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## Dscam1-mediated self-avoidance counters Netrin-dependent targeting of dendrites in *Drosophila*

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### SUMMARY

Dendrites and axons show precise targeting and spacing patterns for proper reception and transmission of information in the nervous system. Self-avoidance promotes complete territory coverage and non-overlapping spacing between processes from the same cell [1, 2]. Neurons that lack *Drosophila* Down syndrome cell adhesion molecule 1 (Dscam1) show aberrant overlap, fasciculation, and accumulation of dendrites and axons, demonstrating a role in self-recognition and repulsion leading to self-avoidance [3–11]. Fasciculation and accumulation of processes suggested that Dscam1 might promote process spacing by counterbalancing developmental signals that otherwise promote self-association [9, 12]. Here we show that Dscam1 functions to counter sensory neuron dendritic targeting signals provided by secreted Netrin-B and Frazzled, a netrin receptor. Loss of Dscam1 function resulted in aberrant dendrite accumulation at a Netrin-B expressing target, whereas concomitant loss of Frazzled prevented accumulation and caused severe deficits in dendritic territory coverage. Netrin misexpression was sufficient to induce ectopic dendritic targeting in a Frazzled-dependent manner, whereas Dscam1 was required to prevent ectopic accumulation, consistent with separable roles for these receptors. Our results suggest that Dscam1-mediated self-avoidance counter extrinsic signals that are required for normal dendritic patterning, but whose action would otherwise favor neurite accumulation. Counterbalancing roles for Dscam1 may be deployed in diverse contexts during neural circuit formation.

### Results and Discussion

#### Elimination of dendritic self-avoidance reveals focal dendrite targets for da sensory neurons

Dendritic territory formation involves both specific dendritic targeting and the spreading of branches across their target region. These processes are likely to be closely coordinated in time but under at least partially distinct molecular control by attractive and repulsive cues.

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#### SUPPLEMENTAL INFORMATION

Supplemental Information includes four figures and Supplemental Experimental Procedures and can be found with this article online at XXXXXXXXX

To examine how these steps in territory formation are coordinated in individual arbors, we assessed the consequences of loss of *Dscam1* on dendritic targeting of dendritic arborization (da) neurons. Sister dendrites of wild-type neurons rarely cross each other and thus spread evenly over their field (Figure 1A)[13], while dendrites of *Dscam1* mutant sensory neurons aberrantly cross, fasciculate, and accumulate (Figures 1B and 1C) [9–11]. Sites of accumulation could correspond to preferred targets of sister dendrites that are revealed absent the spacing constraints of self-avoidance [9]. To characterize the targets that may be responsible for dendrite accumulation we focused on class III da neurons (ddaA and ldaB) that project dendrites near a peripheral sensory organ, the lateral chordotonal organ (LCHO) (Figure 1D) [9, 13]. Labeling of *Dscam1* clones, together with anti-HRP, anti- $\beta$ PS-integrin, or anti- $\alpha$ -tubulin to label surrounding cells, revealed accumulations of dendrites at a distal region of CHO cap cells, a type of support cell that links CHO sensory neurons to the body wall (Figures 1E–1F and S1A) [14]. For simplicity, we refer to this targeted zone of cap cells as the lateral focus (LF). Branch density was higher at LF than the rest of the field in *Dscam1* clones (Figures 1G and 1H). Importantly, CHOs are positioned differently in thoracic and abdominal body segments, nevertheless dendrites of *Dscam1* mutant sensory neurons accumulated at CHO cap cells regardless of their precise position (Figure S1B), strongly suggesting that it is this specific cell type, rather than other nearby cells, that directs targeting. Accumulations of dendrites in *Dscam1* mutant clones represented excessive targeting rather than an abnormal ectopic targeting event, since single dendrites of wild-type ddaA neurons also targeted precisely to LF at each larval instar (Figures 1I–1K and S1C–S1F). These data together suggest that *Dscam1* prevents excessive targeting of dendritic branches at peripheral cells that normally provide targeting instructions to sensory dendrites.

### Targeting of sensory dendrites requires Frazzled

*Dscam1* shows homophilic binding activity [15, 16], and in addition can serve as a receptor for the secreted netrins [17–19]. However, since dendrites showed robust targeting both in the presence and absence of *Dscam1*, we inferred that a signal for an alternate receptor is presented at LF. In a candidate screen of possible attractive receptors for LF targeting, we identified another netrin receptor, Frazzled/Deleted in Colorectal Cancer (Fra/DCC) [20–25]. Whereas wild-type ddaA clones showed robust targeting of a single branch to LF (Figures 2A and 2C), *frazzled*<sup>3</sup> (*fra*<sup>3</sup>) mosaic analysis with a repressible cell marker (MARCM) clones showed significant disruptions in targeting (Figures 2B and 2C). Importantly, coincident loss of Frazzled suppressed the accumulation phenotypes seen in *Dscam1*<sup>-/-</sup> neurons (*Dscam1*<sup>-/-</sup> = 100% accumulated, n=8; *Dscam1*<sup>-/-</sup>, *fra*<sup>-/-</sup> = 0% accumulated, n=13; Figure 2D). These data are consistent with a cell-autonomous role for Frazzled in sensory dendrite targeting and suggest that *Dscam1* prevents excessive Frazzled-dependent targeting by sister dendrites.

We asked whether *Dscam1* and Frazzled control larger-scale patterning of sensory neuron territories in addition to precise targeting. Modified Sholl analysis of the anterior-posterior arborization of ddaA indicated that wild-type arbors are polarized toward LF and have a relatively sparse anterior arborization (Figure 2E). We observed an anterior shift of ddaA territories (increased anterior and decreased posterior arborization) in *fra* mutant neurons (Figure 2E). By contrast, anterior coverage was reduced and arborization toward LF was increased in *Dscam1* mutant neurons (Figure 2E). Loss of anterior coverage could reflect an early role for *Dscam1* in segregation of anterior and posterior processes [5], or *Dscam1*-mediated responses to other, non-netrin, extrinsic cues. Thus, Frazzled and *Dscam1* are both required for proper anterior-posterior positioning of ddaA sensory arbors, but with opposite polarities. Removing activity of both *Dscam1* and Frazzled diminished both anterior and posterior regions of the arbor, and territory footprints were strongly reduced relative to wild-type (Figures 2D, 2E, and S2A). The simplest interpretation of these phenotypes is that

Dscam1 and Frazzled have separable functions in dendritic targeting. It appears that Frazzled promotes directional spreading of dendrites, presumably to sources of guidance cues near LF, while Dscam1 controls territory positioning by promoting dendrite separation and counterbalancing Frazzled-mediated targeting activity through branch-branch repulsion.

We next performed misexpression experiments to identify the domains of Frazzled that are required for precise dendritic targeting. Overexpression of full-length Frazzled in otherwise wild-type neurons did not alter branch targeting at LF (Figures 2F and S2B). By contrast, misexpression of *FraΔC*, which lacks the cytoplasmic tail and acts as a dominant negative [26], strongly suppressed targeting (Figures 2F and S2B). Three conserved domains within the Frazzled cytoplasmic tail - P1, P2 and P3 - are differentially required for axon guidance downstream of Frazzled, but have not been examined for activity in dendrite targeting [24, 26, 27]. We assessed requirements in dendrites by testing the ability of Frazzled transgenes lacking different combinations of these domains to rescue *fra* mutant dendrite targeting phenotypes [26]. Transheterozygous *fra<sup>3</sup>/fra<sup>4</sup>* mutant larvae showed deficits in targeting similar to those seen in *fra* MARCM clones (Figure 2G). Targeting was rescued by full-length *UAS-Fra-myc*, *UAS-FraΔP1-myc*, and *UAS-FraΔP1ΔP2-myc* (Figure 2G). By contrast, *UAS-FraΔP3-myc* did not significantly restore targeting in *fra<sup>3</sup>/fra<sup>4</sup>* mutant larvae ( $p > 0.05$ , Figure 2G). We observed qualitatively similar levels of expression of the different transgenes (data not shown). Thus, sequences within the P3 domain of Frazzled, but not within P1 or P2, appear to be important for dendritic targeting.

### Netrin-B is essential for Frazzled-dependent sensory dendrite targeting

Two closely related *Drosophila* netrins are encoded by the *Netrin-A* (*NetA*) and *Netrin-B* (*NetB*) genes [28, 29]. We analyzed targeting in larvae lacking one or both netrins using *189Y-Gal4>UAS-mCD8::GFP* to label class III neurons. Targeting to LF in *189Y-Gal4>UAS-mCD8::GFP* animals was robust (Figures 3A and 3E), but was strongly disrupted in *NetAB<sup>Δ</sup>* mutant animals (Figures 3B and 3E), implicating netrins in precise targeting. We examined dendrite targeting in mutant animals lacking either *Netrin* gene alone and found that NetB, but not NetA, is required for targeting (Figures 3C–3E). Dendrite targeting in *NetAB<sup>Δ</sup>* mutant larvae was not significantly different from *fra<sup>3</sup>/fra<sup>4</sup>* mutants ( $p > 0.05$ ), suggesting that the activity of netrins during targeting occurs largely, or exclusively, through Frazzled.

We next assessed *NetB* expression in animals carrying a modified *NetB* locus that generates a transmembrane tethered NetB protein tagged with c-myc (*NetB<sup>tm</sup>*) [20]. *NetB<sup>tm</sup>* reflects endogenous NetB distribution in the central nervous system, but because it is not secreted is more closely associated with membranes of cells in which it is produced [20]. *NetB<sup>tm</sup>* was enriched in scolopale cells of LCHO in late embryos (Figure S3A) and accumulated in LCHO cap cells in both abdominal and thoracic segments throughout all larval stages (Figure 3F and S3B–S3D). By contrast, using *in situ* hybridization we detected NetA at segment borders (Figure S3E; [29]) but not in LCHO. Notably, dendritic targeting to LF was disrupted in *ΔNetA NetB<sup>tm</sup>* larvae compared to *ΔNetA NetB<sup>myc</sup>* controls (Figure 3G). Targeting was not significantly different in *ΔNetA NetB<sup>tm</sup>* animals compared to *NetAB<sup>Δ</sup>* null animals ( $p > 0.05$ ), together suggesting that secretion of NetB is required for targeting. These data suggest that LCHO cap cells provide the source of secreted NetB that underlies da neuron dendrite targeting.

Although NetA was dispensable for LF targeting, we cannot exclude a role for NetA in da neuron dendritic patterning outside of LF. Notably, we found that misexpression of Unc5, a repulsive netrin receptor [23], in all da neurons using *109(2)80-Gal4* caused defects in dendritic morphology, including disruption of LF targeting (Figures 3H–3J), and aberrant exclusion of dendrites from segment boundaries (Figures S3F and S3G). These results raise

the possibility that other sources of Netrins, perhaps including NetA at segment borders, could also be important for da neuron dendrite patterning.

### Ectopic expression of Netrin directs dendritic targeting and reshapes sensory neuron territories in a Frazzled-dependent manner

To test whether localized netrin expression is also sufficient for da neuron dendrite targeting, and to further test the respective roles of *Dscam1* and *Frazzled*, we developed an ectopic targeting assay. We used *8-183-Gal4* to drive netrin expression in larval oenocytes, which are clusters of 5–6 large hepatocyte-like cells in the lateral body wall [30]. We assessed directional growth of the *ddaA* dendritic arbor by measuring the relative enrichment of dendrites in either the ‘LF’ quadrant posterior to the cell body or an adjacent ‘oenocyte’ quadrant directly ventral to the cell body (Figures 4A–4D). Wild-type *ddaA* dendrites were polarized toward LF and projected more branches to the LF quadrant than the oenocyte quadrant (Figures 4A and 4B). *NetB* misexpression in the oenocytes (*8-183-Gal4>NetB*) did not significantly alter LF quadrant arborization (Figures 4B, 4D, and S4A), however, in the oenocyte quadrant, overall branch number and proportion of arbor increased relative to control, while the median distance of the closest branch to the oenocytes decreased (Figures 4B, 4D, and S4A–S4B). Both *NetA* and *NetB* had qualitatively similar effects on dendritic targeting in this assay (Figure S4C), suggesting that the selective endogenous role for *NetB* arises due to differential expression rather than differential activity. Dendritic territories of *ddaA* failed to adjust toward ectopic sources of *NetB* in *fra* mutant larvae (Figures 4E and 4F). Furthermore, while we did not find evidence for dendrite accumulation at the oenocytes in *Dscam1* mutant larvae (Figure 4G), or in density plots generated from mutant clones (Figure 1H), *8-183-Gal4* driving *NetB* induced dendritic accumulation at oenocytes in larvae mutant for *Dscam1* (Figure 4H). Thus, localized expression of netrins in peripheral tissue was sufficient to direct the targeting of dendrites and reshape arbor territories, and dendrites from a single cell appeared able to respond to multiple such cues. Since opposite patterning defects arose when guidance activity was provided in the absence of *Frazzled* or *Dscam1*, our data support a role for *Frazzled* in attractive responses to netrin and a role for *Dscam1* in self-repulsion and counterbalancing of *Frazzled*-dependent attraction in normal and ectopic targeting.

### Conclusions

Dendritic targeting is an important component of neural circuit assembly in diverse systems [25, 31–35]. Our data suggest that targeting cues can generate a patterning problem for complex dendrites that must be overcome by self-avoidance signaling. Our findings together suggest a model in which repulsive self-avoidance counters netrin-based attractive signals to permit both precise targeting and full and even territory coverage by sister dendrites. Dendritic branches will be repelled only when their number at any given target becomes excessive. We suggest that control of dendritic targeting by a combination of attractive and self-repulsive cues provides a system for the regulation of branch positioning and spacing that can be implemented dynamically in growing dendrites, and without regard for the exact number and position of guidance cue sources. Indeed, our overexpression results suggest that one advantage of this mechanism is that counterbalancing can be executed in response to multiple, spatially distinct, targeting cues impinging on a single arbor. Conceivably, such action could greatly diversify the shapes taken by dendritic fields. It is worth noting that, in da neurons, counterbalancing is implemented in arbors that are largely restricted to two-dimensional growth. However, in a similar fashion, *Dscam1* appears to prevent local accumulations of dendritic and axonal arbors in regions of the brain where arbors elaborate in three dimensions [5, 8]. These observations raise the possibility that repulsive self-

avoidance functions as a counterbalancing cue more broadly, perhaps revealing a general role for *Dscam1*-based self-recognition and repulsion during the formation of neural circuits.

Recent work has shown that *Dscams* mediate self-avoidance in *Drosophila* and mammals. In *Drosophila*, discrimination between self and non-self dendrites depends on extensive alternative splicing of the *Dscam1* ectodomain and selective binding between identical isoforms [15, 16, 36]. Other *Drosophila* *Dscams*, as well as vertebrate *DSCAM*, lack the same degree of molecular diversity but nevertheless appear to prevent inappropriate associations and fasciculation between like-type isoneuronal or heteroneuronal processes. *Drosophila* *Dscam2* mediates axonal tiling and prevents inappropriate pairings of like-type postsynaptic elements in the visual system [37, 38]. Vertebrate *DSCAM* and the related *DSCAML1* prevent neurite fasciculation and clumping of cell bodies in classes of amacrine cells and retinal ganglion cells [12, 39]. *DSCAM* and *DSCAML1* have been proposed to function in mammalian neurons to mask as yet unknown cell-type specific adhesive cues that promote clumping and fasciculation of processes [12, 39]. An important future goal is to understand how these opposing cues are integrated by dendrites or axons. In sensory neurons, the integration of self-avoidance and targeting cues is apparently biased in favor of repulsion since the initiation of *Dscam1*-mediated self-avoidance appears non-permissive for Frazzled-dependent branch targeting.

In da neuron dendrite targeting, roles for *Dscam1* appear to be separable from those for Netrin and Frazzled, since several of our experiments support roles for *Dscam1* in self-avoidance, and Netrin together with Frazzled in attractive guidance. In addition to Frazzled/DCC, *Dscam1* in *Drosophila* and *DSCAM* in mammals have been shown to act as receptors for netrins during axon guidance [17–19]. In the *Drosophila* CNS, dendritic attraction by the action of netrins appears to depend largely, or entirely, on Frazzled [34], similar to our findings for da neurons. NetA and Frazzled also instruct the development of the v'ch1 chordotonal organ cap cell, which, in turn, influences dendrite orientation of associated chordotonal neurons [40].

Our results demonstrate how analysis of *Dscam1* mutant phenotypes can uncover new sources of signals that affect neurite patterning. Evidence for targeting cues acting on complex dendrites may be difficult to detect because of the dispersed nature of many dendritic fields and constraints imposed by other cues that may exclude arborization, such as repulsive self-avoidance and tiling. Interfering with self-avoidance appears to provide one approach for identifying preferred targets of neurites, which may then be interrogated further to identify patterning cues for highly branched dendritic arbors. The precision of targeting shown by da neuron dendrites should prove useful for dissecting signaling mechanisms of targeting and self-avoidance.

### Highlights

1. The guidance cue Netrin-B and Frazzled control da sensory neuron dendritic targeting
2. *Dscam1* prevents aberrant dendrite accumulation at Netrin-B expressing targets
3. NetB is sufficient for ectopic dendritic targeting in wild-type or *Dscam1*<sup>-/-</sup> neurons
4. Self-avoidance counterbalances uniform responses of sister dendrites to guidance cues

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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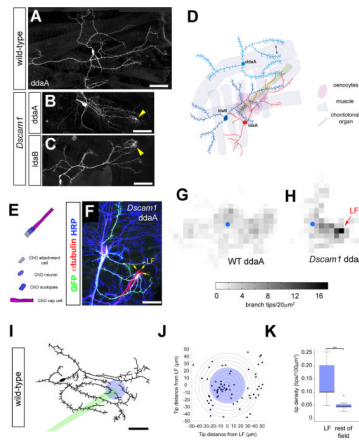
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**Figure 1. Targeting of *Dscam1* mutant neurons to chordotonal organs**

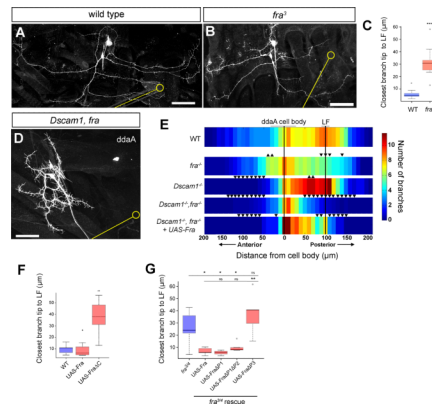
- (A) Wild-type MARCM clone of the class III neuron *ddaA*. Sister dendrites show robust self-avoidance.
- (B) *Dscam1*<sup>21</sup> mutant MARCM clone of class III neuron *ddaA* shows branch overlap and dendrite accumulation (yellow arrowhead).
- (C) *Dscam1*<sup>23</sup> mutant MARCM clone of class III neuron *ldaB* shows branch overlap and dendrite accumulation (yellow arrowhead).
- (D) Schematic of *da* neurons and nearby non-neuronal cells, including muscle, oenocytes, and chordotonal organs. Epidermal cells upon which dendrites grow are not included in the schematic.
- (E) The lateral chordotonal organ (LCHO) consists of a distal cap cell, a scolopale cell, neuron, and proximal attachment cell.
- (F) Labeling of a *Dscam1*<sup>21</sup> mutant *ddaA* MARCM clone with anti-GFP, anti- $\alpha$ -tubulin, and anti-HRP reveals accumulation at the distal margin of LCHO in abdominal segments, termed the lateral focus (LF; yellow arrows).
- (G) Density plot of all *ddaA* dendritic termini for wild-type (WT; n=8) MARCM clones. The average numbers of branch endings per cell that terminate within each 20 $\mu$ m<sup>2</sup> square are plotted, with higher terminal density indicated by increasing grayscale. Blue circle indicates the position of the cell body.
- (H) Density plot of *ddaA* dendritic termini for *Dscam1* MARCM clones (n=4). The average numbers of branch endings per cell that terminate within each 20 $\mu$ m<sup>2</sup> square are plotted as in (G). Blue circle indicates the position of the cell body. The accumulation of dendrite tips at LF in *Dscam1* mutant clones is indicated by a red arrow.
- (I) Tracing of a wild-type *ddaA* neuron labeled with *189Y-Gal4, UAS-mCD8::GFP*. A green bar indicates the trajectory of the chordotonal organ, LCHO. A blue circle with a 25 $\mu$ m radius is centered at LF.
- (J) Dendritic tips of *ddaA* in wild-type larvae are enriched within a 25 $\mu$ m radius of LF when branch tips of multiple cells (n=11) are plotted with respect to LF.
- (K) The density of branch tips in a circle of radius 25 $\mu$ m is significantly greater than the rest of the dendritic field.

All scale bars = 50 $\mu$ m.

Datasets are presented in boxplots as median (thick line), quartiles Q1–Q3 (25%–75% quartiles; shaded box), and data in 1.5 $\times$  quartile range (dashed bars). Data points outside these ranges are indicated as open circles.

\* =  $p < 0.05$ ; \*\* =  $p < 0.01$ ; \*\*\* =  $p < 0.001$  by Wilcoxon rank-sum test

See also Supplemental Figure S1



**Figure 2. Frazzled is required for sensory neuron dendrite targeting**

(A) A wild-type MARCM *ddaA* neuron showing targeting of branches to the chordotonal organ. The chordotonal organ and LF are indicated by a yellow line and yellow circle, respectively in this, and subsequent, panels.

(B) A *fra*<sup>3</sup> MARCM clone of *ddaA* showing disrupted targeting to LF.

(C) Plot of the distance of the closest branch tip to LF in wild-type (WT, n=8) and *fra*<sup>3</sup> (n=9) mutant clones.

(D) Dendrites of *Dscam1*<sup>21</sup>, *fra*<sup>4</sup> double mutant MARCM clone lack enrichment of dendritic termini at LF, and show extensive dendritic clumping near the cell body.

(E) Horizontal Sholl analysis showing mean branch density values at 10µm intervals for wild type (n=8), *fra* (n=8), *Dscam1* (n=5), *Dscam1*<sup>21</sup>, *fra*<sup>4</sup> (n=14) and *Dscam1*<sup>21</sup>, *fra*<sup>4</sup> + *UAS-Fra* (n=6) MARCM clones of *ddaA*. Upwards-pointing arrowheads denote significant increase in branch density relative to control (p < .05), while downwards-pointing arrowheads denote significant decrease in branch density (p < .05) as compared to topmost control strip.

(F) Quantification of the single closest branch tip to LF in control (n=8), *UAS-Fra* (n=7), and *UAS-Fra*ΔC (n=6) FLP-out clones reveals no effect of *UAS-Fra* on targeting, but a disruption of targeting caused by *UAS-Fra*ΔC. The pan-da neuron driver *109(2)80-Gal4* was used in these experiments.

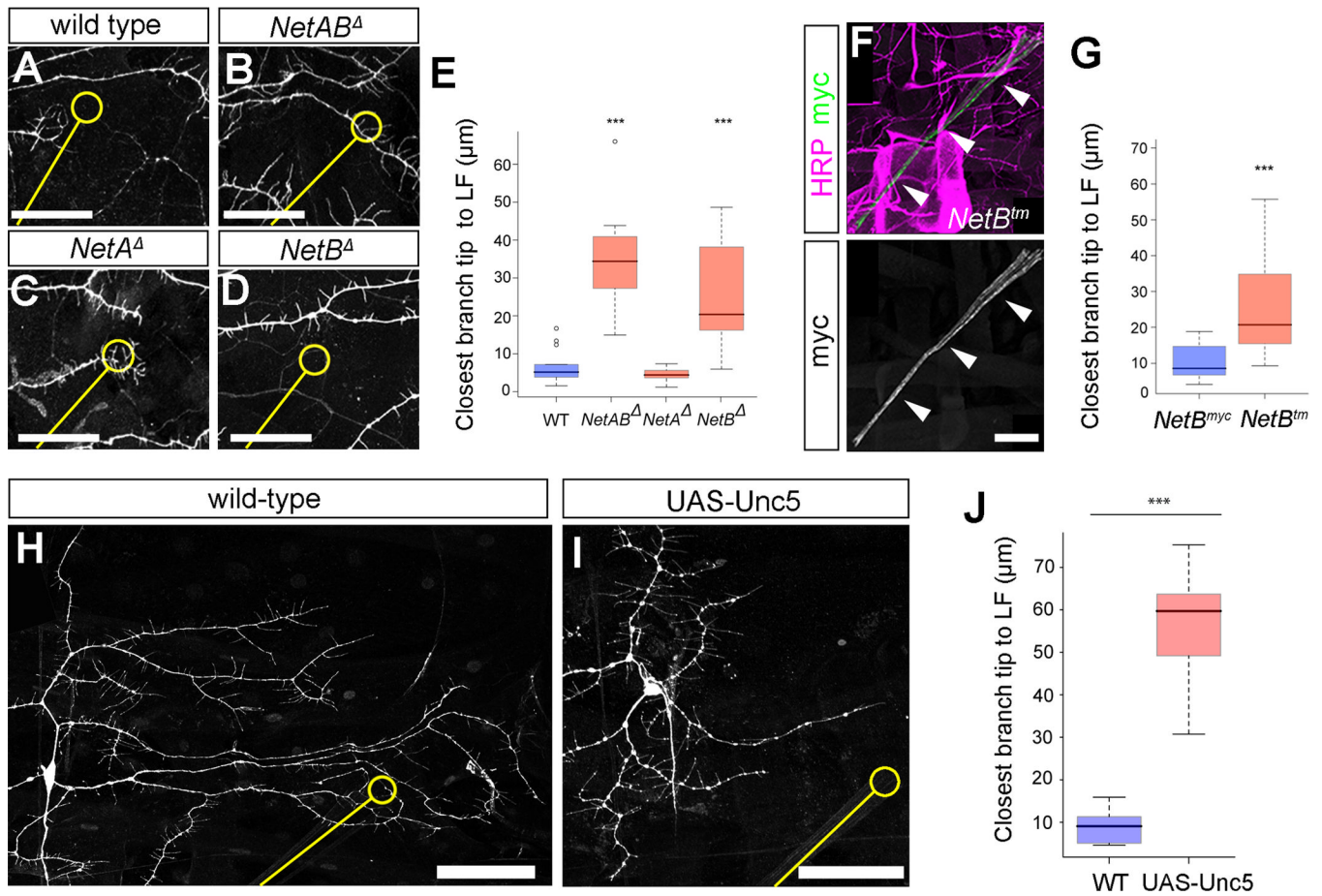
(G) Quantification of closest branch tip to LF in FLP-out rescue experiments in *fra*<sup>3</sup>/*fra*<sup>4</sup> mutant (n=9) background. Introduction of *UAS-Fra* (n=5), *UAS-Fra*ΔP1 (n=3), *UAS-Fra*ΔP1ΔP2 (n=5) significantly rescues the targeting defect seen in *fra*<sup>3</sup>/*fra*<sup>4</sup> mutant larvae, while introduction of *UAS-Fra*ΔP3 (n=5) does not. *109(2)80-Gal4* was used as a driver in these experiments.

Datasets are presented in boxplots as median (thick line), quartiles Q1–Q3 (25%–75% quartiles; shaded box), and data in 1.5× quartile range (dashed bars). Data points outside these ranges are indicated as open circles. In (E), \* = p < .05 by Wilcoxon rank-sum test (no further delineations of significance are presented for clarity of presentation).

\* = p < 0.05; \*\* = p < 0.01; \*\*\* = p < 0.001 by Wilcoxon rank-sum test.

All scale bars = 50µm

See also Supplemental Figure S2



**Figure 3. Function of Netrins in peripheral dendritic targeting and expression of Netrin-B** (A–D) Analysis of targeting of class III da neuron *ddaA* (marked by *189Y-Gal4*, *UAS-mCD8::GFP*) in (A) WT (n=13), (B) *NetAB $\Delta$*  (n=7), (C) *NetA $\Delta$*  (n=7) and (D) *NetB $\Delta$*  (n=13) third instar larvae. A–D show representative images of area surrounding LF. The location of the chordotonal organ, labeled in the HRP channel (not shown), is indicated by a yellow line, and the region corresponding to LF indicated by a yellow circle. Targeting is disrupted in *NetAB $\Delta$*  and *NetB $\Delta$*  mutant larvae.

(E) The distances of the closest major branch tip to LF in WT, *NetAB $\Delta$* , *NetA $\Delta$* , and *NetB $\Delta$*  animals.

(F) *NetB<sup>tm</sup>* levels are highly enriched in the chordotonal organ of third instar larvae (arrowheads).

(G) Quantification of closest tip to LF in  $\Delta$ *NetA*, *NetB<sup>myc</sup>* (n=10) and  $\Delta$ *NetA*, *NetB<sup>tm</sup>* (n=17) larvae.

(H) Morphology and chordotonal organ targeting of single wild-type FLP-out clone of *ddaA*.

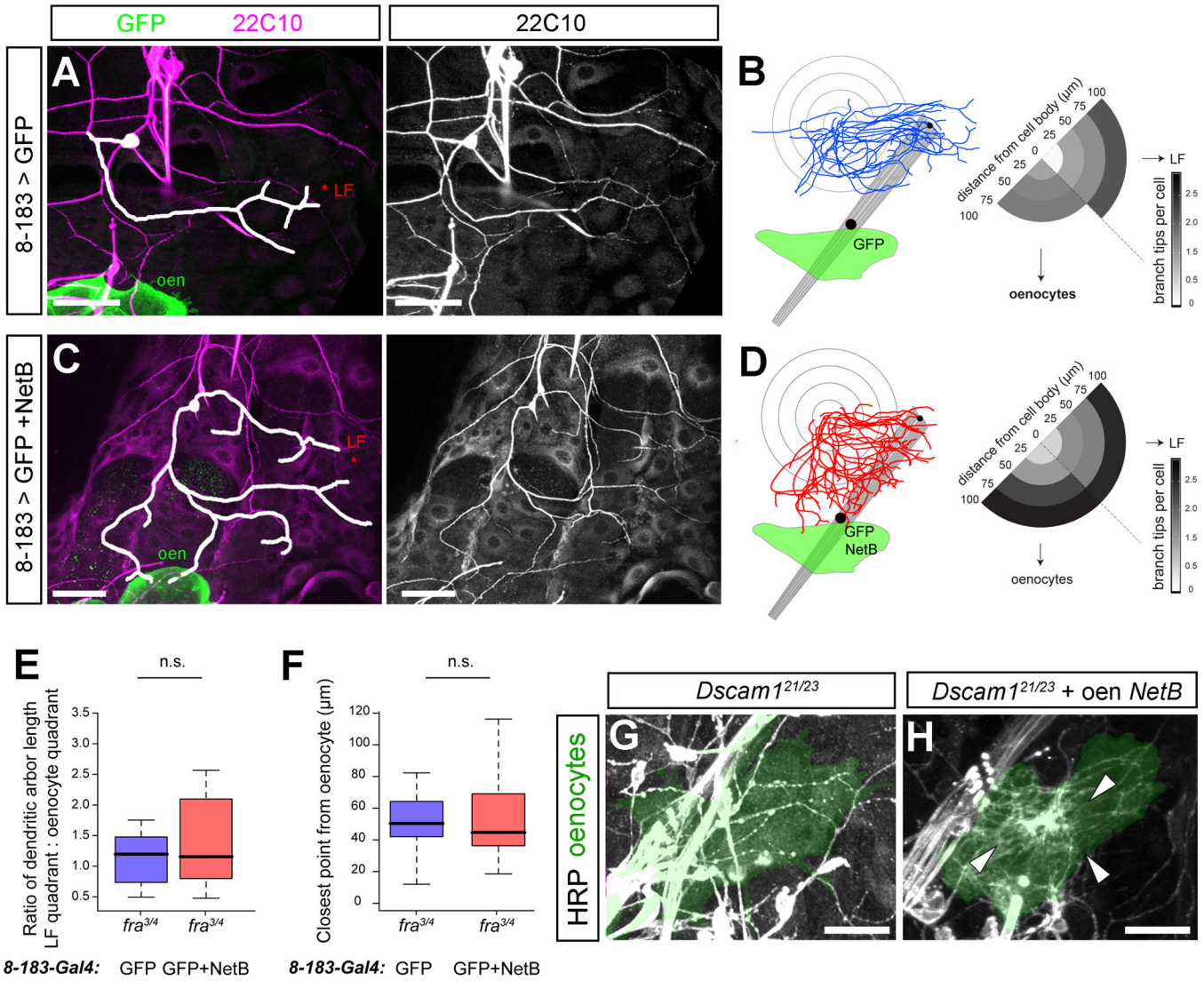
(I) Lack of chordotonal targeting of *ddaA* neuron as revealed by a single FLP-out clone expressing *Unc5* under the control of *109(2)80-Gal4*.

(J) Plot of the closest branch tip to LF in wild-type (WT, n=10) and *UAS-Unc5*-expressing (n=6) *ddaA* neurons.

All scale bars = 50 $\mu$ m.

Datasets are presented in boxplots as median (thick line), quartiles Q1–Q3 (25%–75% quartiles; shaded box), and data in 1.5 $\times$  quartile range (dashed bars). Data points outside these ranges are indicated as open circles.

\* =  $p < 0.05$ ; \*\* =  $p < 0.01$ ; \*\*\* =  $p < 0.001$  by Wilcoxon rank-sum test  
See also Supplemental Figure S3



**Figure 4. Ectopic expression of Netrin-B generates new dendritic targets**

(A) The dendritic arbor of *ddaA* in a wild-type larva, visualized by anti-Futsch/22C10 staining. The ventral and posterior portion of the arbor of *ddaA* is traced in white.

*mCD8::GFP* (GFP) is expressed in oenocytes (oen) driven by *8-183-Gal4*.

(B) A plot of multiple registered *ddaA* arbors (n=12) with concentric circles placed 25, 50, 75, and 100μm from the cell body. Dendritic intersections were segregated into two quadrants, one centered towards LF (LF quadrant) and one centered towards the oenocytes (oenocyte quadrant).

(C) Dendritic arbor morphology of *ddaA* upon expression of *mCD8::GFP* and NetB under the control of *8-183-Gal4*.

(D) A plot of *ddaA* dendritic arbors (n=12) upon misexpression of *mCD8::GFP* and NetB in the oenocytes. Dendrites are plotted as in (B).

(E) Quantification of the ratio of total arbor length within the quadrant centered on LF and the quadrant centered on the oenocytes reveals no change when NetB is expressed under the control of *8-183-Gal4* in *fra<sup>3</sup>/fra<sup>4</sup>* animals.

(F) Quantification of the closest point of each arbor to the oenocytes reveals no change when NetB is expressed under the control of *8-183-Gal4* in *fra<sup>3</sup>/fra<sup>4</sup>* animals.

(G) Dendrites do not accumulate at oenocytes (shaded in green) in *Dscam1<sup>21</sup>/Dscam1<sup>23</sup>* mutant larvae (n=3 animals examined).

(H) Dendrites accumulate at oenocytes (arrows) in *Dscam1<sup>21</sup>/Dscam1<sup>23</sup>* mutant larvae when NetB is expressed under the control of *8-183-Gal4* (n=6 animals examined).

All scale bars = 50µm

Datasets are presented in boxplots as median (thick line), quartiles Q1–Q3 (25%–75% quantiles; shaded box), and data in 1.5× quartile range (dashed bars).

\* =  $p < 0.05$ ; \*\* =  $p < 0.01$ ; \*\*\* =  $p < 0.001$  by Wilcoxon rank-sum test.

See also Supplemental Figure S4