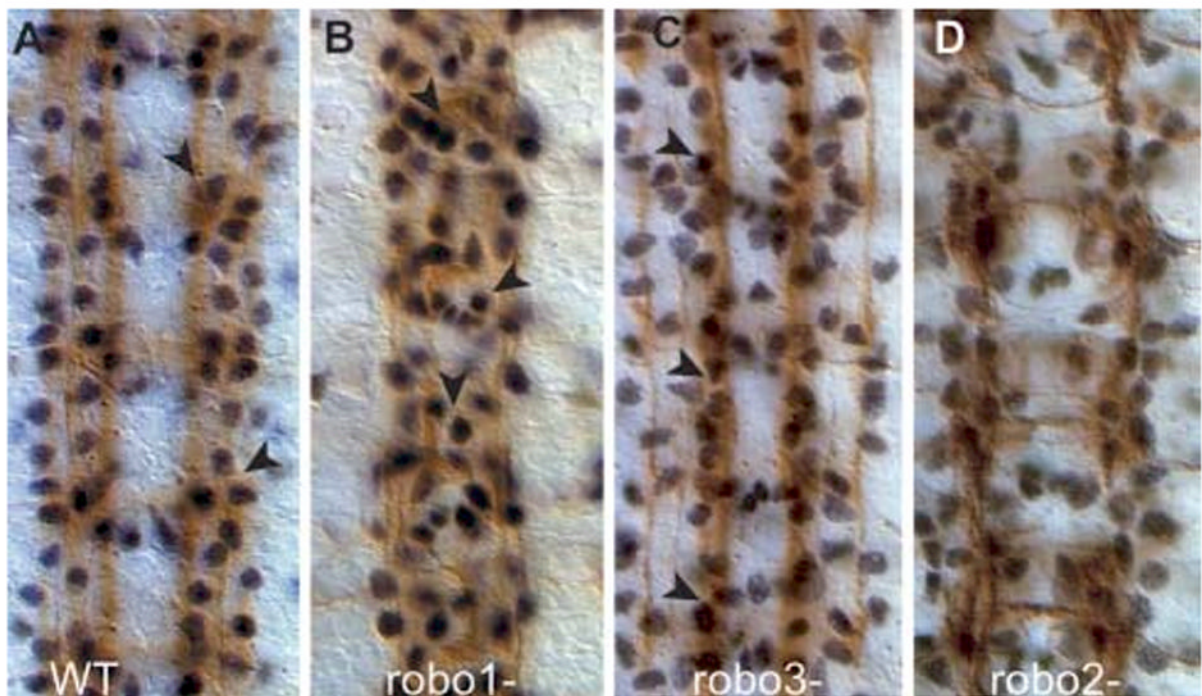
We thank members of our group for helpful discussions. We are grateful to Akira Chiba, Barry Dickson, Corey Goodman, Iowa Hybridoma Bank, Roger Jacobs, Matthias Landgraf, Susana Romani, Guy Tear and Andrew Travers for antibodies and flies. E.F.V.K. is supported by a BBSRC studentship; A.H.'s work is funded by a Wellcome Trust CDF, MRC CEG and EMBO YIP.

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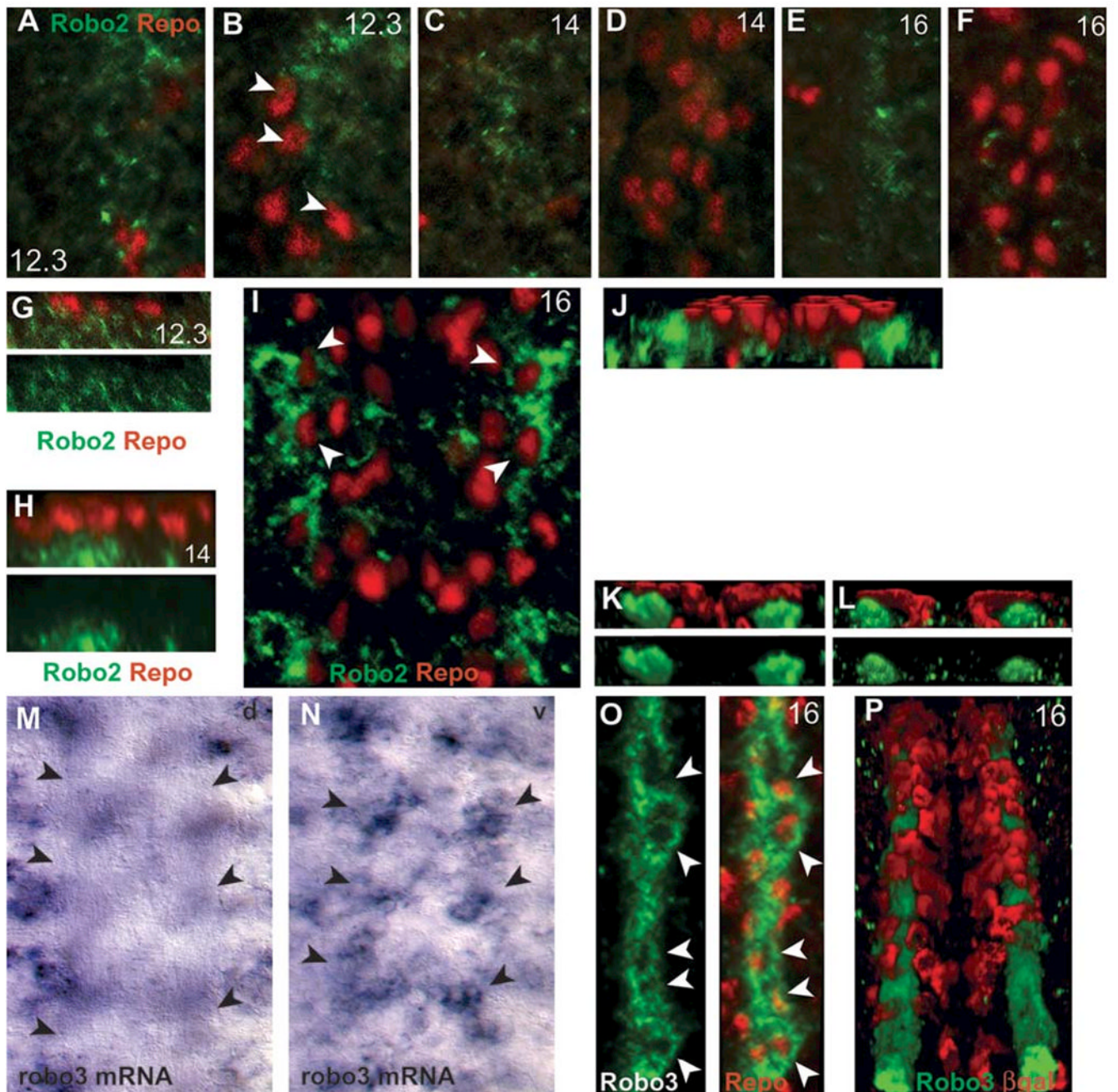
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**Fig. 1.**

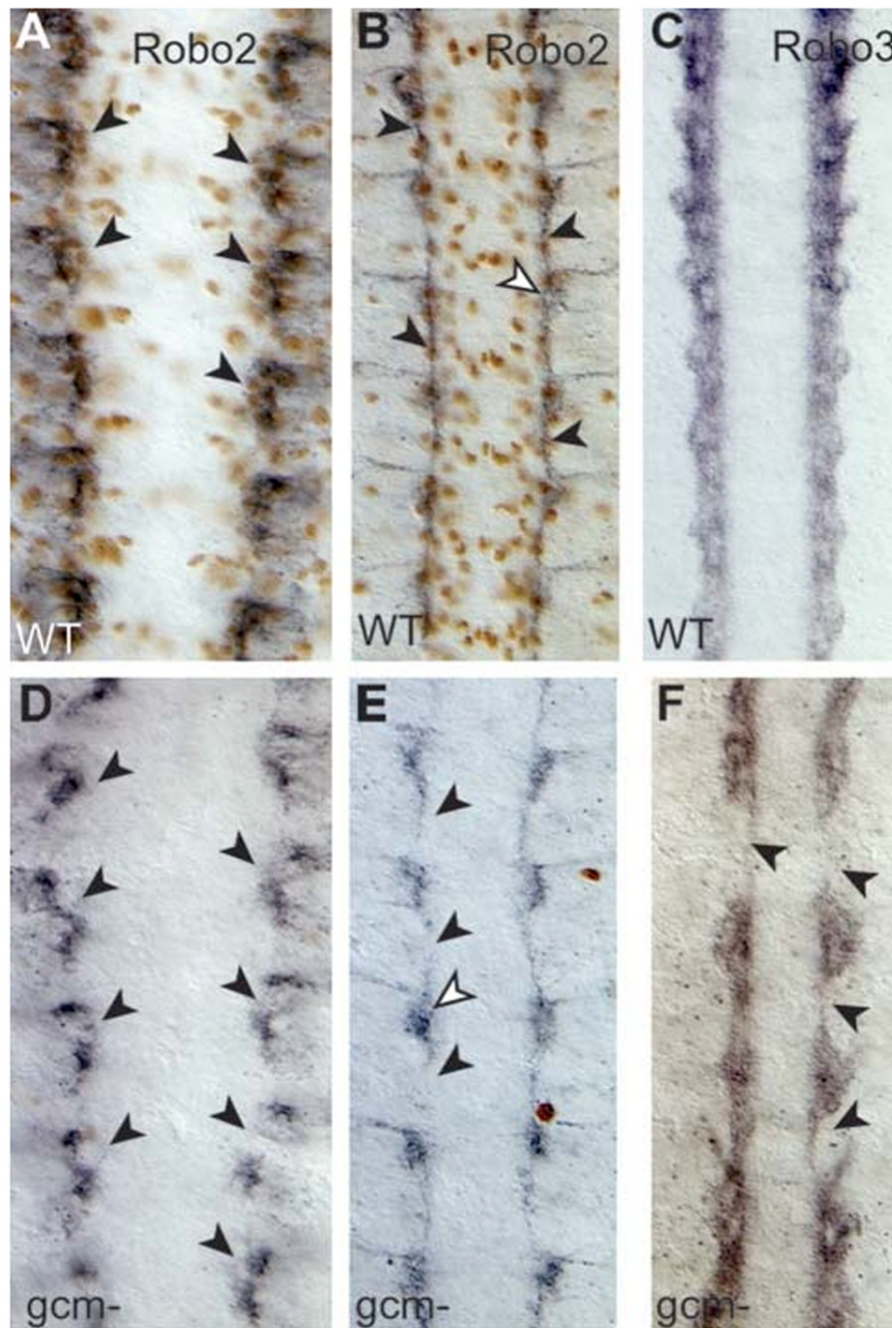
**LG are misplaced in Robo-mutant embryos.** Embryos stained with the glial nuclear anti-Repo (black) and neuronal FasII (brown) antibodies at stage 16. (A) In wild-type embryos glia (arrowheads) are deployed in discrete rows over the three FasII fascicles. (B) In *robo1*-mutant embryos, some glia migrate over the midline (arrowheads) and some remain over the lateral axons. (C) In *robo3*-mutant embryos, many glia are displaced medially and they overlie the fascicle formed by the fused medial and central fascicle (arrowheads). (D) In *robo2* mutants the lateral FasII fascicle and glial patterns are only slightly disorganised. Anterior is to the top.



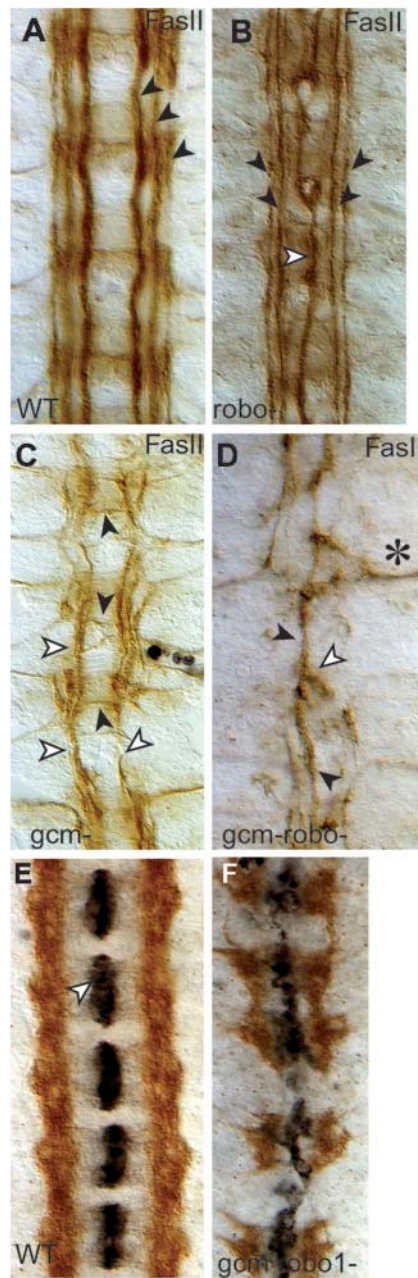
**Fig. 2.**  
**Robo2 and Robo3 are not expressed in LG.** LG nuclei are visualised in wild-type embryos with the glial nuclear marker anti-Repo (red) (A-K,O), and LG cytoplasm with anti- $\beta$ gal on the LG-lacZ reporter (red) (L,P). (A-J) Temporal profile of Robo2 distribution (green) in the ventral nerve cord. Ventral (A) and dorsal (B) longitudinal and (G) sagittal sections of the same stage-12.3 embryo. Robo2 is first detected at stage 12.3 lateral to the LG and mostly in a ventral plane under the LG. Some LG nuclei appear partly surrounded by Robo2 signal (arrowheads in B), but in a transverse view (G) the glial nuclei overlie Robo2 signal. At stage 14, from a frontal view Robo2 signal is ventral (C) and physically separate from the dorsal, Repo-positive LG nuclei (D,H). (G,H) Single channel images of anti-Robo2 are also

shown (green). (E,F,I) At stage 16, Robo2 is lateral to the LG but it appears to surround some Repo-positive LG nuclei (arrowheads in I), however these are in separate planes: ventral Robo2 (E); dorsal Repo glia (F); and a whole-stack projection (I). (J) A tilted, 3D-reconstruction section of the same embryonic stack as in (I), showing that Robo2 lies ventrally and laterally to the LG. (M,N) Distribution of *robo3* transcripts by *in situ* hybridisation with DIG probes. (M) There is no signal in the dorsal plane of the neuropile, axons are seen clear of signal by Nomarski optics (arrows). (N) All DIG signal is in a ventral plane (arrows) below the neuropile. (K,L,O,P) Distribution of anti-Robo3 (green) and either Repo or anti- $\beta$ gal (red) in the LG at stage 16. (O,P) From a frontal view, Robo3 appears to surround the nuclei of LG (arrowheads, whole-stack projections). (O) Single channel image shown in green next to merged image. (K) 3D transverse view of the embryo in (O) reveals that the glial nuclei are dorsal to and separated from the Robo3 axons (bottom, single channel view of Robo3, green; top, merge). (P) LG are visualised with anti- $\beta$ gal (red) using the LG-lacZ reporter overlies the Robo3 axons (tilted 3D reconstruction, whole stack). (L) 3D transverse view of the same embryo as in (P) revealing that the  $\beta$ gal glial signal overlies the Robo3 axonal bundles, but it is not coincident with them. All images were processed with Volocity (Improvision). Anterior is up. A-F,G,H show one hemisegment; O just over one hemisegment; the midline is to the left. I shows one segment; M,N, 4 segments; P, 5 segments; G,H, lateral sections; J,K,L, cross-section views. A-F are projections of sections restricted to or excluding glial nuclei; I,O,P are projections of the whole stack of sections through the neuropile.



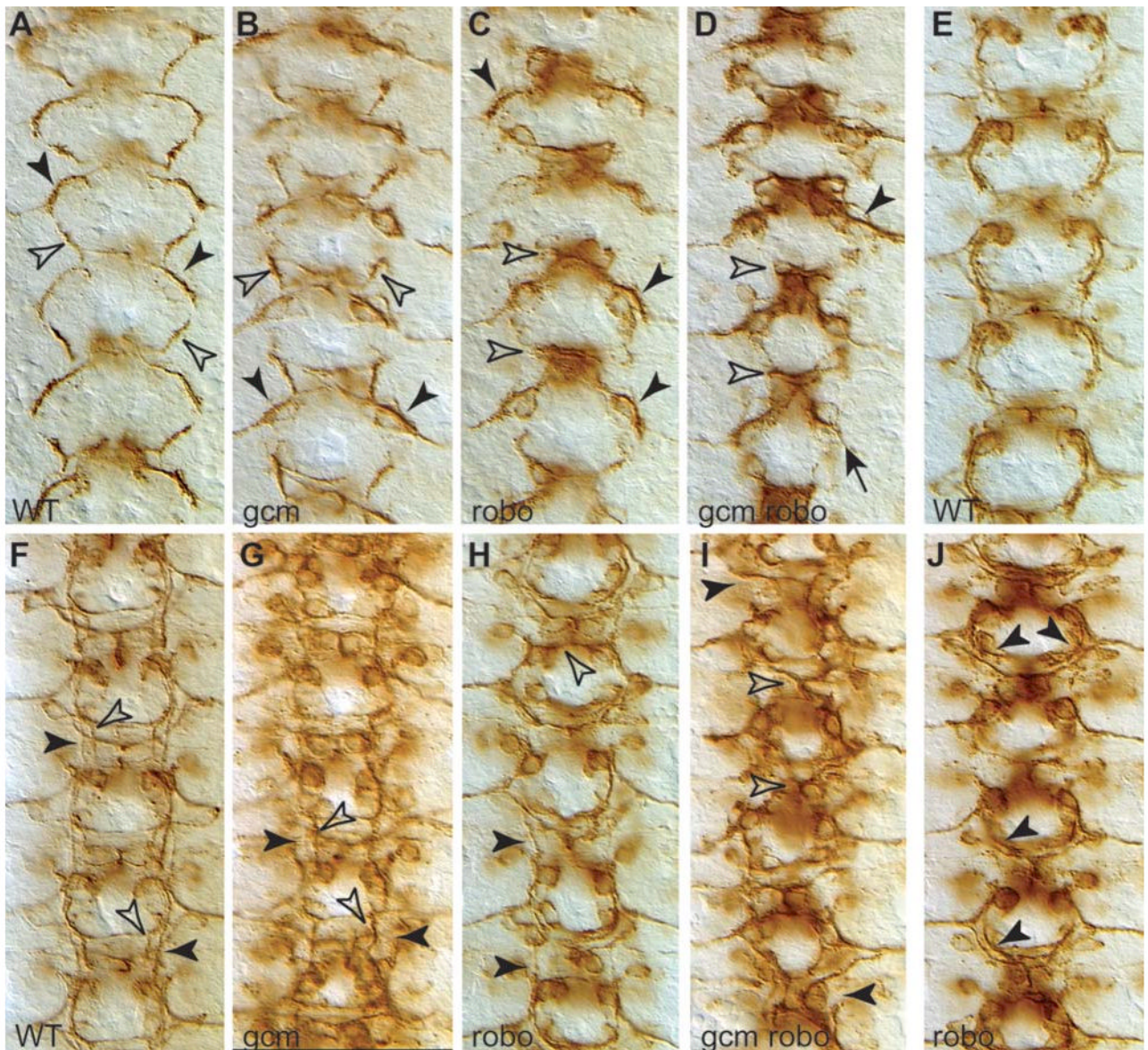


**Fig. 3.**  
**Robo2 and Robo3 in the absence of glia.** Anti-Robo2 (black) and anti-Repo (brown) in wild-type embryos (A,B) and *gcm* mutant (D,E) embryos. Anti-Robo3 in wild-type (C) and *gcm* mutant (F) embryos. At stage 13 in wild-type embryos, Robo2 clusters lie under the LG (A). Robo2 is present but disarranged in *gcm* mutants (D). At stage 16, Robo2 longitudinal axons are frequently missing in *gcm* mutants (E, arrowheads), but Robo2 signal (white arrowhead) is present at the base of the commissures. (F) Robo3 is still present in *gcm* mutants, but the longitudinal projection of axons is severely compromised (arrowheads). Anterior is up.



**Fig. 4.** **Midline collapse of longitudinal axons in *gcm-robo1* double-mutant embryos.** Axons are visualised with anti-FasII antibodies (brown) at stage 16 (A-D). (A) Wild-type embryo. (B) *robo1* mutant. Note that the medial fascicle runs along the midline (white arrowhead) whereas the outer two fascicles are well formed and run parallel to the midline (black arrowheads). (C) In *gcm* mutants, axons sporadically misroute across the midline (black arrowheads) and the lateral fascicles can be fused into one that runs parallel to the midline along the central trajectory (white arrowheads). This embryo has also been stained with anti-Repo (in black, there are a few glia present). (D) In *gcm-robo1* double-mutant embryos, axons can fuse (white arrowhead) into one single fascicle running along the midline (black arrowhead) or exit the CNS (asterisk) and grow towards the muscle, but no axons run along

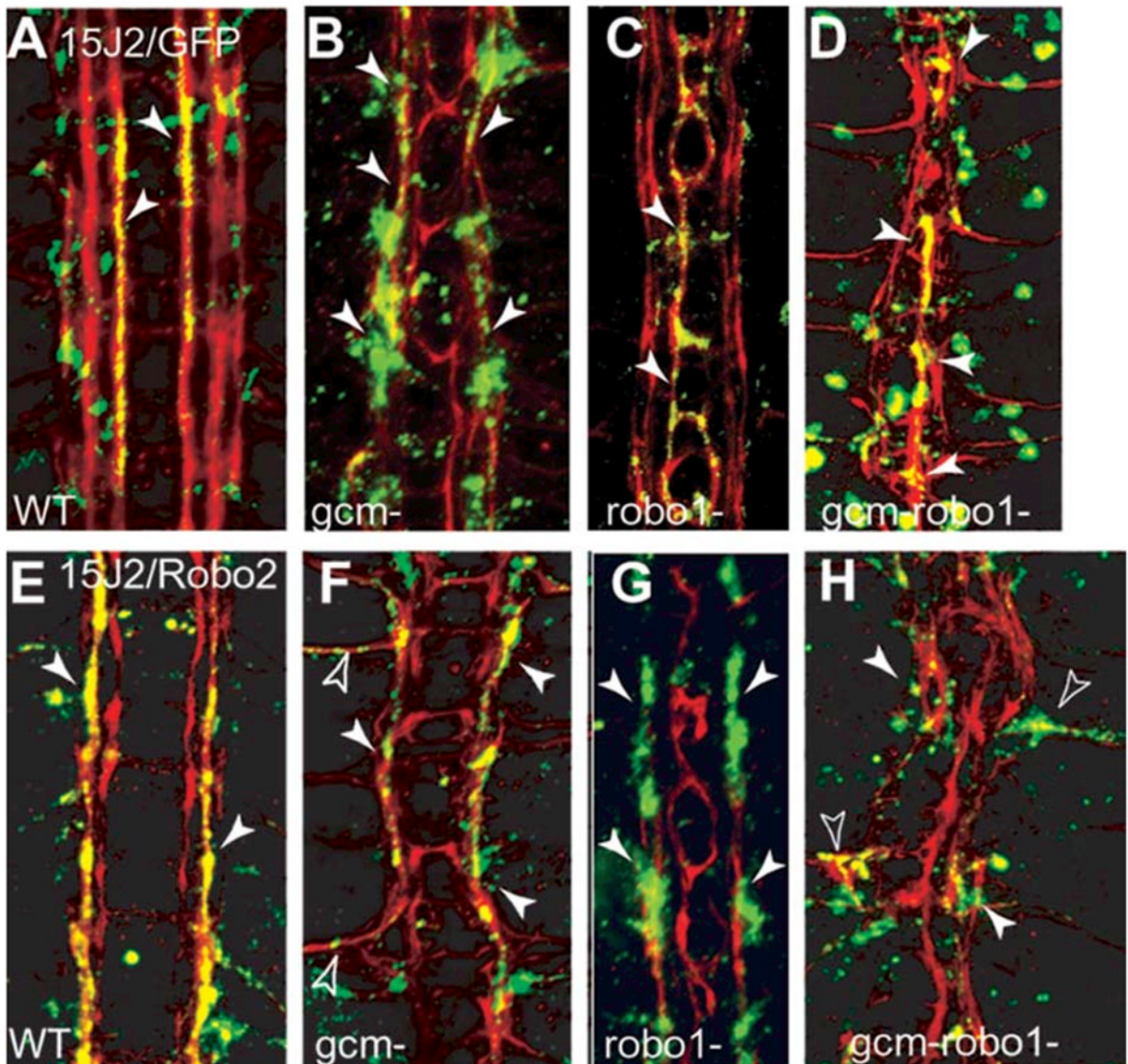
the lateral trajectories. (E,F) Axons are stained with anti-Robo3 (brown) and midline glia with anti-Slit (black) at stage 16. (E) Robo3 axons never cross the midline in wild-type embryos, and anti-Slit stains the midline glia in each segment. (F) *gcm-robo1* double mutant. Slit (black) is still produced by the midline glia, although the midline glia are disorganised. No longitudinal axons remain, instead axons cross the midline despite expressing *robo3*. Anterior is up.



**Fig. 5.**

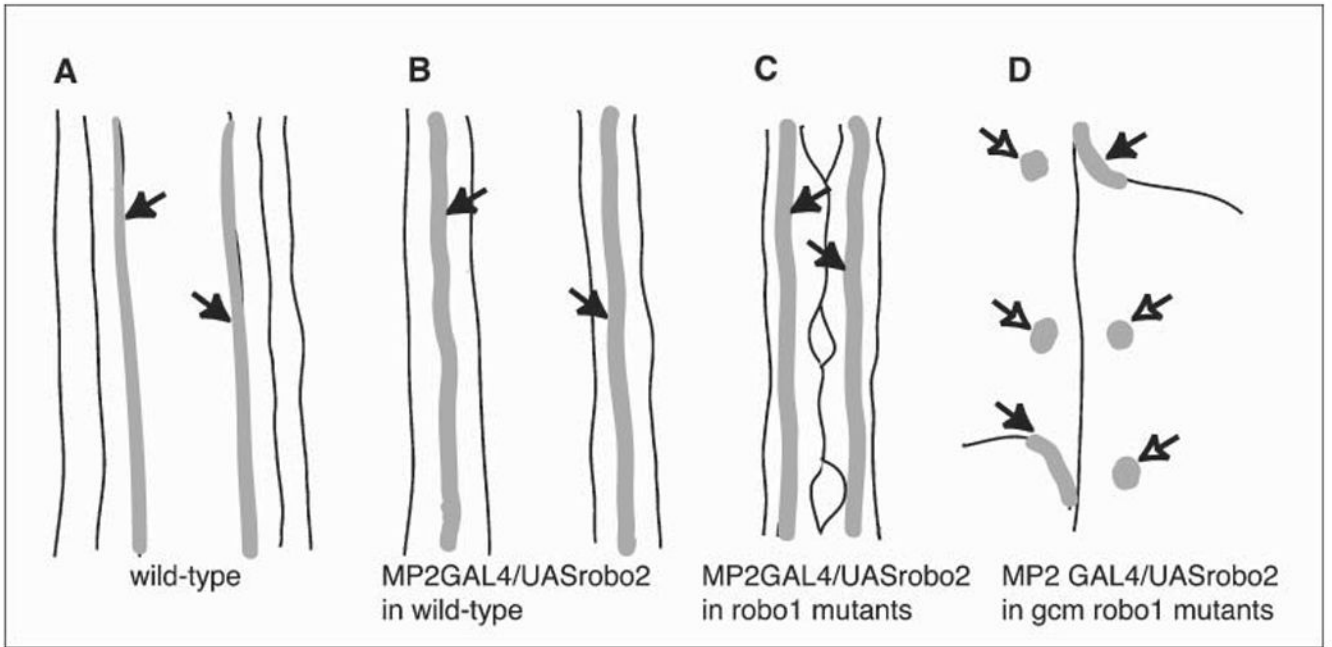
**Axonal defects occur from the earlier phases of pioneer axon extension.** Pioneer axons visualised with 22c10 antibodies in wild-type (A,E,F), *gcm*-mutant embryo (B,G), *robo1* mutant (C,H,J) and *gcm-robo1* double mutant (D,I) at stages 12.1 (A,B,C,D); 13 (E,J) and 14 (F,G,H,I). (A) In wild-type embryos, the descending dMP2/MP1 axons (black arrowhead) meet the ascending vMP2/pCC axons (white arrowhead) at the end of stage 12. (B) In *gcm* mutants, the dMP2/MP1 fascicle misroutes to the muscle, so its descending trajectory is missing (black arrowheads), but the vMP2/pCC fascicle is unaffected (white arrowheads). (C) In *robo1* mutants, the pCC/vMP2 fascicle does not grow but is trapped in the midline (white arrowheads), whereas the dMP2/MP1 fascicle begins its normal descending trajectory. This is also seen slightly later, at stage 13, (J) where the dMP2/MP1 fascicle extends normally all the way down (arrowheads) and only then curves towards the midline as it fails to encounter the ascending vMP2/pCC fascicle, which is missing (compare to E, wild-type). (D) In *gcm-robo1* double-mutant embryos, the pCC/vMP2

fascicle is completely collapsed on the midline (white arrowheads) and the dMP2/vMP2 fascicle is either missing or misrouted to the muscle (black arrowheads). (F) At stage 14, the two fascicles, dMP2/MP1 (black arrowheads) and pCC/vMP2 (white arrowheads) are clearly distinguishable. (G) In *gcm* mutants, only the pCC/vMP2 fascicle is present (white arrowheads) and the dMP2/MP1 fascicle is missing (black arrowheads). (H) In *robo1* mutants, the pCC/vMP2 fascicle collapses on the midline (white arrowhead) but the dMP2/MP1 fascicle (black arrowhead) is normal. (I) In *gcm-robo1* double mutants, the pCC/vMP2 fascicle collapses on the midline, but the dMP2/MP1 fascicle is not discernible. Anterior is up.



**Fig. 6.**  
**Ectopic expression of *robo2* does not displace the MP2 axons along lateral longitudinal tracts in the absence of glia.** FasII axons are visualised in red at stage 16. The MP2 axons are visualised with anti-GFP (green) by the expression of 15J2/UASGAPGFP in multiple genotypes (A-D). Colocalisation of GFP and FasII is seen in yellow. Ectopic expression of *robo2* with 15J2/UAS*robo2-myc*, visualised with anti-Myc (green) antibodies in embryos of different genotypes (E-H). Colocalisation of Myc and FasII is seen in yellow. (A) Wild-type. The MP2 axons run parallel to the midline along the medial FasII fascicle at this stage (arrowheads, 15J2/UASGAPGFP). (E) When expression of *robo2* is driven ectopically in the MP2 axons, their trajectories are displaced to the central fascicle (arrowheads, 15J2/UAS*Robo2myc*). (B) In *gcm*-mutant embryos, the MP2 axons still run along the medial fascicle (arrowheads, *gcm*<sup>ΔP1</sup>/*gcm*<sup>ΔP1</sup>; 15J2/UASGAPGFP). (F) When *robo2* is expressed ectopically in the MP2 neurons in *gcm* mutants (*gcm*<sup>ΔP1</sup>/*gcm*<sup>ΔP1</sup>; 15J2/UAS*robo2myc*), the MP2 axons stay within the fused, single fascicle (arrowheads). This *gcm*-mutant embryo is phenotypically more severe than in B, where the fascicles are defasciculated but not fused. (C) In *robo1* mutants, the MP2 axons run on the medial fascicle, collapsed along the midline

(arrowheads, *robo1/robo1; 15J2/UASGAPGFP*). (G) Expression of *robo2* in the MP2 axons in *robo1* mutants (*robo1/robo1; 15J2/UASrobo2myc*) displaces the MP2 axons laterally onto the central fascicle (arrowheads). (D) *gcm-robo1* double mutant with midline collapse of the longitudinal fascicles, in which the MP2 axons run along the midline (arrowheads, *gcm<sup>ΔPI</sup> robo/gcm<sup>ΔPI</sup> robo; 15J2/UASGAPGFP*). (H) *gcm-robo1* double mutant embryo with midline collapse of the longitudinal fascicles, expressing *robo2* in the MP2 axons (*gcm<sup>ΔPI</sup> robo1/gcm<sup>ΔPI</sup> robo1; 15J2/UASrobo2myc*): the MP2 axons leave the CNS and extend towards the periphery (empty arrowheads) or they do not grow at all (white arrowheads). There are no MP2 axons projecting longitudinally in these embryos. Anterior is up.



**Fig. 7.**

**Schematic of the behaviour of MP2 longitudinal axons in different genetic backgrounds.** In wild-type embryos, MP2 axons extend along the medial fascicle (A, arrows) and are displaced laterally on expression of *robo2* (B, arrows). Lateral displacement still takes place in *robo1* mutant embryos (C, arrows). However, if embryos are double mutant for *robo1* and *gcm* and, therefore, lack functional glia, MP2 axons either do not grow (D, empty arrows) or they misroute out of the CNS towards the muscle (D, arrows).