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Neocortical neurogenesis and neuronal migration

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Abstract

The neocortex, the evolutionarily newest part of the cerebral cortex, controls nearly all aspects of behavior, including perception, language and decision-making. It contains an immense number of neurons that can be broadly divided into two groups, excitatory neurons and inhibitory interneurons. These neurons are predominantly produced through extensive progenitor cell divisions during the embryonic stages. Moreover, they are not randomly dispersed, but spatially organized into horizontal layers that are essential for neocortex function. The formation of this laminar structure requires exquisite control of neuronal migration from their birthplace to their final destination. Extensive research over the past decade has greatly advanced our understanding of the production and migration of both excitatory neurons and inhibitory interneurons in the developing neocortex. In this review, we aim to give an overview on the molecular and cellular processes of neocortical neurogenesis and neuronal migration.

Introduction

A prominent trait of the neocortex is its complex yet well-organized cellular architecture, the formation of which relies on the production and positioning of its diverse neuronal populations. There are two major groups of neurons in the neocortex — glutamatergic excitatory neurons and -aminobutyric acid (GABA)-ergic inhibitory interneurons, which are responsible for generating excitation and inhibition, respectively. Excitatory neurons constitute the vast majority (~70–80%) of neocortical circuit neurons and are responsible for generating the output. On the other hand, inhibitory interneurons provide a rich variety of inhibitions that shape the output of functional circuits. Proper neocortex function critically depends on the production and positioning of a correct number of excitatory and inhibitory neurons, which largely occur during the embryonic stages.

With the advent of improved fate-mapping tools, including genetically engineered mice, excitatory and inhibitory neurons in the neocortex were found to arise from different developmental lineages. In rodents, the progenitor cells of these two neuronal populations are fully segregated in space (Figure 1). Excitatory neurons are generated in the proliferative zone of the dorsal telencephalon and then migrate radially to constitute the future neocortex. In contrast, inhibitory interneurons are produced in the proliferative zone of the ventral telencephalon and migrate tangentially to reach the neocortex, where they co-assemble with excitatory neurons into functional circuits. This spatial segregation in the excitatory and inhibitory neuron progenitors no doubt further complicates the coordinated production and positioning of these neurons.

The production of neocortical excitatory neurons has been extensively studied and various types of neural progenitor cells have been revealed in the proliferative zone of the dorsal telencephalon (Figure 2). In addition, the radial migration of excitatory neurons has been

characterized in great detail. Relatively speaking, our knowledge of the neurogenesis and migration of neocortical interneurons remains limited, in part due to their origin outside the developing neocortex and to their extraordinary diversity.

I. Diversity of excitatory neuron progenitor cells

In the developing vertebrate nervous system, the neural tube is the embryo's precursor to the central nervous system (CNS) and the neuroepithelial cells that constitute the neural tube are the precursors of all future neurons, including those in the neocortex 1.

Neuroepithelial cells (NECs)

In the early stages of embryonic development, most neuroepithelial progenitor cells undergo symmetric division – two neuroepithelial cells are produced at each division, thereby expanding the population of founder cells that will ultimately produce the CNS including the neocortex. Retrovirally-mediated lineage tracing demonstrated that many neuroepithelial cells are multi-potent progenitor cells². They are capable of generating the first group of neurons in the neocortex and radial glial cells (RGCs), the major neural progenitor cells responsible for producing neurons and glia. In the mouse neocortex, the transition of NECs to RGCs appears to occur around embryonic day 9.5–10.5 (E9.5-E10.5). This transition is characterized by the initiation of the expression of astroglial markers such as astrocyte-specific glutamatetransporter, SLC1A3, formerly known as GLAST and fatty acid binding protein 7 (FABP7)^{3,4}, and an alteration in tight junctions including loss of Occludin ^{5,6}.

Radial glial cells (RGCs)

RGCs are a widespread non-neuronal cell type in the developing CNS of all vertebrates examined so far. They were first identified as a transient population that serves as a scaffold for neuronal migration ^{7–9}; however, their fundamental role as neural progenitor cells has been well-demonstrated in the past decade ^{10–12}. RGCs are defined by their characteristic radial bipolar morphology and their astroglial properties. These long bipolar cells expand across the entire thickness of the developing neocortex with a long basal radial process pointing to the pial surface, a short apical ventricular endfoot reaching the ventricular zone (VZ) surface, and the soma located in the VZ ^{13,14}. During the early stages of neurogenesis, RGCs largely divide symmetrically to produce two radial glial progenitor cells after each division ^{15,16}. At later stages, they predominantly undergo asymmetric cell division to selfrenew and, at the same time, to generate a differentiating daughter cell that is either a neuron or an intermediate progenitor cell ^{12,17}, or an outer subventricular zone (OSVZ) progenitor cell ¹⁸. At the end of neurogenesis, RGCs are shown to switch to gliogenesis producing astrocytes and oligodendrocytes ¹⁹.

Intermediate progenitor cells (IPCs)

Besides RGCs, another type of neuronal progenitor cell – the basal or intermediate progenitor cells (IPCs) – emerges at the onset of peak neurogenesis. They accumulate in the subventricular zone (SVZ), a second proliferative region that lies above the VZ. These IPCs are produced by RGCs in the VZ through asymmetric cell division and migrate radially to the SVZ, where they divide symmetrically to generate two postmitotic neurons 17,20,21. IPCs can also divide symmetrically to produce a pair of progenitors and each could subsequently generate a pair of neurons 17,21. The generation of neocortical neurons through IPCs would therefore significantly increase the number of neurons produced by individual RGCs. Hence, it has been postulated that the abundance of IPCs may contribute to the evolutionary expansion of the reacortex 2^2 . Notably, IPCs are molecularly distinguishable from RGCs by their expression of the transcription factor genes *Eomes* (formerly known as *Tbr2*)²³, Cux1/Cux2 $2^{4,25}$, and Satb2 2^{6} .

Short neural precursors (SNPs)

Studies using three-dimensional reconstruction of individually labeled progenitor cells from the VZ surface showed that the vast majority of VZ progenitor cells in the mouse brain (88% at E12.5; 66% at E16.5) have long radial processes extending towards the basal pial surface with a growth cone at their tips ²⁷. Nevertheless, a subpopulation of VZ dividing cells that either possess a short basal process or lack the basal process altogether ('club-shaped') has also been revealed as morphologically, ultrastructurally and molecularly distinct from RGCs and they are termed as short neural precursors (SNPs) ²⁸. Follow-up studies have suggested that SNPs and RGCs cohabit the VZ but display different cell cycle kinetics. They give rise to distinct neuronal lineages because of a differential reliance on IPC amplification ²⁹. While RGCs often give rise to neuronal progeny indirectly through IPCs within the SVZ, most SNPs produce postmitotic neurons directly within the VZ ²⁹. These studies suggest that progenitor cell variety could be a very important factor in neocortical development.

Outer SVZ progenitor cells

Recently, another distinct type of self-renewing progenitor cell – the outer subventricular zone (OSVZ) progenitors – was discovered in developing human and ferret neocortices ^{30–33}. Although they originate from and behave similar to RGCs, OSVZ progenitor cells retain only the basal process and lack the apical process necessary for anchoring to the VZ surface. They were initially recognized as a special population of neural progenitor cells as an evolutionary consequence of neocortex expansion. Strikingly, a very small population of OSVZ progenitor cells, similar to those observed in developing human and ferret neocortices, has of late been observed in the developing mouse neocortex ^{18,34}. They are likely generated by rare oblique RGC divisions ¹⁸.

II. Mechanisms of asymmetric progenitor cell division

It is generally accepted that asymmetric division of progenitor cells is responsible for producing nearly all of the neurons in the developing neocortex either directly or indirectly. However, the molecular and cellular basis of asymmetric progenitor cell division in the developing neocortex remains largely unknown. A number of molecular signaling pathways have been implicated in the regulation of progenitor cell division and neuronal differentiation in the neocortex. For example, NOTCH-, EGFR- and FGF-signaling have all been implicated in the maintenance of RGCs and are hence crucial for the expansion of the neural progenitor pool ^{35–41}. NOTCH signaling in particular has been extensively studied in recent years, given its central role in controlling asymmetric division of neuroblasts in the *Drosophila* nervous system. In the developing neocortex, NOTCH signaling has also been thought to regulate self-renewal versus differentiation of RGCs ⁴², although the precise role of NOTCH signaling in regulating RGC asymmetric division remains unclear ^{43–48}.

A number of models have been proposed to explain the cellular basis of RGC asymmetric division (Figure 3):

Cleavage-plane orientation model

In *Drosophila*, delaminating neuroblasts from the epithelium retain the apical-basal polarity, which orients the mitotic spindle perpendicular to the epithelial layer. This precise orientation of the mitotic spindle ensures asymmetric segregation of the critical fate determinants such as NUMB and Prospero-related family, resulting in distinct fate specification ^{49,50}. In the developing neocortex, RGCs in the VZ display interkinetic nuclear oscillation as they progress through the cell cycle. While their nuclei are located at the top of the ventricular zone during S phase, mitosis occurs only upon arrival of the nuclei at the VZ

surface. It has been postulated that analogous to the *Drosophila* nervous system, critical fate determinants in these cells are enriched at the VZ surface ^{51,52}. Therefore, when the cleavage plane is parallel to the VZ surface, this horizontal division results in asymmetric inheritance of critical fate determinants enriched at the VZ surface and consequently distinct daughter cell fate specification (i.e. asymmetric division). On the other hand, when the cleavage plane is perpendicular to the VZ surface, this vertical division results in symmetric inheritance of critical fate determinants and consequently the same daughter cell fate specification (i.e. symmetric division) (Figure 3A).

While this cleavage plane orientation hypothesis has wielded a dominant influence on the understanding of progenitor cell asymmetric division and neurogenesis in the developing neocortex, it is being challenged by recent experimental data. Based on this hypothesis, asymmetric, neurogenic division should assume the horizontal cleavage plane, while symmetric, proliferative division should assume the vertical cleavage plane. However, recent studies including time-lapse imaging analysis of RGC divisions clearly demonstrated that the vast majority of asymmetric, neurogenic divisions can be asymmetric and neurogenic.

Oblique division model

To resolve the aforementioned discrepancy that vertical divisions can be either symmetric or asymmetric, a new model – oblique division model – has been put forward ¹⁵. This model suggests that dividing progenitor cells possess a small patch of membrane (1-2%) of the total plasma membrane) close to the VZ surface and that the inheritance of this small patch of membrane is critical for fate specification ⁵⁶. When a vertical cleavage plane bisects this small membrane patch, the division is symmetric. On the other hand, given that this membrane patch is so small, a slight tilt in the vertical cleavage plane (i.e. oblique division) would bypass it and result in asymmetric division (Figure 3B) ⁵⁶. The nature of this small membrane patch in individual dividing progenitor cells remains elusive, even though it is largely considered to be the apical domain of the progenitor cells. The first and only molecular marker reported to mark this domain is PROMININ 1 (CD133), a membrane glycoprotein specifically associated with plasma membrane protrusions such as microvilli 57,58. Although PROMININ 1 is usually localized to the apical surface in epithelial cells and is a well-characterized stem cell marker, it is not a typical apical membrane marker. None of the well-characterized apical protein complexes, such as the partition-defective (PAR) protein complex and the PATJ-PALS1-CRBS complex, have been reported to label this domain in interphase and dividing progenitor cells. In fact, it is technically challenging to explicitly reveal the apical/basal polarity of individual dividing progenitor cells at the VZ surface ⁵⁹, even though the polarity of interphase progenitor cells in the VZ has been well-established ⁶⁰. Recent studies suggested that the oblique division of radial glial cells is critical for the production of OSVZ progenitor cells ¹⁸ or IPCs ⁵⁴ in the mouse neocortex. This difference in the daughter cell fate specification further emphasizes the importance to understand the upstream components that control the orientation of the mitotic spindle and the downstream factors inherited by the daughter cells that regulate the fate specification.

Dynamic polarity model

Recently, it has been shown that the distribution of PARD3, an evolutionarily conserved polarity protein that marks the apical domain, in RGCs is dynamic depending on cell cycle progression ⁵⁹. While it is selectively localized at the TJP1 (formerly known as ZO-1)-expressing lateral membrane domain in the ventricular endfeet during interphase, PARD3 becomes dispersed during mitosis. The dynamic distribution of PARD3 in individual dividing RGCs can result in polarized distribution of PARD3 followed by its asymmetric

inheritance by the two daughter cells, which results in differential activation of NOTCH signaling and subsequent distinct cell fate specification. These data suggest the daughter cell that inherits a greater amount of PARD3 develops high NOTCH signaling activity and remains an RGC, whereas the one that inherits less PARD3 has low NOTCH signaling activity and differentiates into a neuron or IPC (Figure 3C) ⁵⁹. Interestingly, while asymmetric inheritance of PARD3 occurs in dividing progenitor cells with different cleavage plane orientations (mostly vertical/oblique, and to a lesser extent horizontal), PARD3 polarity is always perpendicular to the cleavage plane orientation, raising the possibility that the dynamic establishment of polarity in dividing progenitor cells influences the orientation of the cleavage plane and subsequent daughter cell fate specification.PARD3 has also been implicated in progenitor cell asymmetric division in the zebrafish neural tube ⁶¹. However, in this model PAR3 inheritance appears to correlate with a neuronal fate. This difference calls for a closer examination of the behavior and function of endogenous PARD3 across species.

Basal process inheritance model

Being intricately associated with cleavage-plane orientation and apical membrane inheritance, inheritance of the basal process has also been linked to fate specification, despite much controversial data (Figure 3D). For symmetric divisions that expand the progenitor pool at the onset of neurogenesis, the basal process of NECs can be bisected and inherited by both daughter cells ⁶². On the other hand, it has been postulated that in symmetric divisions at later stages, the basal process is inherited by only one of the daughter cells while the other daughter cell has to elaborate a new basal process ^{17,20}. For asymmetric divisions during later neurogenesis, it has been indicated that the neuronal daughter cell inherits the basal process and migrates towards the cortical plate (CP, the future neocortex) through somal translocation, whereas the other daughter cell could grow a new basal radial process and remain in the VZ as a progenitor ^{63,64}. In contrast, time-lapse imaging studies by Noctor and colleagues revealed that the daughter cell inheriting the basal process remains an RGC ^{12,17}. Furthermore, by inducing oblique division in RGCs to split the apical and basal processes to the two daughters, recent studies by Matsuzaki group indicated that the daughter cell retaining the basal process mostly self-renews outside the VZ, similar to the OSVZ progenitors reported in primates and ferrets, while the apical daughter cell that inherits the apical membrane at the VZ surface differentiates ^{18,55}. Only when a daughter cell inherits both the basal process and the apical membrane does it become an RGC. It remains unclear where these discrepancies stem from and future work that identifies the signals associated with the basal process may help clarify these issues.

Cell cycle length model

An increase in cell cycle length, specifically G1 phase, is naturally associated with the progression of neurogenesis ⁶⁵. A recent model, referred to as "cell cycle length hypothesis", has been proposed, largely on the basis of the observation that lengthening the neuroepithelial cell cycle alone is sufficient to promote the transition from symmetric proliferative to asymmetric neurogenic divisions ⁶⁶. This hypothesis emphasizes that unequal inheritance of a cell fate determinant by daughter cells may or may not lead to distinct fate specification. Instead, the length of cell cycle determines the time window in which the cell fate determinant is allowed to act ^{1,66}, thereby resulting in fate specification. This concept has been further supported by more recent observations showing that progenitors undergoing neurogenic divisions ⁶⁷.

Asymmetric centrosome inheritance model

The asymmetric inheritance of the mother versus daughter centrosome has been suggested to play a critical role in RGC asymmetric division. Centrosome duplication during mitosis generates a pair of centrosomes, one retaining the original mature mother centriole (i.e. the mother centrosome) and the other possessing the original less mature daughter centriole (i.e. the daughter centrosome) ⁶⁸. During asymmetric division of RGCs, the mother centrosome is preferentially inherited by the renewing progenitor cell, while the daughter centrosome is inherited by the differentiating daughter cell ⁶⁸. Interestingly, the asymmetric inheritance of the mother versus daughter centrosome has also been described in asymmetrically dividing *Drosophila* male germline stem cells ⁶⁹ and neuroblasts ^{70,71}. In germline stem cells, the mother centrosome is preferentially inherited by renewing stem cells. On the other hand, in neuroblasts the mother centrosome is preferentially inherited by renewing stem cells. On the other hand, in metrical (PCM), while the daughter centrosome gains PCM. This intriguing dynamics of PCM points to the importance of the property, rather than the absolute birth date, of the centrosome in regulating asymmetric division of progenitor/stem cells.

III. Migration of excitatory neurons

After exiting the cell cycle, new-born excitatory neurons need to migrate out of the proliferative zone and into the CP, where they further differentiate and form distinct layers. This stereotypic laminar organization of the neocortex is crucial for its function. Birth-dating studies have shown that layers II–VI of the neocortex are generated in an 'inside-out' fashion, such that neurons generated early reside in the deep layers, whereas later-born neurons migrate past the existing neurons to occupy more superficial layers (Figure 4) ^{72,73}.

New-born excitatory neurons that migrate radially into the CP are thought to do so through two distinct modes: somal translocation or locomotion. Early in neocortical development, the principal mode of neuronal migration appears to be somal translocation ^{64,74,75}, in which neurons typically have a long radial process and maneuver their soma towards the leading edge of the radial process that is attached to the pial surface ⁷⁴. How neurons acquire these long radial processes remains controversial. One possibility is that a newly generated neuron quickly elaborates a new radial ascending process before migration ⁷⁴. Alternatively, the daughter neuron inherits the original radial process through asymmetric, neurogenic division ⁶⁴. At later stages of neocortical development, new-born neurons predominantly adopt a second migratory mode where they make use of the radial processes of RGCs as a scaffold to reach their final positions, and this is known as radial glia-guided migration or locomotion ^{17,76,77}. Time-lapse imaging studies in brain slice cultures have revealed many details of the locomotion of new-born neurons. These neurons do not migrate directly to the CP but instead exhibit four distinct phases of migration as characterized by Noctor, Kriegstein and colleagues ¹⁷. During phase one, new-born neurons assume a bipolar shape and ascend rapidly from the VZ to the SVZ. The second phase consists of migratory arrest for 24 hours or more with a multipolar morphology, followed by a third phase of retrograde migration toward the ventricle with a switch back to bipolar morphology. This retrograde migration is intriguing and raises the possibility that migrating neurons may receive certain signals from the ventricle, although future experiments are needed to confirm that it does occur in native conditions. After coming into contact with the ventricle, the daughter cells enter the fourth phase of migration, when they reverse polarity and migrate toward the CP¹⁷. Notably, once their leading process reaches the marginal zone (MZ), locomoting cells can detach from the radial glial fibers and switch to "terminal translocation" mode ⁷⁴. It has been further suggested that this change in the migratory mode occurs just beneath the primitive cortical zone (PCZ) and is critical for the disabled 1 (Dab1)-dependent inside-out lamination of the neocortex ⁷⁸. Excitatory neuron migration in the neocortex may not be

strictly radial. Lineage studies using retroviral labeling showed that clones of neurons derived from individual progenitors could disperse tangentially ^{79–81}. However, the mechanisms responsible for this tangential dispersion remain unclear. Tangential dispersion may occur between symmetrically proliferating progenitor cells in the VZ/SVZ ^{17,81,82}, and/ or postmitotic neurons in the SVZ/IZ ^{83,84}. Notably, multipolar cells in the SVZ/IZ, often observed at the second phase of radial migration, do not seem to closely associate with the radial fiber and have been shown to occasionally 'jump' tangentially from one radial glial fiber to another ^{17,85}. Serum response factor (SRF), a transcription factor regulated by changes in the ratio of polymerized to unpolymerized actin, has been recently shown to direct excitatory neurons to choose a radial migration pathway along radial glia over a tangential migration mode ⁸⁶. Despite the discovery of other migratory modes, radial glia-guided locomotion remains a central mechanism for establishing the architecture of the neocortex.

Many molecules have been implicated in the regulation of excitatory neuron radial migration in the developing neocortex. Of these, REELIN, an extracellular matrix protein secreted by Cajal-Retzius cells in the marginal zone, appears to be a key player. When mutated spontaneously in *Reeler* mice, the birth date-dependent inside-out fashion of the positioning of excitatory neurons is reversed, such that the layers are generated in an 'outside-in' manner ^{87–93}. Subsequently, other mutants in REELIN signaling such as Scrambler which lacks a functional Dab1 gene have been shown to recapitulate the Reeler phenotype ^{94–96}. Interestingly, in all mutants with abnormal neocortical lamination, the development of the preplate is normal. This observation is consistent with the idea that early-born neurons adopt a different mode of migration – soma translocation – which is unaffected by the cascade of signaling mechanisms that control the radial glia-guided locomotion of later-born neurons occupying the CP. Genetic analysis of mouse mutants and human neurologic disorders associated with disrupted cortical laminar organization has identified other molecules including cyclin-dependent kinase 5 (Cdk5) 97, Cdk5r1 98, transcription factor Pax6^{99,100}, neurotrophin 5 (Ntf5)¹⁰¹, integrins ^{102–104}, and doublecortin¹⁰⁵ as critical regulators of neuronal migration, and thereby layer formation in the developing neocortex.

IV. Interneuron neurogenesis and migration

In contrast to excitatory neurons generated directly in the proliferative zone of the neocortex, most, if not all, neocortical interneurons arise from the ventral telencephalon, predominantly the medial ganglionic eminence (MGE) 106,107 , caudal ganglionic eminences (CGE) 108,109 , and the preoptic area (POA) 110 (Figure 1). Even though the germinal zones of pyramidal neurons and interneurons are spatially segregated, they all harbor RGCs in their VZ 10,111112 . However, the RGCs responsible for producing excitatory neurons and inhibitory interneurons are distinct. RGCs in the VZ of the neocortex express *Pax6* and *Emx1*, two transcription factor genes essential for specifying neocortex features, without which the neocortex is converted into structures normally formed by the ventral telencephalon 113 . Conversely, RGCs in the ventral telencephalon express different sets of transcription factor genes such as *Gsx1/Gsx2* and *Olig2*⁴. Notably, expression analysis has shown that neocortical RGCs are highly heterogeneous and can be divided into multiple antigenically distinct subpopulations that evolve in a stereotypical manner as development proceeds ³. It is possible that FABP7-expressing RGCs in the ventral telencephalon are also heterogeneous, which might be related to the great diversity of neocortical interneurons.

A recent study using retroviral labeling and live imaging analysis clearly demonstrated that RGCs in the VZ of the MGE undergo asymmetric cell division to produce neocortical interneurons ¹¹⁴. In addition, there are abundant IPCs in the SVZ that divide symmetrically

to produce neocortical interneurons (Figure 5). In fact, the SVZ in the ventral telencephalon is substantially larger than that in the neocortex, indicating that symmetrically dividing IPCs represent an important source of neocortical interneurons ^{4,115}.

Neocortical interneurons are remarkably diverse and can be subdivided into distinct subgroups based on neurochemical marker expression, morphology, firing pattern and synaptic connectivity. An open question related to neocortical interneuron neurogenesis is how this bewilderingly diverse group of interneurons is generated. Genetic and transplantation studies have demonstrated that distinct regions of the GEs generate distinct interneuron subtypes ^{109,116–118}. In addition, distinct temporal origins of physiologically defined interneuron subtypes have also been reported ^{109,116–118}. However, the basis of the production of distinct groups of neocortical interneurons at individual progenitor cell level is largely unknown. It appears that individual RGCs in the VZ of the MGE are capable of producing different subtypes of neocortical interneurons ¹¹⁴. It will be interesting in the future to determine the lineage trees of different subtypes of neocortical interneurons.

In contrast to excitatory neurons, neocortical interneurons produced at the ventral telencephalon need to embark on a lengthy tangential migration to reach the dorsal neocortex, and then migrate radially to arrive at their final proper laminar positions ^{106,108,119–128}. On the other hand, new-born interneurons in the MGE and POA, similar to excitatory neurons in the neocortex, initially migrate radially along the mother radial glial progenitor cell before starting their tangential migration ¹¹⁴. The physiological significance of this initial radial migration is unclear. It may allow proper neuronal differentiation of the new-born cells and subsequent proper tangential migration. Previous studies have identified a diverse set of temporally and spatially distinct tangential migratory routes associated with different progenitor zones within the ventral telencephalon 123,129. Early during development (E11.5), neocortical interneurons arise primarily from the MGE and the anterior entopeduncular area (AEP)/POA, and follow a superficial route outside the developing striatum ¹²³. At mid-embryonic stages (E12.5-E14.5), the MGE appears to be the principal source of neocortical interneurons, which migrate either deeply or superficially, relative to the developing striatum. During this period, the LGE and CGE also start to produce interneurons that migrate to the neocortex ¹²⁸. At later stages (E15.5), the CGE reaches its peak of neurogenesis and becomes the more prominent/substantive source of cortical interneurons, which prefer a deeper migratory route ^{128,129}.

Once an interneuron fate has been specified and a tangential migratory pathway is undertaken, there are at least three different types of factors that can regulate this process: 1) motogenic factors that stimulate the movement of interneurons, such as hepatocyte growth factor (also known as scatter factor) ¹³⁰, brain-derived neurotrophic factor (BDNF) and neurotrophin 5 (NTF5) ^{101,131}; 2) extracellular substrates that serve as scaffolds for migration, e.g. migrating interneurons in the lower intermediate zone may use corticofugal axons as the substrate for their migration ^{83,132,133}; 3) guidance factors that direct different migratory streams through appropriate routes towards their targets, such as chemorepulsive SLIT/ROBO ^{134–136} and semaphorin/neuropilin signaling ¹³⁷, and chemoattractive CXCL12/CXCR4 signaling ^{138–141}. Notably, this long-distance tangential migration of individual interneurons is believed to be mostly random ¹⁴², although it is unclear how their random walking behavior could lead to an organized distribution of interneurons in the neocortex for functional circuit assembly, such as birth date-dependent laminar distribution of MGE-derived interneurons. Interestingly, a recent study demonstrated that clonally related interneurons do not randomly disperse but form spatially organized clusters in the neocortex ¹¹⁴, indicating a lineage-dependent organization of interneurons in the neocortex (Figure 5).

After entering the neocortex, interneurons continue to disperse tangentially via highly stereotyped routes in the MZ, the subplate, and the lower intermediate zone/the SVZ ¹⁰⁶. Eventually, interneurons switch from tangential to radial migration to adopt their final laminar position in the neocortex ^{124,142,143}. Proper cell-autonomous development of neocortical interneurons has been indicated as a critical factor in their radial positioning, as defective laminar distribution is often observed in mutant mice with abnormal interneuronal differentiation ^{144–147}. Moreover, neocortical interneurons are intrinsically programmed to modulate expression of the K⁺/Cl⁻ co-transporter SLC12A5 (formerly known as KCC2) in order to sense the ambient extracellular levels of GABA and glutamate as a way to determine when to stop migration in the neocortex ^{148,149}.

Furthermore, many interneurons and excitatory neurons that are born at a similar time end up occupying the same neocortical layer 108,150-152, indicating that the mechanisms controlling the layer acquisition of these two neuronal types are highly coordinated. However, little is known about how this coordination is achieved. Recently, it has been shown that RGCs may serve as a scaffold for both further tangential and radial migration of interneurons within the developing neocortex ^{124,153}. When the radial glia scaffold is disrupted, neither excitatory neurons nor inhibitory interneurons migrate properly into the cortical plate ^{154,155}. Also, gap junctions, which play an important role in the radial migration of excitatory neurons ^{156–158}, have been recently suggested to regulate the switch from tangential to radial migration as interneurons enter the neocortex ¹⁵⁹. In addition, interneurons invade the neocortex only after their excitatory neuron partners, possibly reflecting a need for signals from appropriately located excitatory neurons ¹⁴⁰. Consistent with this idea, interneurons are found to distribute abnormally in the cortex of the Reeler mice, in which pyramidal neuron lamination is nearly inverted ^{160,161}. Further evidence of a significant role of excitatory neurons in influencing interneuron migration and positioning came from a recent study using $Fezf2^{-/-}$ mice, which demonstrated that distinct subtypes of excitatory neurons could uniquely and differentially determine the laminar distribution of neocortical interneurons ¹⁶². Thus, it is possible that the pairing of synchronously-born excitatory neurons and interneurons relies on the expression of complementary molecules, which may have evolved to ensure the coordinated positioning of these two distinct neuronal populations.

Concluding remarks

During the past decade, there has been remarkable progress in our understanding of the production and migration of neocortical excitatory and inhibitory neurons at the mechanistic and cell biological level. However, many key questions remain inadequately addressed. For example, how is symmetric versus asymmetric division of progenitor cells controlled? How is the rich diversity of neuronal types generated? How is the migration of different types of neurons regulated to generate the stereotypic laminar architecture? The greatest challenge presumably lies in the feasibility of systematically addressing these questions, as the regulation of these cellular processes likely involves a comprehensive set of molecular pathways that function in a coherent manner. With the development of new experimental approaches, it is safe to predict that our understanding of neurogenesis and neuronal migration in the developing neocortex will be enriched in the near future.

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Figure 1. Distinct origins of excitatory neurons and inhibitory interneurons in the developing mouse neocortex

Excitatory neurons are generated in the proliferative zone of the dorsal telencephalon and then migrate radially to the cortical plate (red arrows). In contrast, inhibitory interneurons are produced in the proliferative zone of the ventral telencephalon, especially the MGE and AEP/POA, and migrate tangentially to reach the neocortex following two major routes (blue dashed arrows). The colored regions indicate the proliferative zones across the embyronic brain expressing different transcription factors that are essential for proper neurogenesis of distinct neuronal populations. AEP: anterior entopeduncular area; H: hippocampus; LGE: lateral ganglionic eminence; LP: lateral pallium; MGE: medial ganglionic eminence; Ncx: neocortex; POA: preoptic area.



Figure 2. Diverse populations of excitatory neuron progenitor cells in the mouse neocortex During early brain development, neuroepithelial cells (NECs) are the major neural progenitors in the neocortex. NECs divide symmetrically to generate additional NECs, some of which give rise to the first group of neurons (N) through asymmetric division. As the developing brain epithelium thickens, NECs elongate and transit to radial glial cells (RGCs). RGCs can also divide symmetrically to expand the progenitor pool or asymmetrically to generate neurons either directly or indirectly through intermediate progenitor cells (IPCs), which generate neurons directly through symmetric division in the subventricular zone (SVZ). It remains unclear how short neural progenitors (SNPs) are generated. It is possible that they are generated from RGCs or they may be a distinct population originated directly from NECs. Most SNPs produce postmitotic neurons directly in the ventricular zone (VZ). Outer subventricular zone (OSVZ) progenitors likely originate from RGCs through oblique division, which leads to loss of the apical process and ascension of the nucleus towards the cortical plate/intermediate zone (CP/IZ). The minor population of OSVZ progenitors can also generate neurons through asymmetric division outside of the VZ.



Figure 3. Models of symmetric and asymmetric divisions of radial glial progenitor cells in the mouse neocortex

(A) Cleavage-plane orientation model. When the cleavage-plane is perpendicular to the VZ surface, the division of RGCs results in symmetric inheritance of critical fate determinants enriched at the VZ surface (represented by a green line) and consequently the same daughter cell fate specification (i.e. symmetric division). When the cleavage-plane is parallel to the VZ surface, the division results in asymmetric inheritance of critical fate determinants enriched at the VZ surface and consequently distinct daughter cell fate specification - one remains as RGC whereas the other becomes a postmitotic neuron (N) or an intermediate progenitor (I) (i.e. asymmetric division). (B) Oblique division model. Dividing progenitor cells possess a small patch of membrane close to the VZ surface (represented by a short purple line), the inheritance of which is critical for fate specification. Bisection of this small membrane patch leads to symmetric division, whereas a slight tilt in the vertical cleavage plane (i.e. oblique division) bypasses it and results in asymmetric division. (C) Dynamic polarity establishment model. The distribution of the evolutionarily conserved cell polarity protein PARD3 is dynamic depending on cell cycle progression, which becomes dispersed during mitosis. Even distribution of PARD3 in dividing RGCs leads to symmetric division, whereas polarized distribution of PARD3 results in its asymmetric inheritance by the two daughter cells, followed by differential activation of NOTCH signaling and subsequent distinct fate specification. (D) Basal process inheritance model. Inheritance of the basal process has been linked to fate specification though it is still controversial. The basal process can be bisected and inherited by both daughter cells during the symmetric division, although it has also been postulated that the basal process may be inherited by only one daughter RGC while the other has to elaborate a new basal process. For the asymmetric division, earlier studies suggested that the daughter cell that inherits the basal process becomes a differentiating neuron whereas the other daughter grows a new process and remains in the VZ as a progenitor. Recent work indicated that the daughter cell inheriting the basal process remains a progenitor (either a RGC or OSVZ progenitor) while the other differentiates (either a neuron or an intermediate progenitor).



Figure 4. Inside-out layer formation of excitatory neurons in the neocortex

The PP is formed by the production of the first wave of post-mitotic neurons that migrate from the VZ to the pial surface. Then, a second wave of new-born neurons (blue triangles) migrate through the SVZ/IZ and split the PP into the more superficial MZ (neurons in the MZ are represented by orange ovals) and the more deeply located SP (neurons in the SP are represented by green diamonds), creating the CP – the future neocortex. Neurons generated subsequently (represented by triangles with different colors, pink, purple and yellow by birth order) expand the CP in an inside-out fashion, as later-born neurons pass the existing neurons to occupy more superficial layers. The VZ progressively shrinks as the neural progenitor cells decrease during later embryonic development. During the postnatal development towards adulthood, the SP degenerates and leaves behind a six-layered neocortex (panel on the right). CP: cortical plate; IZ: intermediate zone; MZ: marginal zone; PP: preplate; SP: subplate; SVZ: subventricular zone; VZ: ventricular zone; WM: white matter.



