

# Neuronal Migration in the Developing Brain

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## INTRODUCTION

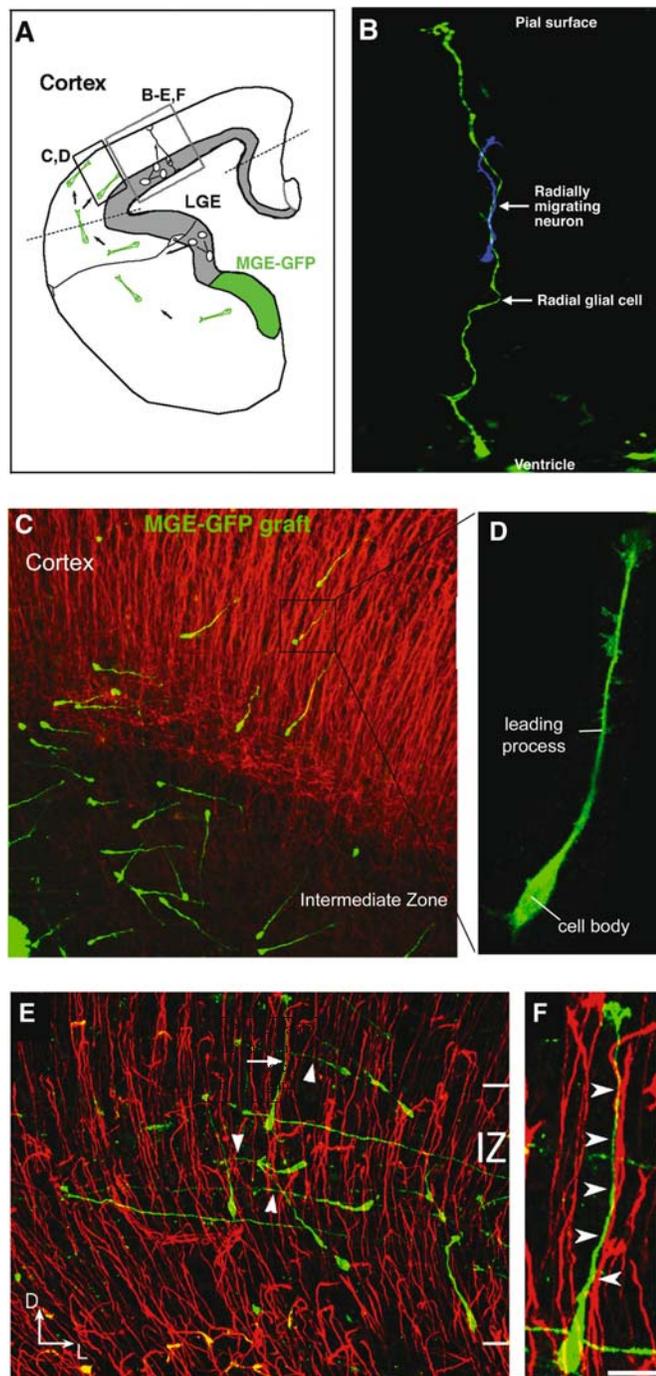
The way in which a nervous system is constructed predisposes and constrains its functions. Thus the study of neuronal cell migration, an elementary step in the histogenesis of any nervous system, is critical if we are to understand how the structure and function of a nervous system come about. Specific neuronal networks emerge as a result of appropriate migration and final placement of neurons during development. In the developing nervous system, most, if not all, neurons undergo their terminal division and terminal differentiation in distinct locations. Specific neuronal populations have to migrate in distinct pathways and patterns over extensive distances to reach their final position. Two main types of migration predominate during the development of the central nervous system: radial vs tangential. Radial migration is characterized by close interactions between migrating neurons and the processes of radial glial cells, which constitute a scaffold bridging the proliferating neuroepithelium and the differentiating zone. Postmitotic neurons migrate radially from the ventricular zone toward the pial surface past previously generated neuronal layers (Rakic, 1971b, 1972a) to reach the top of the cortical plate (CP), where they terminate their migration and assemble into layers with distinct patterns of connectivity. Radial migration of cortical neurons can occur in two distinct modes: locomotion or somal translocation (Nadarajah *et al.*, 2001; Nadarajah and Parnavelas, 2002; Nadarajah *et al.*, 2002). In contrast, tangential migration is referred to as a nonradial mode of neuronal translocation that does not require specific interaction with radial glial cell processes. Observations of tangential dispersion of precursors or postmitotic neurons in the developing cortex suggested the possibility of nonradial migration in the cortex (O'Rourke *et al.*, 1992, 1995, 1997; Walsh and Cepko, 1992; Fishell *et al.*, 1993; Tan and Breen, 1993; Tan *et al.*, 1995; de Carlos *et al.*, 1996). Analysis of *Dlx1/2* double knockout mice has demonstrated for the first time that subpopulations of GABAergic interneurons, originating from the ventral telencephalon (also called the ganglionic eminence [GE]), indeed migrate tangentially into the neocortex (Anderson *et al.*, 1997). Therefore, there is a tight correlation between neuronal subtype

identity (glutamaergic vs GABAergic) and the mode of migration (radial vs tangential) in the developing cortex of mammals (Parnavelas, 2000).

Specific cell–cell recognition and adhesive interactions between neurons, glia, and the surrounding extracellular matrix (ECM) appear to modulate distinct patterns of neuronal migration, placement, and eventual differentiation within cortex. A fundamental challenge in the study of cortical development is the elucidation of mechanisms that determine how neurons migrate and coalesce into distinct layers or nuclei in the developing cerebral cortex. In this regard, several related questions need specific attention: (1) What are the cell-intrinsic and extracellular cues that trigger the onset of neuronal migration following last mitotic cell division? (2) What is the molecular basis and role of glial-independent and glial-guided neuronal migration in cortical development? (3) How do migrating neurons know where to end? and (4) What are the stage-specific genes that determine distinct aspects of neuronal migration in developing mammalian brain? In combination, analysis of these questions may elucidate some of the fundamental rules guiding the development of cerebral cortex.

## PATTERNS OF NEURONAL MIGRATION

Extensive observations of neuronal migration in the past several decades in mammalian cerebral cortex and recent molecular characterization of migration deficits in mice and humans have raised the cerebral cortex as a prototype model for the analysis of migration in the developing mammalian central nervous system. Radial glial cells play a critical role in the construction of the mammalian brain by contributing to the formation of neurons and astrocytes and by providing a permissive and instructive scaffold for neuronal migration. The establishment of radial glial cells from an undifferentiated sheet of neuroepithelium precedes the generation and migration of neurons in the cerebral cortex. During early stages of corticogenesis, radial glial cells can give rise to neurons (reviewed in Fishell and Kriegstein, 2003; Rakic, 2003). Subsequent neuronal cell movement in the developing mammalian cerebral cortex occurs



**FIGURE 1.** Radial vs tangential patterns of neuronal migration. In the developing embryonic cortex (A), radially (B) and tangentially (C) migrating neurons display a unipolar morphology characterized by a prominent leading process (D). These neurons are in intimate contact with either radial glial cells (B, green) or with neurites (C, red) within the developing cortex. Tangentially migrating neurons (arrowhead, E) eventually turn radially (arrow, E) in the intermediate zone and associate with radial fibers (F, red) during final stages of translocation to the cortical plate. Cells shown in B were electroporated with GFP, whereas tangentially migrating neurons (C–F) were isolated from GFP-expressing MGE graft on a slice co-culture assay (Polleux *et al.*, 2002).

mainly along radial glial fibers, though nonpyramidal neurons initially migrate into the cortex in a radial glial-independent manner (Fig. 1). Neurons migrate from the ventricular zone toward the pial surface past previously generated neuronal layers (Rakic, 1971a, b; 1972a, b) to reach the top of the CP, where they terminate their migration and assemble into layers with distinct patterns of connectivity. Radial migration of cortical neurons can occur in two distinct modes: locomotion or somal translocation (Nadarajah *et al.*, 2001; Nadarajah and Parnavelas, 2002; Nadarajah *et al.*, 2002). Somally translocating neurons, prevalent during early phases of corticogenesis, appear to move toward the pial surface by maintaining pial attachment while losing their ventricular attachments. In contrast, radial glial cell fibers serve as the primary migratory guides for locomoting neurons (Rakic, 1971a, b, 1972a, b, 1990; Gray *et al.*, 1990; Hatten and Mason, 1990; Misson *et al.*, 1991). These neurons form specialized membrane contacts variably referred to as junctional domains, interstitial contacts, or punctae adherentia with the underlying radial glial cell substrate (Gregory *et al.*, 1988; Cameron and Rakic, 1994; Anton *et al.*, 1996). Such specialized membrane contacts are hypothesized to be critical elements in the maintenance of directed neuronal cell migration along radial glial cell fibers (Rakic *et al.*, 1994). The radial movement of neurons stops abruptly at the interface between CP and cell sparse marginal zone. The signal to end neuronal cell migration is thought to be provided either by the afferent fibers that migrating neurons encounter near their target location or by the ambient neuronal cell population that had already reached its final position (Sidman and Rakic, 1973; Hatten, 1990, 1993; Hatten and Mason, 1990; D'Arcangelo *et al.*, 1995; Ogawa *et al.*, 1995). Alternately, a change in the cell surface properties of the radial glial substrate may signal a neuron migrating on it to stop, detach, and differentiate.

In contrast to radially migrating neurons, populations of GABAergic interneurons, originating from the GE, migrate tangentially into the neocortex (Anderson *et al.*, 1997; Letinic and Rakic, 2001; Maricich *et al.*, 2001; Tamamaki *et al.*, 1997; Wichterle *et al.*, 2001; see Fig. 1). Some of these neurons migrate ventrally toward the cortical ventricular zone prior to radial migration toward the pial surface (Nadarajah *et al.*, 2002). These distinct patterns of neuronal migration enables several generations of neurons to reach their appropriate areal and laminar positions in the developing CP. Analysis of mutations in mice and humans have revealed several molecular cues controlling different aspects of neuronal migration. Evidently, a dynamic regulation of multiple cellular events such as cell–cell recognition, adhesion, transmembrane signaling, and cell motility events underlies the process of neuronal migration.

## MECHANISMS UNDERLYING RADIAL MIGRATION

### Initiation of Migration

Movement of neuronal cells from their site of birth in the ventricular areas to the specific laminar location involves

a progressive unraveling of three interrelated cellular events: initiation of migration along appropriate pathways or substrates, maintenance of migration through a complex cellular milieu, and termination of migration in the CP at the appropriate laminar location.

In humans with periventricular heterotopia, mutations in actin-binding protein filamin 1 (or filamin- $\alpha$  [FLNA]) results in failure of neuronal migration and accumulation of neuroblasts in the ventricular zone of cerebral cortex (Eksioglu *et al.*, 1996; Fox *et al.*, 1998; Sheen *et al.*, 2001; Moro *et al.*, 2002). FLNA co-localizes to actin stress fibers, highly expressed by neural cells in the ventricular surface, is thought to crosslink F-actin network to facilitate cell motility (Fox *et al.*, 1998; Stossel *et al.*, 2001). The inability of neurons to initiate migration following FLNA mutations is indicative of the significance of actin dynamics in initiation of migration. Whether FLNA's interactions with cell surface integrin receptors ( $\beta 1$  or  $\beta 2$ ), presenilin, and small GTPase RalA is part of the cascade that conveys extracellular signals from the ventricular zone to initiate migration still needs further examination (Sharma *et al.*, 1995; Loo *et al.*, 1998; Zhang and Galileo, 1998; Ohta *et al.*, 1999). However, Filamin A interacting protein (FLIP) is expressed in the ventricular zone and degrades FLNA, thereby inhibiting premature onset of neuronal migration from the ventricular zone (Nagano *et al.*, 2002).

### Maintenance of Migration

Once initiated, a dynamic regulation of multiple cellular events such as cell–cell recognition, adhesion, transmembrane signaling, and cell motility events underlies the process of neuronal migration (Lindner *et al.*, 1983; Grumet *et al.*, 1985; Antonicek *et al.*, 1987; Chuong *et al.*, 1987; Rutishauser and Jessell, 1988; Edmondson *et al.*, 1988; Sanes, 1989; Hatten and Mason, 1990; Stitt and Hatten, 1990; Takeichi, 1991; Misson *et al.*, 1991; Galileo *et al.*, 1992; Grumet, 1992; Komuro and Rakic, 1992, 1993, 1995; Shimamura and Takeichi, 1992; Fishman and Hatten, 1993; Hatten, 1993; Cameron and Rakic, 1994; Rakic *et al.*, 1994; Stipp *et al.*, 1994; Rakic and Komuro, 1995). A migrating neuron attaches itself to the radial glial substrate primarily by its leading process and cell soma. Only the actively migrating neurons form the specialized junctional domains or the interstitial densities with the apposing glial fibers (Gregory *et al.*, 1988; Cameron and Rakic, 1994), whereas the stationary neurons form desmosomes or puncta adherentia. The specialized subcellular accumulations of membrane proteins, such as radial glial based neuron–glial junctional protein 1 (NJPA1) or neuronal astrotactin, at the apposition of migrating neurons and radial glial cells may function in migration by orchestrating cell–cell recognition, adhesion, transmembrane signaling, and or motility. The homophilic or heterophilic nature of the junctional domain antigen interactions are unclear. However, the integrity of neuron–glial junctional complexes appears to depend on their association with microtubule cytoskeleton (Gregory *et al.*, 1988; Cameron and Rakic, 1994). Disruption of microtubules, but not actin filaments, adversely affect neuron–glial adhesion (Rivas and Hatten, 1995).

Junctional domain associated microtubules are thought to play a role in force generation during cell movement, in addition to being vital for the elaboration and maintenance of junctional domains (Gregory *et al.*, 1988). Furthermore, specific cell–cell interactions between migrating neurons and radial glial cells mediated by the junctional domain antigens may also modulate the properties of each other's cytoskeleton, akin to that observed between developing peripheral axons and Schwann cells (Kirkpatrick and Brady, 1994). It is argued that an increase in class III  $\beta$ -tubulin content leads to enhanced microtubule lability, thus allowing the continuous assembly and disassembly of microtubules needed to generate a forward force during cell movement (Falconer *et al.*, 1992; Moskowitz and Oblinger, 1995; Rivas and Hatten, 1995; Rakic *et al.*, 1996).

Significant deficits in neuronal migration were seen following mutations in genes regulating microtubule cytoskeleton (see Table 1 and Fig. 2). In humans, mutations in Lis1 (nonglycosylated subunit of platelet-activating factor acetylhydrolase isoform 1b) Miller–Dieker syndrome, a severe form of lissencephaly (Reiner *et al.*, 1993; Hattori *et al.*, 1994). In mouse, truncation of Lis1 leads to slower neuronal migration and cortical plate disorganization characterized by unsplit preplate (Cahana *et al.*, 2001). Partial loss of Lis1 (i.e., mice with one inactive allele of Lis1) also results in retarded neuronal migration (Hirosune *et al.*, 1998). Lis1 binds to microtubules, microtubule based motor protein, dynein, and related microtubule interactors such as dynactin, NUDEL, and mNudE (Sapir *et al.*, 1997; Efimov and Morris, 2000; Faulkner *et al.*, 2000; Kitagawa *et al.*, 2000; Niethammer *et al.*, 2000; Sasaki *et al.*, 2000; Smith *et al.*, 2000). Loss of Lis1 leads to concentration of microtubules around the nucleus and failed dynein aggregation, whereas overexpression of Lis1 causes transport of microtubule to edges of the cell and aggregation of dynein and dynein (Sasaki *et al.*, 2000; Smith *et al.*, 2000). NUDEL and mNudE appear to control cellular localization of dynein and the microtubule network around the microtubule-organizing centrosome, respectively (Feng *et al.*, 2000; Niethammer *et al.*, 2000; Sasaki *et al.*, 2000). How these associations of Lis1 modify the microtubule network to enable the nuclear translocation of neurons in the developing cortex remains to be elucidated.

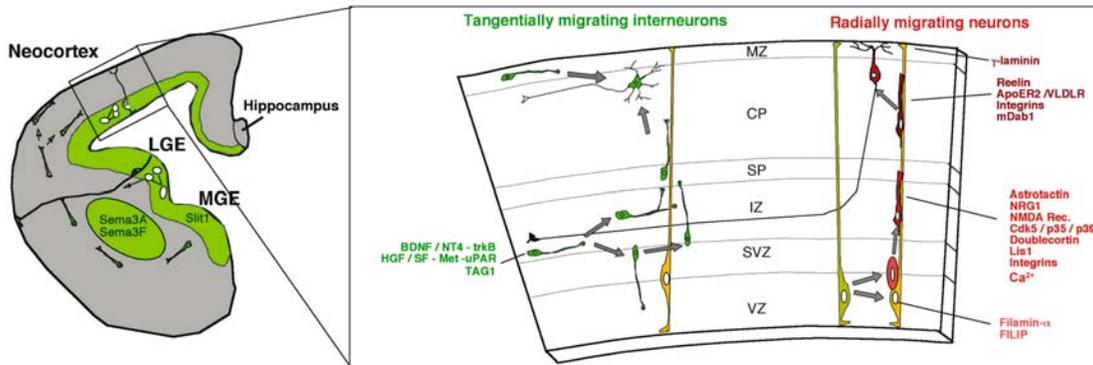
Mutations in another microtubule-associated protein in migrating neurons, doublecortin (Dcx), leads to X-linked lissencephaly (double cortex syndrome) in humans (des Portes *et al.*, 1998; Gleeson *et al.*, 1998). In these patients, neurons that migrated aberrantly are deposited in a broad band in subcortical layers. Dcx is critical for the stabilization of microtubule network (Francis *et al.*, 1999; Gleeson *et al.*, 1999; Horesh *et al.*, 1999). Dcx can associate with Lis1 and promote tubulin polymerization *in vitro* (Gleeson *et al.*, 1999; Caspi *et al.*, 2000; Feng and Walsh, 2001). Overexpression of Dcx results in aggregates of thick microtubule bundles resistant to depolymerization (Gleeson *et al.*, 1999). Structural analysis of Dcx indicates that it contains a  $\beta$ -grasp superfold motif that can bind to tubulin and facilitate microtubule polymerization and stabilization (Taylor *et al.*, 2000). Point mutations within this tubulin-binding motif were seen in patients with double cortex syndrome (Gleeson *et al.*, 1999).

TABLE 1. Molecular Cues Affecting Radial and Tangential Neuronal Migration

Gene	Function and phenotype	References
Reelin	Essential for layer formation Inverted cortical layers	D'Arcangelo <i>et al.</i> , 1995; Hirotsume <i>et al.</i> , 1995; Ogawa <i>et al.</i> , 1995; Hong <i>et al.</i> , 2000
VLDLR/ ApoER2	Reelin receptors, critical for layer formation Inverted cortical layers	Trommsdorf <i>et al.</i> , 1999
mDab1	Adaptor protein component of reelin signaling cascade Inverted cortical layers	Howell <i>et al.</i> , 1997; Sheldon <i>et al.</i> , 1997; Ware <i>et al.</i> , 1997
Astrotactin	Promotes neuron–radial glial adhesion Decreased rate of radial migration in mutants	Adams <i>et al.</i> , 2002
Doublecortin	Critical for the stabilization of microtubule network Mutations lead to X-linked lissencephaly (double cortex syndrome) in humans	des Portes <i>et al.</i> , 1998; Gleeson <i>et al.</i> , 1998, 1999; Francis <i>et al.</i> , 1999; Horesh <i>et al.</i> , 1999
Lis1	Binds to microtubules, microtubule based motor protein, dyenin, and related microtubule interactors such as dynactin, NUDEL, and mNudE Mutations in Lis1 cause Miller–Dieker syndrome, a severe form of lissencephaly	Reiner <i>et al.</i> , 1993; Hattori <i>et al.</i> , 1994; Sapir <i>et al.</i> , 1997; Efimov and Morris, 2000; Faulkner <i>et al.</i> , 2000; Niethammer <i>et al.</i> , 2000; Sasaki <i>et al.</i> , 2000; Smith <i>et al.</i> , 2000; Kitagawa <i>et al.</i> , 2000
Cdk5	Facilitates neuronal migration to CP following splitting of the preplate Inverted layering in mutants	Oshima <i>et al.</i> , 1996; Gilmore <i>et al.</i> , 1998
p35/p39	Regulatory subunits of Cdk5 Similar in function to Cdk5	Chae <i>et al.</i> , 1997; Kwon and Tsai, 1998; Ko <i>et al.</i> , 2001
NRG1	Promotes radial migration Critical for the establishment of radial glial scaffold	Anton <i>et al.</i> , 1999 Schmid <i>et al.</i> , 2003
$\alpha_3$ Integrin	Abnormal neuronal migration and laminar organization of cortex Neuron–glia interaction impaired Premature radial glial transformation	Anton <i>et al.</i> , 1999; Kreidberg <i>et al.</i> , 1996
$\alpha_6$ Integrin	Lethal at birth Disorganized CP Ectopic neuroblasts in embryonic cortex Disorganized basal lamina assembly	Georges-Labouesse <i>et al.</i> , 1996, 1998
$\alpha_3, \alpha_6$ Integrin	No neural tube closure Abnormal basal lamina assembly Multiple neuroblast ectopias in cortex	De Arcangelis <i>et al.</i> , 1999
$\alpha_v$ Integrin	Survive until E14 or birth Intracortical hemorrhage Facilitates basic neuron–glial adhesion	Bader <i>et al.</i> , 1998
$\beta_1$ Integrin (cond.)	Disrupted cortical laminar organization Radial glia end feet and pial basement membrane abnormalities	Graus-Porta <i>et al.</i> , 2001
$\gamma_1$ Laminin	Component of ECM in pial surface Misplaced neurons in CP	Halfter <i>et al.</i> , 2002
Filamin $\alpha$	Actin-binding protein, co-localizes to actin stress fibers, highly expressed by neural cells in the ventricular surface, crosslinks F-actin network to facilitate cell motility Needed to initiate migration from the ventricular zone Mutations cause periventricular heterotopia	Fox <i>et al.</i> , 1998; Sheen <i>et al.</i> , 2001; Moro <i>et al.</i> , 2002
FILIP	Regulates degradation of Filamin $\alpha$ in the ventricular zone Prevents premature onset of migration	Nagano <i>et al.</i> , 2002
Dlx1/2	Transcription factors regulating the differentiation of cortical and hippocampal interneurons from the subpallium Lateral ganglionic eminence (LGE)/medial ganglionic eminence (MGE)	Anderson <i>et al.</i> , 1997; Pleasure <i>et al.</i> , 2000

TABLE 1. (Continued)

Gene	Function and phenotype	References
Nkx 2.1	Transcription factor regulating the migration and differentiation of cortical interneurons from the MGE	Sussel <i>et al.</i> , 1999; Anderson <i>et al.</i> , 2001
TAG1	Neural cell-adhesion molecule expressed in corticofugal fibers Motogenic cue for tangentially migrating interneurons	Wolfer <i>et al.</i> , 1994; Denaxa <i>et al.</i> , 2001
HGF	Motogenic factor expressed in GE Promotes movement of cortical interneurons from MGE toward dorsal pallium	Powell <i>et al.</i> , 2001
u-PAR	Urokinase-type plasminogen activator receptor Enables HGF activation	Powell <i>et al.</i> , 2001
BDNF, NT-4	Motogenic factors for neuronal migration from MGE	Brunstrom <i>et al.</i> , 1997; Ringstedt <i>et al.</i> , 1998; Polleux <i>et al.</i> , 2002
TrkB	Receptor for BDNF/NT-4 Mutation leads to reduced interneuronal migration into cortex	Brunstrom <i>et al.</i> , 1997; Polleux <i>et al.</i> , 2002
Slit 1/2	Chemorepellent for GABAergic interneurons in the GE	Zhu <i>et al.</i> , 1999; Marin <i>et al.</i> , 2003
Sema3A/3F	Chemorepellent expressed in striatal mantle region Helps to channel cortical interneurons toward the cortex	Marin <i>et al.</i> , 2001; Tamamaki <i>et al.</i> , 2003
Nrp1/2	Receptors for class 3 secreted semaphorins Enables cortical interneurons to migrate away from striatum into the cortex	Marin <i>et al.</i> , 2001; Tamamaki <i>et al.</i> , 2003



**FIGURE 2.** Molecular cues influencing distinct patterns of migration into the developing cerebral cortex. In the cerebral wall, neurons migrating tangentially into the cerebral cortex from ganglionic eminence and neurons migrating radially from the ventricular surface to the cortical plate are influenced by different sets of molecules on the right hand side panel. LGE-Lateral ganglionic eminence, MGE-medial ganglionic eminence, VZ-ventricular zone, SVZ-subventricular zone, IZ-intermediate, SP-subplate, CP-cortical plate, MZ-marginal zone.

Regulators of both actin and microtubule network associate with cyclin-dependent kinase-5 (Cdk5), expressed in migrating neurons and axon growth cones of the developing cortex (Nikolic *et al.*, 1998). Both filamin 1 and NUDEL are putative substrates for Cdk5 phosphorylation (Fox *et al.*, 1998; Niethammer *et al.*, 2000; Feng and Walsh, 2001). Mice deficient in Cdk5 and its activating subunits, p35 and p39, display abnormal neuronal migration and placement in cerebral cortex (Ohshima *et al.*, 1996; Chae *et al.*, 1997; Gilmore *et al.*,

1998; Kwon and Tsai, 1998; Ko *et al.*, 2001). Deficits in Brn1 and 2, class III POU domain transcription factors regulating p35 and p39 expression, also lead to cortical migrational abnormalities (McEvelly *et al.*, 2002). Interactions between p35 and  $\beta$ -catenin is thought to enable Cdk5 to regulate negatively *N*-cadherin-mediated adhesion and facilitate neuronal migration through the *N*-cadherin-rich developing cerebral wall up to the CP (Redies and Takeichi, 1993; Kwon *et al.*, 2000).

Transient, intracellular calcium fluxes that modulate neuronal migration *in vitro* can significantly influence the actin and microtubule network of neurons undergoing oriented migration (Rakic *et al.*, 1994). The link between extracellular cues modulating migration and the internal cues involved in mechanics of migration is generated in a highly varied and redundant manner. For example, neurotransmitter receptors such as *N*-methyl-D-aspartate (NMDA) type glutamate receptors, and GABA receptors, or growth factors and their receptors such as neuregulins and its receptors erbB2, erbB3, and erbB4, or BDNF and its high-affinity receptor trkB, can promote radial-guided neuronal migration (Komuro and Rakic, 1993, 1996; Anton *et al.*, 1997; Rio *et al.*, 1997; Behar *et al.*, 2000, 2001). The most direct transmission of external cues via membrane receptors to cytoskeletal changes during migration is provided by integrins. Integrins are heterodimeric cell surface receptors that serve as structural links between the ECM and the internal cytoskeleton. Different integrin receptors display different adhesive properties, regulate different intracellular signal transduction pathways, and thus, different modes of adhesion-induced changes in cell physiology. Integrins are also capable of synergizing with other cell surface receptor systems to finely modulate a cell's behavior in response to multiple environmental cues. Developmental changes in the cell surface integrin repertoire and function may thus modulate distinct aspects of neuronal migration in the developing cerebral cortex by altering the strength and ligand preferences of cell–cell adhesion during development. Different  $\alpha$  integrin subunits dimerize preferentially or exclusively with  $\beta_1$  integrin, which is ubiquitously expressed in the developing cerebral cortex. The varied, yet distinct, cortical phenotypes of integrin subunit null mice provide striking insights into the distinct roles that cell–cell, cell–ECM adhesive interactions play in neuronal migration.

Mice homozygous for a targeted mutation in the  $\alpha_3$  integrin gene die during the perinatal period with severe defects in the development of the kidneys, lungs, skin, and cerebral cortex (Kreidberg *et al.*, 1996; Anton *et al.*, 1999). In the cerebral cortex, the normal laminar organization of neurons is lost, and neurons are positioned in a disorganized pattern. The  $\alpha_3$  integrin modulates neuron–glial recognition cues during neuronal migration and maintain neurons in a gliophilic mode until glial-guided neuronal migration is over and layer formation begins (Anton *et al.*, 1999). The gliophilic to neurophilic switch in the adhesive preference of developing neurons in the absence of  $\alpha_3$  integrin was hypothesized to underlie the abnormal cortical organization of  $\alpha_3$  integrin mutant mice. In contrast to  $\alpha_3$  integrin,  $\alpha_v$  integrins appear to provide optimal levels of basic cell–cell adhesion needed to maintain neuronal migration and differentiation. Substantial disruption of cellular organization in cerebral wall and lateral ganglionic eminence (LGE) is seen at E11–12 in  $\alpha_v$  null mice. Extensive intracerebral hemorrhage in  $\alpha_v$  deficient mice, beginning at E12–13, prevents further evaluation of cortical development in late surviving (until birth)  $\alpha_v$  null mice (Bader *et al.*, 1998). The  $\alpha_v$  integrins expressed on radial glial cell surface can potentially associate with at least five different  $\beta$  subunits,  $\beta_1$ ,  $\beta_3$ ,  $\beta_5$ ,  $\beta_6$ , and  $\beta_8$ . Adhesive interactions involving fibronectin, vitronectin, tenascin, collagen, or laminin, ECM

molecules that are found in the developing cerebral wall, can be mediated through these  $\alpha_v$ -containing integrins (Cheresh *et al.*, 1989; Bodary and McLean, 1990; Moyle *et al.*, 1991; Hirsch *et al.*, 1994). Both transient cell–matrix interactions and cell–anchoring mechanisms that are mediated by different  $\alpha_v$ -containing integrins and their respective ligands are likely to modulate the process of neuronal translocation in cerebral cortex.

In addition to  $\alpha_3$  integrin, some laminin isoforms in the developing cerebral cortex can also interact with  $\alpha_6$  integrin dimers (Georges-Labouesse *et al.*, 1998). The  $\alpha_6$  null mice die at birth (Georges-Labouesse *et al.*, 1996) with abnormal laminar organization of the cerebral cortex and retina (Georges-Labouesse *et al.*, 1998). Analysis of E13.5–E18.5  $\alpha_6$  integrin-deficient embryos revealed ectopic neuronal distribution in the cortical plate, protruding out to the pial surface. The CP was further disorganized by wavy neurite outgrowth of ectopic neuroblasts. Coinciding abnormalities of laminin synthesis and deposition also occurs in the mutant brain. Persistence of glial laminin throughout development may have prevented neuroblasts from appropriately arresting their migration in the developing CP in  $\alpha_6$  null mice. Since cerebral cortex still formed in  $\alpha_6$  mutants, albeit abnormally, other integrin dimers may have overlapping functions with  $\alpha_6$  integrins during early cortical development. The similarities in the ligand preferences of  $\alpha_3$  and  $\alpha_6$  integrins are suggestive of potential functional overlap. The severe and novel cortical abnormalities in  $\alpha_3$ ,  $\alpha_6$  double knockout mutants, that is, disorganization of CP with large collection of ectopias, aberrant basal lamina organization, and abnormal choroid plexus, suggest a synergistic role for  $\alpha_3$  and  $\alpha_6$  integrins during cortical development (De Arcangelis *et al.*, 1999). Deficiency in  $\beta_4$  integrin, which only associates with  $\alpha_6$ , leads to an identical cortical phenotype. Mutations in  $\alpha_6$  or  $\beta_4$  integrin in humans results in skin blistering (epidermolysis bullosa). However, the brain phenotype of the affected patients is unknown.

The  $\beta_1$  integrin in the cerebral cortex can dimerize with at least 10 different  $\alpha$  subunits; thus  $\beta_1$  integrin deficiency leads to lethality from around E5.5 (Fassler and Meyer, 1995; Stephens *et al.*, 1995). Most of the cortical-specific  $\alpha$  subunits seem to dimerize only with  $\beta_1$  integrin. To study the role of  $\beta_1$  integrin in the developing cortex,  $\beta_1$  integrin-floxed mice were crossed with nestin-cre mice, resulting in widespread inactivation of  $\beta_1$  integrins in cortical neurons and glia from around E10.5 (Graum-Porta *et al.*, 2001). Cortical layer formation is disrupted in these mice, in large part as a result of defective meningeal basement membrane assembly, marginal-zone formation, and glial end feet anchoring at the top of the cortex. BrdU birthdating studies suggest that glial-guided neuronal migration is not affected significantly. However, perturbed radial glial end feet development may contribute to the defective placement of neurons in the cortex. The varied cortical phenotypes of  $\alpha_1$ ,  $\alpha_3$ ,  $\alpha_6$ ,  $\alpha_v$ , and  $\beta_1$  null mice may reflect the transdominant, transnegative, or compensatory influences distinct integrin receptor dimers may exert over each other and the ECM ligands in the developing cerebral cortex. *In vitro*, binding of a ligand to a signal transducing integrin or inactivation of signaling through a particular integrin can initiate a unidirectional signaling cascade affecting

the function of the target integrin in the same cell (Simon *et al.*, 1997; Hodivala-Dilke *et al.*, 1998; Blystone *et al.*, 1999). Elucidation of whether such integrin crosstalk regulates patterns of neuronal development and interactions with specific ECM molecules in the developing cortices of different integrin null mice will be informative in fully characterizing the role of integrins in neuronal migration.

### Termination of Migration

Once neurons reach the top of the CP, the movement of neurons stops abruptly at the interface between the CP and the cell sparse marginal zone and cohorts of neurons begin to assemble into their respective layers. This final stage of neuronal migration is the least explored aspect of neuronal migration, in spite of its significance for genetic and acquired cortical malformations (Rakic, 1988; Rakic and Caviness, 1995; Olson and Walsh, 2002). The signal to terminate neuronal cell migration is thought to be provided either by the afferent fibers that migrating neurons encounter near their target location or by the ambient neuronal cell population that had already reached its final position (Sidman and Rakic, 1973; Hatten and Mason, 1990; D’Arcangelo *et al.*, 1995; Ogawa *et al.*, 1995). Alternatively, a change in the cell surface properties of the radial glial substrate at the top of the CP may signal a migrating neuron to stop, detach, and differentiate.

In the *reeler* mouse, deficits in this phase of migration have led to disorganized, inverted cortex, with early-born neurons occupying abnormally superficial positions and later-born neurons adopting abnormally deep positions (Caviness *et al.*, 1972; Caviness and Sidman, 1973; Lambert de Rouvroit and Goffinet, 1998). The inversion of final neuronal positions in the CP of the *reeler* mouse has made it a prototype model for the analysis of mechanisms controlling the final phase of neuronal migration, that is, how neurons disengage from a migratory mode to assemble into distinct layers. The *reeler* locus encodes Reelin, a 388 kDa secreted protein composed of a unique N-terminal sequence with similarity to F-spondin, followed by a series of eight 350–390 amino acid “Reelin repeats” each containing an EGF domain with homology to ECM proteins like Tenascin C (D’Arcangelo *et al.*, 1995; Hirotsune *et al.*, 1995). Reelin acts on noncell autonomously (Ogawa *et al.*, 1997), and the protein is synthesized and secreted in the cerebral cortex predominantly by the Cajal–Retzius (CR) cell of the marginal zone, the outermost layer of the developing cortex (D’Arcangelo *et al.*, 1995; Ogawa *et al.*, 1995).

Mutations in three molecules, VLDLR, ApoER2, and Dab1, have been found to phenocopy almost exactly the effects of the *reeler* gene mutation, suggesting that the corresponding proteins represent a reelin regulated biochemical pathway that mediates proper termination of neuronal migration and formation of cerebral cortical lamination (Gonzalez *et al.*, 1997; Howell *et al.*, 1997; Sheldon *et al.*, 1997; Ware *et al.*, 1997). The *dab1* gene encodes a cytoplasmic adapter protein (Dab1) expressed by neurons in the developing CP, suggesting that Dab1 represents a link in the signaling pathway that receives the Reelin

signal. This idea is confirmed by observation that Reelin expression is normal in the *dab1* mutant cortex (Gonzalez *et al.*, 1997) but Dab1 protein accumulates in the *reeler* mouse brain (Rice *et al.*, 1998) and Dab1 is phosphorylated in response to application of recombinant Reelin (Howell *et al.*, 1999a). Mammalian Dab1 was identified through a two-hybrid screen using the non-receptor tyrosine kinase Src as “bait” (Howell *et al.*, 1997) and found to have homology with *Drosophila disabled* (Gertler *et al.*, 1993). Dab1 has an N-terminal alpha helical structure and the critical amino acids of a protein interaction/phosphotyrosine-binding domain (PI/PTB) (Kavanaugh and Williams, 1994; Borg *et al.*, 1996; Margolis, 1996; Howell *et al.*, 1997). The PI/PTB domain of Dab1 binds proteins that contain an NPXY motif (Howell *et al.*, 1997, 1999b; Trommsdorff *et al.*, 1998) a motif that has been implicated in clathrin-mediated endocytosis (Chen *et al.*, 1990), and integrin signaling (Law *et al.*, 1999).

More recently, mice with compound mutations in both VLDLR and ApoER2 have been found to have a phenotype indistinguishable from *reeler* and *dab1* mutants (Trommsdorff *et al.*, 1999). VLDLR and ApoER2 are members of the low density lipoprotein (LDL) receptor superfamily that interacted with Dab1 in two-hybrid screens through the PI/PTB domain of Dab1 and the NPXY motif of LDL superfamily members (Trommsdorff *et al.*, 1998). The NPXY motif of LDL receptor family members is essential for clathrin-mediated endocytosis (Chen *et al.*, 1990). The implication of VLDLR and ApoER2 as potential Reelin receptors was surprising since LDL superfamily members are well characterized as mediating the endocytosis of specific ligands, but have never demonstrated a direct signaling function. Recent studies, however, have clearly demonstrated that both recombinant ApoER2 and the VLDLR bind Reelin and that this binding leads both to the tyrosine phosphorylation of Dab1 and in the case of VLDLR, the internalization of the receptor and Reelin (D’Arcangelo *et al.*, 1999; Hiesberger *et al.*, 1999). Thus there is compelling evidence that Reelin, VLDLR, ApoER2, and Dab1 function in a common signaling pathway between CR cells and CP neurons, but the downstream molecules that mediate Reelin signaling effect on either migration or adhesion of cortical neurons remains unclear.

Reelin’s effect on cortical layering is hypothesized to result from three distinct cellular effects. First, reelin may regulate CP organization by initiating the splitting of preplate into marginal zone and subplate. Failure of this process in *reeler* mutants leads to the accumulation of cortical neurons underneath the preplate neurons. Second, a reelin gradient may act as an attractant for neurons to the top of the CP, thus enabling newly generated neurons to migrate past earlier generated ones in the developing CP. Third, reelin may induce detachment of neurons from their radial glial guides and thus end neuronal migration at the marginal zone-developing CP interface and initiate the differentiation of neurons into distinct layers.

Cortical neurons in  $\beta_1$  integrin or laminin  $\gamma_1$  nidogen-binding site (Halfter *et al.*, 2002) deficient mice invade the marginal zone in areas devoid of reelin producing CR cells, and in regions with CR cell ectopias, accumulate underneath them, within the CP. Invasion of neurons into areas devoid of

reelin-producing CR cells supports a role for reelin in normal termination of neuronal migration. Furthermore, reelin appears to facilitate detachment of migrating neurons from glial guides *in vitro* and in the rostral migratory stream (RMS) (Hack *et al.*, 2002). The reelin-induced detachment of embryonic cortical neurons from glial guides *in vitro* depends on  $\alpha_3$  integrin signaling. It is hypothesized that during glial-guided migration to the CP neuronal  $\alpha_3$  integrin may interact with glial cell surface molecules such as fibronectin or laminin-2, and at the top of the CP, the ligand preference of  $\alpha_3$  integrins may change from radial glial cell surface ECM molecules to reelin. Different ligands or ligand concentration can determine the surface levels of integrins by regulating the rate at which integrin receptor is removed from the cell surface. Ligands can also regulate polarized flow of integrins toward or away from growth cone membranes. Reelin can also function as serine protease and degrade fibronectin and laminin normally used to maintain glial-based migration (Quattrocchi *et al.*, 2002). Thus changes in the availability, function, and ligand preference of  $\alpha_3$  integrins or reelin proteolytic activity may trigger the decrease in a migrating neuron's bias for gliophilic adhesive interactions and promote neurophilic interactions needed for neurons to detach from radial glial guides and organize into distinct layers. Interestingly, deficiencies in  $\alpha_3$  integrin ligands, laminin-2 and reelin lead to cortical anomalies such as polymicrogyria or lissencephaly (Sunada *et al.*, 1995; Hong *et al.*, 2000).

## TANGENTIAL MIGRATION IN THE FOREBRAIN

As introduced earlier in this chapter, two main types of migration are classically opposed during the development of the central nervous system: radial vs tangential migration. Radial migration is characterized by close interactions between migrating neurons and the processes of radial glial cells which constitute a scaffold bridging the proliferating neuroepithelium and the differentiating zone. By definition, tangential migration is referred to as a nonradial mode of neuronal translocation that does not require specific interaction with radial glial cell processes. Until recently, the predominant view was that the vast majority of neurons in the forebrain were generated through radial migration (Sidman and Rakic, 1973). The first evidence to suggest the need for a revised model came from observations of tangential dispersion of precursors or postmitotic neurons in the developing cortex (O'Rourke *et al.*, 1992a, 1995; Fishell *et al.*,

1993a; Tan and Breen, 1993; Tan *et al.*, 1995; de Carlos *et al.*, 1996). The widespread distribution of clonally related cells also suggested the possibility of non-radial migration in the cortex (Walsh and Cepko, 1992). In an elegant study, Parnavelas and collaborators coupled retroviral-mediated lineage-tracing studies with the determination of neuronal subtype identity and demonstrated a tight correlation between cell dispersion and neuronal subtype (Parnavelas *et al.*, 1991): most excitatory, glutamatergic pyramidal neurons are produced locally by a set of precursors migrating radially in the cortex, whereas most GABAergic, nonpyramidal neurons were produced by a set of progenitors migrating tangentially (Parnavelas, 2000).

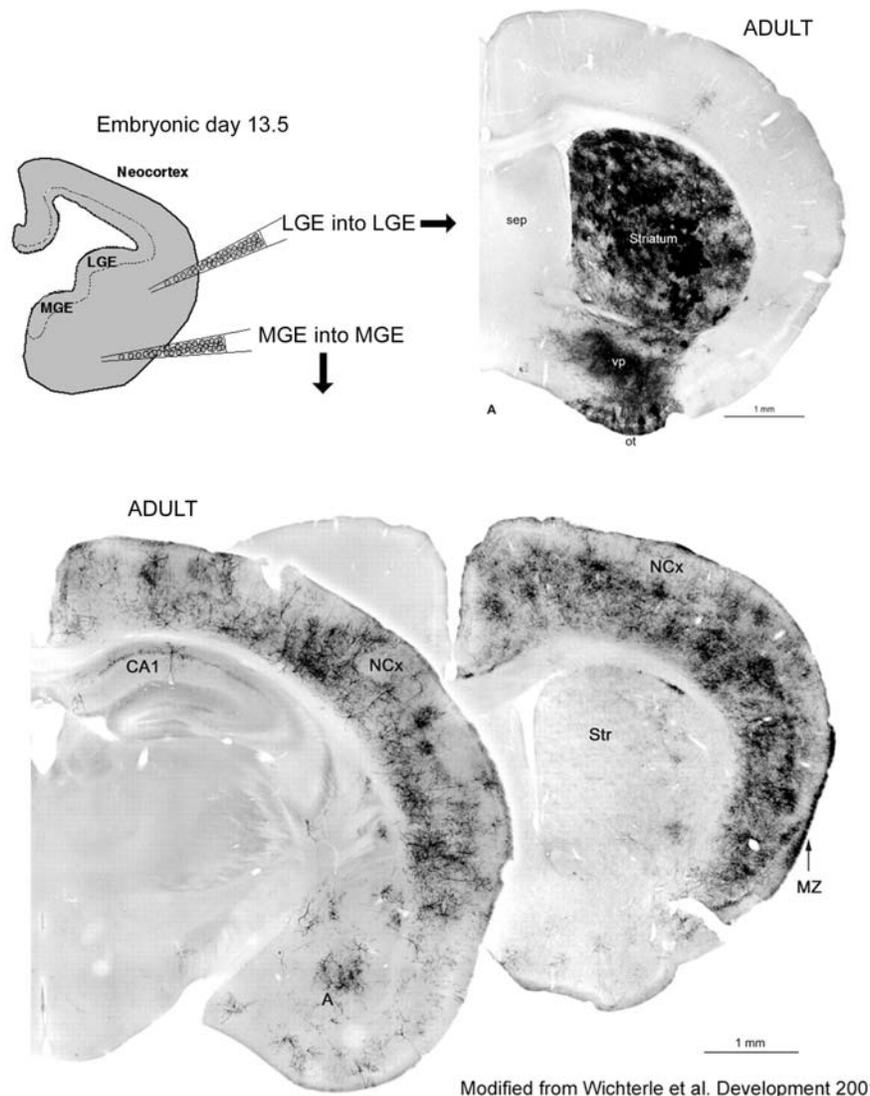
## Origin of Tangentially Migrating Cells in the Forebrain

The source and destination of these tangentially migrating cells, however, remained a mystery until experiments by Anderson *et al.* suggested that neurons migrated from the GE to the cortex where they gave rise preferentially to GABAergic interneurons (Anderson *et al.*, 1997; Tamamaki *et al.*, 1997). This conclusion is based mainly upon the observation that there are virtually no neocortical GABAergic neurons in *Dlx1/2* double knockout mice, two homeobox transcription factors expressed in the ventricular and subventricular zones of the GE (Anderson *et al.*, 1997). The GE is located in the ventral part of the telencephalon and is producing neurons of the basal ganglia (Fentress *et al.*, 1981; Qiu *et al.*, 1995). This ventral structure can be divided into three subregions using neuroanatomical and molecular criteria: the medial, the lateral, and the caudal parts (Corbin *et al.*, 2000). Several transcription factors are differentially expressed in these three regions (Table 2).

Recent *in utero* homotopic transplantation experiments performed in mice have revealed that these distinct regions give rise to specific neuronal populations displaying strikingly different patterns of cell migration (Fig. 3): the *medial* GE gives rise to the majority of GABAergic interneurons of the cortex and hippocampus (Lavdas *et al.*, 1999; Anderson *et al.*, 2001; Wichterle *et al.*, 2001; Polleux *et al.*, 2002) whereas precursors in the *lateral* GE generates projecting medium spiny neurons of the striatum, nucleus accumbens and olfactory tubercle and to the granule and periglomerular cells in the olfactory bulb (Wichterle *et al.*, 2001). The pattern of migration of neurons originating in the caudal GE is less well characterized but it has recently been shown that precursors in this region give rise to interneurons found in layer 5 of the neocortex, various regions of the limbic system and also neurons of the striatum (Nery *et al.*, 2002).

TABLE 2. Transcription Factors Expression in Different Subregions of the Ganglionic Eminence

	<i>Mash1</i>	<i>Dlx1/Dlx2</i>	<i>Nkx2.1</i>	<i>Lhx6</i>	<i>Gsh2</i>
MGE	+	+	+	+	–
LGE	+	+	–	–	+
CGE	+	+	?	?	?



Modified from Wichterle et al. *Development* 2001

**FIGURE 3.** Generation and migration of cortical interneurons from the medial ganglionic eminence. Dissociated neurons (tagged with alkaline phosphatase) isolated from LGE or MGE were transplanted homotopically into LGE or MGE, respectively, at early stages of neuronal migration in cortex. Location and differentiation of transplanted neurons were analyzed in adult brains. Strikingly, MGE cells all migrated into cerebral cortex to become cortical interneurons, whereas LGE cells populated the striatum. LGE, lateral ganglionic eminence; MGE, medial ganglionic eminence. Modified with permission from Wichterle *et al.*, 2001.

### Cellular and Molecular Substrates for Tangential Migration of Cortical Interneurons

Tangentially migrating interneurons display a characteristic unipolar morphology during translocation with a long leading process dragging behind their nucleus (Fig. 2) (Anderson *et al.*, 1997; Tamamaki *et al.*, 1997; Polleux *et al.*, 2002). Interneurons are migrating tangentially through the intermediate zone or the marginal zone, two axon-rich layers located, respectively, deep and superficial, relative to the CP, where all neurons accumulate in a layer-specific manner to undergo their terminal differentiation (O'Leary and Nakagawa, 2002).

During migration to the cortex, tangentially migrating interneurons are not using radial glial cells processes as a scaffold during translocation and these cells do not appear to

fasciculate along a specific cellular scaffold (Polleux *et al.*, 2002) although it has been proposed that they interact with corticofugal axons (Denaxa *et al.*, 2001). *In vitro*, the neural cell-adhesion molecule TAG-1 (also called contactin-2) expressed by corticofugal axons has been shown to play a role in the control of interneuron migration.

### Extracellular Cues Regulating Tangential Migration in the Forebrain

The extracellular cues controlling the tangential migration of interneurons from the GE to the cortex can be classified in three categories: (1) extracellular cues regulating their motility (motogenic cues), (2) directional cues guiding their migration

toward the appropriate territories, and (3) stop-signals abolishing their motility and therefore dictating where interneurons should terminally differentiate.

### Cues Controlling the Motility of Tangentially Migrating Interneurons

Several factors expressed along the migrating pathway of cortical interneurons have recently been shown to be potent stimulators of interneurons motility. Both the hepatocyte growth factor (HGF, also called scatter factor) and the neurotrophin NT4/5 are expressed in the cortex during mouse embryogenesis and are potent stimulators of interneurons migration (Behar *et al.*, 1997; Brunstrom *et al.*, 1997; Powell *et al.*, 2001; Polleux *et al.*, 2002). Neurons migrating tangentially from the MGE to the cortex express c-Met and trkB, the high-affinity receptors for HGF and NT4, respectively. Furthermore, mice mutant for urokinase-type plasminogen activator receptor (u-PAR), a key component of HGF activation, exhibit reduced interneuron migration to the frontal and parietal cortex (Powell *et al.*, 2001). This decreased number of interneurons in the cortex of u-PAR knockout mice has important behavioral consequences on the establishment of the normal cortical circuitry characterized by an imbalanced level of excitation and inhibition which leads to epilepsy (Powell *et al.*, 2003). Mice presenting a targeted deletion of the tyrosine kinase receptor trkB, the high-affinity receptor of NT4, also present a significant reduction of the number of interneurons migrating from the MGE to the cortex (Polleux *et al.*, 2002). The motogen activity resulting from the activation of these tyrosine kinase receptors (c-Met and trkB) is likely to be mediated through their ability to activate phosphoinositide 3-(PI3-)kinase (Polleux *et al.*, 2002), a key regulator of cytoskeleton reorganization and cell motility in nonneuronal cell types (Iijima *et al.*, 2002).

### Guidance Cues (Sema 3A and Sema 3F; Slits)

Several axon guidance cues have been shown to play a role in directing interneuron migration from the GE to the cortex. The diffusible chemorepulsive Sema3A and Sema3F are expressed in the postmitotic mantle region of the developing striatum and migrating interneurons from the MGE express Neuropilin 1 (*Npn1*) and Neuropilin 2 (*Npn2*) (Marin *et al.*, 2001; Tamamaki *et al.*, 2003), Sema3A and -3F respective receptors (Chen *et al.*, 1997; Kolodkin *et al.*, 1997; Giger *et al.*, 1998). *In vitro* experiments demonstrate that MGE-derived interneurons are repulsed by Sema3A and Sema3F in a cooperative manner. Furthermore, the *in vivo* analysis of mice presenting targeted deletion of *Npn1* and *Npn2* demonstrate that they are required for the selective avoidance of the striatum by cortical interneurons and therefore for the directed migration to the cortex (Marin *et al.*, 2001; Tamamaki *et al.*, 2003).

Slit1 and Slit2, another short-range chemorepulsive cue for axons expressed in the ventricular zone of the GE as well as in the medial part of GE, has been shown to repulse MGE-derived interneurons *in vitro* (Zhu *et al.*, 1999). However, Slit 1/2 double knockout mice do not show any defect of guided

migration toward the cortex but nevertheless show a defect in the position of specific interneuronal population within the basal telencephalon, close to the midline (Marin *et al.*, 2003). The cortex exerts a chemoattractive activity on migrating interneurons but these cortex-derived cues remains to be identified.

Finally, membrane-bound cell-adhesion molecules, cadherins, delineate sharp territories of expression restricted to the dorsal telencephalon (*R*-Cadherin) and the LGE (Cadherin-6) in E10–11 developing mouse embryos. Evidence using both electroporation-mediated ectopic expression of cadherins or the *in vivo* analysis of Cadherin-6 knockout mice demonstrate its role in the appropriate sorting of striatal and cortical neuronal populations (Inoue *et al.*, 2001).

### Stop-Signals

Once migrating interneurons have reached the CP, they are targeting specific layers according to their birthdate just as radially migrating neurons do (Fairen *et al.*, 1986). So far, few molecules have been characterized for their capacity to stop the motility of tangentially migrating interneurons and even less is known about the putative cues that coordinate the layer-specific targeting of these two populations of neurons. Interestingly, several studies have shown that tangentially migrating neurons are expressing functional calcium-permeable AMPA receptors (but not NMDA receptors) which could be activated by glutamate released from corticofugal axons (Metin *et al.*, 2000) and/or GABA released from tangentially migrating interneurons themselves (Poluch and Konig, 2002). Both GABA and glutamate have been shown to control the motility of migrating neurons in the developing cortex (Behar *et al.*, 1996, 1998, 1999, 2000) and AMPA receptor activation leads to neurite retraction and is sufficient to stop migration of cortical interneurons in embryonic slice cultures (Poluch *et al.*, 2001). Because the neurotransmitter glutamate is expressed at high levels in the CP (Behar *et al.*, 1999), it could trigger an AMPA-receptor-dependent calcium influx that could act as a stop-signal for tangentially migrating interneurons in their final cortical environment. Further work will be necessary to validate this model meanwhile the identity of the cues leading to the coordinated, layer-specific accumulation of interneurons and excitatory glutamergic neurons remains mysterious and the center of a lot of attention.

### Differences between the Pattern of Tangential Migration in Rodents and Humans

There might be important differences between the pattern of tangential migration of GE-derived interneurons between rodent and human brain. Recent work demonstrate that a contingent of GE-derived interneurons migrate medially from the ventral telencephalon to the diencephalon in the human developing brain but not in nonhuman primate or in mouse embryos (Letinic and Rakic, 2001). Moreover, in the human brain retroviral lineage studies performed *in vitro* suggest that a substantial proportion of cortical

GABAergic neurons are generated in the dorsal telencephalon (Letinic *et al.*, 2002) in contrast with what is observed in the embryonic mouse telencephalon (Anderson *et al.*, 1997a). In the human embryonic brain, dividing precursors located in the dorsal telencephalon are expressing *Mash1* and *Dlx1/2* and have been shown to be competent to generate GABAergic interneurons in the cortical neuroepithelium of human-primates (Letinic *et al.*, 2002) but not in rodents (Fode *et al.*, 2000). This suggests that modifications in the expression pattern of transcription factors in the fore-brain could underlie species-specific programs for the generation of neocortical local circuit neurons (Letinic *et al.*, 2002).

## Other Structures Displaying Nonradial Migration

Many other regions of the developing central nervous system are characterized by nonradial neuronal migration, including the RMS of olfactory interneurons and the tangential migration of granule cells in the cerebellum.

### The Rostral Migratory Stream (RMS)

Precursors of the two populations of olfactory interneurons (periglomerular and granule cells) are not produced within the olfactory bulb but are generated by precursors located in the LGE during embryonic development (Altman, 1969; Lois and Alvarez-Buylla, 1993, 1994; Lois *et al.*, 1996; Dellovade *et al.*, 1998; Sussel *et al.*, 1999; Corbin *et al.*, 2000; Wichterle *et al.*, 2001). This migration is unique because it continues throughout adulthood in rodents providing a constant number of GABAergic neurons to the olfactory bulb. In the adult brain, olfactory interneurons are generated from the subependymal layer lining the lateral ventricles, a proliferative epithelium deriving from the subventricular zone of the embryonic GE (Doetsch and Alvarez-Buylla, 1996; Doetsch *et al.*, 1997, 1999a, b).

The interneurons migrating in the adult RMS are also unique with regard to their neurophilic rather than gliophilic mode of migration, requiring interactions between migrating interneurons (Lois and Alvarez-Buylla, 1994; Lois *et al.*, 1996). When explanted *in vitro*, these interneurons form chains by migrating along each other. This so-called *chain migration* is dependent of the expression of specific cell-adhesion molecules of the immunoglobulin superfamily such as the polysialylated form of neural cell-adhesion molecule (PSA-NCAM; reviewed in Marin *et al.*, 2003).

### Cerebellar Granule Cell Migration

Another population of interneurons migrates nonradially in the cerebellum: the granule cells migrating from the external granule layer (EGL) to the internal granular layer (IGL) during early postnatal stages of rodent development (reviewed in Hatten, 1999). The rate of migration of cerebellar granule neurons also is modulated through the control of intracellular calcium levels by activation of NMDA-, AMPA-, and somatostatin receptors

(Komuro and Rakic, 1992, 1993, 1996; Yacubova and Komuro, 2002). Activation of somatostatin receptors increases the rate of granule cell migration near their site of birth in the EGL, but decreases their rate of migration near their final destination in the IGL. Correspondingly, the size and frequency of spontaneous Ca<sup>2+</sup> fluctuations is enhanced by somatostatin in the early phase of migration, whereas spike-like Ca<sup>2+</sup> transients are eliminated by somatostatin in the late phase (Yacubova and Komuro, 2002).

This mode of migration is characterized by a dynamic switch between tangential and radial mode of migration: after translocation in the superficial EGL, granule interneurons make a sharp 90° turn to migrate along the radial processes of Bergmann glia spanning the molecular layer, to reach the deep IGL where they will undergo terminal differentiation (Komuro and Rakic, 1995). This switch from tangential to radial mode of translocation is not unique to cerebellar granule cells but is also observed for cortical interneurons (Polleux *et al.*, 2002) and is likely to reflect a basic property of migrating interneurons.

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