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Arl13b in Primary Cilia Regulates the Migration and Placement of Interneurons in the Developing Cerebral Cortex

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SUMMARY

Coordinated migration and placement of interneurons and projection neurons lead to functional connectivity in the cerebral cortex; defective neuronal migration and the resultant connectivity changes underlie the cognitive defects in a spectrum of neurological disorders. Here we show that primary cilia play a guiding role in the migration and placement of postmitotic interneurons in the developing cerebral cortex and that this process requires the ciliary protein, Arl13b. Through live imaging of interneuronal cilia, we show that migrating interneurons display highly dynamic primary cilia and we correlate cilia dynamics with the interneuron’s migratory state. We demonstrate that the guidance cue receptors essential for interneuronal migration localize to interneuronal primary cilia, but their concentration and dynamics are altered in the absence of Arl13b. Expression of Arl13b variants known to cause Joubert syndrome induce defective interneuronal migration, suggesting that defects in cilia-dependent interneuron migration may in part underlie the neurological defects in Joubert syndrome patients.

INTRODUCTION

The appropriate placement of neurons in the cerebral cortex, achieved through a coordinated process of oriented neuronal migration, is critical for the emergence of functional neural circuitry underlying higher order brain functions. Neuronal migration enables distinct classes of neurons to navigate from their sites of birth in the progenitor zones to their final laminar and areal destinations within the cerebral cortex (Ayala et al., 2007; Marin et al., 2010; Rakic, 1972; Kwan et al., 2012). An efficient, selective intracellular response to extracellular signaling cues is fundamental in coordinating these diverse migratory processes. Primary cilia are found on cortical neuronal progenitors and neurons and may play a guiding role in the construction of the cerebral cortex (Arellano et al., 2012; Bishop et al., 2007; Han and Alvarez-Buylla, 2010; Händel et al., 1999; Lee and Gleeson, 2011; Wilson et al., 2012). The potential significance of cilia function for cortical development and function is evident in various developmental brain disorders, such as Joubert, Meckel-Gruber, Orofaciodigital, and Bardet-Biedl syndromes (commonly referred to as ciliopathies), where disrupted cilia function and the resultant changes in cortical formation are thought to underlie the cognitive deficits associated with these syndromes (Hildebrandt et al., 2011).

The primary cilium may play a role in the progression of cortical neuronal development. The basal bodies, which reside at the base of the cilium and are required for its formation, are located in the leading process of migrating neurons and are hypothesized to be involved in leading process stability and nuclear translocation (Métin et al., 2008; Higginbotham and Gleeson, 2007; Solecki et al., 2004). Recent studies using ciliogenic mutants indicate that ependymal cilia and primary cilia activity in the neurogenic zones of the developing cortex are required for the appropriate proliferation of progenitors and generation of neurons (Amador-Arjona et al., 2011; Besse et al., 2011; Breunig et al., 2008; Willaredt et al., 2008). Although these studies demonstrate the importance of cilia activity in progenitor dynamics, little is known about how neuronal cilia may influence the migration and differentiation of distinct classes of neurons during corticogenesis. We therefore aimed to identify the essential functions of primary cilia in neuronal migration and differentiation in the developing cerebral cortex.

To study the role of cilia in neuronal migration and differentiation, we used a mouse genetic model in which cilia function can be impaired in a temporal and neuronal type-specific manner. Arl13b, a small GTPase of the Arf/Arl family, is specifically localized to cilia (Caspar et al., 2007). Deletion of Arl13b in mouse results in impaired cilia, which lose their ability to function as conveyors of critical extracellular signals (Caspar et al., 2007). In humans, mutations in ARL13B result in Joubert syndrome (Cantagrel et al., 2008). Thus, neuronal type and developmental stage-specific genetic manipulation of Arl13b function provides a model to disrupt the primary function of cilia in distinct classes of neurons during corticogenesis and evaluate how this affects neuronal migration, placement, and the formation of cerebral cortex. Toward this goal, we conditionally disrupted primary cilia...
Figure 1. Projection Neuron Placement and Axonal Outgrowth in Nex-Cre; Arl13b\textsuperscript{Lox/Lox} Cortex

(A–D) P0 somatosensory cortex of Arl13b\textsuperscript{Lox/+}; Nex-Cre (A and C) or Arl13b\textsuperscript{Lox/Lox}; Nex-Cre (B and D) mice were immunolabeled for Cux1 (layers II–IV), Ctip2 (layer V), Brn1 (layers II–V) and Tbr1 (layer VI). Neuronal positioning was not altered by Arl13b deletion in projection neurons. 

(E and F) Serial coronal sections spanning the rostro-caudal extent of E16 Arl13b\textsuperscript{Lox/+}; Nex-Cre; Tau-mGFP (E) or Arl13b\textsuperscript{Lox/Lox}; Nex-Cre; Tau-mGFP (F) cortex were immunolabeled for mGFP. No differences in the extent of migration of GFP\textsuperscript{+} projection neurons were seen. Major cortical axonal tracts also appear normal in mutants.
function in postmitotic projection neurons and interneurons and define a role for primary cilia activity in the oriented migration and placement of interneurons.

RESULTS

Conditional Deletion of Arl13b in the Developing Cortical Neurons

Arl13b is expressed in the primary cilia of postmitotic cortical neurons during embryonic and postnatal development (Figures S1A–S1H available online). We conditionally ablated Arl13b in distinct subsets of postmitotic neurons of the developing cerebral cortex using the Cre-Lox recombination system (Su et al., 2012). We specifically targeted the two major populations of cortical neurons using the Nex-Cre or the Dlx5/6-Cre-IRESCGFP (Dix5/6-CIE) transgenic lines, which express Cre in postmitotic radially migrating projection neurons or in postmitotic tangentially migrating subcortically-derived interneurons, respectively, and confirmed cell type specific deletion of Arl13b (Figures S1I–S1P) (Goebbel et al., 2006; Stenman et al., 2003; Wu et al., 2005). We also incorporated a Tau-Lox-STOP-Lox-mGFP allele to label these neurons and their projections with mGFP (Hippenmeyer et al., 2005). At E14 and E16, we found the extent of radial migration into the developing cerebral wall was identical in control and Arl13b-deficient neurons, which did possess cilia (Figures 1E, 1F, and 1K). At postnatal day 0 (P0), when projection neurons are terminating migration, we saw no changes in neuronal placement or layer specification using layer-specific markers Cux1 (layers 2–4), Ctip2 (layer 5), Brn1 (layers 2–4) and Tbr1 (layer 6) between control and Arl13b-deficient cortex (Figures 1A–1D). Once neurons reach their laminar destination, they establish functional connections by extending axons and dendrites and forming synapses. We observed generally normal patterns of extension and fasciculation of major subcerebral and cortico-cortical axonal pathways in Arl13bLox/Lox, Nex-Cre; mGFP mice at E16, P0, and P7 (Figures 1G–1R). However, we did notice a consistent decrease in the number of axonal bundles in the internal capsule at P0 and P7 (Figures 1M–1N and 1Q–1R), as well as disruptions in callosal projections (data not shown). Together, these findings suggest that loss of Arl13b function does not affect the migration of cortical projection neurons, but results in subtle disruptions in postmigratory axonal outgrowth and connectivity of selective subsets of projection neurons.

Conditional Deletion of Arl13b Disrupts Interneuronal Placement, but Not Postmigratory Differentiation in the Developing Cerebral Cortex

To determine the role of Arl13b in interneuronal migration, we examined the cortex from Arl13bLox/Lox, Dix5/6-Cre-IRESCGFP mice at postnatal day 10, when most cortical interneurons have terminated their migration and radial distribution within the cortical layers (Miyoshi and Fishell, 2011). By communolabeling Cre-expressing GFP+ interneurons with the interneuronal marker gamma-aminobutyric acid (GABA) in the somatosensory cortex, we saw a change in the positioning GABA/GFP double-positive cells, with a 39% decrease in the distribution of double-positive cells in layer VI with respect to the other layers in Arl13bLox/Lox; Dix5/6-CIE mice (Figures 2A, 2B, and 2G). Further, colabeling with interneuron subtype markers calretinin and somatostatin indicated a 2.5-fold increase in the distribution of calretinin/GFP double-positive cells in layers II and III and a 4.2-fold increase in somatostatin/GFP double-positive cells in layer V of Arl13bLox/Lox; Dix5/6-CIE somatosensory cortex (Figures 2C–2F, 2H, and 2I).

We also deleted Arl13b in the majority of postmigratory cortical interneurons (parvalbumin+) starting around P8, using Parvalbumin-Cre line (Ascoli et al., 2008; Hippenmeyer et al., 2005) but observed no changes in either the numbers or distribution of interneuron subtypes or in the dendritic number, length, or morphology of Arl13b-deficient interneurons compared to controls (Figure S2). Together, these observations indicate that normal Arl13b activity is specifically required for appropriate interneuronal placement within the cortex, but not for early postmigratory interneuronal differentiation.

Conditional Deletion of Arl13b Disrupts Interneuronal Migration in the Developing Cerebral Cortex

We next examined Arl13bLox/Lox; Dix5/6-CIE mice at times when interneurons are actively migrating into the cortex. At E14.5, mutant interneurons clustered at the pallial-subpallial boundary instead of exiting the ganglionic eminence (GE) and migrating into the dorsal cortex like control cells (Figures 3A, 3B, and 3D). The number of GFP+ interneurons entering the dorsal cortex from the GE was significantly reduced in the Arl13bLox/Lox; Dix5/6-CIE mutants (87 ± 3 cells/100 mm3 for Arl13bLox/Lox versus 185 ± 12 cells/100 mm3 for Arl13bLox/Lox). Normally, as interneurons migrate from the GE into the dorsal cortex, streams of migrating neurons can be seen in the marginal zone (MZ), intermediate zone (IZ), and subventricular zone (SVZ). We found fewer mutant interneurons in these migratory streams in mutants (compare Figure 3C to Figure 3D; Movies S1 and S2). We noticed a selective loss of interneurons migrating through the MZ region with a corresponding increase in the percentage of interneurons traversing the IZ (Figures 3E–3G). The extent of interneuronal migration from the pallial-subpallial boundary into the dorsal cerebral wall was significantly reduced in the mutants (Figures 3E–3F; 929 ± 58 μm for Arl13bLox/Lox versus 1,223 ± 47 μm for Arl13bLox/Lox). Moreover, mutant interneurons showed more processes extending from the cell soma as well as increased branching (Figures 3H–3L), but their overall branch length was decreased (Figure 3K). The same patterns of interneuron
migration defects persist at E16 (Figures S3A–S3D). The altered path, decreased distance traveled, and aberrant branching of mutant interneurons indicate that their migration is disrupted in the absence of Arl13b.

We confirmed this phenotype using a different Cre line, Nkx2.1-Cre, which is expressed beginning at E10.5 in both interneuronal progenitors and postmitotic interneurons of medial ganglionic eminence (MGE) and preoptic area (Xu et al., 2008). Similar to Arl13bLox/Lox;Dlx5/6-CIE brains, we observed defects in migration of both calbindin+ and Dlx2+ interneurons into the cortical wall at E14.5 and in the placement of GABA+ interneurons P10 cortex in the Arl13bLox/Lox; Nkx 2.1-Cre mutants (Figures S3E–S3K). We saw no significant changes in cell proliferation or cortical cell death using antibodies against the mitotic marker PH3 or apoptotic marker cleaved caspase3+ (data not shown) indicating the reduction in cortical interneurons migrating in the cortex in either Arl13bLox/Lox;Dlx5/6-CIE mutants or Arl13bLox/Lox; Nkx 2.1-Cre was not a result of changes in either interneuronal generation or survival.

**Figure 2. Interneuronal Placement Is Disrupted after Arl13b Deletion in Interneurons**

(A–F) P10 control Arl13bLox/Lox;Dlx5/6-CIE (A, C, and E) and Arl13bLox/Lox;Dlx5/6-CIE (B, D, and F) cortices were colabeled with GFP (Cre+) and GABA (A and B) calretinin (C and D) and somatostatin (E and F). Insets show higher magnification images of double-positive cells (arrowheads). Scale bar, 100 μm.

(G–I) Quantification of changes in the distribution of double-positive cells across cortical layers. Sections were from somatosensory cortex and counterstained with DRAQ5 (blue). Data shown are mean ± SEM; * indicates significant when compared with controls at p < 0.05 (Student’s t test).

See also Figure S2 and Movies S1, S2, and S3.

To gain insight into the dynamics of the migration defect in Arl13bLox/Lox;Dlx5/6-CIE interneurons in real time, we performed time-lapse imaging of interneuronal migration in E14.5 mutant and control cortices. The streaming of interneurons from the GE into the cortex was altered in Arl13bLox/Lox;Dlx5/6-CIE brains (Movie S1). In medial regions of the cortex that contain the leading front of migrating cohorts of interneurons, we were able to clearly track the migratory behavior of individual GFP+ interneurons. In control slices, we observed robust migration in the MZ and IZ streams. Interneurons frequently exited the IZ stream, migrated radially and entered the MZ stream, where they resumed tangential migration. In the mutant cortices, however, the characteristic interneuron migratory streams were greatly reduced, with a less well-defined stream in the IZ and fewer motile neurons in the MZ (Movie S2).

Our analysis of individual neurons revealed that mutant interneurons underwent the typical mechanics of tangential migration: extension of the branched leading process, movement of the cytoplasmic swelling into the process, followed by somal translocation. However, mutant neurons underwent 49% ± 0.5% fewer translocation events per hour than control cells. The periods of pausing were often characterized by the neurons sequentially extending leading processes in multiple directions without any accompanying somal translocation events. Therefore, we measured the rate of migration and found that Arl13bLox/Lox;Dlx5/6-CIE interneurons...
When interneurons arrive near their final areal position in the dorsal cortex, they turn and radially migrate to their final position within the cortical plate (Nadarajah et al., 2003; Yokota et al., 2007). Live imaging at E16.5 showed clear cortical plate-directed movement of interneurons in the IZ of control slices, but interneurons in mutant slices moved more randomly (Movie S3). We measured the orientation of leading processes in control and mutant interneurons and found that 62% ± 0.6% of Arl13bLox/+;Dlx5/6-CIE interneurons in the IZ displayed a leading process that was oriented toward the cortical plate. In Figure 3. Arl13 Deletion Leads to Interneuronal Migration and Branching Defects

(A and B) GFP-labeled coronal hemisections show interneuron migration defects in Arl13bLox/Lox;Dlx5/6-CIE mutants, with clusters of cells stuck at the pallial-subpallial boundary (SB, arrow).

(C and D) Loss of characteristic interneuronal migratory streams in Arl13bLox/Lox;Dlx5/6-CIE cortex. Arrowheads in C indicate streams of migrating interneurons in the MZ, IZ, and SVZ of control cortex. Arrows in (D) indicate disruptions in these streams in mutant cortex.

(E and F) Higher magnification images of the control (E) and mutant (F) dorsal cortex illustrating disrupted extent and patterns of migration in mutants. Left asterisks mark pallial-subpallial boundary; right asterisks mark the migration front.

(G) Disrupted migration results in altered distribution of INs across the cortical wall in Arl13bLox/Lox;Dlx5/6-CIE cortex, with a larger percentage of cells moving through the IZ and a corresponding loss of cells in other locations. The overall number of interneurons was reduced in mutants, but mutants show a higher percentage of interneurons in the IZ than controls.

(H and I) Camera lucida drawings of sample control (H) and mutant (I) interneurons in the dorsal cortex.

(J–L) Quantification of branching defects in Arl13b-deficient interneurons. (J) Number of processes extending from cell soma. (K) Total process length (n = 506 cells each from control and mutants). (L) Branching index (average number of branching events in a migrating interneuron per hour). Data shown are mean ± SEM; * indicates significant when compared with controls at p < 0.05; ** indicates significant when compared with controls at p < 0.01 (Student’s t test).

Scale bars, 675 μm (A–D); 225 μm (E and F); 100 μm (G and H); 25 μm (J and K). SVZ, subventricular zone; GE, ganglionic eminence; D.CX, dorsal cortex. See also Figure S3 and Movies S1, S2, S3, S4, and S5.

(24.6 ± 1.2 μm/h) migrated 29% slower than control cells (34.2 ± 0.6 μm/h).
Arl13bLox/Lox;Dlx5/6-CIE interneurons, however, only 36% ± 2.7% were oriented toward the cortical plate. Together, these real time observations suggest that Arl13b is critical for the oriented navigation and migration of interneurons in the developing cerebral cortex.

Primary Cilia Function Is Required for Interneuronal Navigation through Gradients of Guidance Cues
To determine whether the defective migration of Arl13b mutant interneurons is partly due to their inability to sense and respond to extrinsic migration guidance cues released by dorsal cortical cells (Valiente and Marín, 2010), we examined the ability of control and mutant interneurons to respond to gradients of such cues using a microfluidics chamber. This microscale device allowed us to present dorsal cortical cell-derived gradients of migration-regulating cues to isolated interneurons and to assay their response to these cues in a defined microenvironment (Taylor and Jeon, 2010). Here, interneurons and dorsal cortical cells are physically separated in two separate channels and interneurons are exposed to a microfluidic gradient of signals secreted by dorsal cortical cells in the opposing channel (Figure 4A). Under control conditions, interneurons migrate toward the dorsal cortical cells (Figure 4B). In contrast, the extent of migration of Arl13bLox/Lox; Dlx5/6-CIE interneurons toward dorsal cortical cells was significantly reduced (Figures 4C and 4D). Further, compared to control interneurons, migration of Arl13b null interneurons was also perturbed (Figures 4E–4G). Together, these results suggest that the disruption of Arl13b in interneurons perturbs their ability to sense and respond to extrinsic guidance cues present in dorsal cortex that are necessary for the appropriate navigation into the cerebral wall.

Disruption of Primary Cilium Localization of Arl13b Disrupts Interneuronal Migration
To investigate whether the requirement for Arl13b in interneuron migration is specifically due to its ciliary role, we generated a nonciliary form of Arl13b, Arl13bV358A (Figure 5A). The Arl13bV358A variant retains its GTPase activity but does not localize to cilia (Figure 5B), due to disruption of its VxPx motif (Deretic et al., 1998). Arl13bV358A could not rescue the migratory defects in Arl13b deficient interneurons (Figure 5C). Likewise, electroporating the Arl13b N-terminal domain (1-190aa; Arl13b-ND), which prevents Arl13b localization to cilia and maintenance of the primary cilium (Hori et al., 2008), impaired the migration of the electroporated interneurons into the dorsal cortex (Figures 5D–5F, 5H, and 5I). Compared to the enhanced green fluorescent protein (EGFP)-electroporated control, 66% fewer Arl13b-ND expressing interneurons entered the dorsal cortex (15 ± 2.46 interneurons entered the dorsal cortex per slice electroporated with EGFP versus 5 ± 0.80 electroporated with Arl13b-ND; n = 35 slices). The migration index,
calculated as the percentage change in interneurons that migrated greater than 350 μm past the GE-dorsal cortex boundary, was reduced by 72% in Arl13b-ND-GFP+ interneurons (Figure S5). These results suggest that primary ciliary localization of Arl13b is required for efficient interneuronal migration into the dorsal cortex.

To confirm that the Arl13b effects on interneurons depend on primary cilia, we analyzed the embryonic cortex of Ift88 null mutants, which lack cilia and die by E11.5 (Pazour et al., 2000). Despite the likelihood of interneuron generation itself being affected in Ift88 mutants, we found the migration of Ift88<sup>−/−</sup> interneurons into the cerebral wall was severely disrupted (Figures 5J and 5K). As conditional deletion of Ift88 in postmitotic interneurons failed to ablate cilia (data not shown), we corroborated this using a small molecule inhibitor (HPI-4) that prevents ciliogenesis and truncates existing cilia (Hyman et al., 2009) (Figure S4). Together, these studies demonstrate that primary cilia activities involving Arl13b are critical for interneuron migration.

**Interneuronal Primary Cilia Show Dynamic Changes in Length and Orientation during Migration**

Primary cilia morphology, length and orientation, are linked to cilia-mediated signaling (Besschetnova et al., 2010; Ishikawa and Marshall, 2011) leading us to investigate the morphological dynamics of primary cilia in migrating interneurons in real time. We labeled interneuronal primary cilia with a Smoothened-tdTomato (Smo-Tom) fusion construct in the MGE of Arl13b<sup>Lox/lox</sup>; Dlx5/6-CIE and Arl13b<sup>Lox/Lox</sup>; Dlx5/6-CIE cortices and used time-lapse imaging on a confocal microscope (Kim et al., 2010). Primary cilia in migrating interneurons are dynamic; they undergo repeated changes in length and position in stationary and moving neurons (Movie S4). We noticed that the cilia in control Arl13b<sup>Lox/lox</sup>; Dlx5/6-CIE interneurons were, on average, 35% longer than in mutant Arl13b<sup>Lox/lox</sup>; Dlx5/6-CIE interneurons (4.78 ± 0.75 μm, Arl13b<sup>Lox/lox</sup>; 3.08 ± 0.70 μm, Arl13b<sup>Lox/Lox</sup>; n = 35 control, 43 mutant cells). In addition, we found that dynamic changes in cilia morphology correlated with the migratory behavior of interneurons. For example, in both Arl13b<sup>Lox/lox</sup>; Dlx5/6-CIE and Arl13b<sup>Lox/Lox</sup>; Dlx5/6-CIE interneurons, cilia predominantly localized to the soma of stationary cells and to the leading process of moving cells (during 66% ± 6.9% of the time spent moving, the primary cilia were found in or near the base of the leading process; during periods of pausing, the cilia were located in the cell soma 62% ± 6.6% of the time). Pausing interneurons displayed dynamic cilia that elongated, shortened, and rotated more than in moving interneurons (Movie S4).

In pausing control interneurons, cilia were, on average, 23% longer than in moving cells (5.07 ± 0.87 μm in pausing neurons; 3.92 ± 1.51 μm in moving neurons, n = 35 cells/group), and the dynamic range of cilia length was 59% greater (4.44 ± 0.73 μm) than in moving neurons (1.83 ± 0.58 μm). Consistent with these live imaging observations, we noticed primary cilia of differing lengths in fixed embryonic interneurons (Figure S5).

In contrast, primary cilia dynamics in Arl13b mutant interneurons were significantly altered (Movie S5). Cilia in pausing cells were only marginally (3.8%) longer than in moving cells (3.13 ± 0.76 μm, stationary; 3.01 ± 1.40 μm, moving; n = 43 cells/group). In addition, the dynamic range of cilia length in pausing cells (2.02 ± 0.41 μm; n = 16) was not significantly different from that of moving cells (2.46 ± 1.39 μm; n = 27), but was significantly different from cilia in control neurons (4.44 ± 0.73 μm). These observations demonstrate that the primary cilia in migrating interneurons are morphologically dynamic, and that the dynamics might correlate with the interneuron’s phase of migration. Importantly, Arl13b mutant interneurons with defective migration display aberrant primary cilia dynamics during migration.

**Localization of Guidance Cue Receptors in the Primary Cilia of Interneurons**

We examined whether the receptors for guidance cues that are known to regulate interneuronal migration localize to the primary cilia of interneurons during migration. We labeled the primary cilia of Arl13b<sup>Lox/lox</sup>; Dlx5/6-CIE and Arl13b<sup>Lox/Lox</sup>; Dlx5/6-CIE interneurons via electroporation with Smo-Tom. We then mapped the interneuronal cilia localization of the following receptors known to be critical for interneuron migration: BDNF receptor TrkB, GDNF receptor GFRα2-1, SDF-1 receptors CXCR4 and CXCR7, NRG-1 receptor ErbB4, serotonin receptor 6-5Htr6, Slit receptors Robo1 and 2, and HGF/SF receptor MET (reviewed in Valiente and Marin, 2010; Sánchez-Alcañiz et al., 2011; Wang et al., 2011, Riccio et al., 2009) (Figures 6A and 6B). All of the guidance cue receptors that we tested localized to primary cilia in both mutant and control interneurons, but with different frequencies (Figure 6C). For each receptor, we observed differences in the percentage of primary cilia showing guidance cue receptor localization in mutant interneurons compared to control interneurons. We also examined the localization of the signaling receptors in ciliated mouse embryonic fibroblasts (MEFs) in which we could induce Arl13b deletion and found similar changes in the frequency of signaling receptor cilia localization. These changes in the distribution of signaling receptors upon Arl13b deletion support the hypothesis that the concentration of critical signaling receptors in primary cilia may promote their ability to function as sensors and conveyors of migration-regulating signals to interneurons during corticogenesis. Consistent with this hypothesis, we detected an increase in cyclic AMP (cAMP) levels and decreased Erk phosphorylation in Arl13b mutant interneurons. These two second messenger pathways are critical for interneuron migration and are downstream of many of the interneurons (IN) migration regulating receptors described above (Garzotto et al., 2008; Grabert and Wahle, 2008; Imamura et al., 2010; Lysoke et al., 2011; Samuels et al., 2008; Sánchez-Alcañiz et al., 2011; Segarra et al., 2006; Wang et al., 2011).

**Analysis of Guidance Cue Receptor Dynamics in Arl13b Mutant Cilia**

In *C. elegans*, the ortholog of Arl13b, ARL-13, regulates intraflagellar transport (IFT) (Li et al., 2010) and is required for the stable localization of ciliary transmembrane proteins (Cevik et al., 2010). To investigate whether Arl13b modulates the transport and localization of transmembrane interneuronal guidance cue receptors into and within the cilium, we used fluorescence recovery after photobleaching (FRAP) to measure the movement of GFP-tagged serotonin receptor 6-5Htr6, which regulates interneuron migration (Riccio et al., 2009), within the primary cilia of control and Arl13b protein null (Arl13b<sup>−/−</sup>) MEFs (Casparry et al., 2007). 5-Htr6-GFP localizes to the primary cilia in both control and Arl13b−/− MEFs (Figure 6C). The primary cilia of Arl13b−/− MEFs displayed aberrant ciliary dynamics, leading to an increase in the percentage of primary cilia showing guidance cue receptor localization in mutant interneurons compared to control interneurons. These changes in the distribution of signaling receptors upon Arl13b deletion support the hypothesis that the concentration of critical signaling receptors in primary cilia may promote their ability to function as sensors and conveyors of migration-regulating signals to interneurons during corticogenesis. Consistent with this hypothesis, we detected an increase in cyclic AMP (cAMP) levels and decreased Erk phosphorylation in Arl13b mutant interneurons. These two second messenger pathways are critical for interneuron migration and are downstream of many of the interneurons (IN) migration regulating receptors described above (Garzotto et al., 2008; Grabert and Wahle, 2008; Imamura et al., 2010; Lysoke et al., 2011; Samuels et al., 2008; Sánchez-Alcañiz et al., 2011; Segarra et al., 2006; Wang et al., 2011).
Figure 5. Ciliary Function of Arl13b Is Critical for the Regulation of Interneuron Migration

(A) Schematic of a nonciliary form Arl13b (Arl13bV358A).

(B) In ciliated IMCD3 cells transfected with control Arl13b-GFP and Arl13bV358A-GFP constructs, Arl13b-GFP localizes to cilia (arrow), but Arl13bV358A-GFP does not. Nuclei were counterstained with DAPI.

(C) Expression of wild-type Arl13b rescued the migratory defect in Arl13b deficient (Arl13bLox/Lox; Dlx5/6-CIE) interneurons. In contrast, expression of Arl13bV358A did not rescue the defect. Data shown are mean ± SEM (n = 42 [Control: Arl13bLox/+; Dlx5/6-CIE], 79 [Arl13b deficient: Arl13bLox/Lox; Dlx5/6-CIE], 104 [Arl13bLox/Lox; Dlx5/6-CIE + Arl13b], 49 [Arl13bLox/Lox; Dlx5/6-CIE + Arl13bV358A]); * indicates significant when compared with controls at p < 0.001; ** indicates significant when compared with Arl13b mutant at p < 0.001 (Student’s t test).

(D–I) Dominant negative N-terminal domain (Arl13b-ND) of Arl13b disrupts selective targeting of Arl13b to primary cilia and inhibits interneuronal migration. (D) Drawing of experimental protocol. Arl13b-ND/GFP or GFP DNA was focally electroporated into the MGE of E14.5 coronal slices. Images of GFP+ interneuronal migration into dorsal cortex were acquired at 24 and 48 hr. (E–H) Interneurons expressing GFP alone leave the MGE at 24 hr (E, left panel) and migrate into the dorsal cortex by 48 hr (E, right panel, and higher magnification image of dorsal cortex in H). Interneurons coexpressing Arl13b-ND mostly remain in the GE at 24 hr (F, left panel) and show less migration into dorsal cortex at 48 hr (F, right panel, and higher magnification image in I). Compared to control (H, arrows), fewer GFP+ interneurons are seen in the dorsal cortex in Arl13b-ND slices (I). (G) Quantification of changes in the extent of interneuronal migration. Migration index indicates...
wild-type and Arl13b<sup>hn</sup> cells (Figures 7A and 7B) and serves as a useful molecular tool to explore the receptor dynamics within primary cilia. Compared to wild-type cells, 27% fewer Arl13b<sup>hn</sup> MEFs localized 5-Htr6-GFP to the cilium. Photobleaching of a subregion of 5-Htr6-GFP-containing cilia revealed that cells lacking Arl13b show a 2-fold increase in recovery half-time (1.76 ± 0.30 s in Arl13b WT versus 3.63 ± 0.44 s in Arl13b<sup>hn</sup>; n = 14 [WT]; Figures 7C and 7D; Movie S6). The increase in recovery time was specific to receptor movement within mutant primary cilia; FRAP rates of 5-Htr6-GFP localized to nonciliary membranes of Arl13b<sup>-/-</sup> cells were similar to wild-type rates (WT, 1.04 ± 0.15 s; Arl13b<sup>-/-</sup>, 1.19 ± 0.09 s). These data suggest that Arl13b is required for the efficient ciliary movement and localization of interneuronal migration guidance cue receptors such as 5-Htr6.

Expression of Mutated Human Arl13b Disrupts Interneuron Migration

Three distinct mutations in Arl13b have been identified in Joubert syndrome patients (Cantagrel et al., 2008). The R79Q substitution interferes with guanosine triphosphate binding, W82X introduces a stop codon in exon 3 and R200C generates a missense (J and K) Disrupted interneuron migration in Ift88<sup>-/-</sup> brains. (I) Calbindin+ interneurons (red, arrow) migrate from the ganglionic eminence into the dorsal cortex. This migration is severely disrupted in Ift88<sup>-/-</sup> brain (K). Insets (J and K) show the presence and absence of Arl13b<sup>+</sup> cilia in wild-type and mutant brains, respectively. Sections were counterstained with DAPI or DRAQ5 (blue).

Scale bars, 34 μm (B and C); 400 μm (E and F); 175 μm (H and I); 180 μm (J and K). See also Figure S4 and Movies S4 and S5.
mutation within the C-terminal coiled-coiled domain of Arl13b (Figure 8A). To test whether expression of the mutated forms of human Arl13b could affect interneuron migration, we focally electroporated wild-type human or mutant human ARL13B constructs tagged with tdTomato into the GE of Dlx-Cre, Arl13bLox/Lox E14.5 cortex. While wild-type human ARL13B rescued the migratory defects in the mutant cells, none of the ARL13B patient variants did (Figure 8), suggesting that the human mutations in Arl13b may affect IN migration in Joubert syndrome patients. Anticaspase labeling indicates that cells expressing human mutant Arl13b are not apoptotic (data not shown). Together, these studies demonstrate that expression of disease alleles of human Arl13b in interneurons can disrupt their migration.

DISCUSSION

Disrupted primary cilia function in humans results in profound cortical abnormalities and cognitive deficits (Hidebrandt et al., 2011; Louvi and Grove, 2011). Nevertheless, the role of cilia function in cortical neuronal organization remains unknown. Here we show that primary cilia play a guiding role in the development and placement of interneurons during the construction of cerebral cortex.

By using Arl13b, we gained an entry point for understanding the role of cilia in cortical development since Arl13b mutants impair but do not destroy the cilia and downstream pathways. Our results provide insights into the etiology of the neurodevelopmental defects found in ciliopathies. It is well established that altered integration of interneurons into the cortical network and the resultant changes in brain’s excitatory/inhibitory balance can lead to developmental neuropsychiatric disorders like autism spectrum disorders, epilepsy and schizophrenia (Batista-Brito and Fishell, 2009; Manzini and Walsh, 2011; Mélin et al., 2008; Valiente and Marín, 2010; Wynshaw-Boris et al., 2010; Yizhar et al., 2011). Importantly, these disorders have been associated with Joubert syndrome (Lee and Gleeson, 2011; Sattar and Gleeson, 2011). Our data suggest that defective interneuron migration and placement following Arl13b deletion may disrupt the appropriate elaboration of cortical inhibitory network, thus contributing to altered cortical inhibition and neurobehavioral deficits in Arl13b-related ciliopathies. Furthermore, appropriate radial distribution and integration of diverse subtypes of GABAergic interneurons into distinct laminar locations occurs during early postnatal development (De Marco García et al., 2011; Lodato et al., 2011; Miyoshi and Fishell, 2011; Rudy et al., 2011). The altered distribution of GABA, calretinin, and somatostatin expressing interneurons in Arl13b mutants is suggestive of a potential role for Arl13b in interneuron subtype migration or the emergence of migration-related interneuron subtype identity.

The distinct roles of Arl13b activity for interneuron and projection neuron migration may reflect the fundamental differences in the way these two major classes of cortical neurons migrate. Projection neurons migrate primarily using radial glial guides as substrates, whereas interneurons rely extensively on extracellular, nonsubstrate-based guidance cues for their migration into the cerebral wall (Valiente and Marín, 2010). Primary cilia activity might be more critical for cell migration that depends on sensing extracellular guidance cues and less so for substrate contact-guided migration.

Our results emphasize a role for cilia in early cortical development. However, the disruption in the major subcerebral axonal projections (i.e., internal capsule) of pyramidal neurons in Arl13b mutants may be related to a role for Arl13b+ cilia in the postmitotically identity and differentiation of subsets of projection neurons. Alternatively, it may also be the result of a yet to be uncovered, noncilia role for Arl13b in axon growth in a subset of projection neurons. Disrupted cortico-spinal axonal connectivity is also seen in Joubert syndrome patients (Engle, 2010; Poretti et al., 2007) and primary cilia-specific activities of Arl13b in pyramidal neurons may underlie some aspects of this defect.

One surprising finding is how dynamic the primary cilia are; not only do they change in length, they also rotate, branch, and appear to probe their surroundings. This is quite distinct from true motile cilia; ependymal cilium beat rhythmically to form a guidance molecule concentration gradient in the CSF that directs the migration of adult neuroblasts (Sawamoto et al., 2006). Intereuronal primary cilia likely bind and transduce signals from gradients of guidance molecules in a cell autonomous fashion that leads to changes in interneuronal morphology and movement. In fact, we found that cilia in wild-type interneurons localized to the leading process during migration, but moved closer to the cell body during periods of pausing when they became much more dynamic in their movement and length. In contrast, primary cilia in Arl13b-deficient interneurons did not display such dynamism. Considering that mutant interneurons paused longer during migration, extended an increased number
of processes in multiple directions, and were unable to respond to microgradients of guidance cues, the reduction in cilia dynamics may reflect an inability to efficiently sense guidance cues in the migratory environment. This is further supported by the disrupted localization of multiple signaling receptors in Arl13b-deficient interneuronal cilia.

The mechanism(s) underlying the misdistribution of guidance cue receptors in Arl13b-deficient cilia remains unclear. Another protein implicated in Joubert syndrome, the transition zone protein, Tctn1, regulates Arl13b localization to the cilium in neurons (Garcia-Gonzalo et al., 2011). Moreover, C. elegans ARL-13 regulates IFT by maintaining the association between IFT sub-complexes A and B and associates with the ciliary membrane via its palmitoylation motifs (Li et al., 2010; Cevik et al., 2010). Together, these data suggest that Arl13b’s links to the IFT complexes, ciliary transmembrane receptor localization, and the ciliary transition zone may help promote the retention and distribution of crucial receptor molecules within the primary cilium of interneurons. Detailed understanding of such a model and its relevance for the regulation of diverse patterns of interneuron subtype migration will await more refined genetic and molecular tools that would allow cilia specific signaling manipulation in specific subsets of interneurons in the early embryonic brain. But consistent with this idea, we detected an increase in

Figure 8. Expression of Mutated Human Arl13b Allele Disrupts Interneuron Migration
(A) Pathogenic ARL13B mutations R79Q, W82X, and R200C cause Joubert syndrome.
(B–E) Panels from time-lapse imaging of control (Arl13b<sup>Lox<sub>129</sub>/Cre; Dlx5/6-CIE) (B), Arl13b deficient interneurons (Arl13b<sup>Lox<sub>129</sub>/Lox<sub>129</sub>; Dlx5/6-CIE) (C), and Arl13b deficient interneurons expressing either WT (D) or W82X (E) human Arl13b (red). Asterisks in (B–E) indicate cell soma of migrating interneurons. Time elapsed between panels is in minutes. Scale bar, 75 µm.
(F) Reduction in the rate of migration in Arl13b deficient interneurons is rescued by expression of human Arl13b. R79Q, W82X, and R200C mutant Arl13b constructs fail to rescue, suggesting these mutations impair interneuron migration. Data shown are mean ± SEM (n = 40 cells for each condition). * Indicates p < 0.001 compared with control; ** indicates p < 0.001 when compared with Arl13b deficient cells (Student’s t test).
cAMP levels along with a decrease in Erk phosphorylation when we examined the total population of Arl13b mutant interneurons. Both of these pathways are downstream of several interneuron migration regulating receptors (Garzotto et al., 2008; Grabert and Wahle, 2008; Imamura et al., 2010; Lysko et al., 2011; Samuels et al., 2008; Sánchez-Alcañiz et al., 2011; Segarra et al., 2006; Stanco et al., 2009; Wang et al., 2011). Increased cAMP levels can disrupt the branching of migrating interneurons, and decreased ERK phosphorylation is associated with retarded interneuron migration (Lysko et al., 2011; Imamura et al., 2010). Although the current evidence on Arl13b suggests that ciliary localization of multiple signaling receptors are perturbed in Arl13 mutants, future identification of IN migration related ligand-receptor combination or receptor subtypes (e.g., GPCRs) that are preferentially affected in Arl13b mutants will help further define the specific signaling pathways that may be defective in Arl13b deficient primary cilia.

This hypothesis raises important questions about how directional information from guidance cues is conveyed through the cilium to lead to changes in interneuron migration. Also, what advantages does the cell gain by relying on such a small organelle to transduce extracellular signals? The primary cilium is structured and positioned to be able to generate a fast and robust response to extracellular signals. The diffusion barrier at the base of the cilium allows a cell to maximize the local concentration of guidance cue receptors and their downstream effectors within cilia. Thus, high levels of second messengers could be sustained more easily within the narrow geometry of the cilium than in other regions of the cell where diffusion through a larger space could lead to a rapid diminishment of the signal. In addition, the cilium’s proximity to the nucleus and its link to the centrosome-associated microtubule network could promote rapid and efficient signal transduction during nucleokinesis associated with interneuron migration. Consistently, pericentrin, a member of the centrosomal complex necessary for neuronal cilia formation (Miyoshi et al., 2009), is crucial for tangential neuronal migration (Endoh-Yamagami et al., 2010). However, it remains untested if primary cilia and the growth cones in the leading processes of migrating interneurons transduce different signals related to directed cell movement. They may differentially affect the chemotactic and haptotactic components of interneuron migration. Understanding the crosstalk that occurs between intracellular signaling cascades that originate in each of these cellular compartments during interneuron movement will help delineate the primary cilia-specific pathways mediating interneuron development. Furthermore, it will be crucial to examine the differences in the activities of receptors that are localized exclusively in one or in both primary cilia and growth cones during interneuron migration. Nevertheless, the evidence we present in this study demonstrates a role for primary cilia in neuronal development and cerebral cortical organization. Moreover, it provides a neurodevelopmental basis for the functional deficits observed in Joubert syndrome patients.

**EXPERIMENTAL PROCEDURES**

**Mice**

Mice were cared for under protocols approved by the University of North Carolina and Emory University. Lines used were: Arl13bhnn/hnn, MGI:3578151 (Caspary et al., 2007), Nex-Cre (Goebel et al., 2006), Dlx5/6-CIE (Stenman et al., 2003), Parv-Cre (Hippemeyer et al., 2005), Tau-Lox-STOP-Lox-GFP (Hippemeyer et al., 2005), Nkx2.1-Cre (Xu et al., 2008), Ift88lox/lox (Haycraft et al., 2007), and CAGG-CreERT2 (JAX: 004882; Hayashi and McMahon, 2002).

**Immunohistochemistry**

Cerebral cortical sections and cortical cells were immunolabeled as previously described (Schmid et al., 2003; Yokota et al., 2007, 2009). List of primary antibodies used are provided in the supplemental information. Secondary antibodies were AlexaFluor 488 or Cy3-conjugated (Invitrogen and Jackson ImmunoResearch). Nuclei were counterstained with DRAQ5 (Biostatus Limited).

**Live Imaging of Interneuronal Migration and Primary Cilia Activity in Interneurons**

GFP” interneurons, Smo-tdTomato” interneuronal primary cilia, or interneurons expressing ARL13B patient variants in embryonic cortices were live imaged using a Zeiss LSM710 inverted confocal laser-scanning microscope or an Olympus Fluvo-View FV1000 inverted confocal laser-scanning system attached to a live-cell incubation chamber.

**Microfluidic Chamber Assay for Interneuronal Migration**

Interneurons seeded onto microfluidic chambers (Taylor and Jeon, 2010) were used probe the ability these neurons to respond to extrinsic migration guidance cue gradients released by dorsal cortical cells. Details of this microfluidics assay are provided in the supplemental information.

**Inducible Deletion of Arl13b in MEFs**

Mouse embryonic fibroblasts were isolated from E12.5 control Arl13bLox/Lox or Arl13bLox/lox, CAGG-CreERT2 embryos as previously described. Mouse embryonic fibroblasts were serum-starved for 24 hr and tamoxifen (2 μM, Sigma H7904) was added in serum-free media. Coverslips were collected 48 hr after tamoxifen treatment, and MEFs were fixed and processed for antibody staining.

**FRAP Analysis**

Wild-type and Arl13bhnn/hnn MEFs were grown in Dulbecco’s modified Eagle’s medium/10% fetal bovine serum and cotransfected with pcAGS-dsRed and pcAGS-HH6-GFP (a gift from Dr. Kirk Mykytyn, Ohio State University). Fluorescence recovery after photobleaching was carried out after 48 hr of serum starvation using the Olympus Fluoview 1000 live cell confocal system. After the acquisition of basal state images, the region of interest (ROI) was photo-bleached with a 800 ms pulse of 488 nm laser, followed by rapid, repetitive acquisition of the recovery of GFP fluorescence in the ROI. Data were analyzed using the Igor Pro 6.12A (Wavevmetrics, Inc.) software to determine the recovery half-time and evaluate the dynamics of receptor movement within primary cilium.

**Measurement of cAMP Levels**

Cyclic AMP levels in the ganglionic eminence extracts from Arl13blox/–;Dlx5/6-CIE or Arl13blox/lox;Dlx5/6-CIE brains (E16) were measured as described in the manufacturer’s instructions (R&D Systems). Extracts contained equal levels of GFP.

**SUPPLEMENTAL INFORMATION**

Supplemental Information includes five figures, six movies, and Supplemental Experimental Procedures and can be found with this article online at http://dx.doi.org/10.1016/j.devcel.2012.09.019.

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